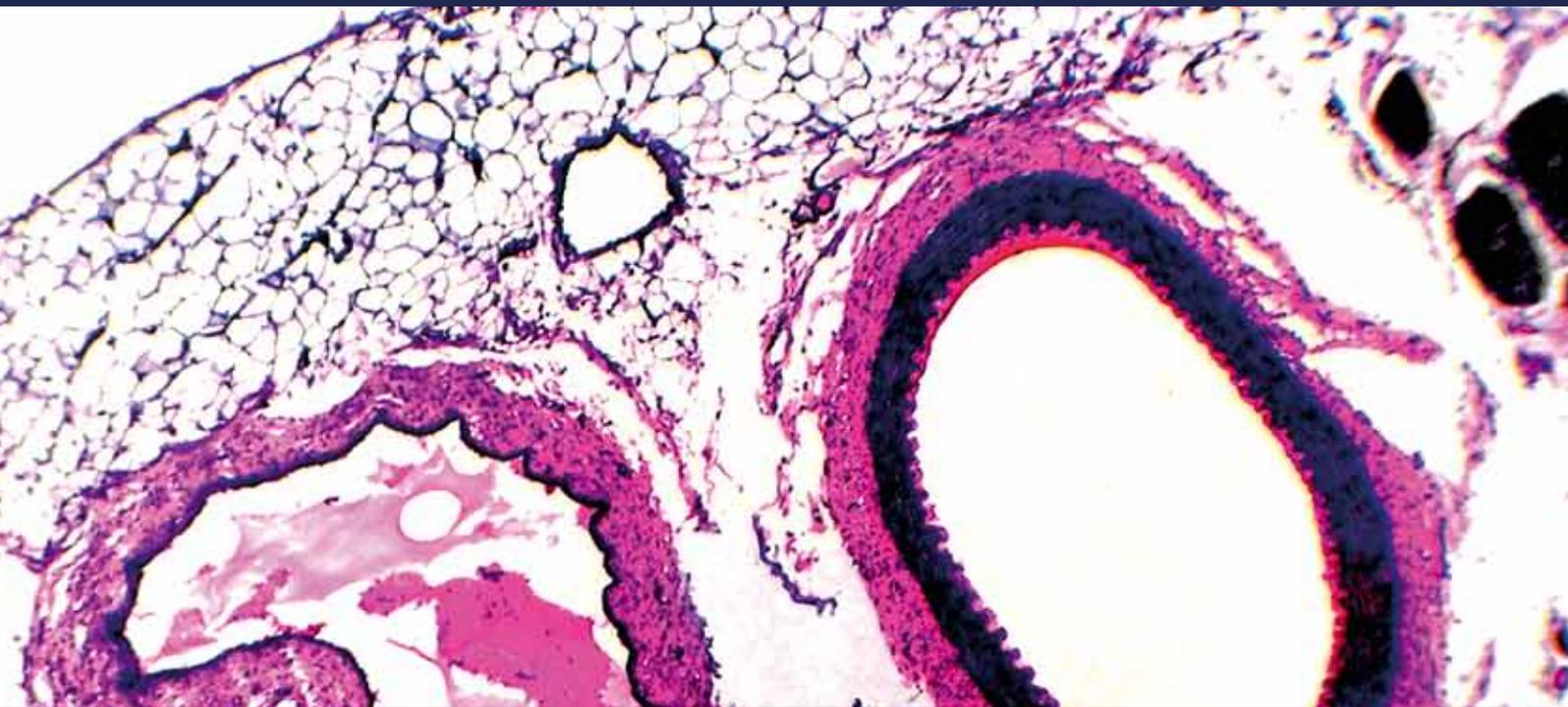


Brain RAS: Hypertension and Beyond

Guest Editors: Marc de Gasparo, Robert C. Speth, Ovidiu C. Baltatu, and Patrick Vanderheyden





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International Journal of Hypertension

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Editorial

Brain RAS: Hypertension and Beyond

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Received 25 February 2013; Accepted 25 February 2013

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Cardiac output and vascular resistance are the cornerstones of blood pressure regulation, which is achieved through neural, humoral, and local tissue factors.

The sympathetic nervous system (SNS) and the renin-angiotensin system (RAS) play a major role in the control of blood pressure. Angiotensins act as endocrine, paracrine, and autocrine regulators. The peripheral physiological and pathophysiological actions of the RAS are well established. The existence and functional relevance of the RAS in the brain is increasingly recognized as a major regulator of the cardiovascular (CV) system and a significant drug target for antihypertensive and other CV therapeutics. All the constituents of the RAS occur in the brain and participate in the regulation of blood pressure through sympathetic activation and vasopressin release. In addition, an interconnection between neurotransmitters and the brain RAS affects behavior and neurological diseases, for example, Parkinson's, and Alzheimer's diseases. Moreover, the clinical efficacy of renin and ACE inhibitors and angiotensin receptor blockers (ARBs) and the presence of their targets in the brain illustrate the synergistic interaction between brain and peripheral RAS.

This special issue illustrates some aspects of the brain RAS pathway and function including its effect on the circadian rhythm of blood pressure.

The RAS has been described in the brain. Using subtype specific antibodies, C. Premer et al. observed selective expression of AT1a, AT1b, and AT2 receptor subtypes in neurons and glia in a large number of brain regions including the subfornical organ, median eminence, area postrema,

paraventricular, and solitary tract nucleus of the rat brain as well as in the pituitary and adrenal.

Ang II formation in the pineal gland and glial cells appears to depend on alternative pathways including chymase (L. A. Campos et al.). One possibility might be that the prorenin receptor (PRR) binds prorenin or renin from circulation to form Ang I and chymase to form Ang II. The brain PRR appears to initiate the brain angiotensin peptide formation (W. Li et al.). Indeed, PRR is expressed ubiquitously in the brain with the highest expression levels in the pituitary and frontal lobe. Recent findings indicate that PRR has RAS independent roles associated with the vacuolar proton-ATPase and the Wnt signaling pathways (W. Li et al.). PRR in the brain could play a pivotal role in neural regulation of blood pressure and body fluid homeostasis.

In addition, AT4/IRAP and Mas receptors are also present in the brain. Aminopeptidases (and other angiotensins degrading enzymes, e.g., ACE2 and endopeptidase), which form fragments such as Ang III, Ang IV, Ang 2–10, Ang 1–9, and Ang 1–7, are also the topic of several reports (A. B. Segarra et al.; M. A. Clark et al.).

Formation of Ang III in the brain may promote hypertension while Ang IV, which inhibits vasopressinase activity and may have a therapeutic value for cognitive function in the brain.

There is still a debate regarding the relative importance of Ang II and Ang III in the brain. Using astrocytes in culture and an inhibitor of aminopeptidase A to prevent conversion of Ang II to Ang III, M. A. Clark et al. demonstrate that both

Ang II and Ang III induce phosphorylation of MAPK and JNK and stimulate astrocyte growth equipotently.

Ang IV binds to the AT₄ receptor. While the AT₄ receptor has been convincingly shown to be the insulin-regulated aminopeptidase IRAP, (also known as vasopressinase and cysteine aminopeptidase), others have suggested that the physiological action of Ang IV may also be mediated through the tyrosine kinase c-Met receptor. Regardless of this controversy, binding of Ang IV causes inhibition of the catalytic activity of the peptidase activity of the IRAP receptor and therefore increases AVP and oxytocin, glucose uptake, and cognitive processes. Intracerebroventricular injection of Ang IV improves memory and learning in the rat. The potential of IRAP inhibitors able to cross the blood brain barrier is discussed by H. Andersson and M. Hallberg.

Clearly, the brain RAS regulates sympathetic activity and norepinephrine (NE) release (K. Tsuda), and hyperactivity of the SNS is clearly involved in the cardiovascular pathology. Ang II through the AT₁ receptor and MAPK stimulation affects noradrenergic nerve terminals in the paraventricular nucleus of the hypothalamus (PVN), inhibiting K⁺ channel and stimulating Ca⁺⁺ channels causing NE release. Also, brain aldosterone-mineralocorticoid receptor- (MR-) ouabain pathway might have a pivotal role in Ang II-induced neuronal activation and pressor responses (K. Tsuda). In contrast, Ang 1-7, a metabolite of both Ang I and Ang II, reduces NE release through BK and NO stimulation (M. Nautiyal et al.).

Regulation of the baroreflex is central to CV regulation, and cardiac autonomic imbalance (decreased cardiovagal and increased sympathetic tone) causes baroreflex dysfunction. Ang II acting through the AT₁ receptor and Ang 1-7 acting through the mas-receptor counterbalance each other in the brain through AMPK and mitochondrial NADPH oxidase stimulation and superoxide production (M. Nautiyal et al.; B. P. Campagnaro et al.). Ang II also stimulates dopamine β -hydroxylase in the striatum regulating the baroreflex (Tsuda). It also stimulates central cholinergic neuronal activity. The frontal cortex and CV system are reciprocally and asymmetrically organized: the frontal cortical activity appears to control CV function. Indeed, whereas there is an increased parasympathetic nervous system (PNS) activity in the left prefrontal cortex, one observes a decreased SNS activity in the right one (A. B. Segarra et al.). This may suggest that stimulation of the frontal cortical activity could lower blood pressure.

Several contributions focus on the AT₂ receptor, the functions of which still remain poorly understood. Interestingly, the hypothalamus expresses both AT₁ and AT₂ receptors that could counterbalance each other's effects (C. Premer et al.). Further, the AT₂ receptor is involved in neuronal disorders and appears to play a major role in the regulation of cognitive function through a balance between phosphorylation and kinase activity versus phosphatase activity (M. O. Guimond and N. Gallo-Payet). AT₂ receptors also stimulate neurite outgrowth, migration and excitability and reduce oxidative stress. Moreover, binding of Ang II to the AT₂ receptor stimulates bradykinin and nitric oxide production and improves cerebral blood flow preventing ischemic damage

and dementia (M. Mogi et al.). In addition, the AT₂ receptor appears to play a role in metabolic syndrome as it regulates appetite and increases glucose uptake. Finally, it is proposed that the AT₂ receptor behaves like a gate keeper of cellular and tissue homeostasis (M. O. Guimond and N. Gallo-Payet).

Some signaling pathways involved in the AT₂ receptor function are described. A family of AT₂ receptor interacting proteins (ATIP) through the C-terminal domain of the receptor affect neurite outgrowth and neuronal differentiation (S. Rodrigues-Ferreira et al.). Such interaction of ATIP with the nicotinic acetylcholine receptor may also affect cognitive function (M. Mogi et al.). In fact, the evidence in favor of a neuroprotective action of AT₂ receptor stimulation is compelling (M. Mogi et al.; M. O. Guimond and N. Gallo-Payet).

Interestingly, the RAS in the brain and in the periphery is differently regulated by sodium (H. Takahashi). Whereas a sodium load suppresses the peripheral RAS and reduces sodium reabsorption from the renal tubules, it activates the brain RAS to retain more sodium. The mechanism involves the brain endothelial Na⁺ channel, activation of brain aldosterone, and endogenous digoxin-like substances, causing stimulation of the sympathetic system and increased blood pressure (H. Takahashi).

Circadian rhythms in neural and endocrine systems regulated by the molecular clock in the suprachiasmatic nucleus affect cardiovascular and metabolic disorders. Blood pressure variations during the day and night derive from the change in the dominant sympathetic tone. L. A. Campos et al. demonstrate here that Ang II acts upon AT₁ receptors in the pineal gland to stimulate tryptophan hydroxylase activity and melatonin formation. This decreases sympathetic activity and increases parasympathetic activity. This reduces oxygen-free radicals and increases nitric oxide availability causing a decrease in blood pressure. Such behavior is typical of the physiological night tone reduction of blood pressure (dipper hypertensive patient), whereas the nondipper hypertensive shows an impaired melatonin formation during the night with an inverse circadian blood pressure profile. Pinealectomy reverses the condition, increasing sympathetic activity and the adrenocorticotropin axis. The brain RAS appears therefore to be a major regulator of circadian variation of blood pressure through its effect on light cycle shifts. Medical treatment coordinated with biological rhythms (chronotherapy) combined with inhibition of the brain RAS may be a means of individualizing treatment of hypertension related to circadian rhythms. Interestingly, increased melatonin release also improves insulin sensitivity. Supporting this finding, decreased melatonin is observed in type 2 diabetes and a melatonin gene receptor mutation was associated with increased risk of type 2 diabetes. Melatonin, a nutritional supplement, may therefore be a good therapeutic complement for diabetic patients. Although pineal Ang II stimulates melatonin (L. A. Campos et al.), a beneficial effect of this approach on type II diabetes remains to be demonstrated.

S. Lattanzi et al. review the effectiveness of ARBs given in acute stroke. In a meta-analysis of three trials, they conclude that there is no place for blood pressure lowering treatment

in the acute phase of stroke. However, there may be a U-shaped relationship between BP and outcome: too low blood pressure may result in brain hypoperfusion and thus in a worse outcome. Two ongoing large studies in acute ischemic and haemorrhagic stroke (ENOS and INTERACT) will help to select the best approach of blood pressure management.

In summary, an abundant body of evidence indicates an important regulatory role for the brain RAS in cardiovascular homeostasis and disease beyond blood pressure regulation. Moreover, accumulating evidence reveals brain RAS roles in brain-specific functions and diseases such as cognitive dysfunction, dementia, Alzheimer's and Parkinson's diseases. Further research is needed to better understand detailed mechanisms of brain RAS in these and other diseases to possibly develop new diagnostic strategies.

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Research Article

Immunohistochemical Localization of AT_{1a}, AT_{1b}, and AT₂ Angiotensin II Receptor Subtypes in the Rat Adrenal, Pituitary, and Brain with a Perspective Commentary

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Received 31 October 2012; Revised 1 February 2013; Accepted 5 February 2013

Academic Editor: Patrick Vanderheyden

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Angiotensin II increases blood pressure and stimulates thirst and sodium appetite in the brain. It also stimulates secretion of aldosterone from the adrenal zona glomerulosa and epinephrine from the adrenal medulla. The rat has 3 subtypes of angiotensin II receptors: AT_{1a}, AT_{1b}, and AT₂. mRNAs for all three subtypes occur in the adrenal and brain. To immunohistochemically differentiate these receptor subtypes, rabbits were immunized with C-terminal fragments of these subtypes to generate receptor subtype-specific antibodies. Immunofluorescence revealed AT_{1a} and AT₂ receptors in adrenal zona glomerulosa and medulla. AT_{1b} immunofluorescence was present in the zona glomerulosa, but not the medulla. Ultrastructural immunogold labeling for the AT_{1a} receptor in glomerulosa and medullary cells localized it to plasma membrane, endocytic vesicles, multivesicular bodies, and the nucleus. AT_{1b} and AT₂, but not AT_{1a}, immunofluorescence was observed in the anterior pituitary. Stellate cells were AT_{1b} positive while ovoid cells were AT₂ positive. In the brain, neurons were AT_{1a}, AT_{1b}, and AT₂ positive, but glia was only AT_{1b} positive. Highest levels of AT_{1a}, AT_{1b}, and AT₂ receptor immunofluorescence were in the subformical organ, median eminence, area postrema, paraventricular nucleus, and solitary tract nucleus. These studies complement those employing different techniques to characterize Ang II receptors.

1. Introduction

The ability of angiotensins II (Ang II) and III (Ang III) to stimulate aldosterone [1, 2] and epinephrine [3] release from the adrenal gland is well established. The central nervous system and adenohipophyseal effects of these peptides are also well documented and numerous. While the effects of Ang II on the adrenal are thought to arise primarily from blood-borne Ang II, it is clear that there is a local

brain angiotensinergic system as illustrated by biochemical, immunohistochemical, behavioral, physiological, and receptor binding studies [4–8] and reviews [9–11]. The anterior pituitary also appears to be subject to both blood-borne and local angiotensinergic systems, as well as receiving indirect regulatory signals from brain angiotensinergic activity [12, 13].

In mammals, there are two primary Ang II receptor subtypes, AT₁ and AT₂ [14–19]. With the discovery of these

multiple subtypes of Ang II receptors, pharmacological studies revealed that the AT_1 subtype mediated both aldosterone [20] and epinephrine [21] release as well as pressor [22, 23], dipsogenic [22–24], and sodium appetite [24–26] responses to Ang II. The localization of AT_1 receptors in the rat brain regions mediating pressor and dipsogenic actions of Ang II, such as the subfornical organ (SFO), median preoptic nucleus (MnPO), organum vasculosum of the lamina terminalis (OVLT) paraventricular nucleus of the hypothalamus (PVN), nucleus of the solitary tract (NTS), and area postrema [27–29] is consistent with this role. In contrast, AT_2 receptors tend to be distributed in sensory, motor, and emotional regions of the brain, for example, superior colliculus, medial geniculate nucleus, locus coeruleus, lateral septum, medial amygdala, subthalamic nucleus, and inferior olivary nucleus [27–29]. It has been suggested that the medial amygdala can mediate salt appetite [30], but beyond that, the functional significance of the AT_2 in the brain and the adrenal has not been established.

The subsequent discovery that rodents express two subtypes or isoforms of the AT_1 receptor, AT_{1a} and AT_{1b} , [31–33] raises the question as to which of these two subtypes may be mediating adrenal hormone release and the physiological effects of Ang II in the brain and pituitary. Pharmacological studies of the ability of angiotensins and AT_1 receptor-selective antagonists to bind to the AT_{1a} and AT_{1b} receptor subtypes reveal little difference in their affinities for these two subtypes [34–37].

PCR amplification of AT_{1a} and AT_{1b} mRNA in female rat adrenal, lung, vascular smooth muscle, pituitary, and brain indicated that the AT_{1a} subtype mRNA was predominant in the lung, vascular smooth muscle, and hypothalamus, while the AT_{1b} subtype was predominant in the adrenal, pituitary, subfornical organ, and organum vasculosum of the lamina terminalis [31, 38]. Both PCR amplification [31, 35, 38–40] and in situ hybridization [39, 41, 42] have been used to compare the expression of mRNA for these two subtypes in the adrenal and brain. However, the expression of mRNA does not always correspond with the expression of the protein it encodes. For example, estrogen treatment can reduce AT_1 receptor expression without altering AT_1 mRNA expression presumably via posttranscriptional inhibition of mRNA translation [43]. Moreover, in neuronal tissues, the receptors may be expressed on axonal terminals distant from their perikaryal mRNA.

Studies of AT_{1a} and AT_{1b} mRNA expression in the adrenal indicate that the AT_{1b} subtype mRNA is predominant in the rat adrenal [35, 38, 39, 44], but that it is absent in the adrenal medulla [44–46]. Studies of AT_{1a} and AT_{1b} mRNA in rodent brain vary considerably along a continuum from a predominance of AT_{1b} expression in the female rat brain [31], to a moderate predominance of AT_{1a} in the male mouse brain [40, 42], a differential distribution of the mRNAs in a two-week-old male rat brain [45], to very low expression of AT_{1b} mRNA in the adult male rat brain [41], and to no expression of AT_{1b} mRNA in rat brain [47]. In comprehensive studies of the distribution of AT_{1a} and AT_{1b} mRNA the rat brain and pituitary [41], the AT_{1a} mRNA was found to be highly expressed in brain regions reported to mediate cardiovascular effects of Ang II, while AT_{1b} expression was very low in

these regions. Conversely, AT_{1b} mRNA was very high in the anterior pituitary while AT_{1a} mRNA was low.

To determine if the distribution of AT_{1a} , AT_{1b} , and AT_2 receptor subtype protein in the rat adrenal, pituitary, and brain corresponds to the distribution of the mRNAs for these subtypes, this study uses fluorescence immunohistochemistry with antibodies directed at unique peptide fragments of each of these three subtypes to localize these receptors.

2. Materials and Methods

2.1. Antibody Preparation. Antipeptide antibodies were generated against fragments of rat AT_{1a} , AT_{1b} , and AT_2 receptors. Peptides candidates were selected by computer analysis of full length receptors retrieved from the NCBI protein database (<http://www.ncbi.nlm.nih.gov/protein>) and by Hopp-Woods analysis [48] for optimal antigenicity. Peptides corresponding to receptor fragments near the carboxy terminal tail of the receptor subtypes where there is a 2 amino acid difference were synthesized by solid phase peptide synthesis. For the AT_{1a} receptor, the peptide was PSDNMSSSACKPASC, which corresponds to amino acids 341–355 of this 359 amino acid protein. For the AT_{1b} receptor, the peptide was SSSAKKSASFFEVE, which corresponds to amino acids 346–359 of this 359 amino acid protein. For the AT_2 receptor, the peptide was CRKSSSLREMETFVS, which corresponds to amino acids 349–363 of this 363 amino acid protein (except that it contained a glutamic acid in position 358 versus an aspartic acid). The peptides were compared with the protein database (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>) to establish the uniqueness of the peptide sequences from other known proteins.

Peptides were conjugated to keyhole limpet hemocyanin (KLH) and injected into rabbits at approximately monthly intervals for 6 months. Serum was obtained from the rabbits and affinity purified. To obtain AT_{1a} -selective antibodies, serum from rabbits immunized with the peptide corresponding to the AT_{1a} receptor subtype was affinity purified using chromatography resin cross-linked with the AT_{1a} peptide. Antibodies retained by this resin were eluted with a high salt solution and the eluate was then applied to an affinity column made by cross-linking the AT_{1b} receptor peptide antigen to chromatography resin. Antibody that was not retained by the AT_{1b} resin was denoted as AT_{1a} receptor selective. Antibody that was retained by both the AT_{1a} and AT_{1b} resins was defined as nonselective for AT_{1a} or AT_{1b} -receptors. A similar strategy was used to derive AT_{1b} selective antibodies except that serum from rabbits immunized with the peptide corresponding to the AT_{1b} receptor subtype was affinity purified using chromatography resin cross-linked with the AT_{1b} peptide initially. Antibodies retained by the AT_{1b} resin were subsequently applied to the AT_{1a} resin. Antibodies retained by the AT_{1b} , but not the AT_{1a} resin, were classified as AT_{1b} selective. AT_2 receptor antibodies were affinity purified using chromatography resin cross-linked with the AT_2 receptor peptide used to generate the antibody. Antibodies retained by the AT_2 resin were eluted with high salt solution and classified as AT_2 selective.

2.2. Animals. Adult male Sprague-Dawley rats (225–300 g body weight; Harlan, Sprague Dawley) were kept in an AAALAC approved vivarium (12:12 Light:Dark). Standard lab chow and water were available ad lib. Animals were kept in the vivarium for at least two weeks prior to use and were housed two per cage. All procedures were approved by the University of Wisconsin, School of Veterinary Medicine Animal Care Committee.

2.3. Western Immunoblotting. Fresh or frozen, whole or dissected rat adrenals ($n = 4$) were employed. A 2 mm slab was cut from the center of the adrenal and the medulla was removed by punch. The cortex was dissected away from the medulla. Tissues were homogenized in one complete mini protease inhibitor tablet (Roche, Indianapolis, IN) dissolved in 7 mL of RIPA buffer (Millipore, Billerica, MD). Lysates were sonicated for 5 minutes and cleared of debris by centrifugation at 15000 rpm for 20 minutes. Samples were normalized so as to amount of protein present via BCA assay (Thermo Scientific, Rockford, IL).

Samples were dissolved 1:1 in loading buffer with beta mercaptoethanol and boiled at 95°C for 4 minutes before loading. Proteins were separated via SDS-PAGE and transferred to PDVF membrane (Bio-Rad, Hercules, CA). Transfer conditions were wet (1 hour at 100 volts). Membranes were incubated for one hour in tris buffered saline containing 0.05% tween-20 (TBST), 5% powdered milk, and 1% bovine serum albumin. Blots were incubated in primary antibodies overnight at 4°C. Primary antibodies (Table 1) were diluted in TBST with 0.2% NaN₃ as a preservative. Blots were incubated in secondary antibody for 45 minutes. Secondary antibody goat anti-rabbit HRP (KPL, Gaithersburg, MD) was diluted 1:100,000 in 20 mL TBST with 2 uL streptavidin HRP (Sigma Aldrich, St Louis, MO). Developing solutions used in this study were LumiGLO immunoblotting reagent (KPL) and Supersignal West Pico Substrate (Thermo Scientific).

2.4. Tissue Preparation. Rats were deeply anesthetized with isoflurane or pentobarbital (65 mg/kg IP) and perfused intracardially with physiological flush solution (Tyrodé's solution) containing heparin and procaine followed by histological fixative (4% paraformaldehyde with 0.05% glutaraldehyde in 0.1 M sodium phosphate, pH 7.5). Brains, pituitaries, and adrenals were removed and immersion fixed at 4°C in the same solution overnight and then stored in saline until sectioning at 50 micron thickness for immunofluorescence microscopy using a Lancer vibratome.

2.5. Immunofluorescence Histochemistry. Adrenals, pituitaries, and brains from 12 rats were used for these studies. Initially all antibodies were screened at dilutions of 1:100 to 1:10,000 in ICC buffer (PBS with 0.25% gelatin, 2% normal goat serum 0.1% thimerosal, and 0.05% neomycin) to determine working dilutions demonstrating the highest signal and lowest background signal for each tissue. Working dilutions of angiotensin II receptor antibodies were (1:500) primary antibody (AT_{1a}, AT_{1b}) and 1:2000 AT₂ for 18–72 hours at 4°C. Control sections were incubated with primary

antibodies incubated with an excess of the antigenic peptides (20 µg/mL of antigenic peptide at the working dilution). Also antibodies were immunoprecipitated from their working dilutions by incubation with 100 µL AT_{1a}, AT_{1b}, or AT₂ affinity gels and then the supernatant was used in place of the antibody solution. Sections were then incubated with Cy3-labeled goat anti-rabbit IgG and then mounted onto poly-L-lysine slides. Slides were viewed and analyzed utilizing a Nikon Eclipse E600 epifluorescence microscope with UV illumination, and a digital camera (Spot RT, Diagnostic Products).

2.6. Immunoelectron Microscopy. Adrenals from 7 rats were used for ultrastructural immunocytochemistry ($N = 4$ rats for immunogold detection and $n = 3$ rats for peroxidase). For both methods, rats were perfused as described above and postfixed for 24 hours in 4% paraformaldehyde with 0.1% glutaraldehyde, washed in PBS and vibratome sectioned at 50 micron thickness. The sections were incubated in 0.1% sodium borohydride 15 minutes, permeabilized in 0.05% triton for one hour, and blocked in either 0.5% BSAc (Aurion, Arnhem, Gelderland, The Netherlands) for one hour for immunogold detection or ICC buffer for immunoperoxidase detection prior to overnight exposure to primary antibody. The primary antibody dilution for AT_{1a} receptors was 1:500 for both immunogold and immunoperoxidase.

For the immunogold method antibody-labeled receptor was detected using ultrasmall gold (Aurion, 0.8 nanometer average size) diluted 1:100 in phosphate buffer and incubated overnight. Tissues were then postfixed in 2.5% glutaraldehyde for 30 minutes. The immunological signal was silver intensified by incubation in R-Gent SE-EM (Aurion) for one hour. For immunoperoxidase detection antibody-bound receptor was incubated with peroxidase labeled goat anti-rabbit IgG-Fab (1:250 overnight in the refrigerator). Peroxidase signal was visualized by incubation in diaminobenzidine (30 mg %) and hydrogen peroxide (0.01%) for 10 minutes in 0.1 M Tris HCL, pH 7.5. Then both immunogold and immunoperoxidase sections were rinsed in 0.1 M sodium phosphate buffer, fixed with osmium, dehydrated through an alcohol series to propylene oxide, and flat embedded in EMBED 812 resin (Electron Microscopy Sciences, Hatfield, PA).

Ultrathin sections were cut and adsorbed to grids coated with Formvar film (Electron Microscopy Sciences), and contrasted with uranyl acetate and lead citrate. All samples were examined and photographed with a Philips CM 120 STEM electron microscope and a Megaview 3 SIs digital camera (Olympus, Munster, Westphalia, Germany) in combination with the software program iTEM (Olympus) at the University of Wisconsin Madison Electron Microscope Facility.

3. Results

Western blotting of protein extracts of the adrenal with the 3 antibodies revealed primary ~69, ~75, and ~71 kD bands for the AT_{1a}, AT_{1b}, and AT₂ receptors, respectively, with secondary bands of ~116, ~126, and ~119, respectively (Figure 1). This suggests that the solubilized receptor was

TABLE 1: Tabulated summary of comparative regional and cellular distribution of Ang II receptor-immunoreactivity in rat brain and pituitary.

Region	AT _{1a}	AT _{1b}	AT ₂
Neocortex			
Lamina I-II	ND	0 ⁺ /5*	1 ⁺ /1*
Lamina III-IV	ND	4 ⁺ /5*	0 ⁺ /4*
Lamina V-VI	ND	4 ⁺ /4*	3 ⁺ /3*
Basal (Anterior) Forebrain			
Entorhinal cortex	2 ⁺ /0*	0 ⁺ /0*	5 ⁺ /1*
Hippocampus			
CA1	1 ⁺ /0*	ND	4 ⁺ /5*
CA3	1 ⁺ /0*	ND	5 ⁺ /5*
Dentate Gyrus	ND	ND	2 ⁺ /2*
Central Amygdala	1 ⁺ /0*	ND	4 ⁺ /3*
Caudate nucleus	3 ⁺ /1*	1 ⁺ /0*	5 ⁺ /3*
Thalamus			
Medial Dorsal Thalamus	0 ⁺ /0*	0 ⁺ /0*	5 ⁺ /2*
Periventricular nucleus of the thalamus	0 ⁺ /0*	0 ⁺ /0*	3 ⁺ /5*
Medial Habenula	1 ⁺ /0*	0 ⁺ /0*	5 ⁺ /3*
Lateral Habenula	0 ⁺ /0*	0 ⁺ /0*	0 ⁺ /0*
Septal Area			
Dorsal Median preoptic nucleus	2 ⁺ /3*	3 ⁺ /4*	4 ⁺ /5*
Medial Septum	0 ⁺ /0*	0 ⁺ /0*	0 ⁺ /2*
Lateral Septum	2 ⁺ /1*	1 ⁺ /1*	2 ⁺ /0*
Hypothalamus			
Anterior Hypothalamic Area	2 ⁺ /3*	1 ⁺ /1*	3 ⁺ /4*
Lateral Hypothalamic Area	3 ⁺ /2 ⁺	4 ⁺ /5*	3 ⁺ /0*
Paraventricular nucleus	4 ⁺ /0*	5 ⁺ /5*	5 ⁺ /3*
Periventricular area	3 ⁺ /0*	2 ⁺ /5	5 ⁺ /0*
Suprachiasmatic nucleus	1 ⁺ /0*	0 ⁺ /0*	2 ⁺ /5*
Arcuate nucleus	5 ⁺ /4*	5 ⁺ /5*	4 ⁺ /0*
Circumventricular Organs/Pituitary			
Median Eminence	0 ⁺ /2*	0 ⁺ /3*	0 ⁺ /4 ⁺
Subfornical Organ	4 ⁺ /5*	0 ⁺ /2*	3 ⁺ /5*
Area Postrema	3 ⁺ /0*	4 ⁺ /4*	3 ⁺ /4*
Posterior Pituitary (pars nervosa)	0 ⁺ /3*	0 ⁺ /2*	ND
Anterior Pituitary (pars distalis)	0 ⁺ /0*	5 ⁺ /0*	5 ⁺ /0*
Stellate cells	0 ⁺ /0*	5 ⁺ /0*	1 ⁺ /0*
Ovoid cells	0 ⁺ /0*	1 ⁺ /0*	4 ⁺ /0*
Cerebellum			
Purkinje Cells	0 ⁺ /0*	5 ⁺ /5*	ND
Hindbrain			
RVLM	4 ⁺ /2*	2 ⁺ /2*	ND
NTS	3 ⁺ /4*	5 ⁺ /2*	3 ⁺ /4*

Key: ⁺ refers to neuronal cell bodies/^{*} refers to fibers. Scored on a scale from 0 to 5. ND is not determined. No AT_{1a} immunoreactivity was observed in glia. AT_{1b} and AT₂ immunoreactivity were observed in glia.

glycosylated since the theoretical molecular weights of the deglycosylated receptors are 40759 Daltons for the AT_{1a}, 40781 Daltons for the AT_{1b}, and 41200 Daltons for the AT₂ receptor. The secondary bands most likely represent dimerized receptors or receptor-protein complexes.

Immunofluorescent staining of the adrenal with the 3 antibodies gave differing discrete staining patterns in the adrenal. Using a working dilution of 1:500 AT_{1a}, immunoreactivity was seen in both the adrenal medulla and the zona glomerulosa (Figures 2(a), 2(d), and 2(g)). The staining

was primarily cytoplasmic in both regions, although in the medulla, localization to the cell membrane is apparent in some cells (Figure 2(g)). AT_{1b} immunoreactivity was present in abundance in the zona glomerulosa of the adrenal (Figures 2(b) and 2(e)). The immunofluorescence was primarily localized to the cell membrane (Figure 2(e)). Weak AT_{1b} immunoreactivity was also present in the zona reticulata (Figure 2(e)). AT_{1b} immunostaining was nearly nonexistent in the medulla (Figure 2(h)).

AT₂ immunoreactivity was abundantly present in both the adrenal medulla and the zona glomerulosa (Figures 2(c), 2(f), and 2(i)). The AT₂ immunofluorescence was also primarily cytoplasmic although a plasma membrane localization was seen in many medullary cells (Figure 2(i)). No immunofluorescent signal was seen in any sections incubated with the antigenic peptide preadsorbed antibodies (not shown).

Immunoelectron microscopic analysis of the subcellular localization of AT_{1a} receptors in the zona glomerulosa and medulla is shown in Figure 3. Both cell membrane and cytoplasmic labeling for AT_{1a} receptors was seen in these cells. AT_{1a} receptor immunogold labeling of endocytic vesicles and mature multivesicular vesicular bodies was seen in glomerulosa cells (Figures 3(b) and 3(c)) and immunoperoxidase labeling of cell membrane and newly forming endocytic vesicles was seen in medullary cells (Figure 3(e)). Intranuclear AT_{1a} receptor immunogold staining was observed in cells of the zona glomerulosa. However, AT_{1a} receptor-immunogold staining was not evident in mitochondria or endoplasmic reticulum of either glomerulosa or medullary cells.

AT_{1b} immunoreactivity was observed in the pars distalis of the anterior pituitary. It was primarily localized to stellate cells, but significant numbers of ovoid cells were also immunopositive. By contrast, AT_{1a} immunoreactivity was not observed in the pituitary (Figure 4). AT₂ receptor immunoreactivity also was observed in the pars distalis of the anterior pituitary, primarily in ovoid cells. AT_{1a} and AT_{1b} receptor immunoreactivity was observed on nerve fibers in the posterior pituitary (Table 1). No Ang II receptor immunoreactivity was observed in the intermediate lobe of the pituitary.

In sections from the brain, neurons were immunopositive for all three receptors, but glial cells showing astrocytic (and microglial, Figure 4 center panel) characteristics were immunopositive only for AT_{1b}. Immunoreactivity for all three angiotensin receptor subtypes was present in abundance in brain regions reported to have high angiotensin receptor density by ligand binding studies and other immunohistochemistry studies (Figures 4–7, Table 1). These regions include the SFO, median eminence, PVN, NTS, and area postrema (Figures 4 and 5, Table 1). In all five of these locations, we demonstrated the presence of all three receptors, although their distribution within each region was not identical (Table 1). Of note, AT_{1b} receptor immunoreactivity was present in the magnocellular division of the PVN while AT₂ receptor immunoreactivity was present in the supraoptic nucleus (SON) (Figure 5). AT₂ receptors were more widely distributed than AT_{1a} and AT_{1b} receptors in the brain, and their immunoreactivity was found

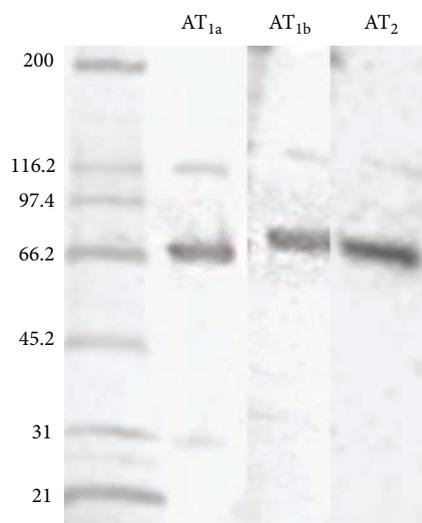


FIGURE 1: Western immunoblots for AT_{1a}, AT_{1b}, and AT₂ receptors of crude extracts of whole adrenals. The three receptors show major bands at ~69–75 kD and well as faint bands at about ~116–126 kD. AT_{1a} receptor-directed antibody (Rabbit 92578-sel), AT_{1b} receptor-directed antibody (Rabbit 92587-sel), AT₂ receptor-directed antibody (Rabbit 92595).

in every region in which AT₁ receptor immunoreactivity was observed (Table 1). AT₂ receptor immunoreactivity was found exclusively in the amygdala, piriform cortex, thalamus, and medial epithalamus (Figures 5 and 6, Table 1).

Angiotensin II receptor immunoreactivity also was found in rat brain regions generally reported to have low expression of Ang II receptors. These include neurons in the cerebral cortex (AT_{1b} and AT₂), hippocampus (AT_{1a} and AT₂), caudate nucleus (AT_{1a}, AT_{1b} and AT₂), and SON (AT₂) (Figure 5).

4. Discussion

4.1. Antibody Development Strategy. The results of these studies unequivocally demonstrate a differential distribution of AT_{1a}, AT_{1b}, and AT₂ receptor immunostaining. This was accomplished by precise epitope targeting within the C-terminus of each receptor, selective antipeptide affinity chromatographic purification methods, Western blotting, and tissue specificity studies in adrenal and pituitary where the distribution of these AT receptor-expressing cells has been established by in situ hybridization and receptor binding studies.

The initial identification of the two subtypes of AT₁ Ang II receptors in rodents demonstrated the presence of mRNA for both the AT_{1a} and AT_{1b} subtype in the rat adrenals [32, 38, 49]. The AT_{1b} was identified as the predominant AT₁ receptor subtype in the rat adrenal based on mRNA expression [38, 49]. While these initial observations have been confirmed in the rat adrenal [50, 51], the AT_{1a} is considered to be the predominant AT₁ receptor subtype in all other rat tissues except the anterior pituitary based on mRNA expression [38, 52].

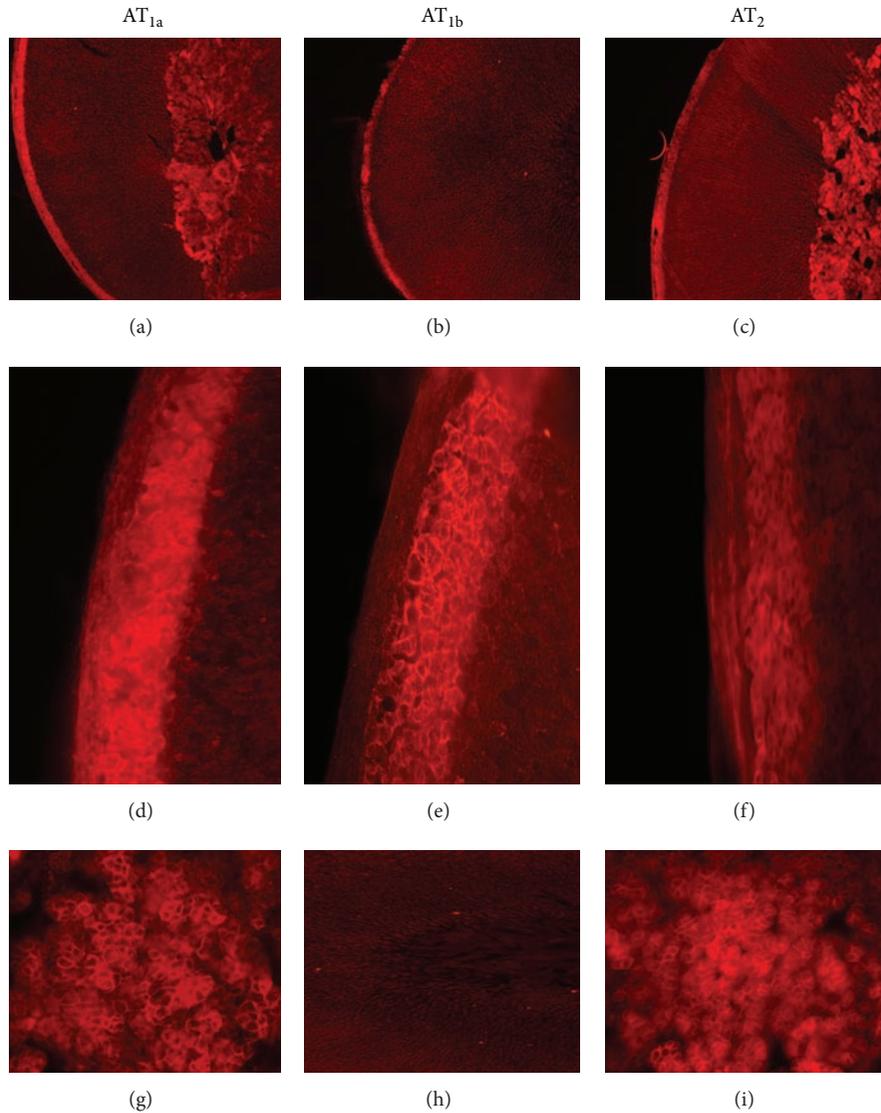


FIGURE 2: Immunofluorescent localization of AT_{1a} , AT_{1b} , and AT_2 receptors in rat adrenals. Survey photomicrographs show positive immunofluorescence for AT_{1a} ((a) and (d); 80x), AT_{1b} ((b) and (e)), and AT_2 ((c) and (f)) in the zona glomerulosa (160x). Positive staining for AT_{1a} ((a) and (g)) and AT_2 ((c) and (i)), but not AT_{1b} ((b) and (h)), is present in the adrenal medulla. The antibodies used were those used in Figure 1.

It is important to be able to discriminate AT_{1a} and AT_{1b} receptor protein expression, because their mRNAs are differentially regulated [31, 39, 49, 52–54]. Furthermore, it is important to determine if the changes in mRNA expression translate into changes in expression of these receptor subtypes, because mRNA expression does not always correlate with protein expression. For example, in the kidney losartan increases AT_{1a} receptor mRNA expression, but decreases AT_1 receptor binding [55]. The existence of miRNAs for angiotensin receptors, for example, miR-155 [56] further erodes the value of mRNA levels as indicators of angiotensin receptor protein expression. Functionality of the subtypes may also differ; AT_{1a} and AT_{1b} can stimulate aldosterone release, while AT_{1a} , but not AT_{1b} , can stimulate corticosterone release in the mouse adrenal [57].

In view of the near identical pharmacological characteristics of the AT_{1a} and AT_{1b} receptor subtypes [34–36], the only way to discriminate these two proteins is to exploit immunological differences arising from differences in their amino acid sequences. While the AT_{1a} receptor (accession no. P25095 <http://www.ncbi.nlm.nih.gov/protein/113493> (accessed 16 March 2012) and AT_{1b} receptor (accession no. NP 112271) <http://www.ncbi.nlm.nih.gov/protein/82524858NP112271> (accessed 16 March 2012) subtypes are encoded by separate genes, they are ~95% identical and are both made up of 359 amino acids [33, 38]. Thus there are only a few regions of these receptors where they differ substantially in amino acid sequence. One of these regions, near the carboxy terminus of the receptor proteins (amino acids 352 to 355), has 2 different amino acids in this 4 amino

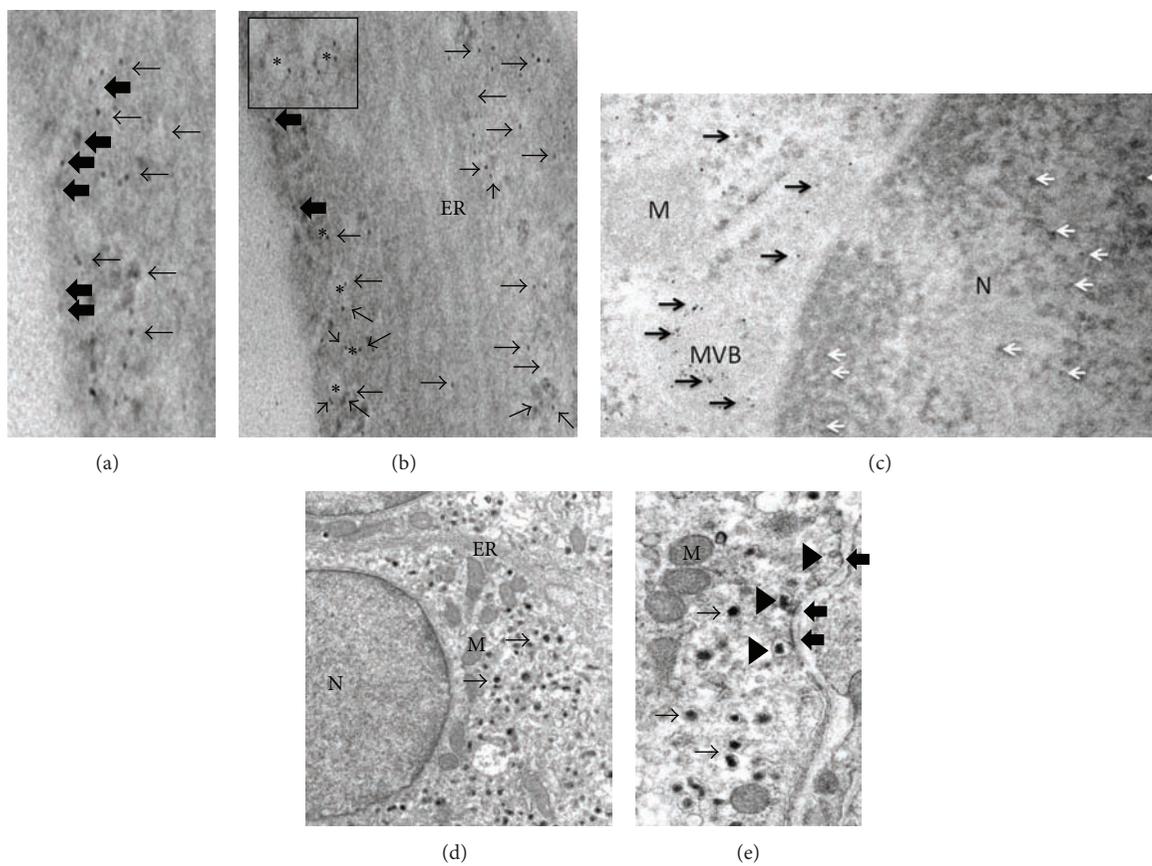


FIGURE 3: Ultrastructural immunocytochemistry of the AT_{1a} receptor (using anti-AT_{1a} receptor #92578-sel in zona glomerulosa (Figures 3(a) through 3(c); 48,000x) and adrenal medulla (Figures 3(d) and 3(e)). Immunogold ultrastructural analysis of zona glomerulosa shows AT_{1a} receptor (showing localization at the cell membrane (bold arrows; Figures 3(a) and 3(b)), in the cytoplasm (line arrows; Figures 3(a) through 3(c)), and on the surface of endocytic vesicles (insert Figure 3(b)). Immunogold particles were also seen in a multivesicular body (MVB) and in the nucleus (Figure 3(c)). Immunoperoxidase staining of adrenal medullary cells reveals a large number of AT_{1a} positive vesicles (line arrows; Figure 3(d); 20,000x and Figure 3(e); 48,000x), patches of membrane receptors (block arrows; Figure 3(e)), and apparent omega body fusion with the cell membrane (arrow heads; Figure 3(e)). Note the lack of localization in the mitochondria (M) and endoplasmic reticulum (ER).

acid stretch. The closest similarities to the sequences of the AT₁ antigenic peptides in the protein database (Protein Blast) <http://blast.ncbi.nlm.nih.gov/Blast.cgi?PROGRAM=blastp&BLASTPROGRAMS=blastp&PAGE=blastp&SHOWDEFAULTS=on&LINK LOC=blasthome>, accessed on February 4, 2013) were the serotonin 5 HT_{2b} subtype with a 7 amino acid identity to the AT_{1a} peptide fragment (score = 24.0 bits) and sestrin 1 with a 7 amino acid identity to the AT_{1b} peptide fragment (score = 24.4 bits).

To generate an antibody to the AT₂ receptor, a similar strategy was applied. A C-terminal domain peptide of 15 amino acids (resembling amino acids 349 to 363) was used as the antigen. The sequence of the AT₂ receptor (accession no. P35351, <http://www.ncbi.nlm.nih.gov/protein/543780> accessed on February 4, 2013) has negligible homology with either of the AT₁ receptor subtypes. The closest similarity to this peptide sequence was an immunoglobulin kappa chain (AAA41415.1) with an 8 amino acid identity to the AT₂ peptide fragment (score = 27.4 bits compared to 49.0 bits for the AT₂ receptor).

4.2. Adrenal AT Receptor Subtype Localization. The presence of AT_{1a}, AT_{1b}, and AT₂ angiotensin receptor subtype immunoreactivity in the rat adrenal was clearly demonstrated in this study. AT_{1a} and AT₂ receptor subtype immunoreactivities were found in both the zona glomerulosa and medulla, which is consistent with receptor binding studies [37, 58–61] and mRNA studies [44, 61–64]. AT_{1b} receptor was not observed in the adrenal medulla, but was present in the zona glomerulosa. This is consistent with in situ hybridization studies of the distribution of AT_{1b} mRNA in the adrenal [39, 44, 46, 54].

Other studies of the localization of Ang II receptor subtype immunoreactivity in the adrenal have given mixed and controversial results. Paxton et al. [65] observed AT₁ receptor immunoreactivity in the zona glomerulosa of the rat adrenal with an antibody prepared against amino acids 15–24 of the rat AT_{1a} and AT_{1b} receptor. However, they did not observe any AT₁ receptor immunoreactivity in the adrenal medulla. Similarly, Lehoux et al. [66] observed AT₁ immunoreactivity in the zona glomerulosa of the rat adrenal

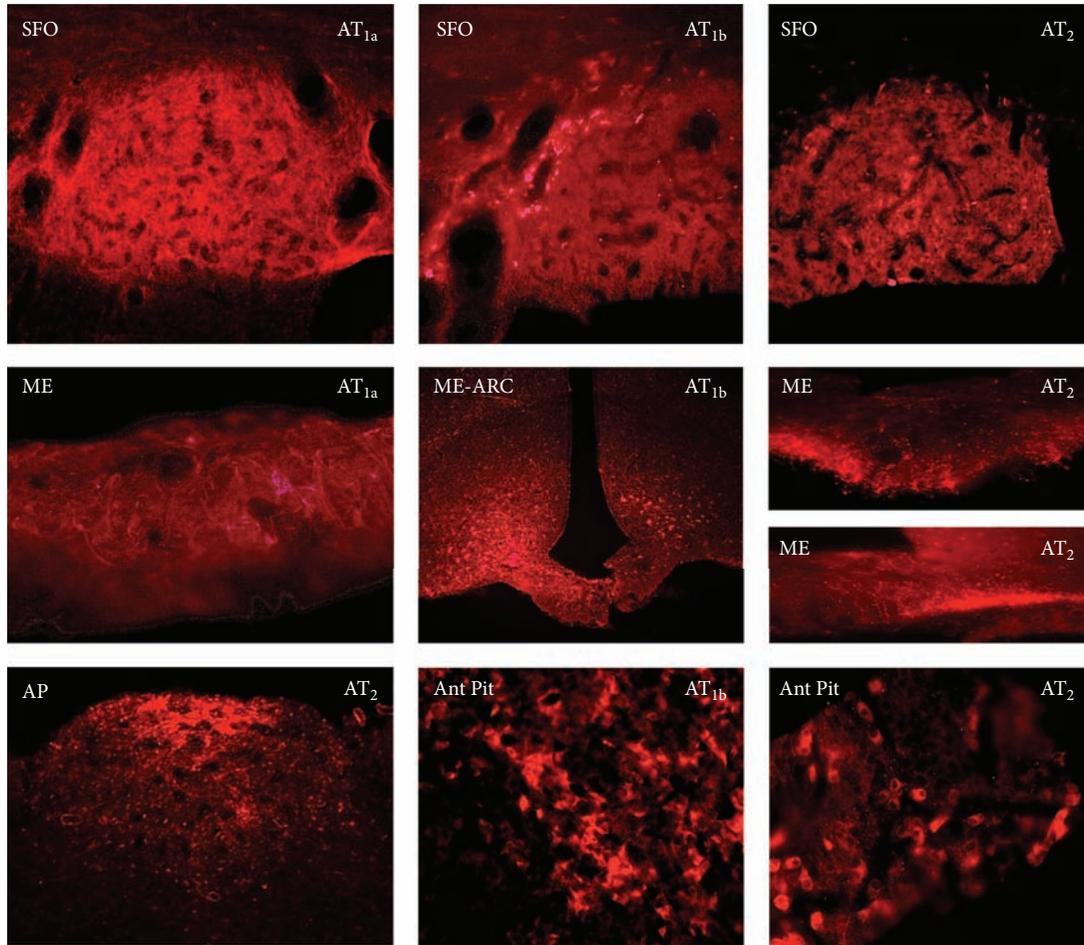


FIGURE 4: Circumventricular organs and pituitary AT receptor immunolocalization. Subfornical organ (top row left to right; 180x) AT_{1a}, AT_{1b}, and AT₂. Median eminence (middle row left to right AT_{1a} (160x), AT_{1b} (100x), and AT₂ (120x)). Bottom row area postrema (AP, 160x) AT₂ and anterior pituitary (pars distalis localization; 120x) of AT_{1b} (middle) and AT₂ (right). No staining for AT_{1a} was seen in the anterior pituitary.

cortex, but not in the medulla using an antibody raised against amino acids 306–359 of the human AT₁ receptor subtype. Of note, adrenals from rats kept on a low sodium diet displayed AT₁ immunoreactivity in other cortical zones (fasciculata and reticularis). The lack of adrenomedullary staining with this human antibody suggests that it may only recognize the AT_{1b} sequence in the rat. Giles et al. [67] observed AT₁ immunoreactivity and AT₁ mRNA in the zona glomerulosa of rat adrenals using an antibody directed against amino acids 350–359 of the rat AT_{1a} subtype plus a small amount of immunoreactivity in the zona fasciculata. However, there was no mention of AT₁ immunoreactivity or mRNA in the adrenal medulla.

Frei et al. [68] observed AT₁ immunoreactivity in the rat adrenal cortex and medulla using a monoclonal antibody raised against amino acids 229–246 of the human AT₁ receptor subtype. Yet, AT₂ immunoreactivity was only observed in the rat adrenal medulla using an antibody raised against amino acids 314–330 of the human AT₂ receptor subtype. On the other hand, Harada et al. [63] observed AT₂ receptor immunoreactivity in immunoblots of the rat adrenal cortex, but not in the medulla using two different

antibodies—one raised against amino acids 21–35 of the rat AT₂ receptor and one raised against amino acids 221–363 of the human AT₂ receptor. However, they did detect a low level of AT₂ receptor-like immunoreactivity in the medulla using the latter antibody for immunohistochemical analysis. Conversely, Yiu et al. [69] reported AT₂ immunoreactivity only in the rat adrenal medulla using an antibody directed against amino acids 341–351 of the rat AT₂ subtype. Notably, they reported that this antibody failed to label brain regions known to express AT₂ receptors. Reagan et al. [70] were unable to demonstrate any AT₂ receptor immunoreactivity in the rat adrenal using a polyclonal antibody developed to recognize AT₂ receptors in N1E-115 cells.

4.3. Subcellular Localization of AT₁ Receptors. Localization of immunofluorescence for all three angiotensin receptor subtypes to the cell membrane as well as the cytoplasm in the adrenal is consistent with the behavior of other G protein coupled receptors that are functionally expressed on cell membranes but undergo receptor-mediated internalization [71]. The electron microscopic localization of AT_{1a}

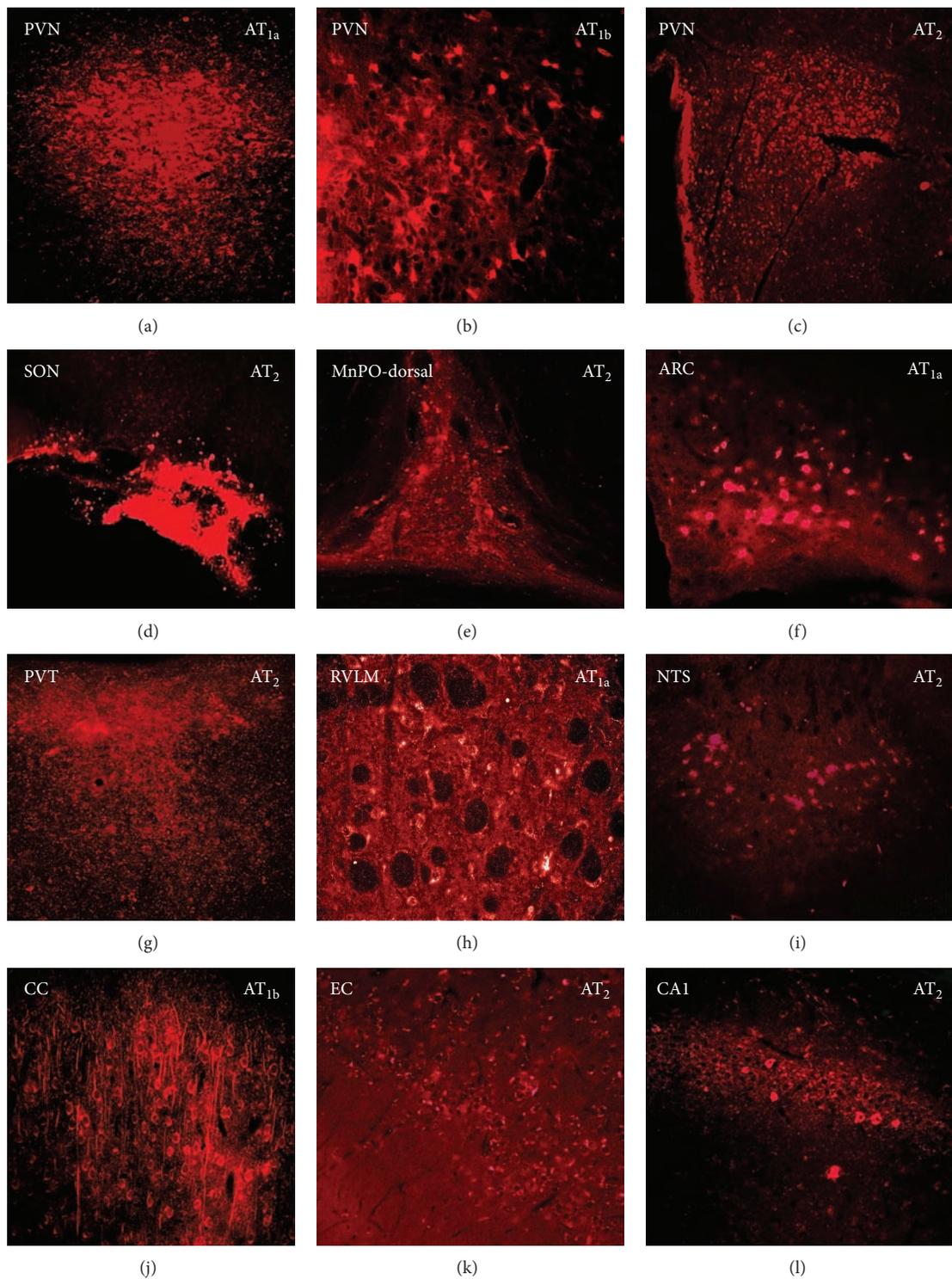


FIGURE 5: Immunofluorescent localization of AT receptors in various brain nuclei. Top row: hypothalamic paraventricular nucleus (PVN, 160x) AT_{1a}, AT_{1b}, and AT₂. Second row: immunofluorescence labeling (left to right) for AT₂ in supraoptic nucleus (SON; 120x) and median preoptic nucleus (MnPO-dorsal part; 80x), and AT_{1a} localization labeling of arcuate nucleus (ARC; 160x). Third row, (left to right): AT₂ in the periventricular nucleus of the thalamus (PVT; 160x), AT_{1a} receptor in the rostral ventrolateral medulla (RVLM, 100x), and AT₂ receptor in nucleus of the solitary tract (NTS; 160x). Bottom row (left to right): AT_{1a} in frontal parietal cortex (160x), AT₂ in entorhinal cortex (80x), and AT₂ in hippocampus CA1 (120x).

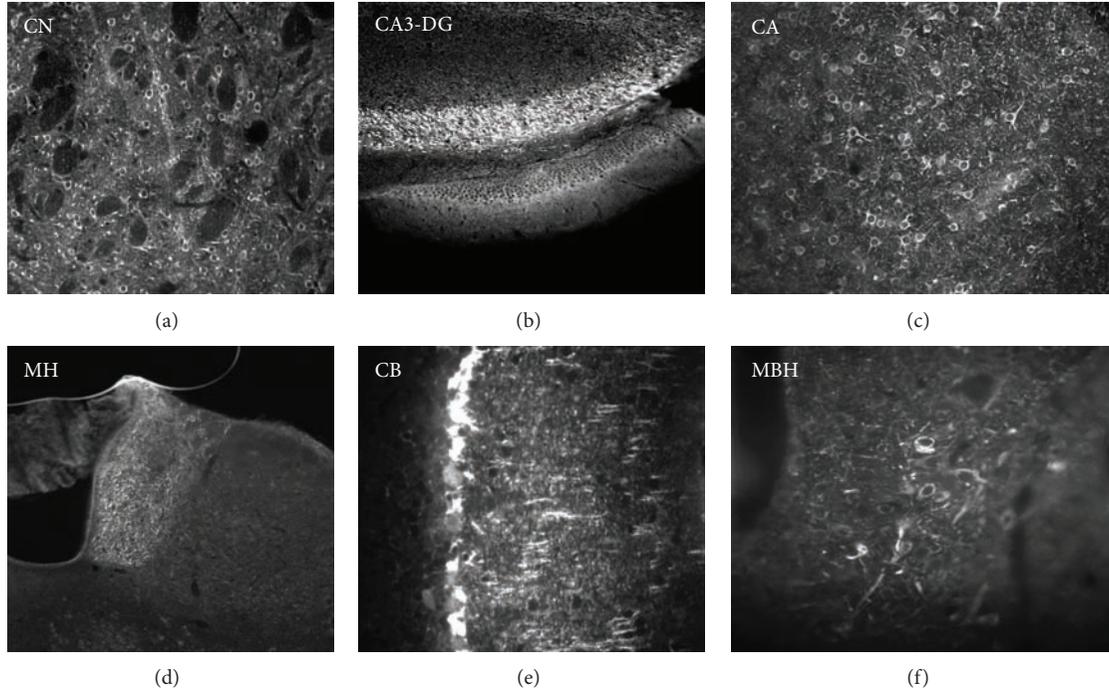


FIGURE 6: AT₂ receptor immunolocalization in caudate nucleus (CN, 400x), CA3-dentate gyrus of the hippocampus (CA3-DG; 80x), central nucleus of the amygdala (CA; 160x), medial habenula (MH, bottom left; 80x), AT_{1b} in cerebellar Purkinje cells (CB; bottom middle; 100x), and AT₂ in mediobasal hypothalamus (MBH; bottom right, 320x).

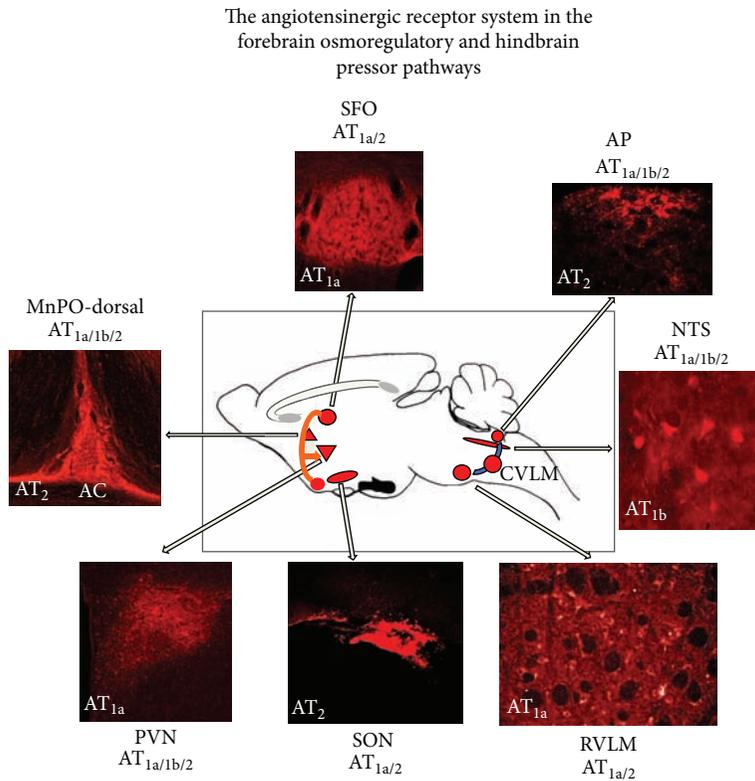


FIGURE 7: Diagrammatic summary of important brain nuclei in the angiotensinergic forebrain osmoregulatory pressor (orange pathway) and hindbrain pressor (blue) pathways. Note that there is more than one AT receptor in each site as given adjacent to each micrograph. A representative AT receptor for each site is shown within the figure.

immunoreactivity to putative developing endosomes still in contact with the cell membrane (Figure 3(e)) is consistent with receptor mediated endocytosis as the mechanism of angiotensin receptor internalization [72]. In addition, there is now a considerable body of evidence supporting the existence of an intracellular RAS which signals via AT₁ receptors [73].

Noteworthy in our study is the nuclear localization of adrenal AT_{1a} receptors. The ability of G protein coupled receptors to localize and signal directly to the cell nucleus is firmly established [74] and likely includes angiotensin receptors. Beginning with the electron microscopic studies localizing ³H-Ang II to myocardial cell nuclei [75], it has been suspected that Ang II receptors are present in cell nuclei. The existence of nuclear Ang II receptors was subsequently documented in isolated hepatic nuclei by Re and Parab [76] who showed that Ang II increased RNA polymerase II activity, increasing RNA synthesis. Notably, they used 4 mM dithiothreitol an inhibitor of Ang II binding to AT₁ receptors [77], suggesting that the Ang II effect might be mediated by AT₂ receptors. Eggena et al. [78] showed that AT₁ receptor subtype binding was present in rat hepatic cell nuclei and that Ang II could specifically induce transcription of mRNA for renin and angiotensinogen in isolated rat liver nuclei. Moreover, hepatic nuclear AT₁ receptor binding and functionality could be dynamically regulated by adrenalectomy and nephrectomy [79]. Re et al. [80] and Eggena et al. [79] reported that nuclear Ang II receptor binding was associated with nuclear chromatin. Of note, Re et al. [80] observed ¹²⁵I-Ang II binding to nuclear chromatin in the presence of 5 mM dithiothreitol, again suggesting that ¹²⁵I-Ang II may be binding to AT₂ receptors [15, 77]. The relative abundance of AT_{1a} binding within the nucleus, but not the nuclear membrane of the glomerulosa cells in this study, is consistent with localization to nuclear chromatin. AT₁ receptor binding sites have also been identified in rat hepatocyte nuclear membranes by Booz et al. [81] and Tang et al. [82]. Interestingly, Tang et al. [82] determined that the majority of the AT₁-like binding of Ang II in hepatocyte nuclei was bound to a soluble intranuclear protein. Licea et al. [83] demonstrated nuclear Ang II receptor binding in nuclei of rat renal cortex. Tadevosyan et al. [84] showed that Ang II could stimulate α -³²P-UTP incorporation into RNA and increase NF-kappaB mRNA expression in isolated rat heart cardiomyocyte nuclei suggesting a nuclear site of action of Ang II.

Additional evidence supporting a nuclear localization of angiotensin receptors includes studies using an AT₁ receptor-GFP fusion construct which translocates to the nucleus in Chinese hamster ovary cells [85] and human embryonic kidney (HEK-293) cells [86], as well as immunohistochemical studies showing colocalization of AT₁ and AT₂ immunoreactivity with the nuclear membrane markers nucleoporin-62 and histone-3 [84]. Moreover, the AT₁ receptor contains a nuclear localization signal motif (KKFVK, 307-11) in its intracellular carboxy terminal tail [87], which promotes its translocation to the cell nucleus. Mutation of one amino acid in this motif (K307Q) in an AT_{1a} r-GFP receptor construct prevents it from localizing to the nucleus of HEK293 cells

[86]. Of note, both agonist induced [87] and agonist independent [71, 88] nuclear localization of AT₁ receptors has been reported.

While there are no published reports of adrenal nuclear angiotensin receptor binding or function, Eggena et al. [78] reported preliminary data suggesting that Ang II could stimulate RNA transcription in isolated adrenal nuclei. In addition, Goodfriend and Peach [89] suggested that Ang III can act intracellularly in the zona glomerulosa to promote aldosterone production.

4.4. Pituitary AT Receptor Subtype Localization. Both AT_{1b} and AT₂ receptor immunoreactivities were present in high amounts in the anterior pituitary. As noted previously mRNA for AT_{1b} receptors is abundant in the anterior pituitary, while AT_{1a} mRNA is much less abundant and AT₂ mRNA is not observed in the anterior pituitary [47, 90]. Autoradiography and radioligand binding studies have demonstrated a high density of Ang II receptors in the anterior pituitary [37, 91–93]. This binding displays AT₁ receptor characteristics, and little or no AT₂ receptor binding has been observed [27, 94]. AT_{1b} expression was highest in stellate cells, while AT₂ expression was highest in ovoid cells. Both AT_{1a} and AT_{1b} immunoreactivity was present on nerve fibers in the posterior pituitary. The ability of Ang II to affect the release of pituitary hormones is well known [95]. There are no reports of Ang II receptor binding in the posterior pituitary of the rat, although there is one report of AT₁ receptor-immunoreactivity in nerve fibers and cell bodies in the posterior pituitary [96] and one report of AT₂ receptor-immunoreactivity in the posterior pituitary as well as in the vasopressinergic magnocellular division of the PVN and the SON [97]. mRNA studies indicate a predominance of the AT_{1b} subtype in the anterior pituitary of the rat [38, 98–100], with little or no AT_{1a} and AT₂ mRNA.

Many of the pituitary hormone-releasing effects of Ang II occur in the hypothalamus and those effects are discussed below. However, some of the pituitary hormone releasing of Ang II occur directly in the pituitary. Systemically administered Ang II stimulates vasopressin release from the posterior pituitary of the dog [101, 102]; however, this may not generalize to the rat. AT_{1a} and AT_{1b} receptors on nerve fibers in the rat posterior pituitary [96] could mediate these effects of Ang II, reminiscent of the mechanism whereby Ang II acts on sympathetic nerve terminals to stimulate norepinephrine release [103, 104].

Radioligand binding studies have revealed high levels of Ang II receptor binding in a lactotroph enriched pituitary preparation [105]. mRNA studies indicate that AT_{1b} receptors appear most often on lactotrophs, being present on more than 50% of all lactotrophs [98]. The appearance of AT_{1b} immunoreactivity in ovoid cells is consistent with these receptors being present on lactotrophs. It has been reported that AT_{1b} mRNA is present in a somatotroph cell line [100]. Somatotrophs are also ovoid in shape and blood-borne Ang II can inhibit growth hormone release [106], although it has also been reported that Ang II synthesized by and released from lactotrophs can stimulate the release of growth hormone from

somatotrophs, [107] suggesting that somatotrophs may have excitatory AT₁ receptors and inhibitory AT₂ receptors.

ACTH release from dissociated corticotrophs in the anterior pituitary is also stimulated by Ang II *in vitro* [108]. The stimulation decreases with supraphysiological estradiol exposure *in vivo* and correlates positively with reductions in Ang II receptor binding caused by *in vivo* supraphysiological estradiol exposure [108]. Autoradiographic studies of AT₁ receptor binding in the anterior pituitary indicate that AT₁ receptor binding varies with the estrous cycle and that exogenous estrogen decreases anterior pituitary AT₁ receptor binding in ovariectomized rats [109]. mRNA for AT_{1b} receptors in the anterior pituitary is also suppressed by estrogen treatment [38, 110]. The appearance of high levels of AT_{1b} immunoreactivity in stellate cells in this study is consistent with these receptors being present on corticotrophs.

There is one report of AT₂ receptor immunoreactivity in pituitary adenoma blood vessels in humans [96], leading to the hypothesis that AT₂ receptors could participate in tumor-induced angiogenesis.

4.5. Brain AT Receptor Subtype Localization. These studies describe a widespread distribution of AT_{1a}, AT_{1b}, and AT₂ receptor immunoreactivity throughout the rat brain. The receptors were expressed abundantly in a number of brain regions that constitute the cardiovascular regulatory circuits of the brain, as well as the noncardiovascular regulatory regions of the brain. There was considerable variation in the degree of expression of the receptors in different regions reminiscent of the profound differences in radioligand binding for Ang II receptors, particularly among the AT₁ receptors. AT₂ receptors displayed an unanticipated widespread distribution throughout the rat brain, which contrasts with their limited distribution as indicated by radioligand binding studies. While AT₁ receptors are considered to play the predominant role of mediating the actions of Ang II in the brain, AT₂ receptors are increasingly recognized as having an important role as physiological antagonists of AT₁ receptor effects. The codistribution of AT₁ and AT₂ receptors in several brain regions as well as the adrenal is consistent with the concept of colocalization of these two subtypes in the same cells as counter regulators to each other at the cellular level as well as on an organismic level [111–113].

The selective expression of AT_{1b} receptors on astrocytes suggests that there is a cell-specific expression of Ang II receptor subtypes in the brain. Functional AT₁ receptors are present in primary cultures of astroglia from rat brain [114], but questions have been raised as to whether this expression could reflect an altered phenotype of cultured cells not seen *in situ* in a living brain [115]. In contrast, Füchtbauer et al. [116] observed AT₁ immunoreactivity (Santa Cruz, sc-579, amino acids 306–359) in astrocytes of the outer molecular layer of the dentate gyrus of the mouse brain, but did not see AT₁ immunoreactivity in the microglia. Of note, retinal astrocytes also express AT₁ receptor immunoreactivity (Alomone, #AAR-011 amino acids 4–18) while amacrine cells in the rat retina display AT₂ immunoreactivity (Alomone, no. AAR12, amino acids 21–35) [117]. These reports and our observations

suggest that glia do express AT₁ receptors and that they are of the AT_{1b} subtype. Since astrocytes are the primary source of angiotensinogen in the brain, the AT_{1b} receptor may play a role in regulating angiotensinogen in the brain.

The expression of AT_{1b} receptor immunoreactivity on cells with the morphological characteristics of microglia suggests that this receptor subtype mediates the proinflammatory effects of Ang II. AT₁ receptor antagonism blocks the activation of microglia in an animal model of brain inflammation [118]. Proinflammatory cytokine participation in the pressor actions of Ang II in the brain is reversible by AT₁ antagonists [119, 120], suggesting that microglial AT₁ receptors may play a role in blood pressure regulation as well as inflammation.

The concept of the presence of Ang II receptors in the brain was firmly established by the cross-perfusion studies of Bickerton and Buckley [121] showing that blood-borne Ang II had sympathoexcitatory effects mediated by the brain. Since that time, a multitude of methodological approaches have been used to map the distribution of Ang II receptors in the brain. Early radioligand binding studies of brain Ang II receptors [122, 123] indicated that Ang II receptors were located in regions within the blood-brain barrier, for example, cerebellum, hypothalamus, thalamus, septum, and midbrain, as well as outside the blood brain barrier. The first receptor autoradiographic study of brain Ang II receptors for blood-borne Ang II clearly demonstrated their presence in 4 circumventricular organs (CVOs): the SFO, OVLT, median eminence, and area postrema [124]. *In vitro* receptor autoradiographic studies of the rat brain confirmed the localization of Ang II receptors in these CVOs and revealed a widespread distribution of discrete populations of Ang II receptors in a large number of brain nuclei [93, 125]. Subsequent receptor autoradiographic studies using Ang II receptor subtype specific competing ligands indicated that both AT₁ and AT₂ receptors were present in the brain and were differentially distributed [27, 58]. Regions containing high densities of AT₁ receptor binding include regions associated with dipsogenesis and cardiovascular regulation, for example, SFO, OVLT, MnPO, PVN, NTS, dorsal motor nucleus of the vagus, area postrema, rostral ventrolateral medulla (RVLM), as well as noncardiovascular regulatory regions, for example, pyriform cortex, subiculum, and spinal trigeminal nucleus. Generally, regions containing high densities of AT₂ receptor binding are unrelated to blood pressure regulation and dipsogenesis, for example, mediodorsal thalamus, inferior olivary nucleus, medial geniculate, and subthalamic nucleus. While many regions have a strong predominance of one or the other subtype, several brain regions show both AT₁ and AT₂ receptor binding, for example, parabrachial nuclei, pedunculo-pontine tegmental nucleus, locus coeruleus, and superior colliculus [126].

Localized injection of exogenous Ang II has been used to map the distribution of brain Ang II receptors. Early studies directed at determining sites of action of Ang II assessed its behavioral and physiological effects. Subsequent studies using iontophoretic or pressure injection of Ang II via micropipettes have focused on its cellular effects. Early mapping of Ang II receptors mediating its dipsogenic

effects indicated a widespread distribution in the forebrain [127]. However, a subsequent study [128] revealed that all the active sites were targeted with a cannula that traversed the anterior cerebral ventricles, and that only when Ang II leaked into the ventricles that a dipsogenic response occurred. Microinjection of Ang II into the SFO and PVN is excitatory to these neurons [129]. Microinjection of Ang II into the RVLM [130], area postrema, and NTS [131] increases blood pressure. Microinjection of Ang II into the periaqueductal gray increases blood pressure via its actions at AT₁ receptors [132], while microinjection of Ang II into the superior colliculus increases blood pressure via its actions at AT₂ receptors [133] consistent with radioligand binding studies indicating the presence of AT₁ or AT₂ receptors in these regions [27]. Lastly, the distribution of angiotensin responsive neurons has been determined using induction of fos expression as a functional marker [134].

A major controversy involves the presence or absence of Ang II receptors on vasopressinergic and oxytocinergic neurons in the SON and the magnocellular division of the PVN. Stimulation of vasopressin and oxytocin release from the posterior pituitary results from stimulation of the magnocellular neurons in the PVN and SON. In this study, all 3 Ang II receptor subtypes were highly expressed in the magnocellular divisions of the PVN. Radioligand binding studies of Ang II receptors reveal high expression of AT₁ receptors in the parvocellular region of the PVN and low expression of Ang II receptors in the magnocellular division of the PVN and SON (as described in the previous section). Similarly, mRNA studies (succeeding section) have failed to demonstrate measurable Ang II receptor synthesizing capacity in these regions. However, electrophysiological studies suggest that neurons in these regions are responsive to Ang II. Nagatomo et al. [135] showed that Ang II inhibited potassium currents in SON neurons using patch clamping in brain slices. Ang II has a direct excitatory effect in the SON, which is consistent with the presence of AT₁ receptors on vasopressinergic and oxytocinergic neurons [136]. The data reported herein is consistent with the presence of functional AT₁ receptors in the PVN and SON.

Parvocellular PVN AT₁ receptors revealed by radioligand binding and mRNA assays are well placed to stimulate CRH neurons in the PVN to release corticotrophin releasing hormone (CRH) from their nerve terminals in the median eminence into the hypothalamo-hypophyseal portal vessels to act upon corticotrophs in the anterior pituitary. In this study, all 3 Ang II receptor subtypes were highly expressed in the parvocellular division of the PVN.

The use of in situ hybridization or PCR for localization of mRNA to determine sites of synthesis of proteins has been widely used to localize Ang II receptor subtypes in the brain. Kakar et al. [31] reported a predominance of AT_{1b} mRNA in the SFO, OVLT, and cerebellum and a predominance of AT_{1b} in the hypothalamus by PCR. Conversely, Jöhren et al. [45] identified AT_{1a} mRNA in the SFO, OVLT, PVN, cerebral cortex and hippocampus, AT_{1b} mRNA in the cerebral cortex and hippocampus, (but not in the SFO or OVLT) and AT₂ mRNA in the medial geniculate and inferior olivary nucleus. Similarly, Lenkei et al. [41] reported a predominance of AT_{1a}

mRNA expression in the SFO, OVLT, PVN, and MnPO as well as the anterior olfactory nucleus with very low AT_{1b} mRNA expression in the SFO and PVN. Lenkei et al. [137] also reported the absence of AT_{1a} and AT_{1b} mRNA in the vasopressin positive neurons and GFAP positive astroglia in the SON and PVN. In the two-week-old rat brains, Jöhren and Saavedra [138] also observed AT_{1a} mRNA in the pyriform cortex, basal amygdala and choroid plexus and AT_{1b} mRNA in the choroid plexus. AT₁ receptor binding has been reported in the choroid plexus [139] although at very low levels [140].

Brain AT₂ receptor mRNA shows both similarities and differences from AT₂ receptor binding in the rat brain. Noteworthy is the presence of AT₂ mRNA in the red nucleus and the absence of AT₂ mRNA in the locus coeruleus, lateral septum, and cerebellum [141]. These discrepancies have been interpreted as indicating that the red nucleus synthesizes AT₂ receptors that are only expressed on its efferent nerve terminals that project to the inferior olivary nucleus and cerebellum, while the AT₂ receptor expressing brain regions devoid of AT₂ mRNA express AT₂ receptors on the nerve terminals of its afferents from other brain regions. Lenkei et al. [142] observed AT₂ mRNA in the red nucleus. However, they also observed AT₂ mRNA in the lateral septum and locus coeruleus, as well as a much greater number of brain regions, including some traditionally AT₁ predominant regions such as the NTS and spinal trigeminal nucleus. Lenkei et al. [47] also did a comprehensive in situ hybridization analysis of the rat brain AT_{1a} receptor mRNA. Overall this is consistent with AT₁ receptor binding, with a few exceptions, for example, the lack of AT₁ mRNA in arcuate nucleus and median eminence, where it is postulated that the AT₁ receptors occur on nerve terminals of hypothalamic neurons that synthesize dopamine or releasing hormones and release them into the hypothalamo-hypophyseal portal system to act upon endocrine cells of the anterior pituitary. There are also some brain regions that express AT_{1a} mRNA, but not AT₁ receptor binding, such as hippocampus CA1 and CA2 and some thalamic and brainstem nuclei [47]. An area of considerable cardiovascular regulatory significance is the RVLM. Chronic Ang II infusion was shown to up-regulate AT₁ mRNA in the RVLM and reduce it in the SFO, suggesting that enhanced activation of the RVLM by enhanced AT₁ stimulation increases sympathetic nervous system activity [143].

There are a large number of studies that have used immunohistochemistry and Western blotting to identify and localize Ang II receptor subtypes in the central nervous system. The receptor antigens are generally peptide fragments from different domains of the receptor protein, although one antibody [144] was generated from a purified AT₂ receptor protein. Some antibodies target an extracellular domain near the amino terminal for example, Santa Cruz Biotechnology, SC-1173 (amino acids 15–24), the transmembrane spanning regions of the receptor, intra- and extracellular domains between the transmembrane spanning domains, the third intracellular loop (amino acids 225–237) of the AT₁ receptor (Chemicon), and the intracellular carboxy terminal domain. Several of these studies have used antibodies directed against

the same carboxy terminal regions of the AT_{1a} (Abcam, AB18801), the AT_{1a} or the AT_{1b} (Advanced Targeting Systems, AB-N25AP, AB-N26AP, or AB-N27AP), and the AT₂ receptors (Abcam, AB19134; Advanced Targeting Systems, AB-N28AP) that were used for generation of these antibodies.

Localization of AT₁ receptor immunoreactivity in the brain was first done by Phillips et al., [145] using the 225–237 antibody directed to the third intracellular loop of the AT₁ receptor. They showed extensive distribution of AT₁ immunoreactivity in areas identified by receptor autoradiography to have Ang II receptors. Cardiovascular regulatory regions that were AT₁ immunopositive included the PVN, OVLT, SFO, area postrema, NTS, RVLM, and nucleus ambiguus. AT₁ immunopositive neurons were also present in the SON, and magnocellular division of the PVN, medial septal nucleus, LC, superior and inferior olivary nuclei, hypoglossal nucleus, ventral horn of the spinal cord and other regions not generally viewed as AT₁ receptor targets of Ang II. Conversely, some areas reported to express Ang II receptor binding sites, for example, pyriform cortex, suprachiasmatic nucleus did not show AT₁ immunoreactivity. They suggested that Ang II via AT₁ receptors may have an expanded role in the CNS beyond that considered at that time.

Other studies also report the presence of AT₁ receptors in the SON and/or the magnocellular division of the PVN using either an amino terminal peptide fragment directed antibody, AB18801 and AB-N27AP [146, 147] and the antibody directed against the 225–237 fragment of the AT₁ receptor [148, 149]. Of note, the number of cells in the magnocellular division of the PVN expressing AT₁ receptor using AB18801 was dramatically increased in rats with induced heart failure [146]. Two other studies observed an increase in total PVN AT₁ receptor immunoreactivity (Abcam unspecified). In the first study, PVN AT₁ immunoreactivity was increased in a rat model of heart failure [150]. In the second study, PVN AT₁ immunoreactivity was increased with chronic intravenous Ang II infusion that was only partially reversed by ICV losartan infusion [151].

Using an antibody against purified AT₂ receptor protein, Reagan et al. [152] immunohistochemically localized AT₂ receptor immunoreactivity in the rat brain. Regions reported to have AT₂ receptor binding and/or mRNA that were immunopositive included the locus coeruleus and several thalamic nuclei. Other regions reported to be AT₂ expressing included the amygdala and the Purkinje cell layer of the cerebellum. In addition, AT₂ immunoreactivity was present in the magnocellular division of the PVN and SON which further confirms observations in our study. However, as noted above, this antibody did not label the adrenal [70].

A series of studies have used the AT_{1a} carboxy-terminal fragment-directed antibody to identify AT₁ receptor immunoreactivity in the area postrema, NTS and RVLM at the electron microscopic level. AT_{1a} immunoreactivity was present in neuronal cell bodies, dendrites, axon terminals, perivascular glial processes of astrocytes, fibroblasts, and vascular endothelial cells in the area postrema and dorsomedial NTS [153]. This AT_{1a} immunoreactivity colocalized with the gp91^{phox} subunit of NADPH oxidase in neuronal cell bodies,

dendrites, and putative vagal afferents in the medial NTS [154]. Dendritic processes of the medial NTS containing AT_{1a} immunoreactivity also were positive for tyrosine hydroxylase (TH) or adjacent to TH containing axons [155]. In the TH positive neurons of the RVLM, AT₁ receptor expression was greater in female rats than in male rats [156], and this increase was associated with a higher estrogen state (proestrus versus diestrus) and increased plasma membrane expression of AT₁ immunoreactivity [157]. This same group has used the AT₂ fragment directed antibody (AB19134) to identify AT₂ receptor immunoreactivity in the PVN and NTS at the electron microscopic level [158, 159]. These studies have colocalized AT₂ immunoreactivity with neuronal nitric oxide synthase (nNOS) in neuronal cell bodies and dendrites in the medial NTS [159], and with vasopressin in neuronal cell bodies and dendrites in the PVN [158]. This latter observation contrasts with the studies of Lenkei et al. [142], who did not find AT₂ receptor mRNA in the PVN.

Extensive studies of AT₁ and AT₂ immunoreactivity in the RVLM and NTS in animal models of heart failure have been carried out by Gao, Zucker and colleagues using AT₁ and AT₂ antibodies, primarily SC-1173 and SC-9040 [160, 161]. AT₁ receptors in the RVLM and NTS showed increased AT₁ immunoreactivity, while AT₂ receptors showed decreased immunoreactivity. Infusion of Ang II into the brain of rabbits to simulate a heart failure model increased AT₁ receptor immunoreactivity in the RVLM [162]. Interestingly, viral transfection of AT₂ receptors into the RVLM, which was documented with increased AT₂ immunoreactivity, suppressed sympathetic activity in normal rats [163]. In a mouse model of hypertension, the RA mouse [164], immunoreactivity for AT₁ (SC-1173) in the NTS and RVLM, was not shown to be up regulated [165].

AT₁ (AB18801) and AT₂ (AB19134) immunoreactivity in the substantia nigra (SN) colocalized with TH in neurons, GFAP in astrocytes and OX-6 and OX-42 in activated microglia [166–168]. Using different carboxy terminal directed AT₁ and AT₂ antibodies for Western blotting, it was shown that estrogen treatment of ovariectomized rats, which was protective against 6-hydroxydopamine induced neurotoxicity in the SN, decreased AT₁ and increased AT₂ expression in the SN [166]. Of note, no change in AT₁ receptor mRNA was observed [166]. These researchers also observed AT₁ and AT₂ immunoreactivity (Santa Cruz, SC-579 and SC-9040) in dopaminergic neurons, astrocytes and microglia in both monkey and human SN [169].

The dorsomedial hypothalamus (DMH), a brain region that exhibits high AT₁ receptor density [170], also displays AT₁ immunoreactivity using the AB-N27AP [147]. This brain region is associated with the cardiovascular manifestations of panic disorder and direct administration of an AT₁ receptor antagonist into the DMH blocks this component of the panic disorder in an animal model of panic disorder [147].

Giles et al., [67] using the 350–359 carboxy terminal peptide directed AT_{1a} antibody, observed strong AT₁ receptor immunoreactivity in numerous brain regions including the SFO, OVLT, MnPO, the parvocellular division of the PVN,

several other hypothalamic nuclei, and the NTS, corresponding well with radioligand binding and mRNA studies of the distribution of brain AT₁ receptors.

4.6. Perspective on the Use of Antibodies for the Study of Angiotensin Receptors. The ambiguity associated with studies of angiotensin receptors using different methods, whether by radioligand binding, receptor autoradiography, mRNA, local application of Ang II, electrophysiology, fos induction, or by immunoreactivity, necessitates considerable stringency in the analysis and interpretation of the data. Strengths of the immunohistochemical studies reported herein are as follows: (1) there is no known peptide sequence that closely mimics those used to generate these antibodies, (2) the antibodies were affinity purified to eliminate antibodies that did not recognize the antigenic peptide, (3) antibody binding is blocked by incubation with an excess of the antigenic peptide (preadsorption control), (4) Western blots indicate that the primary bands of labeled protein have molecular weights within the range of those previously observed for glycosylated, dimerized or chaperone protein linked angiotensin receptors [68, 171–175], and (5) the anatomical pattern of immunoreactivity correlates with radioligand binding for AT receptors [37, 59], agonist-induced c-fos expression [176], and the distribution of mRNA encoding the protein [44].

Weaknesses of this and other immunohistochemical approaches are as follows: (1) one cannot rule out the possibility that another protein could present an epitope similar to that recognized by these antibodies leading to a false positive, (2) there are posttranslational modifications of the receptor proteins that may mask the antigenic sites that they recognize, for example, phosphorylation of serine residues in Ang II receptors by a variety of protein kinases. The C-terminal domains chosen for generation of these antibodies contain several serines which when phosphorylated may mask the epitopes for the antibodies. AT₁ receptors are phosphorylated by G protein receptor kinase GRK2 (formerly known as β adrenergic receptor kinase, BARK1) leading to β -arrestin binding to the intracellular domain of the AT₁ receptors which may also mask the epitopes [177]. An additional post-translational modification is proteolytic cleavage of the receptor into smaller fragments following internalization. Cook et al. [178] demonstrated formation of a 54 amino acid carboxy terminal fragment of the rat AT_{1a} receptor that translocated to the nucleus and induced apoptosis in a variety of cell types. Thus it is possible that the immunoreactivity observed herein is not that of the full length receptor. (3) Receptors undergo protein-protein interactions such as receptor dimerization or interactions with chaperone proteins which have the potential to mask the antigenic site on the receptor; (4) inability to document the loss of immunological reactivity in an animal in which the receptor protein has been eliminated, for example, receptor knockouts. A recent publication [179] using Western blotting and immunofluorescence has challenged the specificity of 6 commercially available AT₁ receptor antibodies, including one previously questioned by Adams et al., [180] based upon the presence of immunoreactive material in mice in which

the AT_{1a} receptor is disrupted. The specificity of 3 AT₁ receptor antibodies, Alomone Labs #AAR-011, Santa Cruz sc-1173, and Abcam 18801, has also been challenged based upon expression of immunoreactivity in AT_{1a} and AT_{1b} knockout mice [181]. A generalized challenge to the ability of antibodies to selectively recognize G protein-coupled receptors (GPCR) based on apparent nonspecificity of 49 GPCR antibodies to 19 different GPCRs (the AT₁ and AT₂ receptors were not among the 19 GPCRs) has called into question the validity of immunological identification of GPCRs [182]. However, Xue et al., [183] using the same antibody as Adams et al., [180] demonstrated knockdown of AT₁ receptor immunoreactivity in the PVN. Of note, the AT_{1a} gene disruption [184] does not eliminate the carboxy terminal coding domain of the receptor that includes the peptide sequences used to generate several of those antibodies. If this portion of the receptor is still expressed it could explain the residual presence of AT_{1a} immunoreactive material in these knockout mice. However, the amino terminal sequence used to generate SC-1173 (amino acids 15–24) is in the deleted part; thus, it remains questionable whether the siRNA knockdown in the rat brain or the knockout of the mouse AT_{1a} receptor gives the correct information regarding the specificity of this and other AT₁ receptor antibodies.

One approach to resolve this question is to determine the identity of the protein in the band that the AT₁ receptor antibodies recognize in both wild-type and AT₁ receptor knockout mice. This has the potential to either (1) validate the immunological identification of AT₁ receptor protein thereby calling into question the efficacy of the AT₁ receptor knockout technology, (2) to discover a heretofore unknown subtype of the AT₁ receptor with an mRNA sequence that somehow evaded recognition by homology cloning approaches, (3) to identify (a) non-AT₁ protein(s) that colocalize(s) with AT₁ receptors and display (a) sufficiently similar epitope(s) as to be recognized by a variety of AT₁ receptor antibodies, (4) to discover (a) proteins with no relationship to AT₁ receptors that coincidentally express the same epitope(s) as the AT₁ receptor antibodies, or (5) to discover (a) novel protein(s) that has/have not yet been identified.

Until such questions are definitively answered, immunohistochemical studies, despite their known and potential limitations, can complement other types of analyses, which are also subject to a variety of differing limitations.

In conclusion, antibodies that can differentiate the 3 different angiotensin II receptor subtypes in the rat were used to immunohistochemically label angiotensin II receptor subtype-like immunoreactivity in the rat adrenal, pituitary, and brain. The pattern of staining corroborates mRNA, radioligand binding, and functional studies of adrenal and anterior pituitary angiotensin receptors. This indicates that AT_{1a} and AT₂ receptor subtypes occur in the zona glomerulosa and medulla of normal rats, the AT_{1b} subtype occurs only in the zona glomerulosa of normal rats while the AT_{1b} is the subtype predominantly expressed in the anterior pituitary. The localization of Ang II receptor immunoreactivity in the brain is in large part consistent with radioligand binding, mRNA, Ang II-induced fos expression, and functional studies; however, differences between these immunoreactivity

observations and observations obtained from some other techniques are yet to be resolved.

Disclosure

R. C. Speth has licensed these antibodies for commercial sale to Advanced Targeting Systems, Inc., San Diego, CA, USA (92121). The immunochemical studies conducted by M. Brownfield did not benefit ImmunoStar (i.e., they do not offer these antibodies).

Acknowledgments

The authors thank Drs. Kevin Grove, Kathryn Sandberg, Julia Cook, and Richard Re for assistance and helpful suggestions in the preparation of this paper. Funding for this work was provided by The Peptide Radioiodination Service Center, Washington State University, and the University of Wisconsin with a research gift from ImmunoStar Corporation.

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Research Article

Distinct Molecular Effects of Angiotensin II and Angiotensin III in Rat Astrocytes

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Received 29 November 2012; Revised 4 January 2013; Accepted 7 January 2013

Academic Editor: Marc de Gasparo

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It is postulated that central effects of angiotensin (Ang) II may be indirect due to rapid conversion to Ang III by aminopeptidase A (APA). Previously, we showed that Ang II and Ang III induced mitogen-activated protein (MAP) kinases ERK1/2 and stress-activated protein kinase/Jun-terminal kinases (SAPK/JNK) phosphorylation in cultured rat astrocytes. Most importantly, both peptides were equipotent in causing phosphorylation of these MAP kinases. In these studies, we used brainstem and cerebellum astrocytes to determine whether Ang II's phosphorylation of these MAP kinases is due to the conversion of the peptide to Ang III. We pretreated astrocytes with 10 μ M amastatin A or 100 μ M glutamate phosphonate, selective APA inhibitors, prior to stimulating with either Ang II or Ang III. Both peptides were equipotent in stimulating ERK1/2 and SAPK/JNK phosphorylation. The APA inhibitors failed to prevent Ang II- and Ang III-mediated phosphorylation of the MAP kinases. Further, pretreatment of astrocytes with the APA inhibitors did not affect Ang II- or Ang III-induced astrocyte growth. These findings suggest that both peptides directly induce phosphorylation of these MAP kinases as well as induce astrocyte growth. These studies establish both peptides as biologically active with similar intracellular and physiological effects.

1. Introduction

Mitogen-activated protein (MAP) kinases constitute a superfamily of serine/threonine protein kinases involved in the regulation of a number of intracellular pathways associated with cellular growth, apoptosis, cellular differentiation, transformation of cells, and vascular contraction [1–4]. We have shown that angiotensin (Ang) II via activation of AT₁ receptors increases the expression of MAP kinases in primary cultures of rat astrocytes [5–7]. ERK1/2 MAP kinases were shown to mediate Ang II-induced astrocyte growth and Ang II-induced c-Fos and c-Myc expression [5, 6, 8]. We have also established that Ang II induces the phosphorylation of stress-activated protein kinase/Jun-terminal kinase (SAPK/JNK) MAP kinases leading to cellular proliferation in cultured rat astrocytes, an effect that was also mediated by the AT₁ Ang receptors [7]. Our findings suggest that Ang II signals through these two different MAP kinase pathways in astrocytes.

More recently, we showed that Ang III also induces the phosphorylation of ERK1/2 and SAPK/JNK MAP kinases in these cells [9, 10]. Moreover, Ang III was equipotent to Ang II in causing these MAP kinases phosphorylation and occurred via interaction with the Ang AT₁ receptor. Ang III also induced astrocyte growth, however, not to a similar extent as Ang II [9].

Similar to our intracellular findings, *in vivo* studies have established that both peptides have similar physiologically relevant effects. For example, intracerebroventricular (ICV) injection of Ang II or Ang III caused a similar dose-dependent increase in blood pressure. Since Ang II is quickly cleaved by aminopeptidase A (APA) into Ang III, the true effector was unknown [11, 12]. In spontaneously hypertensive rats (SHR), injection of both peptides caused a prolonged blood pressure response compared to controls. However, pretreatment with bestatin, an aminopeptidase B (APB) inhibitor, potentiated and prolonged the elevated blood pressure response to Ang III in SHR [13]. Since bestatin

inhibits the breakdown of Ang III, these findings suggest that Ang III was a key player in the blood pressure response elicited by the activation of the renin angiotensin system.

Findings from several studies have implicated Ang III as the active peptide in the central nervous system. Reaux et al. [14] showed that preventing Ang II conversion to Ang III using EC33 (a selective APA inhibitor) blocked the pressor response of exogenous Ang II in rats, suggesting that the conversion of Ang II to Ang III is required to increase blood pressure in these animals. Moreover, ICV injection of EC33 caused a dose-dependent decrease in blood pressure, while ICV injection of PC18 (an APB inhibitor) increased blood pressure, an effect that was prevented by pretreatment with the AT₁ receptor antagonist, losartan [14]. These findings and ours suggest that Ang III is a major effector peptide of the brain renin angiotensin system, and this peptide is involved in central control of blood pressure. In addition, these findings and others implicate Ang III, not Ang II, as the active peptide of the central renin angiotensin system leading to the putative "Ang III hypothesis." Therefore, in this study, we used specific APA inhibitors to establish whether Ang II-mediated phosphorylation of ERK1/2 and SAPK/JNK MAP kinases in astrocytes was a result of Ang II conversion to Ang III. This is highly possible since APA is expressed in astrocytes [15] and neurons [16, 17]. As shown by others, inhibition of this enzyme has attenuated certain effects of Ang II. These studies were conducted in brainstem and cerebellum astrocytes to allow correlation of the current results with our previous findings [9, 10]. Moreover, astrocytes are the main source of brain angiotensinogen [18], the Ang II and Ang III precursors, and thus are ideal brain-derived cells to study the signaling pathways of these peptides.

2. Materials and Methods

2.1. Materials. Tissue culture supplies such as Dulbecco's Modified Eagles Medium (DMEM)/F12 (1:1), fetal bovine serum (FBS), antibiotic solution, and trypsin/EDTA were purchased from VWR (Grand Island, NY, USA). Ang II and Ang III were obtained from Bachem (Torrance, CA, USA). The APA inhibitor glutamate phosphonate (4-amino-4-phosphonobutyric acid, GluP) was generously supplied by Dr. Robert Speth (Nova Southeastern University, FL, USA), while amastatin (AMA) was purchased from Calbiochem (La Jolla, CA, USA). ³H-Thymidine (2000 Ci/mole) was purchased from MP Biomedicals (Solon, OH, USA). The phosphospecific ERK1/2 antibody, the phosphospecific SAPK/JNK antibody (Tyr751), the ERK1/2 antibody, and the SAPK/JNK antibody were purchased from Cell Signaling Technology (Beverly, MA, USA). Protein measurement supplies, gel electrophoresis, and Western blotting supplies including BCA protein reagents, acrylamide, ECL chemiluminescent reagents, and nitrocellulose membrane were purchased from either GE Health Care (Piscataway, NJ, USA) or Biorad Laboratories (Hercules, CA, USA) or from Pierce Biotechnology (Rockford, IL, USA). All other chemicals were purchased from either VWR international (Suwanee, GA, USA) or Sigma (St. Louis, MO, USA).

2.2. Preparation of Astrocytes. Timed, pregnant Sprague-Dawley rats were obtained from Charles River Laboratories (Wilmington, MA, USA) and maintained in the ALAAC-accredited animal facility of Nova Southeastern University. Primary cultures of astrocytes were prepared from the brainstem and cerebellum of 2-3-day-old neonatal pups by physical dissociation as previously described [6, 8, 19]. Cells were maintained in DMEM/F12 with 10% FBS, 100 µg/mL penicillin, and 100 units/mL streptomycin at 37°C in a humidified CO₂ incubator (5% CO₂ and 95% air). Cultures were fed every 3 to 4 days until confluent. Confluent monolayers were placed in DMEM/F12 containing 10 mM HEPES, pH 7.5, 10% FBS, and antibiotics and shaken overnight to remove oligodendrocytes. Astrocytes were detached with trypsin/EDTA (0.05% trypsin, 0.53 mM EDTA), replated at a ratio of 1 to 10, and grown to about 85% confluence prior to use. Isolated cells showed a positive immunoreactivity with an antibody against glial fibrillary acidic protein and negative immunoreactivity with markers for neurons, fibroblasts, or oligodendrocytes.

2.3. Cell Treatments. Cultured brainstem and cerebellum astrocytes growing on 100 mm culture dishes were made quiescent by a 48-hour treatment with serum-free media. Astrocytes were pretreated with either 100 µM GluP or 10 µM AMA for 15 minutes. Following the pretreatments, the cells were subsequently stimulated with 100 nM Ang II or 100 nM Ang III for 10 minutes. For comparative purposes, cells were also incubated with the inhibitors alone. Basal and stimulated levels of the MAP kinases were determined in the presence of DMSO in experiments involving AMA. The concentrations of the GluP and AMA were not cytotoxic based on measurements of the mitochondrial uptake of the tetrazolium dye, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), to its insoluble formazan form (data not shown). Cell lysates from all experiments were prepared by washing the astrocyte monolayers with phosphate-buffered saline containing 0.01 mM NaVO₄ to prevent the dephosphorylation of activated phosphorylated proteins. Cells were solubilized in supplemented lysis buffer (100 mM NaCl, 50 mM NaF, 5 mM EDTA, 1% Triton X-100, 50 mM Tris-HCl, 0.01 mM NaVO₄, 0.1 mM PMSF, 0.6 µM leupeptin, and pH 7.4) for 30 minutes on ice. The supernatants were clarified by centrifugation (12000 ×g for 10 minutes, 4°C), and the protein concentrations were subsequently measured using the BCA method.

2.4. Western Blot Analysis. Western blotting protocols to determine ERK1/2 phosphorylation and JNK phosphorylation were previously described [7, 10]. Essentially, solubilized proteins were separated in 10% polyacrylamide gels and transferred to nitrocellulose membranes. Nonspecific binding to the membranes was prevented by treating the membranes with 5% Blotto and probing with either the specific activated phosphorylated form of the ERK1/2 antibody or the activated phosphorylated form of the SAPK/JNK antibody. After incubating with the primary antibodies, the membranes were probed with goat anti-rabbit antibody coupled to horseradish peroxidase, the immunoreactive bands

visualized using ECL reagents, and the data quantified by densitometry. To quantify protein loading, membranes were also probed with antibodies for the ERK1/2 protein and SAPK/JNK protein. The immunoreactive bands were visualized using ECL reagents and quantified by densitometry.

2.5. Measurement of DNA Synthesis. Subconfluent monolayers of cells growing in 24-well culture dishes were made quiescent by 48-hours treatment with serum-free media. Individual wells were then pretreated for 15 minutes with 100 μ M GluP or 10 μ M AMA. Subsequently, the cells were treated with either 100 nM Ang II or 100 nM Ang III for 48 hours. For comparative purposes, some wells did not receive any peptides or were treated with DMSO (the vehicle for AMA) or were only treated with the inhibitors alone. 3 H-Thymidine (0.25 Ci/mL culture medium) was added during the last 24 hours of treatment. Newly synthesized DNA was precipitated with 5% TCA, dissolved in 0.25 N NaOH, and quantified by liquid scintillation spectrometry as previously described [20].

2.6. Statistical Analysis. All data are expressed as the mean \pm SEM of 6 or more experiments, as indicated. *t*-tests or repeated measures of one-way analysis of variance (ANOVA) with Dunnett's posttest were used to compare treatment groups with groups treated with no chemicals or those treated with the inhibitors, using PRISM (GraphPad). The criterion for statistical significance was set at $P < 0.05$.

3. Results

3.1. Effect of APA Inhibitors on ERK1/2 MAP Kinase Phosphorylation by the Peptides. In previous studies, we showed that Ang II and Ang III interact with Ang AT₁ receptors to significantly increase ERK1/2 MAP kinase phosphorylation to a similar extent [10]. To determine whether the effects observed with Ang II were due to its conversion to Ang III, cerebellar and brainstem astrocytes were pretreated for 15 minutes with 10 μ M AMA or 100 μ M GluP. The cells were subsequently stimulated with either 100 nM Ang II or 100 nM Ang III. As shown in Figure 1, both Ang II and Ang III significantly and equipotently induced ERK1/2 MAP kinase phosphorylation in brainstem astrocytes. The APA inhibitors AMA (Figure 1(a)) and GluP (Figure 1(b)) were ineffective in preventing Ang II- and Ang III-induced ERK1/2 phosphorylation. Similarly, in cerebellar astrocytes, both peptides induced ERK1/2 MAP kinase phosphorylation, effects that were not affected by pretreatment with the APA inhibitors (Figures 1(c) and 1(d)). These findings suggest that the two peptides directly induced phosphorylation of this MAP kinase pathway.

3.2. Effect of APA Inhibitors on SAPK/JNK MAP Kinase Phosphorylation by the Peptides. As shown previously [9] and in Figure 2, Ang II and Ang III via interactions with the Ang AT₁ receptor induced SAPK/JNK MAP kinase phosphorylation similarly in brainstem astrocytes. Pretreatment with the APA inhibitors AMA (Figure 2(a)) and GluP (Figure 2(b)) failed to

prevent Ang II-mediated and Ang III-mediated SAPK/JNK MAP kinase phosphorylation. In cerebellar astrocytes, the two peptides were equipotent in inducing SAPK/JNK MAP kinase phosphorylation (Figures 2(c) and 2(d)). Both APA inhibitors were unable to prevent Ang II-induced and Ang III-induced SAPK/JNK MAP kinase phosphorylation in cerebellar astrocytes as well (Figures 2(c) and 2(d)). Interestingly, brainstem astrocytes were more sensitive to the effects of both peptides, exhibiting higher levels of SAPK/JNK phosphorylation as compared to cerebellum astrocytes. In addition, our findings also suggest that both peptides have direct effects to induce SAPK/JNK MAP kinase in brainstem and cerebellum astrocytes.

3.3. Effects of APA Inhibitors on Ang Peptide Astrocyte DNA Synthesis. Both Ang II and Ang III have been shown to induce cellular proliferation of astrocytes [10]. To determine whether inhibition of APA affected the mitotic effects of Ang II and Ang III, brainstem and cerebellar astrocytes were pretreated with the APA inhibitors (10 μ M AMA and 100 μ M GluP) followed by stimulation with the peptides. As shown in Figure 3, both Ang II and Ang III induced cellular proliferation of brainstem and cerebellum astrocytes. However, Ang II was more potent at inducing astrocyte proliferation in both brainstem and cerebellum astrocytes, a finding that we observed previously [9, 10]. The APA inhibitors AMA and GluP (Figures 3(a) and 3(b)) were ineffective at preventing the proliferative effects of Ang II and Ang III, suggesting that both peptides had direct effects on cell proliferation.

4. Discussion

We and others have established that the angiotensin peptides Ang II and Ang III have important cellular effects in the central nervous system. However, in the intact brain, controversy still surrounds the synthesis, degradation, and physiological effects of Ang II, the primary peptide produced by the renin angiotensin system. One such controversy is known as the "Ang III Hypothesis." The premise of this theory is that the physiologically relevant peptide in the brain that binds to and activates the AT₁ receptor is Ang III, not Ang II. This putative "Ang III Hypothesis" suggests that Ang II is converted to Ang III in order to activate Ang AT₁ receptors [14, 21–25]. Studies have shown that centrally produced Ang II is rapidly degraded by several enzymes (primarily APA), leading to the accumulation of Ang III [26]. It is the degradation of Ang II to Ang III by APA that is the most documented breakdown pathway for Ang II and is the focus of most studies. Since the metabolism of Ang II is relatively rapid, this has helped to fuel the Ang III Hypothesis.

There are a few studies that support the putative Ang III hypothesis. Most of these studies have focused on using APA inhibitors such as AMA to block Ang II conversion to Ang III, then measuring effects of the initial Ang II treatment on blood pressures of normotensive and hypertensive animals [11–14]. A few studies have also focused on the Ang receptor(s) that Ang III interacts with to cause effects on

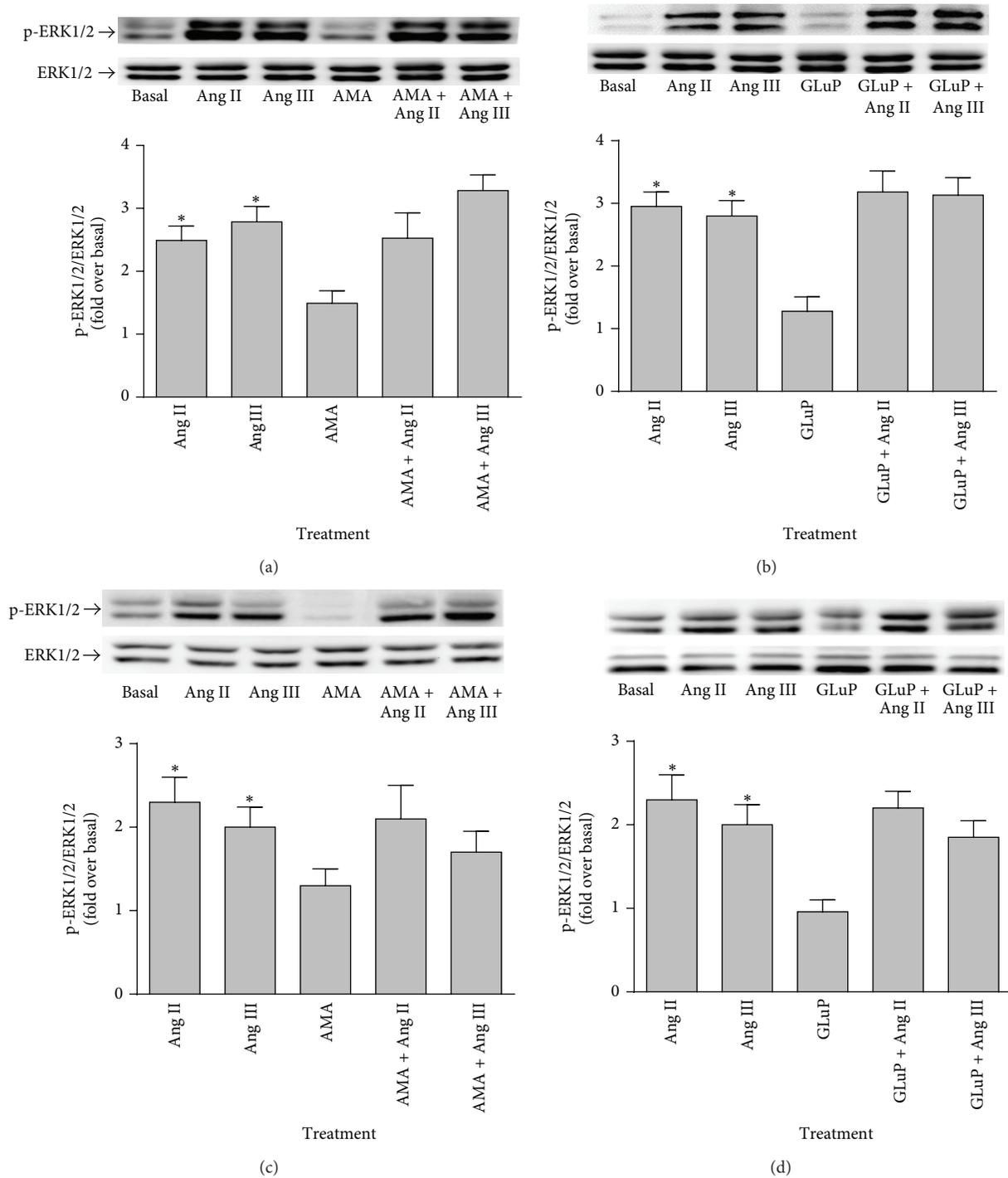


FIGURE 1: Effect of AMA and GluP on Ang II- and Ang III-induced ERK1/2 protein phosphorylation. Quiescent monolayers of brainstem ((a) and (b)) or cerebellum ((c) and (d)) astrocytes were pretreated with 10 μ M AMA or 100 μ M GluP for 15 minutes. The cells were subsequently stimulated with 100 nM Ang II or 100 nM Ang III for 10 minutes. Phosphorylated-ERK1/2 immunoreactive protein levels were measured by Western blot analysis using an antibody specific for the phosphorylated form of ERK1/2. Protein loading was quantified using the ERK1/2 protein antibody. The data were analyzed by densitometry, and the amount of phosphorylation was calculated as the fold increase over basal in the presence of vehicle. Each value represents the mean \pm SEM of preparations of brainstem and cerebellum astrocytes from 6 or more litters of neonatal rat pups. * denotes that $P < 0.05$ as compared to basal levels for ERK1/2 expression in astrocytes prepared from the brainstem and the cerebellum.

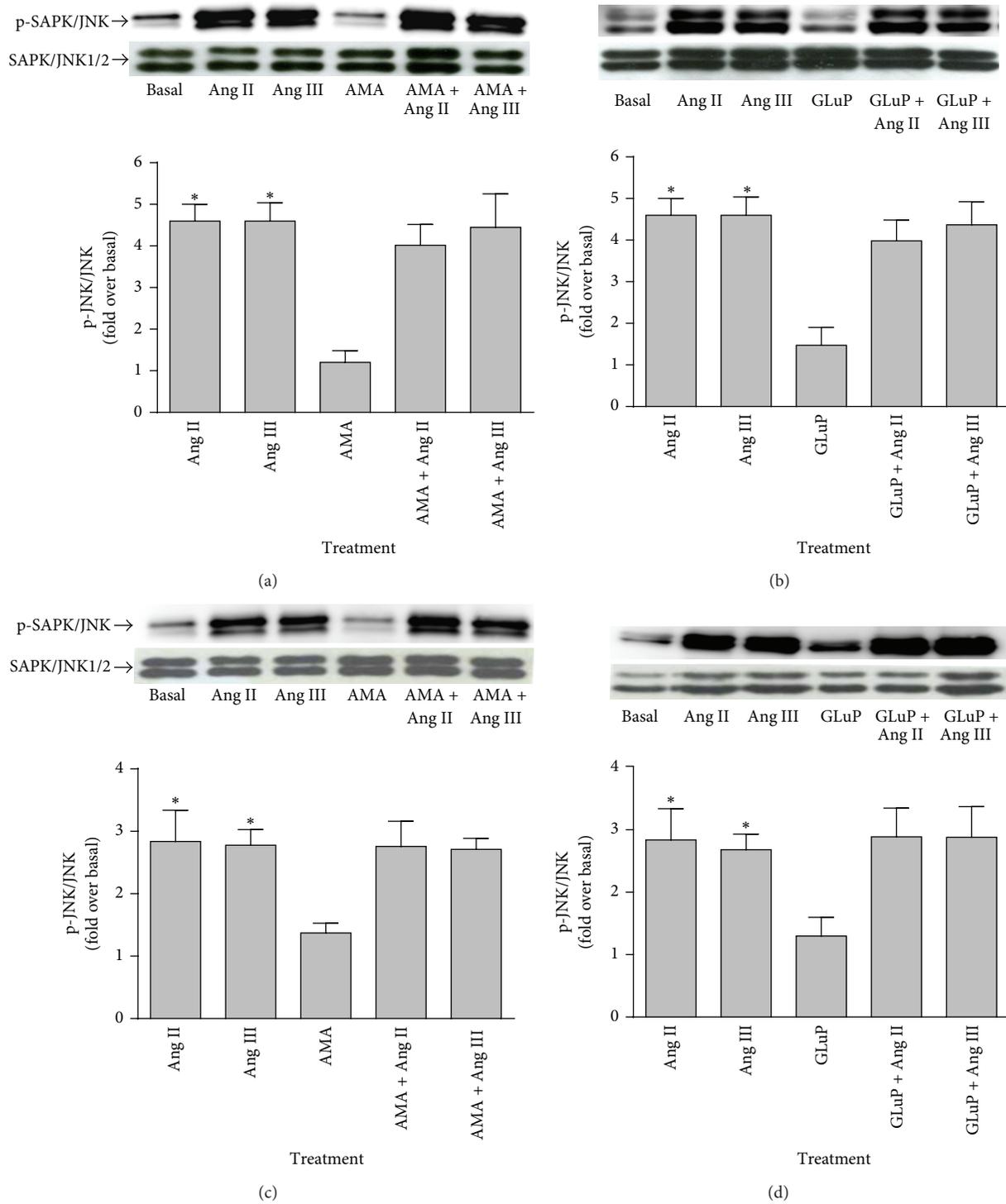


FIGURE 2: Effect of AMA and GluP on Ang II- and Ang III-induced SAPK/JNK protein phosphorylation. Quiescent monolayers of brainstem ((a) and (b)) and cerebellum ((c) and (d)) astrocytes were pretreated with 10 μ M AMA or 100 μ M GluP for 15 minutes. The cells were subsequently stimulated with 100 nM Ang II or 100 nM Ang III for 10 minutes. Phosphorylated-SAPK/JNK immunoreactive protein levels were measured by Western blot analysis using an antibody specific for the phosphorylated form of SAPK/JNK. Protein loading was quantified using the SAPK/JNK protein antibody. The data were analyzed by densitometry, and the amount of phosphorylation was calculated as the fold increase over basal in the presence of vehicle. Each value represents the mean \pm SEM of preparations of brainstem and cerebellum astrocytes from 6 or more litters of neonatal rat pups. * denotes that $P < 0.05$ as compared to basal levels for SAPK/JNK expression in astrocytes prepared from the brainstem and the cerebellum.

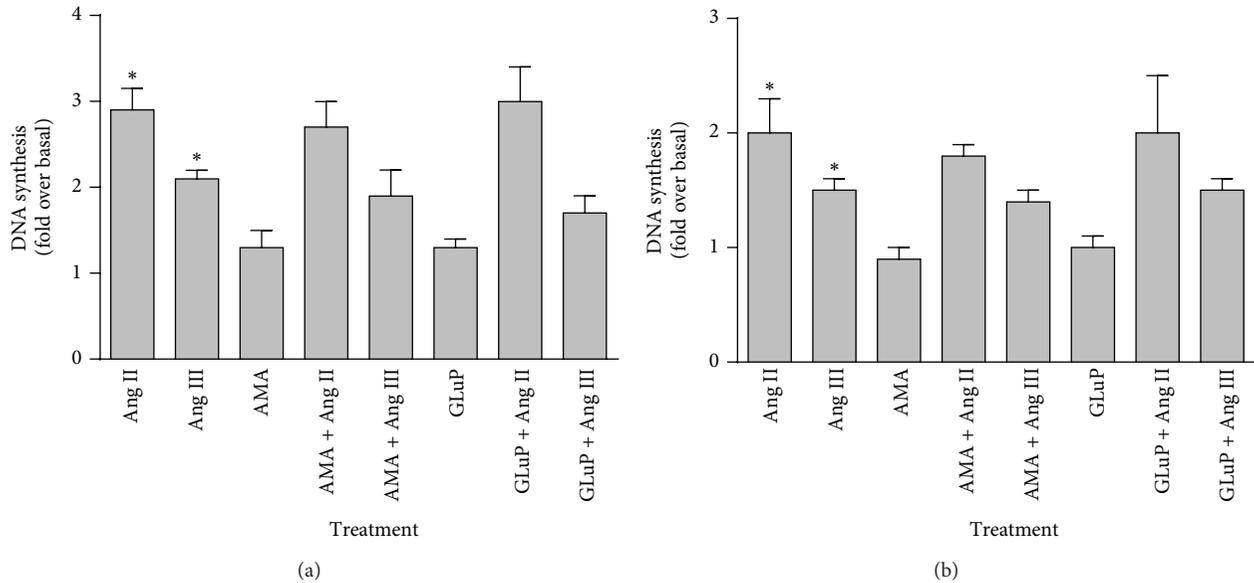


FIGURE 3: Effect of AMA and GluP on Ang II- and Ang III-induced DNA synthesis. Quiescent monolayers of brainstem (a) and cerebellum (b) astrocytes were incubated for 48 hours with 100 nM Ang II or Ang III in the presence and absence of 10 μ M AMA or 100 μ M GluP. During the last 24 hours of treatment, 3 H-Thymidine was added and DNA synthesis measured as described. Cells were also treated with the inhibitors alone. Each value represents the mean \pm SEM of preparations of brainstem and cerebellum astrocytes from 6 or more litters of neonatal rats pups. *denotes that $P < 0.05$ as compared to basal levels of DNA synthesis.

blood pressure. An *in vitro* study by Harding et al. [22] showed that Ang III was more potent than Ang II at eliciting responses from paraventricular nucleus receptors isolated from normal rats and SHR. The Ang II response was blocked by the addition of AMA, the selective APA inhibitor, and was ineffective at preventing the Ang III response [22].

There are a number of studies that refute the theory that Ang II conversion to Ang III is necessary for AT₁ receptor binding. Kokje et al. [27] showed that aminopeptidase-resistant Ang II analogs, when ICV-administered, caused pressor and dipsogenic activities similar to or greater than Ang II. In addition, there were no differences in Ang II, its aminopeptidase-resistant analogs, and Ang III initial peak pressor responses, findings that do not support a requisite formation of Ang III to elicit a response. It has also been established that 125 I-Ang II can bind to brain AT₁ receptors, an effect that does not require conversion to 125 I-Ang III [28].

In the current studies, we sought to determine whether some of the effects that we have previously observed with Ang II are modulated in the presence of the APA inhibitors, AMA and GluP. AMA was selected since it is an APA inhibitor that is used by many investigators to study Ang II metabolism [26]. GluP was selected since it has recently received much attention as a direct APA inhibitor with a better selectivity for APA [28]. This enzyme was selected as a target for inhibition since it is the major enzyme involved in Ang II conversion to Ang III [26, 29].

In these studies, we showed that Ang III and Ang II had similar effects to induce ERK1/2 and SAPK/JNK MAP kinases phosphorylation, effects that we have previously observed [9, 10]. Blocking the ability of APA to convert Ang II to

Ang III did not significantly affect the ability of Ang II to induce phosphorylation of these MAP kinases. In addition, both Ang II and Ang III induced astrocyte proliferation although Ang II was more potent, an effect we have observed previously. Further, blocking the activity of APA failed to prevent the proliferative effects of both peptides. Overall, our findings suggest that Ang II has direct effects to stimulate MAP kinases and to induce astrocyte growth since preventing its conversion to Ang III did not affect the ability of the peptide to induce MAP kinase phosphorylation or its ability to cause astrocyte proliferation. These findings fail to support the Ang III hypothesis and suggest that in astrocytes, Ang II has agonistic and mitogenic properties of its own. Most importantly, these studies also support a role for Ang III as a central peptide.

Ang III agonistic properties were observed a number of years ago [30]; however, the relevance of this peptide as a key player in the renin angiotensin system is not yet appreciated. In astrocytes, our studies support mitogenic properties of Ang III since it directly activates ERK1/2 and SAPK/JNK MAP kinases and causes astrocyte growth [9, 10]. Most importantly, Ang III-induced astrocyte proliferation occurred via the MAP kinase pathways suggesting an important role for these pathways in Ang III effects [9, 10]. It is notable that in most instances, Ang III agonistic effects in the brain are similar to Ang II. Both peptides have been shown to induce similar central-mediated increases in blood pressure or mean arterial pressure [30, 31] and equivalent central effects on thirst and sodium appetite [32]. Centrally administered Ang III was as potent as Ang II in causing pressor and renal effects in rats on normal and high sodium

diets [33]. Ang II and Ang III induced the central expression of the transcription factors c-Fos, c-Jun, and Krox-24 with the same efficacy [34]. Peripherally, Ang III was equipotent to Ang II with respect to blood pressure increases, aldosterone secretion, and renal functions [35]. On the other hand, we have shown that the ability of Ang III to cause proliferation of astrocytes was differentiated from Ang II, exhibiting a similar effect as Ang II at lower doses. At the higher doses of the peptides, astrocytes were more responsive to Ang II [10]. Thus, our studies and others have shown that Ang II and Ang III may have equivalent effects on several physiological processes but differential effects on others. Overall, our findings suggest a relevant physiological role of both Ang II and Ang III in the body, in particular in the brain.

It has also been suggested that the APA inhibitors may be acting as agonist/antagonist of the AT₁ receptor [26]. Thus, in the presence of the inhibitors, the AT₁ receptor responses would be modulated. However, in our hands, pretreating with the APA inhibitors had no effect on both Ang II and Ang III to induce the MAP kinases phosphorylation, and they did not affect the proliferative effects of both peptides. Since both Ang II and Ang III responses are mediated by interaction with the AT₁ receptor, these inhibitors are apparently not acting in an agonist/antagonist capacity in our system. The specificity and the efficacy of the APA inhibitors used in this study may be questionable and/or controversial. However, our findings are credible since most of the studies that are used to support the Ang III Hypothesis use an APA inhibitor (mostly AMA) to prevent Ang II conversion to Ang III [14, 22], suggesting that the results of our current study are directly comparable to others.

Even though a central renin angiotensin system is widely accepted, there are still few controversies that surround its existence, a major one being the Ang III Hypothesis. Our findings suggest that in brainstem and cerebellum astrocytes, both Ang II and Ang III are direct potent central mitogens and that their activities are unaffected by APA inhibitors. Thus, *in vitro*, these findings suggest that both peptides are important modulators of the actions of the brain renin angiotensin system. The relevance of these findings must be corroborated *in vivo* since cell cultures cannot give an overall picture of all actions of these peptides in the brain. It is important that we establish the major peptides that govern the actions of the central renin angiotensin system due to the significance of this system in many physiological and disease processes. Thus, our findings are essential in establishing the relevance of both Ang II and Ang III as direct acting endogenous central peptides.

Conflict of Interests

The authors declare that they have no conflict of interests.

Acknowledgments

The authors would like to thank graduate student Mr. Dhanush Haspula for his excellent technical assistance. This work was supported by a President's Faculty Research and

Development Grant no. 335465 from Nova Southeastern University.

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Research Article

The Brain-Heart Connection: Frontal Cortex and Left Ventricle Angiotensinase Activities in Control and Captopril-Treated Hypertensive Rats—A Bilateral Study

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Received 20 October 2012; Accepted 4 January 2013

Academic Editor: Patrick Vanderheyden

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The model of *neurovisceral integration* suggests that the frontal cortex (FC) and the cardiovascular function are reciprocally and asymmetrically connected. We analyzed several angiotensinase activities in the heart left ventricle (VT) of control and captopril-treated SHR, and we search for a relationship between these activities and those determined in the left and right FC. Captopril was administered in drinking water for 4 weeks. Samples from the left VT and from the left and right FC were obtained. Soluble and membrane-bound enzymatic activities were measured fluorometrically using arylamides as substrates. The weight of heart significantly decreased after treatment with captopril, mainly, due to the reduction of the left VT weight. In the VT, no differences for soluble activities were observed between control and treated SHR. In contrast, a generalized significant reduction was observed for membrane-bound activities. The most significant correlations between FC and VT were observed in the right FC of the captopril-treated group. The other correlations, right FC versus VT and left FC versus VT in controls and left FC versus VT in the captopril group, were few and low. These results confirm that the connection between FC and cardiovascular system is asymmetrically organized.

1. Introduction

Frontal cortex (FC) and cardiovascular functions are reciprocally connected, as part of the model of *neurovisceral integration* [1]. This connection is asymmetric [2] and a neurochemical substrate may underlie this lateralization [3]. Compared with vehicle-treated spontaneously hypertensive rats (SHRs), we recently reported an inverted bilateral behavior of angiotensinase activities between left/right FC and plasma after captopril treatment. The asymmetries between left and right FC markedly increased compared to the control group. We suggested that these results might reflect a systematized lateralized neuroendocrine response between brain and cardiovascular functions involving the autonomic nervous system [4]. There are evidences suggesting that the hyperactivity of the sympathetic nervous system is involved in the cardiac pathologies related to neurological accidents

such as cerebral infarction or head traumas, contributing to their high mortality rates [5]. Similarly, it has been also proposed that the autonomic imbalance in which the sympathetic nervous system predominates over the parasympathetic may be the pathway that connects impaired cognitive processes involving frontal cortex functions and altered heart functions [6, 7]. In addition, it was reported that unilateral prefrontal cortex lesions can alter emotional and cardiovascular autonomic responses, depending on which hemisphere was injured: there was a predominant parasympathetic inhibition by the left prefrontal cortex but a sympathetic activation by the right prefrontal cortex [8]. Aspartyl- (AspAP), glutamyl- (GluAP), alanyl- (AlaAP), and cystinyl-aminopeptidase (CysAP) are aminopeptidases (AP) involved in the metabolism of angiotensin peptides [9]. Based on these evidences of neuroendocrine correlations between brain and cardiovascular function and on our previous report showing

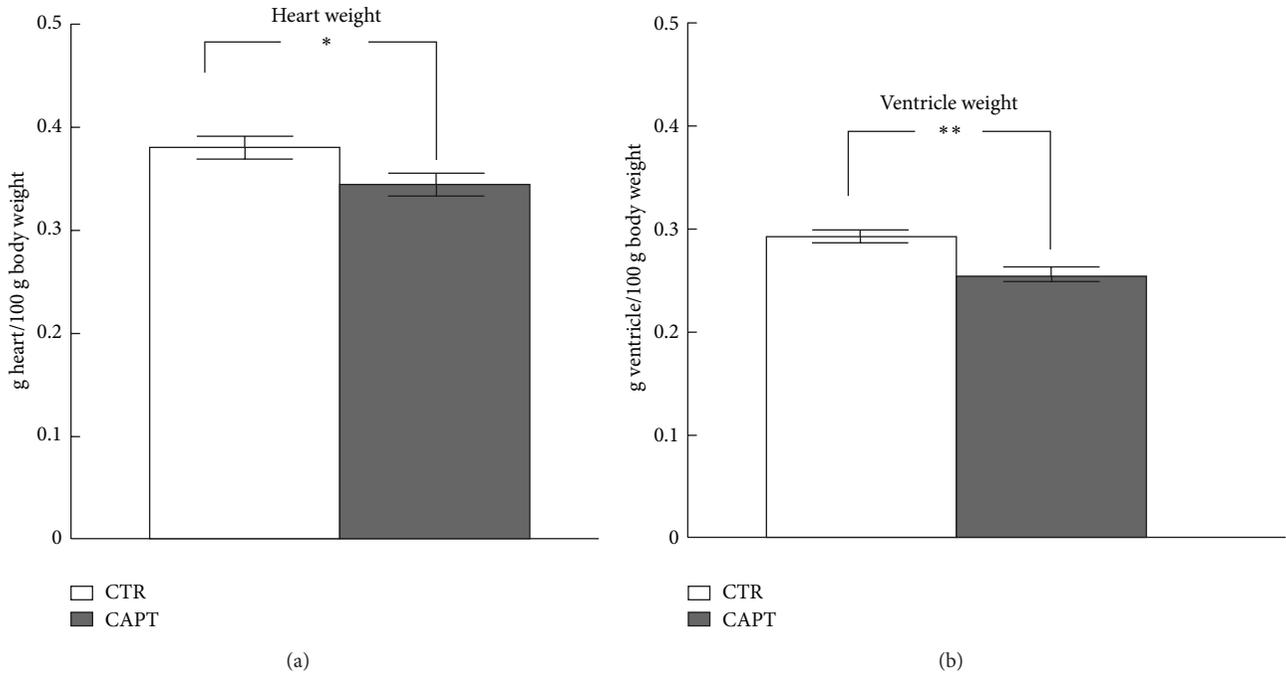


FIGURE 1: Total heart and left ventricle weight (g) in control (CTR) and captopril- (CAPT-) treated animals, as measured at the end of the treatment period (four weeks). The values represent the mean \pm SEM of 10 animals in each group. * $P < 0.05$; ** $P < 0.01$.

an asymmetrical effect of captopril between FC and plasma [4], it is therefore essential to analyze those angiotensinase activities in the heart ventricle (VT) of control and captopril-treated hypertensive rats and to search for a possible relationship between these activities in VT and the same determined in the left and right FC.

2. Material and Methods

All of the experimental procedures involving animals were performed in accordance with the European Communities Council Directive 86/609/EEC and were approved by the Bioethics Committee of the University of Jaén. Twenty adult male SHRs were divided into control ($n = 10$) and captopril-treated ($n = 10$) groups. Captopril (100 mg/kg p.o.) was administered daily in drinking water (0.5 mL/100 mg body weight) for 4 weeks. The systolic blood pressure (SBP) was monitored by the plethysmographic method throughout the experimental period. At the end of the treatment period, after recording the SBP, each rat was perfused with saline under equithesin anesthesia, and the left and right FC and samples from the left VT were obtained as previously described [4, 10]. Briefly, the brain samples were dissected according to the stereotaxic atlas of Paxinos and Watson [11]. For each group, the left and right frontal lobes 11.20 mm anterior to the interaural line were collected separately [12]. In addition, the heart was removed and weighed and the left ventricle was immediately dissected and weighed and a left ventricular sample was obtained. Soluble (SOL) and membrane-bound (MB) Aspartyl- (AspAP), glutamyl- (GluAP), alanyl- (AlaAP), and

cystinyl-aminopeptidase (CysAP) activities were measured fluorometrically using acrylamides as substrates, as previously described [4]. Student's t -test was used to compare the data from control and captopril-treated SHRs, and the paired Student's t -test was used for left FC versus right FC comparisons [4]. The Pearson correlation coefficient of the left or right frontal cortex and plasma AP activities was computed using SPSS13.0 and STATA 90. P values below 0.05 were considered significant.

3. Results

The results of the present research are reported in Figures 1, 2, and 3 and in Table 1. The SBP of captopril-treated SHRs was 47 mm Hg (or 30%) lower than that of control rats ($P < 0.001$) [4]. The weight of total heart decreased significantly after captopril treatment ($P < 0.05$) mainly due to a reduction in the left ventricle weight ($P < 0.01$) (Figure 1). In a previous study [4], we observed that the asymmetries for MB activities markedly increased in frontal cortex after captopril treatment compared to the control group, whereas the bilateral pattern (left versus right differences) of SOL activities did not substantially change. There was a left predominance for GluAP but a right one for AlaAP and CysAP [4].

In the present study, no differences for SOL GluAP, AlaAP, and CysAP activities were observed between the control and treated SHRs in the ventricle (Figure 2). No detectable activity was measured for SOL AspAP. However, a generalized significant reduction ($P < 0.05$) was observed for all MB AP activities ($P < 0.05$) after captopril except for AspAP that

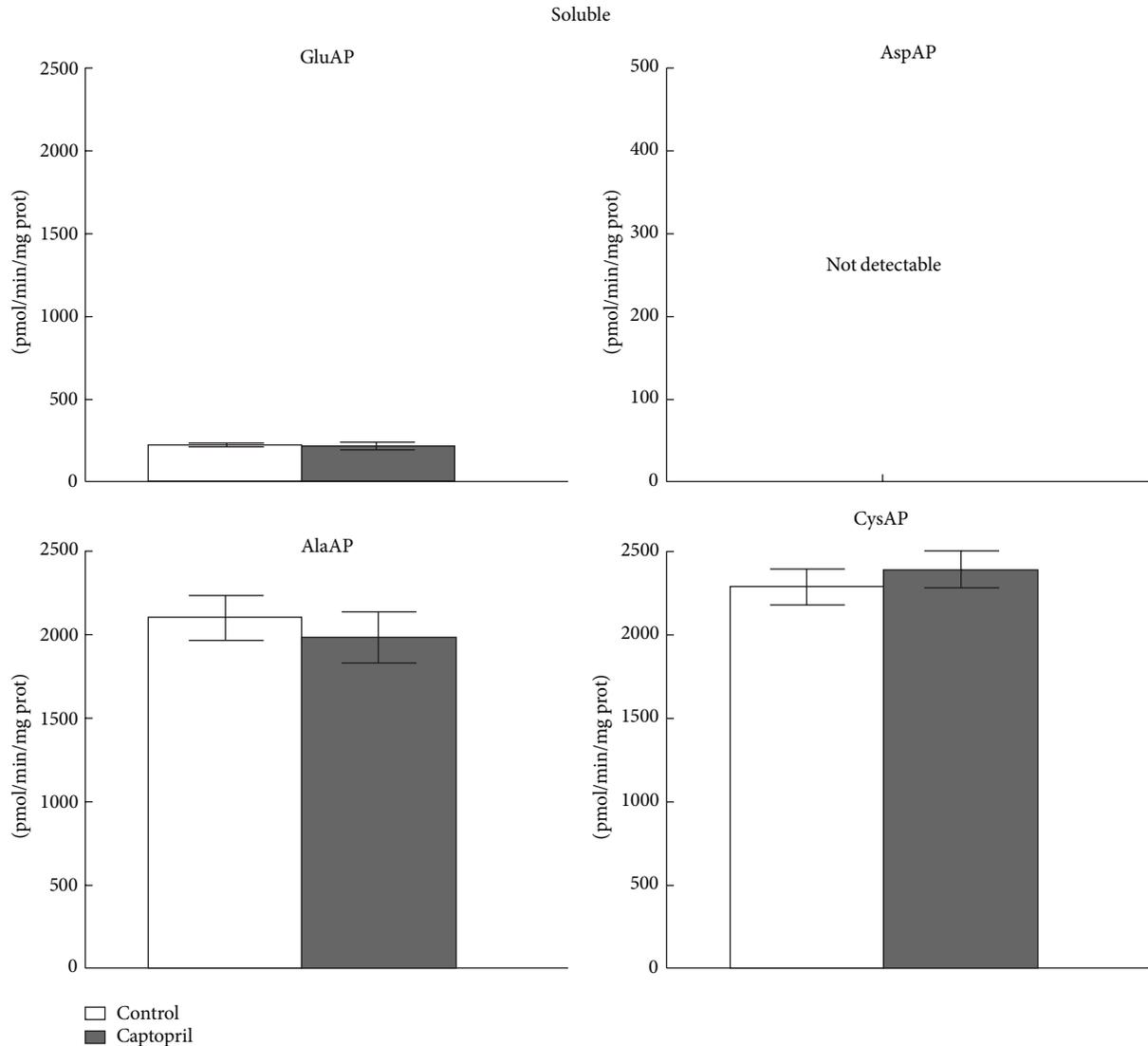


FIGURE 2: Soluble GluAP, AlaAP, and CysAP activities on the left ventricle of control ($n = 10$) and captopril-treated ($n = 10$) spontaneously hypertensive rats. The values represent the mean \pm SEM of specific GluAP, AlaAP, and CysAP activities expressed as picomoles of glutamyl-, alanyl- or cystinyl- β -naphthylamide hydrolyzed per min per mg of protein.

did not reach statistical significance versus the control group (Figure 3).

There are correlations between FC and VT: the majority and most significant correlations were observed with the right FC of the captopril-treated group. Interestingly, the correlations involving MB activities of the right FC were negative for GluAP (left FC predominant) and positive for AlaAP and CysAP (right FC predominant). The other correlations, right FC versus VT and left FC versus VT in controls and left FC versus VT in the captopril group, were few and low. Surprisingly, the opposite was previously observed between FC and plasma [4]. The plasma AP activities correlated significantly with those in the right FC in the control rats, whereas they correlated with activities in the left FC in the captopril-treated group.

4. Discussion

Our results demonstrated that captopril modified angiotensinase activities in heart and in brain, as previously reported [4], and that there is a significant correlation in the levels of these enzymatic activities between both organs.

Several components of a local cardiac RAS can result of an uptake from the circulation but a functional intracrine RAS in the heart also exists [13]. Moreover, ganglionic neurons in human heart express angiotensinogen and are able to generate Ang II [14]. As a whole, the local heart RAS might be involved in cellular hypertrophy and cardiac arrhythmias as well as in the regulation of heart cell volume through the action of Ang II on the AT_1 receptor [15]. In addition, the presence of Ang IV in the left ventricle was previously

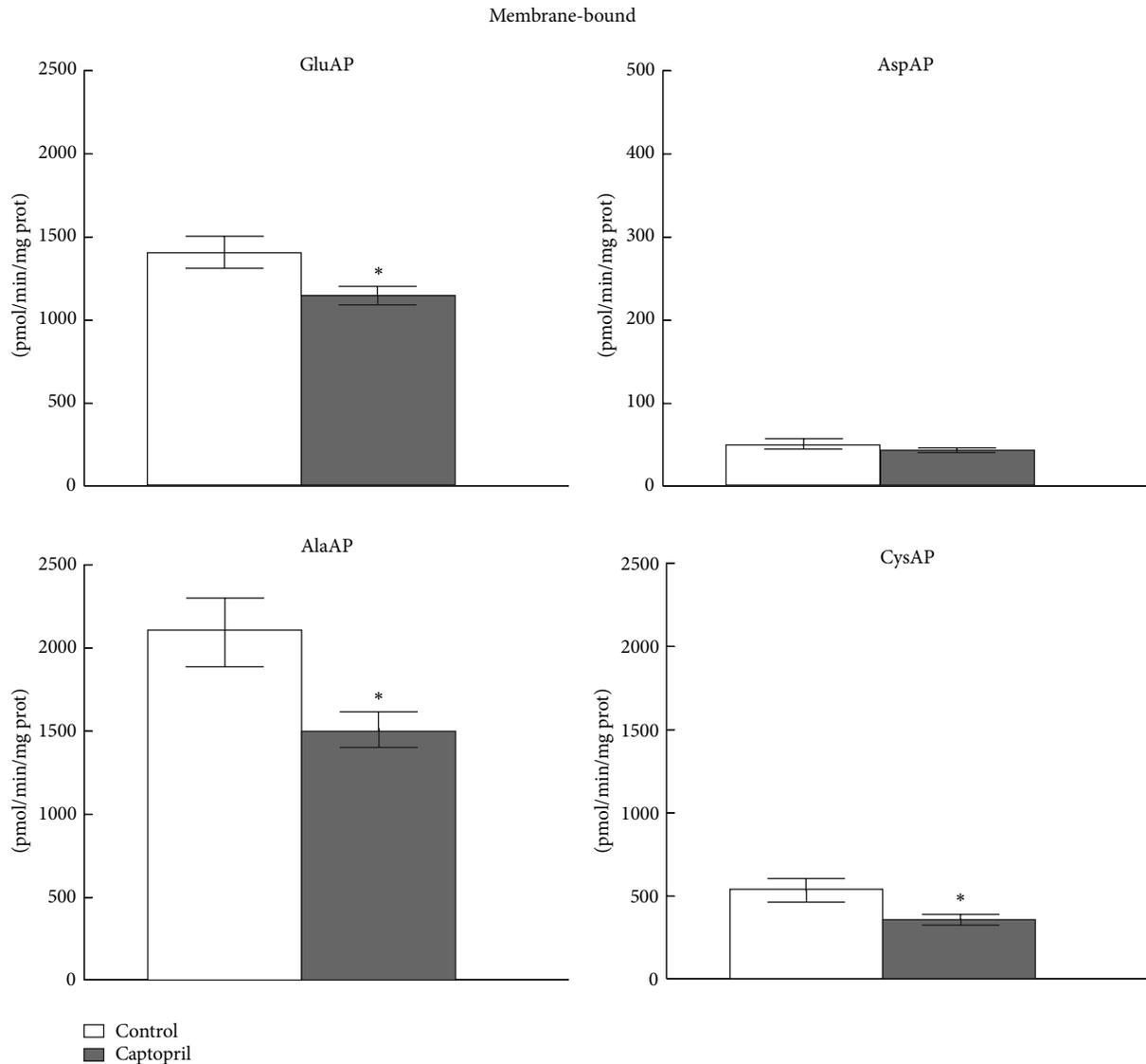


FIGURE 3: Membrane-bound GluAP, AspAP, AlaAP, and CysAP activities in the left ventricle of control ($n = 10$) and captopril-treated ($n = 10$) spontaneously hypertensive rats. The values represent the mean \pm SEM of specific GluAP, AspAP, AlaAP, and CysAP activities expressed as picomoles of glutamyl-, aspartyl-, alanyl-, or cystinyl- β -naphthylamide hydrolyzed per min per mg of protein. * $P < 0.05$.

described [16]. It was also observed that there was 10-fold more AT_4 receptor than AT_1 receptor in rabbit myocardium where they might exert opposite effects to those of Ang II through the binding of Ang II [16].

The present results of lower MB aminopeptidase activities in ventricle may indicate a local reduction in the metabolism of angiotensin peptides after treatment with captopril. More specifically, the results suggest a lower metabolism/higher availability of Ang II (metabolized to Ang III by GluAP), Ang III (metabolized to Ang IV by AlaAP), and Ang IV (also metabolized by AlaAP) in captopril-treated animals. On the other hand, Ang IV binds specifically to the AT_4 receptor, which was proposed to be identical to the insulin-regulated aminopeptidase (IRAP) [17]. Indeed, the high affinity-binding site for Ang IV is absent in IRAP-KO mice [18], and the brains of these mice are protected against

ischemic damage as Ang IV does in control animals [19]. Cystein aminopeptidase (CysAP), also called oxytocinase or vasopressinase (EC 3.4.11.3), is considered the human variant of IRAP [20]. These enzymes can therefore be considered identical. However, the identity of the AT_4 receptor remains in dispute, and its identification with IRAP is still controversial. Indeed, Ang IV has a very rapid effect on signaling molecules [21] at pico/nanomolar concentration, whereas the effect of Ang IV on enzyme inhibition such as on accumulation of endogenous IRAP substrates is slow. Also, the concentration of Ang IV producing a biological effect is below that needed to inhibit IRAP [22]. It was therefore proposed that the physiological action of Ang IV was mediated through the tyrosine-kinase cMet receptor whose function overlap with those of Ang IV/ AT_4 , that is, memory facilitation, cerebroprotection, seizure, neurite

TABLE 1: Frontal cortex versus ventricle.

Left frontal cortex			Right frontal cortex		
FC versus VT	<i>r</i>	<i>P</i>	FC versus VT	<i>r</i>	<i>P</i>
Control					
SOL GluAP versus MB CysAP	-0.733	0.03			
MB GluAP versus SOL AlaAP	+0.703	0.05		No correlations	
MB GluAP versus MB GluAP	+0.693	0.05			
Captopril					
MB GluAP versus MB CysAP	+0.766	0.02	SOL CysAP versus SOL AlaAP	+0.700	0.05
			SOL GluAP versus SOL AlaAP	+0.722	0.04
			SOL AlaAP versus MB CysAP	-0.749	0.03
			SOL GluAP versus MB CysAP	-0.781	0.02
			MB CysAP versus MB AlaAP	+0.692	0.05
			MB AlaAP versus MB AspAP	+0.805	0.02
			MB GluAP versus MB AspAP	-0.870	0.01
			MB CysAP versus MB CysAP	+0.695	0.05
			MB AlaAP versus MB GluAP	+0.742	0.03
			MB CysAP versus MB GluAP	+0.707	0.04

Correlations between the left or right frontal cortex (FC) soluble (SOL) or membrane-bound (MB) AP activities versus SOL or MB left ventricle (VT) AP activities in the control and captopril-treated animals. Pearson's correlation coefficients (*r*) and *P* values are indicated and specify the significance of the differences between these correlations.

outgrowth, cerebral blood flow, depression, and Parkinson's and Alzheimer's diseases [22].

Independently of the discussion on the real nature of the Ang IV binding site, it was proposed that the binding of Ang IV to its receptor results in the inhibition of the receptor's metabolic activity, reducing the catabolism of its substrates and consequently increasing their availability and extending their action [20]. Ang IV could therefore regulate glucose uptake modulating CysAP activity: CysAP is indeed colocalized with the glucose transporter GLUT4. In the presence of insulin, CysAP and GLUT4 are expressed in the plasma membrane, where GLUT4 induces glucose uptake [20]. Therefore, a decrease in CysAP activity as observed here would imply high levels of Ang IV and increased glucose uptake, which would also improve heart function [23].

These results are in agreement with previous reports indicating that the left ventricular expression of the glucose transporter GLUT4 is regulated by insulin at the transcriptional level [24] and that captopril increased myocardial oxygen and glucose uptake [25]. Therefore, considering these possibilities, the general reduction in the metabolism of Ang peptides in ventricle under captopril treatment might lead to a situation in which Ang peptides counteract each other resulting in an improvement of cardiovascular function.

As previously stated, brain asymmetry is not a static but rather a dynamic phenomenon in which both environmental and endogenous factors act as modulators [3]. In addition, this asymmetry extends to the neurovisceral integration through the asymmetrical functioning of the autonomic nervous system [4, 26–28]. For example, unilateral brain manipulations may not only produce alterations in brain bilaterality [27] but also have significant asymmetrical peripheral consequences in plasma and heart function [26, 28]. In addition, drug treatments may also influence brain bilaterality acting

directly and/or indirectly through mechanisms involving the neurovisceral integration model and the asymmetrical functioning of the autonomic nervous system [4]. This hypothesis may agree with the data observed by Chi et al. in 2003 [29] demonstrating that “several genes critical in the establishment of left-right asymmetry were expressed preferentially in venous endothelial cells, suggesting coordination between vascular differentiation and body plan development.” The authors highlighted the importance of the similarity of the migration paths of blood vessels and nerves during development, which reflects their mutual functional interaction [29].

It is particularly interesting that the present results are in contrast with those previously observed in plasma [4] where, in the control rats, the plasma AP activities correlated significantly with those in the right FC, whereas they correlated with those in the left FC in the captopril-treated group. These results strongly support that such responses were due to an asymmetry in the organization of the autonomic nervous system's innervations of the blood vessels and heart [4, 26, 28]. Remarkably, the peripheral autonomic innervation not only uses the classic autonomic neurotransmitters acetylcholine or norepinephrine but also other neuropeptides including angiotensin II that colocalize with them [30]. Thus, neuronal Ang II acts as a neuropeptide and synaptic cotransmitter in the periphery of the sympathetic nervous system [30]. In addition, the present results agree with the notion that essentially the right cortical activity regulates cardiovascular function [2]. Captopril influences heart [31] and brain [32] function and is a clear choice for the treatment of heart failure [33]. In addition, heart function [1] and heart failure [5] affect brain function, and, vice versa, left or right stroke differently influences the heart function [28]. Indeed, the stimulation of the left insular cortex in epileptic patients

before temporal lobectomy causes bradycardia and depressor responses, whereas the opposite happens after stimulation of the right cortex indicating a left parasympathetic and a right orthosympathetic dominance, respectively [34]. Cardiac autonomic derangements and arrhythmias have been mainly described in patients suffering right-sided stroke with insular involvement [35]. A reduced respiratory heart rate variability, a reflex mainly under parasympathetic control, was associated with increased mortality after right-sided stroke, suggesting that the risk of sudden death may be correlated with lateralization and the location of the brain infarct after stroke [36]. These and other evidences have led some authors to recommend a prolonged and intensive cardiovascular monitoring in patients manifesting right-sided stroke [37].

There is therefore a clear reciprocal interaction between heart and brain in physiological and pathological conditions [1, 5] that may be modulated by captopril as suggested by the present data. The brain RAS participates in the development of cardiac hypertrophy and fibrosis through the modulation of the autonomic nervous system. Furthermore, the inhibition of the sympathetic hyperactivity after myocardial infarction through suppression of the brain RAS appears to have a beneficial effect [38]. The blood-brain barrier permeability of captopril is negligible, but it has a marked effect on cerebral blood flow autoregulation when injected intravenously [39]. In addition, it causes a sympathetic inhibition in SHR after chronic administration suggesting that ACE inhibition may be protective for cerebral metabolism against ischemic insult [40]. Therefore, our present results suggest that captopril may centrally modulate the function of the autonomic nervous system leading to a higher connectivity between the right FC and the left ventricle. This could be particularly beneficial after a right-sided stroke.

The asymmetrical organization of the nervous system is the result of evolutionary adaptation [3, 41]. This is a dynamic phenomenon in which both environmental and endogenous physiological or pathologic factors act as modulators [3]. It has been hypothesized that the brain is intrinsically asymmetric. Virtually all brain functions are organized, more or less, in an asymmetrical fashion. It was speculated that imbalances in established brain asymmetries (toward symmetry or toward increasing asymmetry) might lead to neuropathological deviations in the functions of the nervous system, including the peripheral consequences due to changes in the modulation of the autonomic nervous system [3, 42, 43]. A deeper knowledge of how deviations in physiological lateralizations may lead to neuropathological consequences and how some exogenous factors may influence in physiological laterality, is important in order to be able to search for therapeutic tools that balance those deviations.

Captopril asymmetrically modifies angiotensinase activities in frontal cortex and in ventricle and, consequently, affects the functions they are involved in. However, these effects are not independent but according to our results, mutually interdependent. Therefore, but when appropriately characterized and systematized, the present results open new therapeutic consequences in cardiovascular treatment (presumably not only limited to specific drugs) that should

be taken into account in the design of protocols for the evaluation of cardiovascular treatments. The present study provides further molecular support for the involvement of the RAS in the brain frontal cortex-heart connection and adds, for the first time, biochemical evidence on the lateralized functioning of that connection. If this connection is asymmetric and the consequences of unilateral brain insults differ depending on the injured side, the necessity of characterizing appropriately the neurovisceral integration of the RAS is relevant and deserves further preclinical work.

Acknowledgment

This work is supported by the Ministry of Science and Innovation through Project no. SAF-2008 04685 C02 01.

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Review Article

Elevated Blood Pressure in the Acute Phase of Stroke and the Role of Angiotensin Receptor Blockers

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Received 7 October 2012; Revised 31 December 2012; Accepted 2 January 2013

Academic Editor: Marc de Gasparo

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Raised blood pressure (BP) is common after stroke but its causes, effects, and management still remain uncertain. We performed a systematic review of randomized controlled trials that investigated the effects of the angiotensin receptor blockers (ARBs) administered in the acute phase (≤ 72 hours) of stroke on death and dependency. Trials were identified from searching three electronic databases (Medline, Cochrane Library and Web of Science Database). Three trials involving 3728 patients were included. Significant difference in BP values between treatment and placebo was found in two studies. No effect of the treatment was seen on dependency, death and vascular events at one, three or six months; the cumulative mortality and the number of vascular events at 12 months differed significantly in favour of treatment in one small trial which stopped prematurely. Evidence raises doubts over the hypothesis of a specific effect of ARBs on short- and medium-term outcomes of stroke. It is not possible to rule out that different drugs might have different effects. Further trials are desirable to clarify whether current findings are generalizable or there are subgroups of patients or different approaches to BP management for which a treatment benefit can be obtained.

1. Introduction

Raised blood pressure (BP) is common after acute stroke, whether of ischaemic or haemorrhagic type. It exists in more than three quarters of patients, of which about half have a history of hypertension [1], and it declines spontaneously in two-thirds of cases returning to prestroke levels over the first week. Its decrease usually occurs 4–10 days after stroke, but in a significant percentage of patients it falls by about 25–30% just within the first 24 hours; particularly when they are moved to a quiet room, they are allowed to rest and their bladder is empty [2].

Mechanisms and effects of elevated BP in this clinical setting have not been well understood. It might be attributable to either one more of the following conditions: preexisting, inadequately treated or undiagnosed hypertension, stress of hospitalization, raised intracranial pressure, haematoma expansion, damage to autonomic centers and abnormal baroreceptor sensitivity, neuroendocrine response with activation of sympathetic nervous system, renin-angiotensin axis and/or glucocorticoid system, and myocardial changes [3–6].

Most of the studies, although not all, have found that high BP in the acute phase of stroke, whether measured as casual or 24 hours ambulatory readings, is associated with a poor outcome [7–9] and an increased risk of death and dependency [10–14]; a U-shaped relationship between BP values and outcome has been described in different studies [15–17]. Recent evidence suggests that not only BP but even its derived indices and other haemodynamic measures as mid blood pressure, mean arterial pressure, BP variability, heart rate, pulse pressure, and rate-pressure product are related to functional outcome [18–20]. The association is thought to be related to the early stroke recurrence and the development of cerebral edema and greater serious haemorrhagic transformation in ischaemic stroke [21, 22] and to the haematoma expansion in primary intracerebral haemorrhage [23].

While observational studies show that high BP is independently associated with a poor outcome, suggesting that it should be lowered, pathophysiology argues that lowering BP will reduce cerebral blood flow when cerebral autoregulatory mechanisms are impaired. Additionally, in acute ischaemic

stroke the infarcted brain tissue may be surrounded by a “penumbra” zone of underperfused but viable tissue where cerebral blood flow extremely depends on the systemic BP and collaterals until the occluded artery is recanalized. Lowering BP carries the risk of jeopardizing the perfusion of this area leading to an increase of brain infarction or perihematoma ischaemia. Spontaneous thrombolysis may also occur, and the ischaemic area may become hyperaemic; at this stage a very high BP might cause propagation of infarct-related brain oedema or haemorrhagic transformation of the infarct. Unfortunately there is no sure clinical correlate of spontaneous thrombolysis, and in routine clinical care it is not possible to judge when it is better to leave a very high BP untreated or when it is necessary to intervene.

In summary, there is still debate in whether, when and how high BP should be lowered (epidemiological evidence) or not (pathophysiological concerns). Different antihypertensive drug classes might have differential effects [24] considering both their action in lowering BP and specific organ effects: for example β -blockers might be detrimental [25], and the use of calcium channel blockers was associated with a worsening of outcome in some studies, especially those testing intravenous formulation [26], perhaps because of a reduction in cerebral perfusion.

Our purpose is to investigate through a research of the recent literature the effect of the angiotensin receptor blockers (ARBs) administered in the acute phase of stroke on death and dependency.

2. Methods

2.1. Identification and Inclusion of Trials. We used Medline (1985 to December 2011; any language) to identify randomized controlled trials of ARBs in patients within 72 hours of stroke. We selected trials of more than 100 patients which assessed the effect on death or dependency and recorded coronary heart disease events or stroke (irrespective of whether BP lowering was considered the mechanism of action). Search terms were “blood pressure lowering”, “blood pressure reduction”, “antihypertensive”, or “hypertension” or “receptor, angiotensin/antagonists and inhibitors” or the name of all ARBs listed in the British National Formulary as keywords or text words. Limits were Medline publication type “clinical trial” or “controlled clinical trial” or “randomized controlled trial”, or “meta-analysis”. We also searched the Cochrane Collaboration and Web of Science databases and the citations in trials and meta-analysis. Randomized trials were included irrespective of participants’ age, disease status, BP before treatment, and the use of other drugs. Confounded trials (in which ≥ 2 active treatment were compared in the absence of a control arm) were excluded.

2.2. Blood Pressure and Outcome. The authors extracted data independently from identified publications with respect to patients’ number, sex, age, previous medical history, stroke subtype, time from stroke to enrollment, baseline BP, difference in BP between treatment and control groups, follow up period, grade of dependency, and death. Outcome

events included stroke (all, fatal, and nonfatal), myocardial infarction (all), total vascular events (combined stroke, MI, and vascular death), and mortality (all-cause, vascular).

3. Results

3.1. Trials. Three randomized placebo-controlled multicenter studies fulfilled the inclusion criteria (Table 1), each of which had been published [27–29]. The combined sample size was 3728 with almost two-thirds of the data coming from one study (SCAST). In two studies (ACCESS and PROFESS) all patients recruited had ischaemic stroke, while in the SCAST trial 85.4% of patients had ischaemic and 13.5% haemorrhagic stroke; mean time from stroke to enrollment varied from 17.6 to 57.6 hours, and recruitment was limited to patients with high BP. Main characteristics of each trial are summarized in Table 2.

In the ACCESS study, protocol considered a 7-day placebo controlled phase: treatment was started with 4 mg candesartan cilexetil daily or placebo on day 1; on day 2, dosage was increased to 8 or 16 mg candesartan cilexetil or placebo if BP exceeded 160 mmHg systolic or 100 mmHg diastolic; target was a 10–15% BP reduction within 24 hours. On day 7, a 24-hour BP profile was obtained in all patients: in those in the candesartan cilexetil group who showed a hypertensive profile (mean daytime BP > 135/85 mmHg) dosage was increased or an additional antihypertensive drug was added; in placebo arm, candesartan cilexetil was started in patients with hypertensive profile to lower BP to <140/90 mmHg (office BP) or <135/85 mmHg (mean daytime BP), while those with a normotensive profile did not receive antihypertensive medication. Follow-up examinations were performed after 3, 6, and 12 months.

The PROFESS study compared the effect of telmisartan (80 mg daily) versus placebo and combined aspirin (25 mg twice daily) and extended release dipyridamole (200 mg daily) versus clopidogrel (75 mg daily) in a 2×2 factorial design in patients with recent ischaemic stroke followed up for a mean duration of 30 months. The subgroup analysis reviewed included only patients randomized within 72 hours of stroke onset, evaluated at 1 and 3 months.

In the SCAST trial, patients with diagnosis of stroke (ischaemic or haemorrhagic) presenting within 30 hours of symptom onset were allocated in a 1:1 ratio to treatment with candesartan or placebo. There was a fixed-dose escalation scheme: 4 mg on day 1, 8 mg on day 2 and 16 mg on days 3–7. Subsequent evaluations took place on day 7 and at 1 and 6 months; to avoid important differences in treatment during followup, candesartan was the advised antihypertensive agent and was provided free of charge.

In each trial there were no significant differences regarding the use of concomitant medication on hospital admission or during follow-up between the active treatment and placebo groups.

3.2. Baseline Findings. In the ACCESS trial mean BP on hospital admission was 198/103 and on study onset 189/99; baseline severity of stroke was not directly reported but patients had a mean Barthel Index (BI) of 62; no data are

TABLE 1: Included trials.

Study	Year	Study design	Antihypertensive agent
Acute Candesartan Cilexetil Therapy in Stroke Survivors (ACCESS) [27]	2003	Prospective, double-blind, placebo-controlled, randomized, multicenter phase II study.	Candesartan cilexetil
Prevention Regimen for Effectively Avoiding Second Strokes (PRoFESS) [28]*	2008	Prospective, double-blind, placebo-controlled, randomized, multicenter phase III study.	Telmisartan
The angiotensin-receptor blocker candesartan for treatment of acute stroke (SCAST) [29]	2011	Prospective, double-blind, placebo-controlled, randomized, multicenter phase III study.	Candesartan cilexetil

* In the PRoFESS trial a subgroup analysis has been considered.

TABLE 2: Main characteristics of the included trials.

Study	ACCESS [27]	PRoFESS [28]*	SCATS [29]
Main inclusion criteria	Motor deficit, cerebral CT scan excluding intracranial hemorrhage, onset of symptoms within 72 hours, necessity to treat hypertension according to current recommendations [#] .	Ischemic stroke within 72 hours from onset of symptoms, age older than 55 years or age 50–54 years if 2 additional vascular risk factors present, seated systolic BP 121 to 180 mmHg, seated diastolic BP \geq 110 mmHg, neurological and clinical stability.	Patients aged 18 years or older with a clinical diagnosis of stroke (ischaemic or haemorrhagic), presenting within 30 hours of symptom onset and with systolic BP higher than 140 mmHg.
Relevant exclusion criteria	Age \geq 85 years, occlusion or \geq 70% stenosis of the internal carotid artery, malignant hypertension, manifest cardiac failure (NYHA class III and IV), high-grade aortic or mitral stenosis, unstable angina pectoris, contraindications against candesartan cilexetil.	Dysphagia preventing oral medication, mRS $>$ 3 at time of randomization, severe known renal insufficiency or renal artery stenosis or coronary artery disease or recent MI, hyperkalemia, uncorrected volume or sodium depletion, schedule for carotid endarterectomy, currently using or needing ARB	SSS consciousness score \leq 2, premonitory mRS \geq 4, clear indication for or contraindications to or current treatment with an ARB.
Treatment design	Candesartan cilexetil (4–16 mg daily according to BP levels) [†] .	Telmisartan (80 mg daily).	Candesartan cilexetil (4 mg on day 1, 8 mg on day 2 and 16 mg on days 3–7) [†] .
Follow-up evaluation	On day 7, at 3, 6, and 12 months.	On day 7, at 1, and 3 months.	On day 7, at 1, and 6 months.
Primary endpoints	Case fatality and disability (measured as BI) at 3 months.	Combined death or dependency (measured as mRS) at 30 days.	Composite endpoint of vascular death, nonfatal MI or nonfatal stroke and functional status (measured as mRS) at 6 months.
Secondary endpoints	Overall mortality and cerebrovascular and cardiovascular events at 12 months.	Overall mortality and cerebrovascular and cardiovascular events at 7, 30, and 90 days.	Stroke progression [‡] ; neurological status at 7 days (measured as SSS); overall mortality, cerebrovascular and cardiovascular events, functional outcome (measured as BI) at 6 months.

* In the PRoFESS trial a subgroup analysis has been considered.

[#] This was assumed when the mean of at least 2 blood pressure measurements was \geq 200 mmHg systolic and/or \geq 110 mmHg diastolic 6 to 24 hours after admission or \geq 180 mmHg systolic and/or \geq 105 mmHg diastolic 24 to 36 hours after admission.

[†] Candesartan cilexetil 4 mg daily on day 1; on day 2, dosage was increased to 8 or 16 mg if blood pressure exceeded 160 mmHg systolic or 100 mmHg diastolic. At the end of the placebo-controlled 7-days phase, in candesartan cilexetil-treated patients the dosage was increased or an additional antihypertensive drug was added only in the case of a hypertensive profile (mean daytime blood pressure \geq 135/85 mm Hg). In placebo-treated patients candesartan cilexetil was started only in presence of a hypertensive profile.

[‡] Dose adjustments were made if systolic blood pressure was lower than 120 mmHg or when clinically indicated.

[‡] Stroke progression was defined as a neurological deterioration of 2 or more points on the SSS occurring within the first 72 h of stroke onset and believed to be caused by the index stroke, after exclusion of recurrent stroke or systemic reasons for deterioration.

ARB: angiotensin receptor blocker; BI: barthel index; BP: blood pressure; MI: myocardial infarction; mRS: modified ranking scale; NYHA: New York Heart Association; SSS: Scandinavian stroke scale.

available about etiopathogenesis of strokes. In the PROfESS subgroup analysis mean baseline BP was 147/84 mmHg and mean National Institutes of Health stroke scale (NIHSS) at admission was 3. According to TOAST classification, strokes were due to small artery occlusion in 59.5%, large-artery atherosclerosis in 20.9%, cardioembolism in 1.4%, and to undetermined or other determined etiologies in the remaining cases. In the SCAST study mean baseline BP was 171/90 mmHg and patients enrolled had a mean Scandinavian stroke scale (SSS) score of 41, equivalent to a NIHSS of 8 [30]. Strokes have been described following topographic rather than etiological criteria: according to Oxfordshire Community Stroke Project (OCSP) classification, strokes were distributed as total anterior syndrome in 8%, partial anterior syndrome in 48.7%, posterior syndrome in 14%, lacunar syndrome in 28.9%, unknown in <1%. Main baseline findings of the included studies are illustrated in Table 3.

3.3. Blood Pressure and Outcomes. In the ACCESS population no significant difference in BP was evident between the groups during the placebo-controlled phase and in the subsequent followup; at 7 days mean BP was 160/87 mmHg, at 3-months 150/85 mmHg and at 12 months 147/83 mmHg. Only two patients in the placebo arm had a normotensive profile on day 7 and did not receive the antihypertensive drug. In the PROfESS subgroup analysis, telmisartan lowered significantly systolic BP (SBP) by 6 to 7 mmHg and diastolic BP (DBP) by 2 to 4 mmHg over the whole study course: on day 7, BP was 135.3 (17.8)/78.4 (10.8) mmHg in treatment arm and 141.4 (17.0)/81.6 (11.0) mmHg in placebo group, with a difference of 6.1/3.2 mmHg ($P < 0.0001$); on day 30, mean BP was 135.7/79.6 mmHg versus 142.6/83.1 mmHg (treatment versus placebo) with a difference of 6.9/3.6 mmHg ($P < 0.0001$); on day 90, mean SBP was 134.5 (19.9) mmHg (treatment) versus 140.3 (19.0) (placebo) with a difference of 5.8 mmHg ($P < 0.0001$) and mean DBP was 79.2 (11.1) mmHg versus 81.5 (11.2) mmHg (treatment versus placebo) with a mean difference of 2.4 mmHg ($P = 0.0002$).

In the SCAST trial BP fell in both groups during treatment but was significantly lower in patients allocated candesartan than in those on placebo ($P \leq 0.001$ for days 2–7); on day 7, mean BP was 147/82 mmHg (SD 23/14) in the candesartan group and 152/84 mmHg (SD 22/14) in the placebo group. The mean difference in SBP on day 7 was 5 mmHg (95% CI 3–7; $P < 0.0001$) and the mean difference in DBP was 2 mmHg (1–3; $P = 0.001$). During the 6-month followup, mean BP values were similar in the two groups, and at 6 months the mean BP was 143/81 mmHg in both groups.

In the ACCESS the BI revealed no significant difference on study onset and after 3 months (candesartan cilexetil versus placebo, day 0: 60.0 \pm 30.2 versus 64.1 \pm 27.5; 3 months: 87.0 \pm 22.9 versus 88.9 \pm 19.9). The cumulative 12-month mortality (candesartan cilexetil versus placebo: 2.9% versus 7.2%; $P = 0.07$) and the number of vascular events (candesartan cilexetil versus placebo: 9.8% versus 18.7%; $P = 0.026$) differed significantly in favor of the candesartan cilexetil group; the odds ratio was 0.475 (95% CI, 0.252 to

0.895). The clinical benefit was independent of BP values and mainly attributable to a lower incidence of myocardial ischaemic events. Drug tolerance and number or type of undesirable effects did not differ significantly between the groups.

In the PROfESS subgroup analysis combined death or dependency (mRS at 30 days, with adjustment for baseline covariates) did not differ whether analyzed as an ordinal outcome (ordered mRS categories: 0, 1, 2, 3, 4–6 to maintain proportionality) (OR, 1.03; 95% CI, 0.84–1.26; $P = 0.81$) or with dichotomization of the data at the median (mRS 0–1 versus 2–6; OR, 1.00; 95% CI, 0.77–1.29). There was no significant difference between the treatment groups for the distribution of ordinal stroke events (fatal, dependent (mRS 2–5), independent (mRS 0,1), TIA, none), for the time to recurrence ($P = 0.40$) and, similarly, for other events (i.e., death, stroke recurrence, MI, and combined vascular events) at 7, 30, or 90 days. Serious adverse events were similar between telmisartan and control.

In the SCAST trial the analysis of the first co-primary effect variable, the cumulative risk of the composite endpoint of vascular death, stroke or myocardial infarction, showed no significant difference between candesartan and placebo (unadjusted analysis HR 1.09, 95% CI 0.84–1.41; $P = 0.53$; adjusted analysis HR 1.09, 0.84–1.41; $P = 0.52$; per-protocol analysis HR 1.11, 0.85–1.46; $P = 0.46$). Regarding the second co-primary effect variable, the functional outcome at 6 months, no significant difference was seen across the mRS categories (unadjusted ordinal regression analysis, OR 1.13, 95% CI 0.97–1.32; $P = 0.12$); similar results have been obtained in both the fixed dichotomy (mRS 3–6 versus 0–2) analysis (unfavourable outcomes in 35% of patients on candesartan and in 33% of patients allocated on placebo; OR 1.12, 0.90–1.41, $P = 0.32$; RR 1.06, 0.93–1.19, $P = 0.39$) and the sliding dichotomy analysis using the SSS scores at baseline (unfavourable outcomes in 56% of patients on candesartan and 52% of patients on placebo; OR 1.16, 95% CI 0.97–1.38, $P = 0.11$; RR 1.07, 95% CI 0.99–1.16, $P = 0.11$). For all the secondary outcomes assessed there were no significant differences between treatment and placebo: for all events (death from any cause, vascular death, ischaemic stroke, haemorrhagic stroke, all strokes, myocardial infarction, stroke progression, symptomatic hypotension, renal failure, and symptomatic venous thromboembolism: RR 1.47, 95% CI 1.01–2.13; $P = 0.04$), for SSS score at 7 days ($P = 0.13$) and for BI at 6 months (0.47). There were no significant differences between the groups for the adverse events reported by the investigators.

Significant differences in BP values between treatment and placebo were therefore found in the PROfESS and SCAST trials, but in only the first it remained significant during the follow-up; conversely, in the ACCESS study a significant difference was never found.

In all trials no effect of the active treatment was seen on dependency, death and vascular events at one, three or six months; in the ACCESS study a decrease in cumulative mortality and number of vascular, mainly myocardial, events emerged in candesartan group at 12 months.

TABLE 3: Patients characteristics at enrollment.

Study	ACCESS [27] treatment/placebo	PRoFESS [28]* treatment/placebo	SCATS [29] treatment/placebo
Number of patients	173/166	647/713	1017/1012
Age	68.3 (9.3)/67.8 (9.4)	66.8 (8.8)/67.1 (9.2)	70.8 (11.2)/71.0 (11.0)
Male (%)	86 (49.7)/86 (51.8)	420 (64.9)/464 (65.1)	612 (60.2)/564 (55.7)
Clinical history			
Previous stroke/TIA	NA	160 (24.7)/184 (25.8)	252 (24.8)/204 (20.2)
Atrial fibrillation	NA	10 (1.6)/14 (2.0)	190 (18.7)/186 (18.4)
Hypertension	NA	453 (70.0)/503 (70.6)	676 (66.5)/670 (66.2)
Diabetes mellitus	67 (38.7)/58 (35.0)	176 (27.2)/198 (27.8)	163 (16.0)/157 (15.5)
Hyperlipidemia	74 (42.8)/75 (45.2)	264 (40.8)/283 (39.7)	NA
Ischemic heart disease	38 (22)/32 (19.3)	95 (14.7)/104 (14.6)	NA
Time from stroke (hours)	29.9/29.7	57.6 (16.8)/57.6 (16.8)	17.6 (8.1)/17.9 (8.1)
Blood pressure (mmHg)			
Systolic	188 (20.9)/190 (19.7)	146 (16.2)/147 (16.3)	171.2 (19.0)/171.6 (19.2)
Diastolic	99 (14.9)/99 (13.0)	84 (10.1)/84 (10.2)	90.3 (13.9)/90.6 (14.2)
Clinical severity	60.0 (30.2)/64.1 (27.5) at BI	2.9 (2.8)/3.1 (2.9) at NIHSS	40.6 (12.3)/40.5 (12.6) at SSS

* In the PRoFESS trial a subgroup analysis has been considered.

Data are *n* (%) or mean (SD). NA: not assessed. BI: Barthel Index; mRS: modified ranking scale; NIHSS: National Institutes of Health Stroke Scale; SSS: Scandinavian stroke scale.

4. Discussion

While the primary prevention of stroke through the treatment of hypertension is well established and evidence from randomized controlled trials suggests the use of antihypertensive agents for the secondary prevention of vascular events in patients with previous stroke or transient ischaemic attack, the management of BP immediately after stroke has been an enigmatic controversy for more than two decades, exemplified by a debate published back in 1985 [31], and still remains uncertain. In experimental studies on rats, candesartan has been shown to be neuroprotective with both reduction of neurovascular damage, demonstrated by decreased infarct size, hemoglobin content and oedema in the ischaemic brain hemisphere, and improvement of neurological outcome. These favorable effects were evident with an early post ischemic stroke drug administration, at doses that did not affect or moderately lowered BP [32, 33], suggesting a multimodal protective effect, partly due to BP lowering and partly due to pleiotropic, vascular and neuronal, actions. Homeostatic defense processes against postischaemic brain damage, or almost a part of them, are pressure dependent; presumably there are feedbacks and fine interrelationships among BP, the processes limiting the infarct size and the mechanisms involved in the recovery of ischemic tissue. On the other side, competitive inhibition of the binding of angiotensin (AT) II to AT₁ receptors allows an unopposed activation of the AT₂ receptors which has been hypothesized to be protective in focal cerebral ischaemia being responsible of several events as recruitment of cerebral collaterals, normalization of cerebrovascular autoregulation, enhancement of neuronal resistance to anoxia, inhibition of inducible nitric oxide synthetase, reduction of oxidative damage, prevention of apoptosis, promotion of angiogenesis

and attenuation of inflammation, endothelial dysfunction and prothrombosis [34, 35].

Translating these experimental findings to human stroke patients is quite difficult. The clinician who faces a patient in the acute phase of stroke with elevated BP has these main alternatives: he may choose to continue or not a preexisting antihypertensive drug or to introduce or not a new one. In the Continue Or Stop post-Stroke Antihypertensives Collaborative study [36], continuation compared with cessation of preexisting antihypertensive drugs for a two weeks period after acute stroke was not associated with a substantial reduction in two-week death or dependency, cardiovascular event rate or mortality at six months. Besides, clinical trials evaluating the effects of different BP lowering drugs administered in the acute phase of stroke have given conflicting results in respect of functional outcome, and the evidence to guide the practicing clinician is still rudimentary. Sharp reduction in arterial BP has been sought to determine a worse short and long-term prognosis [37, 38], while a moderate and caution reduction might be safe and even improve long term mortality and reduce recurrent vascular events [39, 40]. Not only the degree of BP reduction but even the nature of the pharmacological agent itself, considering its specific mechanism of action and in the perspective of possible drug-class related benefits, has been and still remain matter of debate.

The purpose of this systematic review is to investigate the role of a specific class drug, the ARBs, early administered after stroke. None of the included studies demonstrated a significant benefit of active treatment on functional outcome and stroke recurrence at short and medium term, but some evaluations should be taken in account. Firstly, the mean BP at enrollment was much higher in the ACCESS study (189/99 mmHg) in respect to both the PRoFESS and SCAS

trials (147/84 mmHg and 171/90 mmHg, resp.). Secondly, relevant differences existed in stroke severity: patients enrolled in the PROFESS had very mild stroke in respect to those participating to the SCAST and above all to the ACCESS who had a significantly greater impairment. Thirdly, the effect of treatment on BP: candesartan did not alter BP in the ACCESS while in the PROFESS, despite the fact that BP at baseline was reasonably well controlled, telmisartan further reduced it; also in the SCAST trial BP fell in both treatment and placebo but was significantly lower in patients allocated candesartan although difference disappeared during the followup. Finally, the achievement of endpoints: neither functional outcome nor number of vascular events have been positively affected by treatment but in the ACCESS study, although the primary outcome (disability at 3 months) was neutral, treatment with candesartan was associated with a significant reduction in secondary outcome including the 12-month mortality and vascular events. Taking into account the U-shaped relationship between BP and outcome in acute stroke, the main advantage deriving from lowering BP treatment would be expected in the ACCESS trial which included patients with severely elevated BP. Even in this study, however, 3-month outcome has not been influenced by treatment and the reduction in the number of vascular events observed in favor of the candesartan cilexetil group was mainly due to a lower incidence of myocardial ischaemic events but not of recurrent cerebral ischaemic events which did not significantly contribute to the difference in cardiovascular morbidity and mortality. Much more surprising is the fact that in this study BP has not been affected by treatment, raising the question around the existence of a drug specific effect of the AT₁ receptor blockade, beyond the hemodynamic activity, able to modulate vascular remodeling and to affect cardiovascular survival with a benefit that did not arise immediately but appeared to increase over the time. Analogous data have been acquired from various cardiac intervention studies suggesting the hypothesis that early neurohumoral inhibition has similar beneficial effects in both cerebral and myocardial ischaemia, although the underlying mechanism is not resolved. These findings, however, have not been confirmed in the PROFESS and SCAST trials, and such discrepancies may simply reflect a false positive finding of a small trial which was stopped prematurely on the basis of an interim analysis, requiring more investigations.

Taking their similarities and differences, the studies we have considered raise doubts over the hypothesis of a specific effect of angiotensin receptors blockade in acute stroke; moreover, conclusions are fully compatible with those of a recent meta-analysis [29] and a regularly update Cochrane survey [41] of randomized controlled trials of BP lowering drugs in acute stroke. According to currently available evidence there is no clear evidence of benefit for routine BP lowering treatment in the acute phase of stroke; many pitfalls, however, still remain to explain. Firstly, BP management in this clinical setting may have to be tailored with respect to the underlying etiology, and other parameters than baseline BP may need to be taken into consideration [42, 43]. Patients with atherosclerotic or lacunar strokes are often affected by chronic hypertension with subsequent arterial stiffness and

shift of the cerebral blood flow autoregulatory curve to the right. Consequently, they may tolerate elevated BP in acute stroke more efficiently, and BP values considered “normal” for the general population may be inadequate to perfuse the ischaemic brain. These patients usually present also diffuse atherosclerotic lesions in cerebral vessels which compromise the patency of collateral circulation: high BP may be needed to enhance perfusion of the ischaemic penumbra zone. On the contrary, in the cardioembolic strokes patients may only have atrial fibrillation in absence of arterial hypertension or significant atherosclerotic stenosis and may need only moderately elevated BP to promote perfusion through a patent collateral circulation; moreover, cardioembolic strokes tend to be of larger size with a higher risk of edema and hemorrhagic transformation in case of raised levels of BP. Unfortunately, the trials considered neither reported nor stratified outcome data according to the aetiopathogenesis of strokes. Secondly, the interrelationships between baseline BP, stroke severity, and administration of antihypertensive agents have not been well understood but an interaction could not be excluded; in the PROFESS study the failure to show beneficial effect of telmisartan may reflect that patients had only mild hypertension and mild stroke. Thirdly, favourable effect of treatment may show a significant time interaction; a post hoc analysis of the main PROFESS study indicated that recurrence was lower with telmisartan after the first six months of treatment [44], and in the ACCESS benefit on overall mortality and vascular events did not arise immediately but instead appeared to increase during follow up. The included studies presented followup ranging from 3 to 12 months and have not been designed to investigate long term effects of treatment.

Although a benefit of early treatment has not emerged, it is noteworthy that a substantial safety has resulted without difference in number or type of undesirable effects in treatment and placebo groups; since chronic lowering BP reduces strokes recurrence [45, 46] it may be safe to start such treatment even acutely in selected subgroups. Many other issues like the timing of starting the treatment, the degree, and rapidity of BP reduction, also taking account of the initial level, and the formulation, route of administration, and doses of different pharmacological agents should be further addressed; it is not possible to rule out that different drugs might have different effects.

Our analyses are not ideal in some respects. Firstly, trial-level data rather than individual patients' data were assessed since the latter were not available to us; analyses based on individual patients' data are generally superior and allow subgroup analyses to be performed. Secondly, the inhomogeneity of the patients' population and of the design of the study methods regarding cut offs for BP, stroke severity, time of assessment of the outcome variables; for the PROFESS study, we have considered a subgroup of the patients entered into the main large secondary prevention trial, such that patients' characteristics reflect the inclusion criteria for a study of vascular prophylaxis rather than acute intervention. Thirdly, we could not assess the effect of lowering BP in patients with different subtypes of strokes since trials did not report these data separately.

5. Conclusion and Future Research

The currently available studies did not identify any clear indication that treatment with the ARBs is beneficial in patients with acute stroke and raised BP on functional outcome and stroke recurrence at short and medium term. Many issues, however, should be still considered; two large studies involving more than 2500 patients with acute ischaemic (ENOS) and haemorrhagic stroke (ENOS, INTERACT 2) [47, 48] are ongoing, and it is favorable these and future trials will help to clarify many unresolved questions and whether there are subgroups of patients or different approaches to BP management for which a treatment benefit can be obtained.

Conflict of Interest

All authors have no conflict of interests.

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Review Article

AT2 Receptor-Interacting Proteins ATIPs in the Brain

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Received 29 November 2012; Accepted 22 December 2012

Academic Editor: Patrick Vanderheyden

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A complete renin-angiotensin system (RAS) is locally expressed in the brain and fulfills important functions. Angiotensin II, the major biologically active peptide of the RAS, acts via binding to two main receptor subtypes designated AT1 and AT2. The present paper focuses on AT2 receptors, which have been reported to have neuroprotective effects on stroke, degenerative diseases, and cognitive functions. Our group has identified a family of AT2 receptor interacting proteins (ATIPs) comprising three major members (ATIP1, ATIP3, and ATIP4) with different intracellular localization. Of interest, all ATIP members are expressed in brain tissues and carry a conserved domain able to interact with the AT2 receptor intracellular tail, suggesting a role in AT2-mediated brain functions. We summarize here current knowledge on the ATIP family of proteins, and we present new experimental evidence showing interaction defects between ATIP1 and two mutant forms of the AT2 receptor identified in cases of mental retardation. These studies point to a functional role of the AT2/ATIP1 axis in cognition.

1. Introduction

The renin-angiotensin system (RAS), a major regulator of blood pressure and cardiovascular functions, is now fully recognized as playing important roles in the brain [1–6]. Among the active peptides generated by the RAS, angiotensin II (AngII) stands as the best characterized. This octapeptide binds to two receptor subtypes, namely, AT1 and AT2, that belong to the superfamily of seven transmembrane domains receptors. In most tissues and cell lines, AT1 appears as a driving receptor that mediates most effects of AngII by activating classical heterotrimeric G proteins and intracellular signaling cascades [6, 7]. In contrast, AT2 is generally considered as an AT1-counteracting receptor that involves nonclassical signaling pathways and does not necessarily require exposure to AngII [6, 8]. Over the past few years, several AT2 receptor interacting partners have been identified [8–12], among

which SHP-1 [8, 9], PLZF [13], and ATIP1/ATBP50 [12, 14, 15] that regulate AT2 receptor trafficking, internalization and/or activation. In this paper, we will focus on the family of ATIP proteins and their potential roles in AT2-mediated brain functions.

2. AT2 Receptor in the Brain

In contrast to the ubiquitous AT1 receptor, the AT2 subtype is predominantly expressed during embryonic development and is restricted to few sites in the adult [8, 17, 18]. In the central nervous system, AT2 expression is high during fetal life and remains elevated in the adult in specific areas involved in cognition, behavior, and locomotion [1, 18–20]. AT2 is mainly expressed in neurons and mediates neuronal differentiation [21–27], survival [28–30] and regeneration [31–33] through the regulation of protein kinases and phosphatases

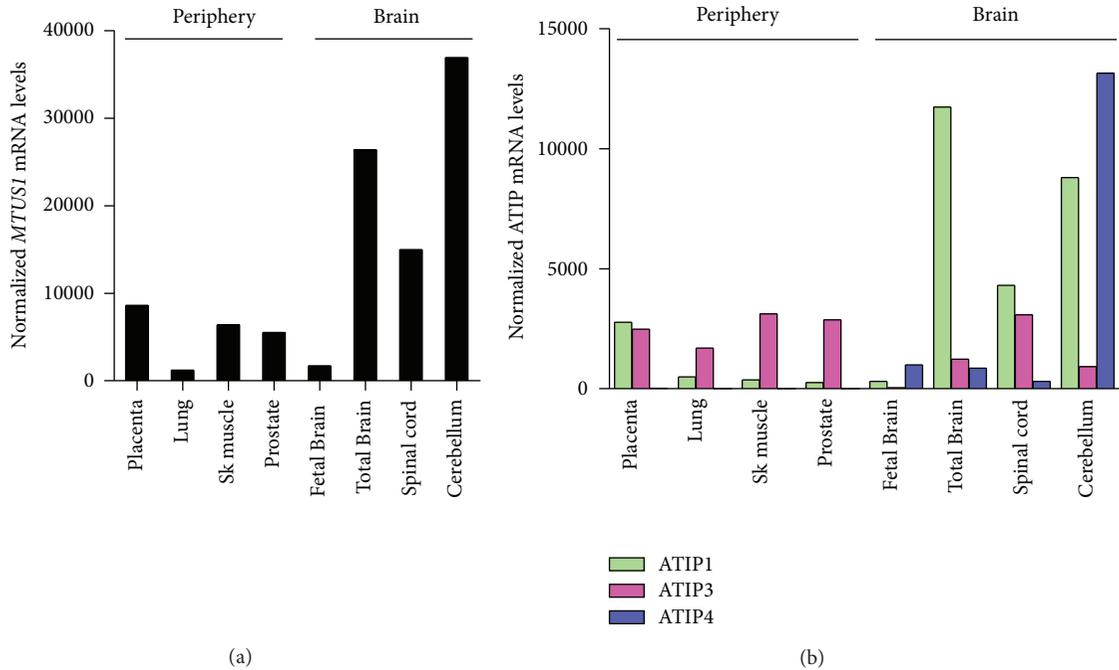


FIGURE 1: Distribution profiles of ATIP transcripts in normal human tissues. (a) Real-time PCR using probes common to all ATIP transcripts (3' exons). (b) Real-time PCR using probes specific for each ATIP transcript (5' exons). Results presented are from Di Benedetto et al., 2006 [16], and are normalized relative to the levels of human acidic ribosomal phosphoprotein P0 (RPLP0). Sk. muscle: skeletal muscle.

[8, 34–38], and the reorganization of the cytoskeleton [39, 40]. Functional *in vivo* studies tend to indicate protective effects of the AT2 receptor against stroke, Alzheimer disease, and cognitive impairment [26, 33, 41–45]. Involvement of AT2 receptors in cognition has also been suggested by the identification of mutations in the corresponding *AGTR2* gene in several cases of mental retardation [46–48]. However these results remain a matter of debate [49–52], and functional alterations of AT2 receptors in mental retardation have still to be demonstrated. Further analyses of AT2 signaling pathways and AT2 interacting partners in the brain [53] following or not receptor activation with compound 21 (M024), a new selective AT2 ligand [54–56], should bring further insights on the effects of AT2 receptors in normal and pathological situations.

3. A Family of AT2 Receptor-Interacting Proteins (ATIP)

A family of AT2 receptor-interacting proteins (ATIPs) has been identified by a two-hybrid system cloning strategy using as a bait the 52 carboxy-terminal residues of the AT2 receptor [14, 15]. Three major human ATIP members (ATIP1, ATIP3, and ATIP4) are encoded by alternative exon splicing [16] and from alternative promoters [57] present on a single gene designated *MTUS1*. This gene contains 17 coding exons encompassing more than 112 kilobases and localizes at chromosomal position 8p22 [16]. All three ATIP transcripts use the same 3' exons of the gene, and therefore encoded proteins are identical in their carboxy-terminal (395 amino

acids) portion, which carries the AT2 receptor-interacting domain [12, 14, 16]. Thus, each ATIP member is in principle able to interact with AT2, although to date, only ATIP1 has been formally demonstrated to bind AT2 in living cells [14, 15, 34]. Expression of all three mRNA species has been detected in nonpathological human tissues by real-time PCR analysis using probes specific for each splice isoform [16] (Figure 1). ATIP1 and ATIP3 are ubiquitous whereas ATIP4 expression is restricted to the central nervous system. All ATIP transcripts were found expressed in every brain area examined [16], ATIP1 being predominant in all brain regions except cerebellum and fetal brain in which ATIP4 represents the major ATIP species.

4. The ATIP1/AT2 Axis in Neuronal Differentiation

ATIP1 (also designated *MTSG1* and *ATBP50* in the mouse) is the first characterized member of the ATIP protein family [14, 15, 58]. ATIP1 is a cytosolic protein that inhibits cell proliferation, receptor tyrosine kinase signaling, and ERK phosphorylation and contributes to the trafficking of the AT2 receptor from the Golgi to the cell membrane.

Real-time PCR analysis of ATIP transcripts in human tissues has revealed that ATIP1 is ubiquitous and the most abundant ATIP mRNA species expressed in the brain [16] (Figure 1). However, only few studies have investigated the effects of ATIP1 in brain functions. In rat fetal neurons, ATIP1 is constitutively associated with the AT2 receptor at the cell membrane and is part of a multimeric complex

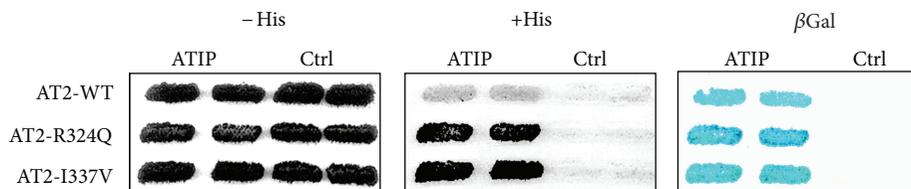


FIGURE 2: Interaction between ATIP1 and wild-type or mutated AT2 receptors. The C-terminus domain of ATIP1 (ATIP) interacts with the C-terminal region of human AT2 receptor, either wild-type (WT) or mutated (R324Q, I337V). The yeast reporter strain HF7 expressing the pairs of indicated hybrid proteins was analyzed for histidine auxotrophy and β -galactosidase expression as described [14]. Transformants were plated on medium with histidine (left), without histidine (His, middle), or replica-plated on Whatman filters and tested for β -galactosidase activity (β Gal, right). Growth in the absence of histidine and blue color in the β -galactosidase assay indicate interaction between hybrid proteins (WT: wild-type sequence; Ctrl; empty vector).

comprising the AT2 receptor and the SHP-1 tyrosine phosphatase [34]. Upon AT2 receptor activation by AngII, ATIP1 and SHP-1 remain associated but detach from the AT2 receptor and translocate from the cell membrane to the nucleus. In the nucleus, the ATIP1/SHP-1 complex activates the transcription of the methyl methanesulfonate sensitive 2 (MMS2) gene, thereby contributing to AT2-mediated neuronal differentiation [34]. These data suggest that detachment of ATIP1 from the AT2 receptor, rather than its association, may trigger activation of AT2 signaling pathways. Accordingly, dissociation of ATIP1/AT2 complexes following AngII stimulation has also been reported in transfected Chinese Hamster Ovary (CHO) cells [14].

5. ATIP1/AT2 Alterations in Mental Retardation

ATIP1 interacts with the C-terminal intracellular portion of the AT2 receptor [14, 15]. Interestingly, two nonconservative amino acid substitutions (R324Q and I337V) in the carboxy-terminal sequence of the human AT2 receptor have been identified in cases of mental retardation [46], prompting us to investigate whether these alterations may impact on the ability of the AT2 receptor to recruit ATIP1. We addressed this question using the two-hybrid system in yeast. The last 52 amino acids of the human AT2 receptor, either wild-type or mutated (R324Q or I337V), were PCR-amplified and subcloned into the pGBT9 vector in frame with the Gal4-DNA binding domain. The AT2-interacting domain of ATIP1 was subcloned into the VP16 vector. Interactions were assayed as previously described in the HF7 yeast strain which contains Histidine and beta-galactosidase reporter genes [14]. Interaction between the C-terminal domain of ATIP1 and the C-terminal region of the human AT2 receptor was confirmed (Figure 2). To our surprise, the interaction of ATIP1 with each of the mutated forms of the AT2 receptor was stronger compared to the interaction with wild-type AT2, suggesting that AT2 mutants may exhibit higher affinity for ATIP1. These data raise the interesting possibility that mutated AT2 receptors may retain ATIP1 at the cell membrane upon AngII stimulation. Further experiments are required to explore this hypothesis. We speculate that AT2 mutations (R324Q and I337V) identified in mental retardation may impair the intracellular activity of the receptor by

preventing the release of ATIP1. These results suggest for the first time that dysfunctions in the AT2/ATIP1 axis may be involved in mental retardation. They point to a role for ATIP1 in brain functions and relaunch the debate on the functional involvement of AT2 receptors in mental retardation.

6. Microtubule-Associated ATIP3

As mentioned before, ATIP3 is identical to ATIP1 in the carboxy-terminal region carrying the AT2-interacting domain; however whether this isoform indeed interacts with the AT2 receptor in living cells remains to be determined. QPCR analyses revealed that ATIP3 transcripts are expressed in all human tissues including in the central nervous system [16]. However, ATIP3 functions in the brain have not yet been investigated. Of interest, ATIP3 closely associates with microtubules (Figure 3) [12, 59, 60], suggesting possible roles of this protein in diverse biological functions associated with cytoskeleton remodeling. Indeed, ATIP3 localizes to the mitotic spindle during cell division and acts as a potent antimitotic protein that inhibits cancer cell proliferation *in vitro* and tumor growth *in vivo*, in line with tumor suppressor effects of ATIP3 reported in breast cancer [59].

In the brain, microtubules play essential roles by regulating neuronal differentiation, neurite outgrowth, and cell migration [61, 62]. Alterations of microtubule-associated proteins such as tau are strongly associated with the occurrence of neurodegenerative pathologies, including Alzheimer disease [63, 64], in which AT2 receptors have also been implicated [43, 44, 65]. Whether microtubule-associated effects of ATIP3 may also contribute to the regulation of brain functions, in response or not to AT2 receptor stimulation, is a question that deserves further studies.

7. Brain-Specific Expression of ATIP4

The cDNA cloning and functional characterization of the ATIP4 isoform have not been undertaken to date. ATIP4 presents two interesting features that make it a good candidate for mediating AT2 functions in the brain. First, expression of the ATIP4 mRNA is restricted to the brain and remains undetectable in peripheral tissues [16] (Figure 1). Of note, ATIP4 mRNA levels are highest in the fetal brain and

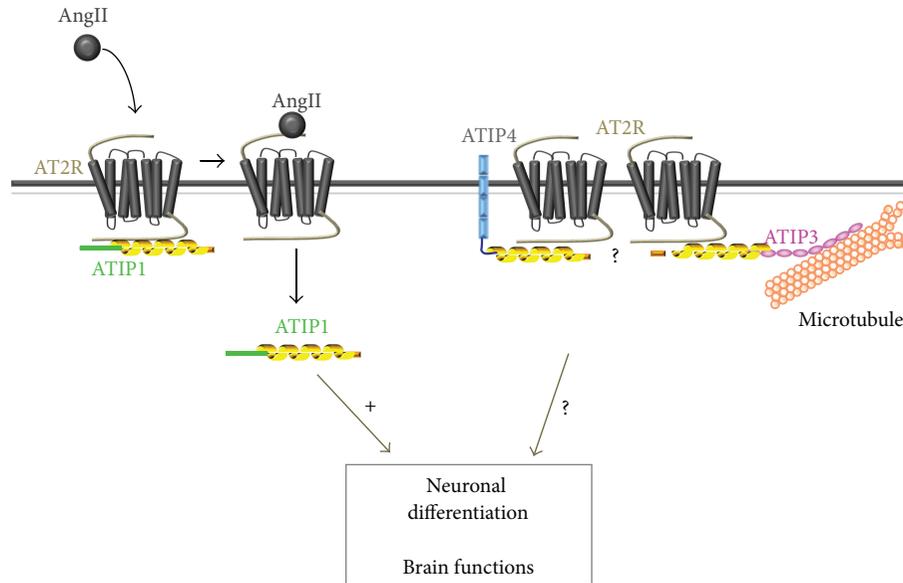


FIGURE 3: Schematic representation of ATIPs localization and interaction with AT2 receptors. ATIP1 is constitutively associated with the AT2 receptor at the cell membrane in rat fetal neurons and dissociates from the receptor upon AngII stimulation [34]. Putative interactions of ATIP3 and ATIP4 with AT2 receptors through their respective carboxy-terminal regions are represented.

in the cerebellum, which are two regions of abundant AT2 receptor expression in human brain [20]. Second, the amino acid sequence of ATIP4 contains a stretch of 24 hydrophobic residues flanked by charged residues, which is the hallmark of intrinsic membrane-spanning domains. Based on these *in silico* observations, it is tempting to speculate that ATIP4 might be structurally organized as a transmembrane protein with a short (36 residues) N-terminal extracellular domain and an intracellular region (456 residues) able to interact with the AT2 receptor (Figure 3). Future studies should be designed to investigate whether ATIP4 and AT2 are indeed colocalized at the plasma membrane in neuronal cells, and whether they functionally interact to regulate important brain functions. Prominent ATIP4 expression in the cerebellum compared to other regions of the brain (Figure 1) may suggest involvement of this ATIP isoform in functions related to locomotion, behavior, and/or cognition.

8. Concluding Remarks

Since the discovery of a new family of AT2 receptor interacting proteins in 2004, the question of whether these polypeptides may play a role in normal and/or pathological brain functions has not been addressed. Notably, all ATIP members are abundantly expressed in the brain and share the same C-terminal domain able to interact with the AT2 receptor, suggesting that each ATIP member may contribute to brain AT2 receptor functions. A functional AT2/ATIP1 axis has been previously reported to be involved in rat fetal neuron differentiation. We present here evidence that AT2/ATIP1 interactions are altered by *AGTR2* mutations identified in cases of mental retardation. These data relaunch the debate on the implication of AT2 receptors in mental retardation and

point to *MTUS1* as an attractive target gene in human brain pathologies.

Conflict of Interests

The authors have no conflict of interests to declare.

Acknowledgments

This work was supported by the University Paris Descartes, Inserm, CNRS, Roche laboratories, the foundation RAJA-Danièle-Marcovici, the association Odyssea, and the association Prolific.

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Review Article

The Brain Renin-Angiotensin System and Mitochondrial Function: Influence on Blood Pressure and Baroreflex in Transgenic Rat Strains

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Received 2 October 2012; Accepted 23 December 2012

Academic Editor: Robert C. Speth

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Mitochondrial dysfunction is implicated in many cardiovascular diseases, including hypertension, and may be associated with an overactive renin-angiotensin system (RAS). Angiotensin (Ang) II, a potent vasoconstrictor hormone of the RAS, also impairs baroreflex and mitochondrial function. Most deleterious cardiovascular actions of Ang II are thought to be mediated by NADPH-oxidase- (NOX-) derived reactive oxygen species (ROS) that may also stimulate mitochondrial oxidant release and alter redox-sensitive signaling pathways in the brain. Within the RAS, the actions of Ang II are counterbalanced by Ang-(1-7), a vasodilatory peptide known to mitigate against increased oxidant stress. A balance between Ang II and Ang-(1-7) within the brain dorsal medulla contributes to maintenance of normal blood pressure and proper functioning of the arterial baroreceptor reflex for control of heart rate. We propose that Ang-(1-7) may negatively regulate the redox signaling pathways activated by Ang II to maintain normal blood pressure, baroreflex, and mitochondrial function through attenuating ROS (NOX-generated and/or mitochondrial).

1. Introduction

The renin-angiotensin system (RAS), and in particular angiotensin (Ang) II, is implicated in the impairment of arterial baroreflex function and reduction of heart rate variability (HRV) commonly associated with hypertension [1-4]. However, more recent studies suggest that a part of the deficit in sensitivity of the baroreflex function (BRS) in hypertension results from a reduction in Ang-(1-7), an alternative product of the RAS, rather than a frank increase in Ang II [5, 6]. Ang II blockade attenuates oxidant production and improves mitochondrial function in peripheral tissues in various experimental models of hypertension [7-10]. The contributions of Ang-(1-7) to the beneficial effects of Ang II blockers are increasingly recognized [11-16], but few studies have directly addressed the role of Ang-(1-7) in mitochondrial function. In this paper, we summarize (1) the

role of Ang II in reactive oxygen species (ROS) generation and (2) the implication of ROS and redox-signaling on blood pressure, baroreflex, and mitochondrial function, with a particular focus on potential mechanisms for the counterbalancing role of Ang-(1-7) (Figure 1). Furthermore, we highlight the recent studies in transgenic rats with altered brain RAS (summarized in Figure 2) as a tool to study changes in brain ROS and signaling pathways in response to Ang peptides [Ang II and Ang-(1-7)] and their effect on BRS and mitochondrial function. The transgenic (mRen2)27 rat strain which overexpresses the murine Ren2 gene is hypertensive and has impaired BRS for control of heart rate (HR) with high levels of Ang II relative to Ang-(1-7) in the brain medullary tissue compared to the normotensive Sprague-Dawley (SD) rats [17, 18]. In contrast, transgenic rats with low glial angiotensinogen (ASrAOGEN) have lower mean arterial pressure (MAP) and HR suggesting decreased

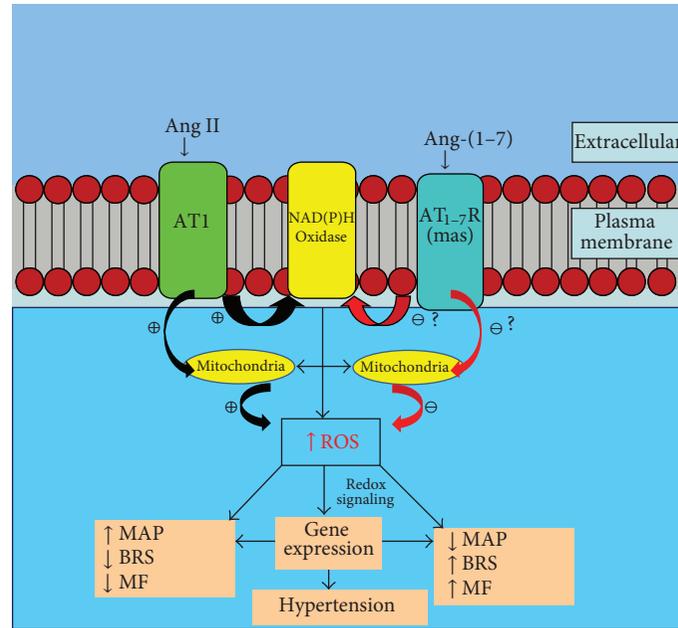


FIGURE 1: Proposed model: Ang-(1-7) through indirect (NADPH oxidase-mediated ROS) and/or direct interactions with mitochondria can attenuate ROS in dorsal medulla resulting in reduced blood pressure and enhanced baroreflex and mitochondrial function. ROS: reactive oxygen species; MAP: mean arterial pressure; BRS: baroreflex sensitivity; MF: Mitochondrial function; ?: not known.

(mRen2)27	Sprague-dawley	ASrAOGEN
Overexpress murine mRen2 gene	Parent control strain	Overexpress antisense oligonucleotide to AOGEN
Hypertensive	Normotensive	Hypotensive
High Ang II/Ang-(1-7) in medullary tissue		Low Ang II/Ang-(1-7) in medullary tissue
Low BRS compared to control SD		High BRS compared to control SD

FIGURE 2: Major characteristics of (mRen2)27, Sprague-Dawley (SD), and ASrAOGEN rat strains with respect to hemodynamic and baroreflex function. AOGEN: angiotensinogen, BRS: baroreflex sensitivity, RAS: renin-angiotensin system.

sympathetic nerve activity and enhanced BRS for control of HR relative to SD rats [19, 20]. While we would expect that both Ang II and Ang-(1-7) would be reduced in the glial cells, nonglial sources (neuronal and/or circulating) of angiotensinogen and Ang peptides appear to be intact in ASrAOGEN rats [21]. Tissue levels of Ang II relative to Ang-(1-7) in the medulla have not been reported; however, blockade of Ang II actions by an AT1 receptor antagonist revealed that there was no Ang II tone attenuating the BRS in anesthetized ASrAOGEN rats and we conclude that glia-derived Ang II is responsible for this action. In contrast, since blockade of endogenous Ang-(1-7) attenuates BRS in both younger and older anesthetized ASrAOGEN rats, a nonglial source of Ang-(1-7) likely contributes to the preservation of BRS in these animals [5, 22]. Thus, there appears to be low Ang II but maintenance of Ang-(1-7) tone contributing to the enhanced BRS seen in the medulla of these animals.

2. Angiotensin Peptides and ROS Generation in the Brain

Overactivation of the RAS in pathological conditions, such as hypertension, results in excessive ROS production through the prooxidant actions of Ang II [24, 25]. The contribution of cytoplasmic NADPH-oxidase (NOX)-generated ROS by Ang II in neurogenic hypertension is well established [26–28]. Ang II also stimulates mitochondrial ROS; both as a result of cytoplasmic NOX-derived ROS or direct effects on mitochondria [29–32]. Scavenging mitochondrial ROS, through agents such as Mito-TEMPO that preferentially targets the mitochondria, prevents Ang II-induced hypertension in mice [31, 33]. Antioxidant therapies targeting mitochondria are suggested to disrupt the mitochondrial ROS-dependent stimulation of cytoplasmic NOX activity, thereby providing beneficial effects in hypertension [34].

However, recent studies show that the NOX isoforms are also present within mitochondria [35–37] and the contribution of mitochondrial NOX to overall mitochondrial ROS in hypertension is unknown. Molecular interventions to target specific NOX isoforms within the mitochondria or other cellular organelle are required to address this issue.

Ang-(1–7) has emerged as a major counter-regulatory peptide to Ang II actions and may serve to inhibit Ang II-stimulated ROS production through inhibiting NOX and/or increased ROS scavenging by augmenting antioxidant enzymes such as catalase [38, 39]. Indeed we find that higher Ang II actions relative to Ang-(1–7) in the brain dorsal medulla of hypertensive (mRen2)27 rats are associated with increased cytoplasmic NOX activity and ROS in isolated brain dorsal medullary mitochondria compared with the hypotensive ASrAOPEN [with higher Ang-(1–7) actions relative to Ang II] or the normotensive SD rats [40]. The levels of ROS were similar in the ASrAOPEN rats compared to SD rats under basal conditions suggesting that Ang-(1–7) may serve to inhibit NOX and/or activate antioxidant enzymes in response to Ang II stimulation. Although this concept is supported by several studies [38, 41], it has not been investigated directly in the brain to our knowledge. Thus, in this respect, it would be interesting to test whether blockade of endogenous Ang-(1–7) in ASrAOPEN rats will result in increased NOX activity/ROS levels in response to Ang II infusion/microinjection in the brain.

3. ROS and Redox-Signaling in the Brain: Influences on Blood Pressure, Baroreflex, and Mitochondrial Function

Excessive ROS in brain contributes to increased sympathetic outflow [42, 43] and impairs mitochondrial [8, 31, 44, 45] and BRS function [46–48]. Our recent studies find that Ang-(1–7) via chronic ICV infusion improved vagal function independent of any blood pressure lowering effect in transgenic hypertensive (mRen2)27 rats [49]. This effect was in contrast to the response to AT₁ receptor antagonist, candesartan, which normalized blood pressure but did not significantly improve the vagal indices of BRS or HRV. We have yet to determine the effect of these treatments on mitochondrial ROS, but neither treatment altered cytoplasmic NOX activity. Central infusion of the ROS scavenger tempol did not lower blood pressure or influence indices of baroreflex function, but significantly reduced cytoplasmic NOX activity, suggesting independence from ROS-related mechanisms for blood pressure lowering and autonomic nervous system balance in the hypertensive (mRen2)27 strain. However, we do not know whether tempol efficiently targets mitochondrial ROS or the extent that alterations in mitochondrial ROS would influence blood pressure and/or BRS in the transgenic rats. Dikalova and colleagues have reported blood pressure lowering effects of Mito-TEMPO in both Ang II-induced and DOCA salt hypertension in mice while a similar dose of tempol alone did not lower blood pressure in this study [33]. Furthermore, mitochondria are in close structural proximity to the endoplasmic reticulum (ER), and ER stress is implicated in

mitochondrial dysfunction [50]. Indeed, the recent study by Young and colleagues link Ang II-induced hypertension to ER and oxidative stress in the brain [51]. These results provide a compelling case to investigate the effects of mitochondrial ROS, independent of cytoplasmic NOX.

AngII/AT₁ receptor/NOX-derived ROS are implicated in the activation of the MAP Kinases (MAPK) p38 and ERK1/2 that contribute to an impaired BRS and the pressor effects of Ang II in the RVLM [52–54]. A role for AT₁ receptors and MAPKs in activation of mitochondrial apoptotic pathways in neural regulation of blood pressure and BRS is also apparent [54]. However, hypertensive (mRen2)27 rats which show an increased NOX activity in the brain dorsal medulla but not activated p38, ERK1/2, or JNK-1 in comparison to SD rats suggesting a lack of association of MAPK signaling pathways with high blood pressure or oxidative stress [40]. In contrast, (mRen2)27 rats have an upregulated phosphoinositol 3 kinase (PI3 K) pathway that contributes to the elevated MAP and impaired BRS [55]. Hypotensive ASrAOPEN rats with normal NOX activity exhibit reduced levels of phosphorylated ERK1/2 and JNK-1 but not p38 in the brain dorsal medulla [40]. These animals have significantly higher expression of MAPK phosphatase-1 [MKP-1, a negative regulator of MAPK signaling [40]] supporting the concept that Ang-(1–7) increases regulatory phosphatases that may buffer against acute Ang II-stimulated signaling. Indeed, ASrAOPEN rats show greater impairments in the BRS for control of HR following acute solitary tract nucleus inhibition of protein tyrosine phosphatase 1b (PTP1b), a negative regulator of the PI3 K pathway, suggestive of increased expression and/or activity of this phosphatase within the dorsal medulla (Figure 3(a)). However, protein expression of total PTP1b (phosphorylated and nonphosphorylated forms) is similar in the dorsal medulla among the three rat strains under baseline conditions (Figure 3(b)). Therefore, given the functional differences observed following inhibition of PTP1b activity, quantification of the phosphorylated active form of PTP1b is necessary to confirm increased PTP1b activity in the ASrAOPEN rats. Differences in ROS [higher in (mRen2)27 versus SD or ASrAOPEN] or the upstream-regulatory kinases/phosphatases can modulate the phosphatase activity by changes in phosphorylation status at a number of different sites, despite lack of changes in the total protein [56]. While an upregulation of phosphatase expression and activity within the dorsal medulla may contribute to the enhanced resting BRS in the ASrAOPEN animals relative to the normal baroreflex function in SD rats [23], the lack of endogenous PTP1b tone in transgenic (mRen2)27 rats (Figure 3(a)) could result in increased PI3 K activity that contributes to an impaired BRS and increased MAP in these animals [55].

An interesting paradox to the beneficial role of these regulatory phosphatases is that both MKP-1 and PTP1b have negative effects on metabolic function [57, 58]. In this regard, global knockdown of these phosphatases improves insulin-sensitivity and prevents diet-induced obesity [59, 60]. MKP-1 is suggested to impair mitochondrial biogenesis in skeletal muscle in response to a high-fat diet through negative regulation of the p38 MAPKs [61]. However, ASrAOPEN rats that

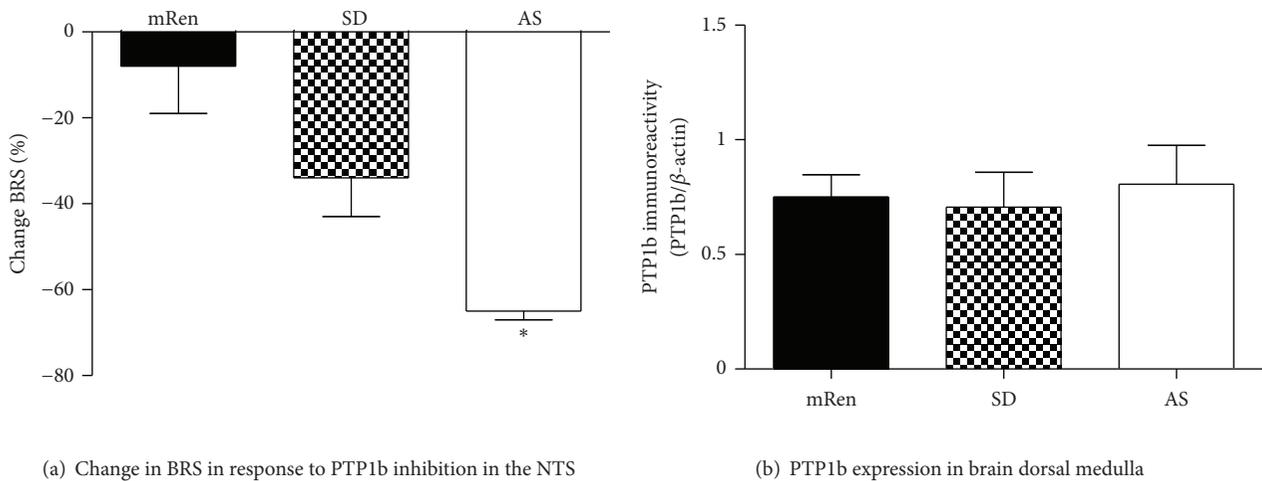


FIGURE 3: ASrAOGEN (AS) rats show significantly greater reduction in the evoked BRS for control of HR following inhibition of PTP1b (a), despite similar levels of the PTP1b protein in dorsal medulla of the three strains (b). Mean \pm SEM ($n = 4-6$ per group for solitary tract nucleus (NTS) microinjections, $n = 6$ for Western blotting); * $P < 0.05$ versus (mRen2)27 rats. Data replotted from [23] for the Sprague-Dawley (SD) rats and original data presented for ASrAOGEN and (mRen2)27 [mRen] rats. Western blotting carried out as published for SD rats [23] and original data presented for ASrAOGEN and (mRen2)27 rats. Note the PTP1b antibody recognizes both phosphorylated and nonphosphorylated forms.

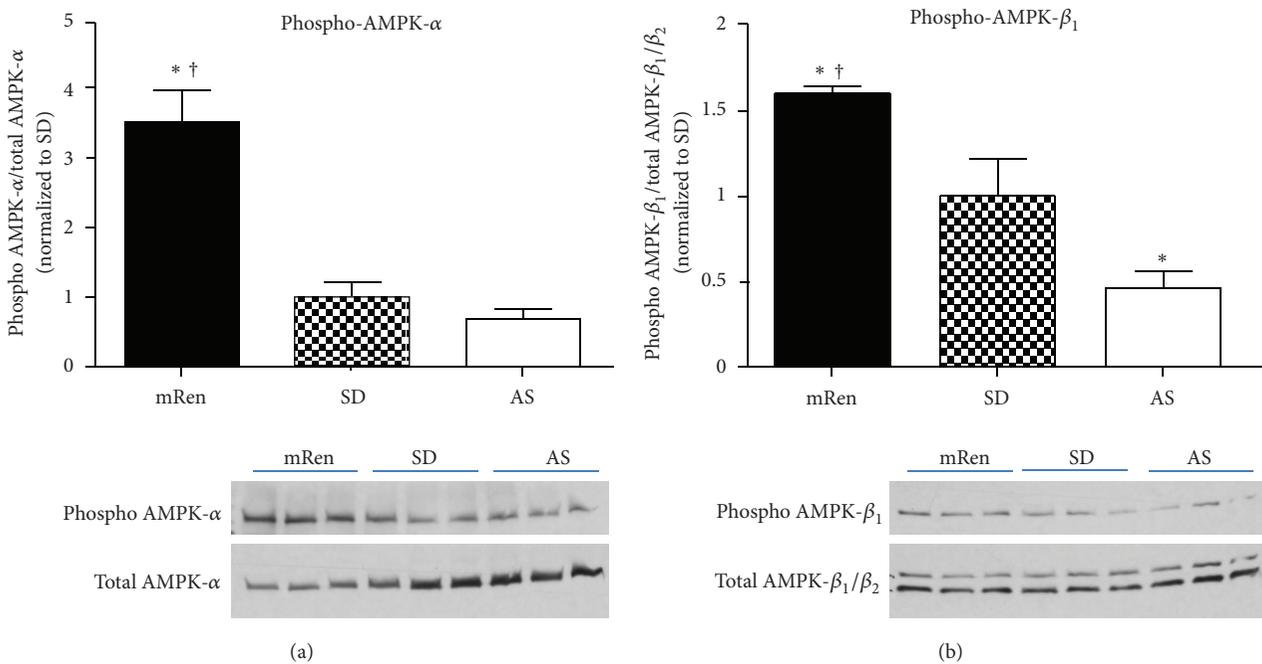


FIGURE 4: Hypertensive (mRen2)27 rats show significantly increased phosphorylated AMP-Kinase (AMPK) in the brain dorsal medulla. AMPK- α (a) and β_1 (b) activities were measured by Western blot hybridization using phospho-specific antibodies (Cell Signaling) in brain dorsal medulla tissues from (mRen2)27 [mRen], Sprague-Dawley (SD), and ASrAOGEN (AS) rats. Top: Densitometry analyses of phosphorylated protein levels normalized to total AMPK- α and β_1/β_2 ; bottom: representative Western blots. Data are mean \pm SEM ($n = 3-6$ per group); * $P < 0.05$ versus SD; † $P < 0.05$ versus AS rats.

have increased activity of these phosphatases at least in dorsal medulla are resistant to diet-induced obesity and spared age-related decline in cardiovascular and metabolic functions [62, 63]. These animals have increased life-span and their phenotype mimics animals with long-term RAS blockade

where improved mitochondrial function is reported [7, 62-64]. Whether the brain-specific actions of these phosphatases contribute to the beneficial metabolic effects in ASrAOGEN rats is of interest and currently unknown. Thus, further studies dissecting the role of these brain signaling pathways

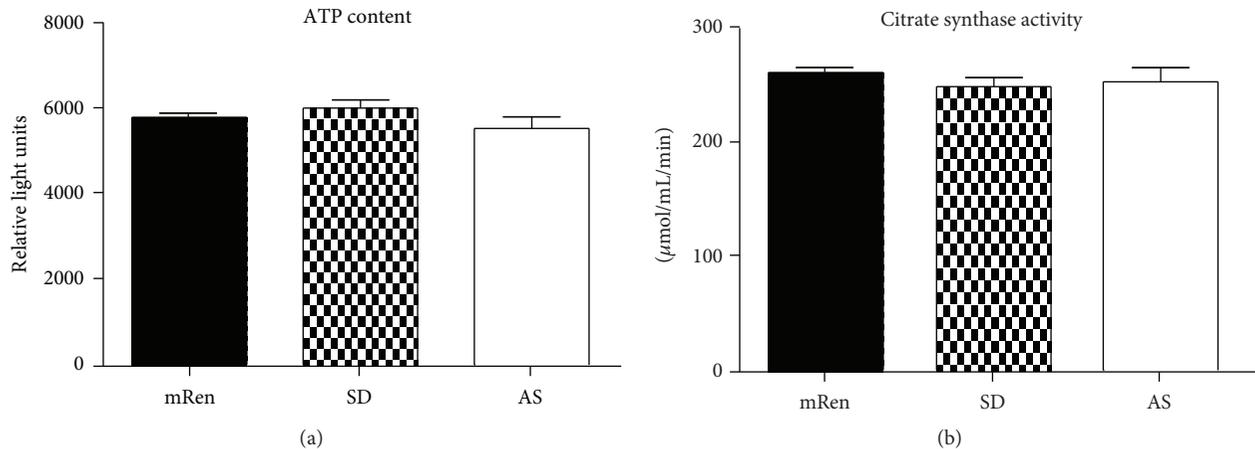


FIGURE 5: Brain dorsal medullary ATP levels (a) and mitochondrial content as measured using citrate synthase (b) was not different among hypertensive (mRen2)27 [mRen], Sprague-Dawley (SD) or hypotensive ASrA0GEN (AS) rats. ATP levels were determined using a chemiluminescent assay (Promega) and citrate synthase activity was measured using the assay kit (Sigma) in tissue homogenates. Mean \pm SEM ($n = 6$ per group).

in regulating MAP, BRS, mitochondrial, and metabolic functions are warranted.

Altered mitochondrial oxidant and/or energy levels are associated with a stimulated AMP-activated protein kinase (AMPK) pathway that is activated in response to depleted cellular energy levels, to restore mitochondrial biogenesis and ATP levels [65, 66]. AMPK was significantly activated (phosphorylated AMPK- α and β_1 subunits, Figures 4(a) and 4(b), resp.) in the dorsal medulla of transgenic (mRen2)27 rats that exhibit increased cytoplasmic NOX activity and ROS levels in the brain dorsal medullary mitochondria relative to SD rats [40]. While we expected lower ATP levels (Figure 5(a)) and the mitochondrial content/number (assessed indirectly using the marker of mitochondrial health or activity, citrate synthase enzyme activity, Figure 5(b)) in the (mRen2)27 rats, these markers were not different in the dorsal medulla of the three strains. Therefore, activation of AMPK in the (mRen2)27 rats may represent a compensatory response to restore normal ATP and mitochondrial activity in the hypertensive strain in the face of increased ROS. Additional studies are necessary to address whether (1) blockade of AMPK activation lowers mitochondrial content and depletes ATP levels and (2) targeting mitochondrial ROS improves MAP, BRS, and mitochondrial function in the hypertensive (mRen2)27 rats.

4. Conclusions and Perspectives

Mitochondria-derived ROS which often accompanies impaired autonomic function is an emerging therapeutic target in hypertension [31, 33, 34, 45–47]. Increased cellular ROS may manifest as impaired BRS for the control of HR and reduced HRV (decreased parasympathetic outflow or vagal tone); and these indices of autonomic imbalance are associated with increased overall mortality, independent of blood pressure. Therefore, determining the key cellular mechanisms underlying the beneficial actions of Ang-(1–7)

(such as altered kinase-phosphatase signaling) in influencing baroreflex function may help elucidate new therapeutic targets for reducing cardiometabolic pathologies. While Ang-(1–7) has been investigated for its role in attenuation of ROS, studies specifically addressing the mitochondria are lacking and few investigators are studying the interactions in brain. Thus, targeting improved vagal and mitochondrial function in addition to MAP may provide better target organ protection than lowering blood pressure alone, leading to reductions in all cause mortality.

Acknowledgments

Funding from COSEHC Warren Trust (MN), American Heart Association Postdoctoral Fellowship (MN), Farley Hudson Foundation of Jacksonville, NC, and National Institutes of Health (Drs. Diz and Chappell: HL-51952 and HL-56973) provided support for these studies.

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Review Article

The Angiotensin-Melatonin Axis

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Received 30 October 2012; Revised 18 December 2012; Accepted 19 December 2012

Academic Editor: Patrick Vanderheyden

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Accumulating evidence indicates that various biological and neuroendocrine circadian rhythms may be disrupted in cardiovascular and metabolic disorders. These circadian alterations may contribute to the progression of disease. Our studies direct to an important role of angiotensin II and melatonin in the modulation of circadian rhythms. The brain renin-angiotensin system (RAS) may modulate melatonin synthesis, a hormone with well-established roles in regulating circadian rhythms. Angiotensin production in the central nervous system may not only influence hypertension but also appears to affect the circadian rhythm of blood pressure. Drugs acting on RAS have been proven effective in the treatment of cardiovascular and metabolic disorders including hypertension and diabetes mellitus (DM). On the other hand, since melatonin is capable of ameliorating metabolic abnormalities in DM and insulin resistance, the beneficial effects of RAS blockade could be improved through combined RAS blocker and melatonin therapy. Contemporary research is evidencing the existence of specific clock genes forming central and peripheral clocks governing circadian rhythms. Further research on the interaction between these two neurohormones and the clock genes governing circadian clocks may progress our understanding on the pathophysiology of disease with possible impact on chronotherapeutic strategies.

1. Introduction

The renin-angiotensin system (RAS) is considered as a major endocrine regulator of cardiovascular homeostasis. The RAS acts in endocrine, paracrine, and autocrine manner in several organs and systems exercising various organ-specific actions with effects on the cardiovascular system [1]. Several lines of evidence from integrative physiology and functional genomics to molecular and genetic levels indicate that the RAS, circulating (endocrine) or tissue (paracrine and autocrine), is one of the major drivers of hypertension and cardiovascular diseases [1–3]. This knowledge led to the successful development of drugs to block the RAS system (angiotensin converting enzyme inhibitors, angiotensin receptor blockers, and renin inhibitors) that proved efficacious in the treatment of hypertension and other cardiovascular diseases [4].

Melatonin is produced by the pineal gland predominantly during night and it is considered as a major hormone regulating the circadian rhythmicity of several biological systems [5]. Research on melatonin functions revealed that this is not only a regulator of the biological circadian clock [5] but also it has a variety of biological functions [6]. Melatonin appears to be involved in various diseases, such as sleep disorders, dementia, mood disorders, cancer, and diabetes [7].

Both angiotensin and melatonin are synthesized in the brain. Angiotensin produced locally in central nervous system in nuclei involved in cardiovascular and fluid-electrolyte homeostasis interacts with other systems, such as sympathetic, vasopressinergic ones [1, 8]. Moreover, there is a local pineal RAS that modulates the synthesis of melatonin, which represents the main hormonal output of the pineal gland [9, 10]. The RAS is classically involved in cardiovascular and metabolic pathophysiology while melatonin deals with

circadian rhythms. In this paper, we aimed at evidencing interference at several levels between the RAS and melatonin to modulate cardiovascular and metabolic pathophysiology (Table 1).

2. Angiotensin-Melatonin Axis

The postulation of a local RAS in the brain has led to discovery of brain-specific roles of angiotensin II (Ang II). A local production of active angiotensins has been documented in several brain nuclei and regions. Of these, the two neuroendocrine glands situated in the brain possess high levels of Ang II-forming activities [9]. This led us to postulate the presence of a local RAS in the pineal gland. Angiotensinogen, the precursor of the RAS, has been identified in pineal glial cells while the receptors type AT1b are localized in pinealocytes [10]. As part of the enzymatic cascade producing Ang II, we identified angiotensin converting enzyme (ACE) and chymase but not renin, indicating the existence of nonrenin pathways [9, 10]. By utilizing both pharmacological and transgenic strategies, we could demonstrate that locally produced Ang II in the pineal can modulate the melatonin synthesis [11]. Melatonin represents the main hormonal output of the pineal gland and it is considered as an important modulator of circadian rhythms. Our studies indicate that Ang II acts on the pinealocyte AT1 receptors to influence the synthesis and activity of tryptophan hydroxylase (TPH), the rate-limiting enzyme of melatonin synthesis. The demonstration of a functional pineal RAS interfering with melatonin synthesis indicates that this may affect melatonin roles, such as in modulation of circadian rhythms (Figure 1).

The circadian system comprised of a group of specialized genes is a key integrator of metabolism and behavior that synchronizes physiological processes. Circadian oscillators are present not only in the suprachiasmatic nucleus (SCN) which is considered to be the master clock but also in peripheral tissues, including cardiovascular organs [12]. Several animal studies are identifying roles for clock genes in cardiovascular and metabolic physiology and pathophysiology [13]. Genetic manipulation of clock genes in transgenic mouse models has uncovered new functions of internal clocks in pathogenesis of cardiovascular diseases [14]. Since our and other studies point to an important role of the RAS in the modulation of circadian rhythms of blood pressure [15, 16], an interaction with the genes governing circadian clock of cardiovascular tissues might be conceivable. An interaction between the RAS and the circadian system has been suggested to contribute to the development of inverted BP profile in transgenic rats harboring the mouse Ren-2 renin gene, TGR(mREN)²⁷ [17, 18].

The interaction between angiotensin and melatonin both centrally and peripherally in hypertension and cardiovascular diseases has to be studied. At pineal level, the challenge is to understand in a hypertensive patient with elevated levels of angiotensin how melatonin release is affected: is there a feedback loop of high levels of circulating angiotensin impacting on either potentiating more melatonin release to counteract hypertension, or if there is a lack of effect, that is,

TABLE 1: Opposing roles of angiotensin II and melatonin in cardiovascular and metabolic pathophysiology.

Angiotensin	Melatonin
Blood pressure: direct effects	
Increase, vasoconstriction	Decrease, vasodilation
Blood pressure: circadian rhythm	
Nondipper/riser hypertension	Decreased levels in nondipper hypertension, chronobiotic
Central clock: suprachiasmatic nucleus	
Precursor and receptors present	Receptors present
Sympathetic nervous system	
Stimulation	Sympatholytic
Oxidative stress	
Increase	Decrease
Inflammation	
Increase	Decrease
Central clock: suprachiasmatic nucleus	
Precursor and receptors present	Receptors present
Insulin	
Insulin resistance	Increase in insulin sensitivity

resistance to central angiotensin, that resets the pineal gland axis and reduces melatonin output, which then may further potentiate hypertension. Further studies are necessary to decipher and dissect possible interactions between the RAS, melatonin, and the clock genes governing circadian clock.

3. Antagonistic Effects of Angiotensin and Melatonin in Cardiovascular and Metabolic Systems

3.1. Cardiovascular System. A cardiovascular role of melatonin has been suggested already 40 years ago by the description of a pinealectomy-induced experimental hypertension model [19–21]. Melatonin receptors are present in the vasculature and mediate vascular constriction and vasodilation through MT1 and MT2 receptors, respectively [22]. Melatonin administration generally induces a decrease in blood pressure [22]. One possible mechanism contributing to the melatonin hypotensive effects is through its sympatholytic properties [23]. On the other hand, a marked reduction of circulating melatonin has been observed in cardiovascular diseases [24]. These findings suggest antagonistic activities of angiotensin and melatonin in the cardiovascular system (Table 1). The mechanisms by which melatonin is antagonizing Ang II actions in cardiovascular and metabolic diseases are comprising its antihypertensive, antioxidant, and anti-inflammatory functions [24]. Melatonin has direct free radical scavenging and indirect antioxidant activity. Through these marked antioxidant properties, melatonin has cardioprotective effects, in particular in myocardial damage after ischemia-reperfusion [25].

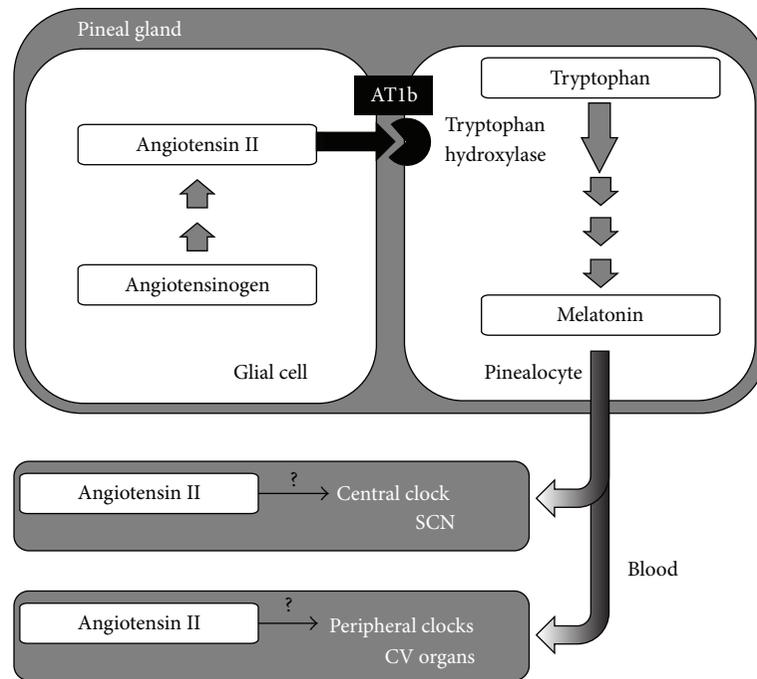


FIGURE 1: Pineal renin-angiotensin system (RAS) interacts with melatonin synthesis. Angiotensin II, produced from angiotensinogen produced by glial cells, acts on AT1b receptors present on pinealocytes to stimulate tryptophan hydroxylase, which is the rate-limiting enzyme in melatonin synthesis. Both angiotensin and melatonin may interact to regulate rhythmicity either centrally in the suprachiasmatic nucleus (SCN) or peripherally in clocks present in several cardiovascular organs.

3.2. Circadian Rhythms of the Cardiovascular System. Multiple clinical studies have implicated blood pressure (BP) and heart rate (HR) variability in the diagnosis and prognosis of arterial hypertension and cardiovascular diseases. In healthy individuals, there is a circadian variation of BP with a nocturnal fall of 10%–20% during the sleep period [26]. In hypertensive patients, this circadian rhythm may disappear or even become inverted. Therefore, according to the BP circadian alterations, patients have been classified as “dippers” when the mean nighttime BP is $\geq 10\%$ lower than the mean daytime BP, as “nondippers” when the reduction is $< 10\%$ or as “risers” when it is higher [27]. Nondippers and risers are at an increased risk for target organ damage and cardiovascular events [28, 29]. Moreover, a circadian pattern becomes quite obvious in the occurrence of acute cardiovascular diseases, such as ischemia, infarction, stroke, and sudden death, and new chronotherapeutic approaches in antihypertensive therapy are trying to exploit the knowledge of circadian rhythms in order to reduce these events [29]. Therefore, a better understanding of the molecular biology and pathophysiology of nondipper hypertension will lead to a better understanding of the disease and possibly lead to new diagnostic tools or therapeutic strategies. Further studies investigating the molecular mechanisms of the circadian regulation of the cardiovascular system should hopefully reveal new diagnostic tools or treatment algorithms for disease.

Our group was the first to do demonstrate that chronic Ang II infusion may induce a shift in the circadian BP rhythm

(Figure 2) [16]. Ang II infused subcutaneously at doses of up to 250 ng/kg per minute that does not produce direct vasoconstriction is described as “slow pressor” or “subpressor” and can induce a gradual increase of BP. Chronic infusion (days to weeks) of subpressor Ang II subcutaneously induces nondipper hypertension similarly with the renovascular and other forms of human hypertension where the circadian variation of blood pressure is altered [16, 30, 31]. Alterations in the circadian BP rhythm are not synchronized with alterations of heart rate or locomotor activity, contributing to the concept that the circadian variability in blood pressure and heart rate are differentially regulated [15, 16]. We further hypothesized that the brain RAS might be involved in the Ang II-induced BP circadian shift. To test this hypothesis, we studied a transgenic rat that has reduced angiotensinogen levels in the brain through expression of an antisense RNA against angiotensinogen, induced by means of the astrocyte-specific glial fibrillary acidic protein promoter [32]. Ang II infusion in the TGR(ASrAOGEN) transgenic rats did not induce a BP circadian shift, indicating that peripheral RAS interacts with the brain RAS to induce not only hypertension [2] but also a BP circadian shift [15, 16]. We employed the TGR(ASrAOGEN) to investigate if the brain RAS is involved in circadian rhythm reentrainment to light phase shifts. The BP and HR acrophases (peak time of curve fitting) in TGR(ASrAOGEN) rats readjusted to light shifts significantly slower than in control (Sprague-Dawley) rats [15, 16]. However, the acrophases of locomotor activity changed similarly in both strains. These data suggest that

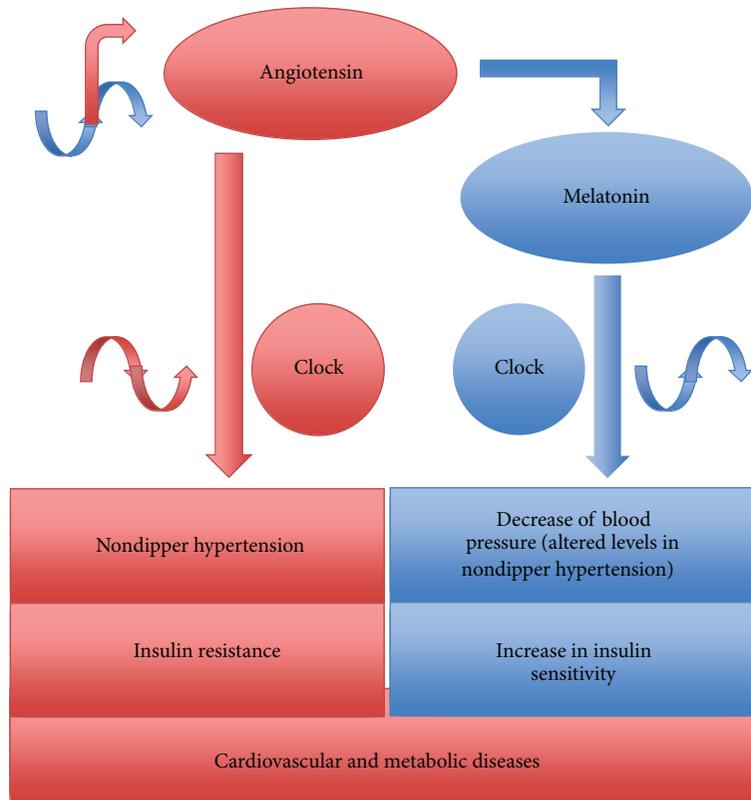


FIGURE 2: Angiotensin versus melatonin in cardiovascular and metabolic diseases. An increase in angiotensin induces nondipper/riser hypertension, which is characterized by a decrease in melatonin. Angiotensin and melatonin have opposing effects on insulin sensitivity.

treatment with RAS blockers with high penetrability of the blood-brain barrier (such as candesartan and valsartan) could slow the resynchronization of cardiovascular system in jet-lag conditions of travelers adapting to a new time zone.

Research on the circadian actions of Ang II has pre-occupied several research groups. Among these, the group of Lemmer has provided several lines of evidence on the importance and the mechanisms regulating cardiovascular circadian rhythms [29]. Lemmer et al. provided significant insights in the pathophysiology of a transgenic rat model, TGR(mREN2)27 [17, 18]. The TGR(mREN2)27 is a well-characterized model of malignant hypertension due to an overactive RAS as it harbors the mouse salivary gland renin gene (mREN2). The TGR(mREN2)27 not only become hypertensive and develop target-organ damage but also exhibit an inverted circadian rhythm of BP, which makes them a valuable model to further study the molecular biology of circadian rhythms. However, detailed mechanisms responsible for the Ang II-induced nondipper hypertension are still not well understood. The circadian rhythms that are a characteristic of most of the physiological parameters are governed by biological clocks. Ablation of the SCN of the hypothalamus that serves as the main zeitgeber for such circadian rhythms eliminates BP, HR, and locomotor activity [33, 34]. Ang II might interact directly or through melatonin to influence the 24-h rhythmic expression of clock genes in SCN (Figure 1). Evidence on a mutual relationship between melatonin and circadian oscillators has recently

been reviewed by Hardeland [35]. Also, an impaired nocturnal melatonin secretion has been detected in nondipper hypertensive patients (Figure 2) [36]. Besides melatonin, the brain RAS might influence the central circadian clock present in the hypothalamic SCN either directly or through vasopressin that is another hormone with demonstrated roles in circadian rhythms [8, 11, 37].

Ang II and melatonin might interact with circadian oscillators present not only in the SCN but also in peripheral tissues, including cardiovascular organs (Figure 1) [12]. Several animal studies are identifying roles for clock genes in cardiovascular physiology [13]. Genetic manipulation of clock genes in transgenic mouse models has uncovered new functions of internal clocks in pathogenesis of cardiovascular diseases [14]. Recent evidence is suggesting that a disruption of central or peripheral clocks may contribute to the progression of cardiovascular diseases [38]. Researching for further insights on the roles of biological clocks in cardiovascular organs shall provide acumens into the relevance of the circadian rhythms in cardiovascular pathology.

3.3. Metabolic Syndrome, Insulin Resistance, and Diabetes.

Several lines of evidence including successful therapies with drugs acting on RAS are demonstrating roles of RAS in diabetes mellitus and metabolic syndrome. Putnam et al. recently reviewed the accumulating evidence describing the RAS as a “target of and contributor to dyslipidemias, altered glucose homeostasis, and hypertension of the metabolic

syndrome" [39]. Angiotensin II causes insulin resistance through activation of the AT1R and increased production of mineralocorticoids [40]. However, the underlying mechanisms of Ang II leading to insulin resistance remain to be fully elucidated. Melatonin on the other hand induces an increased insulin sensitivity [35]. Evidence for a link between melatonin and insulin came from pinealectomized animals that develop diabetogenic syndrome characterized by insulin resistance and a 50% reduction of GLUT4 in adipose and muscular tissue [41]. Moreover, it was demonstrated that the absence of melatonin in pinealectomized animals impairs the temporal organization of several metabolic functions associated to the carbohydrate metabolism, such as daily insulin secretion, adaptation to starvation, and exercise [42–44]. This dramatic picture can be partially or totally restored by melatonin reposition or restricted feeding [45, 46]. Melatonin acting through MT₁ membrane receptors is able to induce insulin receptor phosphorylation at the same time that mobilizes several intracellular transduction steps in common to insulin signaling [47]. Melatonin is able to restore insulin sensitivity and regulates food ingestion and body weight and abdominal adiposity in old rats [48].

Not only melatonin regulates insulin but also insulin can act on *in vitro* pineal glands potentiating the noradrenergic-induced melatonin synthesis, regulating the activity of the enzymes tryptophan hydroxylase and N-acetyltransferase through after-transcriptional mechanisms [49].

The first reports on diabetes and melatonin production showed that diabetic rats and mice, chemically-induced by alloxan (ALX) or streptozotocin (STZ), presented a decrease in melatonin synthesis and plasma levels [50, 51], although Champney et al. [52] reported no alterations in the level of melatonin in diabetic rats. Melatonin was also observed to suppress the onset of type 1 diabetes in nonobese mice, while pinealectomy had the opposite effect [53]. In type 2 diabetic patients and diabetic Goto Kakizaki rats a decreased serum melatonin level was observed [54]. In STZ-induced diabetic animals, melatonin was shown to decrease serum lipid oxidation [55] and protein glycosylation [56], as well as regulate the activity of antioxidant enzymes, improving the protection against the oxidative damage caused by diabetes [57–59]. Despite that, melatonin was not able to normalize hyperglycemia and/or body weight in these animals [55, 60, 61] and lower levels of the indolamine were observed in peripheral tissues like pancreas, kidney, spleen, and duodenum [62].

4. Conclusions and Perspectives

Accumulating evidence suggests that not only angiotensin interferes with melatonin synthesis and release but also both hormones interact at several levels having opposing effects in cardiovascular and metabolic pathophysiology (Table 1). Furthermore, evidence is indicating that not only melatonin but also angiotensin may interfere with circadian rhythms. The intimal regulatory mechanisms of interference between the two systems both centrally and peripherally, at synthesis and action levels, in homeostatic and disease conditions await

further investigation. On how peripheral RAS interacts with the pineal RAS in hypertension and cardiovascular disease and if the sympathetic nervous system or other systems are involved in this interaction are still open questions.

Contemporary progress in chronobiology directs to an important role of clock-associated genes in the progression of cardiovascular and metabolic diseases. Since angiotensin appears to be involved in the modulation of circadian rhythms of blood pressure, an interaction with the clock genes seems likely. Therefore, we believe that further research on the molecular biology of circadian alterations involving interactions between angiotensin, melatonin, and clock genes may have an impact on cardiovascular and metabolic pathophysiology leading to new chronotherapeutic strategies.

Acknowledgment

The authors acknowledge the financial support from the State of São Paulo Research Foundation (FAPESP) (Grants 20011/51410-9 and 2011/50078-0).

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Review Article

RAS in Pregnancy and Preeclampsia and Eclampsia

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Received 22 November 2012; Accepted 14 December 2012

Academic Editor: Marc de Gasparo

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Preeclampsia is a common disease of pregnancy characterized by the presence of hypertension and commitment of many organs, including the brain, secondary to generalized endothelial dysfunction. Its etiology is not known precisely, but it involved several factors, highlighting the renin angiotensin system (RAS), which would have an important role in the origin of multisystem involvement. This paper reviews the evidence supporting the involvement of RAS in triggering the disease, in addition to the components of this system that would be involved and how it eventually produces brain engagement.

1. Introduction

Preeclampsia is a major complication of pregnancy and corresponds to a major cause of both maternal and fetal morbidity and mortality [1–3]. It is a condition that produces a compromise of many organs, including the brain causing seizures, a condition known as eclampsia [4, 5]. The pathophysiology is not well understood, but it involves different factors, such as genetic, immunological, and inflammatory [6, 7]. In recent years there is a series of studies linking the renin angiotensin system (RAS) with preeclampsia [8–10], in the sense that the alteration of this system would be involved in the pathogenesis of this disease, as this could trigger the different characteristics in this pathology, including brain involvement.

2. RAS in Normal Pregnancy

RAS is a system that functions as an important regulator of blood pressure, electrolyte balance, and fluid homeostasis [11]. This system comprises the inactive peptide angiotensinogen, which is converted to angiotensin I and then the active peptide angiotensin II (Ang II) through the action of renin and angiotensin-converting enzyme (ACE) [12]. Ang II exerts its action primarily through the AT1

receptor, located widely in different tissues, including the syncytiotrophoblast [10].

During pregnancy usually occurs overexpression of many components of the RAS, both in the blood and tissues. There is an increase in plasma renin mainly by extrarenal production [13]. There is also a higher-level production of angiotensinogen liver secondary to increased circulating estrogens. ACE is the only component that has been shown to decrease during normal pregnancy, but equally there is a higher plasma concentration of Ang II [8, 13].

There is an upregulation of RAS components during normal pregnancy, but there is also a decrease in sensitivity to Ang II, whereby these women are resistant to the pressor effect of this molecule, requiring twice Ang II by intravenous infusion compared with nonpregnant women to achieve a similarly vasomotor response [14].

It is thought that this might be related to the monomer structure of AT1 during uncomplicated pregnancies, unlike the heterodimeric structure observed in terms of sensitivity to Ang II [15]. In addition, estrogens produce a shift in the formation of angiotensin peptides, reducing the formation of Ang II and increasing the production of Ang-(1–7), which has a vasodilator role [16].

Furthermore, in addition to a systemic RAS, RAS also exists in uteroplacental territory [13]. This unit consists

of a placental portion, corresponding to fetal tissue, and a decidual, which is of maternal origin, and in both all components of the RAS are secreted. Therefore, there are 2 RAS systems: placental and decidual. The latter could be related to the pregnancy-associated vascular remodeling of the spiral arteries [17].

3. Pathophysiology of Preeclampsia: Role of RAS

Preeclampsia corresponds to a multisystem disorder characterized by increased peripheral vascular resistance, increased platelet aggregation, and systemic endothelial dysfunction [18]. Corresponds to a multifactorial disease, involving genetic and environmental components, a defective extravillous trophoblast invasion, an impaired immune tolerance between maternal, fetal and placental tissues and maternal inflammatory disorders [19, 20]. Clinically it is characterized by the presence of hypertension and proteinuria from the 2nd half of pregnancy, and the only effective treatment is the termination of pregnancy [21].

From a physiological point of view, preeclampsia is defined as a disease of two stages [22]. The first is the placental stage that occurs during the first 20 weeks of gestation. In this, the phenomena of remodeling of the vascular walls of the spiral arteries do not develop properly, resulting in abnormal placentation, thus prompting ischemic placenta [23]. The second stage occurs during the second half of pregnancy and is known as the systemic stage. This is the clinical stage of preeclampsia, in which there is an exaggerated maternal systemic inflammatory response and endothelial dysfunction as a central element [24–26]. Between these two stages are some mediators, which are understood as molecules released by the placenta and are capable of transmitting this placental damage and translate into a systemic involvement. Mediators most studied are oxidative stress, microfragments of syncytiotrophoblast (STBM), and antiangiogenic proteins [27].

There is a considerable amount of evidence supporting the role of angiogenic factors in triggering preeclampsia, and these are the tyrosine-like soluble factor (sFlt-1) and soluble endoglin (s-Eng) [28, 29]. These molecules bind to angiogenic proteins such as VEGF and prevent them from joining their membrane receptors on endothelial cells, leading to endothelial dysfunction [30]. It was observed that these factors are elevated about 6–8 weeks before the start of the clinical picture of preeclampsia, and their plasma concentrations are related to the severity of the disease [31, 32]. In animal models it has been found that inoculation of these can produce hypertension, proteinuria, and hepatic involvement, symptoms characteristic of preeclampsia [33, 34]. It is also observed that hypoxia causes increased secretion of these factors [35].

In patients with preeclampsia, dysregulation has been observed in the RAS compared to healthy pregnancies. The levels of renin, Ang I, and Ang II are lower than in uncomplicated pregnancies [36]. Despite this decrease in the expression of RAS components, in patients suffering from

preeclampsia increased sensitivity to Ang II exists, showing an exaggerated pressor response to Ang II.

4. AT1 Receptors Autoantibodies in Preeclampsia

In recent years there is a wealth of evidence supporting the AT1 autoantibodies (AT1-AA) in the pathogenesis of preeclampsia. These correspond to IgG autoantibodies that bind to a seven-amino acid sequence present on the second extracellular loop of the AT1 receptor [37, 38]. They are present in the plasma of patients with preeclampsia and are able to increase the beating rate of the cultured cardiomyocytes [39]. Many research papers show that these autoantibodies are elevated in patients with preeclampsia, but not in uncomplicated pregnancies.

In vitro and *in vivo* studies have determined its role in triggering preeclampsia. These antibodies bind to the AT1 receptors of different cell groups, triggering its pathological action [40]. It has been observed that in human trophoblast cells AT1-AA substances induce the generation of reactive oxygen species (ROS) intracellularly through NADPH oxidase activation [41]. In addition, these same cells stimulate the release of PAI-1, resulting in a decreased trophoblast invasiveness [42, 43], generating a defect of placentation. This increase in PAI-1 is also observed in mesangial cells, which can produce a decrease in extracellular matrix degradation and increased subendothelial fibrin deposition thereby determining renal damage leading to proteinuria and a decreased glomerular filtration rate [44]. It has also been observed that AT1-AA binds to endothelial and vascular cells, causing endothelial damage and vasoconstriction [45, 46].

All these actions could explain endothelial dysfunction, increased peripheral vascular resistance, and impaired coagulation system observed in preeclampsia.

In animal models it has been observed that the inoculation of AT1-AA from patients with preeclampsia is capable of reproducing the characteristics of the disease [47, 48]. Reports in rats showed that surgically induced placental ischemia may cause increased levels of AT1-AA and trigger hypertension and proteinuria [49–51]. It was also observed that these autoantibodies stimulate the release of sFlt-1 and s-eng by the placenta, key proteins in triggering endothelial dysfunction [52, 53].

In the same way pregnant human studies show that placental perfusion abnormality, evaluated with Doppler ultrasound of the uterine arteries, is associated with increased plasma concentrations of AT1-AA before the onset of the disease and that plasmatic levels of these autoantibodies correlate with the severity [54]. The concentration of these autoantibodies is higher in cases of severe preeclampsia, also having a linear correlation with proteinuria and hypertension. This is further confirmed by the fact that in milder cases, as moderate preeclampsia and gestational hypertension, levels of AT1-AA are higher than in normotensive pregnancies, but lower than in severe preeclampsia. Therefore, AT1-AA would be a key element in the pathogenesis of preeclampsia and modulate the secretion

of important factors responsible for the pathophysiology [55].

5. Eclampsia: Loss of Autoregulation of Cerebral Blood Flow and Possible Role of RAS

One of the most serious complications of preeclampsia is eclampsia, which corresponds to the presence of seizures in the context of a patient with hypertension and proteinuria [5, 56]. However, currently the eclampsia is being seen as a manifestation of a much larger entity than pregnancy, known as posterior reversible encephalopathy (PRES), which is produced by other conditions such as hypertensive encephalopathy or use of immunosuppressive drugs. In these cases there is a characteristic increase in blood pressure and/or alteration of endothelial permeability [57–60].

The PRES is characterized by the presence of well-defined signs and symptoms associated with the presence of specific neuroimaging. Among the symptoms are headache, nausea, vomiting, visual disturbances, and seizures [61, 62]. Diagnosis is by observation of symmetrical hyperintense lesions and bilateral parietooccipital level on MRI, suggestive of vasogenic edema [63].

The exact pathophysiology of PRES is not known with certainty, but is thought to be due to alterations of the vasculature and cerebral perfusion [64]. With the increase in blood pressure, brain responds with its vasculature vasoconstriction, which determines an increase of cerebral perfusion pressure (CPP). As this CPP remains persistently high, it will produce a pressure transmission to the distal small cerebral vessels, causing endothelial damage and muscle dysfunction of cerebral vascular territory [65, 66]. This phenomenon is known as barotrauma, corresponding to forced dilation of arterioles and distal opening of endothelial tight junctions, determining a disruption of the blood-brain barrier (BBB), which leads to increased permeability of this, resulting in a vasogenic edema [67].

However, not all cases of PRES relate to alterations in arterial pressure. One study shows that 16% of cases of eclampsia occur in normotensive patients, and only 13% of cases were associated with severe hypertension [68, 69]. Therefore, the loss of autoregulation of cerebral perfusion secondary to hypertension does not explain all cases of PRES, so that alteration of endothelial permeability by disruption of the BBB ought to play an important role [67]. A recent study shows that preeclampsia altered BBB permeability independently of blood pressure. In this report it is determined that plasma from patients with preeclampsia significantly increases the permeability of the BBB, compared to plasma from patients with normal pregnancies [70]. These findings support the concept of the pathophysiology of PRES that is determined by hemodynamic factors and factors altering endothelial function.

Various reports have identified the involvement of RAS in the blood-brain barrier disruption in other medical conditions [71]. Therefore, the presence of AT1-AA in patients with preeclampsia could play a key role in triggering PRES. It is shown that these autoantibodies produced an

increase in peripheral vascular resistance and hypertension, which leads to a systemic endothelial dysfunction via ROS secretion and antiangiogenic proteins [37, 40], so directly involved in the conditions that can trigger PRES.

6. Management of Preeclampsia and Eclampsia Prevention

Currently, the only effective treatment for preeclampsia is the termination of pregnancy, which in very early pregnancies may not be the best alternative, as this results in increased perinatal morbidity and mortality secondary to prematurity [72]. Therefore, it is important to seek management strategies from the physiological point of view, and in that sense, the management of RAS alteration appears to be a logical choice, given the strong evidence about an excessive AT1 receptor activation during preeclampsia.

Regarding eclampsia, the drug of choice for prevention and management is magnesium sulfate. This drug reduces the risk of seizures in patients with severe preeclampsia [73]. Its mechanism of action is not entirely clear, but it has been shown to be capable of reducing the CPP [74]. However, it should be administered intravenously and usually for 48 hrs, in patient hospitalized and only for short periods of time.

Many studies determined that the addition of losartan (AT1 receptors blocker) or neutralizing antibodies of the AT1-AA (7-amino acid peptide epitope or 7-aa) blocks the effect of AT1-AA, further confirming that the action of these antibodies is through activation of AT1 receptors (z). Animal studies with surgically induced placental ischemia demonstrate that the addition of these compounds significantly reduces arterial pressure [49], but this effect was not seen with ACE inhibitors [75]. In another report, AT1-AA purified from pregnant women was injected in mice. They triggered hypertension and proteinuria, were significantly reduced or abolished when administered losartan or 7-aa [47]. Adoptive transfer studies in pregnant mice have demonstrated that release of antiangiogenic factors and proinflammatory cytokines resulting from autoantibody-mediated receptor activation is blocked by the addition of 7-aa or AT1 receptor antagonist [10, 37].

The main problem with AT1 receptors blockers is that they increase the risk of fetal malformations, such as oligohydramnios, pulmonary hypoplasia, transient renal failure, preterm delivery, and Potter syndrome [76–78], so its use during pregnancy should be avoided. However, AT1 receptors blockers can be used postpartum, and considering that currently about 30% of eclampsia occur in postpartum [56, 79] and that AT1-AA levels can remain high for a long time [80], AT1 receptors blockers become an interesting strategy for the management of hypertension during the postpartum period and to decrease the incidence of eclampsia, as well as controlling blood pressure; they block the action of endothelial AT1-AA, which is still present in the postpartum period.

Neutralizing antibodies 7-aa have the advantage of not inhibiting the AT1 receptor; they only block the AT1-AA, without changing completely the action of Ang II.

Therefore, it seems a useful strategy for the management of preeclampsia. However, we must await the development of further studies to determine its safety during pregnancy.

7. Discussion

During preeclampsia there is an alteration of the RAS. The presence of AT1-AA determines triggering a series of actions in tissues and organs that would result in an increase in peripheral vascular resistance, altered coagulation, renal impairment, and systemic endothelial dysfunction. These alterations can generate the commitment of many organs, including the brain, producing a PRES.

Blocking the action of these autoantibodies using losartan or 7-aa substantially decreases the damage caused by AT1-AA, creating an opportunity for the management and prevention of complications of preeclampsia.

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Review Article

The Angiotensin II Type 2 Receptor in Brain Functions: An Update

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Received 31 October 2012; Accepted 29 November 2012

Academic Editor: Marc de Gasparo

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Angiotensin II (Ang II) is the main active product of the renin-angiotensin system (RAS), mediating its action via two major receptors, namely, the Ang II type 1 (AT₁) receptor and the type 2 (AT₂) receptor. Recent results also implicate several other members of the renin-angiotensin system in various aspects of brain functions. The first aim of this paper is to summarize the current state of knowledge regarding the properties and signaling of the AT₂ receptor, its expression in the brain, and its well-established effects. Secondly, we will highlight the potential role of the AT₂ receptor in cognitive function, neurological disorders and in the regulation of appetite and the possible link with development of metabolic disorders. The potential utility of novel nonpeptide selective AT₂ receptor ligands in clarifying potential roles of this receptor in physiology will also be discussed. If confirmed, these new pharmacological tools should help to improve impaired cognitive performance, not only through its action on brain microcirculation and inflammation, but also through more specific effects on neurons. However, the overall physiological relevance of the AT₂ receptor in the brain must also consider the Ang IV/AT₄ receptor.

1. Introduction

A major advance in the field of the renin-angiotensin system (RAS) was the discovery of a complete RAS in the brain, independent from the peripheral system, by Jacques Genest's laboratory in Montreal in 1971 [1] (for reviews see [2–4]). Further studies by Mendelsohn et al. [5, 6] and Unger et al. [7] corroborated these observations, using biochemical, pharmacological, and autoradiographic approaches. It was found that brain levels of angiotensin II (Ang II) are higher than its circulating levels, suggesting independence of the two systems (review in [3, 4]). The various components of RAS (angiotensin-converting enzyme (ACE), Ang II and Ang II receptors) are all found in the adult brain in areas involved in the regulation of fluid and electrolyte balance, in the regulation of arterial pressure and vasopressin release, and regulation of the autonomic system [8, 9]. They are also present in structures involved in cognition, behavior, and locomotion. In particular, over the past 10 years, several advances have been made regarding the role of Ang II in various brain functions, including cerebroprotection, stress, depression, and memory consolidation [3, 4, 10–12].

Moreover, many evidences highlight the potential function of Ang II in the etiology of certain neurodegenerative diseases, including Alzheimer's and Parkinson's disease, seizures, and the development of metabolic syndrome and diabetes (review in [4, 13, 14]).

In the classical view, synthesis of Ang II begins with the conversion of angiotensinogen into Ang I by the enzyme renin. Ang I is then converted by ACE into Ang II (a.a. 1–8), which is then metabolized to Ang III (a.a. 2–8) and Ang IV (a.a. 3–8). Ang II and Ang IV can be further converted into Ang (1–7) and Ang (3–7) (Figure 1). Ang II binds to two main receptors, namely, the angiotensin type 1 (AT₁) and type 2 (AT₂) receptors, both belonging to the G-protein coupled receptor (GPCR) family [4, 12, 13]. Aside from this common link, these two receptors otherwise carry very little similarities. Indeed, their actions are generally opposite and while the AT₁ receptor is expressed abundantly in several tissues, expression of AT₂ receptor is limited to specific tissues and brain areas, where its concentration is generally low compared to the AT₁ receptor. Most of the known effects of Ang II are due to activation of the AT₁ receptor, including vasoconstriction, cellular growth, and proliferation. On the

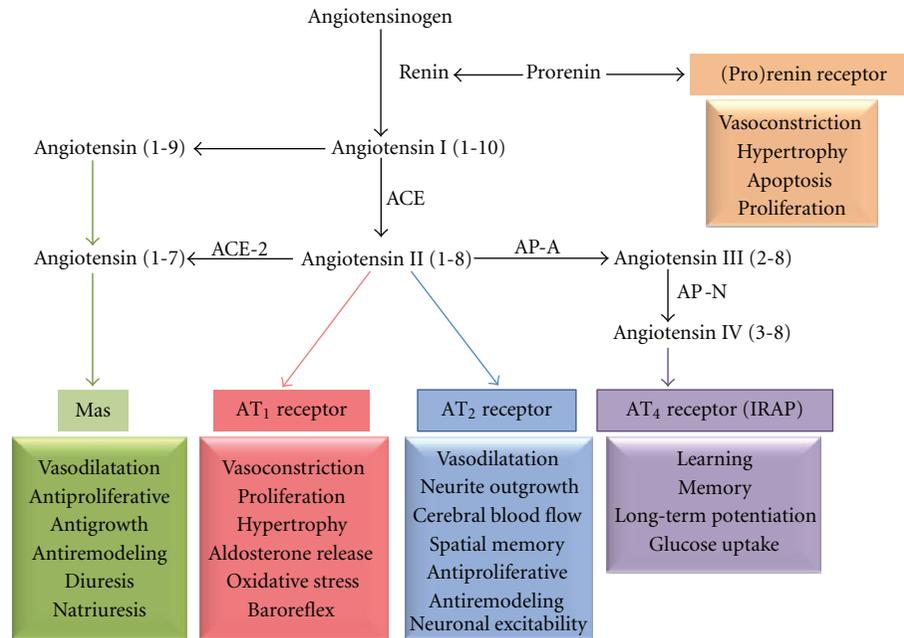


FIGURE 1: Summary of the brain renin angiotensin system (RAS). The figure summarizes the conversion of angiotensinogen to angiotensins I and II through fragments. The biologically active forms include angiotensins II, III, IV, and (1–7). The main enzymatic pathways are mediated by renin and angiotensin converting enzyme ACE or ACE2, AP-N and AP-A. The major brain effects of angiotensins are mediated by AT₁, AT₂, AT₄, prorenin, and Mas receptors. The functions associated with each receptor are indicated. ACE: angiotensin converting enzyme; AP-A: aminopeptidase A; AP-N: aminopeptidase N, adapted from Phillips and Oliveira, 2008 [3] and from Wright and Harding, 2012 [4].

other hand, it is generally assumed that the AT₂ receptor counteracts the action of the AT₁ receptor, promoting vasodilatation, apoptosis, and antigrowth effects. In addition to the classical AT₁ and AT₂ receptors, more recent studies have identified other receptors for RAS components, such as the (pro)renin receptor, the Ang (1–7) Mas receptor, and the Ang IV receptor (AT₄ receptor, called IRAP for insulin-regulated aminopeptidase), all of which are expressed in the brain (for reviews, see [4, 12, 13]) (Figure 1). In particular, several studies have shown that Ang IV-AT₄ receptor/IRAP have important functions in the brain, related to cognition and memory. In addition, in many situations, Ang IV acts as an inhibitor of AT₁ receptor actions [15–17].

The present paper is focused on the known and suggested roles of the AT₂ receptor in brain functions related to neuronal activities and cognitive disorders as well as the potential link between metabolic syndrome and cognitive functions. The role of the AT₂ receptor in the central regulation of blood pressure, thirst and related pathologies, such as hypertension, stroke, or ischemic damage, will not be discussed here; these latter topics are covered in recent reviews (see [11, 18–20]). For further detail, readers are also invited to consult several excellent reviews describing recent up-to-date advances pertaining to the various active Ang II-derived ligands and their receptors [4, 12, 13]. However, considering that there are some similarities between the AT₂ receptor and the Ang IV/AT₄ receptor, comparisons between the two will be included, when appropriate.

2. The Type 2 Receptor—A Nonclassical GPCR

Three important periods highlight the history of the AT₂ receptor: (i) its discovery in the 1980s, (ii) its cloning in 1991–1993, and (iii) the generation of transgenic mice in 1995. However, because Ang II has a similar affinity for both of its receptors, it has been difficult to discriminate between the latter using nonselective ligands such as native Ang II or peptide analogs such as Saralasin or SarIle. At the end of 1980s, tools enabling to distinguish these Ang II receptors became available. The first tools included the nonpeptide antagonists losartan (previously known as DUP753) and PD123,177, and peptide ligands such as CGP42112A and p-aminophenylalanine (review in [2]). Seminal autoradiographic studies [5, 6], followed by biochemical studies by the groups of Marc de Gasparo, Peter Timmermans, and Robert Speth revealed the presence of two binding sites for Ang II, which differed in their expression pattern and biochemical properties [21–25]. However, the conclusive evidence for the existence of two receptors came from their cloning in the early 1990s [26, 27]. Despite the fact that both receptors belong to the large 7-transmembrane domain family of GPCR, AT₁ and AT₂ share only ~34% amino acid sequence identity. Moreover, while the AT₂ receptor displays most of the structural features of a GPCR, it is usually considered as an atypical member of this family. Indeed, although some of its signaling pathways involve a G_i-dependent mechanism, most of the known effects of the AT₂

receptor are independent from G-protein coupling mechanisms (for reviews, see [14, 20, 28–31]). Moreover it fails to induce all of the classical signaling pathways such as cAMP, production of inositol triphosphate, or intracellular calcium release (for reviews, see [28, 29]).

2.1. Selective Ligands of the AT₂ Receptor. In fact, AT₂ receptor functions are still unclear both in many physiological and pathophysiological situations, mainly because research on this receptor has long been hampered by at least three challenges, namely, (i) the low and unusual expression of this receptor, (ii) its nonclassical signaling, and (iii) the absence of appropriate selective ligands. While several nonpeptide ligands with a large spectrum of selectivity for the AT₁ receptor are readily available, only a few have been developed for the AT₂ receptor. Until recently, physiological and pharmacological assessments of the AT₂ receptors were obtained using either the antagonist PD123,319 (a modification of the initial PD123,177) or the more common AT₂ receptor agonist CGP42112A, although two other peptides have also shown agonistic properties on AT₂ receptors, namely, the agonist *p*-aminophenylalanine [22] and novokin [32, 33]. However, the fact that CGP42112A is only a partial agonist, combined with its short half-life, has hampered its further use in *in vivo* studies (review in [20, 34, 35]). Thus, many hypotheses regarding AT₂ receptor functions in physiological situations have emerged from indirect observations, either using transgenic animals, or blockade of the AT₁ receptor. In 2004, Wan and collaborators synthesized the first selective nonpeptide AT₂ receptor agonist, called compound 21 (C21) [36], now renamed M024 [37]. Since then, an increasing number of studies have used C21/M024 for both *in vitro* and *in vivo* studies (see further below) to better investigate the selective role of the AT₂ receptor (for review, see [20, 34]).

3. Overview of the Signaling Pathways of the AT₂ Receptor in the Brain

Signaling pathways associated with the AT₂ receptor primarily involve a balance between phosphatase and kinase activity. The final outcomes vary according to whether the cell is undifferentiated or differentiated and whether angiotensin AT₁ receptors are also expressed or not. Although there is still much controversy surrounding the effects of the AT₂ receptor in peripheral systems, its role and mechanisms of action in the brain have gained much greater consensus [14, 31, 38].

3.1. Signaling Pathways of the AT₂ Receptor Leading to Neurite Outgrowth. In neuronal models, studies on functionality and signaling have usually been conducted simultaneously. Most of the data, including our own findings using NG108-15 cells, indicate that the effects of the AT₂ receptor on neurite outgrowth involve four main complementary signaling cascades (Figure 2). The first cascade entails a decrease in p21^{RAS} [39] and protein kinase C α (PKC α) activities [40], both of which are involved in the switch from proliferation to differentiation. Simultaneously, a second cascade, initiated by Rap1/B-Raf, induces a delayed and sustained phosphorylation of p42/p44^{mapk} [39, 41]. In both NG108-15 [39] and

PC12W [42] cells, this sustained activation is essential for inducing neurite outgrowth. The initial activation of Rap1 by the AT₂ receptor is not direct, but rather mediated by phosphorylation of the tropomyosin-related kinase receptor A (TrkA) [43], through the intervention of a Src family kinase member [44]. The third cascade comprises nitric oxide (NO) and cGMP. In NG108-15 cells, we [45] and others [46] have shown that external application of NO is sufficient to induce neurite outgrowth and elongation. In these cells, neuronal NO synthase (nNOS) activation and cGMP production induced by the AT₂ receptor is dependent on the G α_i protein. However, cGMP is not involved in Ang II-induced activation of p42/p44^{mapk} [45]. More recently, Li et al. [47] have shown that after AT₂ receptor stimulation, the AT₂ receptor interacting protein, ATIP [48], interacts with the tyrosine phosphatase SHP-1. The complex then translocates into the nucleus where it transactivates the ubiquitin-conjugating enzyme variants called methyl methanesulfonate sensitive 2 (MMS2), resulting in neural differentiation and protection (reviewed in [31, 49]). Finally, interaction of AT₂ with certain receptor tyrosine kinases may also induce neurite outgrowth. For example, in a model of fructose-induced insulin-resistant rats, authors also demonstrated that neurite outgrowth of dorsal root ganglia (DRG) neurons induced by the AT₂ receptor was facilitated by activation of the phosphatidylinositol 3-kinase (PI3K)/Akt pathway, suggesting the existence of a crosstalk between the AT₂ receptor and the insulin receptor [50]. In addition, several phosphatases, such as PP2A [51–54] and Src homology region 2 domain-containing phosphatase-1 (SHP-1), are clearly associated with the mechanism of action of the AT₂ receptor. Finally, certain pathways associated with AT₂ receptor activation may be through interaction with protein partners such as the promyelocytic zinc finger (PLZF) protein [55, 56] or the peroxisome proliferator-activated receptor gamma (PPAR γ) [57] (for review, see [29, 31, 58–60]).

3.2. Ang-II-Independent Effects of the AT₂ Receptor. It should be noted that while the main effects of AT₁ and AT₂ receptors are dependent of Ang II binding, some evidences also suggest that they may have certain ligand-independent effects (review in [29, 31]). For example, AT₁ receptors can be activated by mechanical stress, independently from Ang II binding and/or stimulation of p42/p44^{mapk} [61, 62]. Similarly, AT₂ receptor overexpression in CHO cells, R3T3 fibroblasts, and vascular smooth muscle cells enhances apoptosis signaling simply by its overexpression [63]. Another study also observed that the AT₂ receptor, when expressed as a constitutive homooligomer, leads to G-protein dysfunction and symptoms of neurodegeneration without Ang II stimulation [64]. Although the mechanism underpinning this effect still requires further investigation, it appears that the C-terminal portion of the AT₂ receptor is essential, since expression of a mutant AT₂ receptor truncated in its C-terminal region is unable to form oligomers. Aside from its effect on cell survival, Ang II-independent effects of the AT₂ receptor also include modulation of gene expression, at least in human coronary artery endothelial cells, where AT₂ receptor overexpression modulates more genes than

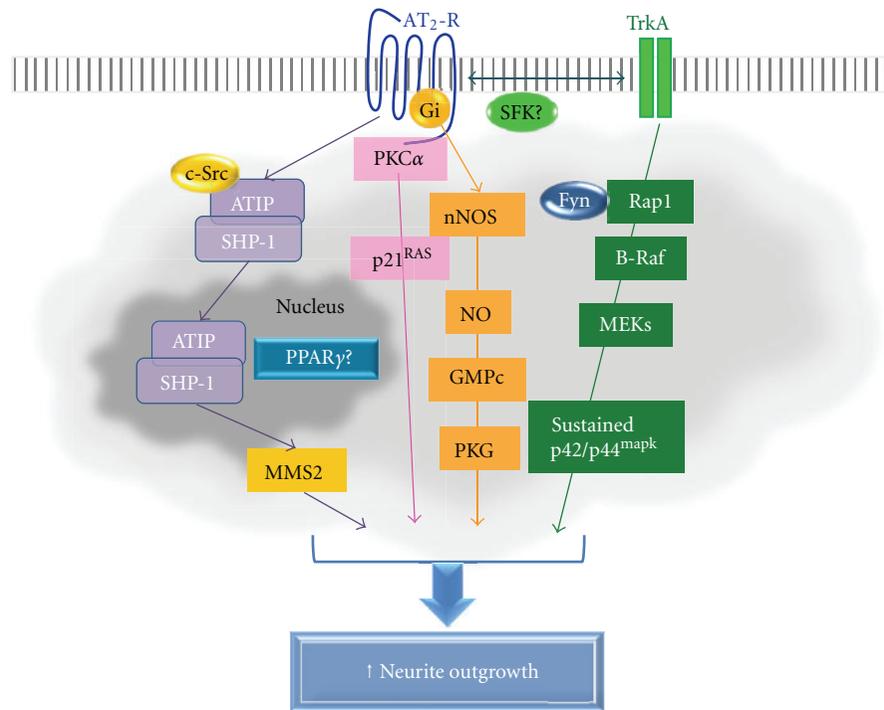


FIGURE 2: Overview of the main signaling pathways implicated in the action of AT_2 receptor leading to neurite outgrowth.

CGP42112A stimulation [65]. Although these AT_2 -regulated genes are associated with many cellular functions, including cell migration, protein processing, intracellular signaling, and DNA repair, it is still unknown whether Ang II-independent effects of the AT_2 receptor are associated with protective effects in neuronal function. Moreover, an Ang II-independent effect of the AT_2 receptor was observed in a model overexpressing the receptor, and thus its relevance in physiological situations is still unknown. Clearly, many questions still remain to be elucidated in order to fully understand how the AT_2 receptor exerts its effects on brain functions. However, new recent insights in AT_2 receptor signaling have been achieved which could partly explain some of the observed discrepancies (see Section 8).

3.3. Signaling Pathways Associated with Ang IV and the AT_4 Receptor. Interestingly, despite the high similarity between central biological effects associated with AT_2 and AT_4 receptor stimulation (memory processing, long-term potentiation facilitation, protective function in cognitive loss) (see Section 5), the signaling pathways induced by these two receptors are clearly different. In contrast with the AT_2 receptor, the AT_4 receptor is a single transmembrane receptor, initially known for its aminopeptidase activity, leading to peptide processing of oxytocin, vasopressin, Ang III, met-enkephalin, somatostatin, and other neuropeptides (review in [4, 12]). Ang IV has also been shown to increase endothelial NO synthase activity via mobilization of intracellular calcium [66, 67] and to stimulate the PI3 K/PKB (protein kinase B) pathway [68] in endothelial cells. A recent

study also showed that Ang IV treatment in diet-induced hyperglycemic mice increased the interaction between IRAP and PI3 K, leading to activation of Akt and the glucose transporter type-4 (GLUT4) translocation. This pathway was also associated with an improvement in glucose tolerance [69], suggesting that Ang IV may be implicated in insulin signaling and development of diabetes. Finally, in some studies, Ang IV was also found to elicit rapid activation of other selected kinases including ERK1/2, p38 kinase, focal adhesion kinase, and paxillin (review in [70]). Nevertheless, it appears that these signaling effects of Ang IV are model dependent, and none of these pathways have been observed in neuronal cells. Therefore, although renewed interest in Ang IV has recently emerged with the recognition that the Ang IV/IRAP/ AT_4 receptor plays an important role in cognition and pain [4, 12, 71], it is still unknown to date whether signaling pathways associated with Ang IV stimulation are implicated in these effects.

4. Expression and Roles of the AT_2 Receptor in the Brain

All of the various RAS components in the brain, and in particular the AT_2 receptor, are highly expressed during fetal life, suggesting that they could play key roles during development. On the other hand, in the adult, AT_2 receptor expression in the brain is limited to certain specific areas (review in [72–75]). For instance, the AT_2 receptor has been found at high levels in the medulla oblongata (control of autonomous functions), septum and amygdala (associated

with anxiety-like behavior), thalamus (sensory perception), and superior colliculus (control of eye movements in response to visual information) as well as in the subthalamic nucleus and cerebellum (areas associated with learning of motor functions). It is also expressed with the AT₁ receptor in areas involved with cardiovascular functions, learning and behavior (cingulate cortex, molecular layer of the cerebellar cortex, superior colliculus, paraventricular nuclei hippocampus) (for extensive mapping, see [72, 73]). More recently, expression of the AT₂ receptor was also detected in the substantia nigra pars compacta, the area involved in dopaminergic signals and associated with Parkinson's disease [76], and in the hippocampus [64, 77]. At the cellular level, the AT₂ receptor is expressed in neurons, but not in astrocytes [28, 72, 78]. Evidence also suggests that the AT₂ receptor is expressed in the vasculature wall, where it acts on cerebral blood flow (review in [31, 49]). In addition, the presence of a non-AT₁/non-AT₂ receptor in the CNS has been suggested, which displays high affinity for Ang I, II, and III [79].

Interestingly, the Ang IV receptor IRAP is also observed in structures classically associated with cognitive processes and sensory and motor functions (including hippocampus, thalamic nuclei, caudate putamen, cerebellum, neocortex, lateral geniculate body, inferior olivary nucleus, superior colliculus, ventral tegmental area, and brain stem) (review in [12, 80, 81]). In contrast, the Ang (1–7) Mas receptor is expressed in brain areas involved in central cardiovascular regulation (in particular, the nucleus of the solitary tract (NTS), rostral ventrolateral medulla (RVLM), caudal ventrolateral medulla (CVLM), inferior olive, parvo- and magnocellular portions of the paraventricular nucleus (PVN), supraoptic nucleus, and lateral preoptic area) (review in [82]). In some instances, these areas also contain AT₁ and AT₂ receptors.

One of the first roles of the AT₂ receptor to be identified was the modulation of neuronal excitability (review in [28, 83]). In cells of neuronal origin, activation of the AT₂ receptor decreases activity of T-type calcium channels [84, 85] while stimulating a delayed-rectifier K⁺ current (I_K) and a transient K⁺ current (I_A) [86]. Using primary cultures of cortical neurons, studies by Grammatopoulos et al. [87] have shown that II neuroprotection against chemical hypoxia was mediated by activation of a delayed rectifier K⁺ channel, an effect exemplified by simultaneous blockade of the AT₁ receptor. Moreover, in preparations of locus coeruleus brain slices [88], angiotensin II, through the AT₂ receptor, was found to depress glutamate depolarization and excitatory postsynaptic potentials. In the superior colliculus, both AT₁ and AT₂ receptors are involved in sensory visuomotor integration. Lastly, Ang II, through AT₂ receptor stimulation with CGP42112A, has also been shown to induce a strong suppressive effect on visual neuronal activity. Together, these results indicate that AT₂ receptor modulation of potassium and calcium channels activity may impact neuronal functions.

The second and best recognized effect of the AT₂ receptor is stimulation of neurite outgrowth in various cell types from neuronal origin (NG108-15 and PC12W cells) as well

as in primary cultures of neurons from retinal explants [89], in neurospheres from mouse fetal brain [90], cerebellar neuronal cells [91], and the cortex [47] (review in [3, 10, 14, 34]). This AT₂-induced neurite elongation is characterized by an increase in mature neural cell markers, such as β III-tubulin and MAP2 [47, 92]. It is also associated with a rise in neuronal migration [54, 91] and neuronal survival following ischemia-induced neuronal injury [93]. These effects may be important not only for developmental differentiation, but also after injury-induced regeneration. Indeed, a beneficial effect of the AT₂ receptor in nerve regeneration has been observed following both optic [89] and sciatic [94] nerve crush or in perivascular nerves implicated in vasodilation [95]. This implication of the AT₂ receptor in neuronal regeneration has even led to the suggestion that Ang II, via the AT₂ receptor, could act as a neurotrophic factor. In summary, the AT₂ receptor, by its function in the modulation of neuronal excitability, neurite elongation, migration, and nerve regeneration, may be an important factor in the regulation of central nervous system activity and cognitive function either following nerve injury or during the development of neurodegenerative disease (Figure 3).

In addition to neuronal differentiation, which is of paramount importance in nerve regeneration, the AT₂ receptor also stimulates differentiation of hematopoietic cells, a key process during regeneration and reconstruction. Indeed, ischemic damage is characterized by infiltration of a number of hematopoietic cells such as leukocytes, platelets, macrophages, and leukocytes [96]. In particular, the AT₂ receptor has the capacity to induce differentiation of human monocytes into dendritic cells [97]. Supporting a protective effect of the AT₂ receptor is the observation that ischemic damages were found to be greater in mice with hematopoietic cells deleted in AT₂ receptor expression [98]. These findings suggest that expression and activation of the AT₂ receptor in hematopoietic cells may be part of its beneficial effect following brain injury, although the mechanism involved remains to be investigated (review in [99]).

Some of the initial hypotheses regarding the potential role of the AT₂ receptor *in vivo* were confirmed in 1995 when two independent groups developed AT₂ knockout mice by targeted gene deletion [100, 101]. Surprisingly, despite the fact that the AT₂ receptor is highly expressed during fetal development in many tissues, including skin, kidney, brain, and heart, it appears that mice lacking the AT₂ receptor do not exhibit any major anatomical defects. However, these mice exhibit markedly reduced exploratory behavior as well as altered thirst reaction, lower body temperature, slightly elevated mean arterial pressure, and a stronger vasoconstrictive response to Ang II [100–102]. Moreover, they exacerbate stronger symptoms when exposed to pathological situations. For example, these mice display an accelerated pathological response when exposed to cardiovascular disease induction. They also exhibit larger cerebral infarct size following medial cerebral artery occlusion (MCAO) [93], stronger cognitive deficits following ischemia [103], and faster progression of atherosclerosis [104]. The various actions of the AT₂ receptor currently documented in the brain are summarized in Figure 3.

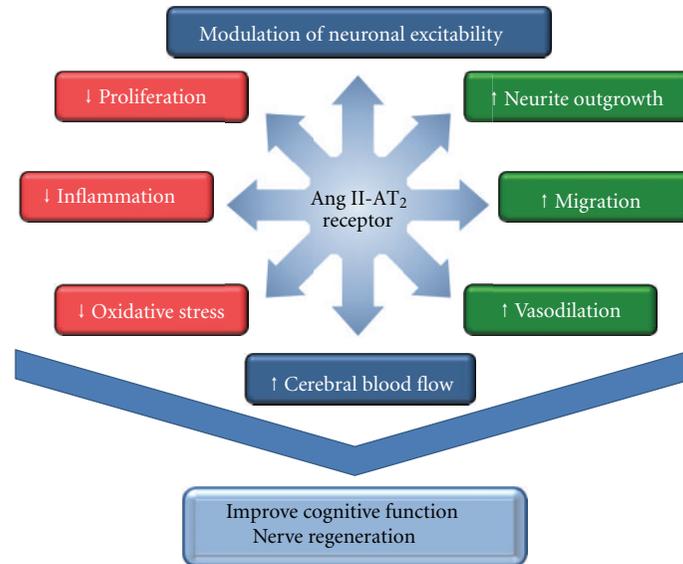


FIGURE 3: Summary of the known cellular effects of the AT_2 receptor. Stimulation of the AT_2 receptor of angiotensin II results in the activation of several intracellular cascades, resulting in a decrease in proliferation and in growth-promoting effects, as well as a decrease in inflammatory cytokines and in reactive oxygen species, hence decreasing oxidative stress. On the other hand, such activation increases neuronal excitability, by acting on K^+ and Ca^{2+} channel activity, and increases neurite outgrowth and migration of neurons, favoring neuronal plasticity. Stimulation of the AT_2 receptor is also known for its effect on vasodilation, thus increasing cerebral blood flow, and to increase glucose uptake, which improves insulin sensitivity. These cellular events in turn decrease or increase important physiological functions, which in general have reached a significant degree of consensus. Together, the AT_2 receptor may appear as a gatekeeper of cellular and tissue homeostasis. Indeed, most current studies suggest that AT_2 receptor activation may have a protective role in various pathological situations.

5. Role of the AT_2 Receptor in Cognitive Function and Neurological Disorders

The first evidence for the implication of the AT_2 receptor in cognition resulted from studies in knockout mice. Indeed, AT_2 -deficient mice suffer from perturbations in exploratory behavior and locomotor activity [100, 101], as well as displaying an anxiety-like behavior [105]. Moreover, in the adult, inhibition of the AT_2 receptor with the AT_2 receptor antagonist PD123,319 has been reported to abolish the Ang II-induced acquisition of conditioned avoidance responses [106]. These results strongly support that, in addition to a role during development, the AT_2 receptor may be involved in cognitive processes in the adult. Furthermore, although the AT_2 receptor is expressed at low levels in many areas of the nervous system, it may be reexpressed in certain pathological conditions such as optic [89] or sciatic [94] nerve transection, stroke [107], and certain neurodegenerative diseases such as Alzheimer's disease [108]. For example, Ge and Barnes [108] found that AT_2 receptor expression is diminished in Parkinson's disease (caudate nucleus and cerebellum) but enhanced in Huntington's disease (caudate nucleus). In Alzheimer's disease, the temporal cortex of the adult brain exhibits an increased expression while the hippocampus displays a decreased expression of the AT_2 receptor. In most of these situations, the AT_2 receptor is described as having beneficial effects in improving neuroprotection by acting not only on neurons, but also on blood

circulation. Indeed, recent studies have clearly shown a protective role of the AT_2 receptor following brain ischemia and demonstrated that expression and activation of the AT_2 receptor may decrease brain damage and restore cognitive loss following middle cerebral artery occlusion (for review, see [11, 14, 49]). Altogether, these data suggest that the AT_2 receptor may play an important role in maintaining functions of the human brain.

5.1. *AGTR2* Mutations in Intellectual Disability. Intellectual disability, previously described as mental retardation [109], affects approximately 1–3% of the population, of which a large number are associated with mutations on chromosome X. Among these, certain mutations on *AGTR2* coding for the angiotensin AT_2 receptor have been identified. Mutations in the *AGTR2* gene correlate with the development of human X-linked intellectual disability [110]. Indeed, 9 patients with X-linked intellectual disability were shown to have mutations in the *AGTR2* gene associated with decreased expression of the AT_2 receptor, including a complete loss of expression in a woman with an IQ of 44. Clinical features of these mutations ranged from moderate to severe intellectual disability, seizure, and manifestations of autism, thus supporting the hypothesis that the AT_2 receptor is required for brain development and for the maintenance of neuronal connections involved in learning and memory. This hypothesis was further corroborated by two other studies reporting mutations of *AGTR2* in patients suffering from intellectual

disability, seizures, restlessness, hyperactivity, and disrupted speech development [111, 112]. However, other studies failed to link any *AGTR2* mutations with intellectual disability, observing no difference in mutation incidence between the latter and control groups [113–115]. Differences among control cohort selection, number of patients per group, and ethnical variations between different studies could explain the discrepancies in these findings. Hence, it remains unclear whether mutations in *AGTR2* are associated or not with intellectual disability.

5.2. *AT₂ Receptor in Alzheimer's Disease (AD).* Amyloid- β ($A\beta$) deposition in senile plaques and the presence of neurofibrillary tangles are the main pathological hallmarks of AD. However, other structural and functional alterations, including inflammation, increased oxidative stress and vascular damage/ischemia, are also associated with AD and other neurodegenerative diseases. These alterations may contribute to neuronal and synaptic dysfunction and loss, as well as the ensuing cognitive deficits and dementia of this disorder [116–121] (for recent review, see [11, 14, 49]). Clinical studies have documented that treatment with antihypertensive drugs is associated with an improvement in cognitive function (review in [122–124]). More recent studies have shown that treatment with angiotensin II receptor blockers (ARBs) is associated with a decrease in AD and dementia progression with a greater efficacy compared to ACE inhibitors [125, 126]. In particular, Tsukada et al. [127] have shown that cognitive deficit induced by $A\beta$ (1–42) in mice was improved by pretreatment with a low dose of telmisartan partly because of peroxisome proliferator-activated receptor- γ (PPAR γ) activation. Corroborating the hypothesis that ARBs could be beneficial in reducing the onset of AD, Kume et al. [128] recently observed that AD hypertensive patients treated with telmisartan presented no decrease in cognitive functions test scores, but an increased cerebral blood flow, suggesting that treatment with this ARB could reduce AD progression. Altogether, studies conducted using various models of cognitive disorders have reported improved memory and cognitive processes and/or attenuation of $A\beta$ 1–42 oligomerization following treatment with ARBs, particularly valsartan [129], losartan [130], telmisartan [128, 131], and olmesartan [132] (now called metabosartans for ARBs with a PPAR γ agonistic effect) (review in [11, 14, 49, 133]). More recently, a study using direct stimulation of the *AT₂* receptor with the selective agonist C21/M024 demonstrated similar effects in an AD mouse model [134]. In this latter study, Jing et al. observed that intracerebroventricular injections of $A\beta$ (1–40) in mice induced a poorer performance in the Morris water maze and that this effect was reversed by coadministration of C21/M024, indicating that direct stimulation of *AT₂* receptors improves spatial memory functions. Stroke is also one of the most important causes of cognitive impairment and dementia (review in [11, 49]). There is increasing evidence suggesting that activation of the *AT₂* receptor could improve cerebral blood flow and microcirculation as well as decrease inflammation [93, 103, 135, 136], both being associated with improvement in cognitive function following cerebral ischemia [90, 93, 103,

137] (for reviews, see [3, 10, 14, 49]). As reported by Iwai et al. [103] and summarized by Horiuchi and Mogi [49] and Mogi and Horiuchi [11], superoxide anion production was found to be more markedly enhanced in *AT₂* receptor-deficient mice compared to wild-type animals in a model of MCAO. These observations suggest that *AT₂* receptor stimulation has a protective effect on ischemic brain lesions, at least partly through modulation of cerebral blood flow and superoxide production. Moreover, beneficial effects of ARBs on these parameters were less evident in *AT₂* receptor-deficient mice [103]. Similar approaches were also used in other studies suggesting a beneficial effect of the *AT₂* receptor on cognitive functions [90, 135, 136].

However, not only blood circulation but also neuronal functions can be improved by activation of *AT₂* receptors. Indeed, activation of *AT₂* receptors in neurons is also associated with a decrease in apoptosis signaling. Grammatopoulos et al. demonstrated in cultures of primary cortical neurons that angiotensin decreased sodium azide-induced apoptosis through *AT₂* receptor activation [138] by reducing caspase-3 activation [139]. Finally, we and others have shown that the *AT₂* receptor can activate the tyrosine kinase Fyn [44] and the phosphatase PP2A [51, 52, 54], both of which are key regulators of the phosphorylation of the microtubule associated protein tau. Hyperphosphorylation of tau is of paramount importance in the development of Alzheimer's disease by forming neurofibrillary tangles and leading to microtubule depolymerization. Thus, *AT₂* receptor activation may participate to the control of equilibrium between tau phosphorylation and dephosphorylation.

Another key observation is the effect of estrogen receptors on *AT₂* receptor functions. In 2008, Chakrabarty et al. [140] demonstrated that estrogen, through its E2 receptor, induced neurite elongation of dorsal root ganglion and that this effect was dependent on *AT₂* receptor activation. Moreover, it has been observed that ischemic damage in *AT₂*-deficient mice was greater in females than in males, while no significant sex-different change was observed in *AT₂*-expressing mice [135]. These results suggest that the existence of some level of crosstalk between *AT₂* receptor and estrogen and that *AT₂* receptor could be necessary for E2 receptors to elicit its full effect on neuronal physiology. Thus, while stimulation of the *AT₂* receptor and/or inhibition of the *AT₁* receptor could lead to potential therapeutic avenues in neurodegenerative disease, the interaction between *AT₂* receptor and estrogen should also be considered. Indeed, there are some differences in response to RAS stimulation according to gender (review in [141]). Nonetheless, the clear demonstration of an *AT₂* receptor effect in Alzheimer's disease remains to be firmly demonstrated, mainly because in most studies to date the presence of *AT₁* receptors and *AT₂* receptors in the hippocampus has not been studied. Indeed, to our knowledge, only two studies have documented the presence of *AT₂* receptors in the hippocampus, one in a model of Alzheimer's disease [64], the other in a model of epilepsy [77]. On the other hand, presence of the *AT₂* receptor was not detected in the hippocampus in various studies from the Llorens-Cortes group [72, 73]. However, these studies were performed with healthy and middle-aged animals.

It should be mentioned that, in addition to the AT₂ receptor, the Ang IV/AT₄ receptor may also have a protective effect on cognitive function. Indeed, Braszko's group was the first to report that intracerebroventricular injections of Ang II and Ang IV were equivalent in facilitating exploratory behavior in rats tested in an open field and improved recall of passive avoidance conditioning in the acquisition of active avoidance conditioning [142, 143], results that were confirmed by others in subsequent studies (review in [71]). This strongly suggests an important function of Ang IV and its receptor in learning and memory processes and could represent a new therapeutic target in the treatment of memory loss associated with dementia (review in [71]). One mechanism proposed to explain this beneficial effect of Ang IV on cognitive function is the colocalization of IRAP with the glucose transporter GLUT4. In hippocampal pyramidal neurons, IRAP and GLUT4 are localized in secretory vesicles responsive to insulin (review in [70, 71, 80]), suggesting that Ang IV, by binding to IRAP, may increase GLUT4 membrane expression thus facilitating glucose uptake, as observed in adipocytes [144] (review in [145]).

5.3. AT₂ Receptor and Parkinson's Disease. In addition to AD and stroke, some evidences also suggest that central RAS could be implicated in the development of Parkinson's disease. Parkinson's is the second most common neurodegenerative disorder and is characterized by the progressive cell death of midbrain dopaminergic neurons in the substantia nigra and the presence of protein inclusions leading to formation of Lewy bodies (review in [146]). Although the mechanisms leading to Parkinson's disease are still unclear, it appears that mitochondrial dysfunction, oxidative stress, and inflammation are key factors to its progression [147]. Recently, ARBs have been shown to reduce lipid peroxidation and protein oxidation while protecting dopaminergic neurons in the substantia nigra in a rat model of Parkinson's disease [76, 148]. However, whether such action is due solely to a blockade of the AT₁ receptor or also from activation of the AT₂ receptor is not yet clearly established (review in [147]). Activation of the AT₂ receptor is able to stimulate differentiation of mesencephalic precursor cells into dopaminergic neurons, suggesting that stimulation of the AT₂ receptor could be useful in increasing the production of dopaminergic neurons in Parkinson's disease [149]. Moreover, Grammatopoulos et al. observed a protection against rotenone-induced oxidative stress and associated cell death in dopaminergic neurons following Ang II stimulation. This protective effect was prevented by the presence of the AT₂ receptor antagonist PD123,319, but was increased in the presence of the AT₁ receptor antagonist losartan [150]. More recently, it has been observed that during the aging process in rats, AT₂ receptor expression is decreased in dopaminergic neurons, as opposed to an increase in AT₁ receptor expression. This was associated with an enhancement of prooxidative and proinflammatory markers in the substantia nigra, leading to an increase in dopaminergic neuronal death [151]. Although these findings do not indicate a role of AT₂ receptor in the development of Parkinson's disease, they strongly suggest that such

modifications in Ang II receptors during the natural aging process may increase the risk of Parkinson's disease.

6. Role of the AT₂ Receptor in the Regulation of Appetite

Obesity, which is characterized by excess body fat accumulation [152], is associated with an increased risk of diabetes, hypertension, and dyslipidemia. It is also one of the major components of the metabolic syndrome. Studies have demonstrated AT₂ receptor expression in tissues associated with glucose metabolism, including pancreatic [153–155] and adipose tissues [13, 30]. For example, expression of the AT₂ receptor in the pancreas has been shown to be important for fetal pancreatic development [156] while in the adult, it is associated with protection against pancreatic fibrosis [157] and a decrease in pancreatic tumor growth [158, 159]. The following is a summary of what is currently known regarding the potential function of the AT₂ receptor in the regulation of appetite, glucose metabolism, and its potential role in metabolic syndrome.

There are some evidences suggesting that Ang II could be implicated in food intake: for example, Ang II suppresses food intake after central infusion [32, 160, 161] while blockade of the AT₁ receptor by telmisartan is associated with a decrease in body weight [162]. Furthermore, both AT₁ and AT₂ receptors are expressed in the hypothalamus, which is implicated in the central regulation of food intake. In 2008, Ohinata et al. [32] observed that the decrease in food intake induced by centrally administered Ang II was inhibited by PD123,319, and absent in AT₂-KO mice, suggesting that this effect was mediated by the AT₂ receptor. A similar effect was observed using novokin, a potent analog of ovokinin with AT₂ receptor agonistic properties [163]. Although the mechanism underlying this AT₂ receptor-associated decrease in food intake remains unclear, it may be linked to its capacity to modulate T-type calcium channels, since a recent study demonstrated that inhibition of these channels inhibits weight gain in mice fed with a high-fat diet [164]. However, this hypothesis remains to be explored. Studies conducted to date with the selective AT₂ agonist C21/M024 do not describe such differences between C21/M024-treated animals compared to the control group [165, 166], suggesting that the duration of C21/M024 treatment may have been too short to induce any modification in body weight. Thus, in this latter instance, even if the AT₂ receptor was observed to decrease food intake, it was probably not sufficient to induce a decrease in body weight, at least following short periods of stimulation (<8 weeks). Moreover, it has also been observed that inhibition of ACE by captopril also reduced body weight of mice fed with a high fat diet [167]. This decrease in body weight, however, was not associated with a decrease in food intake, suggesting that other mechanisms were regulated by Ang II. Since Ang II is no longer available in captopril-treated mice, these results suggest that other members of the RAS, independent of Ang II, could be implicated in the regulation of food intake and body weight. These observations should therefore be considered when interpreting results obtained with ARBs.

7. Link between Metabolic Syndrome and Alzheimer's Disease: Is There a Place for the AT₂ Receptor?

A number of excellent reviews have recently been published regarding the potential link between insulin resistance, metabolic syndrome, and neurodegenerative disorders (both AD and vascular dementia) [168–177], including the involvement of RAS in this process [90, 133, 178]. Development of AD is closely associated with a reduction in cerebral glucose utilization, even in the early stages of the disease. In fact, cerebral metabolism in the AD brain decreases prior to the onset of cognitive decline, suggesting that energy failure could represent one of the earliest hallmarks of AD. Induction of insulin resistance in AD animal models aggravate both amyloid and tau accumulation [179, 180], leading several investigators to refer to Alzheimer's disease as type 3 diabetes (review in [168]). One aspect of this relationship is the loss of insulin signaling in the insulin-resistant brain. In addition to the many peripheral complications associated with dysfunction in insulin sensitivity, it appears that brain insulin signaling plays crucial central functions in the regulation of energy balance (food intake, body weight) as well as in learning and memory (review in [171]). Moreover, inflammation, increase in oxidative stress, and mitochondrial dysfunctions are key features of type 2 diabetes (T2D) that are also shared in Alzheimer's disease.

In T2D patients, results of a major clinical study (Study on Cognition and Prognosis in the Elderly, SCOPE) [181] and a clinical double-blind study [182] have revealed that ARBs have a further therapeutic effect on impaired cognitive function beyond their antihypertensive effects compared with other antihypertensive drugs. Similarly, Tsukuda et al. [131, 183] have demonstrated that candesartan improves impaired cognitive function induced by T2D, with multiple beneficial effects. Two of these effects may be through PPAR γ or through AT₂ receptor activation. Therefore, improvement of metabolic syndrome may also be beneficial in decreasing associated cognitive decline. This would contribute to better insulin signaling in the brain and, therefore, a slowing of cognitive decline associated with brain insulin resistance.

8. New Insights in AT₂ Receptor Knowledge and Perspectives: What Remains to Be Done?

As pointed out recently [18, 20], there is still an ongoing debate as to the putative role of the AT₂ receptor in physiology, and whether this role is deleterious or beneficial. Summarized below are some of the new advances in AT₂ receptor signaling that could have important insights in AT₂ receptor-associated brain functions.

8.1. Homo- and Heterodimerization. Although GPCRs have traditionally been thought to act as monomers (review in [184]), it is now well accepted that many GPCRs can form dimers which could affect both their trafficking and function. In this context, homodimerization of AT₁ and AT₂ receptors, as well as AT₁/AT₂ heterodimer formations, has been reported. For example, the AT₂ receptor is known

to undergo homodimerization, a property which enhances apoptosis [185]. In addition, AbdAlla et al. [186] reported that the AT₂ receptor undergoes heterodimerization with the AT₁ receptor in transfected PC12 cells, in fetal fibroblasts, and in myometrial biopsies. In an animal model of Alzheimer's disease, the same group demonstrated that A β induces the formation of cross-linked AT₂ receptor oligomers [64, 187]. Notably, oligomers of AT₂ receptors have also been observed in prefrontal cortex specimens of Alzheimer's disease patients, while being completely absent in specimens of nondemented control individuals, thus lending further support for a role of the AT₂ receptor in cognitive function. Heterodimerization between the AT₂ receptor and bradykinin has also been described in PC12W cells [188]. It is already known that bradykinin mediates AT₂ receptor-induced NO production [189–191]. This interaction between the two receptors has been shown to enhance phosphorylation of various kinases, including p42/p44^{mapk} and p38^{mapk}. Therefore, it would appear that homo- and heterodimerization of the AT₂ receptor may have a role in its regulation. However, these initial observations still require confirmation before being accepted as important regulatory aspects of Ang II receptor signaling and functions (review in [29, 31]). The use of recently developed methodologies such as FRET/BRET technology has confirmed efficient heterodimerization of the AT₁ receptor with the bradykinin receptor B2 [192], indicating the potential clinical significance of GPCR oligomerization [29, 31]. Moreover, recent studies have identified intracellular crosstalk pathways between the AT₁ receptor and the AT₂ receptor at the gene expression level. Indeed, AT₁ receptor activation enhances AT₂ receptor mRNA degradation, while AT₂ receptor activation increases its own mRNA transcription [193].

8.2. PPAR γ : Could It Be the Missing Link? There is existing confusion regarding the mechanism of action of specific ARBs, since some also have partial PPAR γ agonistic activity (such as telmisartan, irbesartan, and candesartan). There is some evidence suggesting that this PPAR γ activation following blockade of the AT₁ receptor could be part of its anti-inflammatory and antioxidative effects, leading to neuroprotection against ischemia and A β accumulation [127, 194, 195]. Indeed, neuroprotective effects of PPAR γ agonists, such as pioglitazone, have been observed during neural cell differentiation and death, and in inflammatory and neurodegenerative conditions, including amyotrophic lateral sclerosis, Alzheimer's disease and Parkinson's disease models, as well as stroke [196, 197]. PPAR γ is a transcriptional factor regulating the expression of multiple genes, thereby promoting the differentiation and development of various tissues, specifically adipose tissue, brain, placenta, and skin (review in [197]). In addition, certain studies have indicated that AT₂ receptor stimulation increases PPAR γ expression and transcriptional activity, at least in PC2W cells [57] and neurons [131]. The final targets of these pathways are gene expression and phosphorylation of various microtubule-associated proteins, which modulate microtubule stability/dynamics responsible for neurite elongation. This observation is noteworthy, especially with regard to

the implication of PPAR γ in NGF-induced neurite outgrowth [198] which clearly suggests a possible crosstalk between the AT $_2$ receptor and NGF pathways. This hypothesis is furthermore reinforced by the observation that inhibition of the NGF receptor TrkA significantly decreases AT $_2$ receptor-induced neurite outgrowth [43]. Moreover, Iwai et al., using atherosclerotic ApoE-KO mice with an AT $_2$ receptor deficiency (AT2R/ApoE double knockout mice), observed that the lack of AT $_2$ receptor expression decreased the expression of PPAR γ in adipocytes [199]. These observations strongly suggest a link between the AT $_2$ receptor and PPAR γ functions. Considering that similar neuroprotective effects were also associated with the AT $_2$ receptor, it may be hypothesized that activation of PPAR γ could be shared both by ARBs and AT $_2$ receptor stimulation.

9. Could the AT $_2$ Receptor Be an Attractive Therapeutic Target?

One of the biggest challenges in studying the AT $_2$ receptor is applying observations stemming from the use of cell lines to *in vivo* models. Indeed, studies using cell lines expressing the AT $_2$ receptor either endogenously or via transfection have provided paramount information regarding its intracellular mechanisms of action. However, associating these mechanisms with biological functions has proven to be much more difficult. As indicated previously, synthesis and characterization of the selective AT $_2$ receptor selective agonist C21/M024 in 2004 or of the recently developed antagonist [200] has provided long-awaited tools to bypass the difficulty of using traditional AT $_2$ receptor ligands such as CGP42112A or PD123,319. Since then, many studies have allowed significant advances in the understanding of AT $_2$ receptor functions. Nonetheless, one would have thought that this new AT $_2$ receptor ligand would have resolved certain controversies surrounding this enigmatic receptor. However, eight years after the first characterization of this compound, the clear demonstration of an AT $_2$ receptor effect in the brain remains to be unequivocally established. One major input since C21/M024 was first described is that selective stimulation of the AT $_2$ receptor does not decrease blood pressure [134, 165, 166, 201–206]. These results are quite surprising, considering previous reports emanating from the indirect *in vivo* manipulation studies which associated AT $_2$ receptor activation with vasodilation and a decrease in mean arterial pressure. However, blockade of the AT $_1$ receptor with ARBs not only allows stimulation of the AT $_2$ receptor, the only Ang II receptor available in this condition, but also increases the bioavailability of Ang II for ACE2 and aminopeptidase to produce Ang IV and Ang (1–7), both of which exert vasodilatory effects (Figure 1). Nevertheless, beyond its blood-pressure lowering effects, *in vivo* studies using C21/M024 have described a protective role of the AT $_2$ receptor in vascular remodeling [166, 207], in poststroke cardiac [208] and renal function [165, 205] as well as in cognitive functions [134]. Furthermore, observation of AT $_2$ receptors in tissues associated with glucose metabolism, such as the pancreas and adipose tissue, suggests that the AT $_2$ receptor could also be beneficial in metabolic syndrome and associated

cognitive loss. However, the selectivity of C21/M024 for the AT $_2$ receptor has also been recently challenged and differs according to the dosage used and/or route of administration and, importantly, according to experimental conditions. In particular, a recent observation by Verdonk et al. [209], whereby C21/M024 induced vasorelaxation of precontracted iliac arteries from SHR Wistar rats and C57BL/6 mice as well as in AT $_2$ -deficient mice, has rekindled the debate on the relevance of the AT $_2$ receptor and the selectivity of AT $_2$ receptor ligands in physiological functions. Moreover, this effect of C21/M024 in arteries was only partially blocked by PD123,319. These latest findings raise the possibility that C21/M024 could exhibit certain AT $_2$ receptor-independent effects. Therefore, the question still remains: is the AT $_2$ receptor a potential therapeutic target and could it replace or increase beneficial effects associated with ARBs?

10. Conclusion

As described in the aforementioned sections, AT $_2$ receptor activation may act at several stages in the cascade of alterations leading to cognitive impairment and neuronal dysfunction. An increasing number of studies suggest that the protective effects of ARBs on brain damage and cognition may result not only from the inhibition of AT $_1$ receptor effects, but also from the beneficial effect due to unopposed activation of the AT $_2$ receptor. In addition, the relationship between impaired energy metabolism/obesity/insulin resistance and the increased risk of dementia emphasizes the view that the mechanisms of action of the AT $_2$ receptor may have a beneficial protective effect. However, the physiological relevance of the AT $_2$ receptor in the brain will need to be compared with at least two other components of RAS, namely, the ACE2/Ang-(1–7)/Mas complex and Ang IV/AT $_4$ receptor/IRAP in the brain (Figure 1).

Lifestyle-related disorders, such as hypertension, T2D, and obesity, are also implicated as risk factors for dementia. In this regard, two recent publications have established that direct AT $_2$ receptor stimulation with C21/M024 improves insulin sensitivity in a rat model of diet-induced insulin resistance [210] and in type 2 diabetic mice [211]. Thus, any treatment aimed at improving insulin resistance or cognitive functions is likely to slow down symptoms and improve quality of life associated with these age-related disorders. Furthermore, the recent development of selective AT $_2$ receptor agonists should facilitate efforts to elucidate distinct roles of the AT $_2$ receptor in brain physiology, supporting or disproving the hypothesis that the AT $_2$ receptor helps improve a number of brain impairments related to neuronal plasticity and morphology, microcirculation and inflammation (Figure 3), all of which are altered in certain neurological disorders.

There is clearly still much work to be accomplished to fully understand the role of the AT $_2$ receptor in normal versus pathological conditions and to determine whether AT $_2$ receptor agonists could represent an attractive therapeutic target. In this aspect, compounds such as C21/M024 as well as other recently synthesized highly selective nonpeptide AT $_2$ receptor ligands (all leading to neurite outgrowth in NG108-15 cells) and their effectiveness to induce AT $_2$

receptor-dependent effects need to be further explored [212–216].

Acknowledgments

The work presented in this paper was supported by Grants from the Canadian Institutes of Health Research (MOP-82819 to N. Gallo-Payet) and from the Alzheimer's Society of Canada to Dr. Gallo-Payet with Louis Gendron (*Université de Sherbrooke*) and Thomas Stroh (McGill University) and by the Canada Research Chair program to Dr. Gallo-Payet. N. Gallo-Payet is a past holder of the Canada Research Chair in Endocrinology of the Adrenal Gland. M. O. Guimond is a postdoctoral fellow in the laboratory of Dr. Gallo-Payet. They are both members of the FRSQ-funded *Centre de recherche clinique Étienne-Le Bel*. The authors are grateful to Pierre Pothier for critical reading of the paper and editorial assistance (Les Services PM-SYS Enr., Sherbrooke).

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Review Article

The Prorenin and (Pro)renin Receptor: New Players in the Brain Renin-Angiotensin System?

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Received 5 October 2012; Revised 26 November 2012; Accepted 5 December 2012

Academic Editor: Ovidiu C. Baltatu

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It is well known that the brain renin-angiotensin (RAS) system plays an essential role in the development of hypertension, mainly through the modulation of autonomic activities and vasopressin release. However, how the brain synthesizes angiotensin (Ang) II has been a debate for decades, largely due to the low renin activity. This paper first describes the expression of the vasoconstrictive arm of RAS components in the brain as well as their physiological and pathophysiological significance. It then focus on the (pro)renin receptor (PRR), a newly discovered component of the RAS which has a high level in the brain. We review the role of prorenin and PRR in peripheral organs and emphasize the involvement of brain PRR in the pathogenesis of hypertension. Some future perspectives in PRR research are heightened with respect to novel therapeutic target for the treatment of hypertension and other cardiovascular diseases.

1. Introduction

The renin-angiotensin system (RAS) plays an important role in the physiological and pathophysiological regulation of blood pressure (BP) and cardiovascular function. Renin, the rate-limiting enzyme of the RAS, was discovered more than a hundred years ago by Tigerstedt [1]. Angiotensinogen (AGT) is cleaved to angiotensin (Ang) I by renin, which is released from the juxtaglomerular apparatus. Ang I is then further converted into the octapeptide, Ang II, by the angiotensin-converting enzyme (ACE). Ang II is the main effector peptide in this system. Via binding to Ang II type 1 receptor (AT1R), Ang II stimulates vasoconstriction and secretion of the steroid hormone, aldosterone, which mediates sodium reabsorption and water retention [2]. In addition to the classical view of endocrine RAS, local RAS has been identified in various tissues, which interacts with the endocrine RAS [3–6].

The first evidence for the existence of brain RAS was demonstrated by Brickerton and Buckley [7] who showed that administration of Ang II into the brain caused an increase in BP. It is well accepted now that RAS is intrinsic to the brain and plays an integral role in the pathophysiological

regulation of cardiovascular function [8, 9]. The activities of brain RAS are achieved by influencing the autonomic nervous system, the baroreflex sensitivity, vasopressin (AVP) release, and the thirst and salt appetite [10–13]. Brain Ang II also acts as a neuropeptide via AT1R to increase the excitability of neurons in the cardiovascular regulatory centers of the hypothalamus and brainstem [14, 15]. Chronic central infusion of Ang II elicits an increase in arterial pressure and enhances the sensitivity of cardiac sympathetic afferent reflex via AT1R [16]. Human renin and AGT double transgenic mice exhibit Ang II-dependent hypertension, which can be attenuated by AVP antagonist suggesting an interaction between the brain RAS and AVP in hypertension [17, 18].

Despite a multitude of evidence supporting the importance of Ang II action in the brain, conjecture remains about how Ang II forms locally in the brain because brain renin activity is undetectable using currently available methods [19, 20]. Prorenin has been recognized as a precursor of renin, but with very low angiotensinogenase activity [21]. Interestingly, we recently reported the existence of prorenin protein in brain tissues and the brain prorenin levels were ten-fold higher compared with renin levels [22].

The (pro)renin receptor (PRR) is a newly discovered component of the RAS which is able to promote Ang II formation via binding to renin or prorenin [23]. This timely discovery may shed new light on a possible pathway for Ang II formation in the brain regions behind the blood-brain barrier. Our paper focuses on the influence of brain RAS on the regulation of cardiovascular function with a specific emphasis on recent evidence concerning the role of brain PRR in the regulation of BP and cardiovascular homeostasis.

2. The Brain RAS

Most brain areas are separated from the circulation by the blood-brain barrier which is impermeable to Ang II [24]. Therefore, Ang II needs to be generated locally or transported via transcytosis mechanism [25] to interact with its receptors located on neurons and astrocytes of the central nervous system (CNS). Genetic and pharmacological studies have demonstrated the intrinsic existence of brain RAS components [8, 9]. Furthermore, the pathophysiological significance of a brain RAS is supported by observations of increased RAS activities in the brain cardiovascular regulatory areas of hypertensive animal models [26, 27].

3. Renin

Renin is the rate-limiting enzyme for the formation of Ang I. Renin-like enzymatic activity within the brain was first reported by Ganten et al. [28]. However, the isolation of detectable active renin from the brain has remained unsuccessful for decades after its discovery [29]. Although the brain contains a large amount of enzymes, such as cathepsins [30] and tonin [31], which can also generate Ang peptides from AGT, the relevance of these alternative enzymatic pathways for Ang II generation in the brain remains elusive [32]. The most direct evidence supporting renin gene expression in the brain has come from Dr. Sigmund's group [33] who utilized an enhanced green fluorescence protein (eGFP) gene as a reporter for renin expression. They found renin promoter activity in the neurons of the cerebellum and hippocampus and in the cardiovascular regulatory regions such as the rostral ventrolateral medulla (RVLM), the subfornical organ (SFO), the paraventricular nucleus (PVN), and the supraoptic nucleus (SON). This group also discovered two different forms of renin in the brain, the intracellular (icRenin) and secret (sRenin) renin, derived from two different renin transcripts [34]. The relevance of icRenin is still unclear. This renin is a truncated form of prorenin lacking the first third of the prosegment, which remains intracellularly, whereas the sRenin secretes to the extracellular space. Due to the lack of machinery to cleave prorenin to renin in the brain [35], the sRenin in the brain is possibly prorenin. Although true renin activity exists at a finite level in the brain, trypsin or acid treated brain extracts showed a marked increase in renin activity. In addition, the inactive form of renin has been successfully isolated from the brain which could be activated by trypsin and inhibited by antirenin antibody *in vitro* [36]. This inactive

form of renin may be the main source of renin-like activity in the brain. Pharmacological studies have provided strong support for the renin signal in the brain. For example, despite the difficulty of directly measuring renin activity in brain tissue, intracerebroventricular (ICV) infusion of the renin inhibitor, aliskiren, prevented sympathetic hyperactivity and hypertension as well as desensitization of arterial baroreflex function in Dahl salt-sensitive rats on a high salt diet [37]. These *in vivo* studies provide support for the existence of renin or renin-like activity in the brain and its role in hypertension.

4. Angiotensinogen and ACE

In the CNS, the main source of AGT synthesis is from the astroglia [38]. However, AGT expression has also been found in pure neuron cultures [39]. The AGT protein and its transcripts have been reported in the cardiovascular regulatory regions of the brain by different groups [40–42]. Transgenic rats expressing an antisense RNA against AGT mRNA specifically in the brain exhibited lower BP, polyuria, and reduction in plasma AVP [43] indicating that AGT is synthesized in brain and plays an essential role in BP regulation.

The ACE is a zinc metalloprotease which hydrolyzes the carboxyl terminal dipeptide His-Leu of Ang I to form Ang II. High densities of ACE were visualized in the choroid plexus, SFO, the caudate putamen, and the substantia nigra by autoradiography [44]. The ACE activity is present in the renin-containing synaptosomes of the neurons suggesting that intraneuronal synthesis of Ang I and Ang II is possible in synaptosomes [45]. The ICV delivery of the human ACE gene increased sympathetic activity, BP, and heart rate which were accompanied by increased Ang II and AVP production. These effects were abolished by ICV administration of an ACE inhibitor suggesting the importance of brain ACE in Ang II formation [26].

5. Angiotensin II and Angiotensin II Type I Receptor

The actions of Ang II are mediated predominantly by a seven transmembrane domain Gq-protein coupled receptor designated as AT1R. Effects of the AT1R are mediated by multiple intracellular signaling pathways, starting with G-protein and phospholipase activation, followed by an increase in intracellular inositol trisphosphate and calcium, which result in vasoconstriction, cell proliferation, and fibrosis [46, 47]. The AT1R is localized with high densities in the anterior pituitary, area postrema, lateral geniculate body, inferior olivary nucleus, median eminence, nucleus of the solitary tract (NTS), the anterior ventral third ventricle region, PVN, SON, and SFO [48]. The AT1R expression levels are tightly regulated by cardiovascular status and Ang II levels. Higher levels of AT1R were found in the hypothalamus of spontaneously hypertensive rat (SHR) and the SFO, PVN, and NTS of animals with chronic heart failure [27, 49]. Others have shown that chronic Ang II infusion

upregulates AT1R mRNA and protein levels in the PVN and SFO via activation of intracellular mitogen-activated protein kinase (MAPK) signaling pathways [50]. Constitutive AT1R overexpression in the astroglia of RVLM resulted in a chronic increase in BP indicating that increased AT1 receptor activity is a primary determinant of efferent drive from RVLM [51]. Although Ang II is a small octapeptide, it is unable to permeate the blood-brain barrier and the blood-cerebrospinal fluid barrier [24, 52]. However, Ang II immunoreactivity has been found in the cell bodies of magnocellular and parvocellular neurons in the PVN and the magnocellular neurons in the SON [53] and Ang II-stained fibers have been found at all levels of the CNS, from the olfactory bulbs to the spinal cord. Furthermore, brain Ang II content was significantly increased in bilaterally nephrectomized rats despite diminishment of plasma Ang II to a very low level [54]. The body of evidence suggests that brain Ang II is synthesized locally and can be regulated independently of peripheral Ang II.

6. Prorenin

Prorenin, the precursor of renin, is cleaved to its active form by the removal of the 43 amino acid prosegment [21]. Whereas the renal juxtaglomerular cells constitute the most important source of circulating renin [55], a number of extrarenal tissues including the adrenal glands, ovary, testis, placenta, and retina produce prorenin [56]. This also explains the finding that the plasma prorenin level is 10 fold higher than that of renin [57]. However, prorenin has very low enzymatic activity in the plasma. Only two percent of prorenin exists as the “open form”, in which the prosegment of prorenin undergoes a conformational change and exposes the enzyme’s active site [58]. This process is also called nonproteolytic activation of prorenin. The role of non-proteolytic activation of prorenin remains debatable due to the controversial phenotype in prorenin transgenic animal models. Liver-specific transgenic rats with a 400-fold increase in circulating prorenin exhibited severe renal lesions and hypertrophic cardiomyocyte with normal BP [59]. In contrast, transgenic animals with inducible or constitutive overexpression of prorenin, despite expressing 13–179 folds higher circulating prorenin, did not display cardiac or kidney damage. However, they did develop moderate Ang II dependent hypertension, since the BP was reduced by an ACE inhibitor [60, 61]. The reason for the presence of different phenotypes in the prorenin transgenic animal model remains unclear but could be due to the difference in the levels of plasma prorenin or the species. Although Mercure et al. [61] showed that Ang II is responsible for the increase in BP in the prorenin transgenic mice; the high plasma prorenin levels present in his study might not exist in physiological or even pathophysiological states [62]. The formation of Ang II in Mercure’s model may be due to the activity of the two percent “open form” prorenin in the plasma [63]. However, in all three models, the increase in prorenin levels is limited to the circulation and liver. Thus, the role of local or tissue prorenin in cardiovascular regulation remains inconclusive.

7. (Pro)renin Receptor

In the year 2002, a new component of RAS was identified in human mesangial cells and named the (pro)renin receptor (PRR) because PRR binds both renin and prorenin [23]. The PRR gene is identical to ATPase 6 accessory protein2 (ATP6AP2) and is located on the X chromosome. PRR gene encodes a 350-amino-acid and ubiquitously expresses a single transmembrane protein with a large N-terminal extracellular domain which binds both renin and prorenin with affinities in the nanomolar range [64, 65]. Immunofluorescence observed by confocal microscope demonstrated that PRR was located on the cell surface, as well as intracellular compartments especially on the perinuclear space [22, 23, 66]. *In vitro* binding studies have shown that prorenin binds to PRR with a 3-4 fold higher affinity than renin, indicating that prorenin is a preferred ligand for PRR compared with renin [64, 65]. The binding of renin to PRR increases enzymatic activity 5-fold higher than the nonreceptor-bound renin [23]. The binding of prorenin to PRR induces a conformational change; the prosegment is removed from the catalytic cleft and the active site is accessible to AGT leading to full nonproteolytic activation of prorenin [23]. Interestingly, this phenomenon is reversible and prorenin eluted from the receptor reverts to its inactive form. The binding of renin to PRR is independent of the active site and receptor-bound renin or prorenin is not internalized or degraded [64, 67]. Thus, the discovery of the PRR has shed light on an alternate pathway for nonproteolytic activation of prorenin. Although the formation and role of intracellular Ang II has been previously reported [68, 69], there is a lack of attention to whether the intracellular PRR and prorenin contributes to the intracellular Ang II formation. It is likely that in tissues lacking a mechanism to cleave prorenin to renin, activation of prorenin via binding to PRR may contribute to intracellular Ang II formation. Several *in vitro* studies have been completed to assess the ability of prorenin to form Ang II via binding to the PRR. At a concentration of nanomolar range, which is much higher than the circulation level in physiological conditions, prorenin binds to PRR and exhibits enzymatic activity similarly to that of renin [64]. Furthermore, a recent *in vitro* study showed that the PRR activation by prorenin to generate Ang II requires about 800 fold higher prorenin concentration above normal plasma levels; the Ang II-independent activation requires an even higher prorenin concentration [70]. Combining these observations, prorenin may act mainly in the tissues either extracellularly or intracellularly, where its concentration may be high enough to activate Ang II-dependent or independent signals [70].

Transgenic rats with human PRR expression in vascular smooth muscle cells have been shown to exhibit hypertension and increased plasma aldosterone at six months, suggesting a pathological role of PRR in raising BP [71]. This study speculated that the rise in BP may have been caused by increased plasma aldosterone levels in these rats, but whether the increase in aldosterone depends on Ang II was not directly tested by the authors. In another transgenic model, ubiquitous expression of human PRR in rats resulted

in proteinuria, glomerulosclerosis, MAPK activation, and cyclooxygenase-2 upregulation. These rats exhibited normal BP and renal RAS activity suggesting an Ang II independent nephropathy [72]. A previous *in vitro* study has shown that rat prorenin/renin was able to activate human PRR [65]. The phenotype of these transgenic rats could be due to the activation of human PRR transgenes by endogenous rat prorenin/renin. The reason for the differences in the phenotypes of these two models is not clear, but could be attributed to tissue targeting strategy differences. However, these studies clearly suggest that the pathological changes seen in these models might be due to the non-RAS dependent PRR rather than RAS dependent effects. In addition, binding of renin or prorenin to the PRR directly triggered intracellular MAPK signaling cascades in several cell types, which up-regulated the expression of profibrotic genes such as plasminogen activator inhibitor type (PAI)-1, collagens, and fibronectin [73–75]. Despite Ang II type 1 and 2 receptor blockade, renin or prorenin was still able to induce a long-lasting ERK 1/2 phosphorylation by binding to PRR, which was not blocked by aliskiren. Since aliskiren is a renin inhibitor which blocks the breakdown of angiotensinogen by renin, but does not affect the binding of renin to PRR and the PRR mediated signaling, it is clear that PRR mediates RAS independent signaling pathways [73].

Global PRR knockout is lethal in mice indicating an essential role of PRR in embryonic development [76]. Recently, several tissue specific PRR knockout mouse models have been generated; [77–79] these *in vivo* studies provide additional insight into functions of PRR which are independent of RAS activation. PRR knockout in cardiomyocytes of mice led to heart failure and death within four weeks after birth accompanied by deacidification of the intracellular vesicles [77]. This outcome was also reported in the mice with a PRR deletion in the podocytes [78, 79]. In both cases, the defect appears to be associated with an inability to acidify intracellular compartments and a dysfunction of vATPase, suggesting a PRR effect unrelated to RAS *in vivo*. Moreover, PRR functions as a physiological adaptor between vATPase and the Wnt receptors [80]. Wnt via binding to its receptor, low-density lipoprotein receptor-related protein 6 (LRP6), induces receptor aggregation and phosphorylation of LRP6, resulting in the stabilization of β -catenin. The Wnt/ β -catenin signaling is fundamental for a normal patterned embryo. In the adult, however, Wnt signaling is involved in cell proliferation and tissue homeostasis and has been implicated in certain pathologies, such as cancer and diabetes. By employing a genome-wide small inhibitory RNA (siRNA) screen, Cruciat et al. [80] identified PRR as part of the Wnt receptor complex, acting as a specific adaptor between LRP6 and vATPase where PRR and vATPase mediated Wnt signaling during anteroposterior patterning of Xenopus early CNS development.

8. The PRR in the CNS

A mutation of the PRR gene, resulting in frame deletion of exon 4 is associated with X-linked mental retardation

and epilepsy pointing to an important role of PRR in the CNS [81]. PRR mRNA is widely expressed in various regions of the human brain with the highest expression levels in the pituitary and frontal lobe [82]. Immunostaining showed that PRR is colocalized with oxytocin and AVP in the magnocellular neurons of the PVN and SON indicating that PRR may be related to the central control of water-electrolyte homeostasis and BP [82]. Similarly, a wide distribution of PRR mRNA was found in key regions of the mouse brain involved in the regulation of BP and body fluid homeostasis [83]. Furthermore, PRR protein is expressed throughout the brain in cardiovascular regions including the SFO, PVN, nucleus of raphe pallidus, NTS, and RVLM as well as in noncardiovascular regulatory regions [22]. Immunofluorescence staining for PRR, neuron-specific nuclear protein (NeuN, neuron marker), and glial fibrillary acidic protein (GFAP, astroglia marker) revealed that PRR is expressed in astroglia, but with prominent expression in the neurons (Figure 1). Neuronal cells from the hypothalamus and brainstem of normotensive rat brains express PRR with levels 3-fold higher than that seen in astroglial cells from the same brain areas [66]. Moreover, PRR mRNA and protein levels were increased in brain cardiovascular regulatory regions in hypertensive animals [22, 84]. Knockdown of PRR in the SON was associated with attenuation of hypertension and a decrease in plasma AVP in SHR [84]. We recently reported that PRR knockdown in the brain attenuated Ang II-dependent hypertension in human renin and AGT double transgenic mice [22]. This effect was associated with a decrease in AVP levels, sympathetic tone, and improvement of baroreflex sensitivity indicating a role of PRR in the autonomic regulation of hypertension. However, whether PRR regulates BP and cardiovascular homeostasis via Ang II dependent or independent pathways remains inconclusive.

Shan et al. [84] demonstrated that coincubation of human prorenin and AGT in isolated neurons evoked a dose- and time-dependent increase in Ang I and II formation indicating the ability of prorenin to generate angiotensin peptides in neurons possibly by binding to PRR. To address whether PRR mediates Ang II formation in the CNS, our laboratory recently measured Ang II levels in human renin and AGT double transgenic mice following brain-targeted PRR knockdown using PRR short hairpin RNA; Ang II levels were significantly decreased after PRR knockdown in the hypothalamus (Figure 2) and the levels were associated with a decrease of PRR levels in this region, as reported previously [22]. These data show that in hypertension, PRR may be involved in Ang II formation in the CNS. We also recently reported that the prorenin protein exists in mouse brain tissue, and its expression level is 10-fold higher than that of renin. In light of the facts that (1) there is abundant PRR expression in the brain, (2) prorenin exists in the brain despite an extremely low renin activity, and (3) manipulation of PRR modulates the Ang II levels; we propose that the binding of prorenin to PRR may initiate the rate-limiting step for angiotensin peptides formation in the CNS. The limitation of the knock down-based hypothesis is that PRR also acts as an accessory protein for vATPase, which is important for vesicular acidification, an important

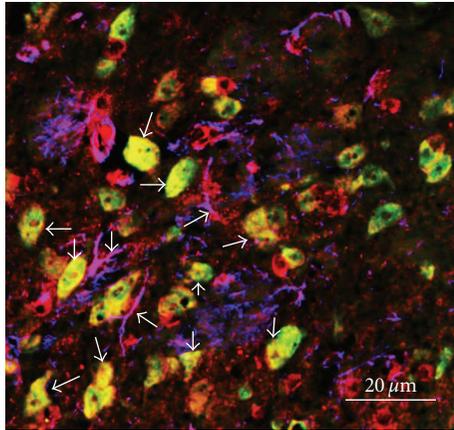


FIGURE 1: (Pro)renin receptor expression in the C57Bl/6J mouse brain. Triple immunofluorescence staining for PRR (red), neuron-specific nuclear protein (green, neuron marker), and glial fibrillary acidic protein (blue, astroglia marker) revealed that PRR is expressed in both neurons and astroglia with prominent expression in the neurons. Arrows indicate the colocalization of PRR with neurons or astroglia.

function in neuronal cells. Although the role of vATPase in Ang II formation has not been reported, it is difficult to distinguish the effects of PRR silencing on RAS-related prorenin inactivation and vATPase dysfunction. So far, the role of RAS independent signaling pathways of PRR in the CNS remains unknown.

9. Conclusion and Perspective

Emerging evidence supports the argument that brain RAS signals contribute to the development of hypertension. Interruption of these signals is beneficial to the control of BP in hypertension. All components of the RAS exist in the brain, but the renin level is extremely low. Prorenin and PRR may be the missing piece of the puzzle of the brain RAS since prorenin is conferred a nonproteolytic activation when binding to the PRR. Although further study is needed to test this hypothesis, evidence which support this case include that: (1) prorenin is the dominant form of total renin (prorenin and renin) in the brain, (2) PRR is highly expressed in the CNS, (3) brain PRR expression levels are increased in several hypertensive animal models, (4) knockdown of the PRR in the hypothalamus is associated with reduction in Ang II formation in this region during hypertension, and (5) reduced PRR expression level is associated with reduced BP in Ang II-dependent hypertension. Conversely, the RAS-unrelated PRR function via vATPase is critical to acidification of the intracellular compartments and might be important in the modulation of autonomic function since vATPase is responsible for neurotransmitter transportation and storage in the neuron vesicles [85]. We conclude that PRR might be an essential component of the tissue RAS, at least in the CNS. On the other hand, the RAS-unrelated PRR function in the CNS needs further investigation. Thus, targeting PRR can be an innovative, new strategy for the

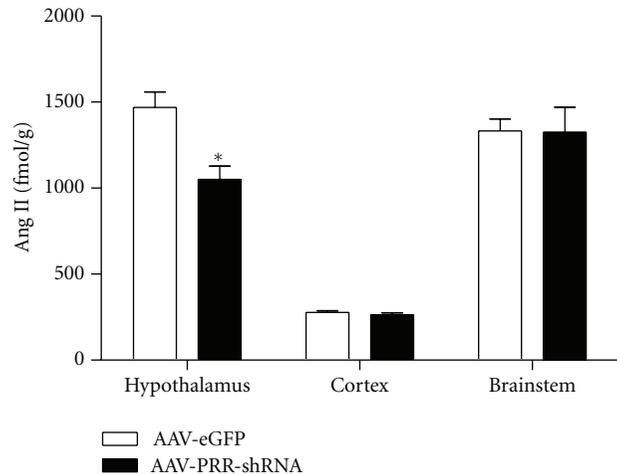


FIGURE 2: ICV delivery of (pro)renin receptor shRNA reduces brain Ang II level in the hypothalamus. The human renin and AGT double transgenic mice were ICV injected with AAV-PRR short hairpin RNA (AAV-PRR-shRNA, 3.5×10^{11} Vg/100nl) or control virus. Two weeks after virus injection, brain tissues were harvested for analysis of Ang II levels in the hypothalamus, cortex, and brainstem. ($n = 6$ /group). * $P < 0.05$ versus AAV-enhanced green fluorescent protein (AAV-eGFP) treatment.

treatment of hypertension. Understanding the physiological and pathophysiological role of PRR in hypertension models would move this possibility forward. Because the PRR plays an essential role in embryonic development, generation of PRR antagonist or inducible tissue-specific PRR knockout animal models would be ideal tools to study the role of PRR in adult diseases models.

Funding

The work was supported by American Heart Association National Center (11SDG4880005) to Y. Feng.

Acknowledgments

The authors thank Nancy B. Busija of the copy-editing for this paper and the Mouse Phenotyping Core at the Tulane Hypertension Center of Excellence for the support to this study.

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Review Article

Upregulation of the Renin-Angiotensin-Aldosterone-Ouabain System in the Brain Is the Core Mechanism in the Genesis of All Types of Hypertension

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Received 4 October 2012; Accepted 23 November 2012

Academic Editor: Marc de Gasparo

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Basic research using animal models points to a causal role of the central nervous system in essential hypertension; however, since clinical research is technically difficult to perform, this connection has not been confirmed in humans. Recently, renal nerve ablation in humans proved to continuously decrease blood pressure in resistant hypertension. Furthermore, when electrical stimulation was continuously applied to the carotid baroreceptor nerve of human adults, their blood pressure lowered. These findings promoted the concept that the central nervous system may actually be involved in the pathogenesis of essential hypertension, which is closely associated with excess sodium intake. We have demonstrated that endogenous digitalis plays a key role in hypertension associated with excess sodium intake via sympathetic activation in rats. Increased sodium concentration inside the brain activates epithelial sodium channels and the renin-angiotensin-aldosterone system in the brain. Aldosterone releases ouabain from neurons in the paraventricular nucleus in the hypothalamus. Angiotensin II and aldosterone of peripheral origin reach the brain to augment sympathetic outflow. Collectively essential hypertension associated with excess sodium intake and obesity, renovascular hypertension, and primary aldosteronism and pseudoaldosteronism all seem to have a common cause originating from the central nervous system.

1. Introduction

Blood pressure (BP) is a physiological phenomenon like body temperature, respiration, and pulse rate; its dysregulation most often results in hypertension. BP regulation is operated via sympathetic and parasympathetic nerves, as well as the release of pituitary hormones. However, antihypertensive agents such as diuretics, beta blockers, alpha blockers, calcium channel blockers (CCBs), mineralocorticoid receptor blockers (MRBs), and antagonists for the renin-angiotensin system (RAS) effectively lower the BP, and their sites of action lie in the peripheral tissues. These antihypertensive treatments seem to have little or no connection with the central nervous system (CNS) regulation of BP; thus, if a central mechanism does exist, its actual regulatory role is believed to be minimal, and, therefore, any such central mechanism of hypertension has not been investigated extensively. However,

evidence from animal models has demonstrated the role of sympathetic nervous system activity (SNSA) in hypertension [1–3]. For example, renal denervation consistently lowers the BP in a variety of animal models of hypertension [4–6]. In line with this, renal denervation by catheter-based radiofrequency ablation technology has been shown to effectively lower BP in patients with resistant hypertension at least for 2 years [7]. As Guyton's model [8] predicted that the renal function to excrete sodium is the infinite determinant of hypertension, denervation natriuresis may be the mechanism of action of renal denervation, demonstrating the crucial role of peripheral SNSA. Furthermore, interventional activation of the carotid baroreflex via continuous electrical stimulation with an implantable device has been shown to effectively lower BP by decreasing SNSA in patients with resistant hypertension [9]. This evidence indicates that the central setting level of BP regulation is elevated at the higher level

in hypertensive patients. These 2 important observations led us to consider the role of SNSA in the genesis of hypertension in humans.

In addition to these findings, other evidence to confirm the role of CNS in the genesis of hypertension has been revealed: a small dose of angiotensin II (Ang II) continuously administered subcutaneously (SC) gradually elevates BP, which can be abolished by intracerebroventricular (ICV) pretreatment with either an aldosterone synthase inhibitor or an MRB in rats [10]. Similar findings have been demonstrated with aldosterone: SC administration of aldosterone with 1% NaCl saline, as drinking water in rats gradually elevates BP, which is blunted by ICV pretreatment with either an Ang II AT-1 receptor blocker (ARB) or MRB [11]. These observations indicate that secondary hypertension such as primary aldosteronism, pseudoaldosteronism, and renovascular hypertension should be classified as centrally induced hypertension. Furthermore, central abolition of BP increases with ARB or MRB treatment, indicating that the major classes of antihypertensive agents may primarily be acting at a central site to lower BP. In fact, reflex tachycardia is absent when treated with the abovementioned antihypertensive agents, indicating that these agents lower BP by acting at a central site to set the BP to a lower level. In this paper, these are the particular points of discussion.

2. Arterial Hypertension and SNSA

The role of the autonomic nervous system activity in the pathogenesis of essential hypertension has been extensively studied [12–14]. Measurement of urinary excretion of norepinephrine has shown increased SNSA in animal models of hypertension, such as spontaneously hypertensive rats [15], Dahl salt sensitive rats [16], and models of deoxycorticosterone acetate (DOCA)-salt hypertension [17], Goldblatt renovascular hypertension [18], and renal mass-reduced hypertension [19]. Furthermore, there are numerous reports showing elevated urinary excretion of catecholamines in humans [20]; however, the difference between normal controls and hypertensive subjects is not high enough to establish the role of SNSA in hypertension.

On the other hand, SNSA is notably elevated in young subjects with labile hypertension who are believed to be in an early phase of essential hypertension [21]. SNSA is clearly increased among obese subjects with metabolic syndrome who are prone to be hypertensive and is decreased with a significant reduction in body weight [22, 23]. Augmented SNSA may be due to increased leptin, which is secreted from fat cells and stimulates SNSA via actions on the arcuate nucleus in the hypothalamus, and suppresses appetite [24]. When rats are fed a high-fat diet, they become resistant to appetite suppression but not resistant to sympathetic activation leading to hypertension [25]. However, SNSA is not much augmented in those hypertensive animals and humans, rather it is nearly normal due to suppression by the BP elevation caused by the initial increase in SNSA; that is, the SNSA converges to a near-normal level after a certain level of increase in the BP caused by an initial increase in

SNSA. The subtle elevation of SNSA will be maintaining the elevated BP.

3. Sodium, SNSA, and Hypertension

Our ancestors living inland during the Stone Age would have consumed natural foods like wild animals do, and sodium intake would have been suitable at the minimum for survival. Epidemiological surveys of Yanomamo Indians in Amazon, Brazil, who, until recently, had been living like the people in the Stone Age did, revealed that their urinary excretion of sodium was only 0.9 mmol/day (equivalent to 0.53 g of sodium chloride) [26]. This is roughly 1/20th of the average sodium intake in modern societies, and this small amount of sodium was proven to be sufficient to live. In fact, all vertebrates, except human beings and domestic animals, consume only natural food without adding sodium salt. The average BP of the Yanomamo Indians was 96.0/60.6 mmHg, and the level did not elevate depending on age, which is in contrast to the observation in modern populations. The average life span of people who lived in the Stone Age is thought to be approximately 30 years, which is too low to cause atherosclerotic vascular complications, even if hypertension existed. Rather, higher BP would be better to supply enough blood to the principal organs and skeletal muscles. Renovascular hypertension may be a good example: when renal blood supply is impaired due to narrowing of renal arteries, the RAS is activated to raise the BP to maintain sufficient blood supply to the kidney. Thus, hypertension might have been a benefit for people in the Stone Age who were often attacked by enemies and wild animals; people who had a prompt elevation in BP could survive and leave offspring. To maintain a high BP, large amount of sodium would be needed, and thus, salt-sensitive subjects would be selected to survive. Those whose bodies promptly elevated the BP upon exposure to stress could succeed in genealogical history. Therefore, their offspring are the people living now, and we are salt-sensitive and prone to hypertension. Mechanisms to retain as much sodium as possible were required, the most powerful of which is the renin-angiotensin-aldosterone system (RAAS) [27]. In subjects who consume low amounts of sodium, the RAAS is working at full strength, as is the case for patients with Bartter syndrome. In addition, increased sympathetic outflow accelerates sodium reabsorption from the renal tubules via renal nerves [28]. Similarly, insulin acts to retain sodium available at renal tubules; this action is exaggerated in the early stage of diabetes mellitus and metabolic syndrome associated with obesity. The precise mechanism of how insulin retains sodium at the renal tubules is now well known.

Together, strong sodium-retaining mechanisms were a desirable trait and selected in later generations. Therefore, hypertension easily develops after excess intake of sodium. This trait was selected to correct our life style with a lower intake of sodium; as a result, the incidence of stroke is markedly decreased in Japan [29]. Understanding the mechanism of induction of hypertension after excess intake of sodium will help to devise efficient measures to control

the BP even after excessive sodium intake. A recent approach to elucidate these mechanisms revealed that the brain RAAS and endogenous digitalis are essentially involved in the induction of hypertension.

4. Endogenous Digitalis underlies the Connection of Sodium and Hypertension

Continuous administration of mineralocorticoids causes sodium accumulation, which results in natriuresis at a certain point, described as an escape phenomenon [30]. Factors associated with natriuresis include the glomerular filtration rate of the kidney, aldosterone, and other factors. The others are referred to as third factors [31]. Since it is well known that endogenous digitalis, an inhibitor for Na^+/K^+ -ATPase, circulates and that the activity of tissue Na^+/K^+ -ATPase decreases during sodium loading [32], endogenous digitalis is the most probable candidate for a third factor. Digoxin has been used for treating patients with congestive heart failure or tachyarrhythmias. Because it may cause intoxication, serum digoxin-like immunoreactivity (DLI) levels have been monitored, and significant levels of DLI have been detected in subjects not consuming digitalis glycosides [33]; this suggests the presence of endogenous digoxin. In fact, digoxin has been detected in human plasma by liquid chromatography and mass spectrometry [34]. Further, the turnover ratio of DLI in the hypothalamus and its plasma concentration increase with increasing sodium loading [35]. This suggests that digoxin may be produced in the hypothalamus and released into circulation. In fact, increased DLI concentrations were detected in the plasma of DOCA-salt hypertensive rats [36]. In humans, urinary excretion of DLI was found to be correlated with urinary excretions of sodium or BP during a medical checkup [33]. However, the DLI concentrations were not high enough to completely explain the pressor mechanism.

Ouabain, another water-soluble cardenolide, is known to be of endogenous origin [37]. Administration of low-dose ouabain causes sustained elevation of BP [38]. The Milan hypertensive rat model is known to have a mutation coding adducin, which augments the renal Na^+/K^+ -ATPase to increase sodium accumulation and hypertension [39]. In this model, plasma ouabain levels are increased. PST2238, or rostauroxin, is an analogue of digitoxigenin, and it antagonizes the action of ouabain and decreases BP in Milan hypertensive rats [40]. Since hypertension induced by low-dose ouabain is abolished by rostauroxin [41], ouabain must be directly involved in causing hypertension. Like Milan hypertensive rats, humans with adducin polymorphisms have elevated circulating ouabain levels, and rostauroxin was found to be effective in lowering their BP [42]. However, results of a large-scale clinical study did not conclusively prove this [43].

Immunohistochemical approach using anti-ouabain or anti-digoxin antibody revealed that neurons in the paraventricular nucleus (PVN) and supraoptic nucleus (SON) showed ouabain- or digoxin-like immunoreactivity (OLI or DLI) [44–46]. Both OLI and DLI are detected not only in

neuronal cell bodies but also in their dendrites and varicosities, which resembles the distribution of neuroendocrine hormones. The nerve fibers densely extend to the median eminence, subfornical organ (SFO), and organum vasculosum of the laminae terminalis (OVLT). Sodium loading increased the turnover ratio of DLI in the hypothalamus [35], and destruction of microtubules by ICV injection of colchicine increased DLI in the hypothalamus [35]. Therefore, DLI is clearly produced in the hypothalamus, PVN, and SON, which are stimulated by sodium loading. Although we did not explore similar experiments concerning the OLI, similar results are expected.

Since a low dose of ouabain injected into the lateral ventricle or the hypothalamus of rats increases the BP along with increasing the peripheral SNSA [47, 48], it seems likely that there are receptors for ouabain and endogenous ouabain acts as a neurotransmitter to increase SNSA.

Immortalized N1 cells are thought to be of PVN origin because these cells release vasopressin and oxytocin when stimulated [49]. Experiments on N1 cell line cultured in a serum-free condition showed that ouabain was released from these cells in a time-dependent manner; therefore, ouabain must be produced in the hypothalamus.

5. Brain RAAS and Sodium Metabolism (Figure 1)

In a clinical setting, BP in essential hypertension can be easily controlled with excellent antihypertensive agents such as diuretics, CCBs, angiotensin I converting enzyme inhibitors (ACEIs), ARBs, and MRBs. Sites where these agents act are crucial for BP regulation. In particular, ACEIs and ARBs are very effective in controlling BP and preventing complications, which indicates that RAAS is essential to the pathogenesis of hypertension. These drugs are effective in high-renin patients but still serve to lower BP even in those with low plasma renin activity (PRA) [50]. The RAAS has not been considered to be involved in sodium-sensitive hypertension because PRA is suppressed with sodium loading. On the other hand, in addition to the renal-renin and adrenal aldosterone system, RAAS was found to exist in other tissues such as the salivary gland and brain [51]. When sodium was loaded, the classical RAAS was suppressed, but the brain RAAS was activated [52]. The classical RAAS regulates sodium balance via a negative feedback, but the brain RAAS forms a positive feedback cycle to retain more sodium via increased renal SNSA. In the CNS, sodium loading upregulates the messenger RNA for renin, ACE, and angiotensin II AT-1 receptor [53]. Therefore, ICV administration of Ang II augmented pressor responses in sodium-loaded rats. Furthermore, aldosterone is present in the brain, and its level increases with sodium loading, leading to increased SNSA [54]. The hypotensive effects of MRB are greater in the low-renin essential hypertensive patients than in the normal renin hypertensives [55], which may be explained by the activated RAAS in the brain. Since Ang II administered into the CNS increases the BP along with increasing the SNSA [56], augmented RAAS in the brain

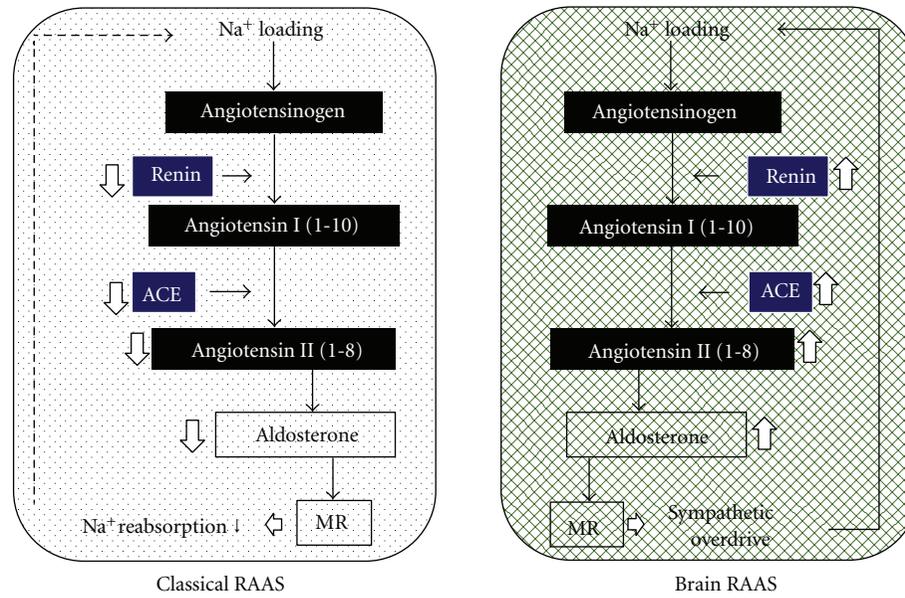


FIGURE 1: The classical RAAS (renal-renin and adrenal aldosterone system) regulates sodium balance via a negative feedback, but the brain RAAS forms a positive feedback cycle to retain more sodium via increased renal SNSA.

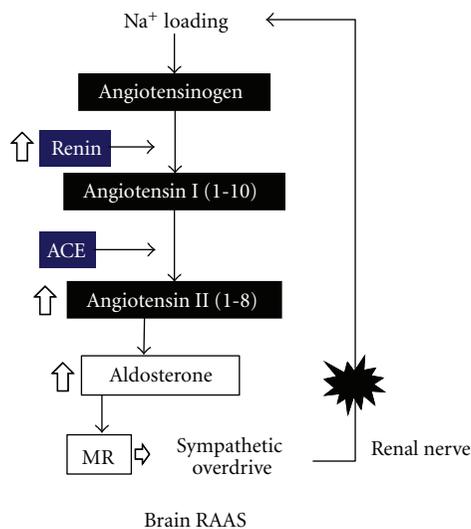


FIGURE 2: The most important component of the positive feedback cycle may be the renal nerve because renal nerve ablation lowers blood pressure even in humans, and inhibitors of RAAS lower blood pressure even in low-renin essential hypertensives.

induced by sodium loading may be a probable cause of essential hypertension. ICV injections of hypertonic saline cause elevation of both BP and plasma DLI, which can be blocked by ICV pretreatment with ARB [57]. Therefore, increased brain RAAS by sodium has been suggested to be involved in hypertensive actions of sodium. Thereby, abdominal SNSA, an upstream event of renal nerve stimulation, is markedly increased, as confirmed by the decreased renal blood flow observed using radioactive microspheres that indicate activated renal SNSA [58]. Increased renal SNSA

decreases renal excretion of sodium by decreasing renal blood flow and increasing renal tubular sodium reabsorption [59]. Thus, increased intracranial sodium concentration forms a positive circuit to retain sodium in the body (Figure 2).

6. Epithelial Sodium Channels May Be a Sensor for Sodium Concentration

The mechanism underlying SNSA activation by sodium intake cannot be explained by osmotic stimuli, since osmotic stimulation by urea does not increase the BP [60]. Epithelial sodium channels (ENaC) were thought to act as sensors for sodium in the brain because these channels sense sodium on the tongue [61]. This was proven because the ENaC-specific inhibitor, benzamil, abolished the pressor responses to ICV injections of hypertonic saline in a dose-dependent manner [61]. Wang et al. [62] explored the connection of ENaC with brain ouabain. They found that ICV administration of hypertonic NaCl with a small dose of aldosterone caused pressor responses accompanied by increased SNSA. This response could be abolished by ICV pretreatment with benzamil. Further, ICV pretreatment with Digibind, an inhibitor of ouabain and digoxin, blocked the pressor effects. These findings indicate that ENaC is a sensor of sodium in the brain, and its downstream effects consist of aldosterone and ouabain/digoxin release, causing an elevation in BP.

7. Relationship between RAAS and Digitalis in the Brain (Table 1)

As mentioned above, ICV pretreatment with ARB can inhibit the pressor responses and DLI release into circulation caused by ICV injections of hypertonic saline in rats. Sodium

TABLE 1: Intracranial pretreatments with one of these agents blunt pressor responses caused by centrally administered sodium, sodium-induced hypertension in Dahl salt-sensitive rats, hypertension caused by subcutaneously injected angiotensin II, or aldosterone.

(i) AT-1 receptor blocker
(ii) Mineralocorticoid receptor blocker
(iii) Aldosterone synthase inhibitor
(iv) Epithelial sodium channel (ENaC) blocker
(v) Antidigitalis blocking antibody
(vi) Anti-oxidative agent

loading upregulates RAS in the brain. ICV pretreatment with spironolactone, an MRB, blocks both BP rises and increased OLI in the hypothalamus and pituitary caused by ICV administration of hypertonic NaCl [63], which indicates that sodium loading increases aldosterone in the brain [54]. In fact, aldosterone-like immunoreactivity in the hypothalamus is increased in rats after sodium loading. Since ICV pretreatment with spironolactone decreases the intrahypothalamic content of OLI, ouabain may be located downstream of aldosterone release. We recently found that aldosterone caused a dose-dependent release of OLI from cultured N1 cells of PVN origin [49].

8. The Central Effects of Ang II or Aldosterone of Peripheral Origin

It has long been known that pressor responses occur due to increased peripheral SNSA after Ang II is administered via vertebral artery [64]. ICV administration of Ang II leads to similar pressor responses [56]. Therefore, Ang II directly acts on the CNS. Because pressor responses can be blocked by ICV pretreatment with ARB, the effect is mediated via AT-1 receptors of Ang II [65]. For example, this response is attenuated in subfornical organ-lesioned rats, which means that one of the regions where Ang II acts is the subfornical organ [65]. SC injection of low-dose Ang II for several days leads to gradual increase in BP in approximately 3 days [10]. ICV pretreatment with an inhibitor of aldosterone synthase abolished the pressor response: it reduced the BP elevation induced by a rather high SC dose of Ang II by approximately 80%. Similar blocking effects can be obtained by eplerenone, an MRB, as well as Digibind. A similar finding has been reported by other researchers [11] who showed that RU28318, an MRB, abolishes the pressor responses caused by SC injection of Ang II. These findings are in absolute contrast to the classical interpretation that Ang II causes hypertension by constricting the arterial beds and increasing cardiac contraction. That MRBs are effective in blocking the central actions of Ang II indicates that Ang II produces aldosterone in the brain. Specifically, the RAAS may be playing a role in the brain similar to its role in the peripheral system [66].

SC injections of aldosterone in addition to 1% saline as drinking water gradually increased the BP by about

30 mmHg; this is similar to the results for Ang II administration [11]. Water intake is concomitantly increased as BP rises. Thereby, ICV administration of irbesartan, an ARB, RU28318, or spironolactone almost completely blocked the pressor responses and water intake. The pressor response can also be suppressed by ICV pretreatments with either apocynin (an NADPH oxidase inhibitor) or tempol (a reactive oxygen species scavenger); this indicates that the response is mediated by oxidative stress caused by aldosterone in the brain. However, water intake was not suppressed by suppression of oxidative stress, suggesting that the pressor mechanism is independent from the 1% saline intake response.

Because ICV pretreatment with MRB abolishes pressor responses to systemic administration of aldosterone, it follows that aldosterone of adrenocortical origin can act at a central site to increase the BP. Like the interpretation for Ang II, conclusion of this unique evidence is entirely different from the classical interpretation that aldosterone causes hypertension by acting at the renal tubule to increase sodium reabsorption. The novel findings that both Ang II and aldosterone primarily act in the CNS to cause hypertension are epoch-making discoveries that radically challenge the classical interpretation (Figure 3).

9. Supposed Common Central Mechanism of a Variety of Models of Hypertension

A hypertensive state means that the set point of the baroreceptor reflex has shifted to a higher level, and the sensitivity is decreased. Thus, when Ang II production is blocked with ACEI, the set point is lowered to the normal level and the sensitivity recovers [67]. It follows, therefore, that Ang II is critically involved in the pathogenesis of hypertension. The lowering of set point is not secondary to normalizing the BP since the set point is still at the lower BP level after treatment with ACEI, even when the BP has been elevated to the hypertensive level with intravenous infusions of phenylephrine. These findings indicate that inhibition of RAAS completely normalizes BP regulation. The previously mentioned evidence indicates that the involved RAAS is not only of peripheral origin, but also of central origin. However, since the vasomotor center exists in the CNS but not in the periphery, Ang II or AT-1 receptors in the brain will be primarily involved in changing the set point of the baroreceptor reflex mechanism.

The epidemiological and experimental evidence previously discussed suggests that a human being who lived in an environment with minimum intake of sodium during long periods of Stone Age acquired a rigid mechanism to retain sodium; this mechanism primarily consists of RAAS and SNSA. Meanwhile, a high BP would be a desirable feature for living in low-salt conditions because people with a high BP could be aroused more rapidly and fight more vigorously than those with lower BP. Atherosclerosis was not a problem, as these people did not live long. However, significant changes in the sodium environment occurred during a short period, and BP fluctuation required

TABLE 2: Sodium retention can be achieved by a variety of causes.

-
- (i) Excess intake of sodium; essential hypertension
 - (ii) Impaired renal excretion
 - (a) Renal insufficiency and renal failure
 - (b) Insulin resistance accompanied by obesity and/or the early stage of type II diabetes mellitus
 - (c) Increased aldosterone production; primary aldosteronism, idiopathic hyperaldosteronism, renovascular hypertension, pheochromocytoma
 - (d) Other mineralocorticoid excess; 17 α -hydroxylase deficiency, 11 β -hydroxylase deficiency, apparent mineralocorticoid excess syndrome and deoxycorticosterone-producing tumor
 - (e) Exaggerated renal sodium reabsorption; Liddle syndrome
-

TABLE 3: A list of secondary hypertension, supposedly caused by direct central actions of angiotensin II or mineralocorticoid.

Angiotensin II
Renovascular hypertension, aortic coarctation, renin-producing tumor, and pheochromocytoma
Mineralocorticoids
Aldosterone; primary aldosteronism, idiopathic hyperaldosteronism, renovascular hypertension, and pheochromocytoma
Other mineralocorticoids; 17 α -hydroxylase deficiency, 11 β -hydroxylase deficiency, apparent mineralocorticoid excess syndrome, and deoxycorticosterone-producing tumor

correction; thus, the renal-renin and adrenal aldosterone systems were equipped to form negative feedback. On the other hand, brain RAAS may have been forming a positive feedback system to retain sodium and maintain the BP at a high level. Specifically, when sodium was loaded, opposing mechanisms may be operational at the periphery and the CNS. Therefore, human beings having a predisposition to hypertension can easily elevate the BP with increased sodium environment, which would result in sodium-sensitive essential hypertension. Similar mechanisms might be playing a role in renal hypertension, obesity-related hypertension, and senile hypertension with reduced renal function. In renovascular hypertension, increased production of Ang II and aldosterone might be directly acting at the central site to cause hypertension. Similarly, in primary aldosteronism and pseudoaldosteronism, MR in the CNS might be mainly involved in the genesis of hypertension. Circumventricular organs surrounding the third ventricle such as SFO, OVLT, and area postrema are outside of the blood-brain barrier, are anatomically different from other brain tissues, and show dense distribution of AT-1 receptors. These areas are thought to sense information from systemic circulation. For instance, when the SFO was electrically destroyed, hypertension caused by chronic SC infusion of Ang II was suppressed [65], indicating that Ang II acts at the SFO. Further, hypotensive effects of losartan, an ARB, are reduced, which means that ARB is actually acting at the SFO to reduce BP. Therefore, Ang II and aldosterone produced in the blood may be acting at these central sites to elicit sympathetic hyperactivity (Tables 2 and 3).

Thus, the central mechanism might be involved in every type of hypertension.

10. Relationship between the Central Vasopressor Mechanism and Antihypertensive Agents (Figure 4)

As mentioned previously, the central cascade causing hypertension may consist of Na⁺, ENaC, RAAS, digitalis, oxidative stress, and SNSA. Therefore, agents acting at one of these components will be excellent antihypertensive agents. In fact, many antihypertensive agents have been identified during the long history of screening for treatment of hypertension, and the currently available antihypertensive agents seem to be acting at any one of the components of this cascade. For example, the role of diuretics is easy to understand, because they reduce sodium loading. Although ENaC inhibitors such as amiloride and triamterene are known to reduce BP [68], these are not used as first-line antihypertensive agents primarily because of their low receptor selectivity. Presently, blockers of RAAS such as ACEIs, ARBs, direct renin inhibitors (DRIs), and MRBs are the most frequently used agents. These may be acting at the core of the mechanism causing hypertension, and therefore, are the most reliable agents. Canrenone and rofloxin (PST2238) are digitalis antagonists, which also reduce BP [69, 70]. However, there remain some concerns particularly in their potency and receptor selectivity. Antioxidants may be effective antihypertensive agents, since tempol is known to reduce BP in hypertensive rats [71]. Centrally acting sympatholytic agents such as α_2 -adrenoceptor agonists or imidazoline receptor agonists are well known to reduce the BP [72], but adverse effects such as drowsiness and dry mouth limit their usage. α_1 -Adrenoceptor antagonists are also used for the treatment of hypertension [73] and are

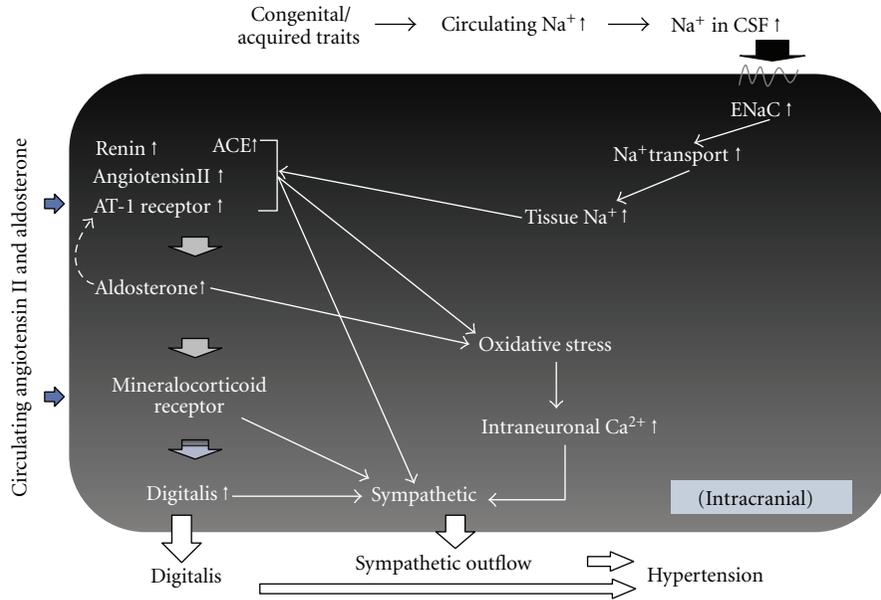


FIGURE 3: Supposed common central mechanism of a variety of models of hypertension. The dotted line indicates the possible actions of aldosterone on the RAS activation in the brain.

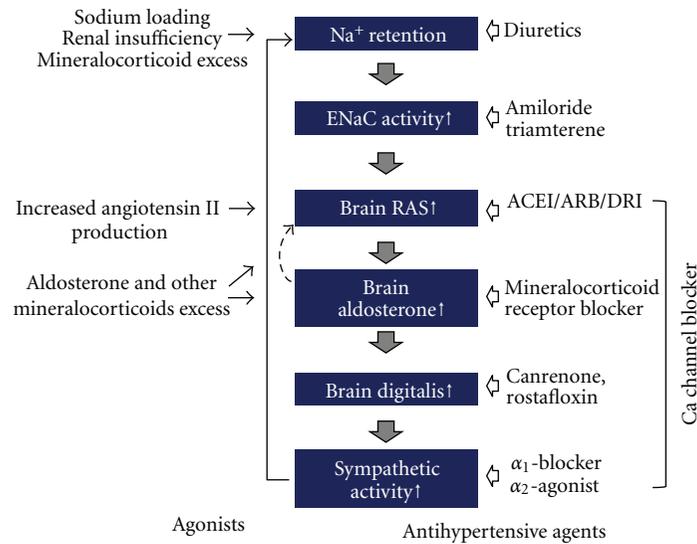


FIGURE 4: A supposed cascade of genesis of hypertension and acting sites of antihypertensive agents. In fact, when we treated hypertensive patients with these agents, reflex tachycardia is missing. This means that these antihypertensive agents are resetting the blood pressure regulatory center to a lower level besides their original actions such as vasodilation and diuresis.

known to act at the CNS level to decrease SNSA in rats [74]. CCBs are widely used in the clinical setting and are known to cause sympathetic inhibition at the CNS in rats [75]. Although short-acting and potent CCBs cause sympathetic activation by the baroreceptor reflex, gradual decreases in BP with slow-acting agents do not cause reflex tachycardia [76]. The fact that reflex tachycardia is not caused by the agents listed before indicates that at least the first and second lines of antihypertensive agents are actually acting at a CNS site to regulate BP and shifting the set point to a lower level.

In general, the blood-brain barrier blocks the entry of agents into the brain tissue. Therefore, concentrations of systemically administered agents, except the lipid-soluble agents, in the brain are very low. Therefore, it is believed that antihypertensive agents, except for α_2 -adrenoceptor agonists, do not affect the brain function. However, as mentioned earlier, these agents act at the circumventricular organs where the barrier is lacking.

Currently available antihypertensive agents have been screened for many years and can reduce the BP comfortably without stimulating the heart and finally improve our

prognosis. This may be because these agents are acting at the core mechanism of hypertension. In other words, the common site of actions of these agents is the cause of hypertension. Thus, it may be desirable to target development of novel agents with selective action at the central mechanism regulating BP in future studies.

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Review Article

Roles of Brain Angiotensin II in Cognitive Function and Dementia

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Received 14 September 2012; Revised 14 November 2012; Accepted 20 November 2012

Academic Editor: Marc de Gasparo

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The brain renin-angiotensin system (RAS) has been highlighted as having a pathological role in stroke, dementia, and neurodegenerative disease. Particularly, in dementia, epidemiological studies indicate a preventive effect of RAS blockade on cognitive impairment in Alzheimer disease (AD). Moreover, basic experiments suggest a role of brain angiotensin II in neural injury, neuroinflammation, and cognitive function and that RAS blockade attenuates cognitive impairment in rodent dementia models of AD. Therefore, RAS regulation is expected to have therapeutic potential for AD. Here, we discuss the role of angiotensin II in cognitive impairment and AD. Angiotensin II binds to the type 2 receptor (AT_2) and works mainly by binding with the type 1 receptor (AT_1). AT_2 receptor signaling plays a role in protection against multiple-organ damage. A direct AT_2 receptor agonist is now available and is expected to reduce inflammation and oxidative stress and enhance cell differentiation. We and other groups reported that AT_2 receptor activation enhances neuronal differentiation and neurite outgrowth in the brain. Here, we also review the effect of the AT_2 receptor on cognitive function. RAS modulation may be a new therapeutic option for dementia including AD in the future.

1. Introduction

The renin-angiotensin system (RAS) in the brain is well known to be involved in systemic blood pressure control, including the regulation of cerebral blood flow [1]. Angiotensin II, a major player in RAS mainly via the angiotensin type 1 (AT_1) receptor, plays an important role in the pathophysiology of tissue dysfunction [2, 3]; therefore, RAS blockade by AT_1 receptor blockers (ARBs) and angiotensin converting enzyme inhibitors (ACEIs), which are widely used as antihypertensive drugs, is expected to prevent multiple-organ damage. Cognitive impairment and dementia are common serious health problems that impair quality of life in the elderly. Previous reports indicate the possibility that treatment with antihypertensive agents prevents the impairment of quality of life including cognitive performance [4, 5]. Possible beneficial effects of RAS blockade on cognitive function are also being highlighted in the clinical field [6, 7]. An epidemiological study by Li et al. recently showed that male subjects treated with ARBs exhibited a significant reduction in the incidence

and progression of Alzheimer disease (AD) and dementia compared with those treated with ACEIs and other cardiovascular drugs [8]. Moreover, Davies et al. also reported that patients diagnosed with dementia had fewer prescriptions for ARBs and ACEIs. Interestingly, the inverse associations with AD were stronger for ARBs compared with ACEIs [9]. In contrast, Ohru et al. demonstrated that long-term use of ACEIs may have a protective role against the development of AD, probably through their direct effects on RAS in the brain [10]. In a subanalysis of the Study on Cognition and Prognosis in the Elderly (SCOPE) trial, hypertensives treated with an ARB, candesartan, showed less decline of specific areas of cognitive function such as attention and episodic memory [11]. However, almost all large clinical intervention trials have shown no significant difference in the incidence of dementia between treatment with ARBs or ACEIs and the placebo group. The Ongoing telmisartan alone and in combination with ramipril global endpoint trial (ONTARGET) and the parallel telmisartan randomized assessment study in ACE intolerant subjects with cardiovascular disease (TRANSCEND) trial showed no clear effects on cognitive

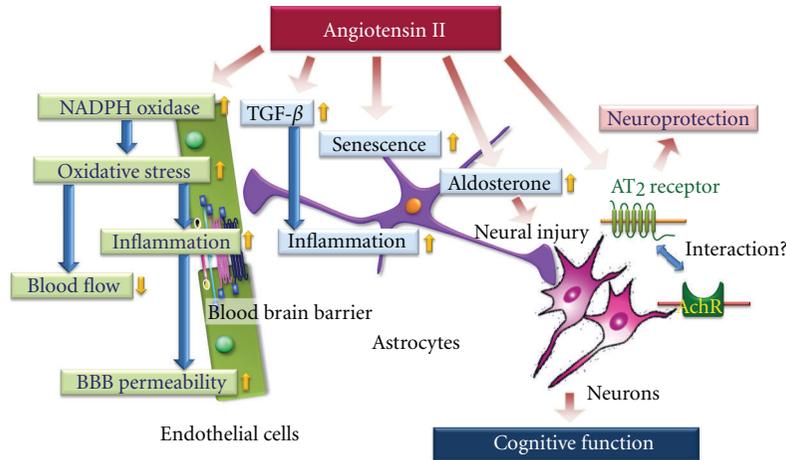


FIGURE 1: Possible effect of angiotensin II on neurovascular unit. AT₂: angiotensin II type 2 receptor, AChR: acetylcholine receptor, BBB: blood brain barrier, and TGF- β : transforming growth factor β .

outcomes [12]. The reason why RAS blockade failed to prevent dementia may be the short-term observation for the long-term preclinical disease stage of dementia; however, the detailed explanation is not clear. Another reason is the selection of hypertensive patients, who have high cardiovascular disease morbidity, in these trials. A large number of these patients are likely to go on to develop dementia, most likely with strong vascular underpinning. In these trials, vascular dementia and AD are not well distinguished because most studies focused on dementia as subanalysis. As described in the review by Kehoe and Passmore, RAS has multifunctional involvement not only in vascular dementia but also in AD [13]. Therefore, in such specific groups with cardiovascular risk, the distinction of dementia subtype is very important in comparing the incidence of dementia.

The effect of angiotensin II on cognition has been examined in basic studies. Although the blood-brain barrier is impermeable for all RAS components, the local brain RAS has possible physiological and pharmacological functions in the neuronal system [14]. Gard reviewed the contradictory role of angiotensin II in memory and learning in animal studies [15]. Angiotensin II enhances memory and learning in rodents [16, 17], but other studies suggest that angiotensin II decreases cognition [18]. To assess the paradoxical effect of angiotensin II on cognitive function, we therefore performed cognitive tests in mice with continuous activation of angiotensin II, using transgenic mice carrying both the human renin and angiotensinogen genes (hRN/hANG-Tg) [19]. Interestingly, the avoidance rate in hRN/hANG-Tg mice did not increase from 14 weeks of age; however, that from 8 to 13 weeks of age tended to be higher than that in wild-type mice. These findings suggest that the acute or subacute effect of angiotensin II may enhance cognitive function, but chronic treatment with angiotensin II may exhaust neural function and result in cognitive impairment. Angiotensin II induces cerebrovascular remodeling, promotes vascular inflammation and oxidative stress, and results in impairment of regulation of cerebral blood flow (CBF)

[20, 21]. Moreover, endothelial function in cerebral vessels was impaired in a genetic model of angiotensin-II-dependent hypertension [22, 23]. On the other hand, Lanz et al. showed that angiotensin II induced sustained central nervous system (CNS) inflammation via transforming growth factor-(TGF-) β in an experimental autoimmune encephalomyelitis (EAE) mouse model [24]. Furthermore, angiotensin II induced astrocyte senescence, which is involved in age-associated neurodegenerative disease via superoxide production [25]. In contrast, a centrally active ACE inhibitor, perindopril, was reported to prevent cognitive impairment in chronic central hypoperfusion rats [26] and Alzheimer disease model mice [27]. These reports indicate that continuous angiotensin II stimulation impairs cognitive function via stimulation of the AT₁ receptor with “environmental degradation of neurons” such as a decrease in CBF and an increase in oxidative stress, CNS inflammation, and cellular senescence in the brain. Such multiple stimuli by angiotensin II induce cognitive impairment following neuronal degeneration.

2. Effects of Angiotensin II on Amyloid β Metabolism and Cholinergic System

There are two major proposed pathomechanisms of AD; the amyloid cascade hypothesis and the cholinergic hypothesis. Amyloid β ($A\beta$) is a 39–42 amino acid peptide, produced by cleavage of amyloid precursor protein (APP) [28]. $A\beta$ (1–42) causes the neurodegenerative abnormalities that lead to clinical AD [29]. Although the effect of angiotensin converting enzyme on $A\beta$ metabolism is one of the hot topics in the relation between RAS and AD [30], it seems that angiotensin II does not directly affect $A\beta$ secretion or secretase activity via activation of the AT₁ receptor [31]. On the other hand, blockade of RAS may affect $A\beta$ metabolism. For example, an ARB, valsartan, was able to attenuate oligomerization of amyloid β peptides into high molecular weight oligomeric peptides [32]. Moreover,

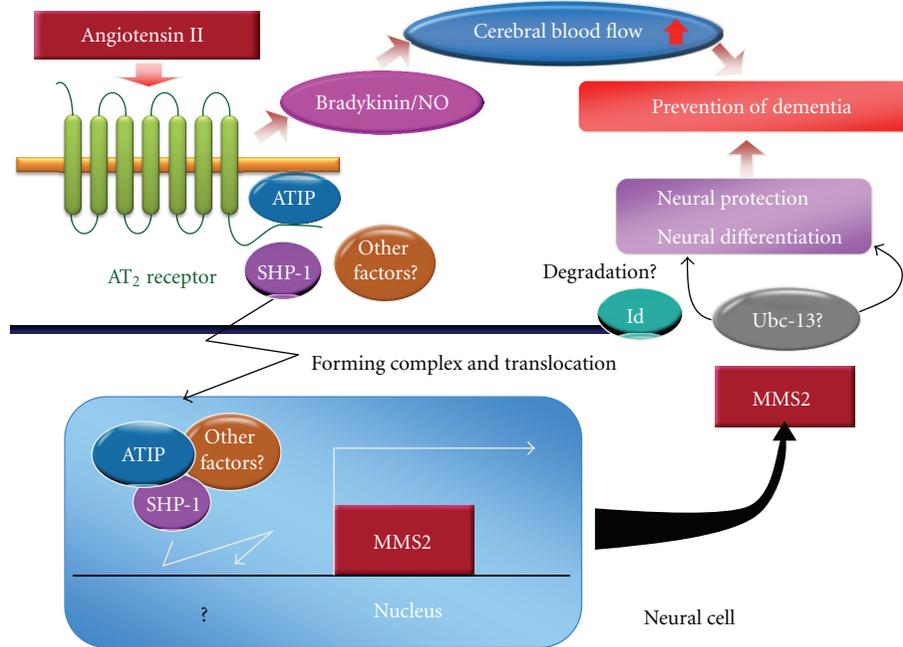


FIGURE 2: Effect of angiotensin II type 2 receptor signaling on cognitive function. AT₂: angiotensin II type 2 receptor, ATIP: AT₂ receptor-interacting protein, Id1: inhibitor of DNA binding protein 1, MMS2: methyl methanesulfonate-sensitive 2, NO: nitric oxide, SHP-1: Src homology 2 domain-containing protein-tyrosine phosphatase 1, and Ubc-13: ubiquitin conjugating enzyme 13.

treatment with valsartan also disrupted the development of amyloid β -mediated cognitive impairment in Tg2576 mice, a model of Alzheimer disease; however, it is reported that this beneficial effect is not observed with treatment with other ARBs. We previously reported that $A\beta$ (1–40) concentration in the brain of ddY mice that underwent intracerebroventricular injection of $A\beta$ (1–40) was significantly decreased by treatment with an ARB, telmisartan [33]. Moreover, Danielyan et al. reported that intranasal administration of losartan exerts direct neuroprotective effects via its $A\beta$ -reducing and anti-inflammatory effects in the central nervous system [34]. These results indicate that treatment with ARBs may have a beneficial effect on $A\beta$ -induced brain injury through unknown mechanisms on $A\beta$ metabolism by angiotensin II inhibition. On the other hand, brain-penetrating ACEIs such as perindopril prevent cognitive impairment in mice with intracerebroventricular $A\beta$ (1–40) injection via attenuation of oxidative stress and hippocampal astrocyte activation [35]. ACE activity is increased in the hippocampus of these AD mice and suppressed by perindopril treatment. Although there is concern that ACEIs may enhance brain $A\beta$ (1–42) deposition from basic research [36] because ACE converts $A\beta$ (1–42), which plays a causative role in the development of Alzheimer disease, to $A\beta$ (1–40) [37], recent pilot clinical trials showed that ramipril inhibits cerebrospinal fluid (CSF) ACE activity, but did not influence CSF $A\beta$ (1–42) and cognition [38]. The effects of other RAS components involving angiotensin-II-generating enzymes on cognition have also been discussed. $A\beta$ clearance is induced by many kinds of degrading enzyme

such as neprilysin (NEP), insulin-degrading enzyme, and endothelin-converting enzyme. Angiotensin II is also generated by degradation of angiotensinogen and angiotensin I by tonins, cathepsins, and chymases as well as ACE. Gene polymorphism in cathepsin G, one of the angiotensin generating enzymes, showed no significant association with AD [39]. In our knowledge, no report has examined the relation between tonin, chymase, and dementia; however, inhibition of angiotensin generating enzymes may also inhibit $A\beta$ degradation. Therefore, it is difficult to assess the effect on $A\beta$ metabolism of drugs that inhibit angiotensin II based on degrading angiotensinogen. Further investigation is necessary to understand the relation among angiotensin II, ACE, other degrading enzymes, and $A\beta$ metabolism.

In the cholinergic hypothesis, AD is also characterized by a loss of neurons, especially those expressing nicotinic acetylcholine receptors (nAChR) [40, 41]. To improve the cognitive deficit in AD, one promising drug target currently under investigation is the neuronal nicotinic $\alpha 7$ acetylcholine receptor ($\alpha 7$ nAChR) [42, 43]. Although there are few reports about the correlation between $\alpha 7$ nAChR and angiotensin II, Marrero's group has demonstrated that angiotensin II blocks nicotine-mediated neuroprotection against $A\beta$ (1–42) via activation of the tyrosine phosphatase, SHP-1 [44]. They also showed that angiotensin II inhibits $\alpha 7$ nAChR-induced activation of the JAK2-PI-3 K cascade in PC12 cells through AT₂ receptor-induced SHP-1 activation [45]. However, AT₂ receptor-induced SHP-1 activation also induces cerebellar development and neural differentiation [46, 47]. Moreover, $A\beta$ triggered AT₂ receptor oligomerization in

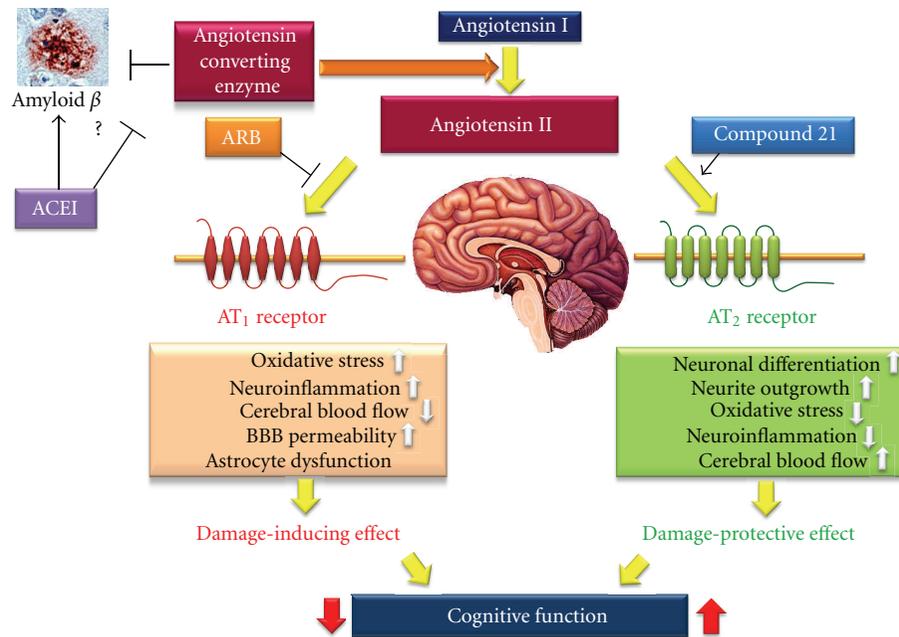


FIGURE 3: Effect of angiotensin II on cognitive function. ACE: angiotensin converting enzyme inhibitor, AT₁: angiotensin II type 1 receptor, AT₂: angiotensin II type 2 receptor, and ARB: angiotensin II type 1 receptor blocker.

the hippocampus [48] and impaired coupling of the muscarinic acetylcholine receptor (mAChR) to heterotrimeric GTP-binding proteins ($G\alpha q/11$) [49]. Therefore, the AT₂ receptor may interact with the cholinergic system; however, the actual effect of angiotensin II mediated by AChRs is still an enigma (Figure 1).

3. Effects of Angiotensin II on Neurovascular Unit

Nonneuronal cells such as vascular cells and glia (astrocytes, microglia and oligodendroglia) comprise the “neurovascular unit” and could play important roles in disease pathogenesis [50]. Especially, CBF functions in concert as a part of the neurovascular unit to maintain homeostasis of the cerebral microenvironment [51]. Iadecola and colleagues demonstrated that angiotensin II increases the production of reactive oxygen species (ROS) in cerebral microvessels via gp91phox (nox-2), a subunit of NADPH oxidase [20, 51]. Moreover, recently they also demonstrated that slow infusion of the pressor angiotensin II induces attenuation of the increase in CBF induced by neural activity (whisker stimulation) and by endothelium-dependent vasodilators, without elevation of mean arterial pressure (MAP) [52]. Such an effect of angiotensin II reduces blood supply and contributes to increased susceptibility to dementia. Interestingly, this angiotensin-induced cerebrovascular dysregulation was attenuated in female compared with male mice [53]. This sexual dimorphism of the cerebral blood-vessel response to angiotensin II may be implicated in the sex difference in cognitive impairment reported in epidemiological studies [54]. On the other hand, Takeda et al. demonstrated that the

ARB olmesartan ameliorates amyloid β -induced impairment of functional hyperemia evoked by whisker stimulation via a decrease in oxidative stress in brain microvessels [55]. Recently, Zhang et al. reported that angiotensin II increases cerebral microvasculature inflammation via induction of oxidative stress and leads to immune-endothelial interaction, resulting in enhancement of BBB permeability [56]. Therefore, angiotensin-II-induced oxidative stress may have a key role in dysfunction of the neurovascular unit (Figure 1).

On the other hand, several reports indicate the effect of angiotensin II on astrocytes to be neuroinflammation, neuronal damage and astrocyte senescence. For example, Lanz et al. clearly demonstrated that angiotensin II acts as a paracrine mediator, sustaining inflammation in the CNS via TGF- β upregulation in astrocytes [24]. We also reported that aldosterone secretion induced by angiotensin II in astrocytes enhances neuronal damage due to angiotensin II [57]. Moreover, Liu et al. showed that angiotensin II induces astrocyte senescence via superoxide production [25]. These findings of astrocyte dysfunction induced by angiotensin II also explain the crucial role of angiotensin II in dysfunction of the neurovascular unit (Figure 1).

4. Effect of AT₂ Receptors on Cognition and Dementia

The major actions of angiotensin II are mediated by the AT₁ receptor, whereas the role of a second receptor subtype known as the angiotensin II type 2 (AT₂) receptor is suggested to be protecting of the brain [58]. In the brain, AT₂ receptors are expressed not only in the vascular wall but also in areas related to learning and control of motor

activity [59, 60]. Mice with deletion of the AT₂ receptor were reported to exhibit worse cognitive function compared with wild-type mice [60]. Reinecke et al. demonstrated the possibility that stimulation of the AT₂ receptor may promote cell differentiation and regeneration in neuronal tissue [61] and that AT₂ receptor stimulation supported neuronal survival and neurite outgrowth in response to ischemia-induced neuronal injury [62]. We also demonstrated that AT₂ receptor signaling enhanced neural differentiation and the repair of damaged DNA through induction of a neural differentiating factor, methyl methanesulfonate-sensitive 2 (MMS2), which is one of the ubiquitin conjugating enzyme variants [47]. Moreover, Gallo-Payet et al. reported that angiotensin II induces neural differentiation and neurite outgrowth via mitogen-activated protein kinase [63] or nitric oxide [64] through AT₂ receptor activation, and is involved in cerebellar development [65]. Therefore, direct AT₂ receptor stimulation is expected to have a beneficial effect on cognitive function. We examined the possibility that direct stimulation of the AT₂ receptor by a newly generated direct AT₂ receptor agonist, Compound 21 (C21), would enhance cognitive function [66]. Daily intraperitoneal injection of C21 for 2 weeks significantly enhanced spatial learning evaluated by the Morris water maze test in C57BL6 mice, but this effect was not observed in AT₂ receptor-deficient mice. C21 treatment increased cerebral blood flow assessed by laser speckle flowmetry and hippocampal field-excitatory postsynaptic potential. Moreover, treatment with C21 prevented cognitive decline in an Alzheimer disease mouse model with intracerebroventricular injection of amyloid β (1–40). AT₂ receptor activation is reported to stimulate the release of NO/cGMP and may mediate vascular relaxation and blood flow indirectly through modulation of bradykinin release [67]. In our model, C21-induced cognitive enhancement was attenuated by coadministration of icatibant, a bradykinin B₂ receptor antagonist. Therefore, direct activation of the AT₂ receptor improves spatial learning via an increase in microcirculation, partly through modulation of bradykinin. The preventive effect of AT₂ receptor signaling on dementia is summarized in Figure 2. Clinical use of C21 is expected to be a new therapeutic option in patients with dementia.

5. Conclusion

Continuous stimulation with angiotensin II may damage neurons via multiple cascades through AT₁ receptor stimulation. On the other hand, stimulation of the AT₂ receptor is expected to prevent neural damage and cognitive impairment (Figure 3). However, it is difficult to perform clinical intervention studies to confirm the results of animal studies because of the long-term progression of cognitive impairment. Moreover, in clinical practice, it is not possible to exclude the antihypertensive effect of RAS blockade on cognition in patients with hypertension. However, RAS modulation may be a new therapeutic option for dementia including AD in the future. Therefore, the hypothesis that RAS regulation affects future cognitive function should be confirmed with carefully designed clinical studies.

Abbreviations

AD:	Alzheimer disease
ARB:	Angiotensin II type 1 (AT ₁) receptor blocker
AT ₁ receptor:	Angiotensin II type 1 receptor
AT ₂ receptor:	Angiotensin II type 2 receptor
ACE:	Angiotensin converting enzyme
CBF:	Cerebral surface blood flow
hRN/hANG-Tg:	Human renin and angiotensinogen genes
RAS:	Renin-angiotensin system
si:	Small interfering
MMS2:	Methyl methanesulfonate-sensitive 2.

Conflict of Interests

The authors declare no conflict of interests.

Acknowledgment

This work was supported by grants from the Ministry of Education, Science, Sports, and Culture of Japan to M. Mogi, J. Iwanami, and M. Horiuchi.

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Review Article

Discovery of Inhibitors of Insulin-Regulated Aminopeptidase as Cognitive Enhancers

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Received 9 September 2012; Accepted 19 October 2012

Academic Editor: Patrick Vanderheyden

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The hexapeptide angiotensin IV (Ang IV) is a metabolite of angiotensin II (Ang II) and plays a central role in the brain. It was reported more than two decades ago that intracerebroventricular injection of Ang IV improved memory and learning in the rat. Several hypotheses have been put forward to explain the positive effects of Ang IV and related analogues on cognition. It has been proposed that the insulin-regulated aminopeptidase (IRAP) is the main target of Ang IV. This paper discusses progress in the discovery of inhibitors of IRAP as potential enhancers of cognitive functions. Very potent inhibitors of the protease have been synthesised, but pharmacokinetic issues (including problems associated with crossing the blood-brain barrier) remain to be solved. The paper also briefly presents an overview of the status in the discovery of inhibitors of ACE and renin, and of AT1R antagonists and AT2R agonists, in order to enable other discovery processes within the RAS system to be compared. The paper focuses on the relationship between binding affinities/inhibition capacity and the structures of the ligands that interact with the target proteins.

1. Introduction

Neuropeptides participate in the transmission or modulation of signals in the central nervous system (CNS) [1]. Hence, these peptides are engaged in neurological functions that include those related to cognition and memory, mood, the experience of pain, stress, reaction to reward, control of the intake of food, and neuroendocrinological regulation. The physiological action of neuropeptides is terminated by proteolytic degradation, and this is most often mediated by extracellular proteases anchored in the cell membrane. In this respect, neuropeptides differ from classic transmitters. Limited hydrolysis of neuroactive peptides may lead to the fragments being formed with either similar or very different biological activities [2]. The conversion of angiotensin II (Ang II) to angiotensin IV (Ang IV) is a good example of the latter. This type of biotransformation results from the action of more

or less specific endoproteases. Several proteases that are capable of releasing bioactive fragments from their substrates have been identified in various CNS tissues [3, 4].

We discuss in this paper the renin-angiotensin system (RAS) and describe briefly how the two proteases, the angiotensin converting enzyme (ACE) and renin, have served and continue to serve as drug targets. We discuss briefly the two major receptors of the parent peptide angiotensin II, AT1R and AT2R, and we describe related antagonists and agonists to these receptors. Finally, we direct our focus to the hexapeptide Ang IV, which plays a central role in the brain. It has been suggested that the insulin-regulated aminopeptidase (IRAP) is the major target for Ang IV in the brain, and we therefore discuss in more detail recent progress in the discovery of inhibitors of IRAP. This paper concentrates on the molecular structures of the ligands that interact with the target proteins.

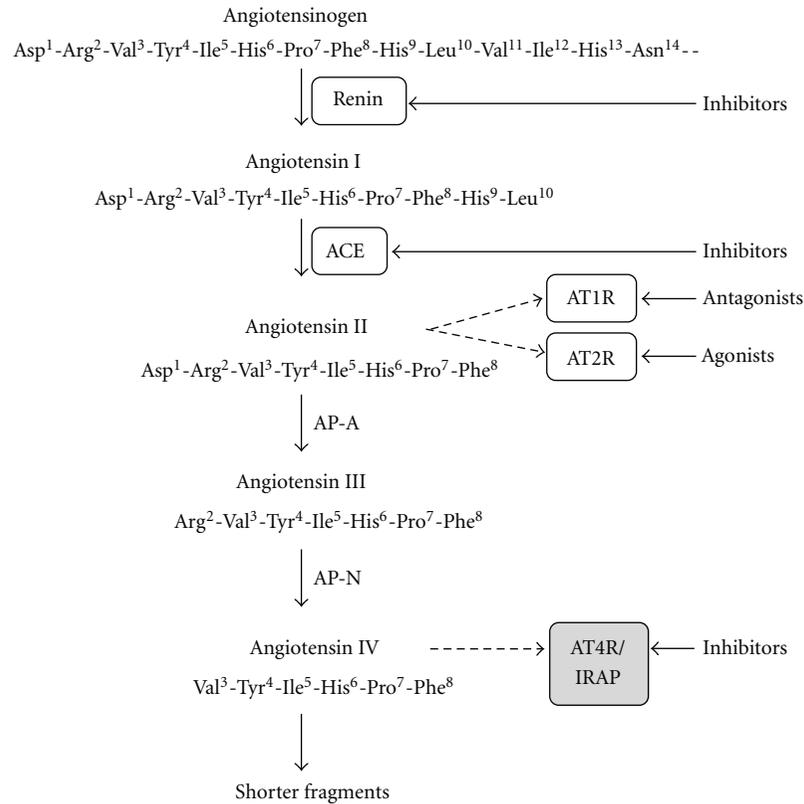


FIGURE 1: A part of the renin-angiotensin system (RAS), including selected degradation products and drug targets.

2. Proteolytic Processing

Angiotensin II (Ang II) is formed from angiotensin I (Ang I), which is an essentially inactive peptide derived from circulating and tissue angiotensinogen (Figure 1). The aspartyl protease renin liberates Ang I from angiotensinogen. The proteolytic cleavage of angiotensin I to produce Ang II is mediated mainly by the metalloproteinase ACE, an established target for drug therapy. Enzymatic cleavage by chymase, carboxypeptidase, cathepsin G or tonin are alternative routes by which Ang II can be produced [5]. As in the cases of the tachykinins and the opioid peptides, metabolism of Ang II results in the formation of several fragments with biological activities that differ from those of the parent peptides. Proteolytic cleavage by glutamyl aminopeptidase A (AP-A) and membrane alanyl aminopeptidase N (AP-N), for example, results in the sequential removal of single amino acid residues from the N-terminal end of the peptide, to form Ang III (Ang II(2–8)) and Ang IV (Ang II(3–8)), respectively [6]. These peptides are important neuropeptide fragments in the CNS [7–10]. Ang IV plays a particularly important role, and its mechanism of action is distinct [11–14]. It is noteworthy that Ang IV can be formed by the action of aminopeptidases on Ang I before it is converted to Ang II [15]. A previously unknown human Ang II-related peptide, denoted Ang A, has recently been discovered [16]. This peptide, (Ala¹)-Ang II, is not a product of proteolysis but is derived from decarboxylation of the aspartic acid residue of Ang II [16]. It acts as a full

agonist with properties that are similar to those of Ang II [17].

Chymotrypsin and dipeptidyl carboxypeptidase can further process Ang IV and the fragment Ang (3–7) to form inactive fragments and amino acid residues [18–23]. Ang (3–7) is formed from Ang IV by carboxypeptidase P (Carb-P) and propyl oligopeptidase (PO) cleavage. Chymotrypsin can hydrolyse bonds to Val, Tyr, and Ile, and this is an important property to consider when designing metabolically stable Ang IV analogues and Ang IV peptide mimetics. Furthermore, Ang II can be converted to the bioactive Ang (1–7) by the proteolytic removal of the C-terminal phenylalanine by Carb-P [12], the action of the mono-peptidase ACE2 [24], and by ACE cleavage of the Phe-His from Ang (1–9) [25].

Figure 1 shows selected degradation products and five major drug targets. There are several other potential targets for drugs in the RAS, such as the Ang (1–7)/Mas receptor [26, 27] and the aminopeptidase A (AP-A), but these will not be discussed here (for a recent paper, see [28]). Neither will antagonists to AT2R as potential future drugs be discussed. This review will focus on the discovery of IRAP inhibitors.

3. Inhibitors of Angiotensin Converting Enzyme and Renin

Angiotensin II (Asp-Arg-Val-Tyr-Ile-His-Pro-Phe) is an important modulator of cardiovascular function and exerts a pronounced hypertensive effect. Forty years ago, it was

discovered that minor modifications of the amino acid residue sequence of Ang II, such as the replacement of the phenylalanine residue at the C-terminal by a residue with an aliphatic side chain, created peptides that block the action of Ang II. Two such peptides, saralasin ((Sar¹, Ala⁸)-Ang II) and sarile ((Sar¹, Ile⁸)-Ang II), in which the N-terminal sarcosine residue enhances the effect, were evaluated in clinical trials [29, 30]. However, due to the peptidic character neither saralasin, approved by FDA for limited applications, nor the more potent sarile found long term use in clinic [31–33]. They did, however, become important research tools and clinical observations from treatment with these two peptides confirmed that the RAS is a very relevant target for drug intervention. Hence, the two major proteases, renin and ACE, that are responsible for the degradation of the precursor protein angiotensinogen to the effector peptide Ang II, became attractive drug targets (Figure 1).

ACE inhibitors were subsequently disclosed [34, 35]. The design of the first ACE inhibitor took advantage of the similarity of ACE to the metalloproteinase carboxypeptidase A, which is inhibited by 2-benzyl succinic acid. It was also known that isolated peptides from extracts of venom from the Brazilian pit viper *Bothrops jararaca* have an antihypertensive effect, inhibiting the conversion of Ang I to Ang II. These two insights allowed the elegant and very rapid discovery of the zinc-binding thiol compound captopril, which entered the market in 1978. The nanopeptide (Pyr-Trp-Pro-Arg-Pro-Gln-Ile-Pro-Pro), known as “teprotide,” had the highest *in vivo* potency among the venom peptides. It contains a proline residue at the C-terminal that is retained in captopril (Figure 2) [36]. Attempts to improve metabolic stability and to minimize the principal side effects of captopril, rashes and loss of taste, that it was hypothesized that the thiol group was responsible for, led to the once daily prodrug enalapril (Figure 2) that entered the market in 1985 [37, 38]. In enalapril, the thiol group had been replaced by a zinc-coordinating carboxyl group. The high-resolution X-ray crystal structure of enalapril (after liberation of the free carboxyl group from the prodrug) is now available [39] and shows that the N-terminal carboxyl group interacts with the catalytic zinc ion at the active site of ACE. The 3D structure of the membrane-bound ACE, however, was not known when the first ACE inhibitors were designed. Subsequently there was a large number of ACE inhibitors introduced, for example, ramipril, quinapril, perindopril, lisinopril, and benazepril, that today are extensively used in clinic [40].

Attempts to inhibit the other major protease involved in the proteolytic processing, the aspartyl protease renin, continued for decades, but it was not until 2007 that the first such inhibitor, aliskiren (Figure 3), reached the market [41–43]. The development of renin inhibitors was often hampered by the peptidic character of the potential inhibitors being studied, leading as it did to limited metabolic stability, poor absorption, and, as a consequence of this, low oral bioavailability. It was, furthermore, difficult to predict effects in humans from results obtained in animal models. The compounds were transition-state analogues, in which the peptide bond to be cleaved had been substituted by a group that mimicked the transition state. The design of the

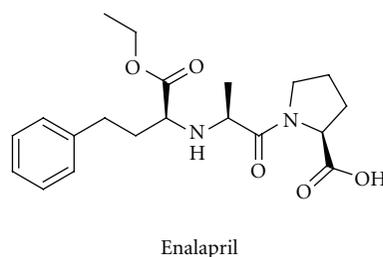
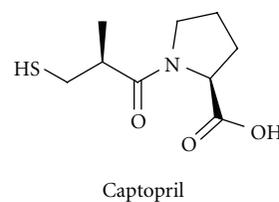


FIGURE 2: Captopril and enalapril, inhibitors of the angiotensin converting enzyme (ACE).

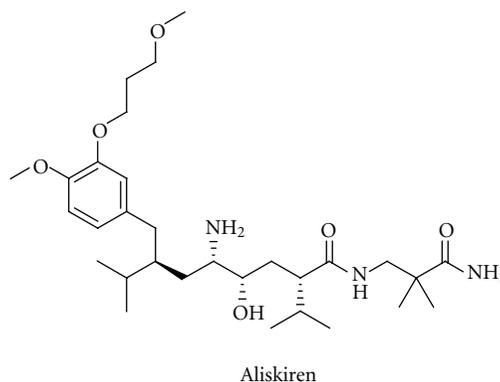


FIGURE 3: The structure of the renin inhibitor aliskiren.

transition-state analogue aliskiren was aided by modelling and the availability of X-ray structures [44]. Experience gained during the renin inhibitor programmes benefited greatly programmes to discover HIV protease inhibitors, which were, in fact, on the market ten years before aliskiren. Significant progress has recently been made in identifying new potent non-peptide “direct renin inhibitors”, but no such inhibitors have progressed to late stage development.

4. Angiotensin II AT1 Receptor Antagonists and AT2 Receptor Agonists

The first angiotensin receptor blocker (ARB) to be used in the clinic, the peptide saralasin, was not orally active and its duration of action was very short. Clinical results obtained with saralasin, however, demonstrated clearly that the Ang II receptor was a suitable target for drugs. The first non-peptide ARB, losartan (Figure 4), was introduced onto the market in 1995. Losartan is characterized by the tetrazole moiety, which serves as a carboxylate bioisostere, at the biphenyl unit and by the readily oxidized hydroxymethyl group. Losartan

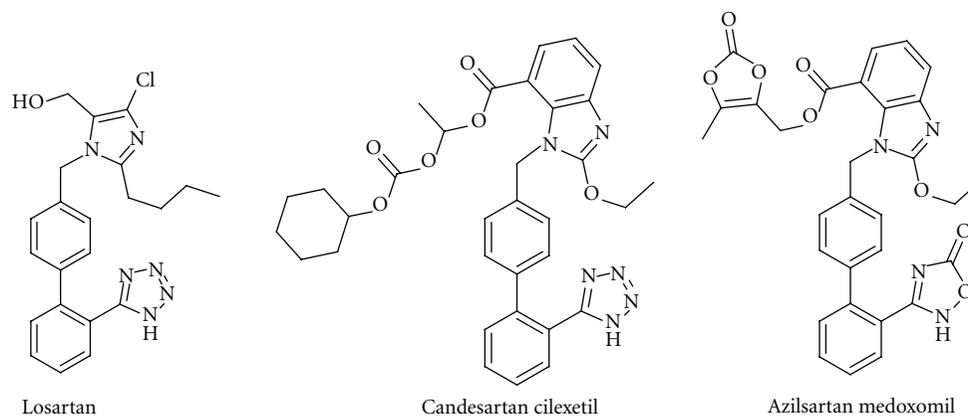


FIGURE 4: Three examples of Ang II AT1 receptor blockers (ARBs).

is converted *in vivo* by oxidation of the hydroxymethyl group to the more potent carboxylic acid metabolite. For a review on the discovery of angiotensin receptor blockers, see [45]. Several other AT1R antagonists are now in clinical use, for example, eprosartan, olmesartan, telmisartan, valsartan, and candesartan. The latter is administered as a prodrug, candesartan cilexetil (Figure 4). This liberates candesartan, which binds strongly to the AT1 receptor [46, 47]. It is believed that an optimal clinical effect of these drugs, known collectively as “sartans,” requires high levels of target occupancy [48, 49]. It should be mentioned in this context that an increasing number of examples in the literature suggest that the *in vivo* duration of drug action of AT1R blockers, for example, depends not only on macroscopic pharmacokinetic properties such as plasma half-life and the time needed to equilibrate between the plasma and the effect compartments, but also on long-lasting target binding and rebinding [50–52]. The prodrug azilsartan medoxomil (Figure 4), which has a less common carboxylic acid bioisostere attached to the biphenyl scaffold, was approved in 2011 and is the latest member of the ARB group [53–55]. It is well established that AT1R antagonists are at least as effective as ACE inhibitors, β -blockers, and calcium channel antagonists in reducing cardiovascular morbidity and mortality [56, 57]. Furthermore, it is reported that losartan [58] and candesartan [59–61] have positive effects on cognition in elderly patients. Interestingly, TRV120023 representing a β -arrestin-biased AT1R ligand has cardioprotective and functional properties *in vivo* which are distinct from losartan. It has been suggested that this novel class of drugs that is G protein independent but β -arrestin selective may provide an advantage over conventional ARBs by supporting cardiac function and reducing cellular injury during acute cardiac injury [62, 63]. Anyhow, efforts to develop new and better chemical entities that block the AT1 receptor are limited. There is strong competition in this research field, which may partly explain this decision, but efforts to develop single compounds that can block both the AT1R and the receptor of the very potent vasoconstrictor endothelin A are in progress [64]. Furthermore, dual antagonism of AT1R and neutral endopeptidase inhibition has recently attracted

interest. Clinical results from the dual inhibitor LCZ696, now in phases II–III, are promising [65].

Recently, the AT2 receptor has emerged as a new target for drug therapy [66–68]. The AT2 receptor is abundant in fetal tissues but in adults this G protein-coupled receptor remains abundant only in certain tissues such as vascular endothelium [69] and brain areas [70, 71]. The AT2 receptor is present in higher density in distinct regions of the brain and it has been suggested that it is involved in growth development and exploratory behaviour [72, 73]. It is expressed in the locus coeruleus, ventral and dorsal parts of lateral septum, the superior colliculus, and subthalamic nucleus, in the nuclei of many cells of the thalamus, and in nuclei in cells of the inferior olive. Both the AT1 and AT2 receptors are expressed in the cingulate cortex, the molecular layer of the cerebellar cortex, the superior colliculus and paraventricular nuclei [70, 74–76]. AT2R RNA and the receptor protein have recently been identified in the substantia nigra pars compacta [77] and in the hippocampus [78, 79]. Thus, the receptor is present in the adult in areas associated with control and learning of motor activity, sensory areas, and selected structures of the limbic system [80]. It has been suggested that modulation of AT2R signalling can improve cognitive performance in persons with Alzheimer’s disease (AD) not only through the action of an AT2R agonist on blood flow/brain microcirculation but also through its more specific effects on neurons [81]. Activation of the AT2 receptor affects neuronal cell differentiation and nerve regeneration [82–84]. Interestingly, several peripheral effects that are mediated through the AT2 receptor oppose other effects that are mediated through the AT1 receptor, suggesting that a similar balance may exist in the CNS [85, 86]. It is worth noting that the AT2 receptor is re-expressed in some disease conditions such as heart failure, renal failure, myocardial infarction, hypertension, and some brain disorders [71, 87–92]. The AT2R mediates vasodilatory, antiproliferative and anti-inflammatory effects [93].

Several potent drug-like and selective AT2R agonists have been disclosed [94–96]. The first of these receptor-selective agonists, M024 or “compound 21” (M024/C21, Figure 5),

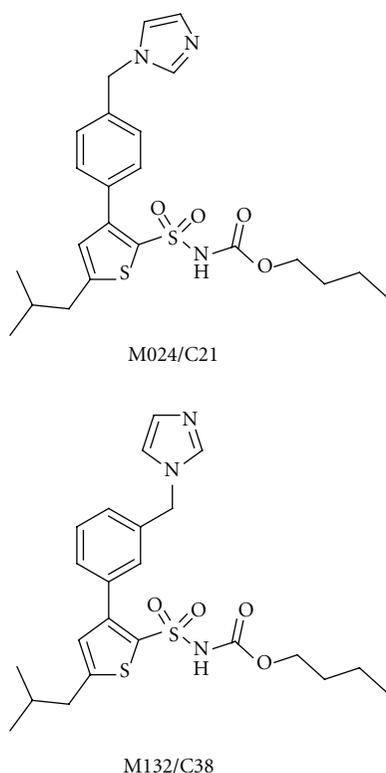


FIGURE 5: The structures of the selective AT₂ receptor agonist M024/C21 and the structurally similar M132/C38, which acts as an AT₂ receptor antagonist.

was first reported in 2004, and has been extensively studied since then [94]. It possesses a sulfonyl carbamate entity as a metabolically stable carboxylic acid isostere and exhibits a striking structural similarity to several AT₁R antagonists. Compound M024/C21, developed through a series of modifications to the non-selective AT₁R agonist L-162,313 [97], stimulates neurite outgrowth in neuronal cells (which express only AT₂R) [94] through the sustained activation of p42/p44 mapk. It decreases dopamine synthesis in the rat striatum, and some results suggest that the AT₁ and AT₂ receptors in the striatum exert opposite effects on dopamine synthesis, rather than dopamine release [98]. The compound enhances cognitive functions in mice [81, 99]. It improves myocardial function independently of blood pressure after myocardial infarction in normotensive Wistar rats [100] and has a pronounced anti-inflammatory effect [101]. M024/C21 gives improved vascular stiffness and lower collagen concentrations in the aorta and myocardium of stroke-prone spontaneously hypertensive rats [102]. It was recently suggested that the combination of M024/C21 with antihypertensive treatment might lead to vasculoprotective effects even beyond the blood-pressure-reducing effect [66, 103].

It is expected that bioavailability in the brain of the drug-like M024/C21 will be low [104]. This is expected to be the case also for M132/C38 (Figure 5), a selective AT₂R antagonist with a very similar structure as M024/C21.

Nevertheless, this AT₂R agonist/antagonist pair, in which the two compounds possess similar pharmacokinetics, should be an important and useful tool in studying the RAS. Furthermore, these molecules may serve as a starting point in medicinal chemistry programmes aimed at discovering molecules that are active as AT₂R agonists in the brain after oral administration. No selective AT₂R agonists have still entered phase I clinical trial. On the contrary, EMA401, a lipophilic structural analogue of the commonly used research tool, the AT₂ receptor antagonist PD123319 is in clinical trials. This antagonist is developed for neuropathic pain [105].

5. Inhibitors of Insulin-Regulated Aminopeptidase

In 1988, Braszko et al. reported that intracerebroventricular (i.c.v.) injection of the Ang II metabolite Ang IV (Val-Tyr-Ile-His-Pro-Phe, Figure 6) improved memory and learning in the rat [106]. They showed that Ang IV affects motor activity, the performance of passive avoidance, and a conditioned avoidance response. Various animal models were subsequently investigated, and results were obtained for Barnes maze, swim mazes, and radial arm mazes [8, 14, 107–110]. Not only Ang IV but also related analogues were studied, such as the endogenous LVV-hemorphin-7 (Leu-Val-Val-Tyr-Pro-Trp-Thr-Glu-Arg-Phe), which has structural similarities to Ang IV at the N-terminal part of the peptide, in that it has a tyrosine residue attached to lipophilic amino acid residues [111]. LVV-hemorphin-7 is a powerful promoter of memory retention and retrieval in rats [107]. The observation that Ang IV improves processes related to memory and learning has attracted considerable interest in recent years. Excellent reviews have been published describing the role of Ang IV in the brain [112–114]. Efficient new chemical entities for the treatment of the cognitive decline associated with Alzheimer's disease, brain trauma, and cerebral ischemia are needed since the clinical studies of the cholinesterase inhibitors and NMDA antagonists used today have been mostly disappointing [115–118]. Thus, new improved enhancers of cognitive functions are desired and the receptor(s) involved in the beneficial effects of Ang IV has emerged as a relevant new target for drug intervention.

A specific binding site for Ang IV was identified in 1992, and was later named the AT₄ receptor [119–121]. Harding et al. found high densities of binding sites in areas of the brain associated with cognitive, sensory and motor functions, including the hippocampus [120].

5.1. Ligands with Affinity to the AT₄ Receptor. Shortly after the discovery of the binding sites, systematic structure-activity relationship studies (SAR) were commenced by Wright and Harding and it became clear that the Val-Tyr-Ile tripeptide motif in the N-terminal part of Ang IV was of critical importance for binding affinity [122]. This conclusion relied on classic glycine and d-amino acid scans and various other alterations of the amino acid residues of

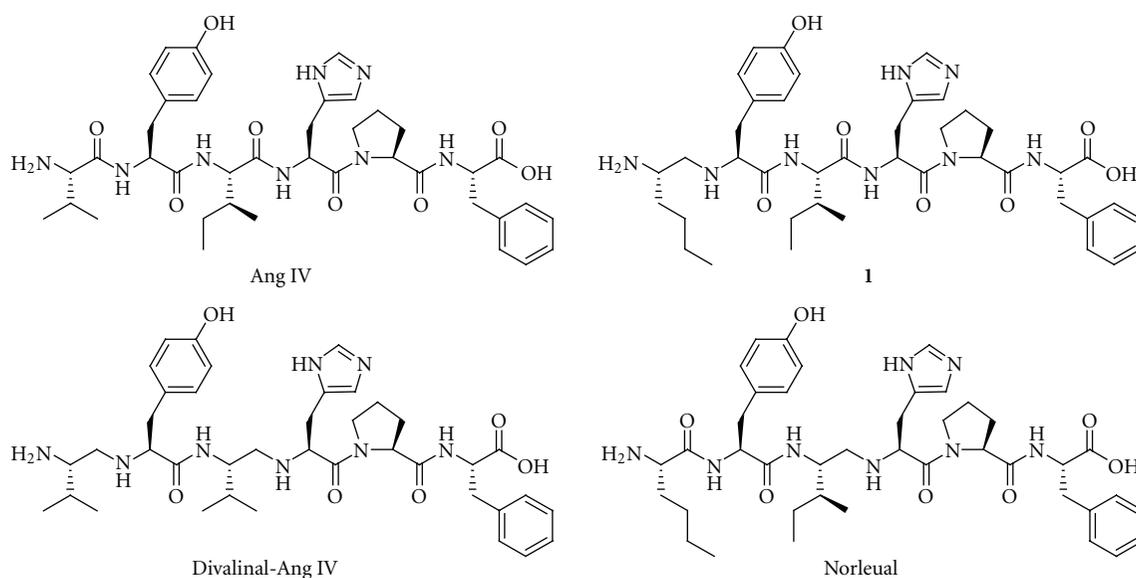


FIGURE 6: Ang IV and three hexapeptide analogues that incorporate one or two reduced peptide bonds ($\Psi[\text{CH}_2\text{NH}]$) as peptide bond bioisostere.

Ang IV [123, 124]. Hydrophobic residues at position one, and norleucine in particular, rendered very high binding affinities, while substitution of the N-terminal amine by acetylation or methylation lead to low affinity ligands. Thus, a residue with straight aliphatic side-chain combined with a primary amine function was found to be preferred in the N-terminal. The C-terminal part, on the other hand, could be altered without affecting the binding affinity dramatically. However, truncations with the exception of removal of the C-terminal phenylalanine residue were found to be unproductive [122].

The hexapeptide Ang IV and its peptide analogues are prone to undergo proteolytic cleavage but a significant improvement of the metabolic stability could be achieved by reduction of the peptide bond between Val¹ and Tyr² of Ang IV with most of the affinity retained. Analogue 1 (Figure 6) constitute another such example, encompassing a reduced peptide bond ($\Psi[\text{CH}_2\text{NH}]$) between residues one and two and thus two amine sites that can be protonated [123]. Divalinal-Ang IV and norleual (Figure 6) are two other pseudopeptides comprising reduced peptide bonds that were studied in some detail and served as important research tools [125, 126].

Kobori et al. at Taisho Pharmaceuticals filed patent applications in the late 1990s that disclosed a series of compounds that bind strongly to guinea pig hippocampus membranes [127, 128]. They had deduced this from competitive experiments with radiolabeled [¹²⁵I]Ang IV. The compounds with the highest binding affinity are characterized by a straight fourcarbon chain at position one and a reduced amide bond ($\Psi[\text{CH}_2\text{NH}]$) between residues one and two. These high-affinity compounds have also a styrene moiety that replaces His-Pro-Phe in Ang IV, (compound 2 and quinoline 3 in Figure 7) and they have IC₅₀ values lower than 1 nM. Ligand 4 in Figure 7 has three basic amino groups and was produced

by reducing the amide bond between residues two and three in compound 2. This compound has a binding affinity for hippocampus membranes that is 40 times lower than that of compound 2. A ligand essentially devoid of affinity was obtained by maintaining the peptide bond between residues one and two intact and reducing the bond ($\Psi[\text{CH}_2\text{NH}]$) between residues two and three. Furthermore, reduction of the *trans* double bond at the C-terminal leads to considerably lower affinities. The potent ligands synthesized by Kobori et al. have to the best of our knowledge not been evaluated as IRAP inhibitors.

5.2. Inhibitors of Insulin-Regulated Aminopeptidase (IRAP).

A receptor for Ang IV was purified from bovine adrenal membranes in 2001. This AT₄ receptor was identified as the insulin-regulated aminopeptidase (IRAP) [129]. The IRAP/AT₄ receptor has attracted considerable interest in recent years as a potential target for pharmaceuticals aimed for the treatment of cognitive disorders [112, 130–133]. IRAP is a single-spanning transmembrane zinc-metallopeptidase that belongs to the M1 family of aminopeptidases. IRAP has been identified as cystinyl aminopeptidase (CAP, EC 3.4.11.3), placental leucine aminopeptidase (P-LAP, soluble human homologue), oxytocinase, gp160, or vp165 [134–137]. The insulin-regulated aminopeptidase has been cloned and characterized in adipocytes in vesicles that contain the insulin-regulated glucose transporter GLUT4 [138]. IRAP contains three domains [136, 139–141]: an intracellular region that is involved in intracellular localization and redistribution, a hydrophobic transmembrane segment, and an extracellular region that contains the catalytic site. The M1 family shares a consensus His-Glu-Xaa-Xaa-His-(Xaa)₁₈-Glu zinc-binding motif that is essential for enzymatic activity, and a Gly-Xaa-Met-Glu-Asn (Xaa = Ala in IRAP)

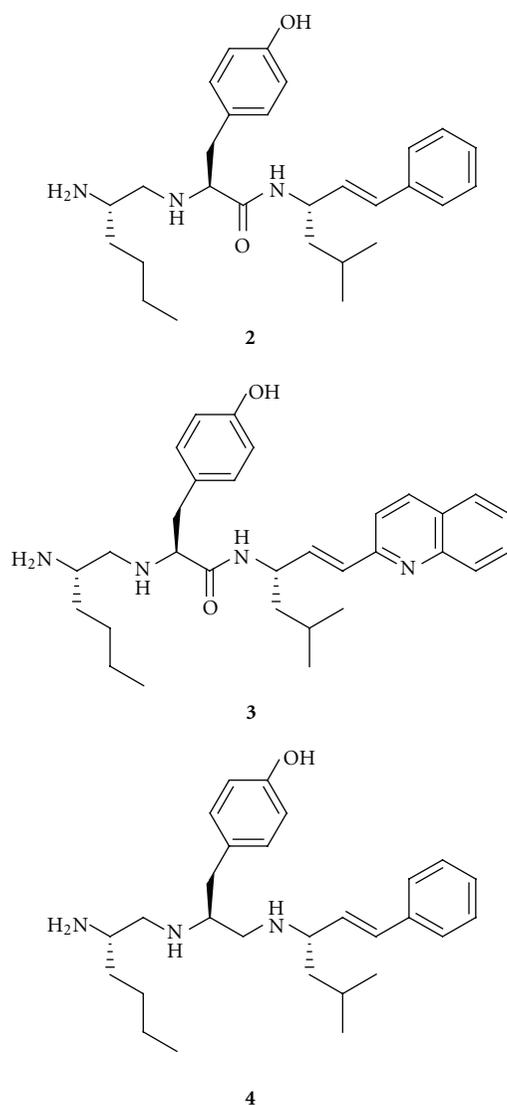


FIGURE 7: Highaffinity Ang IV receptor binding analogues that incorporate one or two reduced peptide bonds ($\Psi[\text{CH}_2\text{NH}]$) and a styrene moiety replacing the C-terminal tripeptide His-Pro-Phe of Ang IV.

exopeptidase motif. The zinc ion is coordinated to the two His residues, the second Glu residue, and a water molecule, which is anticipated to be activated by the other Glu residue during the hydrolytic step [142–144]. Mutational analyses have shown that the Gly-Ala-Met-Glu-Asn motif is important in the recognition of the N-terminal of both substrates and competitive inhibitors [139, 145, 146]. An important and characteristic property of the aminopeptidase is its ability to cleave the N-terminal amino acid residue from several bioactive peptides *in vitro*, including Met-enkephalin and Leu-enkephalin, dynorphin A, neurokinin A, cholecystokinin-8, somatostatin, oxytocin, and vasopressin [135, 147, 148].

Several hypotheses have been presented to explain the ability of Ang IV and its analogues to enhance cognitive

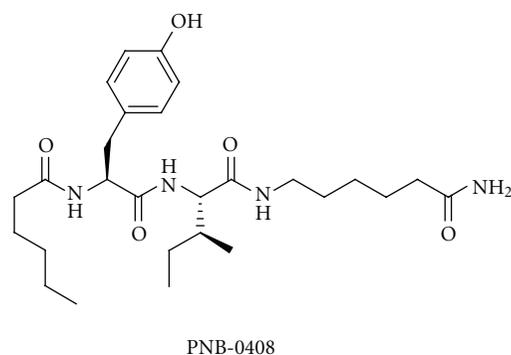


FIGURE 8: The structure of the compound PNB-0408, which crosses the blood-brain barrier.

functions [25, 149, 150]. One hypothesis concerns substrates such as vasopressin, somatostatin, and cholecystokinin, whose half-lives are prolonged when IRAP is inhibited by Ang IV, LVV-hemorphin-7 or analogous derivatives [151, 152]. These substrates improve parameters associated with cognition [153], and vasopressin and oxytocin are considered to be the main substrates of IRAP [148, 154]. It was recently shown that somatostatin has an impact on memory processing through its action on the somatostatin receptor subtype 3 [155]. A second hypothesis proposes that IRAP acts as a classical receptor that transfers information across the cell membrane after receptor binding, while a third is that the Ang IV analogues prolong the localisation of IRAP and GLUT4 at the cell surface, and thereby modulate the uptake of glucose into neurons and other cells [112, 150, 156, 157]. Furthermore, it has been proposed that the metallopeptidases in the same family as IRAP, such as aminopeptidase N (AP-N, EC 3.4.11.2), are targets [158]. Both Ang IV and LVV-hemorphin-7 inhibit AP-N activity [159]. Alternative macromolecular targets for Ang IV have been proposed, such as c-Met, a tyrosine kinase receptor that binds hepatocyte growth factor (HGF) and that is associated with memory and learning consolidation [160, 161]. Norleual (Figure 6) inhibits HGF-mediated effects at picomolar concentrations and blocks [¹²⁵I]HGF binding to c-Met.

The receptor or receptors that are involved in the positive identity identities of the effects of Ang IV and its analogues is still not clear, but both IRAP and c-Met are probably involved [114]. The availability of IRAP/Ang IV receptor ligands that are able to penetrate into the brain is important in order to obtain mechanistic insights. Wright and Harding have synthesised PNB-0408 (N-hexanoyl-Tyr-Ile-N'-(5-carbamoylpentyl)amide, Figure 8) and have shown that it crosses the blood-brain barrier and enhances cognitive activity [133, 162, 163]. This compound should be a very useful research tool. The activity profiles of this modified tripeptide and related analogues have now been established in several models of dementia [133].

Wright and Harding, and their group in USA, have been pioneers in identifying compounds that bind strongly to the

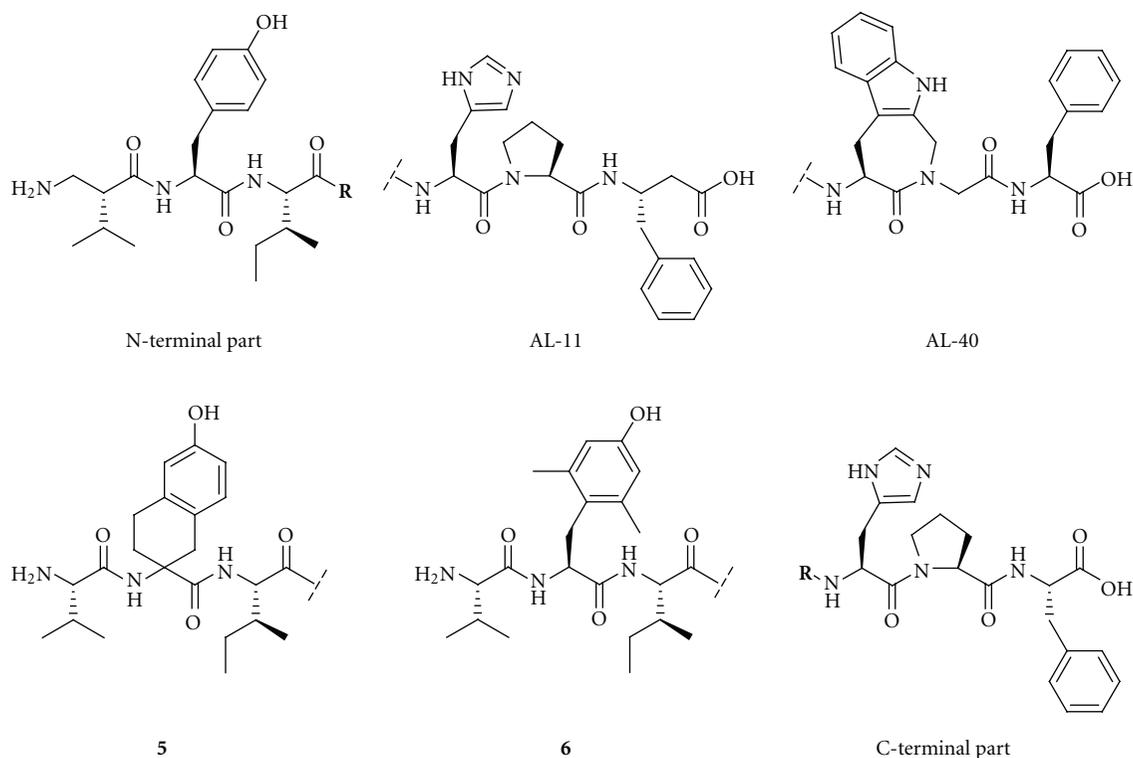


FIGURE 9: Examples of recently identified IRAP inhibitors.

Ang IV receptor, as are also Kobori et al. in Japan. Several other research groups have devoted considerable efforts in recent years to developing small molecules that interact with the Ang IV receptor, for example, in Belgium, Sweden, and (in particular) in Australia. These latter three groups have focussed on making efficient inhibitors of IRAP, and they have used slightly different and complementary approaches. The major objectives have been to identify powerful, selective inhibitors that are metabolically stable, and that resist, in particular, degradation by IRAP itself and related peptidases. To make inhibitors with high bioavailability in brain after oral administration is a tremendous challenge.

Many of the ligands previously identified as high-affinity binders were found to inhibit IRAP [111, 164]. Metal chelators (such as EDTA and phenanthroline) had previously been routinely used in experiments that measured binding affinity. It now became important to determine the ability of ligands to inhibit the hydrolysis of synthetic substrates in the absence of metal chelators. Ligands had different potencies and frequently different rank orders in the IRAP assay in the absence of chelators. It was suggested that the differences were a result of the absence of zinc in the active site when metal chelators were present [159, 164–166]. Thus, it became obvious that chelators must be omitted to obtain physiologically relevant results [167]. Furthermore, the results illustrated also the importance of synthesising inhibitors that are metabolically stable and are not substrates of IRAP.

Lukaszuk et al. in Belgium have recently performed a β -homoamino acid scan of Ang IV [168]. Replacement of

Val¹ by (R)- β^2 hVal and replacement of Phe⁶ by β^3 hPhe led to a metabolically stable and potent IRAP inhibitor, AL-11 (Figure 9). This has a high selectivity for IRAP over AP-N and the AT1 receptor. It has been reported that Ang IV affects blood pressure by a process that is mediated by the AT1 receptor [169]. The His⁴ and Pro⁵ residues were subsequently replaced by other, conformationally constrained, residues [170]. Incorporating (R)- β^2 hVal¹ and Aia⁴-Gly⁵ gave a compound that is highly selective and stable and has a high inhibitory effect (AL-40, Figure 9) [168, 170]. Ang IV itself is only a weak inhibitor of the catalytic activity of IRAP, as has been shown by experiments with the metabolically stable tritiated Ang IV analogue [³H]AL-11, in combination with the selective AP-N inhibitor 7B which is a phosphinic transition-state analogue [171] and in the absence of metal chelators. Adding metal chelators creates the apoform of IRAP [167, 172], and it is important to note that the active form and apoforms of IRAP react differently [173]. Hence, much of the previous results refer to binding to the apoenzyme rather than to the catalytically active enzyme [122–124, 174]. Ascher et al. have recently discussed the regulation of the peptidase activity of IRAP in detail [175].

The Belgian group also performed an extensive study that examined the roles of Tyr², Pro⁵, and Phe⁶ in Ang IV by introducing conformational constraints at the different amino acid residues. The study confirmed that conformational constraints are important in obtaining selectivity. Replacing Tyr² by any one of several conformationally constrained residues impairs the activity of the peptide, while modifications at the C-terminal are more acceptable.

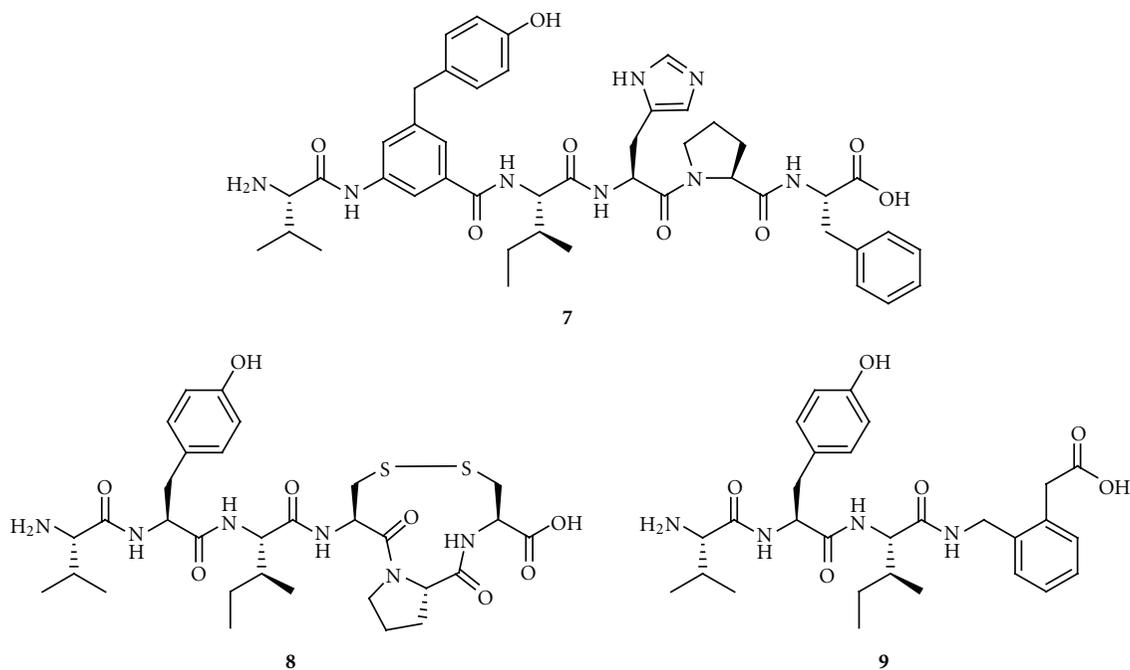


FIGURE 10: The structures of the Ang IV peptidomimetics 7–9.

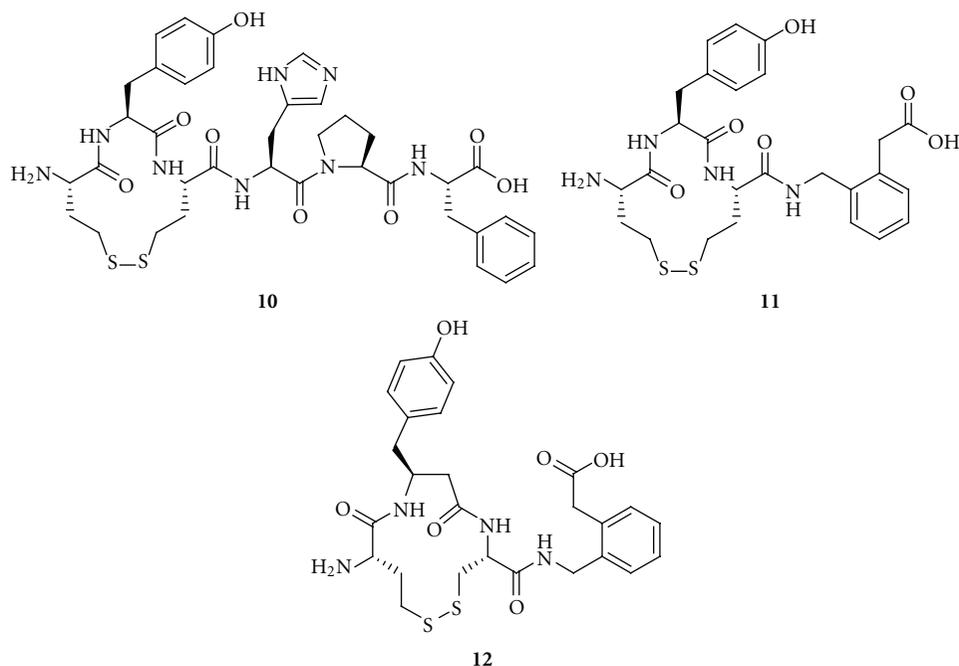


FIGURE 11: The structures of three IRAP inhibitors from a series of disulfide-cyclized Ang IV peptidomimetics. The ability to inhibit IRAP increases from 10 to 12. Both 11 and 12 have 2000-fold selectivity for IRAP over AP-N.

Analogues 5 and 6 (Figure 9), for example, inhibit IRAP to a very low degree. This suggests that the orientation of the Tyr² side chain is critical for activity [176].

Our approach in Sweden to obtain active inhibitors of IRAP has been based on an ambition to determine various bioactive conformations by introducing local steric

constraints. We have also worked to create various secondary structures by side chain cyclizations. Incorporation of a 4-hydroxydiphenylmethane scaffold as a substitute for Tyr² as part of the attempts to create steric constraint, as in Compound 7 (Figure 10), is deleterious for activity. This result again suggested that position two is susceptible

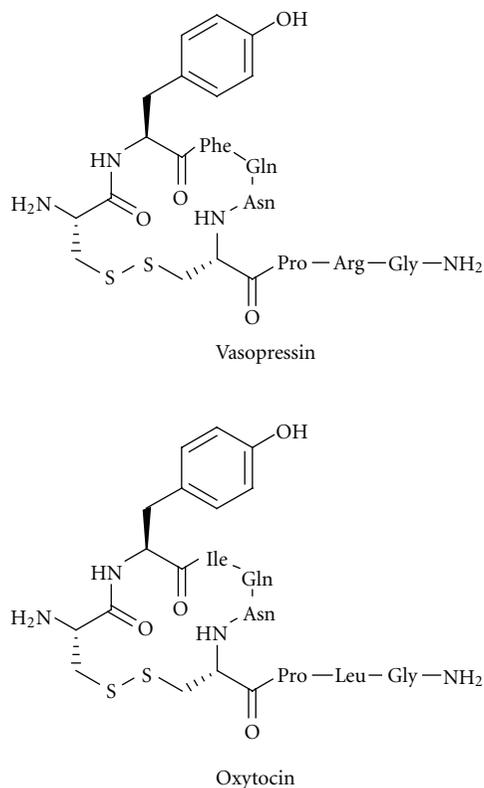


FIGURE 12: The macrocyclic disulfides vasopressin and oxytocin are substrates of IRAP.

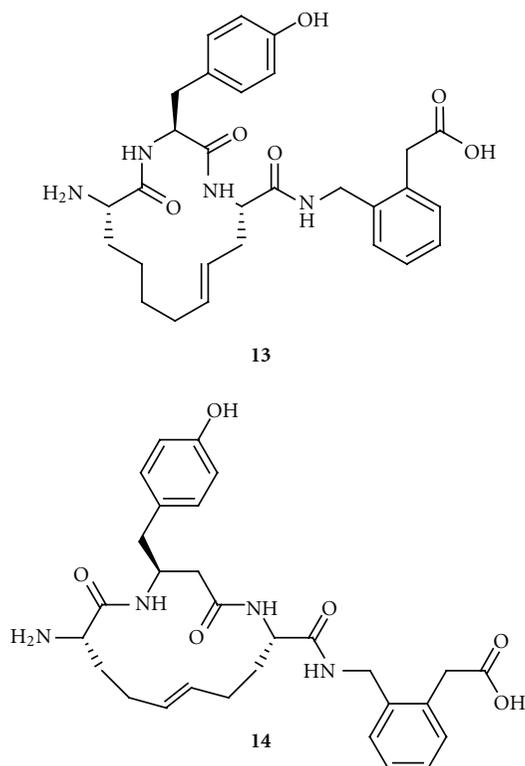


FIGURE 13: Macrocyclic Ang IV peptidomimetics that strongly inhibit IRAP.

to structural manipulation [177]. A series of macrocyclizations was performed in order to obtain improved inhibitors and to better understand how Ang IV binds to IRAP [174, 178–180]. Cyclization of [Cys⁴, Cys⁶]Ang IV to form an 11-membered ring resulted in a compound (**8**, $K_i = 26$ nM, Figure 10) that is more potent than Ang IV ($K_i = 62$ nM) as an IRAP inhibitor. Previous conformational analyses of Ang II suggested that such a macrocyclic system would tend to adopt an inverse γ -turn [181, 182], and the replacement of His-Pro-Phe by a 2-(aminomethyl)phenylacetic acid moiety designed to mimic an inverse γ -turn resulted in compound **9** (Figure 10). This compound has a K_i value of 44 nM and a much simpler structure than previous candidates [171].

Cyclization of [Cys¹, Cys³]Ang IV gives an inactive inhibitor with an 11-membered ring system, while a compound with fair inhibitory potency and a 13-membered macrocycle is obtained using [Hcy¹, Hcy³]Ang IV (**10**, $K_i = 303$ nM, Figure 11). The hybrid of **9** and **10** is a potent inhibitor (**11**, $K_i = 23$ nM, Figure 11), and, importantly, selective for IRAP over AP-N. Thus, it appears that that Ang IV adopts a γ -turn at the C-terminal when binding to IRAP, while an open, less well-defined turn conformation is present at the N-terminal. Further structural optimisation produced compound **12** (Figure 11), which has a β^3 hTyr residue in a 13-membered macrocycle. Its K_i value is 3.3 nM. This compound is 20 times more potent than Ang IV and exhibit a 2000-fold selectivity for IRAP over AP-N [180]. Removal of the carboxyl group at the C-terminal gives less efficient inhibitors of IRAP. Hence, macrocyclizations by oxidative disulfide formations can provide very potent IRAP inhibitors.

Macrocyclizations, obtained by applying the metathesis reaction, are very attractive in efforts to make potent compounds with high oral bioavailability [183, 184]. This observation and the knowledge that compounds such as vasopressin and oxytocin (Figure 12) are macrocyclic in the N-terminal part and substrates of IRAP [148] prompted the synthesis of carba analogues. It was believed that these would be efficient IRAP inhibitors that were more metabolically stable.

A large series of macrocyclic compounds were prepared, of which the compounds labelled **13** and **14** (Figure 13) have the lowest K_i values, 4.1 nM and 1.8 nM, respectively [179]. These carba analogues are also the most metabolically stable analogues that were synthesized in the programme. In the absence of chelators, the binding affinity of the 14-membered macrocyclic compounds to IRAP is 10 times higher than that of Ang IV. N-Methylation of the peptide bond between residues one and two reduces the activity, suggesting that the amide nitrogen and the N-terminal primary amine nitrogen are coordinated to the zinc atom in the active site of the protease [179]. Incorporation of a methylene group adjacent to the N-terminal amino group, as in AL-11 and AL-40 (Figure 9), and replacement of the C-terminal carboxyl group with bioisosteres seem to be the obvious next steps in efforts to improve the inhibitors. The amide bond between residues one and two, which is present in all of the macrocyclic inhibitors that have been reported, is the major target for IRAP. Thus, exchanging this bond

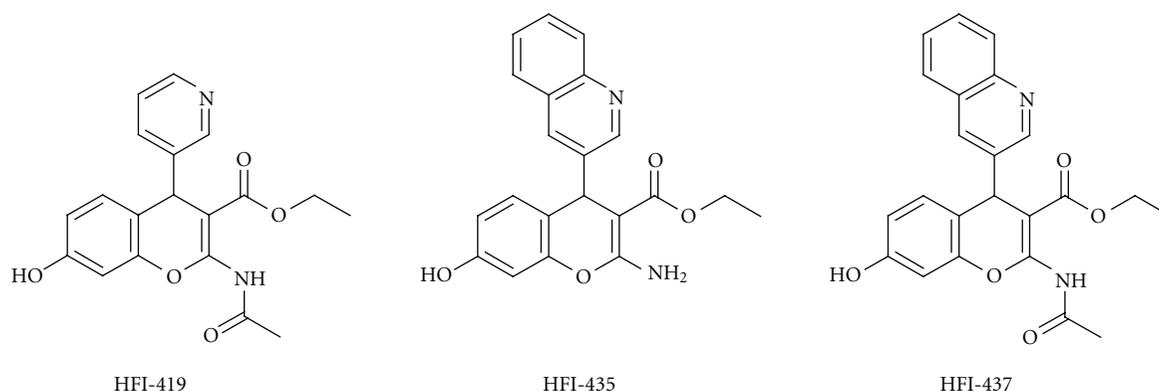


FIGURE 14: Recently identified potent non-peptidic IRAP inhibitors with a benzopyran scaffold.

for a proteolytically inert CH_2NH fragment may improve the properties. Such a CH_2NH fragment should provide also a strong coordination to zinc.

The Australian group discovered a series of IRAP inhibitors based on a benzopyran system as scaffold. These drug-like compounds should be more metabolically stable than the peptidomimetics previously discussed [113]. While compounds **13** and **14** were produced through various modifications of pre-existing compounds, primarily macrocyclizations of Ang IV itself, inhibitors based on benzopyrans originate from a structurebased design process and virtual screening.

The 3D structure of IRAP is not known. Thunnissen et al. therefore used the structurally related human leukotriene A_4 hydrolase [185], which also belongs to the M1 aminopeptidase family, in an *in silico* screening aimed at identifying potential inhibitors [186]. The process involved the *in silico* screening of 1.5 million commercially available compounds against a model structure homologous to IRAP, identification of hits, biological evaluation of hits, and optimisation of structures. Several potent drug-like benzopyrans were identified as IRAP inhibitors. Among those, the racemic pyridine derivative HFI-419, and the quinoline derivatives HFI-435 and HFI-437 (Figure 14), are selective for IRAP, with K_i values of 420, 360, and 20 nM, respectively. The compounds bind with low affinity to structurally related enzymes, such as AP-N and leukotriene A_4 hydrolase itself, despite the latter having been used in the homology modelling [186]. Computational docking suggests that the *S*-isomer is the preferred binding mode in all examples, and two alternate binding conformations for these structurally analogous inhibitors have been proposed [187].

The quinoline compounds HFI-435 and HFI-437 cannot adopt the same binding mode as the pyridinyl compound HFI-419 for steric reasons, but HFI-435 and HFI-437 seem to adopt binding modes that allow a stronger interaction with the zinc ion through coordination with the nitrogen atom in the quinoline heterocycle. It has been predicted that the quinoline compounds are more active than the pyridinyl compounds, partly due to a more favourable coordination to the zinc ion in IRAP. In addition, computational docking

experiments have suggested that Phe⁵⁴⁴ of IRAP provides an important hydrophobic packing point at one side of the active site [187]. No comparative modelling has been carried out, but it is tempting to suggest that the N-terminal of Ang IV, the macrocyclic compound **13** and the quinoline HIF-437 bind to IRAP as shown in Figure 15. The amide nitrogen (or the amide oxygen) and/or the N-terminal nitrogen atom may interact with the zinc atom.

Albiston et al. demonstrated that *i.c.v.* administration of HFI-419 enhances memory in two memory paradigms. The performance of rats in a task involving spontaneous alternation of spatial working memory after administration of compound HFI-419 [186] was similar to the performance after administration of Ang IV and LVV-hemorphin-7 [14]. Results from *in vivo* experiments with this drug-like class of compounds strongly support the strategy of using IRAP as a target for cognitive enhancers. The benzopyran class of IRAP inhibitors provides promising leads for further development.

6. Conclusions

Experimental data from various animal models demonstrate that inhibitors of IRAP facilitate memory. The explanations to the beneficial effects on a biochemical level are not clear and many alternative hypotheses have been proposed. Further studies to elucidate the mechanism of action are needed. In this context, metabolically stable IRAP inhibitors, able to cross the blood-brain barrier and with fair bioavailability in brain should serve as important research tools. Such inhibitors could also provide new chemical entities and enhancers of cognition for future treatments of the memory loss associated with ageing and AD. All data with IRAP inhibitors available today were achieved after *i.c.v.* administration and new inhibitors with oral bioavailability are highly desirable. There are in brief two different approaches addressing the discovery of such inhibitors. One starts from Ang IV itself. After subsequent truncations and macrocyclizations, stabilizing favourable conformations, potent inhibitors could be obtained. However, these are still peptidic in character. The other approach originates from a virtual screening and the utility of a homology model

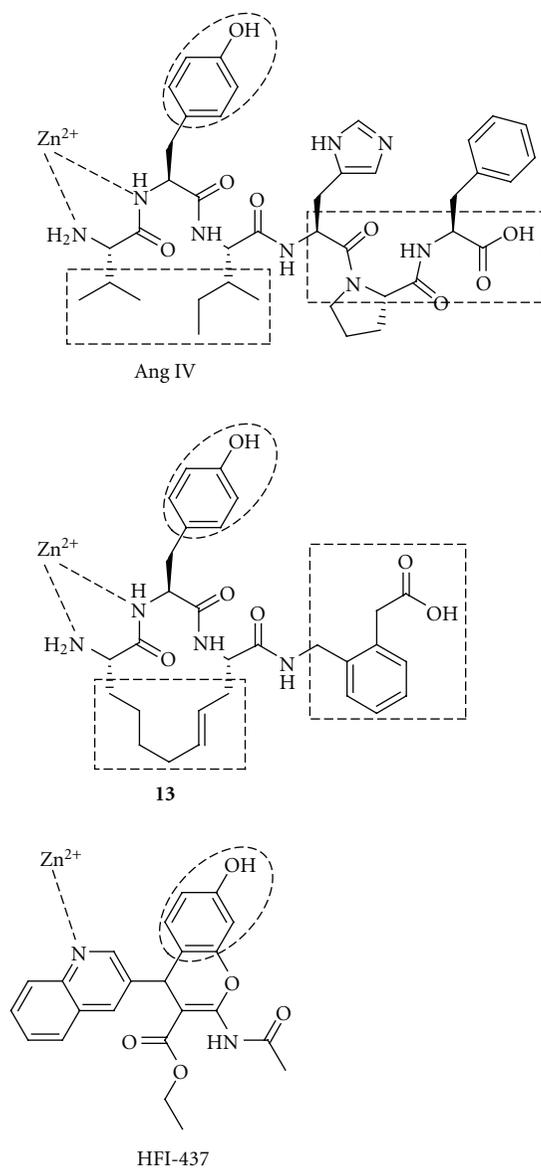


FIGURE 15: Ang IV, a peptidomimetic (**13**) and a non-peptidic (HFI-437) IRAP inhibitor. The rectangles and circles show proposed recognition elements important for interaction with IRAP.

of IRAP. A series of drug-like small inhibitors with high potential for further optimization were identified by this process. The major challenge remaining for these inhibitors is to enable them to cross the blood-brain barrier.

Acknowledgment

This study was supported by the Swedish Research Council.

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Review Article

Renin-Angiotensin System and Sympathetic Neurotransmitter Release in the Central Nervous System of Hypertension

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Received 24 August 2012; Accepted 18 October 2012

Academic Editor: Ovidiu C. Baltatu

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Many Studies suggest that changes in sympathetic nerve activity in the central nervous system might have a crucial role in blood pressure control. The present paper discusses evidence in support of the concept that the brain renin-angiotensin system (RAS) might be linked to sympathetic nerve activity in hypertension. The amount of neurotransmitter release from sympathetic nerve endings can be regulated by presynaptic receptors located on nerve terminals. It has been proposed that alterations in sympathetic nervous activity in the central nervous system of hypertension might be partially due to abnormalities in presynaptic modulation of neurotransmitter release. Recent evidence indicates that all components of the RAS have been identified in the brain. It has been proposed that the brain RAS may actively participate in the modulation of neurotransmitter release and influence the central sympathetic outflow to the periphery. This paper summarizes the results of studies to evaluate the possible relationship between the brain RAS and sympathetic neurotransmitter release in the central nervous system of hypertension.

1. Introduction

There is increasing evidence to suggest that sympathetic nervous activity in both central and peripheral nervous systems may play a major role in the regulation of blood pressure, and that hypertension is accompanied characteristically by increased sympathetic nervous activity in both humans and animal models [1, 2]. Pharmacologic studies have demonstrated that depletion of central and peripheral catecholamine stores could prevent or attenuate the development of hypertension [3, 4]. The concept on the release of sympathetic neurotransmitters from nerve endings has been considerably refined by the demonstration of multiple presynaptic receptors, which were shown to either facilitate or inhibit their release [5, 6]. The renin-angiotensin system (RAS) including angiotensin receptors is widely distributed in the brain [7–9]. It is proposed that the RAS facilitates the sympathetic nervous system and that angiotensin-converting enzyme (ACE) inhibition and angiotensin receptor blockade are antiadrenergic [10, 11]. However, the interaction between the brain RAS and sympathetic nervous system are not fully understood. In the present paper, we discuss the relationship

between the brain RAS and sympathetic neurotransmitter release and further evaluate the role of RAS in the regulation of sympathetic nerve activity in the central nervous system of hypertension.

2. Amount of Norepinephrine Release in the Central Nervous System of Hypertension

Augmented norepinephrine (NE) release and catecholamine synthesis as well as tyrosine hydroxylase gene expression have been reported at central sites related to blood pressure regulation in adult spontaneously hypertensive rats (SHR) [12, 13]. In a study presented earlier, Qualy and Westfall [14] observed that stimulation-evoked NE release from paraventricular hypothalamic nucleus in hypertensive rats was significantly increased in comparison with normotensive rats. We have been demonstrating that the stimulation-evoked NE release from hypothalamic slices was significantly greater in SHR than in normotensive Wistar-Kyoto (WKY) rats, suggesting that the release of NE from the hypothalamus may contribute to the elevated sympathetic nerve

activity in hypertension [15–22]. It was also demonstrated that, by using electrophysiological method, higher discharge rates were detected in neurons of the rostral ventrolateral medulla (RVLM) of neonatal SHR [23]. On the other hand, the stimulation-evoked NE release in the slices of whole medulla oblongata was not significantly different between SHR and WKY rats [24–29]. Recently, Teschemacher et al. [30] demonstrated that although overall electrophysiological characteristics of C1 and A2 catecholaminergic neurons in the brain were compatible between SHR and WKY rats, the angiotensin II- (Ang II-) induced Ca^{2+} -mobilization was reduced in A2 neurons of SHR. Because A2 neurons are a part of an antihypertensive circuit, the reduced sensitivity of the A2 cells to Ang II might further compromise their homeostatic role in SHR. There might be regional differences in the amount of NE release in the central nervous system of hypertensive models.

In human hypertensive subjects, Esler et al. [31] studied the brain NE release and its relation to peripheral sympathetic nervous activity by using transmitter washout method. They showed that overall NE overflow into the internal jugular vein was significantly increased in subjects with essential hypertension compared with normotensive subjects. The finding indicated that central sympathetic tone might be activated in essential hypertension, although precise mechanisms regulating central sympathetic neurotransmitter release in human hypertension is not fully understood.

3. Role of Renin-Angiotensin System in the Regulation of Norepinephrine Release in the Central Nervous System

3.1. Angiotensin II. It was shown that angiotensin I (Ang I) and Ang II injected to the central nervous system significantly elevated blood pressure [32]. Davern and Head [33] showed that the chronic subcutaneous infusion of Ang II caused rapid and marked neuronal activation in circumventricular organs, such as subfornical organ, the nucleus of the solitary tract, paraventricular nucleus, and supraoptic nucleus. In an *in vitro* study presented earlier, Garcia-Sevilla et al. [34] demonstrated that Ang II facilitated in a concentration-dependent manner the potassium-evoked NE release in the rabbit hypothalamus, which was antagonized by saralasin. The result indicated that the increase in NE release might be mediated through presynaptic angiotensin facilitatory receptors on noradrenergic nerve terminals. It was also shown that Ang II increased the potassium-evoked NE release from slices of rat parietal cortex, and that the effect was blocked by saralasin, but not by the Ca channel blocker, nimodipine [35]. Moreover, the facilitative action of Ang II on NE release might be pronounced in the hypothalamus of SHR compared with normotensive rats [36]. In a microdialysis study, Qadri et al. [37] showed that intracerebroventricular administration of 100 ng of Ang II increased blood pressure and NE release in anterior hypothalamus of conscious rats, which was antagonized by the Ang II receptor blocker. By using the similar method,

Stadler et al. [38] also reported that Ang II led to significant dose-dependent increases of NE release in the paraventricular nucleus.

Several lines of evidence demonstrate that function and signaling of the angiotensin type 1 (AT_1) receptors are quite different from the angiotensin type 2 (AT_2) sites and that these receptors may exert opposite effects on blood pressure regulation [39]. Gelband et al. [40] demonstrated that neuronal AT_1 receptors might have a pivotal role in NE neuromodulation, and that evoked NE neuromodulation might involve AT_1 receptor-mediated, losartan-dependent, rapid NE release, inhibition of potassium-channels and stimulation of Ca^{2+} -channels. Furthermore, they proposed that AT_1 receptor-mediated enhanced neuromodulation might involve the Ras-Raf-MAP kinase cascade and lead to an increase in NE transporter, tyrosine hydroxylase, and dopamine β -hydroxylase mRNA transcription. On the other hand, neuronal AT_2 receptors might signal via a G-protein and be coupled to activation of PP2A and PLA2, and stimulation of potassium-channels. Nap et al. [41] showed that prejunctional AT_1 receptors might belong to the AT_{1B} receptor subtype because AT_{1B} receptor inhibition by high concentrations of PD 123319 could suppress the Ang II-augmented noradrenergic transmission. Gironacci et al. [36] demonstrated that Ang-(1-7), which is synthesized by angiotensin converting enzyme 2 (ACE2), significantly decreased the potassium-induced NE release in the hypothalamus of SHR by stimulating the AT_2 receptors. In addition, they showed that the inhibitory effect of Ang-(1-7) on NE release was blocked by the nitric oxide (NO) synthase inhibitor and the bradykinin (BK) B_2 receptor antagonist. The finding indicated that Ang-(1-7) reduced NE release from the hypothalamus of SHR via the AT_2 receptors, acting through a BK/NO-mediated mechanism. Recent findings have also revealed that the Mas oncogene may act as a receptor for Ang-(1-7) [42–44]. It is strongly suggested that activation of the ACE2-Ang-(1-7)-Mas axis might act as a counterregulatory system against the ACE-Ang II- AT_1 receptor axis [42–44].

Bourassa et al. [45] demonstrated that AT_1 receptor binding in both RVLM and caudal ventrolateral medulla as well as dorsomedial medulla was increased in SHR compared with normotensive rats. Conversely, expression of the novel, non- AT_1 , non- AT_2 , Ang II and III binding site, which was recently discovered, might be decreased in the RVLM and dorsomedial medulla of SHR [45]. They proposed that increased AT_1 receptor binding in the RVLM might contribute to the hypertension of SHR, whereas reduced radioligand binding to the novel, non- AT_1 , non- AT_2 , angiotensin binding site in the RVLM of SHR might indicate a role for this binding site to reduce blood pressure via its interactions with Ang II and III.

In this context, the facilitative effect of Ang II on NE release might be an important factor in the excitation of sympathetic tone in the central nervous system, although further studies should be performed to assess more thoroughly the precise roles of the different types of Ang II receptors in the regulation of central sympathetic nerve activity in hypertension.

3.2. Angiotensin-Converting Enzyme Inhibitors. All components of the RAS have been identified in the brain, and the brain RAS might actively participate not only in blood pressure elevation, but also in target organ damages [7–9, 46]. It is proposed that the inhibition of brain ACE activity may be associated with blood pressure reduction induced by ACE inhibitors (ACEIs). Captopril, a widely accepted ACEI, may block the conversion of Ang I to Ang II and has been used as an effective antihypertensive agent in both human and experimental hypertension [47, 48]. Several studies have provided evidence that the distribution of the target sites for ACEIs might be widespread [49, 50]. It was demonstrated that captopril administered centrally significantly lowered blood pressure in intact conscious SHR [51]. Intracerebroventricular administration of captopril significantly suppressed the pressor responses to Ang I given by the same route in SHR [52]. Baum et al. [53] also observed the attenuation of pressor responses to intracerebroventricular Ang I by ACEI in conscious rats. It was shown that oral administration of captopril caused the inhibition of brain ACE activities [50, 54, 55]. It was demonstrated that the ACE activities in various tissues 1 hour after oral administration of several ACEIs and that captopril significantly reduced the ACE activities in aorta, heart, kidney, lung, and brain. Berecek et al. [56] showed that chronic intracerebroventricular injection of captopril attenuated the development of hypertension in young SHR in association with a depression in whole animal reactivity to vasoactive agents and an increased baroreflex sensitivity. It was also observed that central administration of captopril produced a significant depression in vascular reactivity to vasoconstrictor agents in the isolated perfused kidney of SHR in vivo [57]. The findings propose the hypothesis that central action of captopril might be, at least in part, related to vascular relaxation.

The different structures of ACEIs may influence their tissue distribution and routes of elimination. Cushman et al. [55] examined the effects of various ACEIs on brain ACE activity in SHR. They showed that not only captopril, but also zofenopril produced a modest, short-lasting inhibition of ACE activity in the SHR brain. On the other hand, fosinopril, lisinopril, and SQ 29,852 had delayed but long-lasting inhibitory actions, and ramipril and enalapril showed no effects. More studies are necessary to determine the distribution, binding activity to ACE, and metabolism of each ACEI in the brain.

In an *in vitro* study presented previously, we showed that captopril significantly inhibited the stimulation-evoked NE release in slices of rat hypothalamus and medulla oblongata [58], as well as in peripheral tissues, such as rat mesenteric arteries [59]. It might be possible that the inhibition of NE release by captopril might be partially due to a reduction in Ang II formation in the central nervous system. The modulation of central NE release by captopril might reduce the sympathetic outflow to the periphery, which could partially explain the hypotensive effects of the ACEI. Recently, Bolterman et al. [60] showed that captopril selectively lowered Ang II, oxidative stress, and endothelin in SHR. On the other hand, it was demonstrated that captopril had an

asymmetrical effect on the angiotensinase activity in frontal cortex and plasma of SHR [61]. It might be possible that multiple neuroendocrine actions of ACEIs in the brain could, at least in part, contribute to their blood pressure-lowering efficacy in both human and experimental hypertension.

3.3. Angiotensin Receptor Blockers. It was shown that Ang II administered intracerebroventricularly at a dose that induces drinking behavior in rats significantly increased potassium-stimulated release of NE in the hypothalamus [62]. It can be suggested that Ang II is important primarily in pathological states and that NE plays a substantial role in the brain Ang II-induced drinking response. Furthermore, losartan, an angiotensin receptor blocker (ARB), significantly inhibited the potassium-stimulated NE release in the hypothalamus, acting via the AT₁ receptor subtype [62]. Averill et al. [63] showed that losartan attenuated the pressor and sympathetic overactivity induced by Ang II and L-glutamate (an excitatory amino acid) in RVLM of SHR. Huang et al. [64] showed that central infusion of an AT₁ receptor blocker prevented sympathetic hyperactivity and hypertension in Dahl salt-sensitive hypertensive rats on high salt diet.

Previous studies have reported only a limited ability of systemic ARB to cross the blood brain barrier [65–69]. Gohlke et al. [65] reported that orally applied AT₁ receptor blocker candesartan suppressed the central responses of Ang II in a dose- and time-dependent manner in conscious rats, indicating that the AT₁ receptor blocker might effectively inhibit the centrally mediated action of Ang II upon peripheral application. It was also shown that peripherally administered candesartan markedly decreased AT₁ binding areas outside (subfornical organ and area postrema) and inside (paraventricular nucleus of the hypothalamus and nucleus of the solitary tract) the blood-brain barrier [66, 67]. Unger [68] proposed that candesartan might be the most effective ARB in crossing the blood-brain barrier. On the other hand, Pelisch et al. [69] demonstrated that CV-11974, the active form of candesartan, was undetectable in brain tissue after oral candesartan treatment, suggesting that candesartan may not cross the intact blood-brain barrier in its active form, or may reach the brain at undetectable levels. They proposed the possibility that undetectable levels of ARB in the brain tissue would be enough to modulate the brain RAS. Further studies are required to determine the relationship between the molecular characteristics of ARBs and their properties to cross the blood-brain barrier [69].

In human hypertensive subjects, Esler [70] proposed that the ability of ARBs to antagonize neural presynaptic angiotensin AT₁ receptors appears to differ markedly between the individual agents in this drug class. Recently, Krum et al. [71] examined whether ARBs inhibited central sympathetic outflow in human subjects. Using the whole body NE spillover method in humans with essential hypertension, they demonstrated that eprosartan and losartan did not materially inhibit central sympathetic outflow or act presynaptically to reduce NE release at existing rates of nerve firing. They concluded that sympathetic nervous inhibition might not be a major component of the blood

pressure-lowering action of ARBs in subjects with essential hypertension.

On the other hand, de Champlain et al. [72] showed that the hypotensive action of valsartan may be mediated in part by an inhibition of the sympathetic baroreflex in patients with essential hypertension. It was also demonstrated that valsartan not only decreased blood pressure, but also shifted the baroreflex set point in hypertensive subjects [73]. Additional studies are necessary to determine the potential effects of ARBs on sympathetic nerve activity in the central nervous system of both human and experimental hypertension.

3.4. Direct Renin Inhibitors. Recent years have seen the development of the nonpeptide, orally long-term effective direct renin inhibitor (DRI), aliskiren, which may block the initial stages of the RAS and exert a sustained antihypertensive action in hypertensive subjects [74–76].

With regard to the influences of aliskiren on central sympathetic nervous system, Huang et al. [64] examined whether central infusion of aliskiren prevented sympathetic hyperactivity and hypertension in Dahl salt-sensitive hypertensive rats on high salt diet. Intracerebroventricular infusion of aliskiren markedly inhibited the increase in Ang II levels in the cerebrospinal fluid and in blood pressure caused by intracerebroventricular infusion of rat renin. In Dahl-salt sensitive rats, high salt intake increased resting blood pressure, enhanced pressor and sympathoexcitatory responses to air stress, and desensitized arterial baroreflex function. All of these effects were significantly prevented by intracerebroventricular infusion of aliskiren. These results indicated that intracerebroventricular infusions of aliskiren were effective in preventing salt-induced sympathetic hyperactivity and hypertension in Dahl-salt sensitive rats. Because the ARB also exerted similar effects [64], the result strongly confirms the idea that renin in the brain plays an important role in the salt-induced hypertension.

In a clinical study, it was shown that aliskiren significantly reduced sympathetic hyperactivity and blood pressure in patients with chronic kidney disease (CKD) [77]. However, Fogari et al. [78] reported that the increase in plasma NE evoked by Ca channel blocker, amlodipine, was not reduced by aliskiren addition in hypertensive subjects. It is strongly suggested that DRI might directly inhibit sympathetic neurotransmitter release, and that the hypotensive action of DRI might largely depend on its inhibitory effect on the sympathetic nerve activity in the central nervous system.

3.5. (Pro)renin Receptor. Recent evidence indicates that the (pro)renin receptor (PRR), which is a newly discovered member of the brain RAS, might contribute to the pathogenesis of hypertension [79–82]. Li et al. [83] demonstrated that PRR protein was highly expressed in neurons and upregulated in the subfornical organ and the periventricular nucleus in Ang II-dependent hypertensive mice. In addition, they found that PRR knockdown in the brain significantly decreased blood pressure in renin-angiotensinogen transgenic hypertensive mice, which was

associated with a decrease in sympathetic tone and improvement of spontaneous baroreflex sensitivity. Furthermore, PRR knockdown was associated with downregulation of the AT₁ receptors. It is proposed that PRR blockade in the central nervous system might represent a novel approach for the treatment of neurogenic hypertension.

3.6. Mineralcorticoid Receptor. It has been shown that aldosterone as well as Ang II might act within the central nervous system and cause the sympathetic hyperactivity to increase blood pressure [84–87]. Huang et al. [84] proposed that brain aldosterone-mineralcorticoid receptor (MR)-ouabain pathway might have a pivotal role in Ang II-induced neuronal activation and pressor responses. The sympathoexcitatory effect of aldosterone was blocked by intracerebroventricular of MR antagonist [86]. It was also shown that central infusion of aldosterone synthase inhibitor prevented sympathetic hyperactivity and hypertension by central sodium loading [88]. In a clinical study, Raheja et al. [89] showed that MR blockade by spironolactone prevented chlorthalidone-induced sympathetic activation in hypertensive subjects. Blockade of MR receptors and inhibition of aldosterone synthesis in the central nervous system might lead to a reduction in systemic blood pressure, although it remains unclear whether MR receptor blockers might exert greater antihypertensive effects than other RAS inhibitors.

3.7. Oxidative Stress. The involvement of oxidative stress is implicated in the pathogenesis of hypertension. It was demonstrated that superoxide anions in the RVLM, which might generate hydroxyl radicals, were increased in SHR-SP (stroke prone), suggesting that increased oxidative stress would lead to enhancing the central sympathetic outflow [90, 91]. Recent evidence indicates the possible link between Ang II and NADPH oxidase-derived oxidative stress in the central nervous system [92, 93]. Mertens et al. [93] proposed that AT₁ receptors might activate the NADPH oxidase complex which could be the most important source of reactive oxygen species (ROS) and suggested that the produced superoxide anion would be converted into H₂O₂ by superoxide dismutase or combine with NO to generate peroxynitrite, thereby decreasing NO-bioavailability and promoting lipid and protein oxidation. Additional studies are necessary in order to further unravel the implications of oxidative stress in the Ang II-induced neurotoxicity in the brain.

4. Renin Angiotensin System and Other Neurotransmitter Release in the Central Nervous System

4.1. Dopamine Release. There is increasing evidence to suggest that dopaminergic nerve activity in the central nervous system may play a crucial role in the regulation of blood pressure [94–102]. Recent evidence has suggested a functional interaction between brain angiotensin mechanisms and dopamine neurons, although the influences of the

RAS on central dopaminergic activities in hypertension are controversial [94–102].

It has been demonstrated that ACE is localized throughout the brain and, importantly, is found in neurons of striatum and nigrostriatal tract [48]. In addition, there has been increasing evidence in favor of the involvement of nigrostriatal dopaminergic systems in the pathogenesis of hypertension. Linthorst et al. [95] reported that substantia nigra lesions caused a profound attenuation of the development of hypertension in SHR. Brown et al. [103] showed that the Ang II-induced dopamine release was completely blocked by the AT₁ receptor antagonist, losartan, but not by the AT₂ receptor antagonist, PD 123177, suggesting that Ang II acting via the AT₁ receptor subtype might facilitate the release of dopamine, in the rat striatum. By using the microdialysis method, Mendelsohn et al. [104] showed that administration of Ang II into the rat striatum caused an increase in the release of dihydroxyphenylacetic acid (DOPAC), a dopamine metabolite, and proposed that dopamine release was under a tonic facilitative influence of Ang II.

On the other hand, it has been proposed that the nigrostriatal dopaminergic system may mediate baroreflex sensitivity in rats because both striatal dopamine release and baroreflex responses produced by phenylephrine and carotid occlusion were attenuated by lesions of the nigrostriatal dopaminergic pathway [105]. It has also been reported that centrally administered Ang II increased exploratory behavior in rats and that this effect was antagonized by the dopamine receptor antagonist sulpiride. The finding indicated that Ang II potentiated central dopaminergic effects and dopamine-mediated behavioral responses [106]. Banks et al. [107] demonstrated that the ACE inhibitor captopril inhibited apomorphine-induced oral stereotypy in rats. The observation suggests that a decrease in Ang II formation caused by the ACE inhibitor could block nigrostriatal output. In an *in vitro* study, we examined the effects of captopril on the release of dopamine, and further determined a possible role of RAS in the regulation of dopaminergic neurotransmission in the central nervous system [108]. We showed that captopril inhibited the release of dopamine in the rat striatum in a dose-dependent manner [108]. Although the mechanisms underlying the neurosuppressive effects of captopril remain to be elucidated, the finding suggests that the inhibition of dopaminergic neurotransmission may be related to the central action of the ACEI.

Jenkins et al. [109] observed that AT₁ receptors have been identified on dopamine containing cells in the substantia nigra and striatum of human brain using receptor autoradiography. Mertens et al. [110] showed that AT₁ and AT₂ receptors might exert an opposite effect on the modulation of DA synthesis in the striatum. Speth and Karamyan [111] demonstrated that the novel, non-AT₁, non-AT₂ binding site for angiotensin peptides as a mediator of nontraditional actions of Ang II, could have a role in the stimulation of dopamine release from the striatum.

Several studies have been made to elucidate the possible link between the brain RAS and the dopaminergic cell death in Parkinson's disease [93, 112, 113]. One hypothesis is that

AT₁ receptors might play a pivotal role in Parkinson's disease. As mentioned above, the stimulation of AT₁ receptors might lead to the activation of the NADPH oxidase complex and the generation of ROS [92, 93]. Mertens et al. [93] proposed that AT₂ receptor agonists alone or in combination with AT₁ receptor blockers might effectively improve the pathological conditions in Parkinson's disease. It was also shown that chronic treatment of ACEI increased striatal dopamine content in the MPTP-mouse [112]. On the other hand, it was demonstrated that administration of PRR blocker significantly decreased 6-hydroxydopamine-induced dopamine cell death in the cultures, suggesting that the potential neuroprotective strategies for dopamine neurons in Parkinson's disease should address not only Ang II but also PRR signaling [113].

In this context, it can be speculated that the striatal dopaminergic system may actively participate in the control of blood pressure and behavioral responses, although further studies should be performed to assess the precise role of the RAS in the regulation of central dopaminergic nerve activity and its modulation by the RAS inhibitors.

4.2. Acetylcholine Release. Previous studies have shown that the central cholinergic system may also actively participate in blood pressure control [114–123]. Buccafusco and Spector [114] demonstrated that cholinergic stimulation of the central nervous system by direct receptor agonists produced pressor responses involving activation of central muscarinic receptor sites. It was shown that pressor responses induced by intracerebroventricular injection of Ang II were significantly blocked by hemicholinium-3 (an inhibitor of Ach synthesis), which may indicate a possible interaction between the cholinergic nervous system and the RAS in the brain [115]. Vargas and Brezenoff [116] also reported that the decrease in Ach content of the hypothalamus, striatum, and brainstem caused by hemicholinium-3 was associated with a reduction in systemic blood pressure in SHR. In an *in vitro* study, we have shown that captopril significantly inhibited the stimulation-evoked Ach release in rat striatum [108]. The finding may propose the hypothesis that the inhibition of central cholinergic activity could contribute, at least in part, to the hypotensive mechanisms of captopril.

Recently, the possible link between the brain RAS and the pathophysiology of Alzheimer's disease has been documented. Tota et al. [124, 125] observed that perindopril and candesartan significantly ameliorated the scopolamine-induced impairment in memory, cerebral blood flow, and cholinergic function in mice. In addition, they suggested that activation of AT₁ receptors might be involved in the scopolamine-induced amnesia and that AT₂ receptors could contribute to the beneficial effects of the RAS inhibitors. The findings might propose the idea that inhibition of the brain RAS in hypertensive subjects would be neuroprotective against Alzheimer's disease, although further studies are necessary to assess more thoroughly the relationships between the RAS and central cholinergic nerve activity and their roles in the regulation of blood pressure and neurological functions.

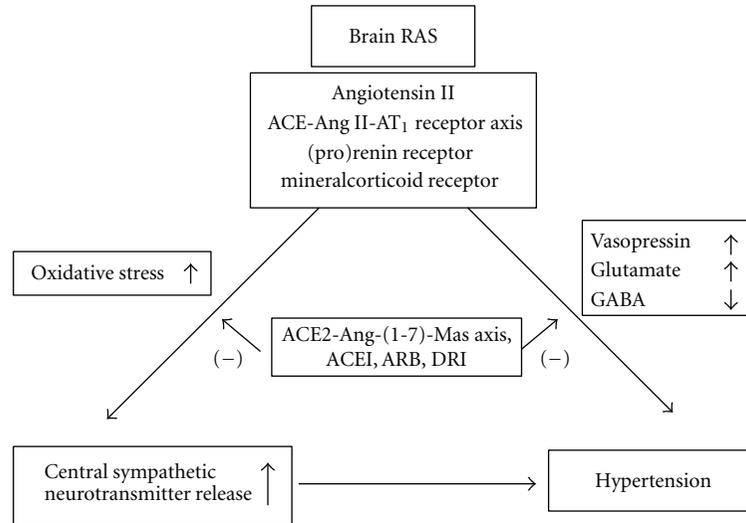


FIGURE 1: Schematic demonstration of the possible relationship between the brain RAS and sympathetic neurotransmitter release in hypertension. RAS: renin-angiotensin system, ACE: angiotensin converting enzyme, ACEI: angiotensin converting enzyme inhibitor, ARB: angiotensin receptor blocker, DRI: direct renin inhibitor, GABA: γ -aminobutyric acid, (-): inhibition.

4.3. Vasopressin Release. Brain RAS also modulates the cardiovascular and fluid-electrolyte homeostasis by interacting hypothalamic-pituitary axis and vasopressin release [126, 127]. It was shown that the angiotensinogen-deficient rats had lower plasma levels of vasopressin and an altered central vasopressinergic system [128, 129]. It was also demonstrated that PRR receptor knockdown significantly reduced AT₁ receptors and vasopressin levels in the renin-angiotensinogen double-transgenic hypertensive mice [83], suggesting that the brain RAS might have a pivotal role in the regulatory mechanisms of vasopressin neurotransmission.

4.4. Glutamate/GABA (γ -Aminobutyric Acid) Release. The role of the brain RAS on glutamate/GABA neurotransmission has also been described [130–132]. It was shown that bilateral microdialysis of the AT₁ receptor blocker, ZD7155, into the RVLM significantly decreased glutamate and increased GABA levels [130]. In contrast, administration of AT₂ receptor blocker, PD 123319, increased glutamate and decreased GABA levels within the RVLM [131]. Fujita et al. [132] demonstrated that administration of candesartan suppressed ischemia-induced increases in the extracellular glutamate with a concomitant reduction in the production of ROS in the retinal ischemia-reperfusion injury model of the rat, indicating that candesartan might protect neurons by decreasing extracellular glutamate after reperfusion and by attenuating oxidative stress via a modulation of the AT₁ receptor signaling. These findings suggested that the RAS might have a crucial role in the regulation of cardiovascular and neurological functions by modulating glutamate/GABA release in the brain.

5. Conclusion

All components of the RAS have been identified in the central nervous system, and the brain RAS may regulate blood

pressure by modulating sympathetic nerve activity. It has been proposed that the RAS may have a stimulatory influence on the sympathetic nervous system. The brain RAS may augment presynaptic facilitation of sympathetic neurotransmitter release and enhance the central sympathetic outflow. In the present paper we discussed the relationship between the brain RAS and sympathetic neurotransmitter release in hypertension (Figure 1). Ang II strongly potentiates sympathetic neurotransmitter release in the central nervous system. In contrast, the inhibitors of the RAS, such as ACEIs, ARBs, and DRIs might suppress sympathetic hyperactivity in the brain. The release of vasopressin, glutamate, and GABA could also be altered by the RAS inhibition. Although the clinical significance of the modulation of central sympathetic neurotransmitter release by the RAS inhibitors is not fully understood, the current findings may be consistent with the idea that the neurosuppressive effect could partially contribute to their hypotensive action in hypertension.

Clearly, more studies are required to further evaluate the precise role of the brain RAS in the control of sympathetic nerve activity, blood pressure, and neurological functions. In addition, better knowledge of the cellular mechanisms in the brain RAS could provide useful information concerning the development of a more specific and more physiological approach to hypertensive research.

Acknowledgments

This study was supported in part by Grants-in-Aid for scientific research from the Ministry of Education, Science, Sports, Culture and Technology of Japan (nos. 15590604, 18590658, 20590710, and 23590901), the Uehara Memorial Foundation (2005), the Takeda Science Foundation (2006), the Salt Science Foundation (2007), and the Mitsui Foundation (2008).

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Review Article

Cardiac-Autonomic Imbalance and Baroreflex Dysfunction in the Renovascular Angiotensin-Dependent Hypertensive Mouse

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Received 3 September 2012; Accepted 10 October 2012

Academic Editor: Ovidiu C. Baltatu

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Mouse models provide powerful tools for studying the mechanisms underlying the dysfunction of the autonomic reflex control of cardiovascular function and those involved in cardiovascular diseases. The established murine model of two-kidney, one-clip (2K1C) angiotensin II-dependent hypertension represents a useful tool for studying the neural control of cardiovascular function. In this paper, we discuss the main contributions from our laboratory and others regarding cardiac-autonomic imbalance and baroreflex dysfunction. We show recent data from the angiotensin-dependent hypertensive mouse demonstrating DNA damage and oxidative stress using the comet assay and flow cytometry, respectively. Finally, we highlight the relationships between angiotensin and peripheral and central nervous system areas of cardiovascular control and oxidative stress in the 2K1C hypertensive mouse.

1. Introduction

The sympathetic nervous system has an excitatory action on the heart and blood vessels, whereas the parasympathetic cardiovascular innervation has an inhibitory action on the heart [1]. Cardiac output and vascular resistance are the main determinants of arterial blood pressure (BP), which is maintained with minimal oscillations by baroreceptors located at the carotid sinus and aortic arch that transmit their signals to integrative medullary areas [1, 2]. Thus, the balanced activity of the efferent autonomic nervous system and arterial baroreceptors is essential for the control of the cardiovascular system to achieve optimal blood flow to the organs of the body.

As recently reviewed [3], conditions of exaggerated and sustained sympathetic activity, reduced parasympathetic activity, and baroreflex dysfunction are important cardiovascular risks. Over the past decades, our laboratory has shown that these pathological conditions are present as a result of

the hypertension induced by the activation of the renin-angiotensin system (RAS) in the rat [4, 5], which are also observed in the RAS-dependent hypertensive mouse [6, 7].

In this paper, we will highlight the characteristics of the murine model of RAS-dependent hypertension, provide new insights into the role played by oxidative oxygen species (ROS) in the integrative brain areas, and discuss which findings are expected to be revealed next.

2. Induction of 2K1C Hypertension in the Mouse

For decades, the rat has been used to study the relationship between RAS and the autonomic nervous system. However, genetic discoveries and advances in molecular biotechnologies have provided the opportunity to develop many mouse models for human diseases. Although a major disadvantage of this animal is the small size, advances in surgical techniques have overcome this limitation, allowing

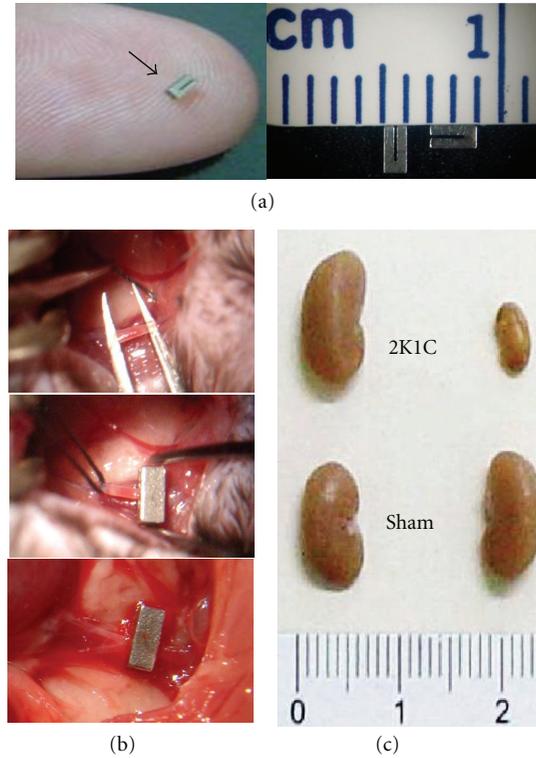


FIGURE 1: The procedure used to induce angiotensin-dependent hypertension in the mouse. A solid, stainless steel clip with an opening width of 0.12 mm (a) is placed around the left renal artery to cause stenosis (b), which results in the atrophy of the clipped kidney and hypertrophy of the contralateral, nonclipped kidney (c) and hypertension.

TABLE 1: Cardiovascular parameters in 2K1C mice compared to Sham mice, two weeks after clipping.

Parameter	Sham	2K1C	Reference no.
Mean arterial pressure (mmHg)	~100–115	~120–135	[13, 14]
Heart rate (bpm)	~500–570	~650	[14]
Cardiac weight/body weight index (mg/g) wet (dry)	~4.1 (~1.0)	~4.5 (~1.3)	[13, 14]
Nonclipped kidney weight (mg)	140–160	170–210	[14, 15]
Clipped kidney weight (mg)	150–160	70–90	[14, 15]

for studies of cerebral [8, 9], cardiac [10], vascular [11], and renal [12] functions.

In our laboratory, we used the procedure established by Wiesel et al. [13] to develop a murine model of two-kidney, one-clip (2K1C) hypertension [6, 7, 11, 14]. To minimize variability, a solid stainless steel clip with an opening width of 0.12 mm is placed around the left renal artery (Figure 1) to constrict it and to chronically reduce the perfusion of the left kidney while leaving the other kidney untouched. A mouse body weight of 23 g and clip lumen size of 0.12 mm allows for the induction of hypertension without causing renal infarction [13]. As illustrated in Figure 1, two weeks after clipping, 2K1C mice show atrophy of the clipped (left) kidney and hypertrophy of the contralateral, nonclipped (right) kidney.

Two weeks after renal artery clipping, 2K1C mice already exhibit arterial hypertension (Table 1) with similar levels observed at four weeks [13]. Similar to the 2K1C hypertensive rat that develops cardiac hypertrophy [4, 5],

our laboratory has shown a similar phenotype in the 2K1C hypertensive mouse [6, 14]. The development of cardiac hypertrophy is thought to be the result of increased angiotensin II levels through the stimulation of protein and DNA synthesis in cardiac cells [16].

3. Systemic and Central Renin-Angiotensin Systems

An advantage to using the C57BL/6 mouse for the induction of RAS-dependent hypertension is that it is a prototype of strains with a single renin gene [17], that is, this mouse does not behave differently from the rat in the 2K1C model of renovascular hypertension. As shown in Tables 1 and 2, the high BP in this model is due to a rapid increase in plasma renin levels (~3-fold) in response to a reduction in the perfusion pressure in the stenotic kidney, which secretes renin from juxtaglomerular cells. This is followed

TABLE 2: Average values of plasma renin, angiotensin I, II and 1-7 in 2K1C mice compared to Sham mice, two weeks after clipping.

Parameter	Sham	2K1C	Reference no.
Renin (ng Ang I/mL/hr)*	~1000	~3000	[13]
Angiotensin I (pmol/mL)	~80	~160	[14]
Angiotensin II (pmol/mL)	~30	~140	[14]
Angiotensin 1-7 (pmol/mL)	~90	~180	[14]

* Measured with a microassay based on angiotensin I trapping by antibody.

by a subsequent increase in plasma angiotensin I, which is further converted to the vasoactive angiotensin II (~4.5-fold). Pressure diuresis and hypertrophy of the contralateral kidney (Figure 1, Table 1) prevents hypervolemia [13, 14]. As recently demonstrated by our laboratory, 2K1C mice also show augmented levels of angiotensin 1-7 (Table 2), which is an angiotensin I metabolite formed by a pathway that is independent of angiotensin-converting enzyme (ACE) [18]. Interestingly, knocking-out the angiotensin 1-7 receptor Mas exacerbates the course of 2K1C hypertension in mice [19]. The observed increase in the level of this peptide in the 2K1C mouse seems to serve as an important endogenous, physiological counterbalancing mechanism that partially attenuates the hypertensinogenic actions of activated RAS [18].

In some brain areas, including the rostral ventrolateral medulla (RVLM), hypothalamic paraventricular nucleus (PVN), and subfornical organ (SFO), a local RAS has been identified to act as a critical mediator of chronic hypertension in the 2K1C mouse model [15, 20, 21]. The SFO is a circumventricular region that has a fenestrated vasculature that could permit the entry of increased circulating levels of angiotensin II in addition to residual locally produced angiotensin, leading to the stimulation of the local production of angiotensin II in other brain areas protected by the blood-brain barrier [20].

4. Imbalance of the Cardiac Autonomic Nervous System

An imbalance of the autonomic nervous system, as often occurs in conjunction with several cardiovascular diseases, affects BP and HR variability [22, 23], which may be associated with targeted organ damage and an increased risk of morbidity and mortality [3]. Central areas that are involved in the autonomic control of the cardiovascular system include the rostral ventrolateral and ventromedial medulla (RVLM and RVMM), the caudal ventrolateral medulla (CVLM), PVN, and SFO [24–26]. The signals that are generated in the sinoaortic baroreceptor endings are transmitted through the afferents of cranial nerves XI and X to the nucleus tractus solitarius (NTS), followed by the CVLM, and are processed in the RVLM. The RVLM also integrates inputs from the SFO and PVN, providing a major input to the preganglionic neurons of the sympathetic nervous system [15, 20, 27]. Thus, through the integrative processing of central areas, the autonomic sympathetic and

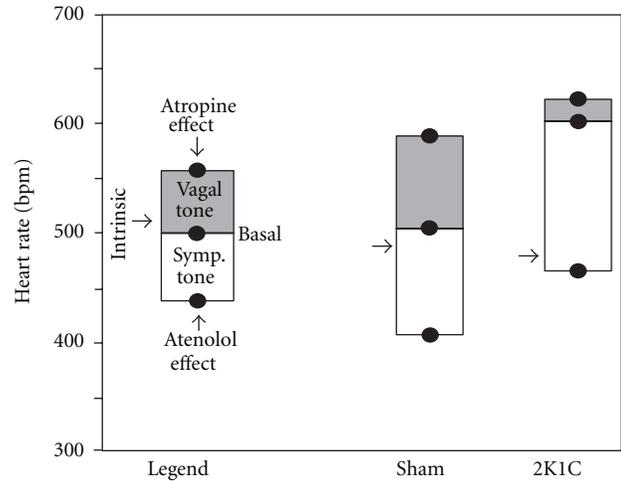


FIGURE 2: Typical imbalance of cardiac autonomic tones in the angiotensin-dependent hypertensive mouse. The cardiovagal tone is represented by the tachycardia observed following the administration of the muscarinic blocker atropine, and the cardiac sympathetic tone is represented by the bradycardia observed after administering the β -adrenergic blocker atenolol. The heart rate after the double blockade indicates the intrinsic heart rate.

parasympathetic nervous systems provide control to the cardiovascular system and the optimal perfusion of organs in accordance with their metabolic needs.

The parasympathetic cardiovagal and sympathetic tones in the mouse have traditionally been assessed through pharmacological methods involving a β_1 -blocker (atenolol), a muscarinic, cholinergic receptor blocker (atropine methyl nitrate) or a double blockade of those receptors [28, 29]. The increase in HR after administering atropine reflects the cardiovagal tone present under baseline resting conditions, and the decrease in HR after atenolol administration reflects cardiac sympathetic tone (Figure 2); a double blockade enables the determination of the intrinsic HR. In the wild-type mouse under resting conditions, a balance between the sympathetic and parasympathetic activities has been reported [7], with a predominance of the sympathetic tone over the cardiovagal tone under special conditions [30].

As shown in Figure 2, the autonomic control of HR in 2K1C hypertensive mice is characterized by an increased cardiac sympathetic tone, whereas the parasympathetic cardiovagal tone is decreased when compared to sham mice [7]. This condition in humans and animal models of cardiovascular diseases represents a major risk factor for cardiovascular mortality [3]. Angiotensin II mediates the increased activity of the sympathetic nerve to the heart in experimental models of RAS-dependent hypertension [32]. In rats, it has been suggested that an infusion of angiotensin II contributes to tachycardia by increasing the intrinsic HR [33]. However, this does not appear to be the case in the 2K1C mouse model, which shows tachycardia without marked changes in this hemodynamic parameter [6]. Considering that the neuronal nitric oxide synthase-(nNOS-) deficient mouse exhibits tachycardia primarily due to abnormal cardiac autonomic control [34] and that

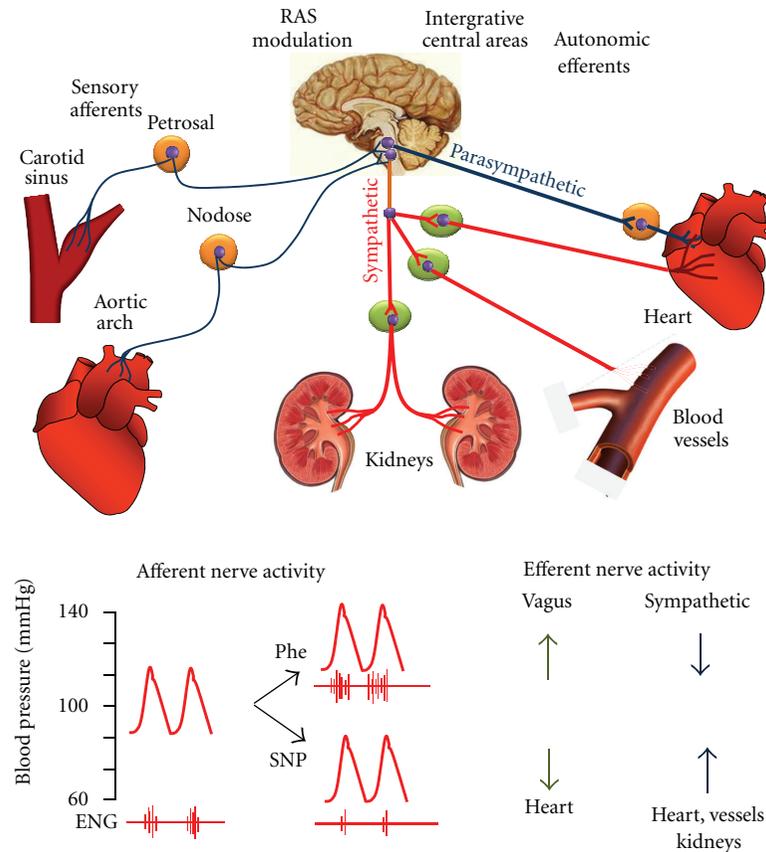


FIGURE 3: Neuroreflex control of circulation. The top panel illustrates the main neural components of the baroreflex arch. The bottom panel shows a schematic illustration of evaluation of the baroreflex function using the vasoactive agent phenylephrine (Phe) and sodium nitroprusside (SNP) in the murine model of renovascular hypertension. ENG, electroencephalogram. The scheme is based on previous publications [3, 31].

endothelial nitric oxide synthase (eNOS) gene therapy restores the basal HR in 2K1C mice [7], it is possible that nitric oxide (NO) plays a role in the autonomic control of HR in this model of RAS-dependent hypertension.

5. Baroreflex Dysfunction

Among the neural systems that control cardiovascular function, the baroreflex is a neural mechanism that acts moment-to-moment to maintain BP with minimal fluctuations [1]. With each arterial systole, mechanosensitive nerve endings located at the carotid sinuses and the aortic arch generate bursts of action potentials that are transmitted to the NTS in the medulla oblongata. Here, the signals are integrated and result in the maintenance of a balanced parasympathetic outflow to the heart and a sympathetic outflow to the heart, vessels and kidneys (Figure 3, top panel). As illustrated in Figure 3 (bottom panel), an immediate rise in BP evokes a reflexive increase in cardiovagal inhibitory activity and a decrease in cardiac and vascular sympathetic excitatory activity, resulting in an immediate correction of BP. Conversely, in response to a rapid decrease in BP, cardiovagal activity is diminished and cardiac and vascular sympathetic activity increase to return the BP to normal values.

In our laboratory, the sensitivity of the baroreflex has been traditionally assessed through pharmacological approaches in conscious animals. An acute, phenylephrine-(Phe-) induced increase in BP leads to an increase in the number of action potentials generated at each discharge and, consequently, to a reflexive increase in parasympathetic and a decrease in sympathetic nerve activities. The opposite is observed during an acute, sodium nitroprusside- (SNP-) induced decrease in BP. Peak values of mean arterial pressure (MAP) and HR in response to Phe and SNP injections are fitted to a sigmoidal logistic equation, which is used to determine the gain (first derivative of the curve) and the maximum reflex tachycardia (upper plateau) and reflex bradycardia (lower plateau) [5, 35]. Considering the small size of the mouse, it is more appropriate to evaluate the baroreflex function in conscious mice by injecting a single dose or by slowly infusing Phe and SNP to avoid volume overloading.

A disruption in the balance between parasympathetic and sympathetic tones, as discussed above, can lead to an impairment in baroreflex sensitivity, as has been demonstrated by our laboratory in different models of hypertension [5, 36–38]. Figure 4 shows representative sigmoidal barocurves of a 2K1C mouse compared to a sham animal. The 2K1C mouse curve is shifted to the right of

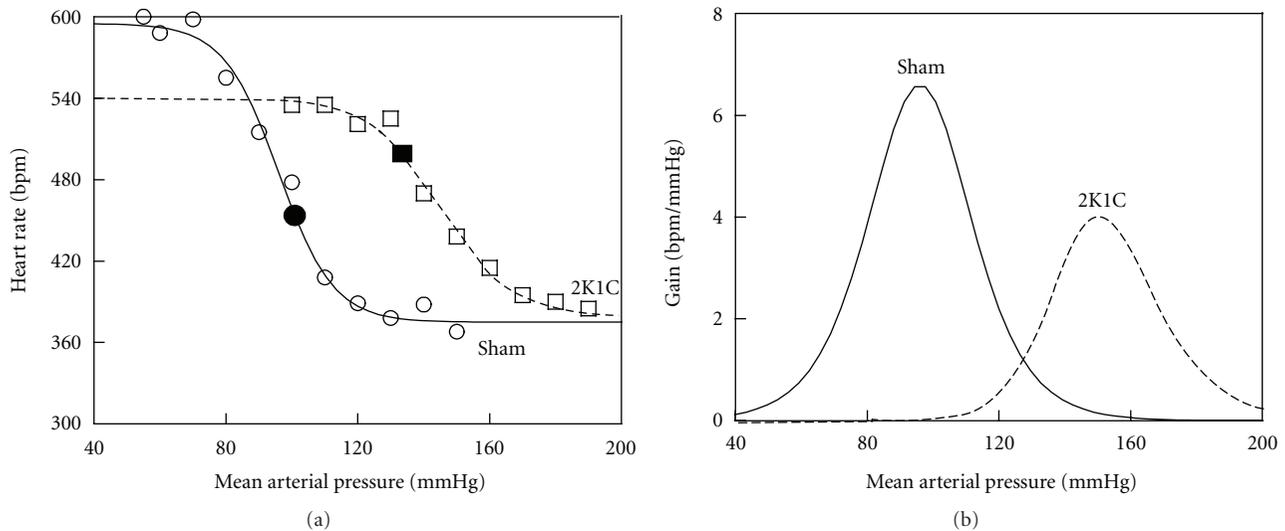


FIGURE 4: Plots showing typical reflex heart rate changes as a function of drug-induced changes in arterial pressure using logistic, sigmoidal-fitting barocurve analysis (a) and baroreflex gains calculated from the first derivative of the sigmoid function (b) comparing 2K1C with sham mice. The small circles and squares indicate individual changes in heart rate in response to every 10 mmHg of drug-induced changes in arterial pressure.

the sham mouse, closely following the high levels of MAP at the midpoint of the curve. The lower slope of the fitting curve indicates impaired baroreflex sensitivity in the 2K1C mouse. We exclude the possibility that the decreased baroreflex sensitivity in 2K1C mice could be due to a limited chronotropic reserve to respond to increases in HR, as the upper plateau of the barocurve of 2K1C mice was below of that observed for sham mice. Based on observations from our laboratory and others, a reasonable explanation for this finding is that, apart from its pressure effect, adventitial angiotensin II and its AT_1 receptors at the aortic arch (and probably at the carotid sinus) act by decreasing the sensitivity of aortic afferents during physiological changes in BP, thus contributing to the impairment of the baroreflex function in cardiovascular diseases [31, 39, 40]. Interestingly, in the rat, central endogenous angiotensin 1-7 has been shown to counterbalance the angiotensin II-induced baroreflex dysfunction [41]. Moreover, a lack of the angiotensin 1-7 Mas receptor-induced baroreflex dysfunction in mice [42]; however, this has not yet been investigated in the renovascular 2K1C mouse model.

6. DNA Damage and Oxidative Stress

There is mounting evidence that increased oxidative stress contributes to increased cardiac and vascular sympathetic tone and decreased baroreflex sensitivity in cardiovascular diseases, including hypertension, as recently reviewed elsewhere [3, 21]. Because ROS play a crucial role in RAS signaling [21, 43, 44], a key mechanism by which angiotensin II influences the heart and vessel function could be through its ability to activate ROS production [20, 45]. ROS have been shown to mediate the actions of angiotensin II at the ganglionic [46] and central nervous system levels, resulting in excessive sympathetic drive to the heart [45, 47, 48]. In

our laboratory, we currently use the comet assay associated with dihydroethidium (DHE) staining to evaluate oxidative stress in different cells and tissues of the 2K1C hypertensive mouse.

The intracellular oxidation of DHE to the fluorescent dye ethidium has been previously used as an indicator of superoxide generation [49]. DHE is freely permeable to cell membranes and can be directly oxidized to ethidium bromide in cell cytoplasm by the superoxide anion [50, 51]. Ethidium bromide becomes trapped in the nucleus by intercalating within DNA, leading to an increase in ethidium fluorescence in the cell nucleus. DHE itself fluoresces blue in the cell cytoplasm, while the oxidized form ethidium fluoresces red following DNA intercalation. Blood cells can be used to assess ROS generation by superoxide detection with DHE.

The most important, biologically active oxidant in the cardiovascular system, superoxide is a highly reactive and short-lived radical responsible for ROS generation. In addition, it can interact with nearby molecules such as DNA, and thus play a key role in inducing DNA oxidative damage [52, 53]. The comet assay is recognized as a versatile and sensitive method for quantifying and analyzing DNA fragmentation in individual cells and can be used to assess DNA exhibiting oxidative damage. The basic principle of the comet assay is the migration of DNA in an agarose matrix under electrophoretic conditions. As a result of this migration, the cells look like comets under microscopic visualization, with a head (intact DNA) and a tail containing DNA fragments. Individual blood cells are embedded in low-melting-point agarose and spread on a common microscope slide. Membranes, soluble cell constituents, and histones are removed by lysing with detergent and high-salt solution. Following the lysis procedure, the slides are placed in an electrophoresis chamber filled with an alkaline

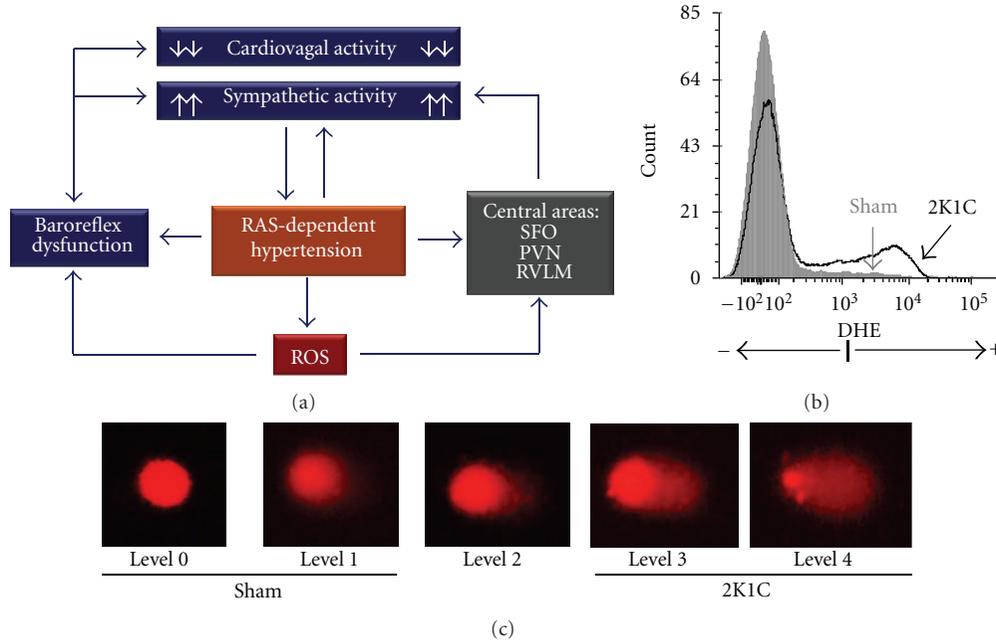


FIGURE 5: Relationship between RAS activation, ROS production, and baroreflex dysfunction in the 2K1C mouse. (a) Effects of renal clipping-induced high plasma levels of angiotensin II on peripheral and central neural areas controlling cardiovascular function mediated by reactive oxygen species (ROS); (b) typical flow cytometric analysis with the dihydroethidium assay (DHE) showing elevated production of superoxide in 2K1C mice; (c) comet assay illustrating the detection of greater levels of DNA damage (comet-tail fragmentation) in the 2K1C mice.

buffer (pH > 13) for DNA unwinding. Then, the DNA undergoes electrophoresis, allowing for the migration of DNA fragments out of the nucleus in an electrical field towards the anode. Staining is usually performed with a DNA-specific fluorescent dye such as ethidium bromide and observed using a fluorescence microscope. The result of this migration is a bright fluorescent head and tail that gives the appearance of a comet. The relative content of DNA in the tail indicates the amount of DNA damage.

As illustrated in Figure 5, the incidence of genomic fragmentation is visually scored into five levels according to the comet-tail size. RAS-dependent hypertensive mice predominantly present comets with elevated DNA damage levels (3 and 4) in whole blood cells. In addition, flow cytometric analysis of blood cells shows an augmentation of DHE staining in these animals, which indicates that 2K1C hypertension increases superoxide generation, in turn, leading to DNA fragmentation in whole blood cells. Ongoing studies are focused on the effects of RAS-induced hypertension in the cells of different tissues.

7. The Relationship between RAS, ROS, and the Autonomic Control of Cardiovascular Function

Our finding of increased DNA damage and ROS production in the 2K1C mouse is in agreement with other studies that found an accumulation of superoxide at the ganglionic level [46] and in different brain integrative areas such as the PVN in this murine model of RAS-induced hypertension

[15, 20, 45]. It is thought that the involvement of the PVN in 2K1C hypertension occurs through the activation of RVLM-projecting parvocellular neurons in this region, leading to increased sympathoexcitation [21, 45]. Based on the above data, a plausible mechanism involved in baroreflex dysfunction and the imbalance of the parasympathetic (diminished) and sympathetic (increased) tones appears to be an excessive generation of ROS in the circulatory system and in both the peripheral and central components of the baroreflex. As recently reviewed, ROS is an insidious and ubiquitous promoter of sympathoexcitation and baroreflex dysfunction that can accelerate or worsen cardiovascular disease processes and cardiovascular risks [3, 21].

8. New Insights in Therapeutic Approaches to Improve Cardiovascular Function in 2K1C Mice

Although therapies have been aimed at nullifying the undesirable effects of angiotensin II, some investigators have focused on demonstrating the importance of the counter-balanced effects of angiotensin 1-7 [18], which is increased in the 2K1C hypertensive mouse [14]. For example, it has been shown that enalapril treatment increases the sensitivity of the baroreflex in the rat, and that this effect was reversed by an i.c.v. infusion of the selective angiotensin 1-7 antagonist, D-Ala7-Ang-1-7 (A-779) [41]. Similar results were observed when A-779 was injected into the CVLM in 2K1C hypertensive rats [54]. Recently, others have shown that the knockout of the angiotensin 1-7 Mas receptor in mice exacerbates

the course of 2K1C hypertension [19]. Considering the identification of the angiotensin-converting enzyme homologue ACE2 as an angiotensin peptide-processing enzyme and of Mas as a receptor for angiotensin 1-7, this axis is a putative target for the development of new cardiovascular drugs [18]. Furthermore, there has been a lack of studies evaluating the effects of peripheral and central manipulation of angiotensin 1-7 on cardiac autonomic tones and the baroreflex function in the 2K1C mouse model.

As recently reviewed [21], there is growing evidence that acute or chronic antioxidant treatment decreases BP and sympathetic activity and improves the baroreflex control of HR in 2K1C rats. Furthermore, tempol or vitamin C administered systemically or into the RVLM or PVN diminishes BP and sympathetic activity [21], highlighting the pivotal role played by central integrative areas controlling cardiovascular function in RAS-dependent renovascular hypertension.

Some gene therapies have also been tested in studies of 2K1C hypertensive mice, but up until now, no favorable results have been observed. For example, Gava et al. [7] used gene therapy in 2K1C mice and observed that it prevented the development of hypertension but not baroreflex dysfunction. Burmeister et al. [15] tested the hypothesis that excessive superoxide anion production in the PVN contributes to the development and maintenance of renovascular hypertension by delivering an adenovirus encoding superoxide dismutase (AdCuZnSOD) to the PVN. They observed that this prevents the elevation in superoxide anions and abolishes renovascular hypertension. However, this approach has not yet been used to evaluate effects on baroreflex dysfunction in the 2K1C mouse.

9. Conclusions and Perspectives

In the past few years, the mouse model of 2K1C hypertension has greatly contributed to the understanding of the relationships between RAS and neural control of cardiovascular function. In addition to the actions of systemic angiotensin II, it has been demonstrated that a local RAS in RVLM, PVN and SFO brain areas act as a critical mediator of chronic hypertension in this experimental model. The 2K1C hypertensive mouse exhibits a cardiac autonomic imbalance characterized by an increased sympathetic tone and a decreased vagal tone, beyond impaired baroreflex sensitivity. In addition to the demonstrations that ROS play a crucial role in RAS signaling at the ganglionic and central nervous system levels, there are growing evidences that DNA damage and increased oxidative stress contribute to the increased cardiac and vascular sympathetic tone and decreased baroreflex sensitivity in the renovascular hypertension. It is well known that angiotensin II increases superoxide production through the activation of NADPH oxidase. Gene therapies by delivering an adenovirus encoding eNOS or enzymes that prevent the elevation of superoxide anions have shown to improve the cardiac autonomic control of HR and baroreflex sensitivity and to prevent renovascular hypertension in the murine model. Although therapies have been aimed at nullifying the undesirable effects of angiotensin II, a putative target for the development of new cardiovascular drugs is

the angiotensin 1-7 which induces the release of NO and diminishes NADPH oxidase activation, counteracting the effects of angiotensin II. Therefore, future studies should address potential strategies to decrease oxidative stress and to prevent or restore the cardiac autonomic balance and the baroreflex function in the mouse model of renovascular hypertension.

Acknowledgments

E. C. Vasquez and S. S. Meyrelles are supported by National Council for the Development of Science and Technology (CNPq, Ref. 302582/2011-8 and 302535/2009-8 Grants, resp.) and the State Agency for the Development of Science and Technology (FAPES, Ref. FAPES/PRONEX 012/2009).

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