# Molecular Determinants of $Ca^{2+}$ Potentiation of Diltiazem Block and $Ca^{2+}$ -Dependent Inactivation in the Pore Region of $Ca_v 1.2$

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## ABSTRACT

Diltiazem block of Ca<sub>v</sub>1.2 is frequency-dependent and potentiated by Ca<sup>2+</sup>. We examined the molecular determinants of these characteristics using mutations that affect Ca<sup>2+</sup> interactions with Ca<sub>v</sub>1.2. Mutant and wild-type (WT) Ca<sub>v</sub>1.2 channels were transiently expressed in tsA 201 cells with  $\beta_{1b}$  and  $\alpha_2\delta$ subunits. The four conserved glutamates that compose the Ca<sup>2+</sup> selectivity filter in Ca<sub>v</sub>1.2 were mutated to Gln (E363Q, E709Q, E1118Q, E1419Q), and each single mutant was assayed for block by diltiazem using whole-cell voltage-clamp recordings in either 10 mM Ba<sup>2+</sup> or 10 mM Ca<sup>2+</sup>. In Ba<sup>2+</sup>, none of the mutations affected the potency of diltiazem block of closed channels (0.05 Hz stimulation). However, frequencydependent block (1Hz stimulation) was eliminated in the mutant E1419Q (domain IV), which recovered more rapidly than WT channels from inactivated channel block. Potentiation of dilti-

Ca<sup>2+</sup> influx via voltage-dependent L-type Ca<sup>2+</sup> channels  $(\alpha_{1C}, Ca_{v}1.2)$  found in cardiac and vascular smooth muscle initiates contraction and contributes to timing of the cardiac action potential (Bers and Perez-Reyes, 1999). Ca, 1.2 channels consist of an  $\alpha_1$  pore-forming subunit and  $\alpha_2\delta$ ,  $\beta$ , and  $\gamma$ auxiliary subunits (Jones, 1998). The  $\alpha_1$  subunit is composed of four homologous domains (I-IV), each of which has six transmembrane segments (S1-S6) (Takahashi et al., 1987; Tanabe et al., 1987). Each domain also includes a pore-lining region, between each S5 and S6 segment, each of which contains a conserved Glu residue (EI, EII, EIII, EIV) that contributes to the Ca<sup>2+</sup>-selectivity filter (Yang et al., 1993). In addition, the Ca<sub>v</sub>1.2  $\alpha_1$  subunit contains an IQ calmodulin binding motif in the intracellular C-terminal tail, which is critical for Ca<sup>2+</sup>-dependent channel inactivation (Peterson et al., 1999; Zuhlke et al., 1999). Ca. 1.2 is sensitive to block by three distinct chemical classes of small-molecule drugs: diazem block of closed Ca<sub>v</sub>1.2 channels in Ca<sup>2+</sup> was abolished in the E1118Q, F1117G (domain III), and E1419Q mutants. Frequency-dependent block in Ca<sup>2+</sup> was reduced compared with WT Ca<sub>v</sub>1.2 in the F1117G, E1118Q, and E1419Q mutants. The C-terminal tail IQ domain mutation I1627A, which disrupts Ca<sup>2+</sup> dependent inactivation, enhanced diltiazem block of closed channels in Ba<sup>2+</sup>. We conclude that, in Ba<sup>2+</sup>, E1419 slows recovery from diltiazem block of depolarized Ca<sub>v</sub>1.2 channels, but in Ca<sup>2+</sup>, E1118, E1419, and F1117 form a Ca<sup>2+</sup> binding site that mediates the potentiation of diltiazem block of both closed and inactivated Ca<sub>v</sub>1.2 channels. Furthermore, Ca<sup>2+</sup>-dependent inactivation, which is impaired in E709Q, E1118Q, E1419Q, and I1627A, is not required for Ca<sup>2+</sup> potentiation of diltiazem block.

hydropyridines (DHPs), phenylalkylamines (PAAs), and benzothiazepines (BZPs) (Hockerman et al., 1997b). All three classes are effective vasodilators, and are therefore used to treat hypertension and angina pectoris (Fleckenstein and Fleckenstein-Grun, 1980). The PAA verapamil and the BZP diltiazem are also used to treat some types of supraventricular arrhythmia since they preferentially block channels that are activated at high frequency, a property known as frequency-dependent block (Lee and Tsien, 1983).

Diltiazem blocks  $Ca_v 1.2$  channels at low micromolar concentrations in both primary cardiac myocytes (Lee and Tsien, 1983) and heterologous expression systems (Hockerman et al., 2000). Based on studies using a quaternary amine derivative, BZPs are thought to approach their binding site on  $Ca_v 1.2$  from the extracellular face of the plasma membrane (Hering et al., 1993). Studies using both point mutants and chimeric channels agree that closed-channel block by diltiazem involves specific amino acid residues in segments IIIS6 and IVS6 (Hering et al., 1996; Kraus et al., 1998; Berjukow et al., 1999; Hockerman et al., 2000). Mutation of amino acid

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**ABBREVIATIONS:** DHP, dihydropyridine; PAA, phenylalkylamine; BZP, benzothiazepine; GFP, green fluorescent protein; V<sub>1/2</sub>, voltage at which half of the channels are inactivated; D888, desmethoxyverapamil.

residues in IVS6 involved in closed-channel block by diltiazem also decreased the accumulation of block during a 1-Hz train of depolarization (Berjukow et al., 1999). Chimeric channels with clusters of amino acid substitutions in transmembrane segment IVS5 of  $Ca_v 1.2$  prevented frequency-dependent block by diltiazem (Motoike et al., 1999) but also severely disrupted voltage-dependent inactivation.

Mutation of EIII or EIV to Gln in Ca<sub>v</sub>1.2 was reported to decrease affinity of the PAA D888 in Ba<sup>2+</sup>, suggesting an interaction between the positively charged amino group in the D888 molecule and the negatively charged Glu residues at physiological pH (Hockerman et al., 1997a). Since diltiazem also contains an essential ionizable amino group, we assayed Ca<sub>v</sub>1.2 channels with single-amino acid mutations of each of the conserved pore Glu residues to Gln [E363Q (EIQ), E709Q (EIIQ), E1118Q (EIIIQ), and E1419Q (EIVQ)] for closed-channel block and frequency-dependent block by diltiazem in Ba<sup>2+</sup>. Diltiazem block of closed channels in Ba<sup>2+</sup> was not significantly affected by any of the E to Q mutations. However, frequency-dependent block of Ca<sub>v</sub>1.2 Ba<sup>2+</sup> current was virtually eliminated in the mutant EIVQ channel.

The conserved glutamates in the pore region of Ca, 1.2 bind Ca<sup>2+</sup> ions as they traverse the channel pore, thus forming a selectivity filter that largely excludes monovalent cations when  $Ca^{2+}$  is present (Yang et al., 1993). However, other divalent ions such as Ba<sup>2+</sup> can pass through this selectivity filter (Hille, 1995). Indeed, the conductance of Ca<sub>v</sub>1.2 is higher for Ba<sup>2+</sup> than for Ca<sup>2+</sup> since Ca<sup>2+</sup> binds more tightly to the pore glutamates (Almers and McCleskey, 1984). Another consequence of  $\operatorname{Ca}^{2+}$  binding in the pore is modulation of drug binding sites. All three chemical classes of L-type channel blockers are more potent when Ca<sup>2+</sup>, rather than  $Ba^{2+}$ , is the permeant ion (Lee and Tsien, 1983). The conserved Glu residues in domains III and IV, as well as an adjacent Phe in the pore region of domain III in Ca<sub>v</sub>1.1, mediate the Ca<sup>2+</sup> potentiation of DHP affinity (Peterson and Catterall, 1995). In addition, Ca<sup>2+</sup> also modulates Ca<sub>v</sub>1.2 channel inactivation via a Ca<sup>2+</sup>/calmodulin binding domain (IQ motif) in the C-terminal tail (Peterson et al., 1999; Zuhlke et al., 1999). We report here that the EIIIQ, EIVQ, and F1117G mutations each abolish Ca2+ potentiation of both closed-channel and frequency-dependent diltiazem block. In contrast, the IQ-domain mutation I1627A potentiated diltiazem block of closed channels in Ba<sup>2+</sup>. Furthermore, Ca<sup>2+</sup>-dependent inactivation, which is markedly reduced in the mutant channels EIIQ, EIIIQ, EIVQ, and I1627A, is not required for the Ca<sup>2+</sup> potentiation of diltiazem block.

# Materials and Methods

**Construction of Wild-Type and Mutant Ca<sup>2+</sup> Channels.** The Ca<sub>v</sub>1.2 (Snutch et al., 1991) and the mutant channels were cloned into pCDNA3 expression vector (Invitrogen, Carlsbad, CA). Mutations were introduced as described (Hockerman et al., 1997a), or using a modification of the QuikChange site-directed mutagenesis procedure (Stratagene, La Jolla, CA). The desired mutations were verified, and the integrity of the clones was confirmed by cDNA sequencing and extensive restriction digest analysis.

**Cell Culture.** Human tsA-201 cells, a simian virus 40 T-antigenexpressing derivative of the human embryonic kidney cell line HEK293, were maintained in monolayer culture in Dulbecco's modified Eagle's medium/F-12 (Invitrogen) enriched with 10% fetal bovine serum (Hyclone Laboratories, Logan, UT) and incubated at 37°C in 10%  $\rm CO_2.$ 

**Expression of Ca<sup>2+</sup> Channels.** tsA-201 cells were cotransfected with WT and mutant Ca<sub>v</sub>1.2 subunits,  $\beta_{1b}$  (Pragnell et al., 1991),  $\alpha_2\delta$  (Ellis et al., 1988), and enhanced green fluorescent protein (GFP) (BD Biosciences Clontech, Palo Alto, CA) such that the molar ratio of the plasmids was 1:1:10.8. Cells were transfected using the Gene-Porter reagent (Gene Therapy Systems Inc., San Diego, CA) and cells were replated at low density for electrophysiological recording 20 to 24 h after transfection. Experiments were conducted 20 to 48 h after replating.

Electrophysiology. Transfected cells were recognized by GFP fluorescence at 510 nm with excitation at 480 nm. Barium and calcium currents through Ca<sup>2+</sup> channels were recorded using the whole-cell configuration of the patch-clamp technique. Patch electrodes were pulled from VWR micropipettes (VWR, West Chester, PA) and fire-polished to produce an inner tip diameter of 4 to 6  $\mu$ m. Currents were recorded using an Axon Instruments Inc. (Union City, CA) Axopatch 200B patch-clamp amplifier and filtered at 1 or 2 kHz (8-pole Bessel filter, -3 dB). Voltage pulses were applied and data were acquired using pClamp8 software (Axon Instruments Inc.). Voltage-dependent leak currents were subtracted using an on-line P/-4 subtraction paradigm. (+)-cis-Diltiazem, dissolved in bath saline, was applied to cells using a Biologic RSC 160 fast perfusion system (Molecular Kinetics, Pullman WA) with constant exchange of the bath solution. Barium current was measured in a bath saline containing Tris (150 mM), MgCl<sub>2</sub> (2 mM) and BaCl<sub>2</sub> (10 mM). Calcium current was measured in the same bath solution, except that Ba<sup>2+</sup> was replaced with Ca<sup>2+</sup> (10 mM). The intracellular saline contained N-methyl-D-glucamine (130 mM), EGTA (10 mM), HEPES (60 mM), MgATP (2 mM), and MgCl<sub>2</sub> (1 mM). (+)-cis-Diltiazem (Fluka Chemical Corp., Ronkonkoma, NY) concentrations were made from a 100 mM stock solution in 70% ethanol. The pH of both solutions was adjusted to 7.3 with methanesulfonic acid. All experiments were performed at room temperature (20-23°C).

**Data Analysis.** Data were analyzed using Clampfit (Axon Instruments Inc.) and SigmaPlot (SPSS Science, Chicago, IL) software. Statistical significance was determined using Student's *t* test.

# **Results**

Structural Features and Sites of Mutations in Ca. 1.2. Figure 1, A and B, illustrates the positions of the amino acids that were mutated for this study. The amino acids E363, E709, E1118, and E1419 were mutated to Gln to create the mutant channels EIQ, EIIQ, EIIIQ, and EIVQ, respectively. As shown in Fig. 1B, the alignment of the pore residues in each domain indicates a high degree of conservation which is also retained across non-L-type channels (Jones, 1998). The structure of diltiazem (Fig. 1C) includes a tertiary amino group that is predominantly charged at physiological pH and could potentially interact with the acidic Glu side chains. The Phe residue at position 1117 in domain III is unique to L-type channels, whereas most non-L-type channels contain a Gly residue at this position. The mutation of the residue corresponding to F1117 to G in Ca. 1.1 disrupted Ca<sup>2+</sup> potentiation of DHP binding (Peterson and Catterall. 1995), as did the EIIIQ and EIVQ mutation. The approximate position of I1627 in the Ca<sup>2+</sup>/calmodulin-binding IQ motif in the C-terminal tail is indicated. Mutation of I1627 to A disrupts Ca<sup>2+</sup>/calmodulin binding to the channel and, consequently, Ca<sup>2+</sup>-dependent acceleration of inactivation (Zuhlke et al., 1999). We examined these Ca<sub>v</sub>1.2 mutants for diltiazem block and potentiation of diltiazem block by Ca<sup>2+</sup>.

Closed-Channel Block of the E to Q Mutants by Diltiazem in Ba<sup>2+</sup>. WT and mutant EIQ, EIIQ, EIIQ, EIIQ, and EIVQ Ca<sub>v</sub>1.2 channels were coexpressed along with  $\alpha_2\delta$ ,  $\beta_{1b}$ , and GFP in tsA-201 by transient transfection. Fortyeight hours after transfection, whole-cell Ba<sup>2+</sup> currents were measured in the cells expressing the GFP marker. Closedchannel block was measured from a holding potential of -60mV using 100-ms steps to +10 mV at 0.05 Hz. Increasing concentrations (5, 10, 50, 100, and 500  $\mu$ M) of diltiazem were applied to cells thus stimulated. Under these conditions, block reached equilibrium rapidly, and little frequency-dependent block was detected (Fig. 2A). The diltiazem IC<sub>50</sub> value for WT was 65.04  $\pm$  13.60  $\mu$ M, and the diltiazem IC<sub>50</sub> values for EIQ, EIIQ, EIIIQ, and EIVQ were not significantly different (Fig. 2, B and C).

Frequency-Dependent Block of E to Q Mutants in  $Ba^{2+}$ . Since none of the E to Q mutants appeared to affect diltiazem block of closed channels, we next examined the effect of these mutations on frequency-dependent block. After diltiazem block (50  $\mu$ M) reached equilibrium at 0.05 Hz as described above, we applied a 20-pulse, 1-Hz train of 100-ms depolarizations to +10 mV from a holding potential of -60 mV. The same 1-Hz train of depolarizations was also applied to cells before the application of diltiazem. The results of these experiments with WT Ca<sub>v</sub>1.2 and each of the E to Q mutations are shown in Fig. 3. Frequency-dependent diltiazem block of WT Ca<sub>v</sub>1.2 channels was not different from



Fig. 1. Structural features of the  $\alpha_1$  subunit of Ca<sub>v</sub>1.2. A, topology of Ca<sub>v</sub>1.2 ( $\alpha_{1C}$ ). Cylinders represent transmembrane segments (1–6) organized into four homologous domains (I-IV). The C- and N-terminal domains are intracellular. Boxes highlight the putative pore-lining regions that contain the elements of the Ca<sup>2+</sup> selectivity filter. B, the amino acid sequence surrounding the Glu residues in each homologous domain that compose the selectivity filter. The conserved Glu residues are in bold italic type, as is the L-type-specific Phe residue directly adjacent to the conserved Glu residue in domain III (box). The approximate location of Ile 1627 and elements of the IQ Ca<sup>2+</sup>/calmodulin binding domain are shown. C, the structure of diltiazem. Note the ionizable tertiary alkylamino group.

that of the mutant EIQ and EIIQ channels in 10 mM Ba<sup>2+</sup> (Fig. 3A). Frequency-dependent diltiazem block by diltiazem was retained in the EIIIQ channel but was significantly reduced compared with WT Ca<sub>v</sub>1.2 (Fig. 3B). In contrast, frequency-dependent diltiazem block was virtually eliminated in the EIVQ mutant, whereas inactivation at the end of



Fig. 2. Diltiazem block of WT Ca $_{\!\!\!\!\!\!\!\!\!}$ 1.2 and the mutant channels EIQ, EIIQ, EIIIQ, and EIVQ at 0.05 Hz in Ba $^{2+}$ . A, representative Ba $^{2+}$  current records from a tsA-201 cell expressing wild-type Ca<sub>v</sub>1.2 in the absence (control) or presence of the indicated concentrations of (+)-cis-diltiazem (inset). The time course of diltiazem block of  $\mathrm{Ca_v}1.2~\mathrm{Ba^{2+}}$  current is shown. Diltiazem was applied at the pulse number and concentrations indicated. Pulses were from a holding potential of -60 mV to +10 mV every 20 s. B. dose-response relationships for wild-type and the indicated mutant Ca<sub>v</sub>1.2 channels. In each case, the averaged, normalized current amplitudes at 5, 10, 50, 100, and 500  $\mu$ M diltiazem (symbols ± S.E., n =3-4) were plotted against the corresponding drug concentration, and the  $IC_{50}$  value was determined by fitting the averaged relative current values at each diltiazem concentration to the equation, relative current = 1 - 1 $\{1/[1+(IC_{50}/[diltiazem])]\}$  (smooth lines). C, the  $IC_{50}$  values of the indicated mutant channels are shown  $\pm$  S.E. (n = 3-5). The values were: WT,  $Ca_v 1.2 = 65.0 \pm 13 \ \mu M; EIQ = 76.8 \pm 17 \ \mu M; EIIQ = 86.8 \pm 19 \ \mu M;$ EIIIQ = 83.4  $\pm$  20  $\mu M;$  EIVQ = 71.5  $\pm$  18  $\mu M.$  None of the mutant channels was significantly different from the WT Ca, 1.2 channel in sensitivity to block by diltiazem under these conditions.

the train of depolarizations in the absence of drug was not significantly different from that of WT (Fig. 3C). Thus, the conserved glutamate residues have distinct effects on the frequency-dependent block of diltiazem. In Fig. 3D, the specific diminution of frequency-dependent, but not closed-channel block in EIVQ is demonstrated in single cells. When 50  $\mu$ M diltiazem is initially applied to either WT or EIVQ Ca<sub>v</sub>1.2 channels stimulated at 0.05 Hz, the same fraction of current is blocked. However, when a 1-Hz train of depolarizations is applied ~7 s after the initial application of diltiazem, the further decrease in current is substantially greater in WT Ca<sub>v</sub>1.2 than in EIVQ. These data suggest that the EIV pore glutamate is required for frequency-dependent diltiazem block of Ca<sub>v</sub>1.2 when Ba<sup>2+</sup> is used as the permeant ion.

Kinetic Analysis of Depolarized Channel Block of WT and E to Q Mutants. To understand the drastic reduction of frequency-dependent diltiazem block in the EIVQ mutant, we examined some of the voltage-dependent properties of the channel, as well as some kinetic parameters of diltiazem block. In Fig. 4A, the current-voltage relationship for EIVQ is compared with that of WT Ca<sub>v</sub>1.2. Although channel activation and the peak of the IV curve is shifted to more positive potentials for EIVQ, the fraction of channels activated at +10 mV is not greatly different from WT and is not likely to account for the sharp decrease in frequencydependent block. In Fig. 4B, the voltage dependence of inactivation for both WT Ca. 1.2 and EIVQ is shown in the absence and presence of 50  $\mu$ M diltiazem. Despite a slight (~3 mV) positive shift in the  $V_{\rm 1/2}$  inactivation for EIVQ compared with WT, the shift in  $V_{1/2}$  inactivation induced by 50  $\mu$ M diltiazem is virtually identical in both channels ( $\sim$ 32 mV). Thus, diltiazem interacts with inactivated WT and EIVQ Ca<sub>v</sub>1.2 channels to a similar extent. Recovery from depolarized channel block was also examined for WT and EIVQ channels (Fig. 4, C and D). Whole-cell Ba<sup>2+</sup> current was measured by depolarizing cells to +10 mV for 1 s followed by recovery intervals of 0.01, 0.10, 0.50, 1, 10, 20, and 30 s at -60 mV. At the end of each recovery period, recovered current was measured by stepping to +10 mV for 50 ms. This protocol was applied in the absence and presence of 50  $\mu$ M diltiazem. When performed in the presence of drug, the protocol was applied following equilibration of drug block at 0.05



**Fig. 3.** Diltiazem block of WT Ca<sub>v</sub>1.2 and the mutant channels EIQ, EIIQ, EIIIQ, and EIVQ at 1 Hz in Ba<sup>2+</sup>. A–C, Whole-cell Ba<sup>2+</sup> currents were recorded in the absence and presence of 50  $\mu$ M diltiazem using depolarizations to +10 mV from a holding potential of -60 mV, at a frequency of 1 Hz. Current measured in the presence of diltiazem followed equilibration of block by 50  $\mu$ M diltiazem at 0.05 Hz. Relative peak current (mean  $\pm$  S.E.) in each successive depolarizing pulse is plotted against pulse number (n = 4-8) in the absence (closed symbols) or presence (open symbols) of 50  $\mu$ M diltiazem for WT channels and the indicated mutants. Asterisk indicates that the fraction of current remaining at the end of the 20-pulse, 1-Hz train was significantly different from that of WT Ca<sub>v</sub>1.2 channels (p < 0.05). D, demonstration of differential effects of the EIVQ mutation on diltiazem block of Ca<sub>v</sub>1.2 at 0.05 and 1 Hz in single cells. Ba<sup>2+</sup> current in cells expressing either WT Ca<sub>v</sub>1.2 or the EIVQ mutant was brought to equilibrium as measured using stimulation at 0.05 Hz as described above. At time 0, 50  $\mu$ M diltiazem was perfused onto the cells. Approximately 7 s after the initiation of drug perfusion, a 20-pulse, 1-Hz train of depolarization was applied (as described above). The fraction of current blocked at the first depolarization after initiation of drug perfusion is virtually identical in the two cells, but subsequent development of block at 1 Hz is greatly reduced in EIVQ compared with WT Ca<sub>v</sub>1.2.

Hz. As shown in Fig. 4C, recovery of WT channels from inactivation in the absence of drug follows a biexponential time course, reflecting a fast and a slow time course of recovery from inactivation as reported previously for  $Ca_v 1.2$ . (Johnson et al., 1996). In the absence of diltiazem, the frac-

tion of current recovering with the fast time constant  $(f_{\rm F})$  was 0.61, with a fast  $(\tau_{\rm f})$  and slow  $(\tau_{\rm s})$  time constant of 0.235 s and 8.8 s, respectively. In the presence of 50  $\mu \rm M$  diltiazem, the fast phase of WT Ca\_v1.2 recovery is substantially slowed  $(\tau_{\rm f}=1.3~{\rm s},~f_{\rm F}=0.43)$  and the time constant of slow recovery is



Fig. 4. Biophysical properties of the mutant Ca, 1.2 channel EIVQ. A, current-voltage relationship for WT Ca, 1.2 and EIVQ channels. Peak amplitude of  $Ba^{2+}$  current elicited by a 100-ms depolarization to the indicated test potential from a holding potential of -60 mV is plotted against the test potential (mean  $\pm$  S.E., n = 4-8). B, voltage dependence of inactivation in WT Ca<sub>v</sub>1.2 and EIVQ channels. Peak  $Ba^{2+}$  current elicited by depolarization to + 10 mV immediately after 10-s conditioning pulses to the indicated potentials from a holding potential of -80 mV are plotted against the amplitude to -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV are plotted against the amplitude of -80 mV and -80 mV and -80 mV are plotted again and -80 mV and -80 mV are plotted again and -80 \text{ mV} and -80 mV and -80 mV and -80 mV and conditioning pulse voltage (mean  $\pm$  S.E., n = 3-8) in the presence or absence of 50  $\mu$ M diltiazem. A 200-s delay was incorporated between each conditioning voltage to allow full recovery of drug-bound channels. The data are fit (smooth lines, WT; broken lines, EIVQ) to the equation, relative current =  $1/(1 + \exp[(V - V_{1/2})/k])$ , where V is the conditioning potential,  $V_{1/2}$  is voltage at which half of the channels are inactivated, and k is a slope factor (potential required for an *e*-fold change). For WT Ca<sub>v</sub>1.2,  $V_{1/2}$  in the absence and presence of 50  $\mu$ M diltiazem was  $-19.5 \pm 1.6$  and  $-51.9 \pm 1.5$  mV, respectively. For EIVQ,  $V_{1/2}$  in the absence and presence of 50  $\mu$ M diltiazem was  $-16.1 \pm 2.4$  and  $-48.8 \pm 1.3$  mV, respectively. Values for k were -10.6 and -14.5 for WT Ca, 1.2 in the absence and presence of drug, respectively. For EIVQ, k values were -18.6 and -13.0 in the absence and presence of 50  $\mu$ M diltiazem, respectively. C and D, recovery of WT (C) and EIVQ (D) Ca<sub>v</sub>1.2 channels from inactivation in the presence and absence of 50  $\mu$ M diltiazem. Recovery from inactivation induced by a 1-s depolarization to +10 mV from a holding potential of -60 mV was measured using 50-ms test pulses to +10 mV after various recovery intervals (0.01, 0.1, 0.5, 1, 10, 20, and 30 s) at -60 mV. Representative traces are shown for both WT and EIVQ (C and D, upper panels). The time points between 300 and 900 ms of the depolarizing pulse are omitted, and the test pulses during the recovery phase are compressed for display purposes. The recovery time before each test pulse is indicated in seconds. The fraction of current recovered is plotted against the recovery interval (mean  $\pm$  S.E., n = 4) (C and D, lower panels). In both cases, the time course of recovery was well fitted with a double-exponential equation. For WT Ca<sub>v</sub>1.2 channels in the absence of diltiazem, the fast time constant of recovery ( $\tau_{\rm F}$ ) was 235 ms, the slow time constant of recovery ( $\tau_{\rm S}$ ) was 8.8 s, and the fraction of channels recovering with the fast time constant ( $f_{\rm F}$ ) was 0.61. In the presence of 50  $\mu$ M diltiazem,  $\tau_{\rm F} = 1.3$  s,  $\tau_{\rm S} = 9.1$  s, and  $f_{\rm F} = 0.43$ . For EIVQ in the absence of diltiazem,  $\tau_{\rm f} = 110$  ms,  $\tau_{\rm s} = 51.9$  s, and  $f_{\rm F} = 0.56$ . For EIVQ in the presence of 50  $\mu$ M diltiazem,  $\tau_{\rm F} = 1.1$  s,  $\tau_{\rm S} = 42.7$  s, and  $f_{\rm F} = 0.83$ .

virtually unchanged ( $\tau_{\rm s} = 9.1$  s). The recovery of EIVQ from inactivation in the absence of diltiazem was also biexponential ( $f_{\rm F} = 0.56$ ,  $\tau_{\rm f} = 0.11$  s,  $\tau_{\rm s} = 51.9$  s; Fig. 4D). However, the recovery of EIVQ from inactivation in 50  $\mu$ M diltiazem was much more rapid than in WT Ca<sub>v</sub>1.2. Rather than a marked difference in the fast time constant of recovery ( $\tau_{\rm F} = 1.1$  s;  $\tau_{\rm S} = 42.7$  s) compared with WT, the acceleration of recovery in EIVQ is likely due to an increase in the fraction of channels recovering with a fast time constant ( $f_{\rm F} = 0.83$ ). This large increase in  $f_{\rm F}$  leads to a "crossover" in the recovery time course, such that after 1 s, the EIVQ channel recovers from inactivation faster in the presence of diltiazem than in its absence.

We also examined the effect of the EQ mutations on diltiazem block rate (Fig. 5). WT and mutant channels were depolarized for 1 s to +10 mV from a holding potential of -60 mV, in the presence or absence of 50  $\mu$ M diltiazem. The resulting current traces (see Fig. 5A) were normalized to peak current, to facilitate comparison of the rate of current decay in the absence and presence of diltiazem. As shown in Fig. 5B, the rate of current inactivation in individual cells expressing either WT or EIVQ channels was fit to either single- or double-exponential equations. The time constants and fraction of channels inactivating with a given time constant are shown in Fig. 5B. Inactivation of WT channels was biexponential, both in the presence and absence of diltiazem. The fast time constant ( $\tau$ -fast) for WT channels was not different in the presence or absence of 50  $\mu$ M diltiazem  $(0.15 \pm 0.01 \text{ s} \text{ and } 0.12 \pm 0.02 \text{ s}, \text{respectively});$  however, the fraction of channels inactivating with the fast time constant was significantly increased in the presence of diltiazem. The slow time constant of WT channel inactivation was accelerated significantly in the presence of diltiazem ( $\tau$ -slow = 0.66 ± 0.03 s versus 0.42 ± 09 s).

In contrast, the inactivation of EIVQ channels followed a single-exponential time course (Fig. 5B) in the absence of diltiazem ( $\tau = 0.76 \pm 0.14$  s) but a double-exponential time course in the presence of diltiazem ( $\tau$ -fast = 0.04 ± 0.01 s). The slow time constant was significantly accelerated ( $\tau$ -slow = 0.28 ± 0.02 s) relative to the time constant of EIVQ channel inactivation in the absence of diltiazem. In addition, diltiazem also significantly increased the fraction of EIVQ channels inactivating with the slower time constant relative to EIVQ channels in the absence of drug. Thus, diltiazem blocks both depolarized WT and EIVQ channels by increasing the fraction of channels inactivating and by decreasing the slow time constant of inactivation.

 $Ca^{2+}$  Potentiation of Diltiazem Block. Diltiazem block of  $Ca_v 1.2$  is potentiated by  $Ca^{2+}$  (Lee and Tsien, 1983). This property is demonstrated in Fig. 6, top panel, for WT  $Ca_v 1.2$ . Using a concentration of diltiazem (50  $\mu$ M) near the  $IC_{50}$  for block of  $Ca_v 1.2$  in  $Ba^{2+}$ , it is clear that more channels are blocked when  $Ca^{2+}$ , rather than  $Ba^{2+}$ , is the charge carrier. To determine the molecular determinants of this modulation of diltiazem block by  $Ca^{2+}$ , we measured the fraction of current remaining after application of 50  $\mu$ M diltiazem to WT



**Fig. 5.** Time course of diltiazem block of depolarized channels. A, representative traces of Ba<sup>2+</sup> current through WT Ca<sub>v</sub>1.2 and EIVQ channels measured during a 1-s depolarization to +10 mV from a holding potential of -60 mV in the absence (control) or presence of 50  $\mu$ M diltiazem (normalized to peak current). B, analysis of inactivation in the absence (Con, filled bars) and the presence (Dilt, open bars) of 50  $\mu$ M diltiazem for Ca<sub>v</sub>1.2 (left panel) and EIVQ (right panel). In the absence of drug, Ca<sub>v</sub>1.2 inactivation is well fit by a double exponential [ $\tau$ -fast = 0.15 ± 0.01 s;  $\tau$ -slow = 0.66 ± 0.03 s; fraction fast ( $F_{\text{fast}}$ ) = 0.40 ± 0.02]. In the presence of 50  $\mu$ M diltiazem,  $\tau$ -fast is not changed (0.12 ± 0.02 s), but  $\tau$ -slow is accelerated (0.42 ± 0.09 s), and  $F_{\text{fast}}$  is increased (0.65 ± 0.03). In the absence of drug, EIVQ channel activation is well fit by a single-exponential function with a time constant similar to  $\tau$ -slow in WT channels ( $\tau$ -slow = 0.76 ± 0.14;  $F_{\text{slow}}$  = 0.50 ± 0.06). In the presence of 50  $\mu$ M diltiazem, EIVQ channel additional fast component of inactivation ( $\tau$ -fast = 0.04 ± 0.01 s;  $F_{\text{fast}}$  = 0.22 ± 0.03),  $\tau$ -slow is accelerated, and the fraction channel inactivating with the slow time course is increased ( $\tau$ -slow = 0.28 ± 0.02;  $F_{\text{slow}}$  = 0.77 ± 0.03). Results are means ± S.E. of individual fits (n = 4-6). #, not detected in EIVQ channels. Asterisks indicate significant differences between the indicated parameter in the presence of 50  $\mu$ M diltiazem (\*, p < 0.05; \*\*, p < 0.01; \*\*\*, p < 0.001).



Fig. 6. Ca<sup>2+</sup> potentiation of diltiazem block: contributions of EIII and EIV. Representative traces demonstrating the enhanced block of WT Ca, 1.2 channels by diltiazem in Ca<sup>2+</sup> compared with Ba<sup>2+</sup> (top panel). A, potentiation of diltiazem block at 0.05 Hz. Diltiazem (50  $\mu$ M) was  $Ca^{2}$ applied to cells expressing the indicated channels under voltage-clamp, as in Fig. 2, using either 10 mM  $Ba^{2+}$  or 10 mM  $Ca^{2+}$  as the charge carrier. The percentage of current remaining (mean  $\pm$  S.E.; n = 3-10) in can be used by the transferred for the transf significant differences between current remaining in Ba<sup>2+</sup> and Ca<sup>2-</sup> ⁺ for each channel (\*, p < 0.05; \*\*, p < 0.01; \*\*\*, p < 0.001) and plus signs indicate significant differences in current remaining in Ca<sup>2+</sup> between WT and mutant channels (++, p < 0.01). B and C, frequency-dependent block in Ca<sup>2+</sup>. Frequency-dependent block of the indicated channels by 50  $\mu$ M diltiazem was measured as described in Fig. 3 except that  $Ba^{2+}$  was replaced with 10 mM  $Ca^{2+}$  (open symbols). For comparison, inactivation during an identical stimulus train in the absence of drug is shown for each channel type tested (closed symbols). There was no significant difference in the fraction of current blocked after 20 pulses between WT and the EIQ and EIIQ mutants. Both EIIIQ and EIVQ channels were blocked to a significantly lesser extent than WT channels under these conditions. Data points are mean  $\pm$  S.E. (n = 3-5) (\*, p < 0.05).

and mutant  $Ca_1.2$  channels in  $Ba^{2+}$  and in  $Ca^{2+}$ . Diltiazem block was allowed to reach equilibrium using 100-ms depolarizing steps to +10 mV from a holding potential of -60 mV at 0.05 Hz. Figure 6A shows the averaged values for these measurements in WT Ca, 1.2. As expected, a greater fraction of WT current was blocked by 50  $\mu$ M diltiazem in Ca<sup>2+</sup> (~40% remaining) than in  $Ba^{2+}$  (~70% remaining) Similar potentiation was observed in Ca<sup>2+</sup> for the EIQ and EIIQ mutants as well. However, in both EIIIQ and EIVQ, the fraction of current blocked was not significantly different in  $Ba^{2+}$  and  $Ca^{2+},$  and was similar to the fraction of WT current blocked in  $Ba^{2+}.$  Similarly,  $Ca^{2+}$  potentiation of frequencydependent diltiazem block was not different from WT in EIQ and EIIQ, but was significantly reduced in EIVQ and EIIIQ (Fig. 6, B and C). Thus, Ca<sup>2+</sup> potentiation of both closedchannel and frequency-dependent block requires the conserved glutamate residues in the pore region of homologous domains III and IV.

We also examined the involvement of two other amino acid residues in the  $\mathrm{Ca}^{2+}$  potentiation of diltiazem block. The Phe residue (F1117) immediately adjacent to EIIIQ is required for Ca<sup>2+</sup> potentiation of DHP binding (Peterson and Catterall, 1995), and the C-terminal tail Ile 1627 is part of a  $Ca^{2+}/$ calmodulin binding site that mediates Ca<sup>2+</sup>-dependent inactivation (Peterson et al., 1999). Mutation of this Ile residue to Ala disrupts Ca<sup>2+</sup>-dependent inactivation (Zuhlke et al., 1999). Therefore, we examined  $Ca^{2+}$  potentiation of diltiazem block of both the F1117G and I1627A mutant channels (Fig. 7). Figure 7A shows that for both F1117G and I1627A, the fraction of channels blocked by diltiazem in  $Ba^{2+}$  and Ca<sup>2+</sup> at 0.05-Hz stimulation was not significantly different. However, for F1117G, neither the fraction of channels blocked by 50  $\mu$ M diltiazem in Ca<sup>2+</sup> nor the fraction of channels blocked by 50  $\mu$ M diltiazem in Ba<sup>2+</sup> is significantly different from the fraction of WT channels blocked by 50  $\mu$ M diltiazem in Ba<sup>2+</sup>. In the I1627A mutant, the fraction of channels blocked by 50  $\mu$ M diltiazem in Ba<sup>2+</sup> was not significantly different from the fraction of WT channels blocked by 50  $\mu$ M diltiazem in Ca<sup>2+</sup>. Thus, whereas F1117G disrupts Ca<sup>2+</sup> potentiation of diltiazem block of closed channels, the I1627A mutation potentiates diltiazem block of closed channels to the same extent as Ca<sup>2+</sup>. Neither F1117G nor I1627A differed from WT channels in the extent of frequency-dependent block at the end of a 20-pulse, 1-Hz train of depolarizations in Ba<sup>2+</sup> (Fig. 7B). Frequency-dependent diltiazem block of the I1627A mutant in Ba<sup>2+</sup> did, however, develop more rapidly than WT or F1117G channels, reaching equilibrium after fewer depolarizations. In Ca<sup>2+</sup>, frequency-dependent block of WT and I1627A channels by 50  $\mu$ M diltiazem was not different, although a 20-pulse train of depolarizations in the absence of diltiazem resulted in potentiation of  $\mathrm{Ca}^{2+}$  current as previously reported for I1627A (Zuhlke et al., 1999) (Fig. 7C). In contrast, frequency-dependent diltiazem block of F1117G in Ca<sup>2+</sup> was significantly reduced compared with WT channels. Thus, the F1117G mutation does not reduce the affinity of closed or inactivated channels for diltiazem in Ba<sup>2+</sup>, but does disrupt the Ca<sup>2+</sup> potentiation of diltiazem block of both closed and inactivated channels. In addition, the I1627A mutation increases the affinity of closed, but not inactivated, channels for diltiazem in Ba<sup>2+</sup>, mimicking Ca<sup>2+</sup> potentiation. However, I1627A does not alter the affinity of diltiazem for either closed or inactivated channels in  $Ca^{2+}$ .

**Inactivation in Ba<sup>2+</sup> and Ca<sup>2+</sup>.** To probe the mechanism that might underlie our observations on the modulation of diltiazem block, we examined the inactivation kinetics of all of the mutations studied in Ba<sup>2+</sup> and Ca<sup>2+</sup>. Figure 8A shows averaged current traces elicited by 1-s depolarizations to +10



Fig. 7.  $Ca^{2+}$  modulation of diltiazem block in F1117G and I1627A. A, diltiazem (50  $\mu$ M) block of WT, F1117G, and I1627A  $Ca_v1.2$  channels in 10 mM  $Ba^{2+}$  or 10 mM  $Ca^{2+}$  at 0.05 Hz. The percentage of current remaining (mean ± S.E., n = 3-10) in each case was: WT as in Fig. 6A, F1117G<sub>Ba</sub> = 59 ± 4, F1117G<sub>Ca</sub> = 68 ± 5, I1627A<sub>Ba</sub> = 42 ± 5, and I1627A<sub>Ca</sub> = 44 ± 6. Plus signs indicate a significant difference in the indicated value between WT and mutant channels. (++, p < 0.001). Frequency-dependent block (1 Hz) of F1117G and I1627A channels in  $Ba^{2+}$  (B) or  $Ca^{2+}$  (C) by 50  $\mu$ M diltiazem (open symbols). The protocol is as in Fig. 3A. For comparison, inactivation during an identical stimulus train in the absence of drug is shown for each channel type tested (closed symbols). Relative current (mean ± S.E., n = 3-6) is plotted versus pulse number. Asterisks indicate a significant difference in the relative current elicited by the 20th pulse in the train between WT  $Ca_v1.2$  channels and the indicated mutant (\*, p < 0.05).

mV from a holding potential of -60 mV for several cells expressing the WT, EIQ, EIIQ, EIIIQ, or EIVQ channels recorded in 10 mM  $Ba^{2+}$ . The extent of inactivation at the end of the 1-s depolarization is not significantly different among all five channels (Fig. 8D). When the same pulse protocol is applied to WT channels in 10 mM Ca<sup>2+</sup>, inactivation in the WT channel is markedly accelerated (i.e.,  $Ca^{2+}$ dependent inactivation is observed). However, in EIIQ, EIIIQ, and EIVQ channels, the extent of inactivation at the end of the 1-s depolarization is not increased in  $Ca^{2+}$  compared with Ba<sup>2+</sup> (i.e., Ca<sup>2+</sup>-dependent inactivation is disrupted) (Fig. 8, B and D). In the I1627A mutant,  $Ca^{2+}$ dependent inactivation is also disrupted (Fig. 8, C and D). However, with I1627A, the extent of inactivation in  $Ba^{2+}$  is markedly increased such that  $Ca^{2+}$  does not significantly increase it further. Finally, the extent of inactivation at the end of a 1-s depolarization observed in EIQ and F1117G is not different from WT Cav1.2 in Ba2+, and inactivation is also significantly accelerated in Ca<sup>2+</sup>. Thus, all of the mutant channels that we assayed except for EIQ and F1117G disrupted Ca<sup>2+</sup>-dependent inactivation. However, in the EIIQ, EIIIQ, and EIVQ mutants, this property was the result of slower inactivation in Ca<sup>2+</sup> than in WT channels, whereas in the I1627A mutant, it was the result of faster inactivation in  $Ba^{2+}$ .

Since the F1117G mutant is deficient in  $Ca^{2+}$  potentiation of diltiazem block but retains Ca<sup>2+</sup>-dependent inactivation that was indistinguishable from that of WT channels, we sought further evidence that F1117G may interact with Ca<sup>2+</sup> ions in the pore. The current traces in Fig. 9A were recorded from the same cell, expressing the F1117G mutant, in 10 mM  $Ba^{2+}$  and then in 10 mM  $Ca^{2+}$ . Interestingly, we found that the peak current amplitude elicited by depolarization to +10 mV was greater in Ca<sup>2+</sup> than in Ba<sup>2+</sup> for the F1117G mutant. In contrast, the WT Ca<sub>v</sub>1.2 channel exhibits a marked decrease in peak current when the extracellular solution is switched from 10 mM Ba<sup>2+</sup> to 10 mM Ca<sup>2+</sup> (Fig. 9C). Despite this apparent increase in permeability for  $Ca^{2+}$  relative to Ba<sup>2+</sup>, the F1117G mutant maintains robust Ca<sup>2+</sup>-dependent inactivation (Figs. 9A and 8D). The current-voltage relationship for F1117G in Ba<sup>2+</sup> versus Ca<sup>2+</sup> (Fig. 9B) shows that the increased permeability for  $Ca^{2+}$  over  $Ba^{2+}$  observed in this mutant is voltage-dependent, occurring only with depolarizations above -10 mV. In Fig. 9C, we compare the change in peak current amplitude, measured at +10 mV, upon switching from 10 mM Ba<sup>2+</sup> in the extracellular solution, to 10 mM Ca<sup>2+</sup> for WT, EIIQ, EIIIQ, EIVQ, F1117G, and I1627A channels. EIQ and EIIQ show decreases in peak current amplitude upon switching from Ba<sup>2+</sup> to Ca<sup>2+</sup>, similar to WT channels. However, peak current amplitude in EIIIQ, EIVQ, and F1117G channels increased upon switching from Ba<sup>2+</sup> to  $Ca^{2+}$ . Thus, the  $Ca^{2+}$  binding site formed by EIII, EIV, and F1117 not only modulates the affinity of closed and inactivated channels for diltiazem but also mediates the lower permeability of Ca<sup>2+</sup> relative to Ba<sup>2+</sup> in Ca<sub>v</sub>1.2 channels.

## Discussion

Selective Effect of EIVQ Mutation on Frequency-Dependent Block of  $Ca_v 1.2$  in  $Ba^{2+}$ . We have shown that amino acid residues in the pore region of  $Ca_v 1.2$  can modulate diltiazem block and that this modulation is distinct,

depending upon the permeant ion used in the experiments. None of the E to Q mutations significantly affected the  $IC_{50}$ for diltiazem block at low frequency (0.05 Hz) in 10 mM Ba<sup>2</sup> (Fig. 2). This result contrasts with a study using the PAA D888, in which the EIIIQ and EIVQ mutations increased the  $IC_{50}$  for D888 by 15- to 20-fold at a stimulation frequency of 0.10 Hz in 10 mM Ba<sup>2+</sup> (Hockerman et al., 1997a). Apparently, closed-channel block of Ca<sub>v</sub>1.2 by diltiazem involves interactions predominantly with amino acid residues from IIIS6 and IVS6 (Hering et al., 1996; Kraus et al., 1998; Hockerman et al., 2000) and does not require an interaction between the alkylamino group and particular pore glutamates as proposed for D888. The near-complete loss of frequency-dependent block by diltiazem in EIVQ in Ba<sup>2+</sup> (Fig. 3C) suggests a role for E1419 either in binding of the drug to the inactivated state of the channel or in modulation of the inactivation properties of the channel.

To distinguish these two possibilities, we investigated the mechanism by which this mutation exerted such a strong effect on frequency-dependent block in Ba<sup>2+</sup>. The voltage dependence of activation and inactivation for WT Ca, 1.2 and EIVQ was not greatly different. Moreover, the shift in  $V_{1/2}$ inactivation induced by 50  $\mu$ M diltiazem was not different for WT Ca<sub>v</sub>1.2 and EIVQ channels (Fig. 4B), suggesting that the EIVQ mutation did not disrupt the ability of diltiazem to bind to the inactivated state of the channel (Li et al., 1999). This characteristic of the EIVQ mutation contrasts with mutations in IVS5 of Ca, 1.2 that disrupted frequency-dependent block but also markedly disrupted voltage-dependent inactivation (Motoike et al., 1999; Bodi et al., 2002). Thus, neither the small changes in the voltage dependence of activation and inactivation nor changes in the ability of diltiazem to bind the inactivated state of EIVQ likely explain the marked loss of frequency-dependent block accumulation in EIVQ.

Recovery from inactivation in Cav1.2 is biexponential (Johnson et al., 1996; Kraus et al., 1998), and diltiazem slowed recovery of WT Ca<sub>v</sub>1.2 channels from inactivation by increasing  $\tau_{\rm F}$  and decreasing the fraction of channels recovering with the fast time constant  $(f_{\rm F})$  (Fig. 4C). The time constant for slow recovery was not different in the presence and absence of drug for WT Ca<sub>v</sub>1.2. Our results differ from those of Kraus et al. (1998), who found that diltiazem decreased  $f_{\rm F}$  but did not appreciably change either  $\tau_{\rm F}$  or  $\tau_{\rm S}$ . This discrepancy may be due to the differences in the  $\alpha_1$  subunits and expression system, since Kraus et al. (1998) used chimeric channels that were predominantly Ca<sub>v</sub>2.1, with L-type sequence inserted into IIIS6 and IVS6, expressed in Xenopus laevis oocytes. Our results with diltiazem are consistent with studies of PAA block of Ca<sub>v</sub>1.2, since PAA binding also decreases the fraction of channels recovering with the fast time constant (Johnson et al., 1996; Hering et al., 1997).

EIVQ channels also demonstrated biexponential recovery from inactivation (Fig. 4D). The  $f_{\rm F}$  for EIVQ in the absence of drug was similar to that of WT Ca<sub>v</sub>1.2, but  $\tau_{\rm F}$  was decreased, and  $\tau_{\rm S}$  was increased 5-fold. In the presence of 50  $\mu$ M diltiazem,  $\tau_{\rm F}$  increased and  $\tau_{\rm S}$  was not changed for EIVQ, as in WT channels. In contrast to WT channels, 50  $\mu$ M diltiazem increased  $f_{\rm F}$  in EIVQ channels such that, after 1s of recovery, EIVQ channels actually recovered faster in the presence of diltiazem than in the absence of the drug. This unexpected result suggests that, in Ba<sup>2+</sup>, diltiazem binding prevents EIVQ channels from entering the slowly recovering, inactivated state. A similar result was observed for PAA block with a mutant channel in which two amino acids in IIIS6 (corre-



Fig. 8. Inactivation properties of WT, and mutant channels in  $Ba^{2+}$  and  $Ca^{2+}$ . Averaged, normalized current traces (n = 2-8)for WT channels, and each of the indicated mutant channels elicited during a 1-s depolarization to +10 mV from a holding potential of -60 mV in extracellular solution containing 10 mM Ba<sup>2+</sup> (A) or 10 mM Ca<sup>2+</sup> (B) are shown. Note that the percentage of channels inactivated at the end of the 1-s pulse is not significantly different from WT for any of the pore mutants in Ba<sup>2+</sup>. C, representative current traces for the I1627A channel in 10 mM Ba<sup>2+</sup> and 10 mM Ca<sup>2+</sup> using the same pulse protocol as in A and B. D, summary of the percentage of channels inactivated at the end of a 1-s depolarization in  $Ba^{2+}$ or  $Ca^{2+}$  (FG = F1117G, IA = I1627A). Whereas WT Ca, 1.2, EIQ, and F1117G channels display significant increases in the percentage of channels inactivated in Ca<sup>2</sup> versus Ba<sup>2+</sup>, Ca<sup>2+</sup> does not significantly increase the percentage of channels inactivated in EIIQ, EIIIQ, EIVQ, or I1627A mutant channels (values are means ± S.E.). Asterisks indicate a significant difference between the percentage of channels inactivated at the end of a 1-s depolarization in 10 mM Ba<sup>2+</sup> versus 10 mM Ca<sup>2+</sup> (\*, p < 0.05; \*\*, p < 0.01; \*\*\*, p < 0.001).

sponding to amino acids I1163 and F1164) were changed to alanine (Hering et al., 1997).

The association of diltiazem with depolarized channels also suggests that diltiazem binds to and modulates the inactivated state of  $Ca_v 1.2$  channels. Figure 5B shows that diltiazem increases the rate of inactivation of WT channels by increasing the fraction inactivating with the fast time constant, and by accelerating the slow time constant of inactivation. In EIVQ channels, diltiazem induces a small fraction of channels to inactivate with a very rapid time constant not observed in the absence of drug (Fig. 5B). However, the major effect of diltiazem on EIVQ channels is the acceleration of the



**Fig. 9.** Permeability of  $Ca^{2+}$  relative to  $Ba^{2+}$  in WT and mutant  $Ca_v 1.2$  channels. A, current traces recorded from a single cell expressing the F1117G mutant recorded in 10 mM  $Ba^{2+}$  or 10 mM  $Ca^{2+}$ . Currents were elicited using a 1-s depolarization to +10 mV from a holding potential of -60 mV. Note the increase in peak current in  $Ca^{2+}$  relative to peak current in  $Ba^{2+}$ . B, current-voltage relationship of F1117G current in 10 mM  $Ca^{2+}$ . Currents at the indicated membrane potentials were recorded as described in the legend to Fig. 4. C, change in peak current amplitude measured at +10 mV from a holding potential of -60 mV when the extracellular solution is switched from 10 mM  $Ba^{2+}$  to 10 mM  $Ca^{2+}$ . The values shown are mean  $\pm$  S.E. (n = 4-12) for WT Ca\_1.2 channels and each of the indicated mutants.

slow time constant of inactivation and an increase in the fraction of channels inactivating with the slow time constant. Thus, it appears that diltiazem can bind to the inactivated state of EIVQ but induces rapid recovery from inactivation that disrupts the accumulation of frequency-dependent block in  $Ba^{2+}$ .

Ca2+ Modulation of Diltiazem Block. Our results indicate that, as for DHP binding, Ca<sup>2+</sup> potentiation of diltiazem block of closed channels is lost in the EIIIQ, EIVQ, and F1117G mutations, whereas it is retained in the EIQ and EIIQ mutants. These mutations do not affect the  $IC_{50}$  for diltiazem in Ba<sup>2+</sup> at 0.05 Hz but selectively abolish the increase in diltiazem potency observed for WT Ca, 1.2 channels in Ca<sup>2+</sup>. These data suggest that EIII and EIV may cooperate to form a Ca<sup>2+</sup> binding site. The role of F1117G may involve participation in  $Ca^{2+}$  binding via a  $\pi$ -cation interaction (Heginbotham and MacKinnon, 1992), or may involve the transduction of a conformational change to the diltiazem binding site. However, the observation that Ca<sup>2+</sup> permeability relative to  $Ba^{2+}$  is altered in F1117G (Fig. 9) argues that F1117 likely interacts directly with Ca<sup>2+</sup> ions in the pore. The extent of frequency-dependent block accumulation in  $Ca_v 1.2$  is also greater in  $Ca^{2+}$  than in  $Ba^{2+}$ (Figs. 6 and 7). As we observed for closed-channel block, the  $Ca^{2+}$ potentiation of frequency-dependent diltiazem block is retained in EIQ and EIIQ but is lost in the EIIIQ, EIVQ, and F1117G mutants. In sharp contrast to our observations in Ba<sup>2+</sup>, frequency-dependent diltiazem block of the EIVQ mutant in Ca<sup>2+</sup> was similar to that of EIIIQ. This result suggests that Ca<sup>2+</sup> binding in the pore of the EIVQ channel prevents the diltiazem-induced acceleration of recovery from inactivation observed in  $Ba^{2+}$ .

In contrast to all the other mutant channels used in this study, the I1627A mutation is not in the putative pore region of the channel but, rather, in the intracellular C-terminal tail. However, this mutation potentiated diltiazem block of closed channels in Ba<sup>2+</sup> compared with WT channels, to an extent similar to that in Ca<sup>2+</sup>, such that the extent of closedchannel diltiazem block in I1627A was not further increased in Ca<sup>2+</sup>. Frequency-dependent diltiazem block of I1627A was not different from WT in either Ba<sup>2+</sup> or Ca<sup>2+</sup>. To understand this observation, we examined the inactivation rates of the channels used in this study in  $Ba^{2+}$  and  $Ca^{2+}$  by measuring the percentage of peak current inactivated at the end of a 1-s depolarization. We found that all of the mutations except I1627A were not different from WT channels in the percentage of inactivated channels in  $Ba^{2+}$ . In the I1627A mutation, inactivation was accelerated in  $Ba^{2+}$  such that  $Ca^{2+}$  did not appreciably increase the percentage of inactivated channels at the end of a 1-s depolarization. The  $V_{1/2}$  inactivation of I1627A in 10 mM  $Ba^{2+}$  (measured as described for Fig. 4B) is not different from that of WT Ca. 1.2 (N. Dilmac and G. H. Hockerman, unpublished data). Thus, it appears that the conformational change that induces faster inactivation of I1627A may also increase the affinity of diltiazem for closed, but not inactivated, I1627A channels. We have observed that this acceleration of inactivation in  $Ba^{2+}$  of the I1627A mutation is  $\beta$  subunit-dependent. In addition to the  $\beta_{1b}$  subunit used in this study, we observed similar acceleration with the I1627A subunit coexpressed with the  $\beta_3$  but not the  $\beta_2$  subunit (G. H. Hockerman and N. Dilmac, unpublished data; also see Peterson et al., 1999). A recent study that examined

the modulation of (-)-gallopamil (a PAA) block of Ca. 1.2 by different  $\beta$  subunits (Sokolov et al., 2001) concluded that channels with an accelerated voltage-dependent inactivation rate were also more sensitive to gallopamil block. Furthermore, Sokolov et al. (2001) proposed that PAA block of Ca, 1.2 depends upon the fast voltage-dependent inactivation observed in Ba<sup>2+</sup> and not the further acceleration of inactivation observed in Ca<sup>2+</sup>. Thus, our data fit well with the model of Sokolov et al. (2001), since the main effect of the I1627A mutation is to shift >80% of channels into the fast inactivating mode in Ba<sup>2+</sup> (i.e., the fast voltage-dependent inactivation of Sokolov et al., 2001; Fig. 8C). Comparing the extent of inactivation in  $Ba^{2+}$  and  $Ca^{2+}$  (Fig. 8), it is clear that EIIQ, EIIIQ, and EIVQ are all deficient in  $Ca^{2+}$ - dependent inactivation compared with WT. However, since the EIIQ mutant shows normal Ca<sup>2+</sup> potentiation of both closed-channel and frequency-dependent diltiazem block, we conclude that Ca<sup>2+</sup>dependent inactivation does not play a critical role in the potentiation of diltiazem block by Ca<sup>2+</sup>. Furthermore, our results suggest that Ca<sup>2+</sup>-dependent inactivation of Ca<sub>v</sub>1.2 may involve Ca<sup>2+</sup> binding in the pore that is modulated by conformational changes in the C-terminal tail.

In conclusion, our results support the notion that Ca<sup>2+</sup> potentiates diltiazem block of both closed and inactivated Ca. 1.2 channels via a Ca<sup>2+</sup> binding site composed of F1117, E1118, and E1419. Thus, our results suggest that diltiazem binds to the same site between transmembrane segments IIIS6 and IVS6 in both the closed and inactivated states, with the inactivated state having a higher affinity for diltiazem. In Ba<sup>2+</sup>, the disruption of frequency-dependent diltiazem block in the mutant EIVQ results from a diltiazem-induced shift of channels into a rapidly recovering, inactivated state. Furthermore, it appears that Ca<sup>2+</sup>-dependent inactivation per se is not required for Ca<sup>2+</sup> potentiation of diltiazem. However, the same conformational change in I1627A that accelerates inactivation in Ba<sup>2+</sup> increases the affinity of the closed channel for diltiazem. It will be interesting to determine whether Ca<sup>2+</sup> potentiation of block of Ca<sub>v</sub>1.2 by PAAs such as verapamil occurs via the same or a distinct mechanism.

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