The $I_{Ks}$ Ion Channel Activator Mefenamic Acid Requires KCNE1 and Modulates Channel Gating in a Subunit-Dependent Manner

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Received July 19, 2019; accepted November 4, 2019

ABSTRACT

The pairing of KCNQ1 and KCNE1 subunits together mediates the cardiac slow delayed rectifier current ($I_{Ks}$), which is partly responsible for cardiomyocyte repolarization and physiologic shortening of the cardiac action potential. Mefenamic acid, a nonsteroidal anti-inflammatory drug, has been identified as an $I_{Ks}$ activator. Here, we provide a biophysical and pharmacological characterization of mefenamic acid’s effect on $I_{Ks}$. Using whole-cell patch clamp, we show that mefenamic acid enhances $I_{Ks}$ activity in both a dose- and stoichiometry-dependent fashion by changing the slowly activating and deactivating $I_{Ks}$ current into an almost linear current with instantaneous onset and slowed tail current decay, sensitive to the $I_{Ks}$ blocker (3R,4S)-(+)-N-[3-hydroxy-2,2-dimethyl-6-(4,4,4-trifluorobutoxy) chroman-4-yl]-N-methylmethanesulfonamide (HMR1556). Both single channels, which reveal no change in the maximum conductance, and whole-cell studies, which reveal a dramatically altered conductance-voltage relationship despite increasingly longer interpulse intervals, suggest mefenamic acid decreases the voltage sensitivity of the $I_{Ks}$ channel and shifts channel gating kinetics toward more negative potentials. Modeling studies revealed that changes in voltage sensor activation kinetics are sufficient to reproduce the dose and frequency dependence of mefenamic acid action on $I_{Ks}$ channels. Mutational analysis showed that mefenamic acid’s effect on $I_{Ks}$ required residue K41 and potentially other surrounding residues on the extracellular surface of KCNE1, and explains why the KCNQ1 channel alone is insensitive to up to 1 mM mefenamic acid. Given that mefenamic acid can enhance all $I_{Ks}$ channel complexes containing different ratios of KCNQ1 to KCNE1, it may provide a promising therapeutic approach to treating life-threatening cardiac arrhythmia syndromes.

SIGNIFICANCE STATEMENT

The channels which generate the cardiac slow delayed rectifier K+ current ($I_{Ks}$) are composed of KCNQ1 and KCNE1 subunits. Due to the critical role played by $I_{Ks}$ in heartbeat regulation, enhancing $I_{Ks}$ current has been identified as a promising therapeutic strategy to treat various heart rhythm diseases. Most $I_{Ks}$ activators, unfortunately, only work on KCNQ1 alone and not the physiologically relevant $I_{Ks}$ channel. We have demonstrated that mefenamic acid can enhance $I_{Ks}$ in a dose- and stoichiometry-dependent fashion, regulated by its interactions with KCNE1.

Introduction

The potassium voltage-gated KCNQ channel subfamily is composed of five known isoforms, KCNQ1–5 (Abbott, 2014). Expression of the first isoform, KCNQ1, has been detected throughout the body, including in the heart, stomach, and ear (Linn et al., 2015). When by itself, KCNQ1 produces a fast activating and deactivating current that has not yet been found to underlie any specific endogenous currents in the body (Abbott, 2014). KCNQ1, however, also coassembles with several $\beta$-subunits, KCNE1–5, which modulate KCNQ1 current kinetics (Bendahhou et al., 2005; Manderfield and George, 2008; Eldstrom and Fedida, 2011). In the heart, the coassembly of KCNQ1 with KCNE1, and perhaps other KCNE subunits, produces a slowly activating and deactivating cardiac delayed rectifier K+ current ($I_{Ks}$), which contributes significantly to cardiac repolarization (Sanguinetti et al., 1996; Lundquist et al., 2005).

There is no general agreement on the stoichiometric ratio of KCNE1 to KCNQ1 subunits underlying $I_{Ks}$, either in vivo or in heterologous in vitro expression systems (Morin and Kobertz, 2008; Nakajo et al., 2010; Plant et al., 2014; Murray et al., 2016), although we know a variable stoichiometry of 4:1 up to 4:4 is possible (Murray et al., 2016). Given that the kinetics of $I_{Ks}$ are greatly affected by the number of KCNE1 subunits, great flexibility in the expressed physiologic and pharmacological properties of $I_{Ks}$ channel complexes is expected from a variable stoichiometry.

This research was funded by the Natural Sciences and Engineering Research Council of Canada [Grant RGPIN-2016-04242], the Canadian Institutes of Health Research [Grant FJT-156181], and the Heart and Stroke Foundation of Canada [Grant G17-0018392] (grants to D.F.). Y.W. holds a Canada Graduate scholarship from a Canadian Institutes of Health Research [Grant PJT-156181], and the Heart and Stroke Foundation of Canada [Grant RGPIN-2016-05422], the Canadian Institutes of Health Research [Grant G17-0018392] (grants to D.F.). Y.W. holds a Canada Graduate scholarship. This article has supplemental material available at molpharm.aspetjournals.org.

ABBREVIATIONS: DIDS, 4,4′-dithiolicynanoato-2,2′-stilbendisulfonic acid; ECmax, maximal effective concentration; G-V, conductance-voltage relationship; HMR1556, (3R,4S)-(+)-N-[3-hydroxy-2,2-dimethyl-6-(4,4,4-trifluorobutoxy) chroman-4-yl]-N-methylmethanesulfonamide; $I_{Ks}$, cardiac slow delayed rectifier K+ current; $I_{\text{max}}$ – $I_{\text{min}}$, peak to steady-state difference currents; $I_{\text{min}}$, minimum current amplitude; I-V, current-voltage relationship; $k$, slope factor; LM, Iltk-mouse fibroblast; LGTS, long QT syndrome; $nH$, Hill coefficient; $V_{1/2}$, voltage at half-maximal activation; VS, voltage sensor; WT, wild type.
The complex also has clinical importance in disease syndromes, including cardiac arrhythmia, with the severity ranging from syncope to sudden death (Saplwaski et al., 2000). Approximately 50% of the mutations seen in long QT syndrome (LQTS) patients are in the KCNQ1 subunit (LQTS type 1) (Hedley et al., 2009), with mutations in KCNE1 causing LQTS type 5.

Activators of $I_{\text{Ks}}$ that can act on the relevant saturated and unsaturated $I_{\text{Ks}}$ complexes are of particular interest, as they may have therapeutic potential in the treatment of LQTS types 1 and 5. To date, although several activators have been reported, some are only effective on KCNQ1 alone with limited efficacy on $I_{\text{Ks}}$. These include ML-277, zinc pyrithione, and L-364,373 (Magyar et al., 2006; Gao et al., 2008; Yu et al., 2013). The known activators of $I_{\text{Ks}}$ are phenylboronic acid (Mruk and Kobertz, 2009) and hexachlorophene (Abitbol et al., 1999), and fatty acids such as lauric acid (Doolan et al., 2002). Phenylboronic acid and hexachlorophene have been shown to increase both KCNQ1 alone and $I_{\text{Ks}}$ current amplitudes, although they are more potent on $I_{\text{Ks}}$ (Mruk and Kobertz, 2009; Zheng et al., 2012). In contrast, lauric acid, DIDS, and 4-acetamido-4'-isothiocyanostilbene-2,2'-disulfonic acid have been shown to only increase $I_{\text{Ks}}$ current (Abitbol et al., 1999; Doolan et al., 2002).

The fenamate mefenamic acid (Fig. 1A) is a nonsteroidal anti-inflammatory drug primarily prescribed to treat menstrual pains (https://www.accessdata.fda.gov/drugsatfda_docs/label/2008/015034s040lbl.pdf). Originally identified as a chloride channel blocker, mefenamic acid has since been shown in various expression systems (Xenopus oocytes, rat mesenteric arteries, canine ventricular myocytes, guinea pig ventricular myocytes, CHO cells, and COS-7 cells) to increase mammalian $I_{\text{Ks}}$ current amplitudes as well as produce a variable amount of instantaneous current and inhibit tail current decay (Busch et al., 1994; Abitbol et al., 1999; Unsöld et al., 2000; Magyar et al., 2006; Toyoda et al., 2006; Chada et al., 2012).

In the present study, using transiently expressed human $I_{\text{Ks}}$ in mammalian cells, we have carried out a more complete biophysical characterization of the effects of mefenamic acid than has been attempted to date. We show that mefenamic acid has a minimal effect on KCNQ1 in the absence of KCNE1 and have quantified drug concentration- and rate-dependent changes in the $I_{\text{Ks}}$ current waveforms, the conductance-voltage relationship, and single-channel conductance and kinetics. Because the stoichiometry of $I_{\text{Ks}}$ may vary and affect its pharmacology and current kinetics (Nakajo et al., 2010; Murray et al., 2016), we have analyzed the dependence of mefenamic acid actions on the stoichiometry of $I_{\text{Ks}}$ channel complexes. Last, through mutational analysis, we identify a specific regulatory site for mefenamic acid on KCNE1. The results suggest that residue K41 on KCNE1 is of particular importance in mediating the effect mefenamic acid has on $I_{\text{Ks}}$.

### Materials and Methods

#### Solutions and Drugs

Unless otherwise stated, all drugs and chemicals used to make solutions were obtained from Sigma-Aldrich (Mississauga, ON, Canada). The control bath solution for whole-cell experiments contained 135 mM NaCl, 5 mM KCl, 1 mM MgCl₂, 2.8 mM NaAcetate, and 10 mM HEPES, pH 7.4 with NaOH. The pipette solution for single-channel experiments contained 6 mM NaCl, 129 mM 2-(4-morpholino)-ethane sulfonic acid, 1 mM MgCl₂, 5 mM KCl, and 10 mM HEPES, pH 7.4 with NaOH. Mefenamic acid and HMR1556 (3R,4S)-(-)-N-[3-hydroxy-2,2-dimethyl-6-(4,4,4-trifluorobutyryl)] chroman-4-yl]-N-methylthanesulphonamide; Tocris Bioscience, Oakville, ON, Canada) were prepared as stock solutions (50, 200, or 500 mM for mefenamic acid and 2 mM for HMR1556) dissolved in 100% dimethylsulfoxide. Stock mefenamic acid solutions were diluted in control whole-cell bath solution to obtain final mefenamic acid concentrations of 10, 30, 100, 300, 500 μM, or 1 mM, which were perfused onto mammalian cells for whole-cell experiments. Stock HMR1556 and mefenamic acid solutions were pipetted directly into the chamber to obtain a final HMR1556 concentration of 1 μM for whole-cell experiments or a final mefenamic acid concentration of 100 μM for single-channel experiments. Concentrations of dimethylsulfoxide in final bath solutions never exceeded 0.2% (v/v). The maximum concentration of mefenamic acid (1 mM) lowered the pH of the final bath solution by 0.15 ± 0.02 (n = 3). This was not corrected.

#### Constructs, Cell Culture, and Transfection

$I_{\text{Ks}}$ is generally understood to be functionally composed of combinations of KCNQ1 and KCNE1 subunits. The stoichiometry of the two subunits may be variable in heterologous expression systems (Murray et al., 2016) and may also vary in vivo (Dvir et al., 2014). In the initial experiments (Figs. 1–4), the initial stoichiometric ratio of KCNQ1:KCNE1 was set at the maximum, 4:4. This was achieved through transfection of a linked KCNQ1 and KCNE1 cDNA (Murray et al., 2016), which is expected to assemble as a tetramer with four KCNQ1 and four KCNE1 subunits. For simplicity, this will be denoted as EQ. In later experiments where the ratio was varied (Fig. 5), cells were transfected with KCNQ1 without KCNE1 (Q1), linked constructs containing one KCNE1 linked with two KCNQ1s (EQQ, expected to assemble in a 2:4 ratio), or one KCNE1 linked with four KCNQ1s (EQQQQ, expected to assemble in a 1:4 ratio). EQ, EQQ, and EQQQQ constructs were generated as previously described (Murray et al., 2016). In all cases, we consider the currents that result from different combinations of KCNQ1 and KCNE1 (except KCNQ1 alone, Q1) to be $I_{\text{Ks}}$, and we use this name interchangeably with the constructs themselves.

tsa201 transformed human embryonic kidney 293 or ltk- mouse fibroblast cells were cultured in modified Eagle’s medium supplemented with 10% fetal calf serum and 100 μg/ml penicillin, 100 μg/ml streptomycin, and 0.25 μg/ml amphotericin and plated for whole-cell and single-channel experiments, as previously described (Murray et al., 2016; Westhoff et al., 2019). Cells were transiently transfected with: 1) GFP-tagged Q1 (Q1-GFP); 2) EQ, EQQ, or EQQQQ and GFP in a 1.5–2.5:1.0–μg ratio; or 3) mutant KCNQ1 and Q1-GFP in a 4.5–6:0. 1.5-μg ratio using Lipofectamine2000 (Thermo Fisher Scientific, Waltham, MA) as per the manufacturer’s protocol. All KCNE1 mutations were generated using site-directed mutagenesis and Pfu Turbo (Agilent Technologies, Santa Clara, CA) followed by sequence confirmation of all mutations. Whole-cell and single-channel experiments were conducted 24–48 hours post transfection.

#### Electrophysiology

An Axopatch 200B amplifier, Digidata 1440A (whole-cell experiments), or Digidata 1200A (single-channel experiments) and pClamp 9 or 11 software (all from Molecular Devices, San Jose, CA) were used to conduct all experiments. For whole-cell experiments, electrodes ranging from 1- to 3-MΩ resistance were first put in from thin-walled borosilicate glass (Sutter Instrument, Novato, CA) using a linear multistage electrode puller (Sutter Instrument) and then fire polished.
before use. Series resistance compensation of 70%–80% was used for all whole-cell experiments. Data were sampled at 20 kHz and filtered at 5 kHz during acquisition. Electrodes for single-channel experiments were pulled from thick-walled borosilicate glass (Sutter Instrument) and fire polished to resistances between 40 and 60 MΩ. Before recording, electrodes were coated with Sylgard (Dow Corning, Midland, MI). Current records were sampled at 10 kHz; low-pass filtered at 2 kHz at acquisition using a −3-dB, four-pole Bessel filter; and digitally filtered at 200 Hz for presentation and analysis (Werry et al., 2013; Eldstrom et al., 2015; Murray et al., 2016; Thompson et al., 2017; Westhoff et al., 2017). For voltage clamp protocols, interpulse intervals refer to start-to-start times between sweeps.

**Data Analysis**

GraphPad Prism 8.1.1 software (GraphPad Software) was used to analyze all data. Where applicable, one-way ANOVA followed by the Bonferroni multiple comparison post hoc test was used to determine statistical significance. A P value less than 0.05 was considered statistically significant. All data in the figures are shown as means ± S.E.M. and reported in the tables as means ± S.D.

**Whole-Cell Experiments.** Mefenamic acid dose-response diary plots (Fig. 1, B and C, right panels) and curves (Figs. 2, A and B; Fig. 5D) were obtained from measurement of the activating $I_{KS}$ current. Specifically, the peak to steady-state difference currents ($I_{max} - I_{min}$) were calculated by subtracting the minimum amplitude of the activating current ($I_{min}$) from the peak amplitude of the activating current ($I_{max}$). This value was then plotted against either the corresponding sweep number or log$_{10}$ concentration of mefenamic acid (Fig. 1; Fig. 2A). Where applicable, the difference current in mefenamic acid was normalized to the maximum control (in the absence of mefenamic acid) difference current and subtracted from 1.0 to obtain the “normalized response,” which was plotted against the

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**Fig. 1.** Mefenamic acid increases EQ current expressed in mammalian cells. (A) Molecular structure of mefenamic acid. Nitrogen and oxygen atoms are depicted in blue and red, respectively. (B) Response of an untransfected cell to mefenamic acid (Mef). Left panel shows representative current traces in the absence of mefenamic acid (control; black) and after the addition of 10 μM to 1 mM mefenamic acid as indicated (gray). Right panel shows $I_{max} - I_{min}$ versus sweep number during the addition of different concentrations of mefenamic acid (solid gray lines). (C) Dose response of EQ currents to mefenamic acid. Left panel shows currents in control (black) and in response (gray) to different concentrations of mefenamic acid (solid gray lines). (C) Dose response of EQ currents to mefenamic acid. Left panel shows currents in control (black) and in response (gray) to different concentrations of mefenamic acid. Right panel shows the data in a diary plot. (D) Response of EQ to 1 μM HMR1556 in the presence of mefenamic acid (100 μM mefenamic acid preincubation for approximately 30 minutes; data not shown). Complete block by HMR1556 is indicated by the red arrows. Left panel shows 100 μM mefenamic acid alone (black) and in response to 1 μM HMR1556 (gray). Right panel shows the data in an initial peak amplitude versus sweep number diary plot. The solid gray line indicates addition of HMR1556. All currents (B–D) were obtained by pulsing to +60 mV for 4 seconds followed by a pulse to −40 mV for 0.9 seconds. The interpulse interval was 15 seconds. Holding potential was −80 mV. Dotted baselines denote the zero-current level.
corresponding log_{10} concentration of mefenamic acid (Fig. 2B; Fig. 5D; Fig. 8B). Dose- and normalized-response curves were fit with a specific binding equation to obtain the EC_{50} and Hill coefficients (n^H) as in Figs. 2, A and B, 5D, and 8B. The HMR1556 response diary plot in the presence of mefenamic acid (Fig. 1D, right panel) was obtained by plotting the initial peak current amplitude against the corresponding sweep number. All current-voltage (I-V) and conductance-voltage (G-V) plots in Figs. 3, 5, and 7 and Supplemental Fig. 1 were obtained from the normalized peak of the 4-second depolarizing pulses (U_{peak} and normalized peak of the initial tail current G/G_{max}, respectively, and plotted against the corresponding voltage. G-V plots were fitted with a Boltzmann sigmoid equation to obtain the voltage at half-maximal activation (V_{1/2}) and slope (θ) values (Tables 1 and 2). In the case of the mutant EQ I_{ks}, the change in V_{1/2} of activation (ΔV_{1/2} = V_{1/2} in the presence of mefenamic acid − V_{1/2} control) was further determined (Fig. 7D; Table 2). In some cells, G-V relationships in the presence of mefenamic acid for wild-type (WT) EQ, K41R, and G40C were essentially linear, and consequently, the V_{1/2} of activation was read from the plots and included in the calculations of the mean values in Tables 1 and 2. All deactivation traces in Fig. 4, A and B were fitted with a single exponential equation to obtain the time constants of deactivation (τ_{deact}), which were plotted against the corresponding membrane potential (Fig. 4C).

**Modeling**

Markov modeling in Fig. 8 was carried out using the IonChannel-Lab software (Santiago-Castillo et al., 2010) incorporating Q-matrix solutions to the differential equations defining the kinetic behavior of rate transitions (Colquhoun and Hawkes, 1995). A balanced model (Zaydman et al., 2014) as revised by Westhoff et al. (2019) was used for simulations, using a 4:4 KCNQ1:KCNE1 stoichiometry. In this model, each voltage sensor (VS) is assumed to undergo two activating transitions, to an intermediate and then activated conformation. Pore subconductance opening can occur as soon as each VS is fully activated conformations. For full model exposition and rates, see Westhoff et al. (2019). To simulate the action of mefenamic acid on I_{ks}, currents, the intrinsic rates of forward VS transitions at 0 mV, between resting and intermediate states (k_{R0}) and between intermediate and activated states (k_{IA0}), were multiplied by the drug concentration (D, micromolar) or log_{10} [D], respectively. This difference reflected the fact that an extreme hyperpolarization of the G-V relation and facilitation of initial steps of activation were caused by mefenamic acid at higher concentrations or rates of activation (Fig. 3), and thus, a greater acceleration of the first VS transition was required to simulate experimental data.

**Results**

**Mefenamic Acid Increases EQ I_{ks} Current Expressed in Mammalian Cells.** Initially, to ensure that mefenamic acid (Fig. 1A) had no effect on the endogenous currents in ltk mouse fibroblast (LM) cells, different concentrations of mefenamic acid (10, 30, 100, 300, 500 μM, and 1 mM) were perfused onto untransfected LM cells (Fig. 1B). At all concentrations of mefenamic acid, no changes were observed in the waveform (Fig. 1B, left panel) or I_{max} − I_{min} (Fig. 1B, right panel) compared with untreated cells (control). A single concentration of mefenamic acid (100 μM) has previously been shown to enhance I_{ks} activity leading to a variable amount of instantaneous current and inhibition of tail current decay in Xenopus oocytes (Busch et al., 1994, 1997) and various mammalian cells (Unsöld et al., 2000; Magyar et al., 2006; Toyoda et al., 2006). The concentration dependence of this enhancement of I_{ks} activity using different concentrations of mefenamic acid perfused onto LM cells transiently transfected with EQ is shown in Fig. 1C. With increasing concentrations of mefenamic acid, the control sigmoidal waveform (indicative of slow activation kinetics) was transformed into an almost linear waveform with significant instantaneous current (Fig. 1C, left panel). The corresponding difference current diary plot of increasing concentrations of mefenamic acid shows the gradual transformation to an instantaneous current over time (Fig. 1C, right panel). Consistent with previous findings, the decay in tail current seen in the control was also inhibited in a dose-dependent manner with increasing concentrations of mefenamic acid (Fig. 1C, left panel).

To confirm that this instantaneous current was produced by mefenamic acid enhancement of I_{ks} activity specifically, we used the I_{ks} blocker HMR1556 (Gögelein et al., 2000). In these experiments, LM cells transiently transfected with EQ were first preincubated in 100 μM mefenamic acid for approximately 30 minutes (data not shown); then, when an I_{ks} positive cell was identified, 1 μM HMR1556 was applied to the bath (Fig. 1D). As is evident in the representative traces (Fig. 1D, left panel) and diary plot (Fig. 1D, right panel), following HMR1556 treatment, the instantaneous current amplitude decreased over time to that of endogenous current amplitudes, suggesting that the instantaneous current was, in fact, flowing through I_{ks} channels.

**Mefenamic Acid Dose-Response Curves for EQ I_{ks}**

Increasing concentrations of mefenamic acid gradually
transformed the sigmoidal activation waveform of EQ $I_{Ks}$ into an almost linear waveform, and this change was quantified by measuring peak to steady-state difference currents (Fig. 2A, upper panel; see Materials and Methods). More specifically, difference currents were calculated by subtracting the initial amplitude of the activating current ($I_{\text{min}}$) from the peak amplitude of the activating current ($I_{\text{max}}$) and plotting the result against the corresponding log concentration of mefenamic acid (Fig. 2A). Normalized-response relationships for EQ were obtained by transformation of $I_{\text{max}} - I_{\text{min}}$ data (see Materials and Methods) and fit with a specific binding equation (Fig. 2B). The $EC_{50}$ and Hill coefficient ($n^H$) for mefenamic acid were 60 $\mu$M and 0.49, respectively. To ensure consistent results and to allow for comparison with previous literature, all subsequent experiments to characterize the gating properties, subunit stoichiometry, and specific regulatory residues were conducted using 100 $\mu$M mefenamic acid, unless otherwise stated.

**Mefenamic Acid Rate-Dependently Hyperpolarizes EQ $I_{Ks}$ I-V and G-V Relationships.** To investigate what happens to EQ $I_{Ks}$ I-V and G-V relationships following treatment with 100 $\mu$M mefenamic acid, a 4-second activation protocol with varying interpulse intervals was used (Fig. 3). Fig. 3A shows representative waveforms of EQ $I_{Ks}$ in both the absence (control; upper panel) and presence of mefenamic acid (lower panel), and shows the characteristic delay of current activation in control and the appearance of an instantaneous current with mefenamic acid treatment. The corresponding I-V (Fig. 3B) and G-V relationships (Fig. 3C) were obtained by plotting the normalized peak amplitudes at the end of the 4-second depolarizing pulses ($I/I_{\text{max}}$) or the normalized peak of the initial tail current ($G/G_{\text{max}}$), respectively, against the corresponding voltage. During exposure to mefenamic acid, the I-V relationship became more linear and hyperpolarized (Fig. 3B). The G-V relationship following treatment with mefenamic acid was also hyperpolarized (control: $V_{1/2} = 24.0$ mV, $k = 24.0$ mV; mefenamic acid: $V_{1/2} = -2.4$ mV, $k = 65.4$ mV; see Table 1). $V_{1/2}$ values for mefenamic acid compared with control were significantly different at all intervals (Table 1).

**Fig. 3.** Mefenamic acid rate-dependently hyperpolarizes EQ I-V and G-V relationships. Currents were obtained using a 4-second step protocol with pulses from $-150$ to $+100$ mV, followed by a repolarizing step to $-40$ mV for 1 second. (A) Currents are shown for EQ in the absence (control; interpulse interval was 15 seconds; upper panel) and presence of 100 $\mu$M mefenamic acid (Mef; interpulse interval was 30 seconds; lower panel). (B) I-V plots of current at the end of the 4-second depolarizing pulses in control (black triangles) and mefenamic acid (gray triangles); data from (A). (C) G-V relationships obtained from peak initial tail currents in control ($V_{1/2} = 24.0$ mV; $k = 24.0$ mV; black circles) and mefenamic acid ($V_{1/2} = -2.4$ mV; $k = 65.4$ mV; gray circles); data from (A). (D and E) Effect of different interpulse intervals on I-V (triangles) and G-V (circles) plots in control and mefenamic acid. Intervals were 7 ($n = 1$; orange) and 15 seconds ($n = 4$; black) in control, and 7 (red), 15 (green), 20 (blue), or 30 seconds (purple) in mefenamic acid ($n = 4$–6). For G-V plots, Boltzmann fits were for 7 ($V_{1/2} = 17.8$ mV; $k = 12.9$ mV) and 15 seconds ($V_{1/2} = 23.9$ mV; $k = 20.4$ mV) in control, and for 7–$V_{1/2} = -108$ mV; $k = 6.9$ mV), 15 ($V_{1/2} = -86.5$ mV; $k = 23.2$ mV), 20–$V_{1/2} = -80.5$ mV; $k = 55.3$ mV), and 30-second intervals ($V_{1/2} = -13.1$ mV; $k = 57.2$ mV) in mefenamic acid (see Table 1), $V_{1/2}$ values for mefenamic acid compared with control were significantly different at all intervals (Table 1).
significantly different from control (Table 1). The slope of the G-V relationship was significantly decreased when the interpulse interval was either 20 (53.3 mV; $P = 0.0045$) or 30 seconds (57.2 mV; $P = 0.0035$) in the presence of mefenamic acid when compared with control (20.4 mV) (Table 1). The slope of the G-V relationship, however, was not significantly different when the interpulse interval was 15 (23.2 mV) or 7 seconds (6.9 mV) in the presence of mefenamic acid when compared with control (Table 1).

Altering the interpulse interval of the activation protocol to 7 seconds, however, did not significantly affect the control (in the absence of mefenamic acid) I-V and G-V relationships (Fig. 3, D and E), although the G-V relationship did become steeper. Overall, these results show that mefenamic acid hyperpolarizes the I-V and G-V relationships of EQ $I_{Ks}$ in a rate-dependent fashion.

**Mefenamic Acid Slows EQ $I_{Ks}$ Deactivation.** To investigate changes in rates of EQ deactivation following treatment with 100 μM mefenamic acid, tail currents were obtained by pulsing to $+60 \text{ mV}$ for 4 seconds to activate $I_{Ks}$ current, followed by a 4-second pulse to a range of potentials from $-40$ to $-150 \text{ mV}$ in 10-mV steps. Holding potential was $-90 \text{ mV}$. (A) Representative tail currents (black) fit with a single exponential curve (blue lines) for EQ $I_{Ks}$, in the absence of mefenamic acid. (B) Representative tail currents (black) fit with a single exponential curve (blue lines) for EQ $I_{Ks}$ in the presence of 100 μM mefenamic acid. (C) Time constants of deactivation ($t_{\text{deact}}$) versus different membrane potentials for EQ $I_{Ks}$, in the absence ($n = 3$, black circles) and presence of 100 μM mefenamic acid ($n = 5$, gray circles). **$P = 0.0089$; ***$P = 0.0003$; and ****$P < 0.0001$.

Altering the interpulse interval of the activation protocol to 7 seconds, however, did not significantly affect the control (in the absence of mefenamic acid) I-V and G-V relationships (Fig. 3, D and E), although the G-V relationship did become steeper. Overall, these results show that mefenamic acid hyperpolarizes the I-V and G-V relationships of EQ $I_{Ks}$ in a rate-dependent fashion.

**Mefenamic Acid Slows EQ $I_{Ks}$ Deactivation.** To investigate changes in rates of EQ deactivation following treatment with 100 μM mefenamic acid, tail currents were

![Fig. 4. Mefenamic acid (Mef) slows EQ deactivation. Tail currents were obtained by pulsing to $+60 \text{ mV}$ for 4 seconds to activate $I_{Ks}$ current, followed by a 4-second pulse to a range of potentials from $-40$ to $-150 \text{ mV}$ in 10-mV steps. Holding potential was $-90 \text{ mV}$. (A) Representative tail currents (black) fit with a single exponential curve (blue lines) for EQ $I_{Ks}$, in the absence of mefenamic acid. (B) Representative tail currents (black) fit with a single exponential curve (blue lines) for EQ $I_{Ks}$ in the presence of 100 μM mefenamic acid. (C) Time constants of deactivation ($t_{\text{deact}}$) versus different membrane potentials for EQ $I_{Ks}$, in the absence ($n = 3$, black circles) and presence of 100 μM mefenamic acid ($n = 5$, gray circles). **$P = 0.0089$; ***$P = 0.0003$; and ****$P < 0.0001$.](#)

![Fig. 5. Effect of $I_{Ks}$ stoichiometry on response to 100 μM mefenamic acid. The currents and conductance-voltage plots (A–C) were obtained using the protocols described in Fig. 3 with a 15 second interpulse interval. For each stoichiometry (see Materials and Methods), representative currents are shown in the absence of control (upper left panel) and in the presence of 100 μM mefenamic acid (lower left panel). Right panels show the corresponding G-V plots in control (black circles) and presence of mefenamic acid (gray circles). Boltzmann fits were: (A) for Q1 in control ($n = 4$): $V_{1/2} = -20.5 \text{ mV}, k = 8.8 \text{ mV}$; and, in mefenamic acid ($n = 4$): $V_{1/2} = -29.6 \text{ mV}, k = 14.9 \text{ mV}$; (B) for EQQQQ in control ($n = 4$): $V_{1/2} = -1.5 \text{ mV}, k = 19.1 \text{ mV}$; and, in mefenamic acid ($n = 4$): $V_{1/2} = -20.8 \text{ mV}, k = 18.3 \text{ mV}$; (C) for EQQ in control ($n = 5$): $V_{1/2} = 15.4 \text{ mV}, k = 20.6 \text{ mV}$; and in mefenamic acid ($n = 4$) $V_{1/2} = -1.6 \text{ mV}, k = 20.3 \text{ mV}$ (See Table 2). (D) Normalized log concentration-response relationships for EQ, EQQ and EQQQQ $I_{Ks}$ were obtained using the analysis method described in Fig. 2. For EQ $I_{Ks}$: $EC_{50} = 60 [38, 89; 95\% CI] \mu M$, $n^0 = 0.49 [0.39, 0.60]; 95\% CI$, green diamonds, data from Fig. 2B; for EQQQQ $I_{Ks}$: $EC_{50} = 615 [422, 955; 95\% CI] \mu M$, $n^0 = 0.47 [0.37, 0.60]; 95\% CI$, blue diamonds; and for EQQQQ $I_{Ks}$: $EC_{50} = 902 [663, 1383; 95\% CI] \mu M$, $n^0 = 0.66 [0.49, 0.87]; 95\% CI$, red diamonds. For each construct, $n = 3–5$ at each concentration. ***, **** and **** denote a significantly different response when compared with EQ $I_{Ks}$ and, where $P < 0.05$, $P < 0.0005$ and $P < 0.0001$, respectively.](#)
obtained in the absence (Fig. 4A) and presence of mefenamic acid (Fig. 4B) and fit with single exponential decay curves. Deactivation time constants were obtained from these fits and plotted against the membrane potential (Fig. 4C). The K+ reversal potential was found to be approximately -80 mV, and therefore, the rate of deactivation at -80 mV was omitted. Treatment with mefenamic acid significantly decreased the rate of deactivation at -70 to -140 mV.

**Effect of IKs Stoichiometry on Response to 100 μM Mefenamic Acid.** To investigate whether the effect of mefenamic acid on IKs dose response and G-V relationships was dependent on the E1:Q1 stoichiometry, mammalian cells (LM and tsA201 cells) were transiently transfected with IKs, EQQIKs, EQQQQIKs, or KCNQ1 alone (Fig. 5). These IKs constructs fix the ratio of E1:Q1 to 4:4, 2:4, or 1:4 through linking the C terminus of KCNE1 to the N terminus of one, two, or four KCNQ1 sequences, respectively.

No change in the KCNQ1 waveform was seen following treatment with 100 μM mefenamic acid (control: Fig. 5A, upper-left panel; mefenamic acid: Fig. 5A, lower-left panel). Mefenamic acid also did not significantly shift the V1/2 of activation or slope of the G-V relationship (Table 2), supporting the conclusion that mefenamic acid has no effect on KCNQ1 alone.

When one KCNE1 subunit was present (EQQQQ IKs), the instantaneous current characteristic of mefenamic acid’s effect on IKs no longer occurred (Fig. 5B, left panel). The EQQQQ IKs waveform was sigmoidal in both the absence (Fig. 5A, upper-left panel) and presence of 100 μM mefenamic acid (Fig. 5A, lower-left panel). Despite mefenamic acid not having a dramatic transformative effect on the activation waveform of EQQQQ, inhibition of EQQQQ tail current decay following mefenamic acid treatment still occurred. A significant leftward shift in the V1/2 of activation also occurred (control: -1.5 mV; mefenamic acid: -20.8 mV; P = 0.0298; Table 2). There were no significant changes in the slope of the G-V relationships.

Similar to EQQQQ IKs, there was no dramatic transformative effect on the EQQ IKs activation waveform following mefenamic acid treatment (Fig. 5C, left panel). Inhibition of EQQ IKs tail current decay, however, still occurred. A significant leftward shift in the V1/2 of activation also occurred (control: 15.4 mV; mefenamic acid: -1.6 mV; P = 0.0472; Table 2). Again, the slope of the G-V relationships did not change. Overall, the leftward shift in the V1/2 of activation (ΔV1/2) for EQQQQ (-19.3 mV) and EQQ (-17.1 mV) was less dramatic than the ΔV1/2 of activation for EQ (-110 mV; Table 2). The normalized responses of the different IKs stoichiometries (normalized difference currents) to different concentrations of mefenamic acid are plotted in Fig. 5D. At all concentrations, the normalized responses of EQQQQ IKs and EQQ IKs were significantly reduced compared with the response of EQ IKs. The EC50 and nH were 902 μM and 0.66 for EQQQQ IKs, 615 μM and 0.47 for EQQ IKs, and 60 μM and 0.49 for EQ IKs, respectively. Since no change in the KCNQ1 waveform or significant shift in the V1/2 was seen following mefenamic acid treatment (Fig. 5A), these data were not included in the log concentration–normalized response plot in Fig. 5D.

**Mefenamic Acid Effects at the Single-Channel Level.** To determine if the enhancement of IKs current upon mefenamic acid treatment was simply a result of an increase in open probability or there were additional effects on conductance, we made single-channel recordings of 4:4 IKs stoichiometry (EQ) in the presence of the drug. Figure 6A shows

### TABLE 1

V1/2 of activation (millivolts) and slope value (k-factor, millivolts) in the absence and presence of 100 μM mefenamic acid for EQ IKs, at different interpulse intervals. The ± denotes S.D., with the P value indicating statistical difference in V1/2 compared with control as determined using a one-way ANOVA and Bonferroni multiple comparisons test.

<table>
<thead>
<tr>
<th></th>
<th>V1/2</th>
<th>k-Factor</th>
<th>n</th>
<th>P Value</th>
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<td>Control</td>
<td></td>
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</tr>
<tr>
<td>EQ</td>
<td>23.9 ± 3.7</td>
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<td>EQQ</td>
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<tr>
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<tr>
<td>Q1</td>
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<td>8.8 ± 2.9</td>
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<tr>
<td>E43C</td>
<td>70.8 ± 3.5</td>
<td>27.0 ± 5.5</td>
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<tr>
<td>L42C</td>
<td>68.9 ± 2.6</td>
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<tr>
<td>K41C</td>
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<td>19.7 ± 4.7</td>
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<tr>
<td>K41E</td>
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<td>24.2 ± 0.8</td>
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<tr>
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<tr>
<td>G40C</td>
<td>39.2 ± 11.8</td>
<td>20.0 ± 2.2</td>
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### TABLE 2

V1/2 of activation (millivolts) and slope value (k-factor, millivolts) in the absence and presence of mefenamic acid for mutant EQ IKs, and different stoichiometrically saturated WT IKs. The ± denotes S.D., with the P value indicating statistical difference in V1/2 compared with control as determined using a one-way ANOVA and Bonferroni multiple comparisons test.

<table>
<thead>
<tr>
<th></th>
<th>V1/2</th>
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<th>ΔV1/2</th>
<th>P Value</th>
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<tr>
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<td>-1.6 ± 2.7</td>
<td>-17.1</td>
</tr>
<tr>
<td>EQQQQ</td>
<td>-1.5 ± 5.0</td>
<td>18.1 ± 1.1</td>
<td>4</td>
<td>-20.8 ± 3.7</td>
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<tr>
<td>Q1</td>
<td>-20.5 ± 1.4</td>
<td>8.8 ± 2.9</td>
<td>5</td>
<td>-25.6 ± 12.9</td>
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<td>E43C</td>
<td>70.8 ± 3.5</td>
<td>27.0 ± 5.5</td>
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<td>L42C</td>
<td>68.9 ± 2.6</td>
<td>21.5 ± 6.4</td>
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<tr>
<td>K41C</td>
<td>17.1 ± 4.6</td>
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<td>14.0 ± 6.2</td>
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<tr>
<td>K41E</td>
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<td>24.2 ± 0.8</td>
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<tr>
<td>K41R</td>
<td>72.7 ± 5.8</td>
<td>20.8 ± 3.2</td>
<td>3</td>
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<td>4</td>
<td>-37.9 ± 23.5</td>
<td>-40.3</td>
</tr>
</tbody>
</table>

NS, not significant.

Mefenamic acid dose was either 100 μM (where applicable, upper row values) or 1 mM (where applicable, lower row values). Given the dramatic effect mefenamic acid has on the G-V relationship for some constructs, Boltzmann curves could not be properly fit in some cases.
three representative active traces of control EQ and three in the presence of 100 μM mefenamic acid. The voltage protocol was analogous to the whole-cell experiments in that a 4-second depolarization was given, with a 0.75-second repolarization period at −40 mV, applied every 10 seconds.

It is clear from these records that channels opened very early upon depolarization in the presence of mefenamic acid (Fig. 6A, bottom-three sweeps), and that channel openings were persistent during the repolarization step to −40 mV. Thus, first latency was shortened and deactivation slowed in the presence of mefenamic acid, as would be expected based on the whole-cell recordings. Note that, during exposure to mefenamic acid, channels were not already activated when initially depolarized, as might have been expected if the effects of mefenamic acid were simply related to a failure of deactivation between pulses. It is clear that channels activate de novo with each pulse. This is confirmed in the ensemble averages of 18 active sweeps of control and 18 sweeps from the same recording taken approximately 23 minutes after mefenamic acid exposure (Fig. 6B), where channel activity was seen earlier during the depolarization in the presence of mefenamic acid (red tracing) than in control (blue tracing), and tail currents persisted to the end of the recording period. The all-points histograms (Fig. 6C) comparing events in the three control and three mefenamic acid sweeps shown in Fig. 6A also show a reduction in the number of closed events, which is indicative of increased channel activity and decreased latency in the presence of mefenamic acid. In addition, the peak open amplitude around 0.4 pA is maintained but shows more events due to prolonged opening bursts seen in the presence of mefenamic acid. Maximum channel conductance was not increased by mefenamic acid; rather, simply more open events were seen at the same levels present in control.

Another important observation from the single-channel experiments using the cell-attached configuration was the delay in action of mefenamic acid. In whole-cell experiments, mefenamic acid effects were observed within approximately 0.5–1 minute of bath application, whereas when the extracellular domain of IKs was shielded by the recording electrode in the cell-attached configuration, the effect took up to 20 minutes to become obvious. This supports previous research pointing to an extracellular binding site for the drug (Abitbol et al., 1999). When mefenamic acid was included in the patch pipette, the effect on open probability was fairly immediate (data not shown).

**Mapping the Mefenamic Acid Regulatory Sites on KCNE1.** Since our data indicate that mefenamic acid has minimal effect on KCNQ1 alone, and the effect of mefenamic acid is dependent upon channel stoichiometry, we further examined potential mefenamic acid regulatory sites on KCNE1. The binding site for mefenamic acid has previously been suggested to lie between residues 39 and 43 on KCNE1 (Abitbol et al., 1999); however, the importance of each residue to mefenamic acid’s subsequent effect on IKs has not previously been characterized. Using mutational analysis, we therefore characterized how singularly mutating residues in this region would affect mefenamic acid’s ability to alter the waveform and G-V relationship of EQ IKs.

Similar to WT EQ IKs, all mutant EQ IKs showed the characteristic delay of current activation in the absence of mefenamic acid (L42C, Fig. 7A; K41C, Fig. 7B; K41R, Fig. 7C; G40C, Supplemental Fig. 1A; current data not shown for K41E and E43C). Additionally, the characteristic appearance of instantaneous current and inhibition of tail current decay induced by mefenamic acid on WT EQ IKs was preserved in the G40C (Supplemental Fig. 1B), L42C (Fig. 7A), and E43C (data not shown) EQ IKs mutants. When an interpulse interval of 15 seconds was used, mefenamic acid also significantly altered the shape of the G-V relationship (Supplemental Fig. 1C) and left shifted the V1/2 of activation for G40C (ΔV1/2: −76.1 mV) (Fig. 7D; Table 2). This effect on the G-V relationship and V1/2 of G40C was also dependent on the interpulse interval—with the shortest interpulse interval (interpulse intervals examined included 7, 15, and 30 seconds) producing the most dramatically altered G-V relationship (Supplemental Fig. 1C) and visually the most leftward shift in V1/2. In contrast,
the effect mfenamic acid had on the G-V relationships of L42C and E43C was not as dramatic (L42C: Fig. 7A, right panel; data not shown for E43C). When an interpulse interval of 15 seconds was used, the shape of the G-V relationship in both the absence and presence of mfenamic acid was sigmoidal for L42C and E43C. In addition, the slope of the G-V relationship was not significantly altered following mfenamic acid treatment; however, a significant hyperpolarizing shift in the $V_{1/2}$ of activation occurred ($\Delta V_{1/2}$ for L42C: $–37.2$ mV; $\Delta V_{1/2}$ for E43C: $–24.4$ mV) (Fig. 7D; Table 2) for both mutants. Therefore, mutating residues L42, E43, and especially G40 on KCNE1 only minimally reduced mfenamic acid’s effect on $I_{Ks}$.

Mutations made at K41 on KCNE1 had a different result. Unlike WT $I_{Ks}$, after treatment with 100 $\mu$M mfenamic acid, the waveform of K41C still showed delayed current activation and WT tail current decay (Fig. 7B, lower-left panel). Moreover, there was also no change in the slope and shape of the G-V relationship (Fig. 7B, right panel) or significant shift in the $V_{1/2}$ of activation (Fig. 7D; Table 2). This effect was more visually less dramatic than that of G40C and WT $I_{Ks}$ (WT $I_{Ks}$ G-V plots in control and 100 $\mu$M mfenamic acid, both with an interpulse interval of 15 seconds, are overlaid on the Fig. 7C right panel for comparison; $\Delta V_{1/2}$: $–110$ mV). Despite this, the effect mfenamic acid had on the G-V relationship of K41R was also dependent on the interpulse interval (data not shown). A shorter interpulse interval of 7 seconds produced a more dramatically altered G-V relationship and more leftward shift in $V_{1/2}$ than when the interpulse interval was 15 seconds. For each $I_{Ks}$ mutant, representative currents are shown in the absence (control; upper-left panel) and in the presence of 100 $\mu$M mefenamic acid (lower-left panel). Right panels show the corresponding G-V plots in control (black circles) and in the presence of 100 $\mu$M mefenamic acid (red triangles), both with a 15-second interpulse interval. Boltzmann fits were as follows: for L42C, $V_{1/2} = 68.9$ mV and $k = 21.5$ mV in control ($n = 3$), and $V_{1/2} = 31.8$ mV and $k = 14.8$ mV in 100 $\mu$M mfenamic acid ($n = 3$) (A); for K41C, $V_{1/2} = 17.1$ mV and $k = 19.7$ mV in control ($n = 6$); $V_{1/2} = 11.3$ mV and $k = 19.3$ mV in 100 $\mu$M mfenamic acid ($n = 4$), and $V_{1/2} = 14.0$ mV and $k = 18.3$ mV in 1 mM mfenamic acid ($n = 4$) (B); and for K41R, $V_{1/2} = 72.7$ mV and $k = 20.8$ mV in control ($n = 3$), and $V_{1/2} = 28.6$ mV and $k = 49.3$ mV in 100 $\mu$M mfenamic acid ($n = 4$) (C) (Table 2). (D) Change in $V_{1/2}$ ($\Delta V_{1/2}$) for each $I_{Ks}$ mutant in control versus mfenamic acid ($n = 3–6$ at each concentration). *$P < 0.05$, **$P = 0.0011$, ***$P = 0.0003$, and ****$P < 0.0001$ (significant change in $V_{1/2}$ comparing control to the presence of mfenamic acid).

**Fig. 7.** Mapping the mfenamic acid (Mef) regulatory sites on KCNE1. The currents and G-V plots (A–C) were obtained using the protocols described in Fig 3. The interpulse interval was 15 seconds. For each $I_{Ks}$ mutant, representative currents are shown in the absence (control; upper-left panel) and presence of 100 $\mu$M mfenamic acid (lower-left panel). Right panels show the corresponding G-V plots in control (black circles) and presence of 100 $\mu$M mfenamic acid (red triangles), both with a 15-second interpulse interval. Boltzmann fits were as follows: for L42C, $V_{1/2} = 68.9$ mV and $k = 21.5$ mV in control ($n = 3$), and $V_{1/2} = 31.8$ mV and $k = 14.8$ mV in 100 $\mu$M mfenamic acid ($n = 3$) (A); for K41C, $V_{1/2} = 17.1$ mV and $k = 19.7$ mV in control ($n = 6$); $V_{1/2} = 11.3$ mV and $k = 19.3$ mV in 100 $\mu$M mfenamic acid ($n = 4$), and $V_{1/2} = 14.0$ mV and $k = 18.3$ mV in 1 mM mfenamic acid ($n = 4$) (B); and for K41R, $V_{1/2} = 72.7$ mV and $k = 20.8$ mV in control ($n = 3$), and $V_{1/2} = 28.6$ mV and $k = 49.3$ mV in 100 $\mu$M mfenamic acid ($n = 4$) (C) (Table 2). (D) Change in $V_{1/2}$ ($\Delta V_{1/2}$) for each $I_{Ks}$ mutant in control versus mfenamic acid ($n = 3–6$ at each concentration). *$P < 0.05$, **$P = 0.0011$, ***$P = 0.0003$, and ****$P < 0.0001$ (significant change in $V_{1/2}$ comparing control to the presence of mfenamic acid).
15 seconds (data not shown). Additionally, mefenamic acid also resulted in a significant leftward shift in the V_{1/2} of activation of K41R (ΔV_{1/2} = -46.2 mV) (Fig. 7D; Table 2). These results clearly show that residue K41, and especially the charge on this residue, is important in facilitating mefenamic acid’s modulation of Ik, gating kinetics.

**Discussion**

Previous studies have shown that mefenamic acid increases activation of Ik, current and slows tail current decay, but curiously, all of these have only used a single concentration of drug (100 μM). Generally, mefenamic acid effects are restricted to complexes of KCNQ1 and KCNE1 (Busch et al., 1994, 1997; Unsöld et al., 2000), although Abitbol et al. (1999) suggested that mefenamic acid may also facilitate KCNQ1 expressed alone in oocytes. Most studies have not shown that facilitation of Ik is accompanied by dramatic changes in the current activation time course, except for Unsöld et al. (2000). In the current study, we confirm that mefenamic acid enhances Ik activity (Fig. 1C) but not KCNQ1 alone (Fig. 5A), and that this effect is specific to Ik given that the large instantaneous current was blocked by HMR1556 (Fig. 1D), and no effect on endogenous currents was seen in untransfected cells treated with increasing concentrations of mefenamic acid (Fig. 1B). Following validation of these previous findings, we further defined the concentration dependence, effect of interpulse interval, stoichiometry dependence, effect on single-channel conductance, and the KCNE1 regulatory sites for mefenamic acid actions on Ik.

**Mefenamic Acid Actions on Saturated Complexes of Ik (EQ).** Increasing concentrations of mefenamic acid (1, 10, and 30 μM) have been used to confirm the functional effect of activating Ik in preconstricted rat mesenteric arteries (E_{max} of 96.1%) (Chadha et al., 2012), but as noted earlier, regardless of the expression system, a single concentration of 100 μM has been used to characterize the electrophysiological actions of mefenamic acid on Ik. One striking effect is the induction of instantaneous current and reduction in the overall time-dependent slow activation of Ik (Fig. 1C). These current changes were used to define the concentration dependence of mefenamic effects on Ik, giving an EC_{50} and n^{H} of 60 μM and 0.49, respectively (Fig. 2). An n^{H} of <1 does not suggest multiple sites of action or positive cooperative binding of mefenamic acid to the channel complex, which is supported by the similar values of n^{H} for different stoichiometric ratios of KCNQ1:KCNE1 (n^{H} = 0.47 and 0.66 for EQ and EQQQQ, respectively). The more commonly reported (in all prior studies) action of mefenamic acid is to slow current deactivation (Busch et al., 1994; Magyar et al., 2006; Toyoda et al., 2006), and our experiments also confirmed the marked slowing of tail currents with time constants increasing across the range of repolarizing potentials (e.g., from 0.52 to 1.24 seconds at -90 mV; Fig. 4C).

Mefenamic acid has a hyperpolarizing effect on the Ik, I-V and G-V relationships, which, in turn, is reflected by a leftward shift in the V_{1/2} of activation. Quantitatively, this hyperpolarization of the V_{1/2} has previously been inconsistently reported (−15 mV in CHO cells and −26 mV in canine ventricular myocytes) (Unsöld et al., 2000; Magyar et al., 2006). In Fig. 3, we demonstrated that this inconsistency may be related to mefenamic acid’s striking effect on the G-V relationship that results in part from slowed channel deactivation at shorter pulse intervals. Prior studies often did not state intervals between pulses, so this cannot be verified.

With a voltage protocol that lasted 5 seconds and an interpulse interval of 7 seconds, there was not enough time to allow for complete Ik deactivation between voltage clamp pulses in the presence of mefenamic acid. As such, an accumulation of current occurred, which partly explains the dramatically altered G-V relationship and instantaneous current at this rate (Fig. 3E). We initially hypothesized that if enough time were given to allow for Ik deactivation, the G-V relationship in the presence of mefenamic acid would mirror that seen in control (in the absence of mefenamic acid). However, even with an interpulse interval of 30 seconds in the presence of mefenamic acid, when deactivation should be complete, the G-V relationship still showed a significantly large departure in shape, slope, and V_{1/2} when compared with control. At interpulse intervals of 15, 20, and especially 30 seconds, there is a flattening of the voltage dependence of the G-V relationship caused by mefenamic acid that suggests a fundamental modification in the way that Ik senses and/or responds to changes in the transmembrane potential in the presence of the drug.

The single-channel data provide support for the whole-cell findings. There was no change in maximum channel conductance (Fig. 6), but openings did group toward higher open subconductance levels (Fig. 6, A and C) (Werry et al., 2013), which points to enhanced channel activation gating in the presence of mefenamic acid. In addition, data showed a large reduction in the first latency to opening of single channels, with channels opening in mefenamic acid de novo, soon after depolarizing pulses were applied. It is important to note that single-channel recordings did not indicate that failure of deactivation between pulses was an important factor in the decreased latency to opening (Fig. 6). Thus, single-channel kinetics provided significant insight into the changes induced by mefenamic acid seen at the whole-cell level. Finally, persistent single-channel openings were seen during repolarizations to −40 mV in the presence of mefenamic acid (Fig. 6A), mirroring the slow decay of macroscopic tail currents in the presence of drug (Fig. 4B).

To test whether actions of mefenamic acid on activation voltage gating could be sufficient to account for the current changes observed, we used an Ik model (Zaydman et al., 2014; Westhoff et al., 2019) to simulate drug action, as described in the Materials and Methods. Increasing VS forward rates for both activation transitions could account quantitatively for the dose- and rate-dependent action of mefenamic acid (Fig. 8). Increasing drug concentration caused the appearance of an instantaneous activating current with a subsequent slower phase, as seen in experimental data (Figs. 1 and 2). At the highest concentrations, the slow activating phase of current was almost abolished in simulations (Fig. 8A). The normalized-response relationship was simulated at a range of drug concentrations and was fit to a Hill equation with an EC_{50} of 78.6 μM, close to that found experimentally (60 μM), and with a Hill slope of 0.81, compared with 0.49–0.66 found for EQ and EQQQQ, respectively. The action of mefenamic acid was also simulated for step voltage clamp pulses from −80 to +100 mV given every 7 and 15 seconds (Fig. 8, C and D). The complete set of simulated currents during the increasing amplitude clamp steps for all four conditions is shown.
in Fig. 8C, with the control tracings in blue and mefenamic acid in red. The 7-second interval protocol lasts ~130 seconds in total, and the longer 15-second interval protocol lasts ~285 seconds. The effect of mefenamic acid was to increase peak current at both rates, especially at more negative potentials, as can be clearly seen. The tail currents from these protocols plotted as a function of voltage (Fig. 8D) give G-V relations, and as seen experimentally (Fig. 3E), the effect of rate was to cause a minor hyperpolarizing shift in the presence of mefenamic acid, which was greater at 7 seconds than at 15 seconds.

Mefenamic Acid Action Is Diminished with Fewer KCNE1 Subunits in the IKs Channel Complex. As the KCNQ1:KCNE1 stoichiometry of IKs is likely variable in vivo (Dvir et al., 2014), understanding the effect of mefenamic acid on different subunit ratios is extremely important, and the use of fixed stoichiometry constructs in the present experiments allowed a quantitative comparison of the action of mefenamic acid on different stoichiometries of IKs (Murray et al., 2016). The V1/2 of activation for KCNQ1 alone was not altered by mefenamic acid, but with the partially saturated IKs complexes, EQQ and EQQQQ, the V1/2 of activation was significantly hyperpolarized (Fig. 5). This hyperpolarization was less dramatic than that seen when IKs was fully saturated (EQ; Fig. 3E). Similarly, in the dose-response curves, the responses of EQ and EQQQQ to mefenamic acid at all concentrations were significantly less than that of EQ, which further supports the idea that the effect of mefenamic acid on IKs is stoichiometrically graded.

Mefenamic Acid Binding to the IKs Complex. The binding site for DIDS and mefenamic acid on IKs has previously been suggested to lie between residues 39 and 43 on KCNE1, with residue E43 specifically identified as critical for the binding of DIDS (Abitbol et al., 1999). Whether this site and/or other residues in the mapped region are critical for the binding of mefenamic acid to KCNE1, and/or IKs, was not studied. Through mutational analysis, we now show that, although mutation of residues E43 and L42 in KCNE1 results in a reduced response to mefenamic acid, residue K41 is critical for the action of mefenamic acid (Fig. 7).

Most of the mutations themselves do have variable effects on the gating of IKs in the absence of mefenamic acid (Table 2), but importantly, K41C has little effect on the position or slope of the IKs G-V relationship compared with WT EQ. This suggests that K41C does not itself destabilize the normal interactions between KCNQ1 and KCNE1 in the IKs channel complex, or its ability to respond to applied changes in potential. These results are particularly interesting when taken in the context of the known importance of interactions between KCNQ1 and this region of KCNE to the pathophysiology of short QT syndrome, in which extremely slow deactivation is a feature (Dvir et al., 2014).
The E43C, L42C, and K41E mutations shift the \( V_{1/2} \) of \( I_{Ks} \) to approximately +70 mV (Table 2), which is opposite to the direction expected if they were inhibiting the interactions between KCNQ1 and KCNE1 (Murray et al., 2016), but which may explain their lesser response to mefenamic acid (Fig. 7D). The G40C mutant responds almost like WT to mefenamic acid (Supplemental Fig. 1), and so defines a proximal limit of the critical region. Taken together, the data indicate the primary importance of K41 in the binding of mefenamic acid to the \( I_{Ks} \) channel complex, and the response of K41R, but not K41E, to mefenamic acid suggests the importance of electrostatic rather than steric interactions in this effect.

Relevance of Mefenamic Acid Activation of \( I_{Ks} \) Channel Currents. Unlike most other activators that have little effect on \( I_{Ks} \) channels with increasingly saturated stoichiometries (Magyar et al., 2006; Gao et al., 2008; Yu et al., 2013), we have shown that mefenamic acid can enhance all \( I_{Ks} \) channel complexes of different stoichiometries, suggesting that molecules like it may represent a therapeutic approach to treating LTQTS types 1 and 5. Although this is well beyond the scope of the present study, we note that mefenamic acid is presently prescribed at a recommended dosage of 500 mg/day, which has been reported to equate to a mean plasma concentration of 82.9 \( \mu \)M (Cryer and Feldman, 1998). This provides little clue toward the amount needed therapeutically to treat LTQTS, but in our experiments such concentrations cause a potent activating effect of mefenamic acid that is also dependent on the stimulus rate, which is important because \( I_{Ks} \) primarily contributes to cardiac repolarization at high heart rates. The definitive stoichiometry of \( I_{Ks} \) in humans as well as the degree of \( I_{Ks} \) channel activation required for a therapeutically beneficial shortening of the QT interval are presently unknown, so we cannot know whether compounds such as mefenamic acid could have a beneficial effect in LTQTS. We do, however, know that, due to the known adverse gastrointestinal effects of COX1 inhibition and block adverse gastrointestinal effects of COX1 inhibition and block"
Unnatural amino acid photo-crosslinking of the IKs channel complex demonstrates a KCNE1:KCNQ1 stoichiometry of up to 4:4. *eLife* 5:e11815.


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