

## RAPID REPORT

# Pathogenic point mutations in a transmembrane domain of the $\epsilon$ subunit increase the $\text{Ca}^{2+}$ permeability of the human endplate ACh receptor

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The  $\epsilon$  subunit of the human endplate ACh receptor (AChR) is a key determinant of the large fraction of the ACh-evoked current carried by  $\text{Ca}^{2+}$  ions ( $P_f$ ). Consequently, missense mutations in the  $\epsilon$  subunit are potential targets for altering the  $P_f$  of human AChR. In this paper we investigate the effects of two pathogenic point mutations in the M2 transmembrane segment AChR  $\epsilon$  subunit,  $\epsilon$ T264P and  $\epsilon$ V259F, that cause slow-channel syndromes (SCS). When expressed in GH4C1 cells, the mutant receptors subunits raise  $\text{Ca}^{2+}$  permeability of the receptors  $\sim 1.5$  and  $\sim 2$ -fold above that of wild-type, to attain  $P_f$  values of 11.8% ( $\epsilon$ T264P) and 15.4% ( $\epsilon$ V259F). The latter value exceeds most  $P_f$  values reported to date for ligand-gated ion channels. Consistent with these findings, the biionic  $\text{Ca}^{2+}$  permeability ratio ( $P_{\text{Ca}}/P_{\text{Cs}}$ ) of the mutant AChRs is also increased. Upon repetitive stimulation with ACh, the mutant receptors show an enhanced current run-down compared with wild-type, leading to a strong reduction of their function. We propose that the enhanced  $\text{Ca}^{2+}$  permeability of the mutant receptors overrides the protective effect of desensitization and, together with the prolonged opening events of the AChR channel, is an important determinant of the excitotoxic endplate damage in the SCS.

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Slow-channel syndromes (SCS) are caused by point mutations in the human endplate acetylcholine receptor (AChR) that greatly prolong the openings of the AChR channel. In a peculiar form of the SCS, tubulofilamentous inclusions, similar to those found in inclusion body myositis (IBM), accumulate in junctional regions of the muscle fibres (Fidzianska *et al.* 2005). This SCS is caused by a valine-to-phenylalanine substitution at position 259 in the M2 transmembrane segment of the AChR  $\epsilon$  subunit ( $\epsilon$ V259F) that lines the channel pore.

Hyperphosphorylation of tau protein leads to formation of tubulofilamentous structures in neuronal diseases as well as in IBM, and is associated with, and possibly caused by, a disrupted  $\text{Ca}^{2+}$  homeostasis (LaFerla, 2002; Pierrot *et al.* 2004). We recently showed that the fractional  $\text{Ca}^{2+}$  current ( $P_f$ ) of human endplate AChR, defined as the percentage of the total ACh-evoked current carried by  $\text{Ca}^{2+}$  ions, is twice as large than that of the homologous mouse

AChR (Fucile *et al.* 2006). The high  $\text{Ca}^{2+}$  permeability of human AChR by itself enhances vulnerability of the human endplate to excitotoxic damage (endplate myopathy) in the SCS. An increased fractional  $\text{Ca}^{2+}$  current would further contribute to  $\text{Ca}^{2+}$  overloading of the postsynaptic region and to the  $\text{Ca}^{2+}$ -dependent pathological effects. Since the human  $\epsilon$  subunit turned out to be a key determinant of the high  $P_f$  at the human endplate (Fucile *et al.* 2006), we postulated that strategically positioned amino acid substitutions in the  $\epsilon$  subunit can alter the  $\text{Ca}^{2+}$  permeability of the AChR channel.

To test this notion, we measured the  $P_f$  of the  $\epsilon$ V259F-AChR and found it almost twice as high as that of wild-type (WT) AChR. To determine whether this effect is mutation specific, we also examined the  $P_f$  for another well-characterized SCS mutation in the M2 segment of the  $\epsilon$  subunit,  $\epsilon$ T264P (Ohno *et al.* 1995). We found that the  $P_f$  of the  $\epsilon$ T264P-AChR is also markedly increased compared

with that of the WT receptor. Our results provide the first direct evidence that mutations associated with SCS alter  $\text{Ca}^{2+}$  permeability as well as channel kinetics of the human endplate AChR.

## Methods

### Cell cultures and transfection

Rat pituitary GH4C1 cells were grown (5%  $\text{CO}_2$ , 37°C) in HAM F10 medium plus 10% fetal calf serum and 1% penicillin/streptomycin. Cells were plated onto 35 mm Petri dishes ( $3 \times 10^5$  cells per dish) 24 h prior to transfection. The cDNAs encoding human wild-type  $\alpha$ ,  $\beta$ ,  $\delta$ ,  $\epsilon$  subunits, or  $\epsilon$  subunits harbouring the  $\epsilon$ T264P or  $\epsilon$ V259F mutations were transiently transfected using Lipofectamine 2000, using 1  $\mu\text{g}$  of cDNA for each subunit per Petri dish. Medium was changed after overnight incubation with cDNAs, and experiments carried out after a further 24–48 h. All media were purchased from Invitrogen.

### Solutions and chemicals

Cells were superfused with a standard external medium containing (mM): 140 NaCl, 2.8 KCl, 2  $\text{CaCl}_2$ , 2  $\text{MgCl}_2$ ,

single-channel recordings) and an Axopatch 200B amplifier (Molecular Devices, Union City, CA, USA). Data were recorded and analysed using pCLAMP 9 (Molecular Devices). All recordings were performed at 25–27°C.

For whole-cell recordings, series resistance was compensated by 80–90%. Cells were voltage clamped at a holding potential of  $-60$  mV and continuously superfused using a gravity-driven fast exchanger perfusion system (RSC-200, BioLogique, France). Current decay and run-down were fitted with single exponential curves [ $i(t) = i_\infty + i_0 e^{-t/\tau}$ ], using Origin 7 (OriginLab Corporation, Northampton, MA, USA).

The relative  $\text{Ca}^{2+}$  permeability in biionic conditions ( $P_{\text{Ca}}/P_{\text{Cs}}$ ) was estimated from the shift of the reversal potential ( $V_{\text{rev}}$ ) of whole-cell ACh-induced current using two different extracellular  $\text{Ca}^{2+}$  concentrations ( $\text{Ca}_{01}$  of 1 mM,  $\text{Ca}_{010}$  of 10 mM). In each cell,  $V_{\text{rev}}$  was calculated at both  $\text{Ca}^{2+}$  concentrations ( $V_{r1}$  and  $V_{r10}$ , respectively), by linear interpolation of the ACh-evoked currents peak amplitude plotted *versus* the test potential. The use of voltage ramp yielded inaccurate results because of the rapid desensitization of  $\epsilon$ V259F-AChR.  $P_{\text{Ca}}/P_{\text{Cs}}$  ratios were obtained from the extended Lewis equation adapted to the chosen ion concentrations (Lewis, 1979; Castro & Albuquerque, 1995):

$$\frac{P_{\text{Ca}}}{P_{\text{Cs}}} = \frac{\text{Cs}_0(1 - e^{F\Delta V_r/RT})}{4\text{Ca}_{01} e^{F\Delta V_r/RT} (1 + e^{FV_{r1}/RT})^{-1} - 4\text{Ca}_{010}(1 + e^{FV_{r10}/RT})^{-1}} \quad (1)$$

10 Hepes-NaOH, 10 glucose, pH 7.3. For cell-attached experiments, patch pipettes were filled with the same solution plus ACh (200 nM).

For whole-cell recordings, patch pipettes were filled with a solution containing (mM): 120 KCl, 5 BAPTA, 10 Hepes-KOH, 2 Mg-ATP, 2  $\text{MgCl}_2$ , pH 7.3.

For  $P_f$  determinations, intracellular solution contained (mM): 140 *N*-methyl-D-glucamine (NMDG), 10 Hepes-HCl, 0.25 Fura-2 pentapotassium salt, 0.001 thapsigargin, pH 7.3. Calibration measurements were performed using an extracellular solution made of (mM): 153 NMDG, 10  $\text{CaCl}_2$ , 10 Hepes-HCl, pH 7.3.

The shift of the reversal potential of ACh-evoked current was measured using an intracellular solution containing (mM): 140 CsCl, 10 Hepes-CsOH, 20 BAPTA, pH 7.3. The external solution contained (mM): 150 CsCl, 10 Hepes-CsOH, 10 glucose, pH 7.3 plus 1 or 10  $\text{CaCl}_2$ .

All chemicals were purchased from Sigma (USA), except for Fura-2 pentapotassium salt (Molecular Probes).

### Electrophysiology

Currents were recorded using borosilicate glass patch pipettes (2–5  $\text{M}\Omega$  tip resistance, Sylgard-coated for

where  $F$ ,  $R$ ,  $T$  are the usual thermodynamic constants and  $V_r = V_{r10} - V_{r1}$ , Ion activity as given by Castro & Albuquerque (1995), was used in calculations. Membrane potential was stepped to the test potential 1 s before ACh application.

ACh dose–current curves were constructed applying to each cell different doses of transmitter (0.1–300  $\mu\text{M}$ , applied at 30–60 s interval) and normalizing the current response to the plateau value. The data were best-fitted using Hill equation by Origin 7 software (Origin Laboratory).

Single-channel data were filtered at 5 kHz and sampled at 25 kHz. Analysis was performed with a 50% threshold criterion, omitting events shorter than 0.12 ms. Slope conductance was calculated by linear fitting of the unitary amplitudes recorded at different pipette potentials (at least three for each cell). The fit was extrapolated to estimate cell resting potential, which was summed to the pipette potential to obtain membrane potential. The critical time used to identify a burst, determined for each cell from the closed time distribution (Colquhoun & Sakmann, 1985), ranged between 1 and 4 ms.

All results are given as mean  $\pm$  s.e.m. Two data sets were considered statistically different when  $P < 0.05$  (ANOVA test).

**Table 1. Single channel properties of WT and mutant AChRs**

	WT-AChR	εV259F-AChR	εT264P-AChR
Conductance (pS)	48.6 ± 2.4 (7)	49.7 ± 1.7 (7)	51.1 ± 2.6 (4)
Burst duration (ms)	1.88 ± 0.19 (5)	47 ± 11 (4)	42.4 ± 7.6 (4)
Critical time (ms)	3.4 ± 0.6 (5)	5 ± 1 (4)	1.8 ± 0.7 (4)
τ <sub>b1</sub> (ms)/weight (%)	0.59 ± 0.06/49 ± 5 (5)	0.25 ± 0.06/32 ± 3 (4)	0.6 ± 0.2/33 ± 4 (4)
τ <sub>b2</sub> (ms)/weight (%)	3.2 ± 0.3/51 ± 5 (5)	3.7 ± 2/18 ± 8 (4)	5.6 ± 2.8/17 ± 9 (4)
τ <sub>b3</sub> (ms)/weight (%)	not present	73 ± 10/50 ± 6 (4)	74 ± 14/50 ± 6 (4)

Data are given as mean ± s.e.m. (number of cells). Burst analysis was performed on cell-attached recordings obtained with bandwidth of 5 kHz;  $T = 25\text{--}27^\circ\text{C}$ , with an estimated membrane potential of  $-90$  to  $-105$  mV (as determined *a posteriori* by the  $i$ - $V$  relation). τ<sub>b1</sub>, τ<sub>b2</sub>, τ<sub>b3</sub>: time constants of the exponential components best fitting the distribution of burst durations.

### P<sub>f</sub> determination

The methods to measure  $P_f$  have been fully reported previously (Fucile *et al.* 2000, 2006). Fluorescence determinations were made using a fluorescence upright microscope (Axioskop 2, Zeiss, Germany), a digital 12-bit cooled camera (SensiCam, PCO, Germany) and a monochromator (Cairn, UK). The system was driven by Axon Imaging Workbench 2 software (Molecular Devices), which also triggered the start of electrophysiological recording. All optical parameters and the setting of the digital camera (50 ms exposure time,  $4 \times 4$  binning) were maintained throughout all measurements. The changes in intracellular calcium were monitored at a single excitation wavelength (380 nm), to achieve a higher time resolution, and expressed as the ratio of time-resolved fluorescence variation over the basal fluorescence ( $\Delta F/F_0$ ).

Only isolated cells with low basal intracellular Ca<sup>2+</sup> ( $F_{340}/F_{380}$  ratio values  $< 2$ ) were considered for recordings. Cells were filled with Fura-2 pentapotassium salt through the patch pipette and measurements performed after obtaining a stable value of basal fluorescence, with  $F_{380} > 200$  arbitrary units (a.u), indicating satisfactory loading. The ratio  $F/Q$  (expressed in  $\text{nC}^{-1}$ ) was defined as:

$$\frac{F}{Q} = \frac{\Delta F/F_0}{\int I_{\text{Nic}} dt}$$

where  $I_{\text{Nic}}$  is the nicotine-evoked whole-cell current. The charge entered into the cell was calculated at each fluorescence acquisition time. The  $F/Q$  ratio was estimated for each cell as the slope of the linear fit of the  $F$  versus  $Q$  plot.  $P_f$  was determined normalizing the  $F/Q$  ratio to the calibration value, obtained in a calibration medium containing only Ca<sup>2+</sup> as permeant ions:

$$P_f = (F/Q_{\text{standard}})/(F/Q_{\text{calibration}})$$

The  $F/Q_{\text{calibration}}$  value obtained in this work was  $6.6 \pm 1.9 \text{ nC}^{-1}$  (mean ± s.e.m.,  $n = 4$  cells).

### Results

#### Characterization of the εV259F and εT264P AChRs in GH4C1 cells

We first determined the main functional characteristics of wild-type (WT) and mutant AChRs expressed in GH4C1 cells. We confirmed the slow-channel properties of both εV259F- and εT264P-AChR at the single-channel level by cell-attached recordings (Fig. 1A). For the WT-AChR, the mean single channel slope conductance was about 50 pS and the mean burst duration was 1.9 ms (Table 1). As shown in Fig. 1B (left), the distribution of burst duration was fitted by two exponential components, with the mean time constants (τ<sub>b1</sub> and τ<sub>b2</sub>) and weights given in Table 1. The duration of the bursts was greatly enhanced by the εV259F as well as the εT264P mutation, while the conductance of unitary events was not significantly affected (Table 1). The distribution of burst durations presented a third exponential component (time constant, τ<sub>b3</sub>) for both mutant AChRs. Typical examples are shown in Fig. 1B (middle and right panels) and mean values of the τ<sub>b</sub>s reported in Table 1. These data confirm that both mutant receptors retain their main functional properties when expressed in GH4C1 cells.

At the macroscopic level, the amplitude of the ACh-evoked currents showed large cell-to-cell variation, ranging from  $-0.2$  nA to over  $-10$  nA at  $-60$  mV (ACh,  $100 \mu\text{M}$ ) for each type of AChR, suggesting that expression of the mutant AChRs was not appreciably impaired compared with that of WT-AChR. Both mutant receptors had a higher apparent affinity for ACh, as dose-response curves showed a decrease of the EC<sub>50</sub> from  $33.5 \pm 1.4 \mu\text{M}$  ( $n = 4$ ) in the WT-AChR, to  $1.2 \pm 0.2 \mu\text{M}$  ( $n = 4$ ) and  $0.9 \pm 0.2 \mu\text{M}$  ( $n = 5$ ) for εV259F-AChR and εT264P-AChR, respectively (data not shown).

During sustained ACh application, whole-cell currents decayed exponentially, reflecting receptor desensitization. At the plateau concentration of  $100 \mu\text{M}$ , the rate of decay was similar for the WT-AChR (τ<sub>decay</sub> =  $207 \pm 28$  ms,

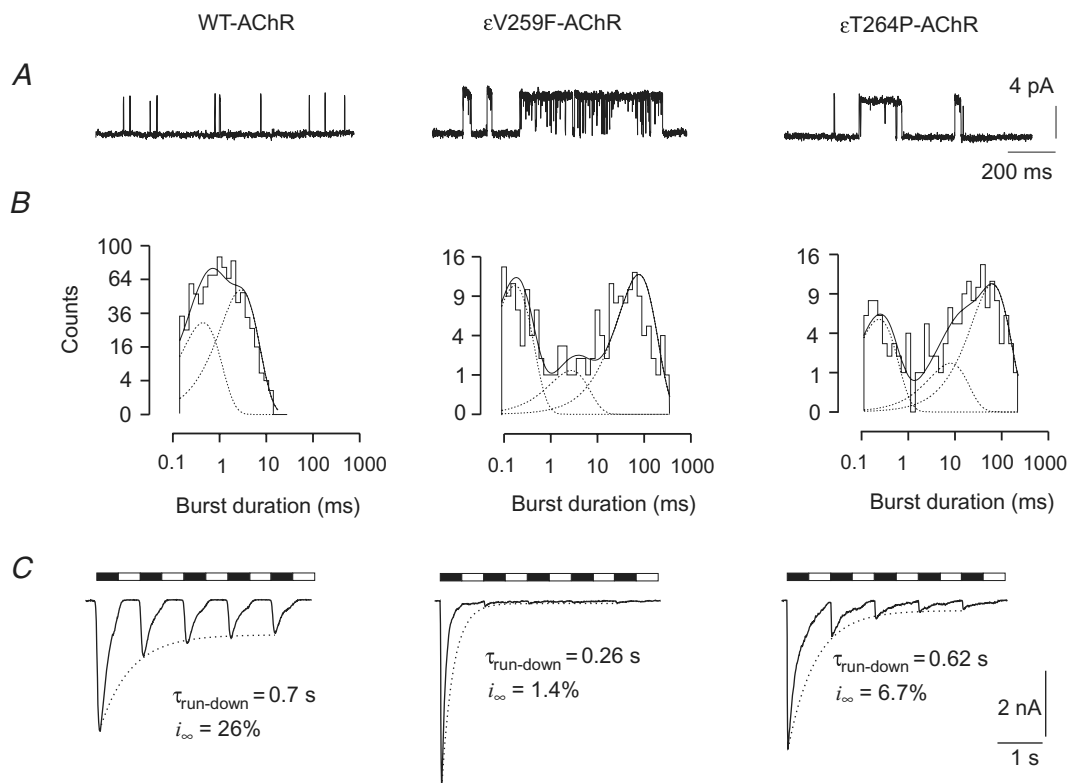
$n = 14$ ) and the  $\epsilon$ T264P-AChR ( $246 \pm 37$  ms,  $n = 7$ ;  $P = 0.4$ ), whereas the  $\epsilon$ V259F-AChR showed a faster decay ( $128 \pm 16$  ms,  $n = 10$ ;  $P = 0.04$ ). Repetitive ACh applications produced currents with decreasing peak amplitude (current run-down) for all three AChRs examined (Fig. 1C). An exponential fit of the consecutive current peaks showed that WT-AChR (Fig. 1C, left) had a relatively slow and limited run-down ( $\tau_{\text{run-down}} = 0.78 \pm 0.08$  s,  $n = 14$ ), with an asymptotic amplitude ( $i_{\infty}$ ) of  $31 \pm 4\%$  of the first response. For the  $\epsilon$ V259F mutant (Fig. 1C, middle), the decay was both faster ( $\tau_{\text{run-down}} = 0.38 \pm 0.06$  s,  $n = 9$ ,  $P = 0.002$ ) and more pronounced ( $i_{\infty} = 4.5 \pm 0.9\%$ ) than for WT-AChR. The  $\epsilon$ T264P mutant AChR (Fig. 1C, right) showed an intermediate behaviour, with a  $\tau_{\text{run-down}} = 0.68 \pm 0.05$  s ( $n = 7$ ) and an asymptote at  $i_{\infty} = 9 \pm 2\%$ , significantly less than that of WT-AChR ( $P = 0.001$ ). Current run-down was not accompanied by an increased rate of desensitization, as, in each cell, the  $\tau_{\text{decay}}$  of the third

current response was not significantly different from that of the first response ( $P > 0.5$ ). Together, these data indicate that the mutant receptors become less responsive than WT, or unresponsive, to ACh at physiological rates of stimulation.

### $P_f$ and $P_{Ca}/P_{Cs}$ measurements

For  $P_f$  determinations we used nicotine because ACh might activate muscarinic receptors, leading to  $Ca^{2+}$  release from internal stores. Currents evoked by nicotine ( $100 \mu\text{M}$ ) were comparable to those elicited by ACh, except for a slower current decay observed for the WT-AChR ( $\tau_{\text{decay}} = 649 \pm 87$  ms,  $n = 14$ ) (e.g. Fig. 2A).

Nicotine-evoked  $Ca^{2+}$  and current ( $I_{\text{Nic}}$ ) responses were simultaneously recorded (Fig. 2A), and the  $F/Q$  ratios calculated (Fig. 2B). In cells expressing WT-AChR (e.g. Fig. 2A, left), normalization to calibration values



**Figure 1. Functional characterization of WT and mutant AChRs in GH4C1 cells**

A, typical cell-attached recordings of channel openings evoked by ACh (200 nM) in cells transfected with WT or mutant AChRs, as indicated, revealing prolonged burst duration in  $\epsilon$ V259F and  $\epsilon$ T264P mutant AChRs. Traces were filtered at 1 kHz for display (5 kHz for analysis). Unitary conductance was: 49.8 pS (WT), 46.0 pS ( $\epsilon$ V259F) and 46.6 pS ( $\epsilon$ T264P). B, distribution of burst durations for the same recordings as in A, best fitted with the sum of two or three exponential components (dotted lines). Time constants (weight) of the fitting distributions in these three cells (at  $-100$  mV membrane potential) were: for WT-AChR,  $\tau_{b1} = 0.46$  ms (42%),  $\tau_{b2} = 2.9$  ms (58%), critical  $\tau = 4$  ms; for  $\epsilon$ V259F,  $\tau_{b1} = 0.17$  ms (41%),  $\tau_{b2} = 2.7$  ms (14%),  $\tau_{b3} = 78$  ms (45%), critical  $\tau = 4$  ms; for  $\epsilon$ T264P,  $\tau_{b1} = 0.24$  ms (34%),  $\tau_{b2} = 7.9$  ms (18%),  $\tau_{b3} = 63$  ms (48%), critical  $\tau = 1$  ms. C, typical whole-cell responses to repetitive ACh (100  $\mu\text{M}$  for 0.5 s) applications (black bars) in cells transfected as indicated. Dotted lines represent the exponential fit of the current peaks with the parameters given on each panel, showing the enhanced run-down of mutant AChRs. Holding potential,  $-60$  mV. Open bars indicate superfusion with standard solution.

yielded  $P_f$  values of  $7.83 \pm 0.69\%$  ( $n = 11$ ), in excellent agreement with previously published values (Fucile *et al.* 2006). The  $P_f$  value measured for  $\epsilon$ V259F-AChR (Fig. 2A, middle) was  $15.4 \pm 1.7\%$  ( $n = 12$ ,  $P = 0.0007$ ), showing that this mutation doubles the Ca<sup>2+</sup> permeability of the AChR-channel. For comparison, we measured the  $P_f$  of the  $\epsilon$ T264P-AChR (Fig. 2A, right) and again found a significantly enhanced Ca<sup>2+</sup> permeability ( $P_f = 11.76 \pm 0.91\%$ ,  $n = 13$ ,  $P = 0.003$ ). Thus, both slow-channel mutations in the M2 segment of the  $\epsilon$  subunit enhance the Ca<sup>2+</sup> permeability of human AChR.

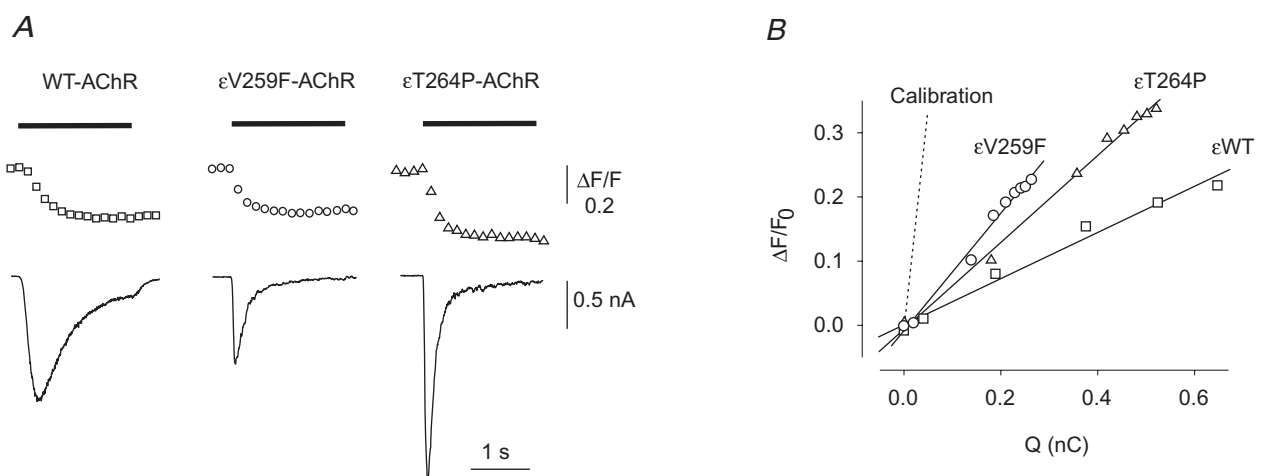
To test by a different approach that the Ca<sup>2+</sup> permeability of the mutant AChRs was enhanced compared with that of WT-AChR, we also assessed the relative Ca<sup>2+</sup> permeability of each isoform by measuring its biionic permeability ratio  $P_{Ca}/P_{Cs}$ . The currents evoked by ACh ( $10 \mu\text{M}$ ) were measured at five or more test potentials between  $-10$  and  $+10$  mV, bracketing the current reversal potential, at two different extracellular Ca<sup>2+</sup> concentrations ( $1 \text{ mM}$  and  $10 \text{ mM}$ ) (Fig. 3A). With  $1 \text{ mM}$  extracellular Ca<sup>2+</sup>, the average values of the reversal potentials were not statistically different for the  $\epsilon$ V259F-AChR ( $0.2 \pm 0.6 \text{ mV}$ ,  $n = 7$ ) or the  $\epsilon$ T264P-AChR ( $-0.5 \pm 0.3 \text{ mV}$ ,  $n = 10$ ) versus the WT-AChR ( $0.1 \pm 0.3 \text{ mV}$ ,  $n = 8$ ;  $P > 0.2$ ). The higher extracellular Ca<sup>2+</sup> concentration induced a right-shift of the reversal potential of each isoform (Fig. 3B). Using eqn (1) (see Methods), we obtained  $P_{Ca}/P_{Cs} = 0.73 \pm 0.07$  ( $n = 7$ ) for the WT-AChR. The value of  $P_{Ca}/P_{Cs}$  obtained for the  $\epsilon$ V259F-AChR ( $1.20 \pm 0.20$ ,  $n = 5$ ) was significantly larger than that of WT-AChR ( $P = 0.038$ ). A further increase over the WT-AChR value was observed

in the  $P_{Ca}/P_{Cs}$  ratio determined for the  $\epsilon$ T264P-AChR ( $1.35 \pm 0.10$ ,  $n = 7$ ,  $P = 0.0007$ ).

## Discussion

This paper provides the first quantitative evidence that the Ca<sup>2+</sup> permeability of the human endplate AChR can be modified by point mutations in the membrane-spanning M2 segment of the  $\epsilon$  subunit. The mutations examined,  $\epsilon$ V259F and  $\epsilon$ T264P, gave rise to a fractional Ca<sup>2+</sup> current of  $\sim 15\%$  and  $\sim 12\%$ , respectively. These  $P_f$  values are among the highest known, being equivalent to, or higher than, those reported for the NMDA receptor (up to  $11\%$ , Burnashev *et al.* 1995;  $13.5\%$ , Jatzke *et al.* 2002), or the homomeric  $\alpha 7$ -AChR ( $11.4\%$ , Fucile *et al.* 2003).

The enhanced Ca<sup>2+</sup> permeability of the  $\epsilon$  SCS mutants additionally increases the Ca<sup>2+</sup> ingress into the junctional sarcoplasm that stems from the prolonged duration of the synaptic current and the intrinsically high Ca<sup>2+</sup> permeability of human AChR and is thus directly relevant to the pathogenesis of the endplate myopathy associated with the observed mutations. An increased permeability to Ca<sup>2+</sup> may also occur with SCS mutations in other AChR subunits, as an increase of  $P_{Ca}/P_{Cs}$  was observed in HEK 293 cells expressing mouse  $\beta$ V299F-AChR, a mutant receptor designed after a recently identified mutation in a SCS patient (Navedo *et al.* 2006). Both  $\epsilon$ V259F and  $\epsilon$ T264P mutations yield a complex effect on receptor function. The affinity of the mutant receptors for ACh is increased by about one order of magnitude, but this increase is accompanied by (or possibly results in) a loss of receptor function upon repetitive stimulation at



**Figure 2. Calcium fractional current of WT and mutant AChRs**

A, typical fluorescence (top trace) and current (bottom trace) responses to nicotine ( $100 \mu\text{M}$ , horizontal bar) simultaneously recorded in cells expressing the indicated AChR. Notice the large Ca<sup>2+</sup> response recorded in the  $\epsilon$ V259 cell in spite of a relatively small current. B, linear fit of the  $\Delta F/F_0$  versus Q in the same three cells as in A. The dotted line represents the averaged slope of the calibration experiments. O,  $\epsilon$ V259F-AChR;  $\Delta$ ,  $\epsilon$ T264P-AChR;  $\square$ , WT-AChR, as indicated.

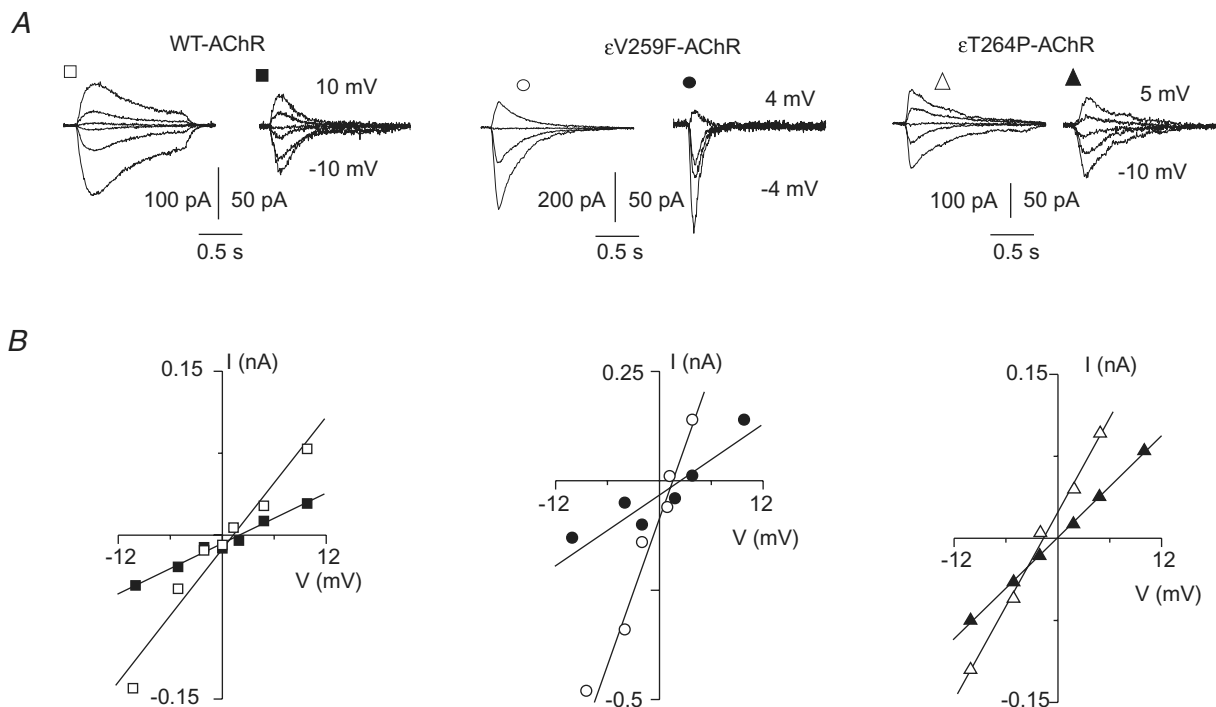
high ACh concentration, as well as an enhanced rate of desensitization in the case of the  $\epsilon$ V259F-AChR. That both mutant AChRs show an increased run-down during repetitive exposure to ACh probably contributes to the abnormal fatigability on exertion observed *in vivo*. Interestingly, the propensity of the mutant receptors to become unresponsive could partially mitigate the adverse effects of the prolonged open time and increased  $\text{Ca}^{2+}$  permeability of the mutant channels, but not enough to protect the endplate from excitotoxic injury.

In the only patient carrying the  $\epsilon$ V259F mutation identified to date (Fidzianska *et al.* 2005), the endplate myopathy was accompanied by the presence of tubulofilamentous inclusion bodies. The formation of these structures might be precipitated by the severe disruption of  $\text{Ca}^{2+}$  homeostasis caused by the concurrent prolongation of synaptic events and doubled  $P_f$ , consistent with the notion that an altered  $\text{Ca}^{2+}$  homeostasis enhances phosphorylation of tau protein and formation of fibrillary structures (LaFerla, 2002).

Several lines of evidence indicate that the  $\epsilon$  subunit is a key determinant of the  $\text{Ca}^{2+}$  permeability of adult muscle AChR. The  $P_f$  is almost 2-fold higher in AChRs containing

the  $\epsilon$  rather than the  $\gamma$  subunit (Villarroel & Sakmann, 1996; Ragozzino *et al.* 1998). Moreover, the  $\epsilon$  subunit is the only determinant of the high  $\text{Ca}^{2+}$  permeability of the human adult AChR (Fucile *et al.* 2006). That a valine-to-phenylalanine mutation in the pore-forming M2 segment of the  $\epsilon$  subunit ( $\epsilon$ V259F) markedly enhances  $P_f$  whereas mutation of the corresponding residue in the  $\alpha$  subunit ( $\alpha$ V249F) has no effect on  $P_f$  (Fucile *et al.* 2006), is also consistent with the important role of the  $\epsilon$  subunit in governing the  $\text{Ca}^{2+}$  permeability of the human receptor. Another mutation in the  $\epsilon$  M2 segment,  $\epsilon$ T264P, also increases the  $P_f$  of the mutant receptor. Thus, the data presented here confirm that the  $\epsilon$  subunit has a strong effect on the  $\text{Ca}^{2+}$  permeability of human endplate AChR.

In our experiments, the alterations in  $P_{\text{Ca}}/P_{\text{Cs}}$  values only partially mirrored those in the  $P_f$ . The rapid run-down of current generated by  $\epsilon$ V259F-AChR during multiple applications of ACh limits the accuracy of the  $P_{\text{Ca}}/P_{\text{Cs}}$  measurements. Furthermore, constant field assumptions leading to eqn (1) may not hold under the actual experimental conditions, as previously reported (Vernino *et al.* 1994; Burnashev *et al.* 1995). For instance, studies that compare muscle and neuronal AChRs reveal



**Figure 3. Relative  $\text{Ca}^{2+}$  permeability ( $P_{\text{Ca}}/P_{\text{Cs}}$ ) of WT and mutant AChRs**

*A*, ensemble whole-cell currents recorded in response to ACh ( $10 \mu\text{M}$ ) applied at various test potentials with low extracellular  $\text{Ca}^{2+}$  ( $1 \text{ mM}$ , open symbols) or high extracellular  $\text{Ca}^{2+}$  ( $10 \text{ mM}$ , filled symbols), in transfected cells expressing the indicated AChRs. In each panel, numbers refer to the most positive and negative test potentials shown. Note the reduction of current amplitude in high  $\text{Ca}^{2+}$  for all the AChR isoforms. *B*, current peak amplitude plotted versus membrane test potential in the same cells as in *A*. Lines represent the linear fit of each data group. Note positive shifts of current reversal potentials ( $0.7 \text{ mV}$  for WT-AChR;  $1.5 \text{ mV}$  for  $\epsilon$ V259F-AChR;  $1.6 \text{ mV}$  for  $\epsilon$ T264P-AChR) at  $10 \text{ mM}$   $\text{Ca}^{2+}$  (filled symbols).

a 7-fold difference in the  $P_{Ca}/P_{Cs}$  ratio (Vernino *et al.* 1994), but only a 2-fold difference for the  $P_f$  values (Vernino *et al.* 1993). Thus, biionic permeability ratios often provide qualitative rather than quantitative estimates of the relative Ca<sup>2+</sup> permeability of a given channel whereas  $P_f$  determinations require no *a priori* assumptions (Zhou & Neher, 1994) and thus yield a more reliable measure of the Ca<sup>2+</sup> permeability of the AChR channel (Burnashev *et al.* 1995).

In conclusion, our studies provide further insight into the pathogenic mechanisms whereby single amino acid substitutions in AChR subunits cause disease, which results from the delicate balance between the deleterious effects of the increased Ca<sup>2+</sup> permeability and the dampening of the synaptic response by run-down of the investigated SCS mutant receptors at physiological rates of stimulation, by loss of AChR from the degenerating folds, and by the altered endplate geometry.

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