

Key roles of hydrophobic rings of TM2 in gating of the $\alpha 9\alpha 10$ nicotinic cholinergic receptor

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1 We have performed a systematic mutagenesis of three hydrophobic rings (17', 13' and 9') within transmembrane region (TM) 2 of the $\alpha 9\alpha 10$ nicotinic cholinergic receptor (nAChR) to a hydrophilic (threonine) residue and compared the properties of mutant receptors reconstituted in *Xenopus laevis* oocytes.

2 Phenotypic changes in $\alpha 9\alpha 10$ mutant receptors were evidenced by a decrease in the desensitization rate, an increase in both the EC₅₀ for ACh as well as the efficacy of partial agonists and the reduction of the allosteric modulation by extracellular Ca²⁺.

3 Mutated receptors exhibited spontaneous openings and, at the single-channel level, an increased apparent mean open time with no major changes in channel conductance, thus suggesting an increase in gating of the channel as the underlying mechanism.

4 Overall, the degrees of the phenotypes of mutant receptors were more overt in the case of the centrally located V13'T mutant.

5 Based on the atomic model of the pore of the electric organ of the *Torpedo* ray, we can propose that the interactions of side chains at positions 13' and 9' are key ones in creating an energetic barrier to ion permeation.

6 In spite of the fact that the roles of the TM2 residues are mostly conserved in the distant $\alpha 9\alpha 10$ member of the nAChR family, their mechanistic contributions to channel gating show significant differences when compared to other nAChRs. These differences might be originated from slight differential intramolecular rearrangements during gating for the different receptors and might lead each nAChR to be in tune with their physiological roles.

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Abbreviations: ACh, acetylcholine; BAPTA-AM, 1,2-bis(2-aminophenoxy)ethane-*N,N,N',N'*-tetraacetic acid-acetoxymethyl ester; nAChR, nicotinic acetylcholine receptor; TM, transmembrane domain

Introduction

Nicotinic acetylcholine receptors (nAChRs) are members of the 'Cys-loop' family of neurotransmitter-gated ion channels, which also includes GABA_A, GABA_C, glycine, 5-HT₃ and some invertebrate anionic glutamate receptors (LeNovere & Changeux, 1995). They are complexes of protein subunits which co-assemble to form an ion channel gated through the binding of the neurotransmitter. The transmembrane domain 2 (TM2) lines the channel pore and is involved in determining ion selectivity. Residues that participate in channel gating as well as the location of the gate within TM2 has been the subject of numerous studies for several members of the family. In the case of the muscle nAChR, the use of the substituted cysteine accessibility method has suggested that the gate is

located on the cytoplasmic end of TM2 (Akabas *et al.*, 1994; Wilson & Karlin, 1998). Using this method, a more centrally located gate within TM2 has been identified for the 5-HT₃ receptor (Panicker *et al.*, 2002). Moreover, a peptide backbone mutagenesis study has suggested that the central to extracellular residues, 13', 16' and 19', are involved in channel gating (England *et al.*, 1999).

Mutagenesis studies of a conserved leucine (L9') have implicated the middle of TM2 in gating of the nAChRs, GABA_A and GABA_C receptors (Revah *et al.*, 1991; Filatov & White, 1995; Labarca *et al.*, 1995; Chang & Weiss, 1998, 1999). A similar approach has been performed to show the participation of 13' valine (V) within TM2 of the $\alpha 7$ nAChR (Galzi *et al.*, 1992; Corringer *et al.*, 1999). On the other hand, the use of rate equilibrium linear free energy has suggested that the TM2 α helix of muscle nAChRs bends or swivels about its central residues during gating, with the conformational change of the extracellular half preceding the movement of the intracellular half upon opening (Cymes *et al.*, 2002). Finally,

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a 4 Å atomic model of the closed pore of the nAChR of the electric organ of the *Torpedo* ray has indicated that the gate is a constricting hydrophobic girdle that includes L9' and V13' at the middle of the membrane (Miyazawa *et al.*, 2003).

Within the nAChR family, the $\alpha 9$ and $\alpha 10$ subunits are the latest that have been cloned (Elgoyhen *et al.*, 1994; 2001). They are distant members of the nAChR family and form a distinct phylogenetic subfamily (Elgoyhen *et al.*, 1994; 2001; LeNovere *et al.*, 2002). Moreover, heteromeric receptors assembled from these subunits exhibit a peculiar mixed nicotinic-muscarinic pharmacological profile which is distinct from that of other nAChRs and shares properties with GABA_A, glycine and 5-HT₃ receptors. A hallmark of this receptor is that it is not activated by nicotine, the prototypic agonist of the family. Given its lower conservation and distinct properties when compared with other members of the nAChR family, experimental evidence is necessary to probe if residues that have been shown to exert key roles in channel gating are functionally conserved in the $\alpha 9\alpha 10$ receptor.

We have now performed a mutagenesis study of the TM2 region of the $\alpha 9\alpha 10$ nAChR, including three hydrophobic rings of amino acids (outer 17', valine 13' and equatorial 9') proposed to face the lumen of the channel in the stratified organization of an α -helical ionic pore (Bertrand *et al.*, 1993; Karlin & Akabas, 1995; Miyazawa *et al.*, 2003). Hydrophobic residues were mutated to the hydrophilic residue threonine. We have compared the phenotypes of receptors assembled from mutated $\alpha 9$ and $\alpha 10$ subunits at each position. We present evidence indicating that, as reported for other nAChRs, the centrally located amino acids at 9' and 13' are involved in gating of the $\alpha 9\alpha 10$ nAChR. However, different from that reported for muscle nAChRs (Filatov & White, 1995; Labarca *et al.*, 1995), the 9'T mutation is not independent, equivalent or multiplicative in its effect on the responses to acetylcholine (ACh) probably indicating subunit asymmetry in the role these leucines play in activation. Moreover, it is the 13'T and not the 9'T mutation, as observed for the $\alpha 7$ nAChRs, that renders a more drastic phenotype. Based on the proposed atomic model for the gate of nAChRs (Miyazawa *et al.*, 2003), this result could suggest that hydrophobic interactions at 13' might contribute the most to create an energetic barrier to ion permeation. Further mutations to residues other than threonine would be needed in order to prove this notion.

Methods

Generation of mutant receptors

Site-directed mutagenesis of the $\alpha 9$ and $\alpha 10$ rat cDNAs, subcloned in a modified pGEMHE vector (Elgoyhen *et al.*, 1994; 2001), was performed with the QuickChange Site-Directed Mutagenesis kit (Stratagene, La Jolla, CA, U.S.A.). Mutations were confirmed by sequence analysis to verify that only the desired nucleotide changes were present.

Expression of recombinant receptors in *Xenopus laevis* oocytes

Capped cRNAs were *in vitro* transcribed from linearized plasmid DNA templates using the mMessage mMachine T7

Transcription Kit (Ambion Corporation, Austin, TX, U.S.A.). The maintenance of *X. laevis*, as well as the preparation and cRNA injection of stage V and VI oocytes, has been described in detail elsewhere (Katz *et al.*, 2000). Typically, oocytes were injected with 50 nl of RNase-free water containing 0.01–1.0 ng of cRNAs (at a 1:1 molar ratio) and maintained in Barth's solution at 17°C.

Electrophysiological recordings were performed 2–6 days after cRNA injection under two-electrode voltage clamp with a Geneclamp 500 amplifier (Axon Instruments Corp., Union City, CA, U.S.A.). Both voltage and current electrodes were filled with 3 M KCl and had resistances of ~1–2 MΩ. Data acquisition was performed using a Digidata 1200 and the pClamp 7.0 software (Axon Instruments Corp., Union City, CA, U.S.A.). Data were analyzed using Clamp Fit from the pClamp 6.1 software. During electrophysiological recordings, oocytes were continuously superfused (~10 ml min⁻¹) with normal frog saline comprised of (mM): 115 NaCl, 2.5 KCl, 1.8 CaCl₂ and 10 HEPES buffer, pH 7.2. When the effect of nominally zero Ca²⁺ was assessed (Figure 2), experiments were carried out in oocytes injected with 7.5 ng of an oligonucleotide (5'-GCTTTAGTAATTCCTGCCAT GTTTC-3') antisense to connexinC38 mRNA (Arellano *et al.*, 1995; Ebihara, 1996), in order to minimize the activation of the oocyte's nonselective inward current through a hemigap junction channel in response to the reduction of the external divalent cation concentration. However, in the case of the V13'T mutant, current due to both the activation of the nonselective hemigap junction channel plus the constitutively active mutant channels rendered leak currents that were too big to compensate. Therefore, the concentration–response curves of Figure 2c were performed (both those with and without Ca²⁺) in a saline solution that contained 0.5 mM Mg²⁺, in order to minimize the activation of the nonselective inward current. As reported previously, Mg²⁺ does not potentiate ACh currents through $\alpha 9\alpha 10$ receptors; however, it does produce a channel block (Weisstaub *et al.*, 2002). Drugs were applied in the perfusion solution of the oocyte chamber. To minimize activation of the endogenous Ca²⁺-sensitive chloride current (Elgoyhen *et al.*, 2001), all experiments were performed in oocytes incubated with the Ca²⁺ chelator 1,2-bis(2-aminophenoxy)ethane-*N,N,N',N'*-tetraacetic acid-acetoxymethyl ester (BAPTA-AM, 100 μM) for 3–4 h prior to electrophysiological recordings.

Concentration–response curves were normalized to the maximal agonist response in each oocyte. The mean and standard error of the mean of peak current responses are represented. Agonist concentration–response curves were iteratively fitted with the equation: $I/I_{\max} = A^n/(A^n + EC_{50}^n)$, where I is the peak inward current evoked by the agonist at concentration A ; I_{\max} is current evoked by the concentration of agonist eliciting a maximal response; EC_{50} is the concentration of agonist inducing half-maximal current response and n is the Hill co-efficient.

Current–voltage (*I*–*V*) relationships

I–*V* relationships were obtained by applying 2 s voltage ramps from –120 to +50 mV, 10 s after the peak response to ACh from a holding potential (V_{hold}) of –70 mV. Leakage correction was performed by digital subtraction of the *I*–*V* curve obtained by the same voltage ramp protocol prior to the

application of ACh. Generation of voltage protocols and data acquisition were performed using a Digidata 1200 and the pClamp 6.1 or 7.0 software (Axon Instruments Corp., Union City, CA, U.S.A.). Data were analyzed using Clamp fit from the pClamp 6.1 software.

Single-channel recordings

Single-channel currents were recorded in the cell-attached and outside-out patch configuration (Hamill & Sakmann, 1981) at 20°C. Before the experiments, the oocytes were incubated for 4 h in a buffer containing BAPTA-AM in order to block endogenous calcium-activated chloride channels. The vitelline membrane was removed with fine forceps from the oocytes just before the experiments. The bath and pipette solution contained 150 mM NaCl, 0.5 mM CaCl₂, 5.6 mM KCl and 10 mM HEPES (pH 7.4). Solutions free of magnesium and with low calcium were used in order to minimize the channel block (Weisstaub *et al.*, 2002). Patch pipettes were pulled from Kimax capillary tubes (Kimble, Vineland, NJ, U.S.A.) and coated with Sylgard (Dow Corning, Midland, MI, U.S.A.). Pipette resistance ranged from 5 to 7 M Ω . ACh was added to the pipette solution.

The resting potential of the oocytes in this bath solution varied from -30 to -50 mV. Currents were recorded using an Axopatch 200 B patch-clamp amplifier (Axon Instruments Corp., Union City, CA, U.S.A.), digitized at 5 μ s intervals with the PCI-611E interface (National Instruments, Austin, TX, U.S.A.), recorded to the hard disk of a computer using the program Acquire (Bruxton Corporation, Seattle, WA, U.S.A.), and detected by the half-amplitude threshold criterion using the program TAC 4.0.10 (Bruxton Corporation, Seattle, WA, U.S.A.) at a final bandwidth of 8 kHz (Bouzat *et al.*, 1994; 2002). Open-time histograms were plotted using a logarithmic abscissa and a square root ordinate and fitted to the sum of exponentials by maximum likelihood using the program TACFit (Bruxton Corporation, Seattle, WA, U.S.A.).

Statistical significance was evaluated by the Student's *t*-test (two-tailed, unpaired samples). Multiple comparisons were performed with a one-way ANOVA followed by Tukey's test. *P* < 0.05 was considered significant.

Materials

ACh chloride and choline chloride were bought from Sigma Chemical Co. (St Louis, MO, U.S.A.). ICS 205,930 HCl, (-)-nicotine-di-*d*-tartrate and (+)-muscarine chloride were obtained from RBI (Natick, MA, U.S.A.). Drugs were dissolved in distilled water as 10 mM stocks and stored aliquoted at -20°C. BAPTA-AM (Molecular Probes, Eugene, OR, U.S.A.) was stored at -20°C as aliquots of a 100 mM solution in dimethyl sulfoxide, thawed and diluted 1000-fold into saline solution shortly before incubation of the oocytes.

All experimental protocols were carried out in accordance with the National Institute of Health Guide for the Care and Use of Laboratory Animals (NIH Publications No. 80-23) revised 1978.

Results

Mutant receptors exhibit a decreased desensitization rate and an increased ACh EC₅₀

Figure 1a shows an alignment of the TM2 region of different nAChRs and the positions of amino acids (one-letter code) that have been mutated. The numbering used is the one that has been adopted to allow comparison of homologous amino acids from different types of neurotransmitter-gated channels. Position 1' corresponds to the start of the TM2 region. The residues that have been mutated to threonine are the three hydrophobic rings (outer 17', 13' and equatorial 9') of amino acids proposed to face the lumen of the channel in the stratified organization of an α -helical ionic pore (Bertrand *et al.*, 1993; Karlin & Akabas, 1995; Miyazawa *et al.*, 2003). Residues were mutated to threonine, since the introduction of this amino acid has rendered drastic phenotypes in other ligand-gated ion channels (Bertrand *et al.*, 1992; Labarca *et al.*, 1995; Chang & Weiss, 1998).

We first examined whole-cell responses of *Xenopus* oocytes expressing wild-type and mutant receptors. All mutants yielded functional receptors that responded to ACh (Figure 1b). Peak responses to ACh of double mutants ($\alpha 9^*\alpha 10^*$) were similar to those obtained in the wild-type receptor, except in the case of the L9/T, where a reduction was observed (Table 1). Representative responses to increasing concentrations of ACh are shown in Figure 1b and their respective concentration-response curves in Figure 1c. EC₅₀ and Hill coefficient values from a fit of the Hill equation to these data are provided in Table 1. The table also includes data derived from concentration-response curves performed in receptors assembled from single-mutant subunits, that is either $\alpha 9$ or $\alpha 10$ were mutated at each position and co-injected with the wild-type partner subunit. In the case of the double mutants, mutations increased the sensitivity for ACh, with no change in Hill coefficients, as evidenced by a reduction in the EC₅₀. The rank order of potency of ACh for mutated receptors was: 13'T > 9'T > 17'T (*P* < 0.05). Thus, the major shift in the EC₅₀ for ACh, 86-fold, was observed for the 13'T mutant. In spite of the fact that a sole detectable population of receptors is found in oocytes with a ($\alpha 9$)₂($\alpha 10$)₃ stoichiometry (Plazas and Elgoyhen, unpublished observations), no differences in EC₅₀ values were observed at positions 13' and 9' when comparing the $\alpha 9^*\alpha 10$ to the $\alpha 9\alpha 10^*$ mutant receptors, suggesting that at these positions both types of subunits contribute in an asymmetric and nonadditive manner to a pentameric assembly. On the other hand, differences in EC₅₀ values were observed at position 17' (Table 1), where a bigger shift in the EC₅₀ was observed when mutating the $\alpha 10$ subunit. It should be noted that at the concentrations of ACh used in the present study (maximum of 30 μ M for wild-type receptors) we did not find evidence for channel block produced by ACh, as assessed by a rebound in currents after washing. All following experiments were performed with the double-mutant receptors.

Mutant receptors exhibited a decrease in the rate of desensitization at prolonged applications of 100 μ M ACh (Figure 1d), that can be quantified by the percent of maximal peak current remaining after a 30 s application of ACh (Table 1). Moreover, while wild-type currents quickly decayed after removal of the agonist, a substantial residual current could be recorded for several seconds in the case of V13/T

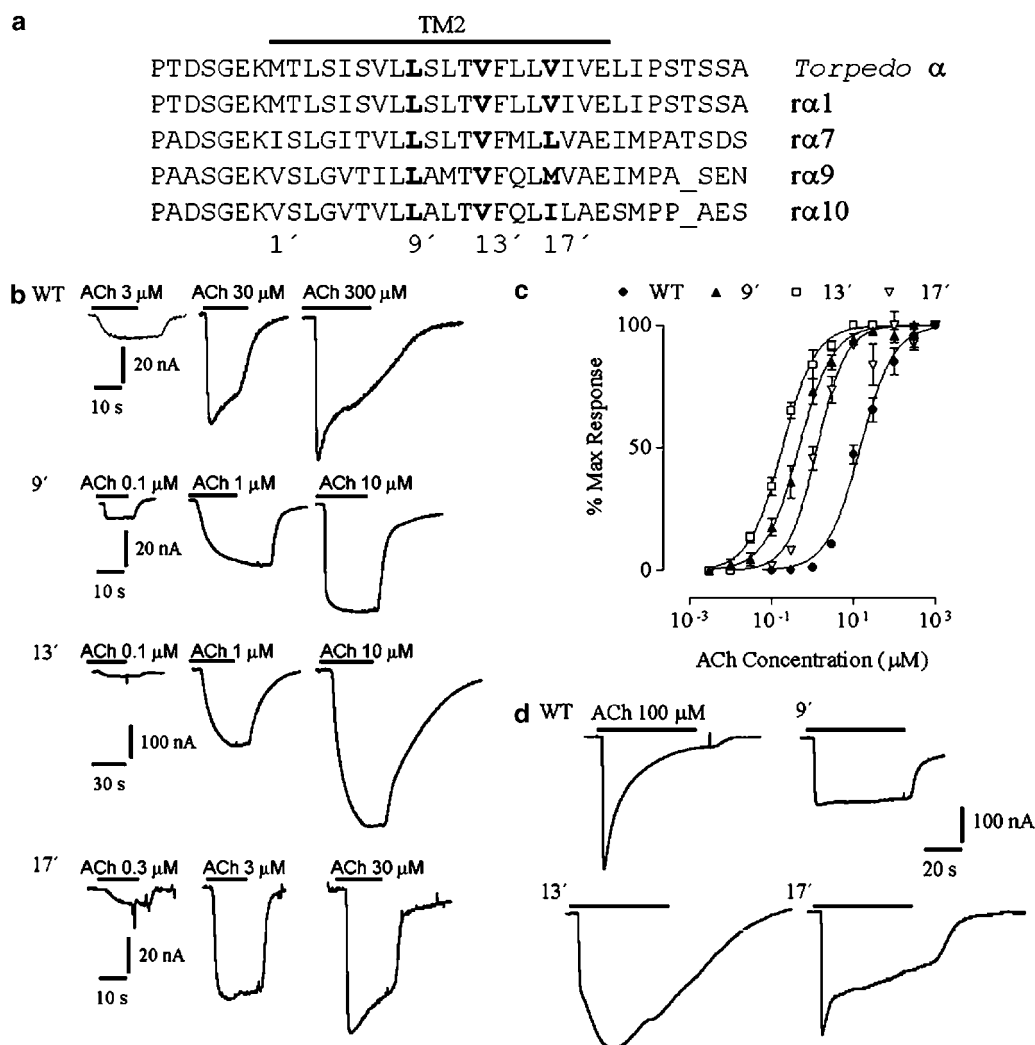


Figure 1 Responses of mutant receptors to ACh. (a) Alignment of the amino-acid sequences of the *Torpedo* α , rat $\alpha 1$, $\alpha 7$, $\alpha 9$ and $\alpha 10$ nAChR subunits. Residues that have been mutated are shown in bold. (b) Representative responses to increasing concentrations of ACh for wild-type and each mutant receptor. (c) Concentration–response curves to ACh. Peak current values were normalized and referred to the maximal peak response to ACh in each case. The mean and s.e.m. of four to five experiments per group are shown. (d) Representative responses of wild-type and mutant receptors to a 1-min application of 100 μ M ACh.

(Figure 1d). In addition, activation of macroscopic currents in V13/T had a fast and a slow component.

Allosteric modulation by extracellular Ca^{2+} is reduced in mutant receptors

It has been shown that external Ca^{2+} modulates the activity of several nAChRs. In particular, the $\alpha 9\alpha 10$ receptor is highly permeable to Ca^{2+} , and it is both potentiated and blocked by physiological concentrations of external Ca^{2+} (Weisstaub *et al.*, 2002). Potentiation is voltage-independent and results in a decrease in the EC_{50} of the receptor for ACh, thus suggesting that Ca^{2+} interacts at an extracellular binding site to allosterically modulate coupling between ligand binding and gating. On the other hand, blockage is voltage dependent, suggesting that the site of action of this ion lies within the channel pore and might result from the permeation process.

The bar diagram of Figure 2a shows the responses to a fixed concentration of ACh at varying concentrations of extracellular Ca^{2+} for wild-type, 9'T and 13'T mutant receptors at a

holding potential of -90 mV. The concentration of ACh used in each case was near the corresponding EC_{50} value derived from the concentration–response curves of Figure 1: 10 μ M for the wild type, 0.5 μ M for the 9'T and 0.1 μ M for the 13'T mutant. Responses were normalized to the value obtained at 1.8 mM Ca^{2+} for each case. As previously shown for the $\alpha 9\alpha 10$ wild-type receptor (Weisstaub *et al.*, 2002), responses to ACh were potentiated by Ca^{2+} up to 0.5 mM, and blocked by higher concentrations of this ion. Moreover, in oocytes that had a low level of subunit expression, responses in the absence of Ca^{2+} were often too small to be detected. On the other hand, in the case of the 9'T and 13'T mutants, the potentiating effect of Ca^{2+} was not observed. Responses were highest at low Ca^{2+} concentrations, suggesting that channel gating by ACh became independent of the presence of extracellular Ca^{2+} . Blockage by Ca^{2+} was still observed. However, the magnitude of block by Ca^{2+} of the 13'T mutant was reduced when compared to that of the 9'T mutant (percentage of response at 3 mM Ca^{2+} compared to 0.1 mM Ca^{2+} : 9'T, $29 \pm 2\%$ and 13'T, $51 \pm 2\%$, $n = 4-10$, $P < 0.001$).

Table 1 Properties of mutant receptors

	Peak current (nA)	I_{30s}/I_{max} (%)	EC_{50} (μ M)	n_{Hill}
WT	600 \pm 100 (40)	16 \pm 3 (11)	14.6 \pm 5.3 (5) ^a	1.1 \pm 0.1
<i>L9'T</i>				
$\alpha 9^*\alpha 10$	547 \pm 145 (30)	89 \pm 2 (12) ^b	5.30 \pm 0.32 (8)	1.7 \pm 0.1
$\alpha 9\alpha 10^*$	155 \pm 35 (12) ^c	88 \pm 5 (6) ^b	5.62 \pm 0.40 (8)	1.8 \pm 0.2
$\alpha 9^*\alpha 10^*$	235 \pm 40 (22) ^c	92 \pm 2 (12) ^b	0.46 \pm 0.04 (5) ^a	1.1 \pm 0.1
<i>V13'T</i>				
$\alpha 9^*\alpha 10$	632 \pm 269 (5)	80 \pm 6 (8) ^b	0.42 \pm 0.02 (5)	1.1 \pm 0.1
$\alpha 9\alpha 10^*$	237 \pm 77 (10)	91 \pm 2 (8) ^b	0.46 \pm 0.02 (5)	1.4 \pm 0.1
$\alpha 9^*\alpha 10^*$	531 \pm 116 (16)	74 \pm 4 (23) ^b	0.17 \pm 0.01 (5) ^a	1.1 \pm 0.1
<i>M117'T</i>				
$\alpha 9^*\alpha 10$	101 \pm 12 (15) ^d	80 \pm 1 (18) ^b	5.47 \pm 0.71 (5)	1.2 \pm 0.2
$\alpha 9\alpha 10^*$	660 \pm 281 (15)	73 \pm 3 (9) ^b	2.86 \pm 0.60 (9)	1.1 \pm 0.2
$\alpha 9^*\alpha 10^*$	430 \pm 67 (10)	67 \pm 3 (11) ^b	1.09 \pm 0.13 (4) ^a	1.5 \pm 0.2

All parameters were determined as described in Methods.

^awt > 17' > 9' > 13', $P < 0.05$, one-way ANOVA followed by Tukey's test.

^b $P < 0.01$ with respect to wild type.

^c $P < 0.05$.

^d $P < 0.01$ with respect to wild type, Student's *t*-test.

Figure 2b shows representative *I*-*V* curves for 9'T and 13'T mutant receptors, obtained by applying 2-s voltage ramps (-120 to +50 mV), 10 s after the peak response to ACh at different Ca^{2+} concentrations. The apparent reversal potentials at 1.8 mM Ca^{2+} , -11 ± 1 , $n = 22$, for wild-type receptors (Elgoyhen *et al.*, 2001), -16 ± 4 , $n = 6$, for the 9' and -9 ± 3 , $n = 6$, for the 13' mutants were not significantly different. Near the reversal potential, the 9'T mutant showed a marked rectification, similar to that previously reported for the wild-type receptor (Elgoyhen *et al.*, 2001; Weisstaub *et al.*, 2002). In contrast, almost linear *I*-*V* curves were observed in the case of the 13'T mutant. Different from what had been previously described for the $\alpha 9\alpha 10$ wild-type receptor (Weisstaub *et al.*, 2002), in the case of mutant receptors, responses were smaller the higher the Ca^{2+} concentration, at all concentrations of the ion tested (Figure 2b). This result is consistent with that shown in Figure 2a, and indicates that whereas the allosteric potentiation by Ca^{2+} is lost in the mutants, its blocking effect is still maintained. Calcium block was clearly voltage-dependent in the case of the 9'T receptor, where no blocking effect was observed at potentials positive to 0 mV. On the other hand, while block by Ca^{2+} was still voltage-dependent in the case of the 13'T mutant, the dependency upon membrane potential diminished. This is evidenced when comparing the ratio of current at 3 mM to 0.2 mM Ca^{2+} at -110 and +40 mV, respectively: 9'T, 0.31 ± 0.08 and 0.98 ± 0.05 ($n = 6$); 13'T, 0.27 ± 0.09 and 0.59 ± 0.10 ($n = 6$).

The bar diagram of Figure 2a shows the effect of Ca^{2+} at only one concentration of ACh. In order to analyze if the potentiating effect of Ca^{2+} is lost at all concentrations of the agonist, full concentration-response curves to ACh were performed and compared at nominally zero and 1.8 mM Ca^{2+} (Figure 2c). In all cases, responses were normalized to the maximal response at Ca^{2+} 1.8 mM. In the case of the wild-type receptor, responses to all concentrations of ACh tested were potentiated in the presence of Ca^{2+} . The maximal response achieved in nominally zero Ca^{2+} was $31 \pm 2\%$ of that at 1.8 mM, and a decrease in potency without changes in the

Hill coefficient was observed (nominally zero Ca^{2+} : EC_{50} , $83.1 \pm 9.2 \mu$ M; n_{Hill} , 1.2 ± 0.3 , $n = 5$; 1.8 mM Ca^{2+} : EC_{50} , $22.6 \pm 2.4 \mu$ M; n_{Hill} , 1.1 ± 0.2 , $n = 5$). In the case of the 9'T mutant, responses were potentiated by Ca^{2+} at concentrations of ACh below 1 μ M and blocked at higher concentrations of the agonist. Potentiation was significant at 0.3 μ M ACh, where responses at zero Ca^{2+} were $1.3 \pm 0.4\%$, $n = 5$, and at 1.8 mM Ca^{2+} $38.7 \pm 12.3\%$, $n = 5$, of the maximum obtained at 1.8 mM Ca^{2+} . On the contrary, responses of the 13'T mutant to ACh were not potentiated by Ca^{2+} and the blocking effect of this ion was evidenced at low and high concentrations of ACh. Note that in a previous work we have shown that responses in zero Ca^{2+} achieve the maximal response at high concentrations of ACh (Weisstaub *et al.*, 2002). However, those experiments were performed in a solution devoid of divalent cations, whereas the present experiments were carried out in the presence of Mg^{2+} , for reasons explained in experimental procedures. Moreover, parameters derived from Figure 3c are not comparable to those of Table 1 because of the different ionic composition of the saline solution.

Mutant receptors retained the high Ca^{2+} permeability previously described for the $\alpha 9\alpha 10$ wild-type receptor (Weisstaub *et al.*, 2002); $P_{Ca}/P_{monovalents}$: wt, 9 ± 1 , $n = 5$; 9'T, 12 ± 6 , $n = 5$; 13'T, 11 ± 5 , $n = 7$; 17'T, 12 ± 6 , $n = 7$, data not illustrated).

Choline, a partial agonist of wild-type receptors is a full agonist of mutant receptors

Choline, the metabolite of the enzymatic degradation of ACh, has been shown to activate several nAChRs, including $\alpha 7$ and $\alpha 9$ (Papke *et al.*, 1996; Verbitsky *et al.*, 2000). Figure 3 indicates the concentration-response curves to choline, for both wild-type $\alpha 9\alpha 10$ and V13'T mutant receptors. Responses were normalized to the maximal responses to ACh in each case. Curves for other mutations are not shown, but rendered similar results. Choline was a partial agonist of wild-type receptors, with a maximal response that reached 36% of the maximum to ACh and an EC_{50} of 538 μ M (Table 2). On the other hand, in the case of the mutant receptors, choline behaved as a full agonist, reaching maximal responses similar to those of ACh. As reported in Figure 1 and Table 1 for ACh, mutations increased the sensitivity for choline, as evidenced by a reduction in the EC_{50} . The rank order of potency of choline for mutated receptors was: 13'T > 9'T > 17'T ($P < 0.05$). Thus, the major shift in the EC_{50} for choline, 49-fold, was observed for the V13'T mutant.

Classical antagonists of wild-type receptors are agonists of mutant receptors

The wild-type $\alpha 9\alpha 10$ nAChR is blocked by the classical nicotinic receptor agonist, nicotine, as well as by the classical muscarinic receptor agonist, muscarine. Moreover, ICS 205, 930, a classical 5-HT₃ receptor antagonist, is one of the most potent blockers of $\alpha 9\alpha 10$ nAChRs (Elgoyhen *et al.*, 2001). This pharmacological profile is a hallmark of the peculiar $\alpha 9\alpha 10$ nAChR. As shown in Figure 4, these antagonists of wild-type receptors behaved as agonists of mutant 13'T and 9'T receptors. For all the three compounds, a higher efficacy was obtained in the case of the V13'T mutant receptor when compared to the L9'T (Table 2 and Figure 4). Moreover, while

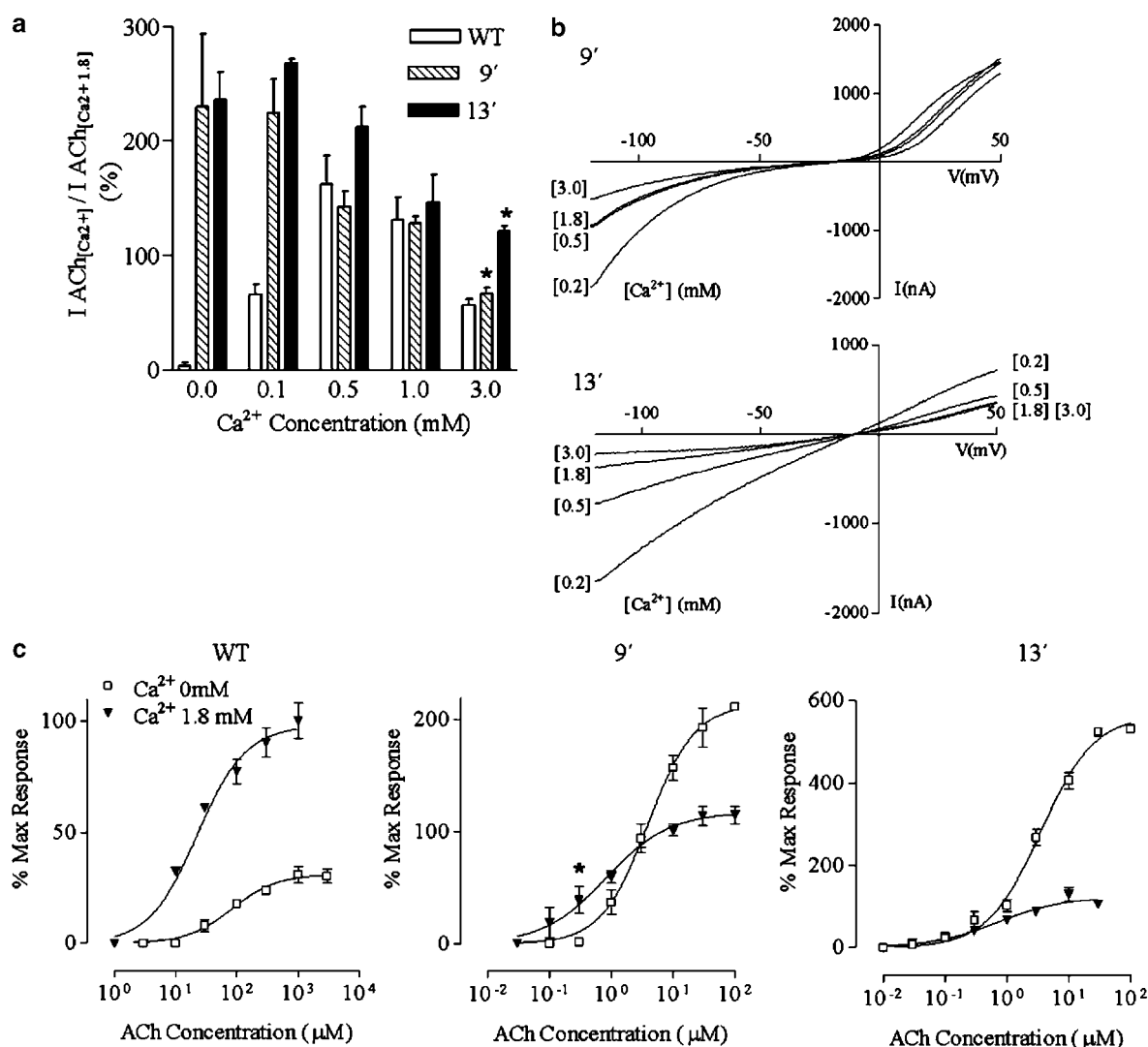


Figure 2 Modulation of ACh responses by extracellular calcium. (a) Bar diagram illustrating the effects of extracellular Ca^{2+} on responses to ACh in wild-type and mutant receptors at a membrane holding potential of -90 mV. The concentration of ACh used in each case was near the corresponding EC_{50} value derived from the concentration–response curves of Figure 1: 10μ M for the wt, 0.5μ M for the 9' and 0.1μ M for the 13' mutant. Current amplitudes obtained at different Ca^{2+} concentrations in each oocyte were normalized with respect to that obtained at 1.8 mM in the same oocyte. Each bar represents the mean and s.e.m. of the normalized response obtained in different oocytes ($n=4$ – 10 per bar). * $P<0.05$ with respect to the corresponding value at 0.1 mM Ca^{2+} . (b) Representative $I-V$ curves, obtained by application of a voltage ramp protocol (-120 to $+50$ mV, 2 s) 10 s after the peak response to either 0.5μ M ACh for the 9' (upper panel, $n=6$) or 0.1μ M ACh for the 13' mutant (lower panel, $n=6$). Oocytes were voltage-clamped at -70 mV, and ramps were performed at different Ca^{2+} concentrations. (c) Concentration–response curves to ACh, performed either at nominally zero or 1.8 mM Ca^{2+} . Responses were normalized to the maximum obtained at 1.8 mM Ca^{2+} for each case. The mean and s.e.m. of four to ten experiments per group are shown. * $P<0.05$ with respect to the corresponding value at nominally zero Ca^{2+} .

ICS 205,930 became a full agonist of V13'T with an EC_{50} of 8 nM, it behaved as a partial agonist of L9'T, achieving 50% of the maximal response to ACh with an EC_{50} of 40 nM. ICS 205,930, nicotine as well as muscarine also became agonists of receptors mutated at the 17' position. However, maximal responses were too small (4%, $n=8$, of the maximal response to ACh in the case of nicotine) and therefore full concentration–response curves could not be performed.

Mutant receptors exhibit spontaneous openings

Mutations to threonine at positions 13' and 9' of the $\alpha 7$ nAChR induce spontaneous openings of the receptor (Bertrand *et al.*,

1997; Corringer *et al.*, 1999). This is evidenced by a decrease in the leak current in the presence of the competitive antagonist methyllycaconitine. In the case of the $\alpha 9\alpha 10$ nAChR, classical antagonists that did not behave as agonists of mutant receptors as those described in Figure 4 did cause a deflection of the baseline current in the positive direction, that is, a reduction of the leak current (Figure 5). This was the case for compounds such as strychnine, D-tubocurarine, atropine, bicuculline and serotonin. No modification of the holding current was observed in noninjected oocytes or in oocytes injected with wild-type receptors. This result can be interpreted as a closure of receptors that are spontaneously opened in the absence of agonist (Bertrand *et al.*, 1997; Corringer *et al.*, 1999).

Figure 5a shows representative traces of responses of mutant 9'T and 13'T receptors to strychnine, including a maximal response to ACh in each case for comparison. The maximal response to strychnine reached 87% ($n=9$), 20% ($n=12$) and 0.6% ($n=5$) of the maximal current evoked by ACh in the 13'T, 9'T and 17'T mutants, respectively. Thus, a more pronounced phenotype was observed when mutating V13' to threonine. This was correlated with the fact that receptors injected with the 13'T mutant subunits exhibited unusually large holding currents when the membrane potential was voltage clamped at -70 mV (wt: 77 ± 5 nA, $n=55$; 9': 65 ± 6 nA, $n=55$; 13': 600 ± 57 nA, $n=55$, 17': 60 ± 2 nA, $n=40$).

The midpoint of the concentration–response curve, EC_{50} , is an empirical parameter that depends on the rate constants for ligand binding and unbinding, as well as those for channel opening and closing, that is, channel gating (Colquhoun, 1998). Thus, changes in the EC_{50} for ACh could derive from changes in the channel gating properties. This seems to be the case for the mutations in the TM2 region of the $\alpha 9\alpha 10$ nAChR, since the ACh EC_{50} values for mutant receptors were inversely correlated to the degree of spontaneous openings (gating) of the channels (Figure 5b).

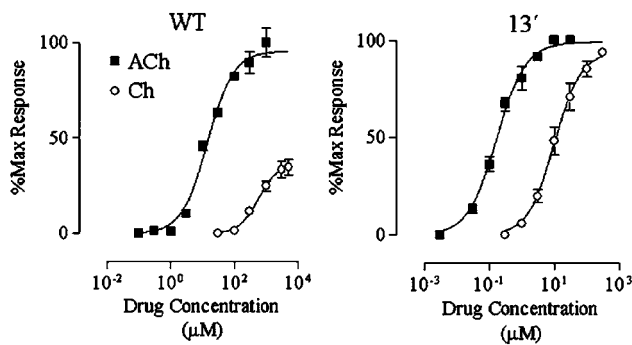


Figure 3 Choline is a full agonist of mutant receptors. Concentration–response curves to choline were performed. Peak current values were normalized and referred to the maximal peak response to ACh in each case. The mean and s.e.m. of four and seven experiments for wild-type and V13'T receptors, respectively, are shown. The EC_{50} and Hill coefficients are shown in Table 2.

Single-channel recordings reveal an increased apparent mean open time of mutant receptors

In order to characterize at the single-channel level the properties of the $\alpha 9\alpha 10$ wild-type and mutant receptors, we performed single-channel recordings in the cell-attached patch configuration in oocytes injected with wild-type, L9'T and V13'T $\alpha 9$ and $\alpha 10$ subunits.

To first determine the basal channel activity of the oocytes, we recorded channels from noninjected oocytes. Channel openings were detected in more than 90% of the patches (Figure 6a). Channel activity was similar to that corresponding to stretch-activated channels previously described (Taglietti & Toselli, 1988). At a pipette potential of $+120$ mV, at which the membrane potential is about -150 mV, the amplitude histogram showed a main component of 7.1 ± 1.2 pA ($n=3$). At positive potentials (70 to 120 mV), the I/V curve was linear. The conductance, calculated by the slope of the curve, was 76 pS. The open-time histogram could be well fitted with one component of 730 ± 170 μ s ($n=3$).

After characterizing the basal activity, we recorded single-channel currents under the same conditions, from oocytes injected with $\alpha 9$ and $\alpha 10$ wild-type and mutant subunits. The

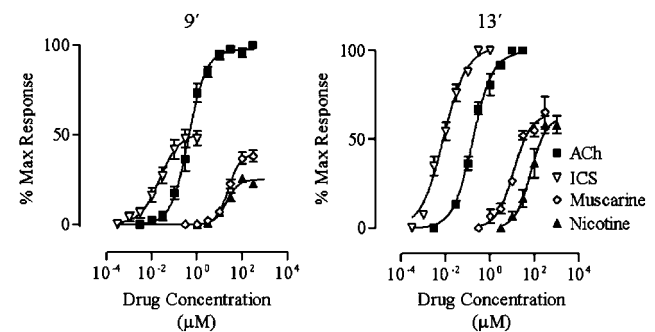


Figure 4 Effect of classical antagonists of the $\alpha 9\alpha 10$ nAChR. Concentration–response curves to ACh, ICS 205,930, muscarine and nicotine were performed. Peak current values were normalized and referred to the maximal peak response to ACh in each case. The mean and s.e.m. are shown. The number of experiments for each set of data is shown in Table 2. The EC_{50} , Hill coefficients and maximal responses are shown in Table 2.

Table 2 Responses of mutant receptors to different agonists

	WT	L9'T	V13'T	M/I17'T
Ch	$EC_{50} = 538 \pm 140$ μ M Max Resp = $36 \pm 3\%$ $n=4$	$EC_{50} = 20 \pm 2$ μ M Max Resp = $97 \pm 3\%$ $n=5$	$EC_{50} = 10 \pm 1$ μ M Max Resp = $98 \pm 2\%$ $n=7$	$EC_{50} = 27 \pm 3$ μ M Max Resp = $94 \pm 2\%$ $n=4$
ICS	$IC_{50} = 0.04 \pm 0.01$ μ M $n=4$	$EC_{50} = 0.04 \pm 0.01$ μ M Max Resp = $49 \pm 4\%$ $n=8$	$EC_{50} = 8.2 \pm 0.6$ nM Max Resp = $98 \pm 3\%$ $n=7$	
Nic	$IC_{50} = 4.6 \pm 1.0$ μ M $n=4$	$EC_{50} = 20 \pm 4$ μ M Max Resp = $25 \pm 2\%$ $n=6$	$EC_{50} = 70 \pm 14$ μ M Max Resp = $63 \pm 7\%$ $n=4$	
Musc	$IC_{50} = 40.9 \pm 5.1$ μ M $n=5$	$EC_{50} = 24 \pm 1$ μ M Max Resp = $39 \pm 3\%$ $n=7$	$EC_{50} = 12 \pm 3$ μ M Max Resp = $60 \pm 4\%$ $n=7$	

All parameters were determined as described in Methods. IC_{50} values for the wild-type receptor are included for comparison and have been extracted from Elgoyhen *et al.* (2001).

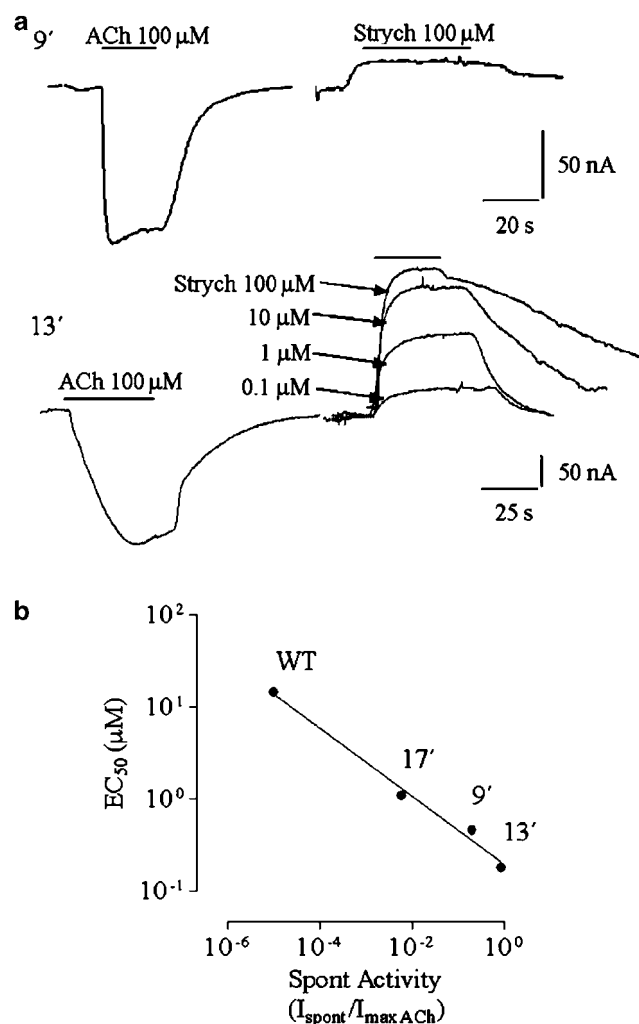


Figure 5 Block of leak current by strychnine. (a) Representative responses ($n = 5$ per mutant) to strychnine of oocytes injected with either the 9' (upper panel) or the 13' (lower panel) mutant receptors. Note the deflection of currents in the upward direction in the presence of the drug. A maximal response to ACh in each oocyte is shown for comparison. (b) Correlation of the EC_{50} values for ACh for each receptor with the degree of spontaneous activity calculated as the percentage of the maximal response to strychnine compared to that of ACh ($r^2: 0.989$).

concentration of ACh used was near the one that produces a maximal response for each receptor, derived from macroscopic currents. Single-channel recordings from oocytes expressing wild-type $\alpha 9\alpha 10$ nAChRs in the presence of 60 μ M ACh revealed a new population of channels that were neither observed in the absence of agonist nor in oocytes injected with the mutant subunits (Figure 6a). These channels were observed in only one of 76 seals. At a pipette potential of +120 mV, the mean amplitude of these channels was 18.4 pA, and thus could be well distinguished from the stretch-activated channels. As shown in the histograms, a single class of conductance was observed for both wild-type and mutant channels. The open-time histogram showed two components, a main one of 90 μ s (relative area 0.77) and a longer one of 320 μ s (relative area 0.24). The channel activity appeared in clusters of openings at 60 μ M ACh (Figure 6a).

In three of 33 recordings from oocytes injected with $\alpha 9$ and $\alpha 10$ subunits both carrying the L9'T mutation, we observed single channels activated by 1 μ M ACh, with a mean amplitude of 13.8 ± 2.1 pA ($n = 3$), similar to that determined in wild-type receptors (Figure 6a). The open-time histograms showed two components, which were both longer than those observed in wild-type receptors. Channel activity appeared in clusters. At 1 μ M ACh, the open components and relative areas were: 165 ± 30 μ s (0.63 ± 0.07) and 1.1 ± 0.4 ms (0.37 ± 0.07).

In three of 29 recordings performed in oocytes injected with $\alpha 9$ and $\alpha 10$ subunits both carrying the V13'T mutation, we observed single channels activated by 1 μ M ACh with a mean amplitude of 13.5 ± 2 pA ($n = 3$), similar to that observed in wild-type $\alpha 9\alpha 10$ receptors (Figure 6a). Channel activity was observed in very tight clusters. Clusters contained openings which were dramatically prolonged with respect to those of wild-type and 9'T mutants, as well as briefer closings. In two recordings, the open-time histograms were similar and were fitted with two components. The averages of the mean durations and relative areas for both recordings were 390 ± 160 μ s (0.38 ± 0.3), and 5.5 ± 0.5 ms (0.63 ± 0.3). In the third recording, in addition to the first two open components shown before, 205 μ s (0.4) and 3.7 ms (0.3), an additional significantly prolonged component was also observed (57.7 ms, relative area 0.3).

To further determine that the observed channels corresponded to $\alpha 9\alpha 10$ nAChRs, we performed recordings from outside-out patches before and after exposure to a 30-s pulse of ACh. We were able to detect channels in only two patches. One patch was obtained from oocytes injected with wild-type and the other with V13'T subunits. As shown in Figure 6b, the channels were indistinguishable from those recorded in the cell-attached patch configuration for each type of receptor (Figure 6a). Again, the V13'T channels were dramatically prolonged with respect to wild-type nAChRs. Each type of channel was not detected either before the application of ACh or in oocytes injected with other subunits. Thus, these results further support the fact that the channels observed in the cell-attached configuration correspond to $\alpha 9\alpha 10$ nAChRs.

Although the heterologous expression of some channels leads to a differential concentration in the animal pole of the oocyte (Grigoriev *et al.*, 1999), we did not find any difference in the number of positive seals when the patches were performed in either pole. Thus, assuming that receptors are homogeneously distributed in the oocyte surface, and considering an average single current of 10 pA, a macroscopic current of 600 nA at -70 mV, and a probability of opening of 0.5, the number of receptors per μ m² would be as low as 0.04. Owing to the low rate of success of finding channels other than the stretch-activated ones, we were not able to perform a thorough characterization of these channels in order to unequivocally determine their kinetic properties. However, the following lines of evidence allowed us to suggest that the shown channels should correspond to $\alpha 9\alpha 10$ nAChRs: (i) the channels were never observed in noninjected oocytes; (ii) the kinetics of the channels were different for wild-type, L9'T and V13'T mutants, and the changes were the ones expected from observations of the macroscopic currents; (iii) in two outside-out patches, channels were detected after application of ACh and showed the expected amplitudes and kinetics.

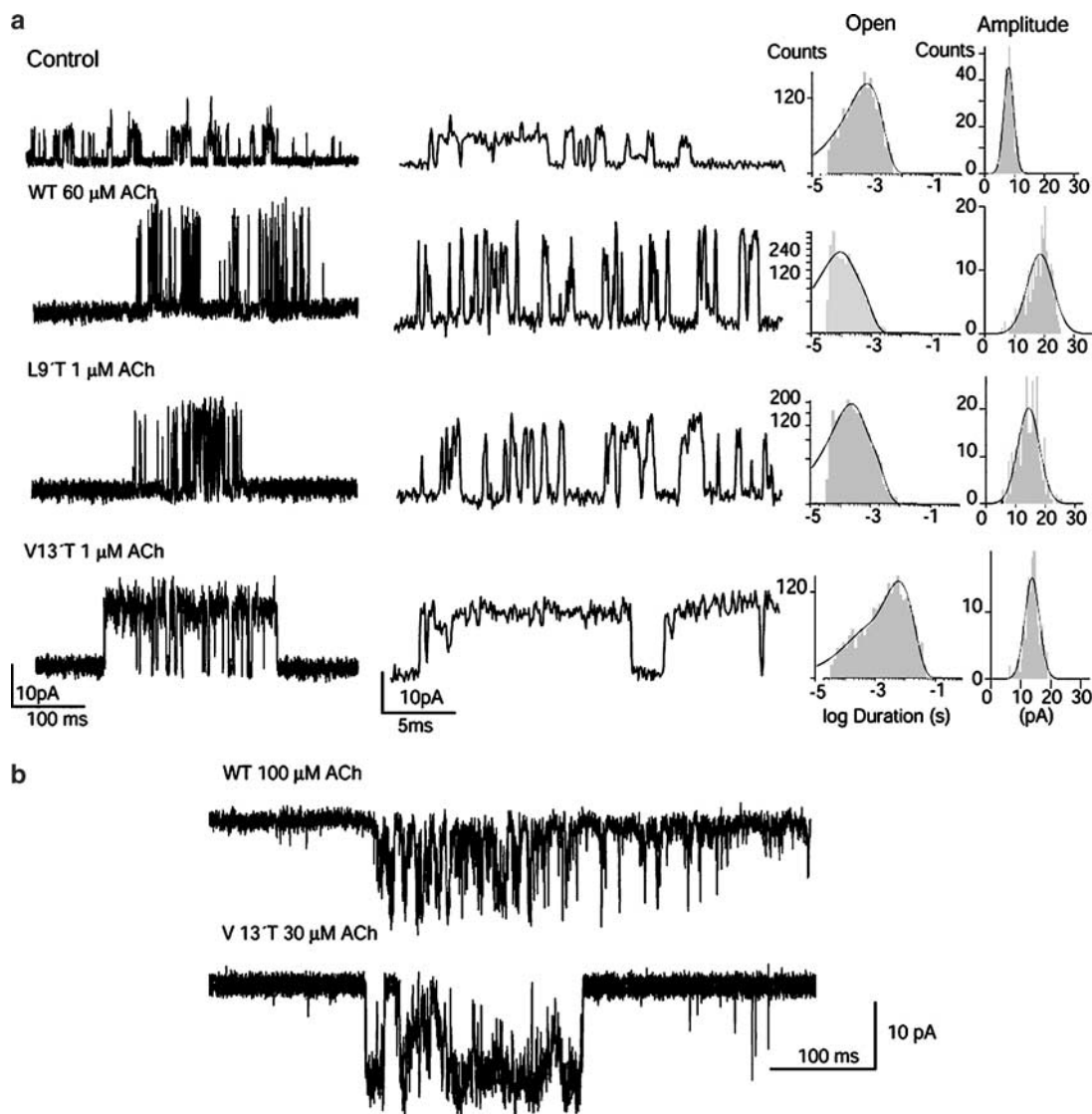


Figure 6 Single-channel recordings of wild-type and mutant $\alpha 9\alpha 10$ nAChRs. (a) (left) Channel traces recorded in the cell-attached configuration from oocytes injected with wild-type, L9'T and V13'T subunits. As a control, endogenous channels were recorded from noninjected oocytes. Traces are shown at two different time scales for each recording. Currents are displayed at a bandwidth of 5 kHz with channel openings as upward deflections. Pipette potential: 120 mV. To the right, open-time and amplitude histograms of the corresponding recordings are shown. (b) Channel traces obtained after application of ACh to outside-out patches from oocytes injected with wild-type or V13'T $\alpha 9\alpha 10$ subunits. Currents are displayed at a bandwidth of 5 kHz with channel openings as downward deflections. Pipette potential: -70 mV.

Discussion

Centrally located residues are involved in channel gating

The present results provide functional evidence suggesting that the centrally located positions 9' and 13' of TM2 are major contributors to channel gating, in the case of the $\alpha 9\alpha 10$ nAChR. Based on the atomic model described for the electric organ of the *Torpedo* ray (Miyazawa *et al.*, 2003), which suggests a tight hydrophobic girdle around the pore at positions 9' and 13', and on the fact that at these positions residues along the entire family are highly conserved, we can propose that the interactions of side chains at 13' and to a lesser extent at 9' are key ones in creating an energetic barrier to ion permeation. This conclusion is based on the observation that the magnitude of the phenotypes observed was more overt

in the case of the 13'T mutants and on the prediction that perturbation of the hydrophobic contacts by the introduction of a polar residue should increase the relative stability of the open pore (Miyazawa *et al.*, 2003). However, further mutations to residues other than threonine would be needed in order to prove this notion.

A major effect of the threonine substitution at 13', and to a lesser extent at 9', was to create constitutive opened receptors, to enhance ACh sensitivity, and at the single-channel level to increase the apparent mean open time and consequently to stabilize the open state of the mutant receptors, with no major changes in channel conductance. This is consistent with the hypothesis that the conserved centrally located TM2 residues are important for ACh receptor gating (Labarca *et al.*, 1995; Miyazawa *et al.*, 2003), and that the mutations weaken the contacts that hold the channel in the closed state.

In the case of wild-type $\alpha 9\alpha 10$ nAChRs, responses to ACh are tightly dependent upon the presence of external Ca^{2+} and responses at nominally zero Ca^{2+} are 5% of the maximal response obtained at 0.5 mM (Weisstaub *et al.*, 2002). It has been suggested that Ca^{2+} probably interacts at an extracellular binding site to allosterically modulate coupling between ACh binding and gating (Galzi *et al.*, 1996a; Weisstaub *et al.*, 2002). One could expect that a mutation that increases ACh receptor gating should eventually result in mutant receptors that are less dependent upon the presence of extracellular divalent cations to favor transitions to the open state. This is consistent with the experimental data obtained for mutations at the 9' and 13' TM2 positions shown in Figure 2.

A mutation that increases the ability of a receptor to change its conformation, that is an increase in gating, should enhance the efficacy of partial agonists (Colquhoun, 1998). This is consistent with the observation that the partial agonist choline became a full agonist of mutant receptors. On the other hand, weak partial agonists may actually behave as antagonists of wild-type receptors, since their ability for driving the receptor to the open state once bound is very low; they lead to very few channel openings and do not give rise to measurable macroscopic currents (Raya *et al.*, 2004). Increased channel gating, introduced by the mutations, would lead to measurable macroscopic currents. This could account for the observation that antagonists of wild-type receptors became agonists of mutant receptors.

Although one might expect a mutation in the pore of the channel to affect its gating properties but not its ligand-binding site, there is no reason that this should be necessarily true in an allosteric protein like an ion channel (Colquhoun, 1998). Propagated conformational changes to the extracellular N-terminal domain of the protein could eventually subserve some of the observed changes. However, the fact that at the single-channel level mutant receptors exhibited a drastic increase in the apparent mean open time, together with the observation that the ACh EC_{50} values for mutant receptors were inversely correlated to the degree of spontaneous openings of the channels, argues in favor of a direct effect on channel gating as the main underlying mechanism.

Single-channel recordings in $\alpha 9\alpha 10$ wild-type and mutant receptors

The single-channel recordings are the first ones to be reported for the $\alpha 9\alpha 10$ receptors. In spite of the fact that the number of patches in which we observed channels was extremely low, these channels were easily distinguished from the oocyte's endogenous channel activity due to their higher amplitude, which was enhanced by the use of solutions free of magnesium and low calcium to avoid channel block of $\alpha 9\alpha 10$ receptors (Weisstaub *et al.*, 2002). One possible explanation for the low rate of success in finding $\alpha 9\alpha 10$ channels is that the density of receptors is very low in oocytes. In this regard, the average peak current obtained for wild-type $\alpha 9\alpha 10$ nAChRs, 600 nA, is about ten-fold lower than that observed for the muscle nAChR, 9 μA (Labarca *et al.*, 1995), where successful single-channel recordings can be performed. A higher density of channels, with a very low probability of opening, could also explain the low rate of successful single-channel recordings.

Comparison to other Cys-loop receptors

Similar mutagenesis approaches have been undertaken for other receptors of the 'Cys-loop' family (Revah *et al.*, 1991; Filatov & White, 1995; Labarca *et al.*, 1995). These experiments have implicated the middle of the TM2 in the gating of the nAChR. Although *a priori* the phenotypes obtained in mutated $\alpha 9\alpha 10$ receptors might only appear as confirmatory of those already published, important differences when compared to those of other nAChRs were observed.

The present results differ from those reported for the neuronal $\alpha 7$ nAChRs, where the magnitude of the phenotypes assessed by the shift in ACh EC_{50} , and the activation by classical antagonists is more overt for the L9'T than the V13'T (Galzi *et al.*, 1996b). Moreover, in the case of $\alpha 7$ nAChRs, an additional conductance appears in the 9'T and 13'T mutants (Revah *et al.*, 1991; Galzi *et al.*, 1992). The fact that this was not observed in $\alpha 9\alpha 10$ indicates that, different from that proposed for the $\alpha 7$ receptor (Bertrand *et al.*, 1997), we do not need to invoke a conducting desensitized state to explain the observed results. However, we cannot disregard the fact that an additional conducting state might exist and could not be detected, either due to the low rate of success in finding channels or due to the fact that the new state has a low conductance that cannot be distinguished from that of the oocyte's endogenous stretch-activated channels. It is also possible that although amplitude histograms show a single-channel population, different populations may not be distinguishable due to closely spaced conductances and unequal relative areas. The latter may result in broad amplitude histograms, as shown in Figure 6.

Hydrophilic substitutions of the L9' of muscle nAChR are nearly independent, equivalent and multiplicative in their effects on the ACh EC_{50} value (Filatov & White, 1995; Labarca *et al.*, 1995). On the contrary, the EC_{50} values obtained for single and double mutants of the $\alpha 9\alpha 10$ nAChRs at 9' and 13' indicate that at these positions both types of subunits contribute in an asymmetric and nonadditive manner. This can be inferred from the fact that receptors containing only the $\alpha 9$ or the $\alpha 10$ mutant subunits showed the same decrease in the EC_{50} values, despite the fact that they cannot be in the same proportion in the pentameric assembly of a nAChR, and that a sole population of receptors is observed in oocytes with a $(\alpha 9)_2(\alpha 10)_3$ stoichiometry (Plazas & Elgoyhen, unpublished observations). Consequently, the EC_{50} values of $\alpha 9\alpha 10$ receptors containing all mutant subunits ($\alpha 9^* \alpha 10^*$, Table 1) are different from the ones calculated by multiplying the shifts in EC_{50} for receptors containing only $\alpha 9$ or $\alpha 10$ mutant subunits. Whether this reveals differences in the dynamics of the subunits during gating or results from an asymmetric orientation of the five subunits in the pore of the pentameric complex still has to be determined. However, following the atomic model of the pore, the five subunits of the receptor come into a close proximity at positions 9' and 13' and interaction of side chains from different subunits takes place (Miyazawa *et al.*, 2003). Thus, it is not surprising that a conformational change in one subunit can influence other subunits.

Different results were obtained at the more extracellular 17' position, where mutations to threonine in either $\alpha 9$ or $\alpha 10$ lead to 2.6- and 5.1-fold decreases in the EC_{50} values, respectively. The expected shift for the double-mutant receptor if both types

of subunits contribute independently to ACh sensitivity would be 13.3-fold. Interestingly, this agrees with the experimentally calculated decrease in the EC_{50} value (13.4-fold, Table 1). Therefore, it can be postulated that at this position both types of subunits contribute independently and symmetrically to ACh sensitivity, a result which could be explained if, as suggested in the atomic model (Miyazawa *et al.*, 2003), the 17' residues are not located at the girdle of the pore. Again, this differs from that described for the neighboring 16' residue of the muscle nAChR, where the contributions of the different subunits to gating are nonsymmetrical (Labarca *et al.*, 1995).

Conclusion

The present experimental data analyze for the first time the participation of different TM2 residues in gating of the $\alpha 9\alpha 10$ nAChR. We show evidence indicating that the centrally located amino acids at 9' and 13' are involved in activation

of the $\alpha 9\alpha 10$ nAChR. In general, our results show that although the roles of the TM2 residues are mostly conserved in the distant $\alpha 9\alpha 10$ member of the nAChR family, their mechanistic contributions to channel gating show significant differences. These differences might be originated from slight differential intramolecular rearrangements during gating for the different receptors and might lead each nAChR to be in tune with their physiological roles. Thus, results obtained from one type of receptor cannot be necessarily directly extrapolated to other receptors of the same family.

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