## Supplementary Figure1

A
Mus musculus angiotensin II receptor, type 1a
MALNSSTEDGIKRIQDDCPRAGRHSYIFVMIPTLYSIIFVVGIFGNSLVVIVIYFY MKLKTVASVFLLNLALADLCFLLTLPLWAVYTAMEYRWPFGNHLCKIASASV SFNLYASVFLLTCLSIDRYLAIVHPMKSRLRRTMLVAKVTCIIIWLMAGLASLPA VIHRNVYFIENTNITVCAFHYESRNSTLPIGLGLTKNILGFLFPFLIILTSYTLIWK ALKKAYEIQKNKPRNDDIFRIIMAIVLFFFFSWVPHQIFTFLDVLIQLGVIHDCK IADIVDTAMPITICIAYFNNCLNPLFYGFLGKKFKKYFLQLLKYIPPKAKSHSSL STKMSTLSYRPSDNMSSAAKKPASCSEVE

B
Homo sapiens angiotensin II receptor, type 1
MILNSSTEDGIKRIQDDCPKAGRHNYIFVMIPTLYSIIFVVGIFGNSLVVIVIYFY MKLKTVASVFLLNLALADLCFLLTLPLWAVYTAMEYRWPFGNYLCKIASASV SFNLYASVFLLTCLSIDRYLAIVHPMKSRLRRTMLVAKVTCIIIWLLAGLASLPAI IHRNVFFIENTNITVCAFHYESQNSTLPIGLGLTKNILGFLFPFLIILTSYTLIWKA LKKAYEIQKNKPRNDDIFKIIMAIVLFFFFSWIPHQIFTFLDVLIQLGIIRDCRIAD IVDTAMPITICIAYFNNCLNPLFYGFLGKKFKRYFLQLLKYIPPKAKSHSNLSTK MSTLSYRPSDNVSSSTKKPAPCFEVE

C
PDownload - Graphics Sortby: Evalue
unnamed protein product
Sequence ID: Ic||Query_71157 Length: 359 Number of Matches: 4
Range 1: 165 to $190 \underset{\text { Expect }}{\text { Graphics }}$ Identities Positives Match $\triangle$ Previous Match
$\begin{array}{lllll}83.3 \text { bits(189) } & \text { Expect } & \text { Identities } & \text { Positives } & \text { Gaps } \\ & 3 e-24 & 24 / 26(92 \%) & 25 / 26(96 \%) & 0 / 26(0 \%)\end{array}$
Query 1 ITRRNPTIEMTMTYCABHYEQQIST 26


## Supplementary Figure 2

WB: AT1-AA AT1-mAb

AT1R


