

Title

Evidence for a common pharmacological interaction site on K_{Ca}2 channels providing both selective activation and selective inhibition of the human K_{Ca}2.1 subtype

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Running title:

Positive and negative gating modulation of K_{Ca}2.1

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Abbreviations:

B-TPMF, *N*-{7-[1-(4-*tert*-butyl-phenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidinium; CaM, calmodulin; CaMBD, calmodulin binding domain; CM-TPMF, *N*-{7-[1-(4-chloro-2-methylphenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidinium; CyPPA, cyclohexyl-[2-(3,5-dimethyl-pyrazol-1-yl)-6-methylpyrimidin-4-yl]-amine; DCEBIO, 5,6-dichloro-1-ethyl-1,3-dihydro-2*H*-benzimidazol-2-one; 1-EBIO, 1-ethyl-2-benzimidazolinone; GW542573X, 4-(2-methoxyphenylcarbamoyloxymethyl)-piperidine-1-carboxylic acid *tert*-butyl ester; HEK293, human embryonic kidney 293; K_{Ca}2, small

conductance Ca^{2+} -activated K^{+} channel (SK channel); $\text{K}_{\text{Ca}}3.1$, intermediate conductance Ca^{2+} -activated K^{+} channel (IK channel); NS309, 6,7-dichloro-1*H*-indole-2,3-dione 3-oxime; NS4591, 4,5-dichloro-1,3-diethyl-2,3-dihydro-1*H*-1,3-benzodiazol-2-one; NS8593, (*R*)-*N*-(benzimidazol-2-yl)-tetrahydro-1-naphtylamine; S, transmembrane segment; SKA-31, naphtho[1,2-*d*]thiazol-2-ylamine.

ABSTRACT

We have previously identified Ser293 in transmembrane segment 5 as a determinant for selective $K_{Ca2.1}$ channel activation by GW542573X (4-(2-methoxyphenylcarbamoyloxymethyl)-piperidine-1-carboxylic acid *tert*-butyl ester). Now we show that Ser293 mediates both activation and inhibition of $K_{Ca2.1}$: CM-TPMF (*N*-{7-[1-(4-chloro-2-methylphenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine) and B-TPMF (*N*-{7-[1-(4-*tert*-butyl-phenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine), two newly identified and structurally related [1,2,4]triazolo[1,5-*a*]pyrimidines, act either as an activator or as an inhibitor of the human $K_{Ca2.1}$ channel. While (-)-CM-TPMF activates $K_{Ca2.1}$ with an EC_{50} value of 24 nM, (-)-B-TPMF inhibits the channel with an IC_{50} value of 31 nM. In contrast, their (+)-enantiomers are 40-100 times less active. Both (-)-CM-TPMF and (-)-B-TPMF are subtype-selective with 10-20 fold discrimination towards other K_{Ca2} channels and the K_{Ca3} channel. Co-application experiments reveal competitive-like functional interactions between the effects of (-)-CM-TPMF and (-)-B-TPMF. Despite belonging to a different chemical class than GW542573X the $K_{Ca2.1}$ selectivity of (-)-CM-TPMF and (-)-B-TPMF depend critically on Ser293 as revealed by loss- and gain-of-function mutations. We conclude that compounds occupying the TPFM site may either positively or negatively influence the gating process depending on their substitution pattern. Notably, (-)-CM-TPMF is 10 times more potent on $K_{Ca2.1}$ than NS309 (6,7-dichloro-1*H*-indole-2,3-dione 3-oxime), an unselective but hitherto the most potent K_{Ca3}/K_{Ca2} channel activator. (-)-B-TPMF is the first small molecule inhibitor with significant selectivity among the K_{Ca2} channel subtypes, and in contrast to peptide blockers like apamin and scyllatoxin, which preferentially affect $K_{Ca2.2}$, (-)-B-TPMF exhibits $K_{Ca2.1}$

selectivity. These high-affinity compounds, which exert opposite effects on $K_{Ca2.1}$ gating, may help define physiological or pathophysiological roles of this channel.

Introduction

Small and intermediate conductance Ca^{2+} -activated K^{+} channels (the KCNN family: $\text{K}_{\text{Ca}2.1}$, $\text{K}_{\text{Ca}2.2}$, $\text{K}_{\text{Ca}2.3}$, and $\text{K}_{\text{Ca}3.1}$) contain six transmembrane segments (S1-6) arranged as a tetramer around a central pore. Each subunit associates with one calmodulin (CaM) molecule, acting as a β -subunit at the C-terminal CaM binding domain (CaMBD) (Xia et al., 1998; Khanna et al., 1999). Ca^{2+} binding to CaM rearranges the CaM/CaMBD region and the S6/S5 domains, thereby opening the channel. The nature of the gate remains elusive but cysteine scanning experiments have narrowed down its possible location to residues in the inner pore vestibule close to the selectivity filter. In contrast to K_v channels, KCNN channels are assumed to exhibit “deep pore gating” (Bruening-Wright et al., 2007; Bruening-Wright et al., 2002; Garneau et al., 2009; Klein et al., 2007), which is important for the understanding of the molecular pharmacology of positive and negative KCNN channel gating modifiers.

The $\text{K}_{\text{Ca}2}$ channel subtypes are widely and distinctly expressed in the CNS, controlling somatic excitability, pacemaker firing rates, and synaptic plasticity (Pedarzani and Stocker, 2008; Bond et al, 2005). Clinically used drugs with $\text{K}_{\text{Ca}2}$ activating properties are the centrally acting muscle relaxants zoxazolamine and chlorzoxazone, which are used for treating spasticity, and riluzole, which is registered for amyotrophic lateral sclerosis. The $\text{K}_{\text{Ca}2}$ -facilitating properties (probably $\text{K}_{\text{Ca}2.2}$) of both chlorzoxazone and riluzole have recently been suggested to account for their beneficial effects in rodent ataxia models (Alviña and Khodakhah, 2010; Janahmadi et al., 2009) and in patients (Ristori et al., 2010). Arguably, $\text{K}_{\text{Ca}2}$ activation (probably $\text{K}_{\text{Ca}2.3}$) also underlies reduced craving and dampening of excessive alcohol intake observed with chlorzoxazone in rats (Hopf et al., 2011). Important hallmarks for a therapeutic $\text{K}_{\text{Ca}2}$ -facilitating mechanism of these drugs are that 1-EBIO (1-ethyl-2-benzimidazolinone), the prototype KCNN channel activator,

exerts a similar anti-ataxic action when applied locally into the cerebellum of tottering mice (Walter et al., 2006) and anti-addictive effect against alcohol when applied into rat nucleus accumbens core region (Hopf et al., 2010). In contrast, inhibition of K_{Ca2} channels has not been demonstrated to account for clinical efficacy of any drug. The anti-depressant fluoxetine is a weak blocker of cloned K_{Ca} channels (Terstappen et al., 2003) and $K_{Ca2.3}$ channel block may mediate an anti-depressant mechanism (Galeotti et al., 1999, Jacobsen et al., 2008). However, it is doubtful whether this mechanism for fluoxetine plays a role compared to serotonin transporter inhibition.

Validation of specific K_{Ca} channel subtypes as targets for CNS drug discovery is hampered by lack of selective and potent molecules that pass the blood brain barrier. Classic K_{Ca2} channel blockers like UCL1684 (6,10-diaza-3(1,3),8(1,4)-dibenzena-1,5(1,4)-diquinolinacyclodecaphane) mimic the inhibition by the K_{Ca2} selective peptide apamin, and are unselective among the K_{Ca2} subtypes. Like apamin, they further contain basic and charged moieties that limit their passage into the CNS. 1-EBIO and the K_{Ca2} -activating drugs are non-selective K_{Ca2} enhancers and act more potently on $K_{Ca3.1}$. Unfortunately this is also the case for the more potent positive modulators DCEBIO, NS309, NS4591 and SKA-31, even though these have emerged from dedicated chemical optimization programs (Wulff and Zhorov, 2008).

Subtype-selective positive gating modulators and non-selective negative gating modulators have recently been identified. These have been instrumental in defining new sites and new modes-of-actions for gating modulation of K_{Ca2} channels. The negative gating modulator NS8593 ((*R*)-*N*-(benzimidazol-2-yl)-tetrahydro-1-naphtylamine), causes a reduction in the apparent Ca^{2+} -sensitivity for channel activation (Strøbæk et al., 2006, Sørensen et al., 2008). This effect depends on Ser507 and Ala532 ($K_{Ca2.3}$ numeration) positioned in the S5 pore helix and S6

transmembrane segment just below the selectivity filter and was therefore interpreted as NS8593-interaction with deep pore gating structures (Jenkins et al., 2011). These amino acids are conserved among the K_{Ca2} channels, thus explaining the lack of K_{Ca2} subtype selectivity of NS8593 and close analogues (Sørensen et al., 2008). The $K_{Ca2.1}$ selective activator GW542573X, which acts dually by directly opening the channel and by increasing the apparent Ca^{2+} -sensitivity (Hougaard et al., 2009), depends on Ser293 in the S5 transmembrane helix, a $K_{Ca2.1}$ specific amino acid. These mixed effects differ significantly from the mode-of-action of non-selective activators, but also from the $K_{Ca2.2/3}$ selective modulator CyPPA (cyclohexyl-[2-(3,5-dimethyl-pyrazol-1-yl)-6-methylpyrimidin-4-yl]-amine), which increases the apparent Ca^{2+} -sensitivity via interaction with the C-terminal (Hougaard et al., 2007; Li et al., 2009; Pedarzani et al., 2001).

Here we describe two novel and closely related [1,2,4]triazolo[1,5-*a*]pyrimidines, (-)-CM-TPMF and (-)-B-TPMF, which both exhibit high-potency (~30 nM) and subtype-selective interactions with $K_{Ca2.1}$ via Ser293. These compounds show opposite functions with (-)-CM-TPMF acting as an activator and (-)-B-TPMF as an inhibitor.

Materials and Methods

Chemistry

B-TPMF ((*N*-{7-[1-(4-*tert*-butyl-phenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine) and CM-TPMF (*N*-{7-[1-(4-chloro-2-methylphenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine) were purchased as racemates from Key Organics Ltd, Camelford, Cornwall, UK. The compounds were resolved into their enantiomeric pairs by preparative chiral chromatography using for B-TPMF a Chiracel OD-H column (solvent 40% ethanol in hexane, flow rate 1.0 ml/min) and for CM-TPMF a Chiralpak AD-H column (solvent 40% ethanol in hexane, flow rate 1.0 ml/min). Determination of chiral purity was performed on an analytical Waters Thar Investigator instrument equipped with a PDA detector: for B-TPMF using a Phenomenex Lux2 column, 250 x 4.6 mm, 5 μ m particle size (solvent super critical CO₂ with ethanol as co-solvent and isocratic elution with 45% co-solvent and a flow rate of 3.2 ml/min); for CM-TPMF using a Phenomenex Lux1 Column, 250 x 4.6 mm, 5 μ m particle size (solvent super critical CO₂ with methanol as co-solvent and isocratic elution with 40% co-solvent and a flow rate of 3.2 ml/min). Chiral purity was determined by integration of the PDA (total absorbance) trace. Optical rotations were measured on a Perkin Elmer polarimeter (model 314). Absolute configurations were not determined. NMR spectra were recorded on a Bruker ultra shield plus 400 MHz (Advance II) instrument. Characterization of (-)-B-TPMF (C₁₉H₂₄N₆O₂): Specific optical rotation $[\alpha]_D^{25} = -291^\circ$ (CHCl₃); enantiomeric excess (ee) >99%; HR-MS(ES+) *m/z* 369.20362 ([*M* + 1]⁺, 100%), calcd. 369.20328; ¹H NMR (CDCl₃) δ 1.26 (s, 9H), 1.78 (d, 3H, *J* = 6.6 Hz), 3.92 (s, 3H), 5.89 (q, 1H, *J* = 6.5 Hz), 6.75 (m, 2H), 7.10 (m, 2H), 7.25 (m, 2H), 7.82 (d, 1H, *J* = 10.4 Hz), 7.98 (d, 1H, *J* = 10.4 Hz); 8.60 (d, 1H, *J* = 4.7 Hz). Characterization of (-)-CM-TPMF (C₁₆H₁₇ClN₆O₂): Specific optical

rotation $[\alpha]_D^{25} = -275^\circ$ (CHCl_3); ee = 96.4%; HR-MS(ES+) m/z 361.11775 ($[\text{M} + 1]^+$, 100%), calcd. 361.11738; ^1H NMR (CDCl_3) δ 1.80 (d, 3H, $J = 6.5$ Hz), 2.34 (s, 3H), 3.93 (s, 3H), 5.89 (q, 1H, $J = 6.4$ Hz), 6.44 (d, 1H, $J = 8.8$ Hz), 6.95-7.04 (m, 2H), 7.17 (m, 1H), 7.81(d, 1H, $J = 10.4$ Hz), 7.97 (d, 1H, $J = 10.2$), 8.61 (d, 1H, $J = 4.7$ Hz).

Molecular biology.

The generation of point mutated $\text{hK}_{\text{Ca}2.3\text{L476S}}$ has been described previously (Hougaard et al., 2009) and $\text{hK}_{\text{Ca}2.1\text{S293L}}$ was generated by mutagenesis on $\text{hK}_{\text{Ca}2.1}$. Briefly, the plasmid encoding $\text{hK}_{\text{Ca}2.1}$ was uracilated by the *E. Coli* RZ1032 and used as template in a mutagenesis reaction utilizing the oligonucleotide GTGCTGCTGGTCTTCtcgATaTCCctCTGGATCATCGCAGC, T7 DNA polymerase and T4 DNA ligase. The mutation was verified by sequencing.

Cell cultures

For patch clamp experiments with $\text{hK}_{\text{Ca}2.1}$, $\text{hK}_{\text{Ca}2.2}$, $\text{hK}_{\text{Ca}2.3}$ and $\text{hK}_{\text{Ca}3.1}$, HEK293 cell lines stably expressing these channels were used (Hougaard et al., 2009). Point mutated channels ($\text{hK}_{\text{Ca}2.1\text{S293L}}$ and $\text{hK}_{\text{Ca}2.3\text{L476S}}$) were transiently transfected into HEK293 cells using Lipofectamine (Invitrogen, Carlsbad, CA) and standard transfection methods. Electrophysiological measurements were performed 2-3 days after transfection. Cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM, Sigma-Aldrich, Brøndby, Denmark) enriched with 10% fetal calf serum (FCS, Gibco, Invitrogen, NY, USA) at 37°C and 5% CO_2 . At approximately 75% confluence, the cells were washed once with phosphate buffered saline (PBS), harvested by trypsin/EDTA (Sigma-Aldrich) treatment and transferred to petri dishes containing cover slips (\varnothing 3.5 mm, purchased from VWR international, Herlev, Denmark).

Electrophysiology

Membrane currents were recorded using the whole-cell or the inside-out configuration of the patch-clamp technique. Cells seeded on cover slips were transferred to a 15 μ l recording chamber and continuously superfused at 1 ml min⁻¹. An integrated Ag/AgCl pellet electrode served as reference. Experiments were conducted at room temperature. Patch pipettes (approx. 2 M Ω) were pulled from borosilicate tubes with an outside diameter of 1.32 mm (Vitrex Medical, Herlev, Denmark) using a horizontal electrode puller (Zeitz Instruments, Augsburg, Germany). An electronically controlled micromanipulator (Eppendorf, Radiometer, Denmark) was used for the positioning of pipettes and the experiments were controlled by an EPC-9 amplifier (HEKA, Lambrecht, Germany). Data were filtered at 3 kHz. Currents were elicited by applying a 200 ms linear voltage ramp from -80 to +80 mV every 5 s from a holding potential of 0 mV. In whole-cell experiments, the cell capacitance and series resistance (R_s below 8 M Ω , 80% compensation) were updated before each voltage ramp.

In all experiments a solution with a high K⁺ concentration was applied to the extracellular side of the membrane (in mM): 154 KCl, 2 CaCl₂, 1 MgCl₂ and 10 HEPES, pH adjusted to 7.4 with 1 M KOH. The intracellular solutions contained (in mM): 154 KCl, 10 HEPES, 10 EGTA, or a combination of EGTA and NTA (10 mM in total). Concentrations of MgCl₂ and CaCl₂ required to obtain the desired free concentrations (Mg²⁺ always 1 mM, Ca²⁺ 0.01 – 10 μ M) were calculated (EqCal, Cambridge, UK) and added. In the nominal Ca²⁺-free intracellular solution, no Ca²⁺ was added. The intracellular solutions were adjusted to pH 7.2 with 1 M KOH.

Calculations and statistics

In inside-out experiments EC_{50} and IC_{50} as well as n_{Hill} values for compounds and Ca^{2+} were estimated from equilibrium concentration-response relationships by fitting to the Hill-equation. In whole-cell experiments, EC_{50} values for current activation are not readily determined due to poor clamping conditions at higher degrees of channel activation. Instead the concentrations of compound giving 100 % current increase (called the SC_{100} value, see Table 1) were estimated. In some experiments K_d values were calculated from single concentration applications by fitting to the kinetics of inhibition (see Strøbæk et al., 2006).

All results are presented as the mean \pm standard error of the mean (S.E.M.) with the number of experiments indicated. Significance testing was performed by Student's t-test for unpaired samples.

Results

The chemical structures of the two new compounds B-TPMF (*N*-{7-[1-(4-*tert*-butylphenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine) and CM-TPMF (*N*-{7-[1-(4-chloro-2-methylphenoxy)ethyl]-[1,2,4]triazolo[1,5-*a*]pyrimidin-2-yl}-*N'*-methoxy-formamidine) as well as GW542573X are shown in Fig. 1. The effects of the isolated enantiomers of B-TPMF and CM-TPMF (Materials and Methods, chemistry section) were determined in whole-cell voltage-clamp experiments by applying 10 nM of the compounds to HEK293 cells expressing human $K_{Ca2.1}$ channels. Representative current traces are shown in Fig. 2A and the current recorded at -75 mV is plotted vs. time in Fig. 2B. The (-)-enantiomer of B-TPMF *inhibited* the $K_{Ca2.1}$ current, whereas the (+)-enantiomer was without effect. In contrast, (-)-CM-TPMF, induced a robust *activation*, again with the (+)-enantiomer exerting no effect. The time course of this experiment (Fig. 2B) underscores the reversibility of these actions and the I-V curves show preservation of the normal inward rectification in presence of all compounds. Fig. 2C summarizes this series of experiments and specifically documents that 10 nM of (-)-B-TPMF significantly ($p < 0.001$, students t-test for unpaired samples) inhibits the $K_{Ca2.1}$ current, while (-)-CM-TPMF enhances it. The corresponding (+)-enantiomers of both compounds in contrast are inactive at this concentration. Whole-cell experiments were also performed in order to quantify the enantiomer selectivity (experiments not illustrated): K_d values for (+)-B-TPMF and (-)-B-TPMF were $1.6 \pm 0.5 \mu\text{M}$ ($n = 5$) and $15 \pm 7 \text{ nM}$ ($n = 5$), respectively. Similarly, (+)-CM-TPMF activated $K_{Ca2.1}$ with an SC_{100} of $200 \pm 44 \text{ nM}$ ($n = 5$) compared to $5 \pm 0.2 \text{ nM}$ ($n = 6$) for (-)-CM-TPMF. Thus, for B-TPMF and CM-TPMF the activity on $K_{Ca2.1}$ channels primarily resides in the (-) enantiomers with enantiomer selectivities of ~100 fold and ~40 fold.

To investigate the subtype selectivity of the more active (-)-enantiomers experiments were conducted on HEK293 cells stably expressing hK_{Ca}2.1, hK_{Ca}2.2 or hK_{Ca}2.3 channels (Fig. 3). Since (-)-B-TPMF and (-)-CM-TPMF act equally well from both sides of the membrane, the inside-out configuration of the patch-clamp technique was chosen in order to achieve a well defined [Ca²⁺]_i. At a concentration of 30 nM, (-)-B-TPMF inhibited the K_{Ca}2.1 current by approximately 50% whereas conversely (-)-CM-TPMF induced a 3-fold increase. At the same concentration only minor effects were detectable on K_{Ca}2.2 and K_{Ca}2.3 (Fig. 3A) indicating that both compounds exhibit selectivity for K_{Ca}2.1 over the closely related isoforms. Full concentration-response relationships were obtained for the three K_{Ca}2 subtypes and results for both compounds are summarized in Fig. 3B and 3C. The data generated for (-)-B-TPMF-induced inhibition conformed well to the Hill-equation with an IC₅₀ value for K_{Ca}2.1 of approximately 30 nM and a complete inhibition of the current at 1 μM, whereas K_{Ca}2.2 and K_{Ca}2.3 both were inhibited with IC₅₀ values around 1 μM (see Table 1, and text to Fig. 3). Hence (-)-B-TPMF is a selective inhibitor of K_{Ca}2.1 which displays approximately 30-fold selectivity over the other K_{Ca}2 channels. The concentration-response relationships obtained from applications of (-)-CM-TPMF to marginally activated K_{Ca}2 channels (0.2 μM Ca²⁺, degree of activation ~ 15%) yielded an EC₅₀ value of 24 nM for K_{Ca}2.1 and a maximal degree of activation of approximately 80 % compared to a [Ca²⁺]_i of 10 μM. K_{Ca}2.2 and K_{Ca}2.3 were activated with EC₅₀ values of 290 and 250 nM showing that (-)-CM-TPMF is 10-fold more selective for K_{Ca}2.1. Similar selectivity ratios were obtained against K_{Ca}3.1 except for (-)-CM-TPMF, which acted as a (very weak) activator (Table 1). Table 1 also summarizes the effects of (-)-CM-TPMF and (-)-B-TPMF on more distantly related ion channels. Only marginal inhibitory effects were observed for both compounds at 10 μM on members of other K⁺ channel gene families (K_{Ca}1.1, K_v7.2+K_v7.3,

and K_v11.1) as well as Na_v1.2 channels. A broader selectivity profile was obtained for the two racemates by testing at concentrations of 10 μ M in the 69 receptor LeadProfilerScreen (PT# 1107076) at MDSPharma (now Ricerca Biosciences LLC), Taiwan. Neither of the racemates reached the pre-defined significance criteria of > 50 % effect in any assay.

We next investigated the mode-of-actions of (-)-CM-TPMF and (-)-B-TPMF. Classic enhancers of K_{Ca}2 channel activity increase the apparent Ca²⁺-sensitivity and are unable to activate the channels in the absence of intracellular Ca²⁺. In contrast, GW542573X (see Fig. 1), a recently described K_{Ca}2.1 selective compound, acts by a dual mechanism, increasing the apparent Ca²⁺-sensitivity and inducing a small Ca²⁺-independent channel activation. Fig. 4A shows the time course of an inside-out experiment where the [Ca²⁺]_i was varied (0, 0.2, and 10 μ M) and the ability of 100 nM (-)-CM-TPMF to activate the K_{Ca}2.1 current at each Ca²⁺ concentration was investigated. Fig. 4B shows the corresponding I-V relationships. Addition of (-)-CM-TPMF clearly enhanced the K_{Ca}2.1 current in the nominal absence of Ca²⁺ and shifted the I-V relationship from a small, non-specific linear leak current to an inwardly rectifying current typical for K_{Ca}2 channels recorded under symmetrical K⁺ conditions. At an intermediate Ca²⁺ concentration of 200 nM, where the K_{Ca}2.1 channels are partially Ca²⁺-activated, as well as at saturating Ca²⁺ concentrations of 10 μ M, where the channels are maximally activated, application of (-)-CM-TPMF induced an increase in the current level. At all [Ca²⁺]_i the current measured in the presence of (-)-CM-TPMF maintained the characteristic inward rectification. Fig. 4C shows the Ca²⁺-response relationship of K_{Ca}2.1 in the absence of compound (Ctrl) as well as in the presence of 0.1 μ M or 1 μ M of (-)-CM-TPMF. Notably, (-)-CM-TPMF exerts a dual action on the K_{Ca}2.1 channel, consisting of a leftward shift in the Ca²⁺-response curve (i.e. positive modulation) as well as a substantial direct activating effect, which is independent of

Ca^{2+} (close to 40% at low Ca^{2+} and in the presence of 1 μM of (-)-CM-TPMF). Thus, from a mode-of-action perspective, (-)-CM-TPMF belongs to the GW542573X functional class of $\text{K}_{\text{Ca}2.1}$ activators. Quantitatively, however, (-)-CM-TPMF is considerably more potent (> 100 fold) than GW542573X and the fraction of Ca^{2+} -independent activation induced by the compound is much larger (~40 % vs. 5 %).

$\text{K}_{\text{Ca}2}$ channels are inhibited by peptides like apamin and scyllatoxin as well as small molecules carrying positive charges mimicking the charges on apamin. These molecules bind from the outside, and despite an allosteric of apamin (Lamy et al., 2010), their inhibition is independent of the degree of channel activation by Ca^{2+} . In contrast, negative gating modulators like NS8593 interfere with the gating mechanism of $\text{K}_{\text{Ca}2}$ channels, causing reduced apparent affinity for Ca^{2+} . A characteristic feature of this mode-of-action is potent inhibition at low $[\text{Ca}^{2+}]_{\text{i}}$ and reduced effect at higher Ca^{2+} concentrations. The following experiment was designed in order to elucidate whether the inhibition by (-)-B-TPMF is dependent on the $[\text{Ca}^{2+}]_{\text{i}}$. Fig. 5A shows the time course of an inside-out experiment where the ability of 30 nM of (-)-B-TPMF to inhibit the $\text{K}_{\text{Ca}2.1}$ was evaluated at two different Ca^{2+} concentrations: 0.3 μM - equal to approximately 30% of maximal channel activity - and 10 μM Ca^{2+} corresponding to maximal activity. (-)-B-TPMF inhibited approximately 75 % of the current at the lower Ca^{2+} concentration, whereas it only inhibited about 20 % at 10 μM Ca^{2+} . In Fig. 5B the degree of inhibition by 30 nM (-)-B-TPMF at three Ca^{2+} concentrations (0.3-, 0.5-, and 10 μM) is quantified. The effect is clearly dependent on the intracellular Ca^{2+} concentration/channel activity and (-)-B-TPMF accordingly fulfills the criteria for being a negative gating modulator.

Due to the very similar structures of (-)-B-TPMF and (-)-CM-TPMF we envisioned that they might interact with a common binding site on $\text{K}_{\text{Ca}2.1}$. To test this hypothesis we conducted a

series of patch clamp experiments in order to elucidate possible competitive-like functional interactions. In the first series of experiments we investigated whether $K_{Ca2.1}$ activated by different concentrations (1 and 10 μM) of (-)-CM-TPMF would affect experimentally determined IC_{50} values for (-)-B-TPMF. These concentrations of (-)-CM-TPMF were chosen as they both induce near maximal activation of $K_{Ca2.1}$ (see Fig. 3C) at a permissive fixed $[\text{Ca}^{2+}]_i$ of 200 nM. Fig. 6A shows the I-V relations obtained in the presence of 1 μM (left panel) and 10 μM (right panel) of (-)-CM-TPMF. When 1 μM (-)-B-TPMF was co-applied, an approximate 60% reduction in the current activated by 1 μM (-)-CM-TPMF was obtained, whereas the same concentration of (-)-B-TPMF only inhibited 15% of the current activated by 10 μM (-)-CM-TPMF. Inhibition curves for (-)-B-TPMF were generated and Fig. 6B clearly shows that increasing the concentration of (-)-CM-TPMF from 1 to 10 μM resulted in an increase in the IC_{50} -value of (-)-B-TPMF from 0.43 μM to 2.96 μM . Although no rigorous comparison is possible it is noteworthy that the IC_{50} value for (-)-B-TPMF at 200 nM $[\text{Ca}^{2+}]_i$ in the absence of (-)-CM-TPMF is expected to be in the low nM range (Fig. 5). In the second series of experiments we attempted to prove that the increase in IC_{50} values in the presence of (-)-CM-TPMF is dependent of the presence of this compound *per se*, rather than being an effect of the high open state probability prevailing under the experimental conditions in Fig. 6. Therefore, we compared the inhibition by (-)-B-TPMF at high $[\text{Ca}^{2+}]_i$ (10 μM) with the inhibition at semi-maximal $[\text{Ca}^{2+}]_i$ (0.5 μM) in combination with 3 μM (-)-CM-TPMF resulting in similar degrees of activation (Fig. 7). The time-course of the experiment (Fig. 7A) shows that at 10 μM Ca^{2+} , (-)-B-TPMF (0.1 μM) inhibits the current reversibly by 75%. However, after wash-out and reactivation with 0.5 μM Ca^{2+} + 3 μM (-)-CM-TPMF, no effect was observed of 0.1 μM (-)-B-TPMF, and even a 100 times higher concentration was needed in order to get the same degree of inhibition as when

the channel was similarly activated by Ca^{2+} alone. Note also the strongly reduced rate of inhibition compared to the faster rate obtained at 0.1 μM (-)-B-TPMF at the first addition. We conclude from these experiments that (-)-B-TPMF and (-)-CM-TPMF interact functionally on the $\text{K}_{\text{Ca}2.1}$ channel in a manner likely to reflect direct competition on a common binding site.

The action of both positive and negative modulators of $\text{K}_{\text{Ca}2}$ channels depend on very specific amino acid residues. 1-EBIO and CyPPA interact with the intracellular C-terminal CaMBD, whereas NS8593 depends on transmembrane amino acids. The $\text{K}_{\text{Ca}2.1}$ -selective activator GW542573X acts via a single amino acid, Ser293, located in the S5 transmembrane region of $\text{K}_{\text{Ca}2.1}$. Due to the similar mode-of-action of GW542573X and (-)-CM-TPMF we conducted a series of experiments in order to investigate the possible involvement of Ser293 in the actions of both (-)-CM-TPMF and (-)-B-TPMF. Two point-mutated $\text{K}_{\text{Ca}2}$ channels were used, one ($\text{K}_{\text{Ca}2.1\text{S}293\text{L}}$) in which Ser293 in $\text{K}_{\text{Ca}2.1}$ is substituted with the equivalent leucine (Leu321 in $\text{K}_{\text{Ca}2.2}$ and Leu476 in $\text{K}_{\text{Ca}2.3}$) and one with the reverse mutation (Leu476 in $\text{K}_{\text{Ca}2.3}$ changed to serine, $\text{K}_{\text{Ca}2.3\text{L}476\text{S}}$; see Fig. 8A for approximate position in S5 and amino acid alignment of the S5 region of $\text{K}_{\text{Ca}2.1}$ and $\text{K}_{\text{Ca}2.3}$). Both channel constructs were activated by $[\text{Ca}^{2+}]_{\text{i}}$ within the normal range with EC_{50} values of $0.44 \pm 0.05 \mu\text{M}$ ($\text{K}_{\text{Ca}2.1\text{S}293\text{L}}$, $n = 3$) and $0.47 \pm 0.06 \mu\text{M}$ ($\text{K}_{\text{Ca}2.3\text{L}476\text{S}}$, $n = 4$). At 30 nM, a compound concentration that significantly modulated $\text{K}_{\text{Ca}2.1}$ but had limited effect on $\text{K}_{\text{Ca}2.3}$ (see Fig. 3), both (-)-CM-TPMF and (-)-B-TPMF failed to affect $\text{K}_{\text{Ca}2.1\text{S}293\text{L}}$ (Fig. 8B) but, on the other hand, significantly modulated the $\text{K}_{\text{Ca}2.3\text{L}476\text{S}}$ current level (Fig. 8C). Full concentration-response relationships were generated for (-)-B-TPMF (Fig. 8D) and (-)-CM-TPMF (Fig. 8E) on $\text{K}_{\text{Ca}2.1\text{S}293\text{L}}$ and $\text{K}_{\text{Ca}2.3\text{L}476\text{S}}$ in excised inside-out patches. For comparison, the best fit curves for $\text{K}_{\text{Ca}2.1}$ and $\text{K}_{\text{Ca}2.3}$, obtained from Fig. 3, are included as dotted lines. As seen from the figure, data obtained on $\text{K}_{\text{Ca}2.1\text{S}293\text{L}}$ was similar to those generated

on K_{Ca}2.3, thus demonstrating full loss-of-sensitivity both with respect to inhibition by (-)-B-TPMF and activation by (-)-CM-TPMF. Likewise, data obtained on K_{Ca}2.3_{L476S} was virtually identical to those on K_{Ca}2.1, demonstrating full gain-of-sensitivity of the K_{Ca}2.3_{L476S} construct for both compounds.

Discussion

We have described two structurally related compounds as new pharmacological modulators of Ca^{2+} -activated K^+ channels belonging to the KCNN gene family. Both are [1,2,4]triazolo[1,5-*a*]pyrimidines, a chemical series hitherto not described as having ion channel modulating properties. Both (-)-CM-TPMF and (-)-B-TPMF potently modulated the $\text{K}_{\text{Ca}2.1}$ subtype while having considerably weaker activity on $\text{K}_{\text{Ca}2.2}$, $\text{K}_{\text{Ca}2.3}$ and $\text{K}_{\text{Ca}3.1}$. (-)-CM-TPMF is a mixed opener/positive gating modulator, whereas the analogue (-)-B-TPMF is a negative gating modulator. Co-application experiments clearly revealed competition-like interactions between (-)-CM-TPMF and (-)-B-TPMF and the high-affinity actions of both compounds were shown to depend critically on the Ser293 positioned in the S5 segment as demonstrated by loss-of-sensitivity ($\text{K}_{\text{Ca}2.1\text{S293L}}$) as well as gain-of-sensitivity ($\text{K}_{\text{Ca}2.3\text{L476S}}$) mutations. Structurally, CM-TPMF and B-TPMF are remarkably similar and the pharmacology leading to either positive or negative gating modulation is determined entirely by the substituents on the terminal phenyl ring, a 4-*tert*-butyl group in B-TPMF or 4-chloro-2-methyl substitution in CM-TPMF. We are currently further exploring the structural components leading to either activation or inhibition and the general structure-activity relationship associated with chemical modification within this compound class (Eriksen et al., 2010a, b, and c). Interestingly, for each of these compounds the chirality introduced by the carbon atom in the linker separating the phenyl and the bicyclic heteroaromatic groups, led to pairs consisting of a highly potent (-)-enantiomer and a 40-100 fold less active (+)-isomer. Overall we conclude that $\text{K}_{\text{Ca}2}$ channels - in particular the $\text{K}_{\text{Ca}2.1}$ isoform - possess a common high-affinity binding-site for (-)-[1,2,4]triazolo[1,5-*a*]pyrimidines, here exemplified by (-)-CM-TPMF and (-)-B-TPMF, that pivotally influences the gating process in a facilitating or dampening way depending on minor changes in the substitution pattern of the

interacting molecules. This is the first demonstration of a common site for both positive and negative pharmacological modulation of gating in K_{Ca} channels, a phenomenon which is well established for the dihydropyridine pharmacology of L-type Ca^{2+} channels (i.e. nifedipine vs. Bay-k-8644, Greenberg et al., 1984) as well as for the benzodiazepine pharmacology of $GABA_A$ receptors (i.e. diazepam vs. DMCM, Sieghart, 1994).

It is interesting to compare the chemical structures of CM-TPMF/B-TPMF with the previously described (Hougaard et al., 2009) non-chiral and lower-potency $K_{Ca2.1}$ activator GW542573X (Fig. 1), which also depends on Ser293, and which has a qualitatively identical opener/positive modulator mode-of-action as described here for (-)-CM-TPMF. Both structural classes are relatively flexible, small-molecules of similar sizes and molecular weights, both classes contain a terminal phenyl ring, although with different substitution pattern, and both GW542573X and the [1,2,4]triazolo[1,5-*a*]pyrimidines have a single potential hydrogen bond donating group and 7-8 hydrogen bond acceptors. Still, it is evident that GW542573X represents a structurally very different compound class containing two carbamate functionalities of which one contains a terminal *tert*-butyl group also present in B-TPMF. Despite this apparent point of similarity it should be noted that GW542573X is an activator, whereas a closer structure activity analysis comprising further structural analogues (not shown) revealed that the *tert*-butyl group is crucial for the effect of B-TPMF as an inhibitor. Overall, based on the structural differences described above and also the fact that the small structural difference arising from chirality has such detrimental effects on the pharmacology in CM-TPMF and B-TPMF, we consider it unlikely that the [1,2,4]triazolo[1,5-*a*]pyrimidines described herein, would have exactly identical interaction points as GW542573X and share an identical binding site. Despite the parallels between GW542573X and (-)-CM-TPMF in terms of mode-of-action and interaction with Ser293, we

therefore cannot exclude that it actually interacts with a neighbouring site to the [1,2,4]triazolo[1,5-*a*]pyrimidines and just shares the Ser293 and the positive coupling to the gate with (-)-CM-TPMF as down-stream functional effects. From this follows also uncertainty, whether Ser293 *per se* actually constitutes part of the physical binding site for the [1,2,4]triazolo[1,5-*a*]pyrimidines.

Very recently a series of experiments exploring the apparent Ca^{2+} affinity of rK_{Ca}2.2 through mutational analysis demonstrated the complexity and the extended distribution of amino acids participating in or influencing the gating process in rK_{Ca}2.2 channels (Li and Aldrich, 2011): Charged residues in S6 were found to form a ring of positivity near the inner pore mouth that causes intrinsic rectification, but also influences the gating via an electrostatic mechanism, in particular by keeping the open state probability very low at zero Ca^{2+} . Charge reversal mutations of these amino acids strongly increase the open state probability at zero Ca^{2+} and very significantly left-shift the Ca^{2+} activation curve. Without in anyway suggesting a common causality, these findings are phenomenologically very similar to the effects of (-)-CM-TPMF and GW542573X, and serve to illustrate the point that amino acids in the transmembrane segments, far away from the CaM/CaMBD in the C-terminal, are pivotally important for gating and determining apparent Ca^{2+} -affinity of K_{Ca}2 channels.

The physiological role of the K_{Ca}2.1 subtype remains obscure in contrast to the K_{Ca}2.2 and K_{Ca}2.3 subtypes, which, through their contribution to action potential afterhyperpolarizations and functional coupling with the NMDA receptor, are involved in controlling firing-precision in pacemaking neurons and determining synaptic plasticity (Bond et al., 2005). K_{Ca}2.1 is predominantly expressed in the cortical/limbic structures of the brain, a distribution overlapping to a large extent with the expression of K_{Ca}2.2, but is essentially absent from the basal ganglia

and monoaminergic neurons, where $K_{Ca2.3}$ is the dominating subtype. Importantly, $K_{Ca2.1}$ is also the KCNN subtype with the most specific CNS expression and appears confined to neurons (Rimini et al., 2000). In contrast, the other KCNN members are also broadly expressed in the periphery on both neurons and somatic cell types and $K_{Ca2.3}$ is also present on glia cells (Armstrong et al., 2005). A serious obstacle towards exploration of the physiology of the $K_{Ca2.1}$ channel is the inability of the rat (and mouse) isoforms to express recombinantly in oocytes as well as mammalian cells. Furthermore, the phenotype of the $K_{Ca2.1}$ knock out mouse is still largely unknown, although it has been reported that the mAHP in CA1 is unaltered in these mice (Bond et al., 2004). An additional complication is the lack of subtype selective tools targeting $K_{Ca2.1}$. In contrast to GW542573X, which is not very potent ($EC_{50} \sim 7 \mu M$) both (-)-CM-TPMF and (-)-B-TPMF are high potency $K_{Ca2.1}$ modulators (~ 30 nM) which display acceptable selectivity over $K_{Ca2.2}$ and $K_{Ca2.3}$ and good to excellent selectivity over other ion channels and receptors. In particular, we envision that the “package” of the two very close analogues with opposite effects on $K_{Ca2.1}$ may constitute a useful combination for many experimental situations: A detailed look at the concentration-response curves in Fig. 3 show that (-)-CM-TPMF and (-)-B-TPMF can be applied up to 200 nM (causing 90 % inhibition and 40 % enhancement of $K_{Ca2.1}$) without significant functional activity on the other subtypes. (A note of caution: Due to the Ca^{2+} -dependent modulatory actions of both compounds the corresponding percentages for natively expressed channels may deviate slightly). The reversibility properties of both compounds (Fig. 2) are also promising with respect to their use in a sequential manner on the same cell, at least in isolated neuronal preparations. We tentatively suggest that the opposite effects of (-)-CM-TPMF and (-)-B-TPMF on biological responses obtained from complex in vitro or in vivo biological systems - possibly combined with lack of effects of their respective

(+)-enantiomers - would be a strong indication of the participation of the K_{Ca}2.1 isoform in that particular process.

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Legends to Figures

Fig. 1. Chemical structures of the negative $K_{Ca2.1}$ channel modulator B-TPMF and the positive modulators CM-TPMF and GW542573X.

Fig. 2. (-)-CM-TPMF acts as an activator and (-)-B-TPMF acts as an inhibitor of $K_{Ca2.1}$ channels. A, whole-cell current recorded from HEK293 cells stably expressing $K_{Ca2.1}$. I-V relationships were obtained at symmetrical $[K^+]$ and with the free $[Ca^{2+}]$ in the pipette solution buffered to 0.3 μ M. I-V relationships were measured upon application of 200-ms voltage ramps (-80 to +80 mV), elicited every 5 s from a holding potential of 0 mV in the absence (Ctrl) or in the presence of 10 nM of the compounds, (+)-B-TPMF, (-)-B-TPMF, (+)-CM-TPMF and (-)-CM-TPMF, as indicated. B, whole-cell $K_{Ca2.1}$ current at -75 mV obtained from the voltage ramps (A) as a function of time. The cell was exposed to 10 nM of the (-)- and (+)-forms of CM-TPMF and B-TPMF as indicated by the bars. C, average currents (mean \pm S.E.M.) in the presence of (-)- and (+)-forms of CM-TPMF and B-TPMF (10 nM). Current measured in the presence of compound is shown relative to the current level prior to compound addition (control; $n = 4 - 11$). (-)-B-TPMF was more potent than (+)-B-TPMF ($p < 0.001$, students t-test) in inhibiting $K_{Ca2.1}$ current, while (-)-CM-TPMF was more potent than (+)-CM-TPMF in potentiating the current ($p < 0.001$, students t-test).

Fig. 3. (-)-B-TPMF is a $K_{Ca2.1}$ subtype selective inhibitor and (-)-CM-TPMF a $K_{Ca2.1}$ selective activator. A, I-V relationships measured from inside-out patches obtained from HEK293 cells stably expressing h $K_{Ca2.1}$, h $K_{Ca2.2}$ or h $K_{Ca2.3}$. Each I-V plot shows a control trace, a trace obtained in the presence of 30 nM (-)-B-TPMF and a trace in the presence of 30 nM (-)-CM-

TPMF. The free $[Ca^{2+}]$ in the bath/intracellular solution was 0.3 μM (resulting in approximately 30% of the maximal current level) to allow for both positive and negative modulation. B, concentration-response relationships of (-)-B-TPMF on the three K_{Ca2} subtypes. Residual current is depicted as a function of the (-)-B-TPMF concentration. Currents were measured from inside-out patches at 0.5 μM free Ca^{2+} (approximately 90% of the maximal current level) and each data point is the mean \pm S.E.M. of 5-7 experiments. The solid lines are the fit of data to the Hill equation yielding the following IC_{50} values and Hill coefficients: $K_{Ca2.1}$: 0.031 μM and -1.2; $K_{Ca2.2}$: 1.05 μM and -1.7; $K_{Ca2.3}$: 1.32 μM and -1.7. C, concentration-response relationships of (-)-CM-TPMF on $K_{Ca2.1}$, $K_{Ca2.2}$ and $K_{Ca2.3}$. Currents were measured from inside-out patches at 0.2 μM free Ca^{2+} (approximately 15% activation) and shown relative to the current level obtained at 10 μM Ca^{2+} in the absence of compound. Each data point is the mean \pm S.E.M. of 6-8 experiments and the solid lines are the fit of data to the Hill equation yielding the following EC_{50} values and Hill coefficients $K_{Ca2.1}$: 0.024 μM and 0.8; $K_{Ca2.2}$: 0.29 μM and 1.5; $K_{Ca2.3}$: 0.25 μM and 1.1. Efficacy with respect to saturating Ca^{2+} was in the range 83-87%.

Fig. 4. (-)-CM-TPMF is a K_{Ca2} channel opener and a positive modulator. A, $K_{Ca2.1}$ current at -75 mV, measured from an inside-out patch and depicted as a function of time. The patch was exposed to 0.1 μM (-)-CM-TPMF either in the absence of Ca^{2+} (0), at 0.2 - or at 10 μM Ca^{2+} . B, I-V relationships at 0, 0.2 - or 10 μM Ca^{2+} in the absence (Ctrl) or presence of 0.1 μM (-)-CM-TPMF as indicated. C, Ca^{2+} -response curves for $K_{Ca2.1}$ measured from inside-out patches either in the absence (Ctrl) or in the presence of (-)-CM-TPMF (0.1 μM or 1 μM). Currents from individual patches were normalized with respect to the effect of 10 μM Ca^{2+} . Data points are the mean \pm S.E.M. of at least 6 experiments. The lines are the fit of data to the Hill equation yielding

the following EC_{50} values and Hill coefficients: Ctrl: 0.32 μ M and 4.4, 0.1 μ M (-)-CM-TPMF: 0.15 μ M and 2.9, 1 μ M (-)-CM-TPMF: 0.13 μ M and 2.8. Note the substantial Ca^{2+} -independent opener effect of (-)-CM-TPMF at very low Ca^{2+} concentrations ($14 \pm 3\%$ and $32 \pm 5\%$ of maximal activity at 0.1 μ M and 1 μ M (-)-CM-TPMF) and the higher efficacy at saturating Ca^{2+} ($I/I_{max} > 1$) in the presence of (-)-CM-TPMF.

Fig. 5. (-)-B-TPMF inhibits the $K_{Ca2.1}$ current in a Ca^{2+} -dependent manner. A, $K_{Ca2.1}$ current at -75 mV plotted versus time. Current was measured from an inside-out patch exposed to 30 nM (-)-B-TPMF either at 0.3 or 10 μ M Ca^{2+} . As control for the background/leak current level, periods with 0 free Ca^{2+} were included. B, % current inhibition induced by (-)-B-TPMF (30 nM) at the different Ca^{2+} concentrations. Data points are the mean \pm S.E.M. of 5-6 experiments at each Ca^{2+} concentration.

Fig. 6. Maximal (-)-CM-TPMF activation concentration-dependently decreases the apparent affinity of the inhibitor (-)-B-TPMF. A, I-V relationships measured from inside-out patches obtained from HEK293 cells stably expressing $hK_{Ca2.1}$ at a free $[Ca^{2+}]$ of 0.2 μ M. I-V curves were obtained in the presence of 1 or 10 μ M (-)-CM-TPMF alone or in combination with 1 μ M (-)-B-TPMF. B, concentration-response relationship of (-)-B-TPMF in the presence of 1 or 10 μ M (-)-CM-TPMF. Residual current is depicted as a function of the (-)-B-TPMF concentration. Each data point is the mean \pm S.E.M. of 4-9 experiments and the solid lines are the fit of data to the Hill equation resulting in the following IC_{50} values and Hill coefficients: 0.43 μ M and -0.9 (at 1 μ M (-)-CM-TPMF) and 2.96 μ M and -1.4 (in the presence of 10 μ M (-)-CM-TPMF).

Fig. 7. The reduced apparent affinity of (-)-B-TPMF is due to (-)-CM-TPMF competition, not the degree of channel activation. A, $K_{Ca2.1}$ current measured at -75 mV as a function of time. The inside of the patch was exposed to a $[Ca^{2+}]$ of 0, 0.5 or 10 μM as indicated. (-)-B-TPMF was added once steady current activation was obtained by 10 μM Ca^{2+} alone or by the combination of 0.5 μM Ca^{2+} and 3 μM (-)-CM-TPMF. B, I-V relationships measured at the time points indicated by letters in A. Data is from a single experiment representative of 4 independent experiments.

Fig. 8. The activity of (-)-B-TPMF and (-)-CM-TPMF is critically dependent on Ser293 in $K_{Ca2.1}$. A, Schematic illustrating the approximate S5 location of the single amino acid that is mutated in the channel constructs. In $K_{Ca2.1}$, serine is replaced with leucine, which is the equivalent amino acid in $K_{Ca2.3}$, and in $K_{Ca2.3}$ leucine is replaced with serine. Amino acid alignment of the S5 region of $K_{Ca2.1}$ and $K_{Ca2.3}$ is shown on the right. B and C, I-V relationships measured from inside-out patches obtained from HEK293 cells transiently transfected with $K_{Ca2.1S293L}$ (B) or $K_{Ca2.3L476S}$ (C). Each plot shows a trace in the absence of compound (Ctrl), a trace in the presence of 30 nM (-)-B-TPMF and one in the presence of 30 nM (-)-CM-TPMF. The free $[Ca^{2+}]$ in the bath/intracellular solution was 0.3 μM to allow for both positive and negative modulation of the current level. D, concentration-response relationship of (-)-B-TPMF on $K_{Ca2.1S293L}$ and $K_{Ca2.3L476S}$. Residual current in the presence of (-)-B-TPMF is depicted as a function of the (-)-B-TPMF concentration. Currents were measured from inside-out patches at 0.5 μM free Ca^{2+} and each data point is the mean \pm S.E.M. of 5-6 experiments. The solid lines are the fit of data to the Hill equation yielding the following IC_{50} values and Hill coefficients: $hK_{Ca2.1S293L}$: 0.81 μM and -1.2, $K_{Ca2.3L476S}$: 0.05 μM and -1.6. For illustrative purposes, data obtained on wild type $K_{Ca2.1}$ and $K_{Ca2.3}$ is included as dotted lines. E,

concentration-response relationship of (-)-CM-TPMF on $K_{Ca2.1S293L}$ and $K_{Ca2.3L476S}$. Currents were measured at 0.2 μM free Ca^{2+} and depicted relative to the current level obtained at 10 μM Ca^{2+} and in the absence of compound. Data points are the mean \pm S.E.M. of 4-5 experiments and the solid lines are the fit of data to the Hill equation: $hK_{Ca2.1S293L}$: 0.7 μM and 1.3, $K_{Ca2.3L476S}$: 0.03 μM and 1.2. Efficacy with respect to maximal Ca^{2+} was 89% ($hK_{Ca2.1S293L}$) and 85% ($K_{Ca2.3L476S}$). Again, for illustrative purposes, data on $K_{Ca2.1}$ and $K_{Ca2.3}$ are included as dotted lines.

Table 1 Effects of CM-TPMF and B-TPMF at various ion channels

^a at 0.2 μM $[\text{Ca}^{2+}]_i$ ^b at 0.5 μM $[\text{Ca}^{2+}]_i$		(-)-CM-TPMF		(+)-CM-TPMF	(-)-B-TPMF		(+)-B-TPMF
		EC ₅₀ (μM) ^a	Hill slope		IC ₅₀ (μM) ^b	Hill slope	
Inside-out	hK _{Ca} 2.1	0.024	0.8		0.031	-1.2	
	hK _{Ca} 2.2	0.29	1.5		1.05	-1.7	
	hK _{Ca} 2.3	0.25	1.1		1.32	-1.7	
	hK _{Ca} 2.3 _{L476S}	0.03	1.2		0.05	-1.6	
Whole cell		SC ₁₀₀ in μM (n)		SC ₁₀₀ in μM (n)	K _d in μM (n)		K _d in μM (n)
	hK _{Ca} 2.1	0.0053 \pm 0.00021 (6)		0.19 \pm 0.044 (5)	0.015 \pm 0.0067 (5)		1.6 \pm 0.54 (5)
	hK _{Ca} 2.2	0.17 \pm 0.05 (4)		2.6 \pm 0.50 (4)	0.29 \pm 0.039 (4)		13 \pm 3.2 (4)
	hK _{Ca} 2.3	0.047 \pm 0.0049 (6)		1.5 \pm 0.07 (4)	0.23 \pm 0.05 (4)		10 \pm 2.3 (8)
	hK _{Ca} 3.1	0.12 \pm 0.0049 (5)		2.1 \pm 0.12 (4)	SC ₁₀₀ = 5.2 \pm 1.0 μM (6)		35 \pm 3.5 (3)
		% inhibition at 10 μM (n)			% inhibition at 10 μM (n)		
	hK _{Ca} 1.1	39 \pm 3.0 (3)			20 \pm 5.2 (3)		
	hK _v 7.2+7.3	1.3 \pm 1.3 (4)			12 \pm 2.5 (4)		
	hK _v 11.1	37 \pm 3.9 (3)			36 \pm 4.0 (3)		
	rNa _v 1.2	8.3 \pm 2.8 (4)			26 \pm 3.1 (3)		

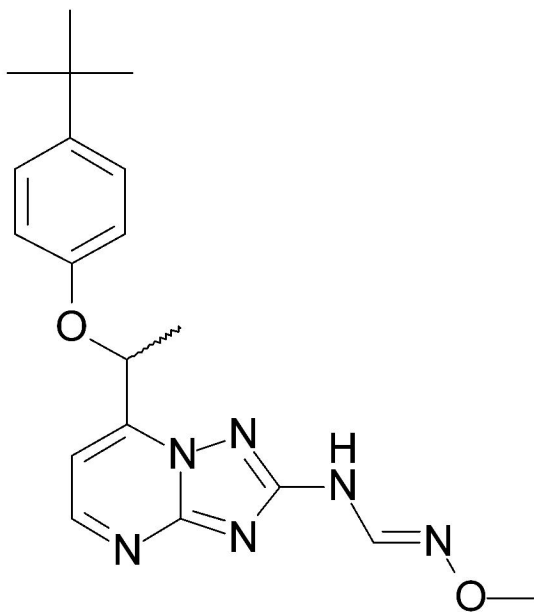
Compiled effects of (-) and (+) enantiomers of both CM-TPMF and B-TPMF on different cloned ion channels and constructs.

Different potency/effect measures (EC₅₀, IC₅₀, K_d, SC₁₀₀, and % inhibition) have been applied as explained in Materials and Methods.

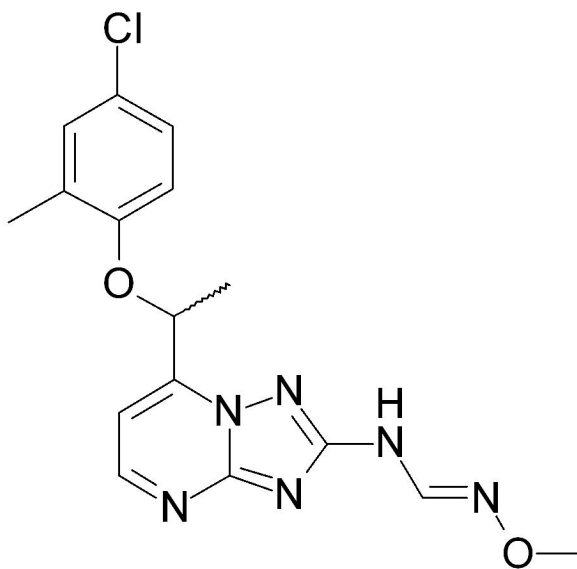
^a and ^b denotes the calculated free Ca²⁺-concentrations used in the inside-out experiments. The h and r prefixes mean human and rat.

The values are means \pm S.E.M, n is the number of independent experiments.

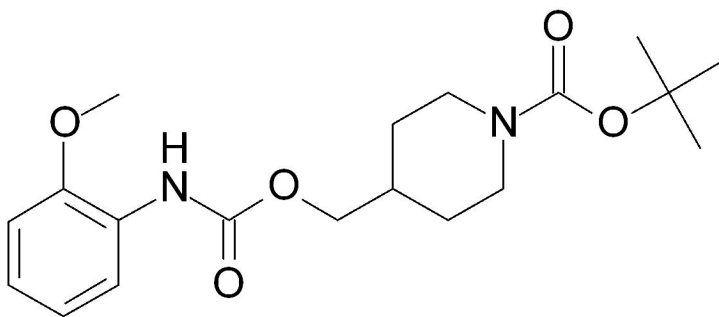
Figure 1



B-TPMF
inhibitor



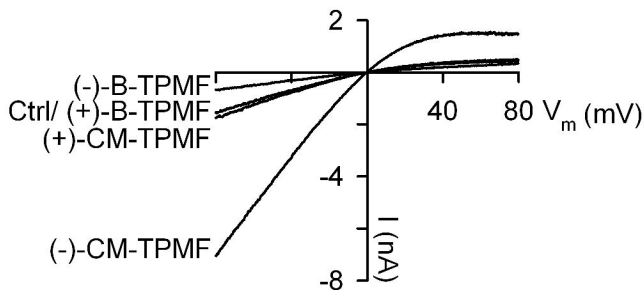
CM-TPMF
activator



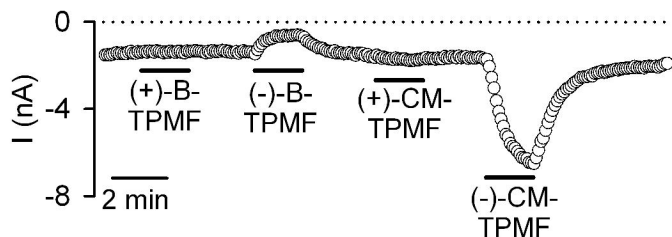
GW542573X
activator

Figure 2

A



B



C

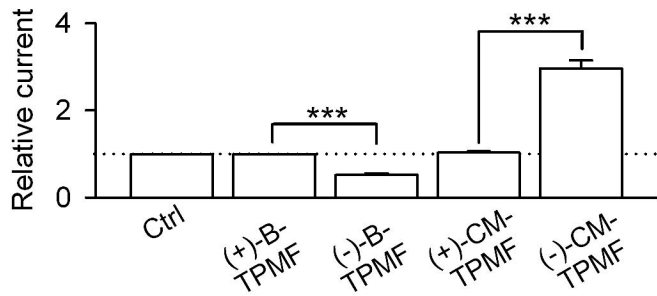
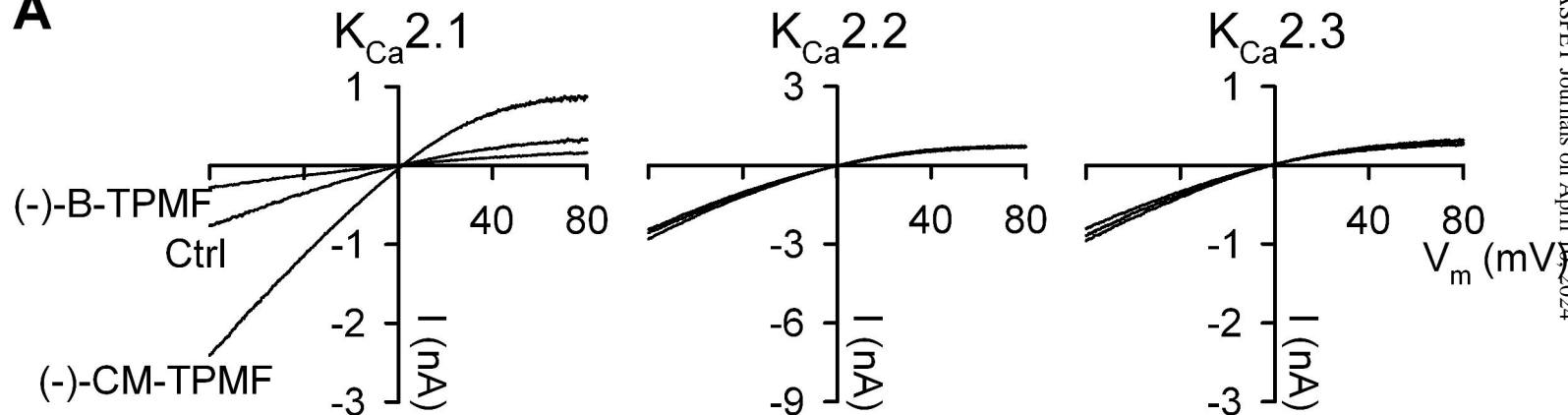
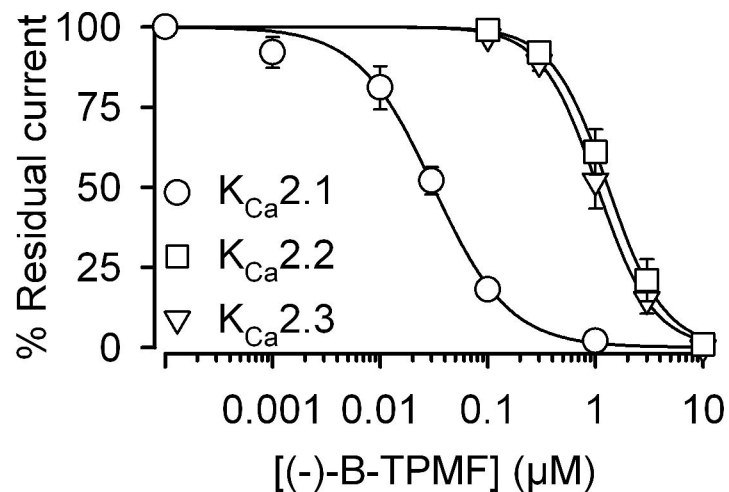


Figure 3

A



B



C

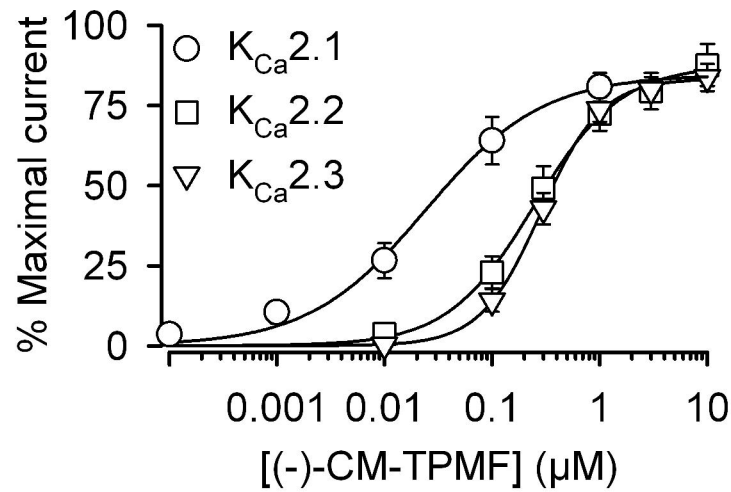


Figure 4

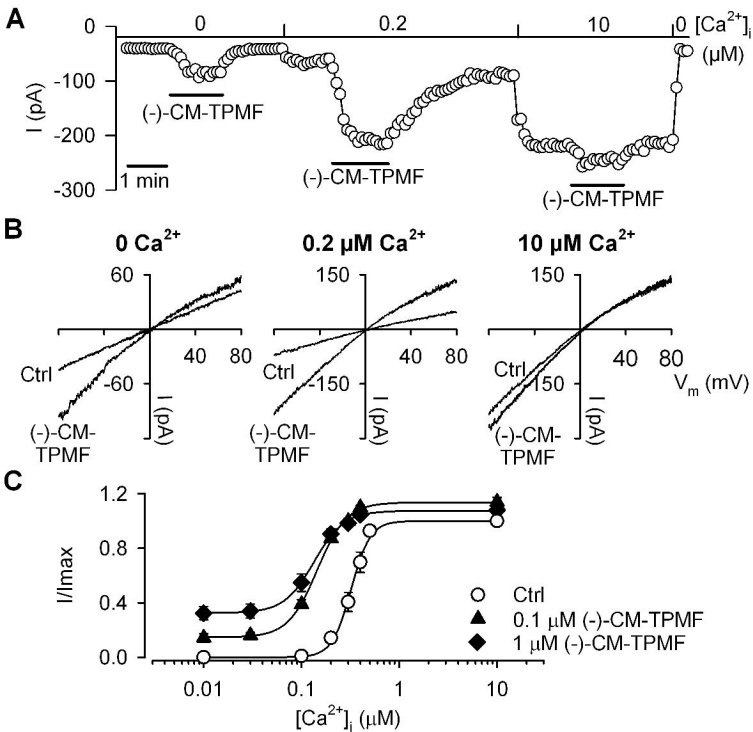
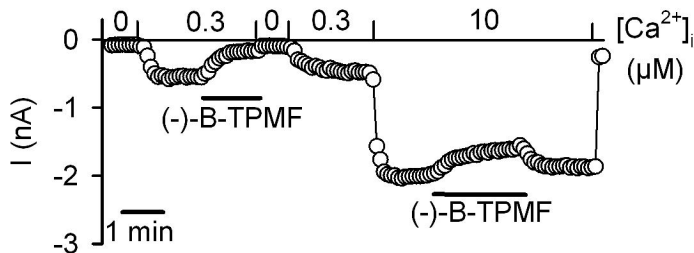


Figure 5

A



B

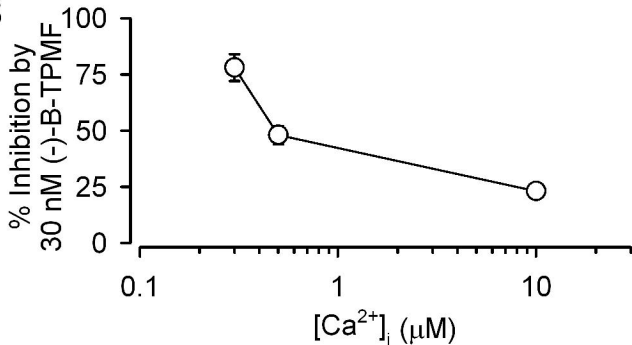


Figure 6

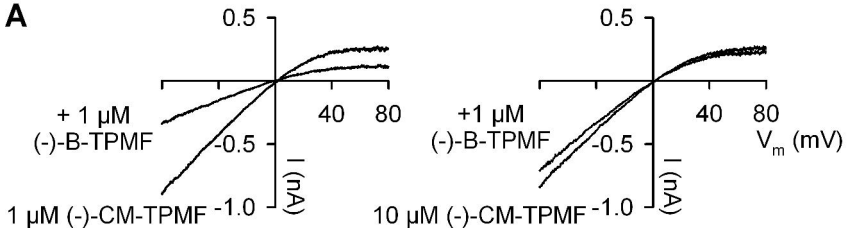
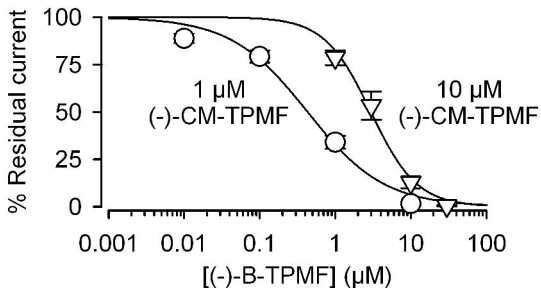
A**B**

Figure 7

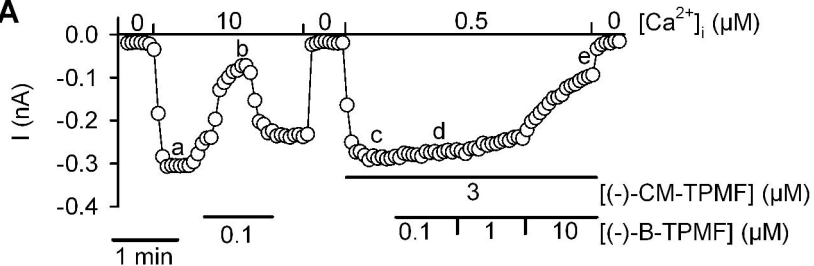
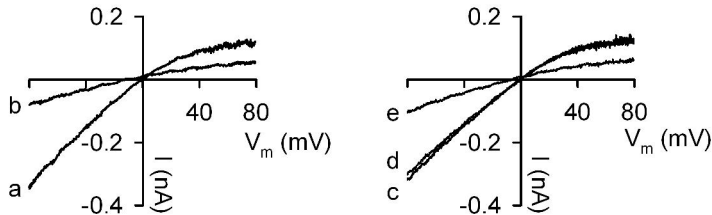
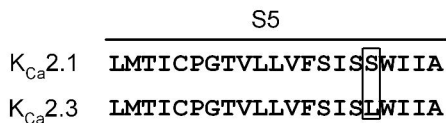
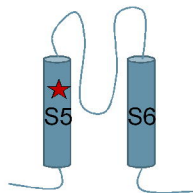
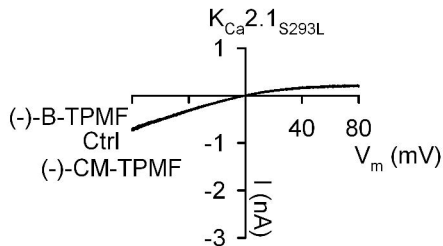
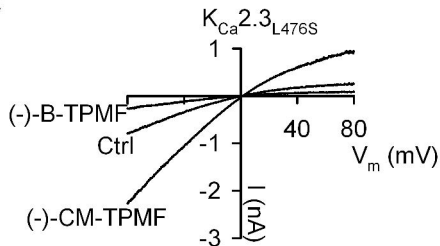
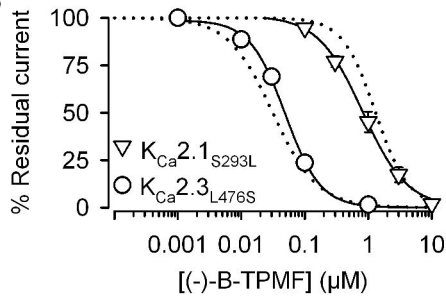
A**B**

Figure 8

A**B****C****D****E**