

## Facilitation of rabbit $\alpha_{1B}$ calcium channels: involvement of endogenous $G\beta\gamma$ subunits

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1. The  $\alpha_{1B}$  (N-type) calcium channel shows strong G protein modulation in the presence of G protein activators or  $G\beta\gamma$  subunits. Using transient expression in COS-7 cells of  $\alpha_{1B}$  together with the accessory subunits  $\alpha_2$ - $\delta$  and  $\beta_{2a}$ , we have examined the role of endogenous  $G\beta\gamma$  subunits in the tonic modulation of  $\alpha_{1B}$ , and compared this with modulation by exogenously expressed  $G\beta\gamma$  subunits.
2. Prepulse facilitation of control  $\alpha_{1B}/\alpha_2$ - $\delta/\beta_{2a}$  currents was always observed. This suggests the existence of tonic modulation of  $\alpha_{1B}$  subunits. To determine whether endogenous  $G\beta\gamma$  is involved in the facilitation observed in control conditions, the  $\beta$ ARK1  $G\beta\gamma$ -binding domain (amino acids 495–689) was overexpressed, in order to bind free  $G\beta\gamma$  subunits. The extent of control prepulse-induced facilitation was significantly reduced, both in terms of current amplitude and the rate of current activation. In agreement with this, GDP $\beta$ S also reduced the control facilitation.
3. Co-expression of the  $G\beta_1\gamma_2$  subunit, together with the  $\alpha_{1B}/\alpha_2$ - $\delta/\beta_{2a}$  calcium channel combination, resulted in a marked degree of depolarizing prepulse-reversible inhibition of the whole-cell  $I_{Ca}$  or  $I_{Ba}$ . Both slowing of current activation and inhibition of the maximum current amplitude were observed, accompanied by a depolarizing shift in the mid-point of the voltage dependence of activation. Activation of endogenous  $G\beta\gamma$  subunits by dialysis with GTP $\gamma$ S produced a smaller degree of prepulse-reversible inhibition.
4. The rate of reinhibition of  $\alpha_{1B}$  currents by activated G protein, following a depolarizing prepulse, was much faster with  $G\beta_1\gamma_2$  than for the decay of facilitation in control cells. Furthermore,  $\beta$ ARK1 (495–689) co-expression markedly slowed the control rate of reinhibition, suggesting that the kinetics of reinhibition depend on the concentration of free endogenous or exogenously expressed  $G\beta\gamma$  in the cells. In contrast, the rate of loss of inhibition during a depolarizing prepulse did not vary significantly between the different conditions examined.
5. These findings indicate that, in this system, the voltage-dependent facilitation of  $\alpha_{1B}$  that is observed under control conditions occurs as a result of endogenous free  $G\beta\gamma$  binding to  $\alpha_{1B}$ .

Voltage-dependent G protein-mediated inhibition of neuronal  $Ca^{2+}$  channels is characterized by an inhibition of current amplitude and a slowing of the activation kinetics (for review, see Dolphin, 1996). G protein-mediated inhibition of neuronal N-type ( $\alpha_{1B}$ ) and P/Q-type ( $\alpha_{1A}$ ) calcium currents is mediated by  $G\beta\gamma$ , and not  $G\alpha$ , subunits (Herlitze, Garcia, Mackie, Hille, Scheuer & Catterall, 1996; Ikeda, 1996). The main binding sites for  $G\beta\gamma$  subunits on  $\alpha_{1A}$  and  $\alpha_{1B}$  have been localized to two domains on the intracellular loop that links transmembrane domains I and II (De Waard, Liu, Walker, Scott, Gurnett & Campbell, 1997; Zamponi, Bourinet, Nelson, Nargeot & Snutch, 1997). Although a binding site for  $G\beta\gamma$  has also been found on the C-terminal tail of human  $\alpha_{1E}$  (Qin, Platano, Olcese, Stefani &

Birnbaumer, 1997), this was not observed for  $\alpha_{1B}$  (Zamponi *et al.* 1997).

G protein modulation is most pronounced at potentials just supra-threshold for activation; this has been interpreted as G protein-modulated channels requiring stronger depolarizations to open (Bean, 1989). Thus, inhibition is weaker at more depolarized potentials. Early models involved binding of activated G protein to both closed and open states of the channel (Elmslie & Jones, 1994; Boland & Bean, 1993). More recently, it has been suggested that these phenomena are due to binding of activated G protein to only the closed state of the channel (Patil, de Leon, Reed, Dubel, Snutch & Yue, 1996). The finding that G protein modulation is voltage dependent provided the basis for the use of large

depolarizing prepulses to reverse G protein-mediated inhibition (Elmslie & Jones, 1994). Enhancement of calcium currents in response to prior depolarization is termed facilitation, but it does not necessarily involve reversal of G protein modulation (for review, see Dolphin, 1996). It has also been observed for both cloned and native cardiac calcium channels (Sculptoreanu, Rotman, Takahashi, Scheuer & Catterall, 1993a; Sculptoreanu, Scheuer & Catterall, 1993b; Bourinet, Charnet, Tomlinson, Stea, Snutch & Nargeot, 1994), which do not show direct modulation by inhibitory G proteins.

In the present study, we have examined the G protein modulation of  $\alpha_{1B}$  calcium channels in the presence of  $\beta_{2a}$ , a subunit that dramatically slows inactivation, in order to examine changes in the activation properties in isolation. Under these conditions, we observed facilitation of  $\alpha_{1B}$  under control conditions, both in terms of an increase in activation kinetics following a depolarizing prepulse and an increase in current amplitude. We have therefore compared the effects of co-expressing exogenous  $G\beta_1\gamma_2$  and of activating endogenous G proteins with GTP $\gamma$ S, and conversely of either reducing endogenous G protein activation with GDP $\beta$ S or reducing free endogenous  $G\beta\gamma$  concentrations by exogenous expression of  $\beta$ ARK1 (495–689), which contains only the C-terminal  $G\beta\gamma$ -binding domain of  $\beta$ ARK (Koch, Inglese, Stone & Lefkowitz, 1993; Pitcher, Touhara, Payne & Lefkowitz, 1995). Expression of  $\beta$ ARK1 (495–689) has been shown previously to block several  $G\beta\gamma$ -mediated signalling pathways in COS-7 cells (Koch, Hawes, Inglese, Luttrell & Lefkowitz, 1994). We now show that the observed control facilitation of  $\alpha_{1B}$  is reduced by expression of exogenous  $\beta$ ARK1 (495–689), and is thus likely to be due to endogenous free  $G\beta\gamma$  subunits.

## METHODS

### Materials

The following cDNAs were used: rabbit  $\alpha_{1B}$  (accession number, D14157; Fujita *et al.* 1993) was provided by Dr Y. Mori (Seriken, Okazaki, Japan); rat  $\beta_{2a}$  (M80545; Perez-Reyes *et al.* 1992) was provided by Dr E. Perez-Reyes (Loyola University, Chicago, IL, USA);  $\beta_{1b}$  (X11394) was provided by Dr T. Snutch (UBC, Vancouver, Canada); the full-length rat  $\alpha_2$ - $\delta$  (neuronal splice variant, M86621; Kim, Kim, Lee, King & Chin, 1992) was provided by Dr H. Chin (National Institutes of Health, Bethesda, MA, USA). Bovine  $G\beta_1$  (M13236; Fong *et al.* 1986) and bovine  $G\gamma_2$  (M37183; Gautam, Baetscher, Aebersold & Simon, 1989) were provided by Dr M. Simon (CalTech, Pasadena, CA, USA). The C-terminal minigene of  $\beta$ ARK1 (*BTARKB*) was provided by Dr R. Lefkowitz (Duke University, Durham, NC, USA). The S65T mutant of green fluorescent protein (GFP) was a gift from Dr S. Moss (University College London, UK) and the mut-3 GFP mutant was a gift from Dr T. Hughes (Yale University, New Haven, CT, USA). All cDNAs were subcloned, using standard techniques, into the pMT2 expression vector (Genetics Institute, Cambridge, MA, USA) for transient expression in COS-7 cells, except that encoding  $\beta$ ARK1 (495–689), which was expressed from pRK5 (Koch *et al.* 1994).

### Transfection of COS-7 cells

COS-7 cells were cultured and transfected, using electroporation, essentially as described previously (Campbell, Berrow, Brickley, Page, Wade & Dolphin, 1995a). For transfection, 15, 5, 5 and 1  $\mu$ g of  $\alpha_1$ ,  $\alpha_2$ - $\delta$ ,  $\beta_{2a}$  (or  $\beta_{1b}$ ), and either S65T or mut-3 GFP constructs were used. When used, 2.5  $\mu$ g of both  $G\beta_1$  and  $G\gamma_2$  cDNA or 5  $\mu$ g  $\beta$ ARK1 minigene (*BTARKB*) were included. Cells were maintained at 37 °C, then replated using a non-enzymatic cell dissociation medium (Sigma), and kept at 25 °C prior to electrophysiological recording. Successfully transfected cells were identified by expression of GFP. Maximum GFP fluorescence and Ca<sup>2+</sup> channel expression were observed between 2 and 4 days post-transfection (Brice *et al.* 1997; Berrow, Brice, Tedder, Page & Dolphin, 1997; Stephens, Page, Burley, Berrow & Dolphin, 1997).

### Antibodies

Polyclonal antisera to the calcium channel subunits  $\alpha_{1B}$  and  $\beta$  were raised in rabbits using standard methodology. The calcium channel  $\beta_{common}$  antiserum was raised against a peptide corresponding to amino acids 65–79 of the rat brain  $\beta_{1b}$  sequence (SRPSDSDVSLIEDRE), as previously described (Berrow, Campbell, Fitzgerald, Brickley & Dolphin, 1995). The  $\alpha_{1B}$  antiserum was raised against a peptide corresponding to amino acids 851–867 (RHHRHRDRKDKTSASTPA) of the rat brain  $\alpha_{1B}$  sequence (rbB-1; Dubel, Starr, Ahlijanian, Enyeart, Catterall & Snutch, 1992). This epitope forms part of the intracellular loop between transmembrane segments S6 of domain II and S1 of domain III, and is 65% homologous to the corresponding region of the rabbit brain  $\alpha_{1B}$  sequence used for this study (BILL, Fujita *et al.* 1993). Cross-reactivity of this antiserum with the rabbit  $\alpha_{1B}$  was confirmed by expression and immunostaining of the rabbit  $\alpha_{1B}$  subunit in COS-7 cells.

### Immunocytochemistry

COS-7 cells used for immunocytochemistry were transfected as for electrophysiological experiments, but omitting the GFP. Immunocytochemistry was performed essentially as described previously (Brice *et al.* 1997). Briefly, the cells were fixed in 4% paraformaldehyde 2–3 days post-transfection and permeabilized with 0.02% Triton X-100. The cells were then incubated overnight with  $\alpha_{1B}$  antiserum (1:500 dilution), calcium channel  $\beta_{common}$  antiserum (1:500 dilution), or  $G\beta_{common}$  antibody (Santa Cruz, California, USA; 250  $\mu$ g ml<sup>-1</sup>, used at 1:500 dilution).  $\beta_{1b}$  was used in this study rather than  $\beta_{2a}$ , as the antibody, although raised against a peptide common to all  $\beta$  subunits, recognizes  $\beta_{1b}$  better than  $\beta_{2a}$  (Brice *et al.* 1997). Bound antibodies were detected using biotin-conjugated goat anti-rabbit IgG antibody and streptavidin-conjugated Texas Red (Molecular Probes). Immunostaining was observed using an MRC 600 confocal microscope (Biorad, Hemel Hempstead, UK). Arbitrary optical density units were calculated using the Kontron KS400 program (Kontron Elektronik, Eshing, Germany).

### Electrophysiology

Recordings were made at room temperature (20–22 °C) from COS-7 cells which had been replated between 1 and 16 h previously. Cells were viewed briefly using a fluorescein filter block, and only fluorescent cells expressing GFP were used in experiments. The internal (pipette) and external solutions and recording techniques are similar to those described previously (Campbell, Berrow, Fitzgerald, Brickley & Dolphin, 1995b). The patch pipette solution contained (mM): caesium aspartate, 140; EGTA, 5; MgCl<sub>2</sub>, 2; CaCl<sub>2</sub>, 0.1; K<sub>2</sub>ATP, 2; GTP, 0.1 (when used); and Hepes, 10; pH 7.2, 310 mosmol l<sup>-1</sup> with sucrose. GTP $\gamma$ S (100  $\mu$ M) or GDP $\beta$ S

(0.5–2 mM) were included, where stated, in place of GTP. The external solution contained (mM): TEA-Br, 160; KCl, 3; NaHCO<sub>3</sub>, 1.0; MgCl<sub>2</sub>, 1.0; Hepes, 10; glucose, 4; and BaCl<sub>2</sub> or CaCl<sub>2</sub>, 1; pH 7.4, 320 mosmol l<sup>-1</sup> with sucrose. Pipettes of resistance 2–4 M $\Omega$  were used. Whole-cell currents were elicited from holding potentials ( $V_h$ ) of -100 mV. Cells used had a capacitance of  $31.9 \pm 4.1$  pF ( $n = 10$ ),  $27.1 \pm 1.8$  pF ( $n = 15$ ) and  $37.3 \pm 2.1$  pF ( $n = 10$ ) for control,  $G\beta_1\gamma_2$  cDNA-transfected and  $\beta$ ARK1 (495–689) cDNA-transfected cells, respectively. Cells were only used where series resistance was compensated to 80%, and space clamp was adequate as judged by graded activation of  $I_{Ca}$  or  $I_{Ba}$ . The voltage errors from the residual uncompensated series resistance were less than 1 mV for the largest currents, and no further correction was made. An Axopatch-1D amplifier (Axon Instruments) was used, and data were filtered at 0.5–5 kHz and digitized at 1–10 kHz. Analysis was performed using pCLAMP 6 (Axon Instruments) and Origin 3.5 (Microcal Software, Northampton, MA, USA). The junction potential between external and internal solutions was 6 mV; the values given in the figures and text have not been corrected for this. Current records are shown following leak and residual capacitance current subtraction ( $P/4$  or  $P/8$  protocol). Data are expressed as means  $\pm$  s.e.m. Statistical analysis was performed using Student's paired or unpaired  $t$  tests, as appropriate.

## RESULTS

### Biophysical properties of $\alpha_{1B}$ subunits co-transfected with $\beta_{2a}$ and $\alpha_2$ - $\delta$

Transient transfection of  $\alpha_{1B}$  cDNA together with  $\beta_{2a}$  and  $\alpha_2$ - $\delta$  cDNAs resulted in inward  $I_{Ca}$  with a similar current density to that which we have previously described for  $I_{Ba}$ . In most experiments described here, Ca<sup>2+</sup> was used as the charge carrier, but a number of the experiments were performed with Ba<sup>2+</sup> (where stated), and no major differences were observed. A minor difference was that the mid-point of voltage dependence of activation ( $V_{1/2}$ ) (see legend to Fig. 1) was  $-10.5 \pm 1.8$  mV ( $n = 7$ ) in 1 mM Ba<sup>2+</sup>, whereas it was  $-6.6 \pm 2.1$  mV ( $n = 11$ ) in 1 mM Ca<sup>2+</sup>, as expected for differences in charge screening. The  $\beta_{2a}$  subunit acts to retard entry of channels into the inactivated state (Olcese *et al.* 1994), and was used here to limit any confounding effects of voltage-dependent inactivation on the observation of G protein modulation (see Jones & Elmslie, 1997, for discussion of this problem). Co-expression of  $\beta_{2a}$  dramatically slowed the voltage-dependent inactivation of  $\alpha_{1B}$ , compared with that previously observed when these subunits were co-expressed with  $\beta_{1b}$  subunits (Page, Stephens, Berrow & Dolphin, 1997). The inactivation kinetics of  $\alpha_{1B}/\alpha_2$ - $\delta/\beta_{2a}$   $I_{Ba}$  at -10 mV, measured using 8 s voltage steps, could usually be described by a single exponential, with a time constant of  $4980 \pm 800$  ms ( $n = 4$ ), together with a plateau phase representing approximately 30% of the current.

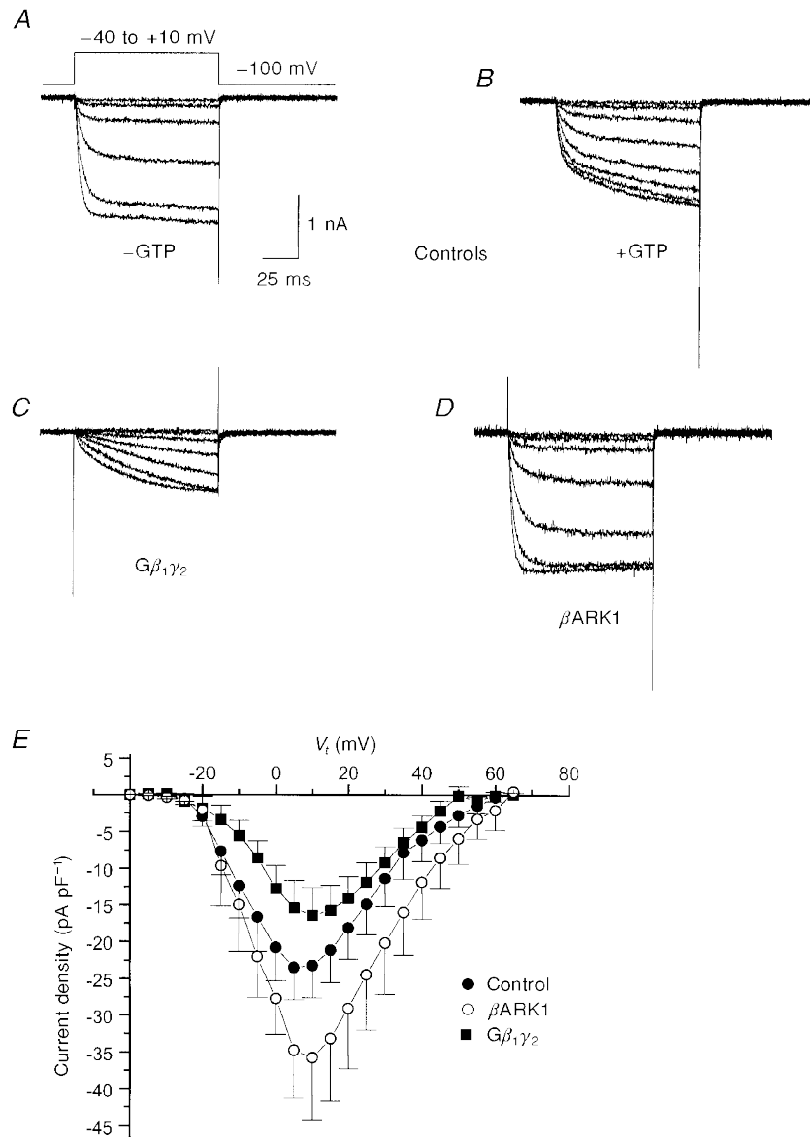
Control  $I_{Ca}$  recorded in an intracellular solution without added GTP was normally rapidly activating (Fig. 1A). A few examples of currents with a clearly distinguishable additional slowly activating component were also observed (3/13 cells), and this was also observed when  $I_{Ca}$  was recorded with GTP in the intracellular medium (2/4 cells) (Fig. 1B). In contrast,

in the presence of co-transfected  $G\beta_1\gamma_2$  subunits, the inward currents usually consisted of a slowly activating current only (Fig. 1C, representing 12/14 cells, with 2/14 also having a rapidly activating component). To determine whether the slowed activation observed in some control cells was a result of the effect of endogenous free  $G\beta\gamma$  subunits, currents were examined in the presence of co-expressed  $\beta$ ARK1 (495–689) to bind free  $G\beta\gamma$  (Koch *et al.* 1993), and in the absence of GTP. Under these conditions, currents were always rapidly activating ( $n = 15$ , Fig. 1D). From current–voltage relationships, current density was increased by  $\beta$ ARK1 (495–689), compared with control currents (Fig. 1E), whereas  $G\beta\gamma$  co-expression reduced current density and shifted the mid-point of activation to more depolarized potentials compared with the control (Fig. 1E).

### Facilitation of $\alpha_{1B}$ following a large depolarizing prepulse

Current facilitation was examined by comparing current amplitudes and activation rates during a 50 ms test pulse before (P1) and 10 ms after (P2) application of depolarizing prepulses to +120 mV (Fig. 2, see left panel for protocol). The depolarizing prepulse produced a small but significant effect on control  $\alpha_{1B}$   $I_{Ca}$  current density (Fig. 2A, middle) and activation kinetics (Fig. 2A, right). The facilitation ratio of the current amplitudes was also calculated as P2/P1 at 11 ms after the start of the step (Fig. 3). This was designed to measure only rapidly activating current, as an estimate of current that was not modulated at -100 mV. Control facilitation ratios were greater than 1 particularly at the voltages that are just supra-threshold for activation (Fig. 3A,  $\square$ ; e.g. P2/P1 =  $2.2 \pm 0.4$  at -20 mV). Currents were also examined using Ba<sup>2+</sup> as the charge carrier (Fig. 3B) to determine whether the facilitation was Ca<sup>2+</sup> dependent, and in the presence of GDP $\beta$ S to limit any tonic activation of endogenous  $G\alpha\beta\gamma$  by endogenous GTP. Control facilitation was still present, although to a smaller extent, in the presence of Ba<sup>2+</sup> (Fig. 3B,  $\blacksquare$ ; P2/P1 =  $1.5 \pm 0.2$  at -20 mV), and was reduced, particularly at -30 mV, by the inclusion of GDP $\beta$ S (Fig. 3B,  $\boxtimes$ ).

As described above, expression of  $G\beta_1\gamma_2$  with  $\alpha_{1B}$  caused a prominent slowing of activation kinetics and a reduction in current amplitude (Fig. 2B). Both of these effects were readily reversed by a depolarizing prepulse (Fig. 2B, middle and right), and the facilitated current activated at more hyperpolarized potentials than that before the prepulse (Fig. 2B, middle). The facilitation ratio was  $5.9 \pm 2.0$  ( $n = 6$ ) at -20 mV (Fig. 3A,  $\boxtimes$ ). The effect of GTP $\gamma$ S on facilitation was much less marked than that of  $G\beta\gamma$  overexpression (Fig. 3B,  $\boxtimes$ ), the facilitation ratio being  $3.0 \pm 0.7$  at -20 mV. In contrast, co-expression of  $\beta$ ARK1 (495–689) almost completely abolished prepulse facilitation (Fig. 2C), both of the current amplitude (Fig. 2C, middle), and of the activation kinetics (Fig. 2C, right). Only a non-significant degree of facilitation remained at submaximal activation potentials (Fig. 3A,  $\boxtimes$ ; P2/P1 =  $1.2 \pm 0.4$  at -20 mV).



**Figure 1. Calcium channel currents in COS-7 cells transfected with  $\alpha_{1B}$ , and the effects of  $G\beta_1\gamma_2$  and  $\beta$ ARK1 (495-689) co-expression**

A-D, examples of traces of  $I_{Ca}$  recorded from cells transfected with  $\alpha_{1B}$  cDNA together with  $\alpha_2$ - $\delta$  and  $\beta_{2a}$  cDNAs. Steps of 100 ms duration were applied to increasing test potentials ( $V_t$ ) to maximally activate  $I_{Ca}$  from a holding potential ( $V_h$ ) of  $-100$  mV.  $V_t$  values are example traces between  $-40$  and  $+10$  mV. A and B, cells recorded with (A) or without (B)  $0.1$  mM GTP included in the intracellular medium. C, cells were co-transfected with  $G\beta_1$  and  $G\gamma_2$  cDNAs, and  $I_{Ca}$  was recorded in the absence of added intracellular GTP. D, cells were co-transfected with  $\beta$ ARK1 (495-689) cDNA, and  $I_{Ca}$  was recorded in the absence of added intracellular GTP. E, current density-voltage relationships for cells recorded under conditions A, C and D. Results are given as means  $\pm$  s.e.m. for 10, 11 and 15 cells, respectively. Individual current density-voltage plots were also fitted with the equation:

$$\text{Current density} = G_{\max}(V - V_{\text{rev}}) / \{1 + \exp[(V - V_{1/2})/k]\},$$

where  $G_{\max}$  is maximum conductance (nS pF<sup>-1</sup>),  $V_{1/2}$  is the voltage at which 50% of the current is activated,  $V_{\text{rev}}$  is the null potential and  $k$  is the slope factor. From this analysis,  $V_{1/2}$  was  $-6.6 \pm 2.1$  mV for control,  $-4.9 \pm 1.6$  mV for  $\beta$ ARK1 (495-689) and  $+0.1 \pm 0.2$  mV for  $G\beta\gamma$  co-expression ( $P < 0.05$  vs. control);  $G_{\max}$  was  $0.6 \pm 0.1$  nS pF<sup>-1</sup> for control,  $0.8 \pm 0.1$  nS pF<sup>-1</sup> for  $\beta$ ARK1 (495-689) and  $0.5 \pm 0.1$  nS pF<sup>-1</sup> for  $G\beta\gamma$  co-expression;  $k$  was  $4.0 \pm 0.4$  mV for control,  $3.1 \pm 0.2$  mV for  $\beta$ ARK1 (495-689) and  $4.5 \pm 0.3$  mV for  $G\beta\gamma$  co-expression; and  $V_{\text{rev}}$  was  $49.5 \pm 3.5$  mV for control,  $53.1 \pm 3.7$  mV for  $\beta$ ARK1 (495-689) and  $51.8 \pm 2.4$  mV for  $G\beta\gamma$  co-expression.



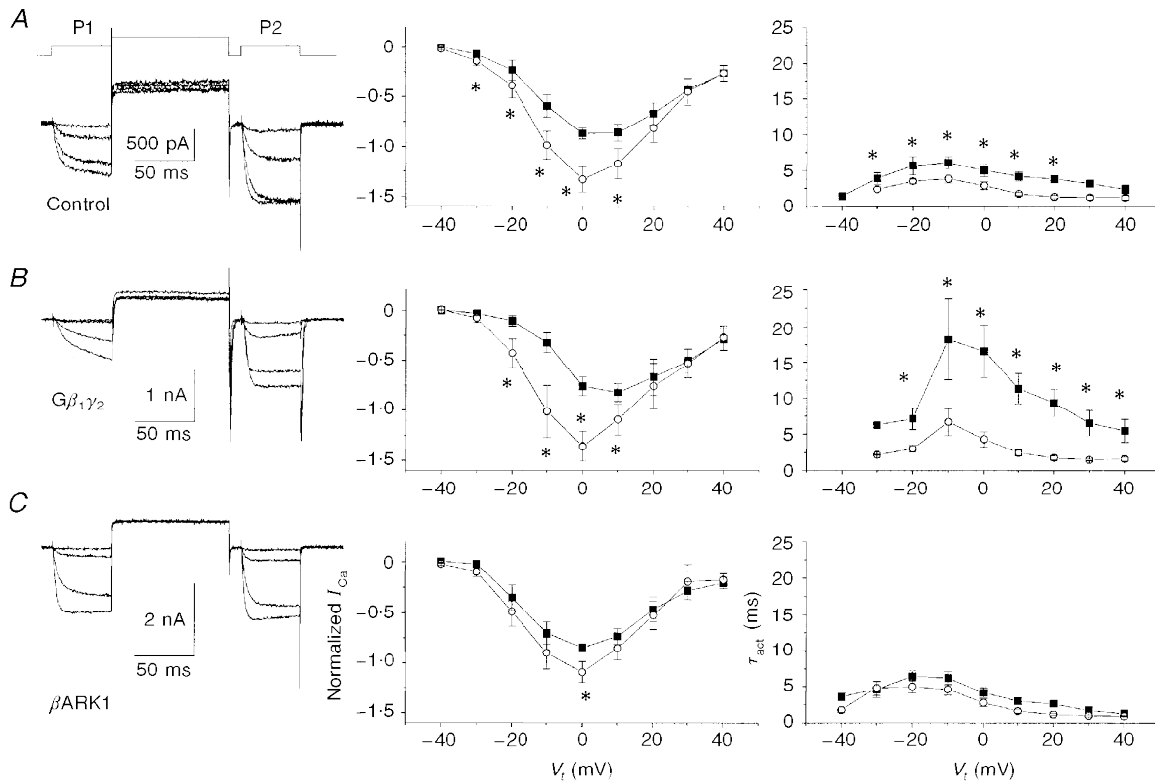
**Kinetics of reinhibition of  $\alpha_{1B}$  following a depolarizing prepulse**

A prominent model of G protein interaction with  $\alpha_1$  subunits is one where channels exist in either a normal ('willing') or a 'reluctant' gating mode; the latter requiring stronger depolarization to open (Bean, 1989; Elmslie, Zhou & Jones, 1990). One possibility is that the transition between these states is determined by binding of activated G protein ( $\beta\gamma$ ) subunits to the channel in a bimolecular reaction. Previous reports have examined the reversal of modulation (possibly representing actual dissociation of  $G\beta\gamma$  from the channel), and the reinhibition following reversal (possibly representing re-association of  $G\beta\gamma$  and the calcium channel), by using depolarizing prepulse protocols.

In order to characterize further the tonic- and G protein-mediated inhibition of  $\alpha_{1B}$ , we examined the kinetics of reinhibition by increasing the delay between the depolarizing

prepulse and the subsequent test pulse (P2). This analysis was performed for control facilitation, for that in the presence of GTP $\gamma$ S or co-expressed  $G\beta_1\gamma_2$ , and for the small degree of facilitation remaining in the presence of  $\beta$ ARK1 (495–689).

Reinhibition (re-association) kinetics at  $-100$  mV were examined using the protocols shown in Fig. 4. Following the prepulse, the time ( $\Delta t$ ) before the subsequent test pulse (P2) was incremented until current was reinhibited to the same extent as during P1, prior to the prepulse. Data were recorded at levels of maximal prepulse potentiation (usually  $-20$  mV). Under control conditions (Fig. 4A), reinhibition occurred with kinetics that were well described by a single exponential (Fig. 4D). The time constant of reinhibition ( $\tau_{\text{re-inhibition}}$ ), determined by exponential fits to data from individual cells, was  $350 \pm 35$  ms ( $n = 6$ ). This was not affected when  $Ba^{2+}$  was used as the charge carrier



**Figure 2. Prepulse potentiation of  $\alpha_{1B}$  calcium channel currents**

$I_{Ca}$  was recorded from cells transfected with  $\alpha_{1B}$  cDNA, together with  $\alpha_2\text{-}\delta$  and  $\beta_{2a}$  cDNAs in the absence of GTP in the intracellular solution, in control conditions (A,  $n = 9$ ), with  $G\beta_1\gamma_2$  co-expression (B,  $n = 7$ ), or with  $\beta$ ARK1 (495–689) co-expression (C,  $n = 10$ ). Activation of  $I_{Ca}$  was examined from  $V_h = -100$  mV, immediately before (P1) and 10 ms after (P2) application of a 100 ms depolarizing prepulse to  $+120$  mV, according to the voltage protocol given. Example traces are given in the left panel:  $V_t$  values are between  $-40$  and  $0$  mV. Current–voltage relationships are given in the middle panel from the amplitudes at 50 ms in P1 (■) and P2 (○). The currents have been normalized for each cell to the maximum current in P1 (normally at  $0$  mV in A and C, and  $+10$  mV in B). Activation was described by a single exponential fit starting at the beginning of the negative current deflection for the currents in P1 and P2, to obtain the time constant of activation,  $\tau_{\text{act}}$ .  $\tau_{\text{act}}$ –voltage relationships are given in the right panel, with the same symbols as for the middle panel. All data are given as means  $\pm$  s.e.m., and statistical significances were examined using Student's  $t$  test for paired data; \*  $P < 0.05$ .

( $\tau_{\text{reinhhibition}} = 361 \pm 124$  ms,  $n = 3$ ). For  $G\beta_1\gamma_2$  over-expression, the reinhibition kinetics of  $I_{\text{Ca}}$  were much more rapid (Fig. 4B and D), being  $41.3 \pm 8.9$  ms ( $n = 7$ ;  $P < 0.01$  compared with control). Similarly rapid reinhibition kinetics were obtained using  $\text{Ba}^{2+}$  as the charge carrier ( $\tau_{\text{reinhhibition}} = 25 \pm 5$  ms,  $n = 5$ ;  $P < 0.05$  compared with control). Reinhibition kinetics of  $I_{\text{Ba}}$  in the presence of GTP $\gamma$ S were slower than for  $G\beta_1\gamma_2$  ( $\tau_{\text{reinhhibition}} = 210 \pm 30$  ms,  $n = 5$ ;  $P < 0.05$ , compared with the  $\tau_{\text{reinhhibition}}$  in the presence of  $G\beta\gamma$ ). To determine whether the slow reinhibition kinetics in control conditions were due to a low rate of reassociation with free  $G\beta\gamma$  subunits, the time constant of reinhibition was further examined in the presence of expressed  $\beta\text{ARK1}$  (495–689) C-terminal polypeptide. Although the amount of facilitation was small, as shown above, reinhibition of  $I_{\text{Ca}}$  was extremely slow (Fig. 4C),  $\tau_{\text{reinhhibition}}$  being  $653 \pm 102$  ms ( $n = 9$ ;  $P < 0.05$  compared with control) (Fig. 4D).

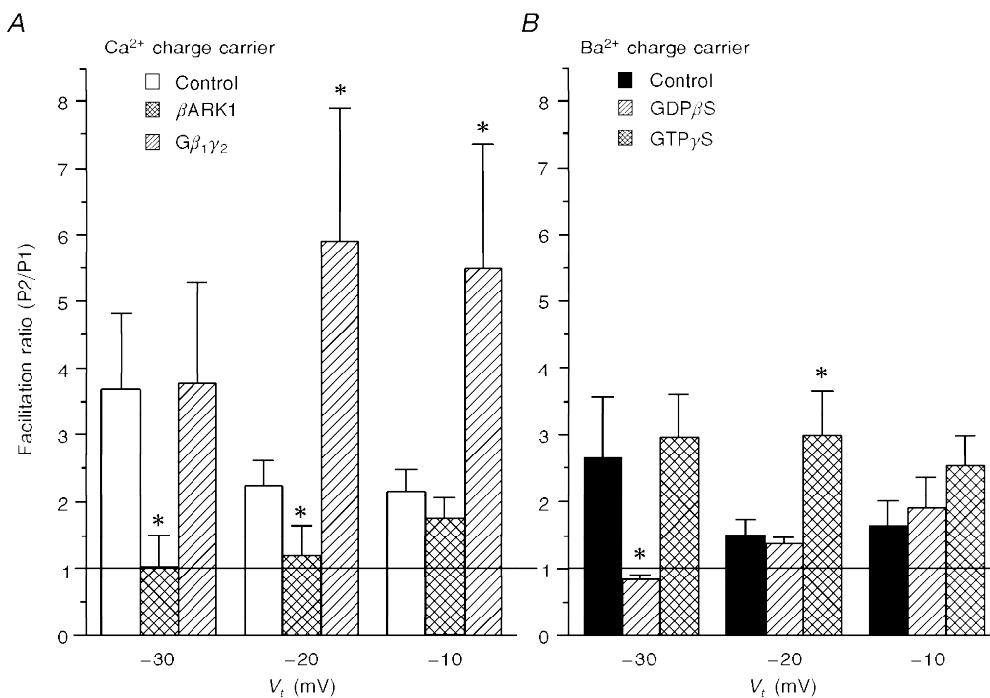
#### Kinetics of loss of G protein-mediated inhibition of $\alpha_{1B}$ during a depolarizing prepulse

Dissociation of G protein subunits from  $\alpha_{1B}$  at +120 mV was measured using the protocol shown in Fig. 5A. Two test pulses, P1 and P2, were given, with a depolarizing prepulse

to +120 mV of 0.25–100 ms duration 10 ms before P2. In the absence of a prepulse, the P1 and P2 currents are identical. The length of the depolarizing prepulse was incremented until the current elicited by the subsequent voltage step had saturated at a maximal level; data were again taken at levels of maximal prepulse potentiation (usually –20 mV). Current amplitude was measured 11 ms after the start of the test pulse, to provide an estimate of only the rapidly activating current component. The time constant of dissociation ( $\tau_{\text{diss}}$ ) was then determined by single exponential fits to data from individual cells (see Fig. 5B for examples from three of the conditions: control,  $G\beta_1\gamma_2$  and  $\beta\text{ARK1}$  (495–689) overexpression). The  $\tau_{\text{diss}}$  did not differ significantly between the different conditions examined here (Fig. 5C), being between 18 and 24 ms, in agreement with the view that relief of block should be independent of free  $G\beta\gamma$  subunit concentration at large depolarizations, where rebinding of  $G\beta\gamma$  is minimal (Jones & Elmslie, 1997).

#### Immunolocalization of calcium channel and G protein subunits

The calcium channel  $\beta$  subunit and the  $G\beta\gamma$  subunits both have binding sites located on the intracellular I–II loop of



**Figure 3. Comparison of facilitation ratios for  $\alpha_{1B}$  calcium channel currents**

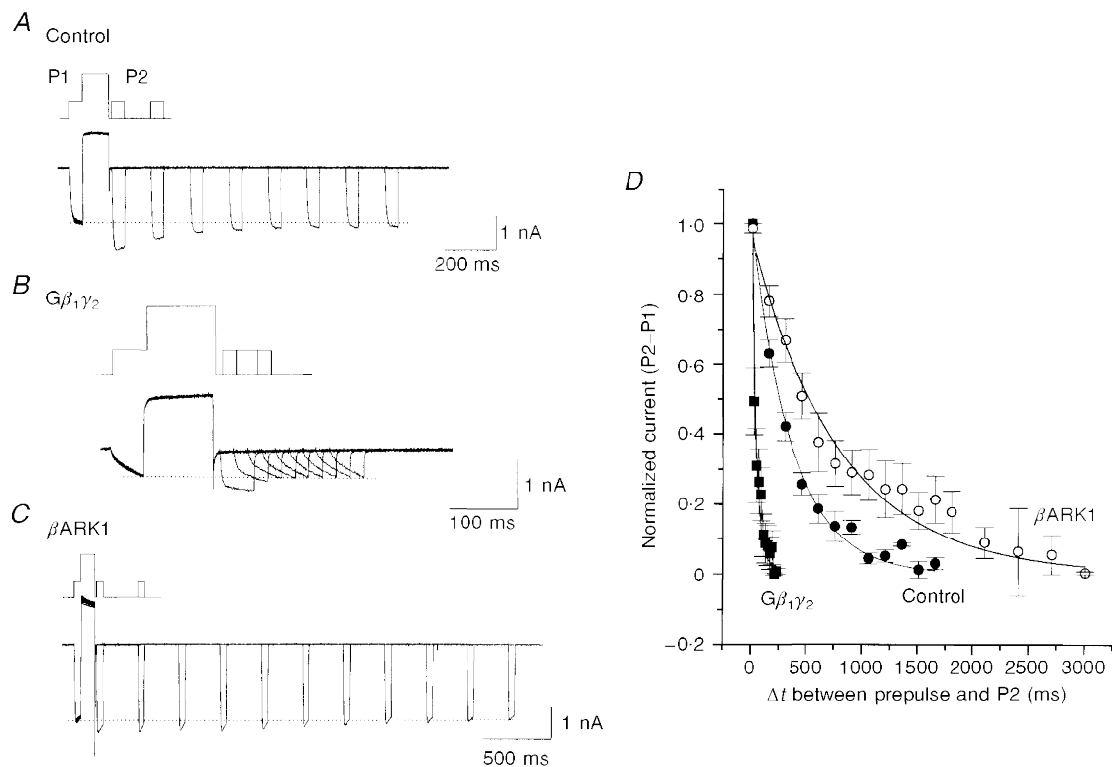
The facilitation ratio at the stated potentials is the ratio P2/P1 of the current amplitudes in P1 and P2 measured 11 ms after the start of the step when the rapidly activating current component had saturated. This was to gain a measure of the steady-state inhibition at the holding potential. P2/P1 was determined for each cell under the conditions given. A,  $\text{Ca}^{2+}$  as charge carrier: control without GTP (□),  $\beta\text{ARK1}$  (495–689) co-expression (▣) and  $G\beta_1\gamma_2$  co-expression (▨);  $n = 9, 10$  and  $6$ , respectively. B,  $\text{Ba}^{2+}$  as charge carrier: control without GTP (■),  $\text{GDP}\beta\text{S}$  (▨),  $\text{GTP}\gamma\text{S}$  (▣);  $n = 7, 4$  and  $12$ , respectively. All data are given as means  $\pm$  s.e.m., and statistical significances compared with respective controls were examined using Student's  $t$  test; \*  $P < 0.05$ .

the  $\alpha_{1B}$  subunit (Pragnell, De Waard, Mori, Tanabe, Snutch & Campbell, 1994; De Waard *et al.* 1997; Zamponi *et al.* 1997). As one of the  $G\beta\gamma$ -binding sites overlaps directly with that of the calcium channel  $\beta$  subunit, it is important to demonstrate that in the present experiments the results obtained in experiments in which  $G\beta\gamma$  is co-expressed are not due to interference with the role of the calcium channel  $\beta$  subunit in localizing the  $\alpha_1$  subunits to the plasma membrane (Chien *et al.* 1995; Brice *et al.* 1997). In support of the electrophysiological data, Fig. 6 shows that  $\alpha_{1B}$  (and also the calcium channel  $\beta$  subunit) was still predominantly membrane localized in the presence of  $G\beta_1\gamma_2$ ; in addition, a pan-specific antibody to  $G\beta$  indicated that significant amounts of this subunit also reached the cell membrane. Therefore,  $G\beta\gamma$  co-expression does not suppress the calcium channel  $\beta$  subunit function to traffic  $\alpha_1$  subunits to the plasma membrane. This is in agreement with electrophysiological results that the maximum conductance ( $G_{\max}$ ) obtained from current density–voltage relationships (see

legend to Fig. 1) was not significantly reduced in the presence of  $G\beta\gamma$  compared with the control. In 1 mM  $\text{Ca}^{2+}$ , the control  $G_{\max}$  was  $0.6 \pm 0.1$  nS  $\text{pF}^{-1}$ , whereas for  $G\beta\gamma$  overexpression it was  $0.5 \pm 0.1$  nS  $\text{pF}^{-1}$  (Fig. 1E). In 1 mM  $\text{Ba}^{2+}$ , the control  $G_{\max}$  was  $0.62 \pm 0.11$  nS  $\text{pF}^{-1}$  ( $n = 7$ ), whereas for  $G\beta\gamma$  overexpression it was  $0.65 \pm 0.13$  nS  $\text{pF}^{-1}$  ( $n = 7$ ).

## DISCUSSION

Calcium channel facilitation is a phenomenon that has been observed in a number of different systems, for both cloned and native calcium channels (for review, see Dolphin, 1996). In the case of cloned  $\alpha_{1C}$  calcium channels, facilitation has been shown to depend on the presence of a  $\beta$  subunit, and possibly to involve phosphorylation (Sculptoreanu *et al.* 1993*a,b*; Bourinet *et al.* 1994). In contrast, for native neuronal calcium channels, it has been suggested that facilitation represents the reversal of tonic G protein

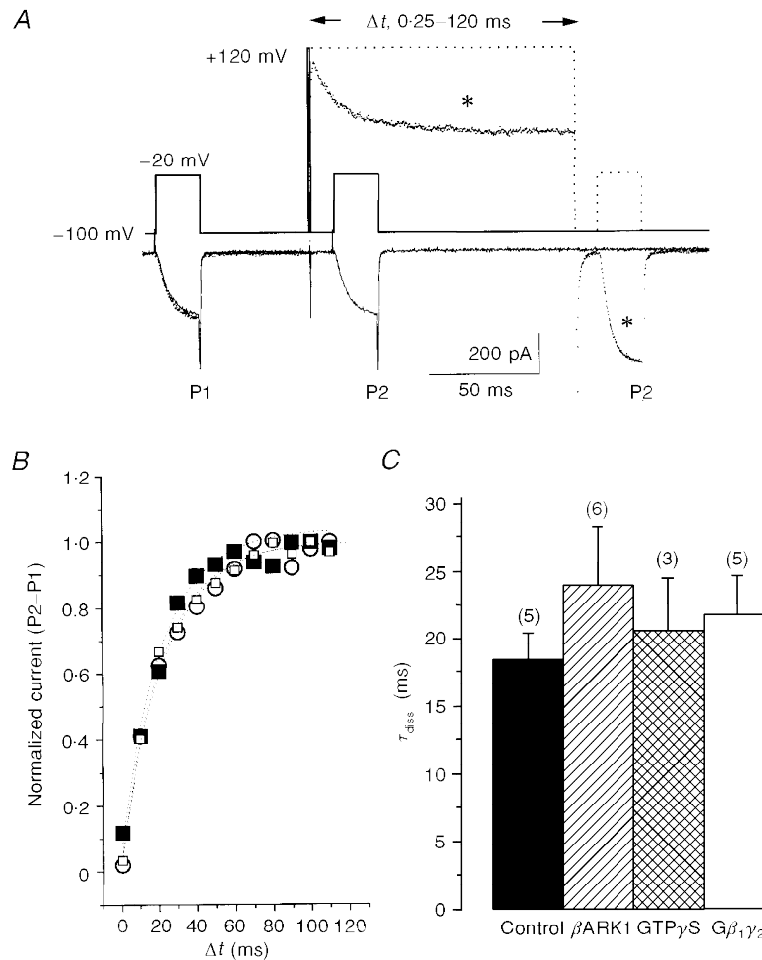


**Figure 4. Reinhibition of  $\alpha_{1B}$  at  $-100$  mV following a depolarizing prepulse to  $+120$  mV**

$I_{\text{Ca}}$  was recorded from cells transfected with  $\alpha_{1B}$  cDNA together with  $\alpha_2$ - $\delta$  and  $\beta_{2a}$  cDNAs in control conditions (no GTP in the patch pipette, *A*), and with  $G\beta_1\gamma_2$  (*B*) or  $\beta\text{ARK1}$  (495–689) co-expression (*C*). *A–C*, facilitating prepulses to  $+120$  mV were applied after a control depolarization (P1), and the time ( $\Delta t$ ) between the prepulse and the second test pulse, P2, was incremented on successive recordings until reinhibition was complete. The  $\Delta t$  increments were 150, 20 and 300 ms in *A*, *B* and *C*, respectively, and  $V_t$  was to  $-20$  mV where maximal prepulse potentiation was observed. In each case the first two voltage traces are shown above the set of current traces. A dotted line on the current traces indicates the current level at which reinhibition is complete. *D*, reinhibition of  $\alpha_{1B}/\alpha_2$ - $\delta/\beta_{2a}$  under control conditions ( $\bullet$ ,  $n = 9$ ), with  $G\beta_1\gamma_2$  ( $\blacksquare$ ,  $n = 7$ ) or with  $\beta\text{ARK1}$  (495–689) ( $\circ$ ,  $n = 9$ ). Current amplitude was measured at the end of the 50 ms step at P2 for each  $\Delta t$  value and normalized to that at maximum potentiation (i.e. P2–P1 at  $\Delta t = 10$  ms). The continuous lines represent single exponential fits to the group data.

modulation (Doupnik & Pun, 1994; Dolphin, 1996; Albillos *et al.* 1996). The present study has examined the role of endogenous and exogenous  $G\beta\gamma$  subunits in the tonic and G protein-mediated modulation of  $\alpha_{1B}$  calcium channels, which underlie the neuronal N-type current. In order to separate any effects on voltage-dependent inactivation

(Zhang, Ellinor, Aldrich & Tsien, 1994; Herlitzke, Hockerman, Scheuer & Catterall, 1997; Page *et al.* 1997) from voltage-dependent G protein-mediated slowing of activation kinetics, the auxiliary  $\beta_{2a}$  subunit was co-expressed in order to limit inactivation (Olcese *et al.* 1994). Under control conditions, amplitude facilitation (P2/P1) ratios significantly



### Figure 5. Relief of G protein inhibition of $\alpha_{1B}$ at +120 mV

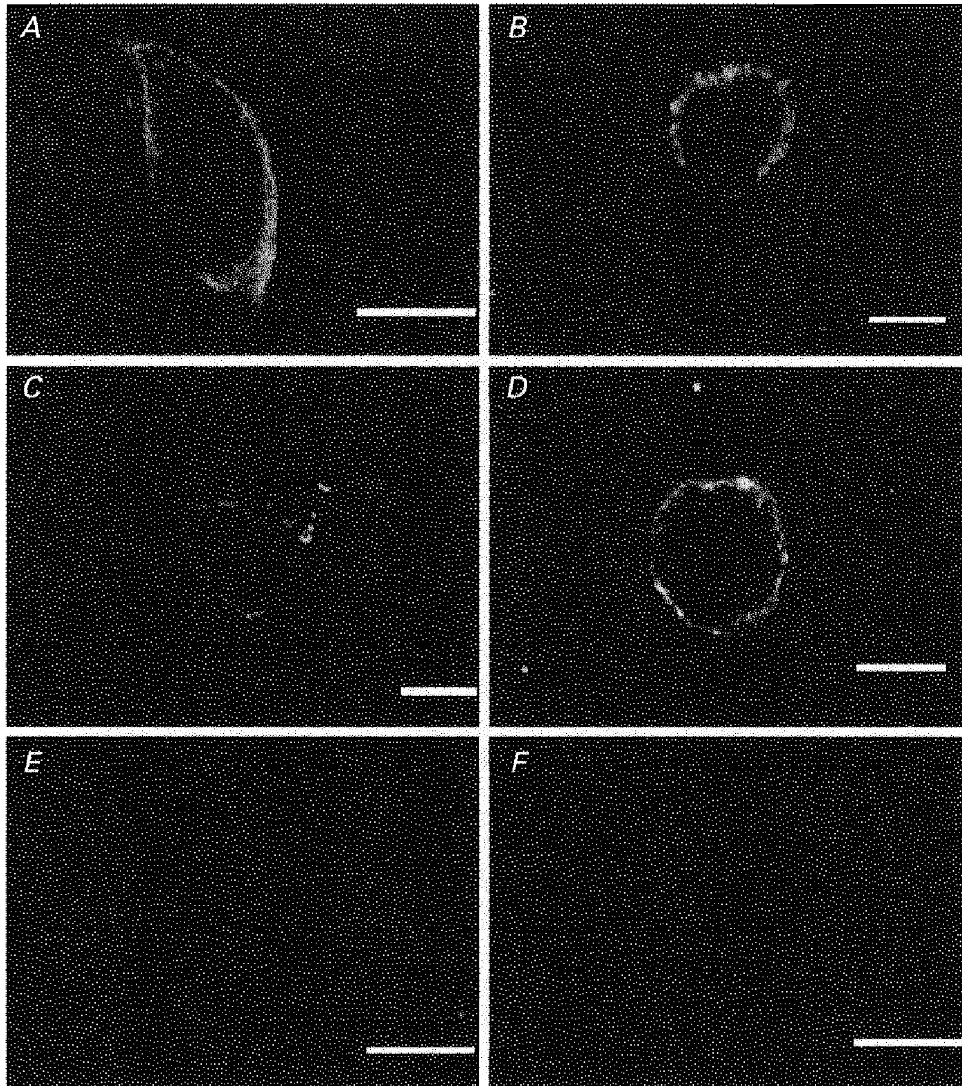
Calcium channel currents were recorded from cells transfected with  $\alpha_{1B}$  cDNA together with  $\alpha_2$ - $\delta$  and  $\beta_{2a}$  cDNAs under control conditions ( $I_{Ca}$ ), in the presence of 100  $\mu$ M GTP $\gamma$ S in the patch pipette ( $I_{Ba}$ ) or with  $G\beta_1\gamma_2$  ( $I_{Ca}$ ) or  $\beta$ ARK1 (495–689) co-expression ( $I_{Ca}$ ). *A*, two examples of the first (continuous line) and last (dotted line, trace marked with \*) currents and the corresponding voltage protocols used are shown for  $\alpha_{1B}$  under control conditions. Two test pulses, P1 and P2, were applied before and 10 ms after a depolarizing prepulse to +120 mV, respectively. The prepulse duration ( $\Delta t$ ) was incremented from 0.25 ms (thick line) to 120 ms (dotted line).  $V_t$  was at levels of maximal prepulse potentiation; for this example,  $V_t = -20$  mV. As the length of the prepulse ( $\Delta t$ ) was incremented, current at P2 increased and saturated. *B*, relief of G protein-mediated block of  $\alpha_{1B}/\alpha_2$ - $\delta/\beta_{2a}$  in typical examples in control conditions ( $\square$ ), and with  $\beta$ ARK1 (495–689) ( $\circ$ ) or  $G\beta_1\gamma_2$  ( $\blacksquare$ ) overexpression. Current amplitude was measured 11 ms after the start of the P1 and P2 steps for each  $\Delta t$  value, and normalized to that at maximum potentiation (i.e. maximum P2–P1 value). The time constant of dissociation ( $\tau_{diss}$ ) of the active G protein from the  $\alpha_{1B}$  channel was determined from a single exponential fit to data from individual experiments (dotted lines). The  $\tau_{diss}$  values for the examples given are 22.2, 21.2 and 23.3 ms for control,  $G\beta_1\gamma_2$  and  $\beta$ ARK1 (495–689), respectively. *C*, mean  $\tau_{diss} \pm$  s.e.m. for the different conditions shown, with the number of experiments given in parentheses. The rate of dissociation was similar for all conditions. There was no significant difference between control ( $\blacksquare$ ) and either decreasing free  $[G\beta\gamma]$  with  $\beta$ ARK1 (495–689) expression ( $\text{hatched}$ ), increasing free  $[G\beta\gamma]$  with GTP $\gamma$ S ( $\text{cross-hatched}$ ) or transfection of  $G\beta_1\gamma_2$  ( $\square$ ).



less than 1 were never seen with  $\beta_{2a}$ , either in the presence or absence of a depolarizing prepulse between the two test pulses. This is in contrast to results obtained with  $\beta_{1b}$ , where the significant degree of inactivation during the prepulse protocol could mask facilitation (G. J. Stephens & A. C. Dolphin, unpublished observations).

### G $\beta\gamma$ subunits mediate both G protein-induced and tonic inhibition of the $\alpha_{1B}$ calcium channel

The prepulse facilitation data for the calcium channel  $\alpha_{1B}$  subunit under control conditions suggest some tonic facilitation in COS-7 cells (Fig. 3), as previously reported for expression in HEK 293 cells (Toth, Shekter, Ma, Philipson



**Figure 6. Immunolocalization of  $\alpha_{1B}$ ,  $\beta$  and G $\beta$  subunits in COS-7 cells**

Cells were transfected with  $\alpha_{1B}$ ,  $\alpha_2\text{-}\delta$ ,  $\beta_{1b}$  and G $\beta_1\gamma_2$  cDNAs, except for *A* and *F* where the G $\beta_1\gamma_2$  subunits were omitted, and *E* which represents transfection of pMT2 vector alone. The cells were labelled with polyclonal antibodies directed against  $\alpha_{1B}$  (*A*, *B* and *E*),  $\beta_{\text{common}}$  (*C*), G $\beta_{\text{(common)}}$  (a pan-specific antibody that recognizes all G $\beta$  isoforms, *D*) or with  $\alpha_{1B}$  pre-immune serum (*F*). Scale bars are 15  $\mu\text{m}$ . The staining at the plasma membrane (M) and in the intracellular compartment (C) of a random sample of cells from each group were compared using confocal microscopy. The location of membrane and intracellular compartment were determined from an overlaid phase image. For each of the four groups, immunostaining against the underlined subunit (in arbitrary units (a.u.) determined using the Kontron KS400 program) was significantly higher for the plasma membrane than the intracellular compartment: *A*,  $\alpha_{1B}/\alpha_2\text{-}\delta/\beta_{1b}$  (M =  $117 \pm 10$  a.u., C =  $93 \pm 10$  a.u.;  $n = 8$ ,  $P < 0.005$ ); *B*,  $\alpha_{1B}/\alpha_2\text{-}\delta/\beta_{1b}$ , G $\beta_1\gamma_2$  (M =  $119 \pm 13$  a.u., C =  $69 \pm 9$  a.u.;  $n = 8$ ,  $P < 0.005$ ); *C*,  $\alpha_{1B}/\alpha_2\text{-}\delta/\beta_{1b}$ , G $\beta_1\gamma_2$  (M =  $98 \pm 8$  a.u., C =  $62 \pm 8$  a.u.;  $n = 17$ ,  $P < 0.005$ ); *D*,  $\alpha_{1B}/\alpha_2\text{-}\delta/\beta_{1b}$ , G $\beta_1\gamma_2$  (M =  $98 \pm 16$  a.u., C =  $64 \pm 12$  a.u.;  $n = 8$ ,  $P < 0.05$ ). Values are means  $\pm$  s.e.m. No staining was observed in cells transfected with pMT2 vector alone or cells stained using  $\alpha_{1B}$  pre-immune serum.

& Miller, 1996). Dialysis with GDP $\beta$ S reduced control facilitation at  $-30$  mV, indicating that rather than being an intrinsic property of the  $\alpha_{1B}$  calcium channel, as may be the case for  $\alpha_{1C}$  (Sauptoreanu *et al.* 1993a), it may be a result of tonic modulation by endogenous free G $\beta\gamma$  subunits, whose level could be influenced by the endogenous GTP concentration, and possibly by tonic activation of endogenous receptors (Koch *et al.* 1994). It may also reflect the voltage dependence of the influence of the calcium channel  $\beta$  subunit, whose binding may compete with G $\beta\gamma$  (Campbell *et al.* 1995b; Dolphin, 1996; Qin *et al.* 1997). To examine further the possibility that  $\alpha_{1B}$  is modulated by free G $\beta\gamma$  under control conditions, we examined the effect of co-expression of  $\beta$ ARK1 (495–689) (Koch *et al.* 1994). This is the C-terminal G $\beta\gamma$ -binding domain of  $\beta$ ARK1, comprising amino acids 495–689, which has no kinase activity. It has an affinity for G $\beta\gamma$  sufficient to interfere with G $\beta\gamma$ -mediated signalling in other systems (Koch *et al.* 1994). We observed a marked reduction in the amount of prepulse potentiation in cells co-expressing  $\beta$ ARK1 (495–689), presumably because it lowered the amount of free G $\beta\gamma$  able to bind to  $\alpha_{1B}$ .

In contrast, dialysis with GTP $\gamma$ S potentiated the observed facilitation of  $\alpha_{1B}$ , in agreement with our previous studies in which the auxiliary  $\beta$  subunit was  $\beta_{1b}$  (Page *et al.* 1997). Co-expression of  $\alpha_{1B}$  with G $\beta_1\gamma_2$  caused classical modulation of  $I_{Ca}$ , in which G protein inhibition is associated with a depolarizing shift in  $V_{1/2}$ . This inhibition was readily reversed by a large depolarizing prepulse, both in terms of facilitation of current levels and increase in current activation rate. These findings confirm the involvement of G $\beta\gamma$  in N-type calcium channel inhibition (Herlitze *et al.* 1996; Ikeda, 1996). It is likely that the action of GTP $\gamma$ S is to recruit endogenous G protein subunits, and in all cases co-expression of G $\beta_1\gamma_2$  led to a greater modulation of activation kinetics than GTP $\gamma$ S. These differences are probably due to a higher effective concentration of the active G protein subunits being present when G $\beta_1\gamma_2$  subunits are co-expressed, as discussed below. In addition, GTP $\gamma$ S produces, in equal proportion, G $\alpha$ -GTP $\gamma$ S and free G $\beta\gamma$ , whereas when G $\beta_1\gamma_2$  subunits are transfected they are presumably present in excess over G $\alpha$  subunits.

#### Kinetics of reinhibition of $\alpha_{1B}$ by G $\beta\gamma$ subunits following a depolarizing prepulse

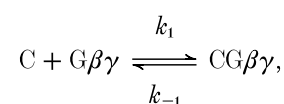
We further examined the interaction between  $\alpha_{1B}$  and activated G protein subunits in terms of reinhibition characteristics for  $\alpha_{1B}$  calcium current under control conditions, and compared this with  $\alpha_{1B}$  current in the presence of GTP $\gamma$ S (as a measure of maximal modulation of endogenous G proteins), co-expressed with G $\beta_1\gamma_2$  (as a measure of a saturated system), or co-expressed with  $\beta$ ARK1 (495–689) to bind endogenous free G $\beta\gamma$  subunits (Koch *et al.* 1994). Consistent with an increased concentration of active G protein subunits, G $\beta_1\gamma_2$  cDNA transfection produced a much faster reinhibition of  $\alpha_{1B}$  current than GTP $\gamma$ S, and this in turn was faster than the reinhibition

following facilitation under control conditions. Presumably the action of GTP $\gamma$ S is dependent on the concentration of endogenous G $\beta\gamma$  generated by GTP $\gamma$ S-induced activation of G $\alpha$  subunits. There is no G $\alpha_o$  in COS-7 cells, but they do contain G $\alpha_{11}$  and G $\alpha_q$  (see Lee, Park, Wu, Rhee & Simon, 1992) and also G $\alpha_i$  and G $\alpha_s$  (Daaka, Pitcher, Richardson, Stoffel, Robishaw & Lefkowitz, 1997), all of which could generate endogenous G $\beta\gamma$  subunits in this system. This suggests either that the level of G $\beta\gamma$  attainable from endogenous G $\alpha\beta\gamma$  heterotrimers is much lower than the level reached when G $\beta\gamma$  cDNA is co-transfected into this system, or that the particular endogenous G $\beta\gamma$  subunits generated do not couple well to the calcium channel  $\alpha_{1B}$  subunit present. The former explanation seems more likely for our results with  $\alpha_{1B}$ , in that N-type calcium channels seem to be fairly promiscuous in the G $\beta\gamma$  isoforms with which they couple (Ikeda, 1996); in agreement, we see similar effects on  $\alpha_{1B}$  when G $\beta_1\gamma_2$  is replaced by the G $\beta_2\gamma_3$  combination (G. J. Stephens & A. C. Dolphin, unpublished results). The prolongation of the reinhibition time course in the presence of  $\beta$ ARK1 (495–689) suggests that this has lowered the resting level of free G $\beta\gamma$ . Thus, during the depolarizing prepulse, residual G $\beta\gamma$  that was bound to  $\alpha_{1B}$  is removed, and rebinding is slow because it occurs from the bulk phase. The fact that rebinding does occur indicates that, at  $-100$  mV, G $\beta\gamma$  binding to the  $\alpha_{1B}$  calcium channel is of high affinity. The affinity of G $\beta\gamma$  binding to the interaction domain (AID) on the  $\alpha_{1A}$  I–II loop *in vitro* has been measured to be 63 nM (De Waard *et al.* 1997). The present results provide evidence that actual dissociation of G $\beta\gamma$ , rather than just a reduction in its effectiveness, occurs during depolarization.

The rate of calcium current reinhibition has been shown previously to be dependent on the concentration of agonist (Elmslie & Jones, 1994; Zhou, Shapiro & Hille, 1997). In oocytes, reinhibition of  $\alpha_{1B}$  following a depolarizing prepulse during modulation by a receptor agonist had a time constant of about 77 ms (Zhang, Ellinor, Aldrich & Tsien, 1996). These studies were performed in the presence of both the agonist somatostatin and exogenously expressed G $\beta\gamma$ , although G $\beta\gamma$  expression may not be maximal as the ligand was still able to inhibit currents. In general, reinhibition rates in native neurons following inhibition of the calcium current by agonist application have time constants of between 50 and 140 ms (see Dolphin, 1996; Zhou *et al.* 1997). It seems likely that transient overexpression of G $\beta_1\gamma_2$  in the present study causes maximal inhibition of the  $\alpha_{1B}$  current.

#### Estimation of free G $\beta\gamma$ concentrations

For the reaction:



C is one of the closed states of the calcium channel  $\alpha_{1B}$  subunit,  $k_1$  is the association rate constant and  $k_{-1}$  is the

dissociation rate constant, both of which may be (indirectly or directly) voltage dependent. At equilibrium, from the law of Mass Action:

$$k_1[G\beta\gamma][C] = k_{-1}[CG\beta\gamma]$$

(where  $[G\beta\gamma]$  and  $[CG\beta\gamma]$  are the concentrations of free and channel-bound  $G\beta\gamma$ , respectively), at  $-100$  mV

$$1/\tau_{\text{reinhhibition}} = k_1[G\beta\gamma] + k_{-1}, \quad (1)$$

and

$$\text{Steady-state inhibition} = k_1[G\beta\gamma]/(k_1[G\beta\gamma] + k_{-1}); \quad (2)$$

therefore,

$$P1/P2 = k_{-1}/(k_1[G\beta\gamma] + k_{-1}). \quad (3)$$

Substituting into eqn (1) the  $\tau_{\text{reinhhibition}}$  values for control,  $G\beta\gamma$  or  $\beta$ ARK1 (495–689) overexpression obtained from Fig. 4, and substituting into eqn (3) the facilitation ratios P2/P1 determined in Fig. 3A for a step to  $-20$  mV,  $k_{-1}$  at  $-100$  mV can be calculated to be  $1.25 \text{ s}^{-1}$  for control,  $4.26 \text{ s}^{-1}$  for  $G\beta_1\gamma_2$  overexpression, and  $1.28 \text{ s}^{-1}$  for  $\beta$ ARK1 (495–689) overexpression. It is of interest that the  $k_{-1}$  value is similar in the two cases involving endogenous  $G\beta\gamma$ , but is larger for co-expressed  $G\beta_1\gamma_2$ , suggesting that the dissociation rate from the  $\alpha_{1B}$  channel for different  $G\beta\gamma$  subunits may not be identical. However, if we assume the affinity of  $G\beta\gamma$  binding to  $\alpha_{1B}$  at  $-100$  mV to be similar to that calculated for  $G\beta\gamma$  binding to the  $\alpha_{1A}$  I–II intracellular loop (63 nm; De Waard *et al.* 1997), for all the conditions described here, we can estimate  $k_1$  to be 19.8, 67.6 and  $20.3 \mu\text{M}^{-1} \text{ s}^{-1}$  for control,  $G\beta_1\gamma_2$  and  $\beta$ ARK1 (495–689) overexpression, respectively, at  $-100$  mV. Taking these values, the free  $[G\beta\gamma]$  under these three conditions is 50, 295 and 11.8 nM, respectively.

### Kinetics of dissociation of $G\beta\gamma$ from $\alpha_{1B}$ during a depolarizing prepulse

The time constants for relief of G protein-mediated inhibition of  $\alpha_{1B}$  currents by varying the depolarizing prepulse are very similar under control conditions, or in the presence of GTP $\gamma$ S, or with  $G\beta_1\gamma_2$  or  $\beta$ ARK1 (495–689) co-expression ( $\tau_{\text{diss}} = 18\text{--}24$  ms). The similarity of these dissociation rates under conditions of varying free  $G\beta\gamma$  concentration indicates that there is little reassociation of  $G\beta\gamma$  during the depolarizing prepulse. Thus, there is a strong voltage dependence of  $k_1$ , and dissociation will be favoured at this potential (see Patil *et al.* 1996). Assuming  $k_1$  to be negligible at  $+120$  mV, then  $k_{-1}$  under control conditions is approximately  $50 \text{ s}^{-1}$  at this potential, about 50-fold more rapid than at  $-100$  mV.

Dissociation time constants, measured by varying the depolarizing prepulse, of between 5 and 12 ms have been reported for neuronal N-type calcium channels in the presence of GTP $\gamma$ S (Currie & Fox, 1997; Jones & Elmslie, 1997), and 3 ms for somatostatin inhibition of  $\alpha_{1B}$  currents in oocytes (Zhang *et al.* 1996), which are somewhat faster than those measured in the present study. One major

difference between this and the study of cloned  $\alpha_{1B}$  expressed in oocytes (Zhang *et al.* 1996) is that  $\alpha_{1B}$  was co-expressed with the  $\beta_{2a}$  subunit here. The role of this subunit in the kinetics of G protein modulation requires further comparative investigation; however, it may be speculated that the kinetics of interaction between  $G\beta_1\gamma_2$  and  $\alpha_{1B}$  are dependent on the  $\beta$  subunit present, since there are binding sites for both on the I–II loop. Some preliminary data support this hypothesis, in that transmitter-induced slowing of calcium channel kinetics was prominent with  $\beta_{2a}$ , but not seen with  $\beta_{1b}$  (Patil, Brody, Snutch & Yue, 1997). The molecular composition of neuronal calcium channels, and in particular  $\beta$  subunit diversity, may thus have important consequences not only for voltage-dependent inactivation, but also in the fine-tuning of modulation by  $G\beta\gamma$  subunits.

We also show that  $G\beta\gamma$  co-expression does not suppress calcium channel  $\beta$  subunit function to traffic  $\alpha_1$  subunits to the plasma membrane. Conversely, current amplitude and activation rates were clearly inhibited by  $G\beta_1\gamma_2$  in the presence of  $\beta$  subunits; and the reversal of inhibition by depolarizing prepulses suggests that the nature of the interaction of  $G\beta\gamma$  with the channel is highly dynamic. In contrast, the  $\alpha_1$ – $\beta$  interaction might be expected to be much more stable with very high affinity binding reported *in vitro* (De Waard, Witcher, Pragnell, Liu & Campbell, 1995). Nevertheless, it is likely that this binding is also voltage dependent, and it will be of interest to determine the effects of the  $\beta$  subunit on  $G\beta\gamma$  binding to the I–II loop, to examine more fully the hypothesis that  $\beta$  subunits and G proteins may interact competitively with calcium channel  $\alpha_1$  subunits (Campbell *et al.* 1995b; Bourinet, Soong, Stea & Snutch, 1996; Qin *et al.* 1997).

In conclusion, the data shown here indicate that under control conditions there is sufficient free  $G\beta\gamma$  present in the vicinity of the  $\alpha_{1B}$  calcium channel to produce a significant tonic block of  $\alpha_{1B}$  current. If this also occurs in native neurons and neurosecretory cells, it would lead to frequency- or depolarization-dependent facilitation of calcium entry through N-type channels, as has been observed in several cell types (e.g. Doupnik & Pun, 1994; Albillos *et al.* 1996).

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