

Review Article

Animal Models of Cardiovascular Diseases

**Carlos Zaragoza,¹ Carmen Gomez-Guerrero,² Jose Luis Martin-Ventura,²
Luis Blanco-Colio,² Begoña Lavin,¹ Beñat Mallavia,² Carlos Tarin,² Sebastian Mas,²
Alberto Ortiz,² and Jesus Egido²**

¹ *Department of Epidemiology, Atherothrombosis and Cardiovascular Imaging, Fundacion Centro Nacional Investigaciones Cardiovasculares Carlos III (CNIC), Sinesio Delgado 3, 28029 Madrid, Spain*

² *Renal and Vascular Research Laboratory, IIS-Fundacion Jimenez Diaz, Universidad Autonoma, Avda Reyes Catolicos 2, 28040 Madrid, Spain*

Correspondence should be addressed to Jesus Egido, jegido@fd.es

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Cardiovascular diseases are the first leading cause of death and morbidity in developed countries. The use of animal models have contributed to increase our knowledge, providing new approaches focused to improve the diagnostic and the treatment of these pathologies. Several models have been developed to address cardiovascular complications, including atherothrombotic and cardiac diseases, and the same pathology have been successfully recreated in different species, including small and big animal models of disease. However, genetic and environmental factors play a significant role in cardiovascular pathophysiology, making difficult to match a particular disease, with a single experimental model. Therefore, no exclusive method perfectly recreates the human complication, and depending on the model, additional considerations of cost, infrastructure, and the requirement for specialized personnel, should also have in mind. Considering all these facts, and depending on the budgets available, models should be selected that best reproduce the disease being investigated. Here we will describe models of atherothrombotic diseases, including expanding and occlusive animal models, as well as models of heart failure. Given the wide range of models available, today it is possible to devise the best strategy, which may help us to find more efficient and reliable solutions against human cardiovascular diseases.

1. Introduction

Cardiovascular diseases are the first leading cause of death and morbidity in developed countries. Cardiac and vascular complications are complex multifactorial pathologies, in which both genetic and environmental factors are implicated, thus making them very difficult to prevent. The development of animal models of cardiovascular disease (CVD), including cardiac and atherothrombotic diseases, has provided us today with important insights into the pathophysiology, and they were found to be essential tools to evaluate new therapeutic strategies to predict and to prevent these complications.

Here, we will summarize the most common models of cardiovascular diseases, including those implemented in both large and small animals, designed for helping to cover with more precision and to better understand every single

aspect related to these human pathologies. In particular, we will describe models of atherothrombotic diseases, including expanding abdominal aortic aneurysms (AAA), thoracic aneurysms, and occlusive atherosclerotic diseases, as well as models of heart failure. These situations constitute today a significant challenge since predictors to evaluate early detection and forecast progression are crucial in these pathologies, yet they are poorly explored.

2. Animal Models of Atherothrombotic Disease

2.1. Mouse Models. Atherosclerosis is a complex multifactorial disease with different etiologies that synergistically promote lesion development. Mouse models have proved to be useful to study development and progression of atherosclerotic lesion, and several reviews have extensively discussed the different available models [1–3]. In particular,

knockout and transgenic mouse models for atherosclerosis have been instrumental in understanding the molecular and cellular mechanisms involved in atherogenesis, and in evaluating the effectiveness of new and existing atherosclerotic drugs [4].

As wild-type mice are resistant to lesion development, the current mouse models for atherosclerosis are based on genetic modifications of lipoprotein metabolism with additional dietary changes. Among them, low-density lipoprotein receptor-deficient mice (LDLR^{-/-} mice) and apolipoprotein E-deficient mice (apoE^{-/-} mice) are the most widely used. Atherosclerotic lesions seen in these models can be exacerbated by the addition of risk factors such as hypertension or diabetes. Mice have become widely used as models of human atherosclerosis as they offer advantage compared with other species (Table 1).

2.2. LDLR^{-/-} Mice. The LDLR^{-/-} mouse represents a model of familial hypercholesterolemia due to one of the mutations affecting the LDLR, and the plasma lipoprotein profile resembles that of humans. Mice, which are genetically deficient in LDLR manifested delayed clearance of VLDL and LDL from plasma. As a result, LDLR^{-/-} mice exhibit a moderate increase of plasma cholesterol level and develop atherosclerosis slowly on normal chow diet [5, 6]. Interestingly, the severity of the hypercholesterolemia and atherosclerotic lesions in LDLR^{-/-} mice can be accelerated by feeding a high-fat, high-cholesterol diet [5–7], by mutating the apoB gene into an uneditable version [8, 9], and by crossing either with leptin-deficient mice [10] or with apoB100 transgenic mice [11]. Under these conditions, the lesions in the aorta can progress beyond the foam-cell fatty-streak stage to the fibroproliferative intermediate stage.

In addition to LDLR^{-/-} mice, the LDLR and apoE double-deficient mouse (LDLR^{-/-}apoE^{-/-}) which develops severe hyperlipidemia and atherosclerosis even on a regular chow diet, has been proposed as a suitable model to study the antiatherosclerotic effect of compounds without having to feed the animals an atherogenic diet [12, 13]. However, the response of both LDLR^{-/-} and LDLR^{-/-}apoE^{-/-} mice to the treatment with hypolipidemic drugs varies from lowering of plasma cholesterol without atherosclerosis decrease to a weak lesion reduction with or without lower plasma cholesterol [4, 14]. By contrast, those mice effectively respond to agonists of peroxisome proliferator-activated receptor (PPAR) or liver X receptor [15, 16]. This great variability indicates that LDLR^{-/-} is probably not well-suited for analyzing the cholesterol-lowering and antiatherogenic effects of drugs.

2.3. ApoE^{-/-} Mice. In 1992 two different groups simultaneously generated the apoE^{-/-} mice by homologous recombination in embryonic stem cells [17, 18]. Homozygous deficiency in apoE gene results in a marked increase in the plasma levels of LDL and VLDL due to a failure in their clearance through the LDLR and LDLR-related proteins. The apoE^{-/-} mouse contains the entire spectrum of lesions observed during atherogenesis and was the first mouse model

described to develop lesions similar to those of human [17, 18].

Under normal dietary conditions, apoE^{-/-} mice have dramatically elevated plasma levels of cholesterol, and they develop extensive atherosclerotic lesions widely distributed throughout the aorta [18–20]. This process can be exacerbated on a high-fat diet, with female mice more susceptible than male mice [19]. A chronological analysis of atherosclerotic lesions in apoE^{-/-} mice revealed that the sequential events involved in lesion formation in this model are strikingly similar to those in larger animal models and in humans. Predilection sites for atherosclerotic lesions in apoE^{-/-} mice are the aortic root, followed by the aortic arch, the brachiocephalic trunk, the left carotid, and subclavian and coronary arteries [6, 21]. Aortic lesions rapidly develop from initial fatty streaks comprised primarily of foam cells with migrating smooth muscle cells to more complex lesions in middle-aged mice. These advanced lesions are heterogeneous but typically composed of a necrotic core surrounded by proliferating smooth muscle cells and extracellular matrix proteins [20, 22].

The apoE^{-/-} mice are currently the most widely utilized animal model for the study of atherosclerosis. In fact, the effect of many genes on the development of atherosclerosis has been examined by crossing the apoE^{-/-} mice with other genetically manipulated animals. Furthermore, the apoE^{-/-} mouse serves as a useful tool to: (i) identify atherosclerosis-susceptibility-modifying genes, by the candidate-gene and gene-mapping methods, (ii) decipher molecular mechanism and cell types involved in atherogenesis, (iii) search into the drug effects on atherosclerosis, and (iv) assess novel therapies that prevent lesion progression. In this sense, the apoE^{-/-} mouse model was used to test additional therapeutic effects of statins beyond those attributable solely to cholesterol-lowering. One of the first observations was the paradoxical effect of simvastatin on atherogenesis in both apoE^{-/-} and LDLR^{-/-} mice [23]. In contrast to the atheroprotective effect of simvastatin in LDLR^{-/-} mice, age-matched apoE^{-/-} showed elevated serum total cholesterol and increased aortic plaque area, thus suggesting that the therapeutic effect of simvastatin may depend on the presence of a functional apoE [23]. In spite of this, the antiatherosclerotic effects of other statins have been effectively proven in apoE^{-/-} mice [24, 25]. Several other compounds, such as angiotensin II receptor antagonists or PPAR agonists [26] also reduced the extent and severity of atherosclerotic lesions without lowering plasma cholesterol in apoE^{-/-} mice. However, the recent finding of increased atherogenesis in apoE^{-/-} mice treated with PPAR alpha and PPAR gamma agonists is consistent with clinical findings of the adverse cardiovascular events of dual therapy [27].

Nevertheless, a major limitation of the apoE^{-/-} mouse model is the infrequency of plaque rupture and thrombosis, two common complications of human atherosclerosis. Ischemic cardiomyopathy has been occasionally found in aged mice [20], but interestingly, rapid coronary artery occlusion, myocardial infarction, and even premature death occur when apoE^{-/-} mice were crossed with mice deficient in scavenger receptor class B type I or its adaptor protein

TABLE 1: Animal models of atherosclerosis: advantages and limitations.

Mouse	(i) Rapid development of atherosclerotic plaques	(i) Only partial resemblance to humans
	(ii) Short reproductive cycle	(ii) More atherosclerotic than atherothrombosis model
	(iii) Large litters	(iii) Very high levels of blood lipids
	(iv) Well-known genome	
	(v) Relative ease of genome manipulation	
	(vi) Relatively cheap	
	(vii) Useful for noninvasive imaging (MRI, PET, CT, ultrasound)	
	(viii) Large experience	
Rat	(i) Easy, available, and cheap	(i) Do not develop atheroma
	(ii) Useful for restenosis analysis	
Rabbit	(i) Medium size	(i) Need for high blood cholesterol levels
	(ii) Fibroatheroma lesions	(ii) No plaque rupture model
	(iii) Useful for restenosis models	(iii) A model of neointima formation rather than atherosclerosis
	(iv) Affordable	
Porcine	(i) Lesions more similar to human disease	(i) High cost
	(ii) Valid for restenosis studies	(ii) Difficult handling
		(iii) Few genomic tools

[28, 29], thus mimicking many cardinal features of human coronary heart disease.

2.4. Transgenic Mice. Transgenic technologies have provided a series of very useful mouse models to study hyperlipidemia and atherosclerosis. Among them, mice expressing mutant forms of apoE, such as apoE3Leiden (E3L) and apoE (Arg 112 → Cys → 142) transgenic mice, are the more widely studied. These mice display a lipoprotein profile comparable to that of patients with dysbetalipoproteinemia, in which plasma total cholesterol and triglycerides are mainly confined to (V) LDL [30]. The E3L transgenic mice develop atherosclerotic lesions with all the characteristics of human vasculopathy, varying from fatty streak to mild, moderate, and severe plaques [30, 31]. Furthermore, E3L transgenic mice and the more recently developed E3L/Cholesteryl ester transfer protein (CETP) transgenic mice have been shown to be more sensitive to a variety of hypolipidemic drugs and PPAR agonists than apoE^{-/-} and LDLR^{-/-} mice [4, 32].

2.5. Mouse Models of Diabetes-Accelerated Atherosclerosis. Diabetes is a high risk factor of cardiovascular disease. The cardiovascular complications of diabetes are manifested

primarily as ischemic heart disease caused by accelerated atherosclerosis, and also as cardiomyopathy. Several models are available to study atherosclerosis and cardiomyopathy associated with diabetes, including apoE^{-/-} and LDLR^{-/-} mice in which type 1 diabetes is induced by streptozotocin or viral injection [33, 34]. In both mice, diabetes induction did not markedly change plasma lipid levels, thereby mimicking the accelerated atherosclerosis seen in patients with type 1 diabetes. Importantly, streptozotocin-injected atherosclerotic mice exhibited increased atherosclerosis in the aortic sinus, carotid artery, and abdominal aorta, as well as calcifications in the proximal aorta [34, 35].

In brief, mouse models have been very useful to unveil the importance of inflammatory and immunological mechanisms in the formation and progression of atheroma plaque. Recently, an enormous interest for the use of noninvasive magnetic resonance imaging (MRI) in mouse models of atherosclerosis has arisen [36], since MRI accurately characterizes the location, the size and the shape of lesions. In addition, MRI allows the differentiation between fibrous and lipid components of regress in plaques in mice. In combination with noninvasive imaging technologies, mouse models of atherosclerosis today also serve to test novel

contrast agents, and to design and target specific molecules involved in high-risk plaque.

2.6. Rabbit Models. The high-cholesterol diet rabbit model has been widely used for experimental atherosclerosis. Back in 1913, cholesterol was found to cause atherosclerotic changes in the rabbit arterial intima, which is very similar to human atherosclerosis. Atherosclerotic lesions also develop in normolipemic rabbits as a result of repeated, or continuous intimal injury by an indwelling aortic polyethylene catheter, balloon angioplasty or nitrogen exposure. Therefore, many studies have used the rabbit model with high-cholesterol diet, arterial wall injury, or, most commonly, a combination of these two methods. In all these models, the observed lesions resemble, at least in part, those seen in human plaques, mainly regarding the inflammatory component, though the vascular smooth muscle cell proliferation determines for a great deal the lesion.

The rabbit model has largely been used to study the influence of lipid lowering (by diet or statins) on the plaque formation and “stabilization.” Those studies have contributed to unveil the mechanisms by which lipid lowering reduces macrophage accumulation and other aspects of atheroma inflammation [37, 38].

Recently, we have set up a novel rabbit model to examine the influence of inflammation on atherosclerotic plaque. The aim was to study some mechanisms by which atherosclerosis is particularly severe in patients with rheumatoid arthritis. Briefly, the model consists in a combination of femoral injury in hyperlipidemic rabbits and induced acute knee arthritis. Those animals had more intensive vascular lesions than animals without inflammation. This model could represent a novel approach to the study, inflammation-associated atherosclerosis [39].

A model for plaque rupture has been also developed in rabbits. Shimizu and coworkers [40] have developed a simple rabbit model of vulnerable atherosclerotic plaque, with the combination of aggressive vascular injury associated to a hyperlipidemic diet. The histological findings showed that an aortic plaque had the three features of “vulnerable plaque”: lipid-rich core, accumulation of macrophages, and a thin fibrous cap. In addition, a low-density lipoprotein (LDL) receptor-deficient animal model (the WHHL rabbit) has been developed. This model resembles to human familial hypercholesterolemia and shows evidence of progressive disease of the aorta with accumulation of birefringent lipids in intimal lesions and plaques, as well as in the media from birth to 1 year of age.

Although rabbit aortic arteries are smaller in vessel diameter than human carotid artery, they allows the studies with endovascular therapeutic devices. In addition, the rabbit model has also been used for the quantification of atherosclerotic aortic component by MRI. This technique accurately quantifies fibrotic and lipid components of atherosclerosis in the model and may permit the serial analysis of therapeutic strategies on atherosclerotic plaque stabilization [41].

2.7. Porcine Models. The prevention of heart attack and stroke depends on the detection of vulnerable plaques and development of plaque-stabilizing therapies. Animal models are essential for testing mechanistic hypotheses in a controlled manner, they should be representative of a human disease, and at the same time be easy to manipulate. However, vulnerable plaque recreation is one of the toughest tasks in animal model design. Plaque rupture is an additional complication of an already complex process, and the precise mechanisms involved remain hypothetical. A plethora of experimental approaches are available for growing atherosclerotic lesions in various animal species as mentioned above (Table 2).

Currently, there is no single and golden standard animal model of vulnerable plaque, but pig models are probably the best way to recreate human plaque instability. The combination of diabetes and hypercholesterolemia constitutes a good model of accelerated atherosclerosis [42], and it was relevant study the role of certain biomarkers, such as the Lp-PLA2 since these animals share a similar plasma lipoprotein profile humans. In this regard, the selective inhibition of Lp-PLA2 by darapladib decreased progression to advanced coronary atherosclerotic lesions and confirmed a crucial role of vascular inflammation not associated to hypercholesterolemia, in the development of lesions implicated in the pathogenesis of myocardial infarction and stroke [43].

Several porcine models of advanced human-like coronary atherosclerosis have been employed to analyze the development and validation of coronary imaging technologies. In the evolving era of technological development, the availability and use of such animal models will become critical for the development of emerging technologies in interventional cardiology [44], and for the study of drug-eluting stents [45]. In addition, the porcine models of coronary atherosclerosis allow examining the impact of adventitial neovascularisation, on atherosclerotic plaque composition and vascular remodelling [46].

3. Animal Models of Abdominal Aortic Aneurysms (AAAs)

Animal models of atherothrombotic AAA are essential tools for the preclinical evaluation of new therapeutic strategies for the suppression of aneurysmal degeneration (Table 3). Recent insights into the mechanisms of human AAA have come from the studies in mouse models, and elastase-induced AAA in particular appears to recapitulate many features of human AAA. Here we briefly outline the most frequently used models of AAAs, and refer the reader to recent comprehensive reviews regarding additional animal models [47–52].

3.1. Rat Models

3.1.1. Localized Aortic Perfusion with Elastase. This model consists of exposing a segment of the abdominal aorta and infusing it with elastase [53]. The degradation of elastic fibers triggers an inflammatory response that develops into

TABLE 2: Animal models of plaque rupture and plaque associated thrombosis.

Spontaneous	Induced
39–54-month-old pigs with inherited hyper=LDL cholesterolemia.	ApoE ^{-/-} mice after squeezing the aorta supplemented between forceps.
42–54-week old ApoE ^{-/-} mice.	Hypercholesterolemic rabbits subjected to balloon injury.
Dahl salt-sensitive hypertensive transgenic rats for human cholesteryl ester transfer protein.	Atherosclerotic ApoE ^{-/-} mice subjected to photochemical injury.
—	Intraperitoneal injection of Russell's viper venom in New Zealand White rabbits intermittently fed with high cholesterol diet.
—	Intraperitoneal injection of Russel's viper venom in Watanabe heritable hyperlipidemic rabbits, in combination with the administration of serotonin or angiotensin II.

an aneurysm [54, 55]. The severity of the induced AAA can be increased by adding plasmin to the infusion. This model has been adapted for use in several other species, including rabbit, mouse, and large animals.

3.1.2. Decellularized Xenografts. This model was based on the observation that chronic rejection of arterial allografts and xenografts, results in arterial wall dilatation and rupture. Michel and coworkers decellularized a section of abdominal aorta from one species (e.g., guinea pig), and the resulting tube of intact extracellular matrix was grafted into another morphologically compatible species, usually rat [58]. The xenogenic extracellular matrix triggers the destruction of host matrix, leading to aneurysm formation. The model has been successfully used to evaluate therapeutic targets [64–69], although the heterogeneity of the aneurysms formed and the lack of vessel rupture are significant limitations.

3.2. Mouse Models. The mouse has become the preferred model for cardiovascular research for several reasons, including the ease of handling, low procedure costs, and the ability to manipulate the mouse genome. Consequently, of all animal models of AAA, mouse models have provided most of the insights into the mechanisms of human AAA. The following models are the most widely-used to date.

3.2.1. Calcium-Chloride-Induced AAA. In this method, initially developed in rabbits [62], calcium chloride is applied periaortically in the region between the renal arteries and the iliac bifurcation. Significant dilatation of the aorta is observed within 14 days, and the severity is significantly increased if calcium chloride is applied together with thioglycolate and if animals are fed a high-cholesterol diet [56]. Unlike other models, calcium chloride application induces AAA without the need for mechanical intervention.

3.2.2. Elastase-Induced AAA. The elastase-induced model was adapted for mice by Pyo et al. [57]. Elastase perfusion in mouse aorta causes a mild-to-moderate dilatation initially, which subsequently develops to a >100% increase in aortic diameter within 14 days. In this model, the degradation of medial elastin is delayed, and the subsequent aortic wall inflammation consists of mononuclear phagocytes

throughout the adventitial and medial layers, with relatively few polymorphonuclear cells localized to the adventitia [57]. Elastase-induced injury increases the expression of MMPs, cathepsins, and other proteases [70], with MMP-9 being localized to aneurysm-infiltrating macrophages [71]. Elastase-induced AAAs thus appear to recapitulate many features of human AAAs, and this model has become a valuable and convenient tool for systematically evaluating the roles of individual gene products in aneurysmal degeneration [71–80].

When compared to calcium-chloride-induced AAA, the main limitation of this method is in the mechanical stress required to recreate medial elastic degradation. However, the protocol resembles the time course of events leading to human AAA, including initial recruitment of leukocytes and mast cells, the development of a transmural aortic wall inflammatory response, and finally the upregulation of extracellular matrix metalloproteinases and other proteases, which induce a progressive degradation of the medial elastin and collagen, leading to the final aortic dilatation.

3.2.3. Angiotensin II-Induced AAA. This procedure was initially developed to define whether increased plasma concentrations of Angiotensin II (Ang II), have a direct effect on the atherogenic process in hyperlipidemic old apoE^{-/-} mice. Unexpectedly, Ang II also produced large suprarenal abdominal aortic aneurysms in these animals [81]. In this model, inflammation of the vessel wall is associated with signaling through AT1a receptors [82], nuclear factor- (NF-) kappaB-mediated induction of proinflammatory genes, including MCP-1, M-CSF, iNOS, COX-2, inhibition of PPARs [83], activation of the NADPH oxidase p47phox [84], c-JUN N-terminal [85], Rho kinases [86], and enhanced recruitment of macrophages [87, 88] and extracellular matrix components and degrading enzymes [89–91], leading to vessel dissection, and rupture. The severity of AAA is higher in hyperlipidemic apoE^{-/-} or LDLR^{-/-} male mice (~60% of mice), when compared to normolipidemic C57Bl/6 mice, although in these models neither hyperlipidemia per se nor atherosclerosis is considered major determinants [92–94].

The model contributed to evidence the implication of the rennin-angiotensin (RAS) system in aneurysmal disease. However, two main limitations should be considered: the

TABLE 3: Current procedures for inducing AAA in animals.

Species	AAA models
Murine	<i>Calcium Chloride</i> [56].
	<i>Elastase infusion</i> [57].
	<i>Angiotensin-II-infused AAA</i> : used in hyperlipidemic (ApoE ^{-/-} or LDLR ^{-/-}) male mice or in wild-type C57BL/6 mice in conjunction with repeated administration of neutralizing TGF- β antibodies [50].
	<i>Decellularized xenografts</i> : grafting of abdominal aortic extracellular matrix from one species to a compatible recipient of a different species [58].
	<i>Spontaneous</i> [59].
Rabbit	<i>Elastase-induced AAA</i> : similar to the murine model [60], also applicable to the carotid artery [61].
	<i>Calcium chloride</i> [62].
Pig	<i>Surgical model</i> : dilatation of the infrarenal aorta with an angioplasty balloon followed by infusion of pancreatic porcine elastase [63].

suprarenal location of the aneurysm (in contrast to the infrarenal location in humans) and the clinical relevance of RAS inhibition, since the association of RAS in human AAA has provided controversial results, pointing towards necessary large population studies.

3.2.4. Spontaneous Mouse Mutants. The blotchy mouse is a mouse strain containing a spontaneous mutation on the X chromosome which leads to abnormal intestinal copper absorption. These animals have weak elastic tissue due to failed crosslinking of elastin and collagen, and develop aortic aneurysms mainly in the aortic arch, thoracic aorta, and occasionally in the abdominal aorta [59]. However, results from this model are difficult to interpret, since the mutation produces many severe additional effects.

3.3. Rabbit Models. Several of the same interventions used in mice are also implemented in rabbits, including elastase infusion [60] and calcium chloride application to the abdominal aorta [56]. Another intervention used in rabbits is elastase infusion in the right carotid artery [61]. The main advantage of rabbits over other animal models is that rabbit aneurysms more closely resemble human aneurysms hemodynamically and histologically. Rabbit models also combine several of the attractive features of small animals, such as the easy housing and handling. In addition, similarly to large animals, rabbit aneurysms can be monitored by accessing through the femoral artery, thus providing an excellent model for testing endovascular therapies [95, 96].

3.4. Porcine Models. Porcine models of AAA have provided significant information about the changes that occur after AAA induction and about the responses to stent deployment. A recently-developed porcine model combines mechanical dilatation by balloon angioplasty with enzymatic degradation by infusion of a collagenase/elastase solution. The model is characterized by gradual AAA expansion associated with degradation of aortic wall elastic fibers, an inflammatory cell infiltrate, and persistent smooth muscle cell loss [63]. A broad number of similarities were found between this model and human AAA, and the procedure may also represent an excellent method to evaluate endovascular

related procedures. Despite the benefits, however, pigs have significant disadvantages, including complex animal handling, the requirement of special housing and surgical room facilities, the elevated cost of the animals, and the reduced sample sizes per assay.

3.5. Thoracic Aortic Aneurysm (TAA). Elastic tissue degradation is also related to the development of thoracic aortic aneurysm (TAA), and mouse models have significantly advanced the understanding of this pathology. TAA is a characteristic feature of Marfan syndrome (MFS), a disorder caused by mutations that affect the structure or expression of the extracellular matrix protein fibrillin-1, a glycoprotein that is associated of extracellular proteins, including integrin receptors and insoluble elastin [97]. Fibrillin-1 mutations in MFS decrease ECM sequestration of latent TGF β , thus rendering it more prone to or accessible for activation [98–100]. TAA progression in MFS is driven by elastic fiber calcification, vascular wall inflammation, intimal hyperplasia, structural collapse of the vessel wall, impaired activation of MAP kinase signaling, and altered synthesis of ECM proteins and matrix-degrading enzymes (MMPs) [101]. Systemic administration with TGF β antagonists has been successfully used to mitigate vascular disease in mouse models of MFS and in children with severe and rapidly progressive MFS [97, 102]. Moreover, studies in mouse models have shown that fibulin-4 and LRP1 are also associated with TAA [103, 104].

4. Animal Models of Heart Failure

Models of heart disease in small animals, particularly rats, have been very useful for the assessment of pharmacological therapies. In addition, several target genes have been identified in genetically modified mouse models. Many of these genes have proved to be crucial in the initiation and progression of heart disease. Below, we describe the animal models currently used to study heart failure, which are also summarized in Table 4.

4.1. Rat Models. Rat models have dominated research into heart damage because, while rats share many of the benefits of mice (low cost, ease of handling, etc.), their larger

TABLE 4: Current procedures for inducing heart damage in animals.

Species	Heart failure models
Murine	<i>Myocardial damage</i> <i>Chemical</i> : isoproterenol administration [106]. <i>Electrical</i> : overlapping burns [107]. <i>Surgical</i> : coronary artery ligation [105].
	<i>Ischemia Reperfusion</i> : Transient occlusion of the left coronary artery [108]. <i>Cryoinfarction</i> : cryo-injuries to the epicardium [109]. <i>Pressure overload</i> : aortic constriction and banding. Aortic stenosis [110, 111]. <i>Transgenic models of dilated cardiomyopathy</i> : mutation of cardiac α -actin protein [112].
	<i>Spontaneous</i> : WHHLMI rabbits [113].
	<i>Coronary artery occlusion</i> : similar to the murine model, also applicable to the carotid artery. This is an excellent model for testing endovascular therapies [114].
	<i>Pressure overload</i> : aortic banding [115], valvular stenosis [116].
	<i>Microembolization</i> [117].
Rabbit	<i>Tachycardia</i> : ventricular pacing [118].
	<i>Aortic stenosis</i> [119].
Dog	<i>Surgical model</i> : balloon occlusion of coronary arteries [120].
Pig	<i>Tachycardia</i> : pacing-induced supraventricular tachycardia [121].

size greatly facilitates surgical and postsurgical procedures. Myocardial damage in rat hearts is induced by three procedures: surgical, pharmacological, or electrical.

The surgical method, first developed by Pfeffer and coworkers, consists of ligating the left coronary artery [105]. In this procedure, left thoracotomy is performed on the anesthetized rat, and the heart is rapidly exteriorized by gentle pressure on the right side of the thorax. The left coronary artery is either ligated or heat cauterized between the pulmonary artery outflow tract and the left atrium. The heart is then returned to its normal position and the thorax immediately closed. Several modifications have been introduced to improve performance and to reduce animal mortality, and left coronary artery ligation is the most common method used to induce acute myocardial damage in rat and other animal models. One important modification is temporary occlusion followed by reperfusion, allowing flow recovery through the previously occluded coronary artery bed. Left coronary artery ligation can thus be used to evaluate diverse parameters resulting from either permanent ischemia or ischemia/reperfusion.

Pharmacological induction of heart damage was first implemented by Bagdon and coworkers in 1963 and is achieved by treatment with the beta-one adrenergic receptor (B-AR) agonist isoproterenol [106]. Isoproterenol administration before ischemia exerts a cardioprotective action in rats, but at the right dose it induces cardiac myocyte necrosis and extensive LV dilatation and hypertrophy.

Isoproterenol treatment and left coronary artery ligation in rats are efficient and reproducible methods that provide valuable information about the underlying mechanisms implicated in human heart disease.

The electrical method consists of generating overlapping burns in exposed rat hearts by applying a 2-mm-tipped soldering iron to the epicardium of the left ventricle [107].

While this is also a valid method, the degree of heart damage produced is not consistent among laboratories, limiting the reproducibility of the results obtained with this procedure.

4.2. Mouse Models. Against the many general advantages of working with mice (ease of handling, low pregnancy times, etc.), investigators choosing them as models of heart failure must consider two important limitations: the small size of the heart and the structural differences with respect to the human cardiovascular system. Nonetheless, the availability of transgenic and knockout strains and the relative ease with which new genetic modifications can be introduced make the mouse one of the most attractive models for research into the molecular basis of heart failure.

One of the most widely used models of heart failure in mice is the left coronary artery ligation procedure, adapted from rat. In some protocols the artery is occluded permanently, but recently procedures for temporary occlusion have been introduced to reproduce human ischemia/reperfusion injury [108]. In this method the left anterior descending coronary artery is occluded and then reperfused, allowing flow recovery through the previously occluded coronary artery bed. Reperfusion is monitored visually, and the infarct can be analyzed by histopathological techniques, and can be documented in real time by non invasive high frequency. The areas at risk and the infarct size are revealed by staining with Evans blue dye and triphenyltetrazolium chloride and are assessed by computerized planimetry. This model has confirmed the benefits of reperfusion, since infarct size was found to be significantly lower than after permanent occlusion of the coronary artery.

The method has been further modified to analyze ischemic preconditioning of the heart. In this method, the left coronary artery is repeatedly occluded to subject the heart to several rounds of brief ischemia and reperfusion,

followed by permanent occlusion. This approach has identified several ischemia-induced genes that confer tolerance to a subsequent ischemic event [122].

More recently, a model of myocardial infarction was developed, in mice and rats, in which a series of cryo-injuries is generated in the heart. This new model is yielding promising results [109].

4.3. Large Animal Models of Heart Failure. Small animal models have provided significant insights into human cardiac pathophysiology. However, rodent and human hearts differ in their architecture, heart rates, oxygen consumption, contractility, protein expression, and even stem cell populations, and there is therefore an obvious need for models of heart failure in large animals.

The first large animals used to study heart failure were dogs, in which models of myocardial infarction and serial microembolization of the coronary artery were developed [117]. However, the preferred large animal model of heart damage is the pig, because the collateral coronary circulation and arterial anatomy of pigs and humans are very similar and infarct size can be accurately predicted [123]. Among several models of MI in pigs, one of the most widely used is balloon occlusion of the left anterior descending coronary artery. In this model, a catheter is inserted through the femoral artery, positioning an angioplasty balloon over a guide wire at a position distal to the second largest diagonal branch of the artery, and infarction is induced by balloon inflation [120]. The similar size and cardiac physiology of pigs and humans mean that this model offers major advantages over models in other species. However, the method requires specialized equipment, dedicated surgical facilities and skilled personnel, limiting the number of laboratories able to conduct these studies.

The rabbit, much less expensive than pig, offers a compromise solution. Rabbit models of heart failure, including coronary artery occlusion models [114], have major advantages over other species. For example, the composition of sarcomeric proteins in rabbits is similar to that in humans, and the sarcolemmic reticulum contributes about 70% of calcium elimination. In addition, the WHHLMI rabbit strain provides a model of spontaneous myocardial infarction requiring no surgical intervention. This model was developed by selective breeding of coronary atherosclerosis-prone WHHL rabbits [124]. The main limitation of the WHHLMI model is that it does not feature plaque rupture, whereas in humans coronary plaque rupture and subsequent intravascular thrombosis are the major causes of acute myocardial infarction. Despite this limitation, the model is valid for the study of atherosclerosis-related heart complications [113, 125].

An additional model of heart failure in large and small animals is pressure overload of the left ventricle, induced by transverse aortic constriction in mice [110] and aortic banding in rats and rabbits. Left ventricle hypertrophy can also be recreated by ventricular pacing in dogs [115, 118, 126], valvular stenosis in rabbits [116], and renal artery constriction or aortic stenosis in rats, hamsters, mice, rabbits and dogs [111, 119].

Another model of heart failure is the dilated cardiomyopathy. Human dilated cardiomyopathy has been modeled in rabbits and pigs by inducing chronic tachycardia with a pacemaker [121, 127]. Transgenic mouse models, involving mutations that predispose to dilated cardiomyopathy, have also proved very useful. These models have identified an association of cytoskeletal and contractile proteins with this pathology, and very recently a transgenic model expressing a mutated cardiac alpha-actin gene was provided, in which calcium sensitivity of myofilaments is decreased and the expression of calcium/calmodulin-dependent kinase II δ (CaMKII δ) is increased [112]. Inhibition of CaMKII δ in these animals prevented the increase in p53 and apoptotic cardiomyocytes and ameliorated cardiac function.

5. Conclusion

Animal models of cardiovascular disease yield important insights into the genetic basis of human cardiovascular diseases and provide a test bed for pharmacological and treatments. Nonetheless, investigators need to carefully consider their choice of model: no single method perfectly recreates the human disease, and there are related considerations of cost, infrastructure and the requirement for specialized personnel. Taking these considerations into account, experimenters therefore need to select models that best reproduce the aspect of disease being investigated. In particular, when moving from bench to bedside it is essential to test procedures in relevant models that yield highly reproducible results, but despite these limitations, given the range of animal models available today it will always be possible to devise an appropriate strategy, and animal models remain the best tools for advancing the understanding of the mechanism of human cardiovascular disease.

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