

nAChRs (species)	Expression system	Assay	NIC	ACE	CLO	IMI	Agonist effect	References
$\alpha 3\beta 4$ (rat)	Xenopus oocyte	Membrane current	$EC_{50} = 4.62 \mu M$	$EC_{50} = 0.13 M$	$EC_{50} = 8.39 mM$	n/a	weak	This study
$\alpha 4\beta 2$ (human)	HEK 293	Membrane current	n/a	n/a	$EC_{50} \sim 0.100 mM$	$EC_{50} \sim 0.05 mM$	weak	Li <i>et al.</i> , 2011 (41)
$\alpha 7$ (rat)	Xenopus oocyte	Membrane current	n/a	$EC_{50} = 0.74 mM$	$EC_{50} = 0.74 mM$	n/a	strong	Cartreau <i>et al.</i> , 2018 (42)

n/a: not available.

Table S1

nAChR subunit	Primer designation	Primer sequence (5'-3') ^a	Purpose
$\alpha 3$	nAChSA	GCACCAGAGTGTCTCCCTCCC	Full-length cDNA
	nAChRB	AGCCTCGATGAACAGTGACAG	
	nAChS1	ttaaaagaattcgccaccATGGGTGTTGTGCTGCTCCC	pGEMHE-ORF
	nAChR2	tgatataagcttCTATGTGTCATCTCTGGCCATCA	
$\beta 4$	nAChSC	GGCTGCCACCCGGCTGGCC	Full-length cDNA
	nAChRD	ACGCCGGGTAGCCTAGGAGTC	
	nAChS3	ttaaaagaattcgccaccATGAGGGGTACGCCCCTG	pGEMHE-ORF
	nAChR4	tgatataagcttCTAGGAGTCCTTGGAGGGTGC	

^aThe restriction sites and Kozak sequence are indicated by single (EcoRI; HindIII) and double underlining, respectively.

Table S2

nAChR subunit	Nucleotide variation (CNS vs AM)	Location ^a	Amino-acid modification
α3	T816C	M2	-
	C903T	M3	-
	G1011A	M3-M4 intracellular loop	-
	G1102A	M3-M4 intracellular loop	G368S
β4	T411C	E loop	-
	G1261C	M3-M4 intracellular loop	G421R

^a Functional miscellaneous features of each nAChR subunit transcript are extracellular loops (A to F loops) involved in the ACh binding site and transmembrane domains (M1 to M4) linked by intracellular and extracellular loops and forming the channel pore.

Table S3

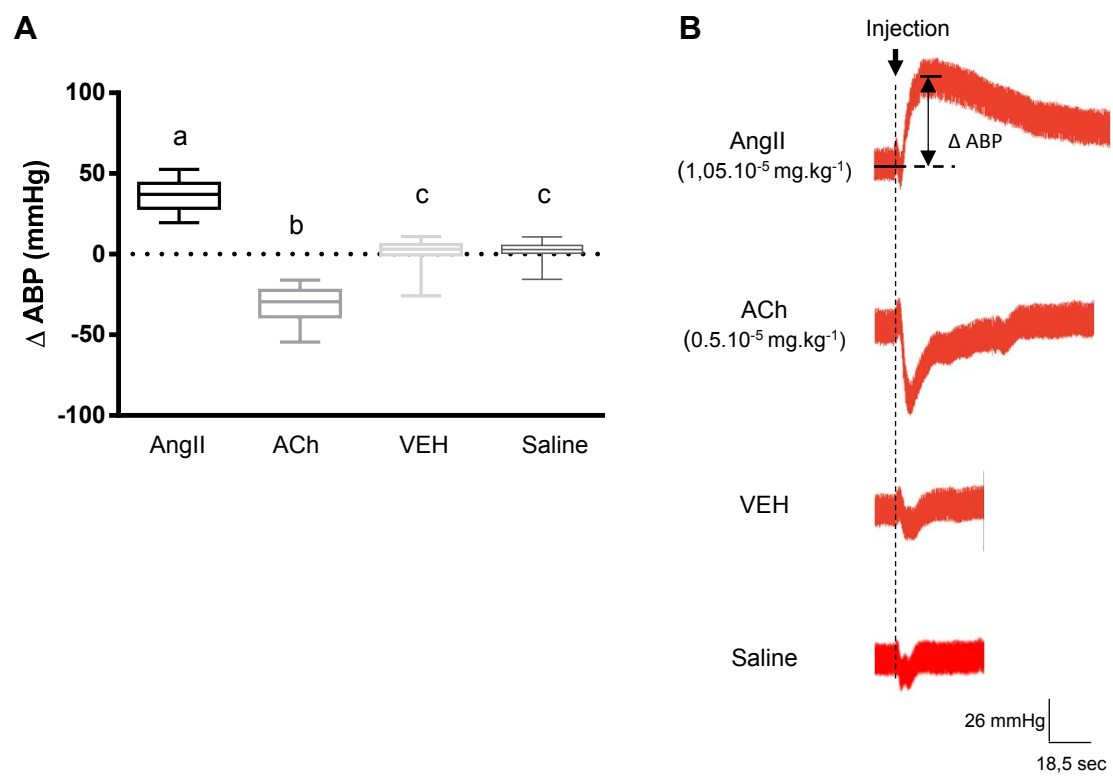


Figure S1

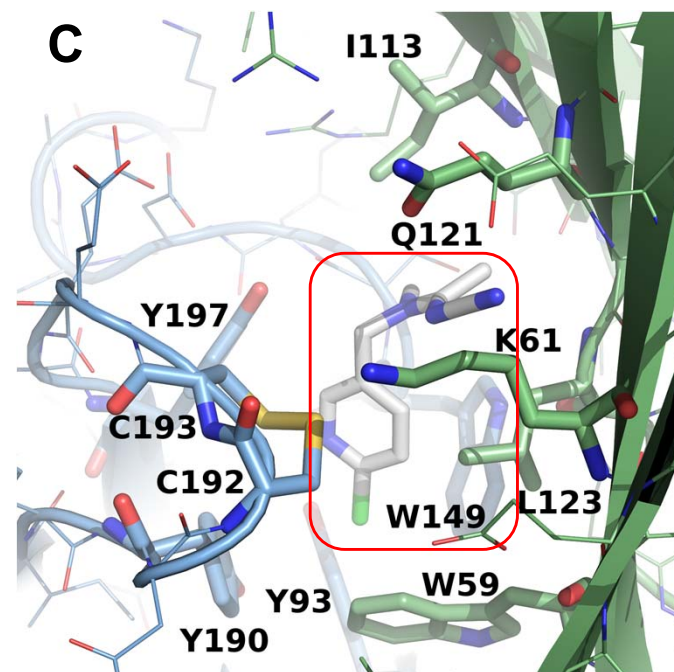
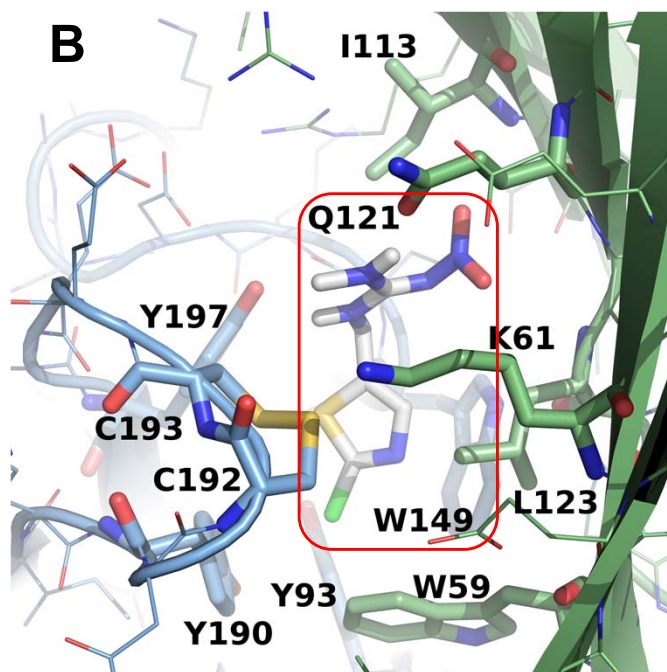
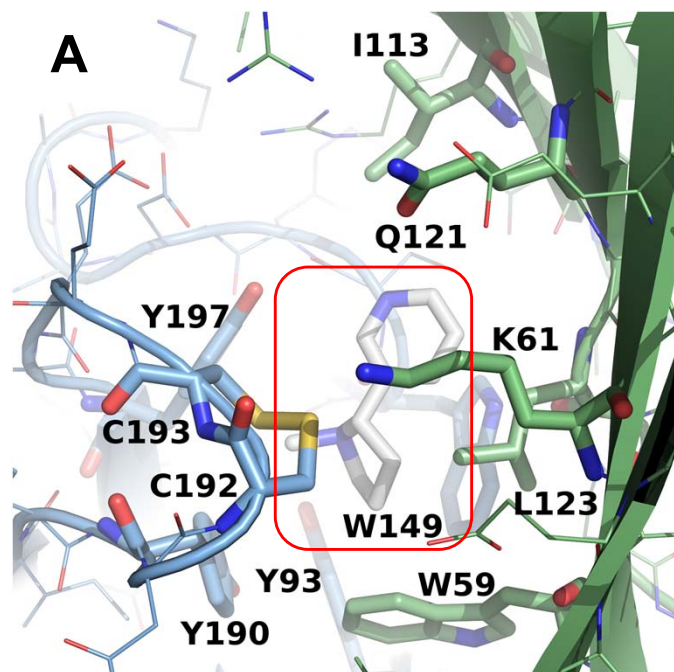


Figure S2

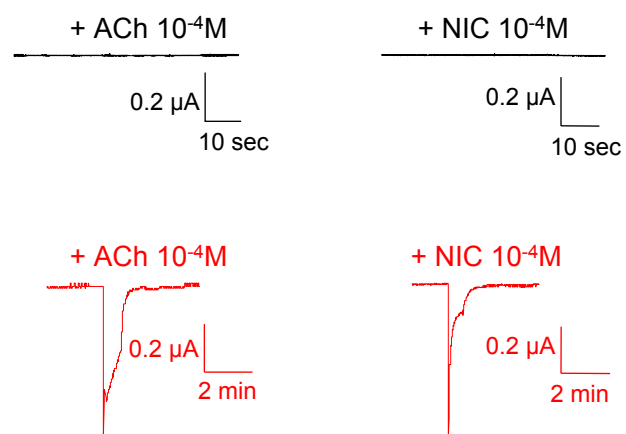


Figure S3

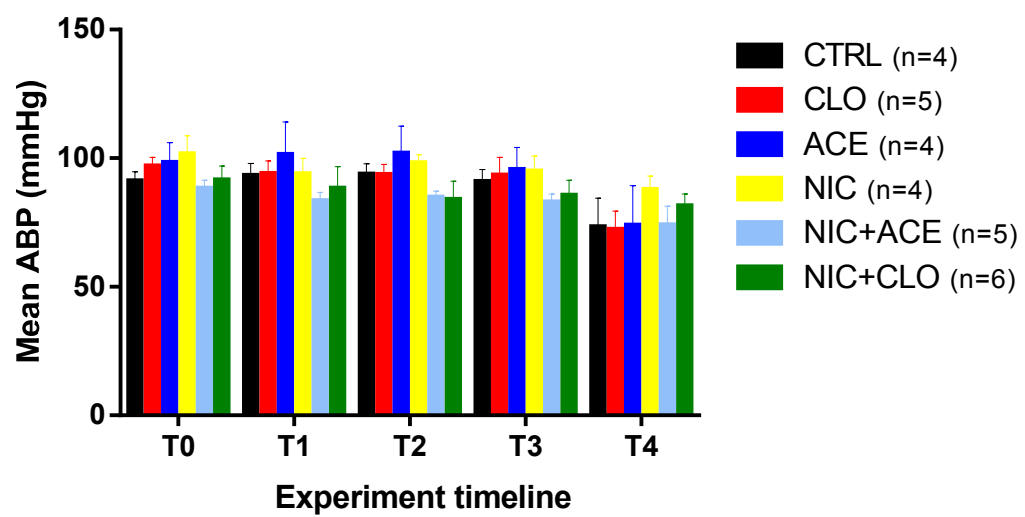


Figure S4