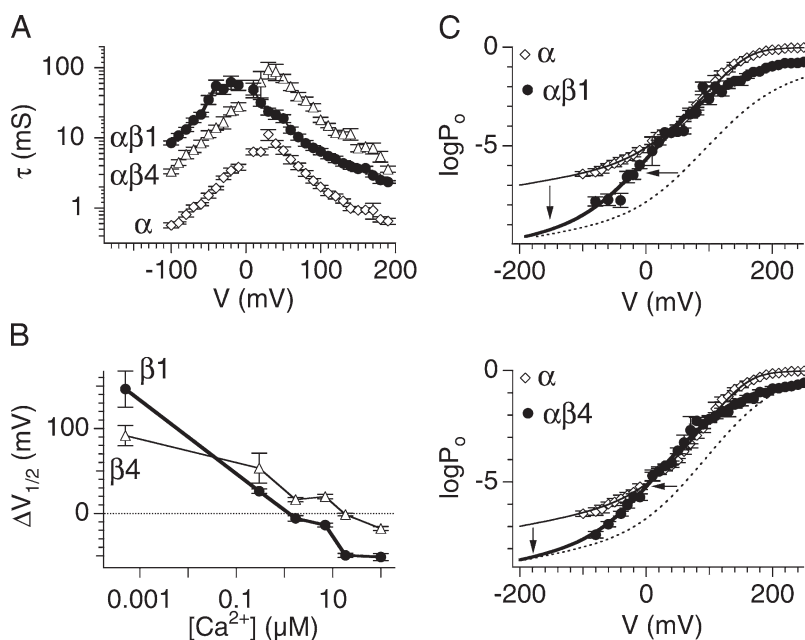
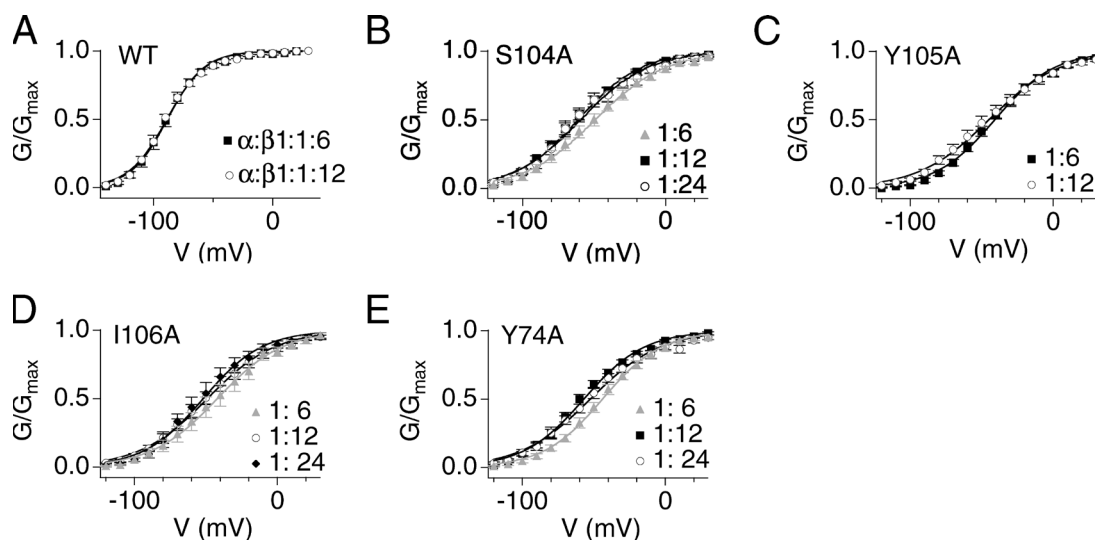
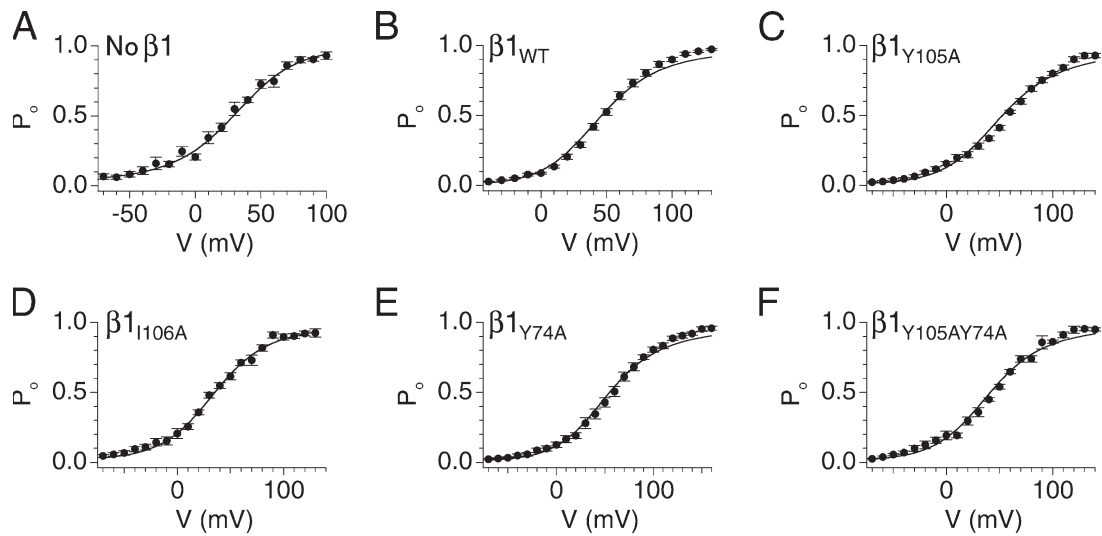


Gruslova et al., <http://www.jgp.org/cgi/content/full/jgp.201110698/DC1>

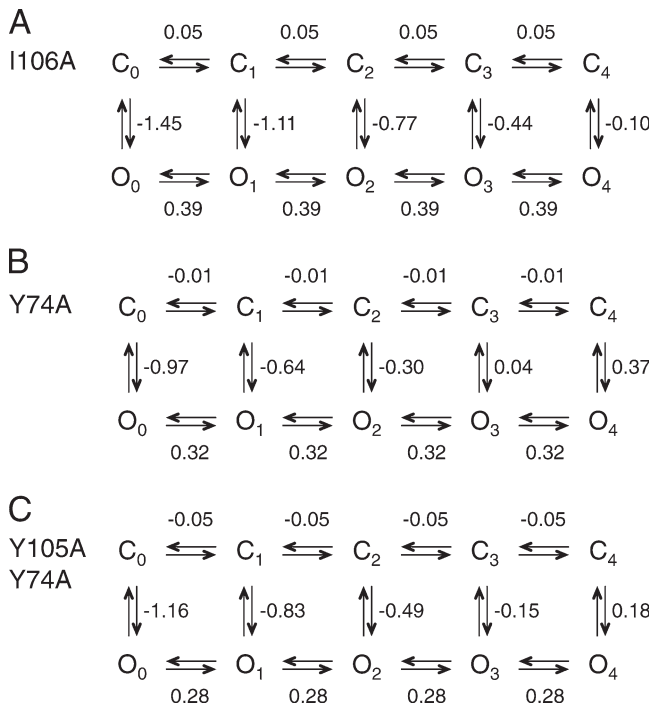
**Figure S1.**  $\beta 1$  and  $\beta 4$  share common gating effects. (A) Both  $\beta 1$  and  $\beta 4$  slow gating kinetics. Averaged activation and deactivation time constants at  $7 \mu\text{M Ca}^{2+}$ .  $\alpha$ ,  $n = 10$ – $13$ ;  $\alpha\beta 1$ ,  $n = 13$ – $27$ ;  $\alpha\beta 4$ ,  $n = 12$ – $26$ . (B) Both  $\beta 1$  and  $\beta 4$  increase steady-state opening at high  $\text{Ca}^{2+}$  but reduce opening in lower  $\text{Ca}^{2+}$ .  $\Delta V_{1/2}$  represents differences between  $\alpha\beta$  channels and  $\alpha$  alone channels.  $\alpha$ ,  $n = 8$ – $44$ ;  $\alpha\beta 1$ ,  $n = 7$ – $39$ ;  $\alpha\beta 4$ ,  $n = 10$ – $26$ . (C)  $0 \text{ Ca}^{2+}$   $\log P_o$ - $V$  relations reveal similar dual gating effects of  $\beta 1$  and  $\beta 4$ : reducing intrinsic gating (downwards arrow) and stabilization of voltage sensor activation (leftwards arrow). These data have been reported in our previous publications (Wang and Brenner, 2006; Wang et al., 2006).  $\alpha$ ,  $n = 3$ – $12$ ;  $\alpha\beta 1$ ,  $n = 4$ – $22$ ;  $\alpha\beta 4$ ,  $n = 4$ – $11$ . Error bars represent SEM.



**Figure S2.** Alanine substitutions of four  $\beta 1$  residues alter G-V relations. (A–E) Averaged G-V relations of  $\alpha\beta 1_{\text{WT}}$ ,  $\alpha\beta 1_{\text{S104A}}$ ,  $\alpha\beta 1_{\text{Y105A}}$ ,  $\alpha\beta 1_{\text{I106A}}$ , and  $\alpha\beta 1_{\text{Y74A}}$  at indicated  $\alpha/\beta 1$  molar ratios. For  $\alpha\beta 1_{\text{WT}}$  and  $\alpha\beta 1_{\text{Y105A}}$  channels, G-Vs largely overlap at both 1:6 and 1:12  $\alpha/\beta 1$  molar ratios, suggesting that  $\beta 1$  expression is saturated. For  $\alpha\beta 1_{\text{S104A}}$ ,  $\alpha\beta 1_{\text{I106A}}$ , and  $\alpha\beta 1_{\text{Y74A}}$  channels, reducing  $\alpha/\beta 1$  molar ratio from 1:6 to 1:12 caused negative G-V shifts; however, further reduction to 1:24 did not produce further negative G-V shifts. For the 1:12  $\alpha/\beta$  molar ratio,  $\alpha\beta 1_{\text{WT}}$ ,  $n = 14$ ;  $\alpha\beta 1_{\text{S104A}}$ ,  $n = 9$ ;  $\alpha\beta 1_{\text{Y105A}}$ ,  $n = 15$ ;  $\alpha\beta 1_{\text{I106A}}$ ,  $n = 9$ ;  $\alpha\beta 1_{\text{Y74A}}$ ,  $n = 14$ . For the 1:24  $\alpha/\beta 1$  molar ratio,  $\alpha\beta 1_{\text{S104A}}$ ,  $n = 8$ ;  $\alpha\beta 1_{\text{I106A}}$ ,  $n = 5$ ;  $\alpha\beta 1_{\text{Y74A}}$ ,  $n = 10$ . Error bars represent SEM.



**Figure S3.** Segment A and B residues contribute to stabilization of open-channel voltage sensor activation. (A–F) Averaged  $P_o$ -V relations (circles) and best fits to the Horrigan-Aldrich model (black curves). The number of patches is the same as in Fig. 7. Error bars represent SEM.



**Figure S4.** Effects of I106A, Y74A, and Y105AY74A on intrinsic gating and voltage sensor activation. Numbers indicate effects of the mutation on free energies associated with C-C transitions (right to left), O-O transitions (left to right), and C-O transitions (downward).

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