Selective Activation of the SK1 Subtype of Human Small-Conductance Ca²⁺-Activated K⁺ Channels by 4-(2-Methoxyphenylcarbamoyloxymethyl)-piperidine-1-carboxylic Acid tert-Butyl Ester (GW542573X) Is Dependent on Serine 293 in the S5 Segment

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ABSTRACT

A new small molecule, 4-(2-methoxy-phenylcarbamoyloxymethyl)piperidine-1-carboxylic acid tert-butyl ester (GW542573X), is presented as an activator of small-conductance Ca²⁺-activated K⁺ (SK, K_{Ca}2) channels and distinguished from previously published positive modulators of SK channels, such as 1-ethyl-2benzimidazolinone (1-EBIO) and cyclohexyl-[2-(3,5-dimethylpyrazol-1-yl)-6-methyl-pyrimidin-4-yl]-amine (CyPPA), in several aspects. GW542573X is the first SK1-selective compound described: an EC₅₀ value of 8.2 \pm 0.8 μ M (n = 6, [Ca²⁺]_i = 200 nM) was obtained from inside-out patches excised from hSK1expressing HEK293 cells. Whole-cell experiments showed that hSK2 and hSK3 channels were more than 10 times, and hIK channels even more than 100 times, less sensitive to GW542573X. The Ca²⁺-response curve of hSK1 was left-shifted from an EC₅₀ (Ca²⁺) value of 410 \pm 20 nM (n = 9) to 240 \pm 10 nM

(n = 5) in the presence of 10 μ M GW542573X. In addition to this positive modulation, GW542573X activated SK1 in the absence of Ca²⁺ and furthermore induced a 15% increase in the maximal current at saturating Ca2+. Thus, GW542573X also acts as a genuine opener of the hSK1 channels, a mechanism of action (MOA) not previously obtained with SK channels. The differential potency on hSK1 and hSK3 enabled a chimera approach to elucidate site(s) important for this new MOA and selectivity property. A single amino acid (Ser293) located in S5 of hSK1 was essential, and substituting the corresponding Leu476 in hSK3 with serine conferred hSK1-like potency (EC $_{50} = 9.3 \pm 1.4 ~\mu M, n = 5$). GW542573X may activate SK channels via interaction with "deeppore" gating structures at the inner pore vestibule or the selectivity filter in contrast to 1-EBIO and CyPPA that exert positive modulation via the intracellular calmodulin binding domain.

Small-conductance Ca2+-activated K+ channels (SK channels) are voltage-independent channels that are gated solely by intracellular Ca²⁺. Three subtypes of SK channels have been described previously (SK1-SK3, K_{Ca}2.1-K_{Ca}2.3; Stocker, 2004), which, together with the related Ca²⁺-activated K^+ channel of intermediate conductance (IK, $K_{Ca}3.1$),

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acquire their Ca2+-sensitivity from C-terminally attached calmodulin (CaM) (Köhler et al., 1996; Xia et al., 1998; Khanna et al., 1999). All SK subtypes possess nearly identical Ca²⁺-response curves (EC₅₀ \sim 400 nM; $n_{\rm H}$ \sim 5). The high Hill value may indicate cooperative binding of Ca²⁺, because each of the four SK α -subunits is associated with CaM capable of binding up to two Ca²⁺ (Schumacher et al., 2001). Fast and high-affinity responses to changes in the intracellular Ca²⁺ concentration ([Ca²⁺]_i) are the basis for functional cou-

ABBREVIATIONS: SK channel, small conductance Ca²⁺-activated K⁺ channel; BMB, bicuculline methobromide; CaM, calmodulin; CaMBD, calmodulin binding domain; CHO, Chinese hamster ovary; CK2, casein kinase 2; CyPPA, cyclohexyl-[2-(3,5-dimethyl-pyrazol-1-yl)-6-methylpyrimidin-4-yl]-amine; 1-EBIO, 1-ethyl-2-benzimidazolinone; GW542573X, 4-(2-methoxy-phenylcarbamoyloxymethyl)-piperidine-1-carboxylic acid tert-butyl ester; HEK, human embryonic kidney; IK channel, intermediate conductance Ca2+-activated K+ channel; MOA, mode of action; NS309, 6,7-dichloro-1H-indole-2,3-dione 3-oxime; NS8593, (R)-N-(benzimidazol-2-yl)-1,2,3,4-tetrahydro-1-naphtylamine; PP2A, phosphatase 2A; S, transmembrane segment; SKA-31, naphtha[1,2-d]thiazol-2-ylamine; VIPR, voltage ion probe reader; PCR, polymerase chain reaction; ANOVA, analysis of variance; DC-EBIO, 5,6-dichloro-1-ethyl-1,3-dihydro-2H-benzimidazol-2-one; ChTX, charybdotoxin; MTSEA, 2-aminoethyl methane thiosulfonate hydrobromide.

pling of SK channels to specific localized Ca²⁺ transients in different neurons and for their participation in neurophysiological processes like spike-frequency adaptation, pacemaking, and synaptic integration and plasticity (Bond et al., 2005; Pedarzani and Stocker, 2008).

Gating of SK channels can be modulated by physiological and pharmacological means. SK2 and SK3 channels are part of multiheteromeric protein complexes consisting of CaM, casein kinase 2, phosphatase 2A (PP2A), and scaffolding proteins. Casein kinase 2 and PP2A oppositely regulate the apparent Ca2+ sensitivity by phosphorylating/dephosphorylating CaM at threonine 80 (Bildl et al., 2004; Allen et al., 2007). This system has been shown to respond to the activation of noradrenaline receptors causing phosphorylation of CaM, reduced SK channel Ca²⁺ sensitivity, and functional uncoupling to Ca²⁺ influx mediated by P/Q Ca²⁺ channels (Maingret et al., 2008). Similar processes may underlie the σ -1 receptor-mediated inhibition of SK channels that facilitate the induction of long-term potentiation in the hippocampus (Martina et al., 2007). Small molecules like NS8593 (Strøbaek et al., 2006) and its analogs (Sørensen et al., 2008) also possess the ability to inhibit SK channels by shifting the Ca2+-response curve to the right (apparent reduced Ca²⁺ sensitivity), an inhibitory principle called negative-gating modulation. This differs mechanistically from the action of apamin, a selective-pore blocker of SK channels. In contrast, 1-EBIO, NS309, and SKA-31 positively modulate IK and SK channels (IK > SK1 = SK2 = SK3) by shifting their Ca²⁺-response curves to the left (apparent increased Ca²⁺ sensitivity) without exerting agonism, per se (Pedarzani et al., 2001; Hougaard et al., 2007; Sankaranarayanan et al., 2009). Even though clear molecular and mechanistic differences exist, positive gating modulation may qualitatively resemble the physiological effects of PP2A-mediated increased Ca2+ sensitivity described for SK2/SK3 channels. Despite the highly homologous gating mechanisms shared by all SK subtypes, pharmacological modulation can be subtype-selective, as illustrated by the compound CyPPA, which is active on hSK3 and hSK2 channels, whereas it is inactive on hSK1 or hIK ($hSK3 > hSK2 \gg hSK1 = hIK$) (Hougaard et al., 2007). The effects of CyPPA and 1-EBIO have been shown, from their actions on hSK3/hSK1 or SK2/IK chimeras, to depend on the CaMBD in the C terminus.

Here we describe a new activator of hSK channels, GW542573X, which is unique in several respects. First, this compound demonstrates a novel selectivity profile, being selective for hSK1 over hSK2 and hSK3 and virtually inactive on hIK (hSK1 > hSK2 = hSK3 > hIK). Second, it partially activates hSK1, even in the absence of intracellular Ca²⁺. Third, this compound does not act via the intracellular CaMBD but via a transmembrane segment (S5). Fourth, remarkably, loss of sensitivity is achievable by substituting the hSK1 serine residue with the equivalent hSK3 leucine residue (S293L), whereas gain of sensitivity is achievable by the reverse mutation in hSK3 (L476S). Thus, GW542573X is the first representative of a new chemical and functional class of SK channel activators. The identification of a molecular determinant for SK channel activation positioned close to the inner pore cavity and selectivity filter is discussed in relation to the proposed "deep pore" mechanism for gating of SK channels (Bruening-Wright et al., 2002, 2007).

Materials and Methods

Chemistry. GW542573X was prepared at GlaxoSmithKline (Uxbridge, Middlesex, UK) by the following procedure. 4-Hydroxymethyl-piperidine-1-carboxylic acid tert-butyl ester (1.0 equivalent) in anhydrous dichloromethane at 0°C under a nitrogen atmosphere was treated with sodium hydride (1.05 equivalent) and then stirred at 0°C for 1 h. 2-Methoxyphenyl isocyanate (1.0 equivalent) was added, and the reaction was stirred for 18 h at ambient temperature. The reaction was quenched with saturated aqueous sodium bicarbonate solution, diluted with an equal volume of dichloromethane, and washed with saturated aqueous sodium bicarbonate solution. The organic layer was separated, dried over anhydrous MgSO4, and evaporated in vacuo. The crude product was purified in small quantities by mass-directed high-performance liquid chromatography, or in larger quantity by silica chromatography, eluting with cyclohexane/ethyl acetate (9:1). The purified product was obtained as a colorless oil, which slowly crystallized to a white solid on standing: MH^{+} 365 (100%), MNH_{4}^{+} 382 (100%), 265 (40%) (Dale et al., 2004).

NS309 (Strøbaek et al., 2004) was synthesized at NeuroSearch A/S (Ballerup, Denmark). GW542573X and NS309 were dissolved in dimethyl sulfoxide and diluted to the final concentration in the experimental solution. Apamin (Sigma-Aldrich, Brøndby, Denmark) was dissolved in the experimental solution and bicuculline methobromide (Biomedicals Inc., Illkirch, France) in dimethyl sulfoxide.

Cell Cultures. For the patch-clamp experiments, HEK293 cell lines stably expressing hSK1, hSK2, hSK3, and hIK channels were used. cDNA encoding hSK1 was constructed as described by Strøbaek et al. (2000) (Swiss-Prot Q92952; UniProtKB, Geneva, Switzerland) and was subcloned into the custom-made expression vector pNS1n derived from pcDNA3. Stable hSK1 HEK293 cell lines were isolated using G418 as a selection marker. pcDNA3_hSK2, a kind gift from B. Attali (Desai et al., 2000; Swiss-Prot Q9H2S1, UniProtKB), was used to generate HEK293 cells stably expressing hSK2. hSK3-expressing HEK293 cell lines were based on the plasmid pNS3n_hSK3 described previously (Strøbaek et al., 2004; Swiss-Prot Q9UG16, UniProtKB). cDNA-encoding hIK (Jensen et al., 1998; Swiss-Prot O15554, UniProtKB) was subcloned using EcoRI and XhoI into pNS1n, and stable HEK293 cell lines were selected using G418. Expression of hSK1, hSK2, hSK3, and hIK were verified by patch-clamp measurements. Channel chimeras and point-mutated channels (see below) were transiently transfected into HEK293 cells using Lipofectamine (Invitrogen, Carlsbad, CA) and standard transfection methods. Electrophysiological measurements were performed 2 to 3 days after the transfection. Cells were cultured in Dulbecco's modified Eagle's medium (Sigma-Aldrich) supplemented with 10% fetal calf serum (Invitrogen) at 37°C and 5% CO2. At approximately 75% confluence, the cells were washed once with phosphate-buffered saline, harvested by trypsin/EDTA (Sigma-Aldrich) treatment, and transferred to Petri dishes containing coverslips (3.5-mm diameter; VWR International, Herley, Denmark).

For the voltage ion probe reader (VIPR) experiments, hSK1-CHO cells were used (Dale et al., 2002). Stable cell lines were cultured under selection for resistance to 500 $\mu \rm g/ml$ Geneticin in α -minimal essential medium (Invitrogen) with 10% heat-inactivated fetal bovine serum (Invitrogen) at 37°C and 5% CO $_2$. Two to 4 days before the experiment, cells were plated in black-walled clear-bottomed sterile 96-well plates (Costar; Corning Life Sciences, Acton, MA) at a density of 50,000 cells/well.

Molecular Biology. Three unique restriction sites were introduced in conserved regions of hSK1 and hSK3 at the following amino acid positions: SalI at hSK1 Leu148 and hSK3 Leu331; HinDIII at hSK1 Ser190 and hSK3 Ser373; and AgeI at hSK1 Thr350 and hSK3 Thr533. In brief, hSK1 and hSK3 cDNAs were subcloned into the pcDNA3-derived vectors pNS3n and pNS3h, respectively, using EcoRI and BamHI. All restriction enzymes were purchased from New England Biolabs (Ipswich, MA). These plasmids were uracilated via *Escherichia coli* RZ1032 (Strat-

agene, La Jolla, CA) and used as templates in mutagenesis reactions. Six oligonucleotides (MWG Biotech, Ebersberg, Germany) were used to introduce the unique restriction sites: hSK1SalI, ATGCCTCATCAGCCTgTCgACGGCCATCCTGCTGGG; hSK1HinDIII, GCGCGTGTTCCTCATaagct-TAGAGCTGGCAGTGTGC; hSK1AgeI, GTGTGTGCCTGCTCACcGGtATCATGGGAGCTGGC; hSK3SalI, CCTTATCAGTCTGTCgaccatcatccttttg; hSK3HinDIII, GAGCGCATCCTGTACATaAGCtTGGAGATGCTGGTG; and hSK3AgeI, GTGTCTGTCTCCTCACcGGt-ATCATGGGTGCAGGC.

The mutagenesis reactions were performed using T7 DNA polymerase and T4 DNA ligase (New England Biolabs). E. coli XL1-Blue (Stratagene) was transformed with an aliquot of the reaction mixture, and the resulting plasmid DNA was purified using standard methods. Chimera 1 and 2 were assembled using the AgeI site, and chimera 3 was generated by first introducing the hSK3 part between SalI and HinDIII into chimera 2 and then to introduce the N-terminal tail of hSK3 with overlapping PCR (Expand High-Fidelity PCR System; Roche, Mannheim, Germany) using the following oligonucleotides: hsk3-hSK1s, agtgactatgctCTCATTTTCGGCAT-GTTT; and hsk1-hSK3as, gccgaaaatgagAGCATAGTCACTCAGTCG, introducing the overlapping PCR product using an upstream polylinker restriction site and SalI. Chimera 4 was generated by joining N-terminal hSK3 and chimera 2 in the HinDIII site. Chimeras 5 to 9 were made by performing mutagenesis on Chimera 3 using the following oligonucleotides: chimera 5, CATAAGCTTAGAGCTactAGTGTGCGCCATTCACCCGaTcCCCGGCgAgTACCGCTTCACGTGG; chimera 6, GCGCGGCTGGCCTTCtCaTAtGCGCCCTCGGTGGCCG-AGGCCGACGTGGAtaTcaTaCTGTCCATCCCCATG; chimera 7, CCTGCGCCTCTACCTGaTcGcCCGGGTGATGCTACTG; chimera 8, GGTGATGCTACTGCAttcgAAAcTCTTCACGGACGCCTC and GGCCCTCAACAAGATtAatTTCAACACGCGCTTC: chimera 9. GTGCTGCTGGTCTTCtcgATaTCCctCTGGATCATCGCAGC; and chimera 10 was generated by mutagenesis on hSK3 using the oligonucleotide CTGTGCTGCTCGTGTTCtcgATaTCTtcGTGGATCAT-TGCTGCCTG. All constructs were verified by sequencing (MWG Biotech).

Membrane Potential Assays. The membrane potential of hSK1-CHO cells was monitored using fluorescence resonance energy transfer between a phospholipid-anchored coumarin donor CC-2-dimyristoyl phosphatidylethanolamine and oxonol acceptor bis-(1,3-diethylthiobarbituric acid) trimethine oxonol on a VIPR (Aurora Biosciences, San Diego, CA) (González et al., 1999). Experiments were performed at room temperature by first washing the cells with an external buffer containing 145 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, 20 mM HEPES, and 10 mM glucose, pH 7.4, and then incubating with CC-2-dimyristoyl phosphatidylethanolamine (8 µM diluted in external buffer) for 30 min. Cells were then washed to remove unbound coumarin before the addition of bis-(1,3-diethylthiobarbituric acid) trimethine oxonol (8 µM in external buffer for 45 min) and ESS-Cy4 (350 μM), a background signal fluorescence quenching reagent. The coumarin fluorophore was excited at 405 nM using a xenon lamp, and dual emission was measured each second at 460 nm (coumarin) and 580 nm (oxonol). The signal ratio at the two wavelengths was calculated as a measure of the fluorescence resonance energy transfer between the two fluorophores. After five consecutive baseline readings, GW542573X (100 nM to 100 µM) was added online, and fluorescence was monitored for up to a further 5 min. To determine final Rf/Ri ratios, fluorescence values from cellfree wells of the microtiter plate were first subtracted.

Electrophysiology. Membrane currents were recorded under voltage-clamped conditions using the whole-cell or the inside-out configuration of the patch-clamp technique. Cells seeded on coverslips were transferred to a 15- μ l recording chamber continuously perfused at 1 ml/min at room temperature. The chamber was grounded by an integrated Ag/AgCl pellet. Patch pipettes (approximately 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. Experiments were controlled by an EPC-9 amplifier (HEKA, Lambrecht/Pfalz, Germany) connected to a Macintosh computer (Apple Computer, Cupertino, CA) by an ITC-16 interface. 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 ($R_{\rm s}$ less than 8 $M\Omega,\,80\%$ compensation) were updated before each voltage ramp.

In all experiments, a high K^+ solution was applied to the extracellular side of the membrane 154 mM KCl, 2 mM CaCl $_2$, 1 mM MgCl $_2$, and 10 mM HEPES, pH adjusted to 7.4 with KOH. The intracellular solutions contained 154 mM KCl, 10 mM HEPES, and 10 mM EGTA, or a combination of EGTA and nitrilotriacetic acid (10 mM in total). Concentrations of MgCl $_2$ and CaCl $_2$ required to obtain the desired free concentrations (Mg $^{2+}$ always 1 mM, Ca $^{2+}$ 0.01–10 μ M) were calculated (EqCal, Cambridge, UK) and added. In the nominal Ca $^{2+}$ -free intracellular solution, no Ca $^{2+}$ was added. Intracellular solutions were adjusted to pH 7.2 with concentrated HCl or KOH.

Calculations and Statistics. EC_{50} and $n_{\rm H}$ values for compounds and Ca^{2+} were estimated from equilibrium concentration-response relationships by fitting to the Hill equation:

$$B(C) = \frac{(B(\inf) - B(0)) \cdot C^{n_{\rm H}}}{\mathrm{E}C_{50}^{n_{\rm H}} + C^{n_{\rm H}}} + B(0),$$

where B(C) is the response (current or Rf/Ri) at the concentration C; $B(\inf)$ is the response fitted at saturating concentrations; B(0) is the response at 0 concentration; and $n_{\rm H}$ is the Hill coefficient. All results are given as the mean \pm S.E.M. Tests for significant difference were performed by one-way analysis of variance (ANOVA; and Bonferroni post hoc comparison) or Student's t test, as appropriate.

Results

Figure 1A shows the chemical structure of the novel compound GW542573X that is chemically unrelated to published SK channel modulators. The activating effects GW542573X on hSK1 channels were first identified in a VIPR membrane-potential screen. As seen in Fig. 1, B and C, GW542573X evoked concentration-dependent membrane hyperpolarizations with a mean EC₅₀ value of 4.0 \pm 0.8 μM (range, 1.6–12.5 μ M) and a Hill coefficient of 2.1 \pm 0.3 (range, 0.64-4.3, n=23). These effects were observed with three independent synthetic batches of compound. No responses were observed in untransfected CHO cells. To confirm that the hyperpolarizing effects of GW542573X observed in the VIPR experiments truly represent the activation of SK channels, patch-clamp experiments were performed. Figure 1D shows a study with a multichannel inside-out patch pulled from an hSK1-expressing HEK293 cell. Current-voltage (I-V) relationships were measured at symmetrical [K+] and with the $[Ca^{2+}]$, buffered at 0.2 or 10 μ M (solid lines), yielding 5 or 100% activation, respectively. Applying GW542573X (0.1–50 μ M, broken lines) to the inside of the patch at a $[Ca^{2+}]_i$ of 0.2 μ M resulted in a concentrationdependent increase in hSK1 current. The characteristic inward rectification of the SK1 current was maintained when the channels were activated by a combination of low Ca²⁺ and GW542573X. Figure 1E shows the GW542573X concentration-response relationship for hSK1 based on experiments similar to that shown in Fig. 1D. Data were read at -75 mV after attaining a stable current level at each concentration

and were analyzed by normalization to the maximal obtainable current measured at 10 μ M Ca²⁺. Fitting data to the Hill equation yielded an EC₅₀ value of 8.2 \pm 0.8 μ M (n = 6), a Hill coefficient of 1.4 \pm 0.1, and an efficacy of 76 \pm 5% relative to the current level at infinite [Ca²⁺]_i. For comparison, the SK channel enhancers NS309, DC-EBIO, and 1-EBIO activated hSK1 with EC₅₀ values of 0.6 \pm 0.06 (n = 4), 25 \pm 2 (n = 4), and 650 \pm 158 μ M (n = 4), respectively (data not shown). The efficacies with respect to saturating Ca²⁺ of these compounds were close to 100%.

To investigate the selectivity profile of GW542573X, whole-cell patch-clamp experiments were performed using HEK293 cells stably expressing hSK1, hSK2, hSK3, or hIK channels. The whole-cell configuration was chosen because, in our hands, the hIK current exhibited substantial rundown after patch excision. The cells were exposed to symmetrical [K⁺] at a nominal [Ca²⁺]_i of 0.3 μ M, and currents at -75 mV were plotted as a function of time. Figure 2A shows that GW542573X enhanced the hSK1 current in a concentration-dependent manner when applied to the extracellular solution at concentrations ranging from 0.1 to 10 μ M. At the beginning of the experiment, 0.1 μ M NS309 was added as a control, and the experiment was terminated by the addition of the fast SK channel blocker bicuculline methobromide (BMB, 300 μ M). Figure 2, B to D, show data from HEK293 cells

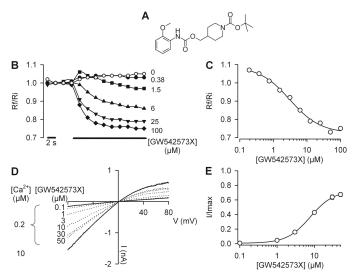


Fig. 1. GW542573X, a new positive modulator of SK channels. A, chemical structure of GW542573X. B, VIPR membrane potential measurements from hSK1-expressing cells in the absence (0) and presence of increasing concentrations (0.38-100 μM) of GW542573X. The abscissa shows time, and the solid bar shows the duration of the compound application. The ordinate shows the calculated Rf/Ri values, which reduce in size upon membrane hyperpolarization. C, representative concentration-response curve for GW542573X-evoked peak changes in membrane potential. The derived EC_{50} value from the fit to the Hill-equation was 3.1 μM with an n_H value of 0.96. D, an inside-out patch obtained from an HEK293 cell stably expressing hSK1. The patch was exposed to symmetrical [K $^{+}$] (154 mM), and 200-ms voltage ramps (-80 to +80 mV) were elicited every 5 s from a holding potential of 0 mV. The solid lines are I-V relationships obtained at a bath/intracellular [Ca2+] of 0.2 or 10 μM in the absence of GW542573X, whereas the dotted lines represent I-V relationships in the presence of GW542573X (0.2 μ M Ca²⁺, GW542573X concentration from 0.1 to 50 μM as indicated). E, concentration-response relationship for GW542573X performed in the presence of 0.2 μM Ca²⁺. Currents were normalized with respect to the effect of 10 μ M Ca²⁺, which induces maximal SK channel activity, and data points represent mean ± S.E.M. of six experiments. Data were analyzed at -75 mV. The solid line is the fit of the averaged data to the Hill equation, yielding an EC_{50} value of 7.8 $\mu\mathrm{M}$ and an n_{H} value of 1.4. The efficacy was 72%.

expressing hSK2 (B), hSK3 (C), or hIK (D, experiment terminated by 30 nM charybdotoxin, ChTX). Figure 2E summarizes the results from these experiments. The -fold increase in current levels induced by NS309 was not statistically different between the SK subtypes (hSK1, 5.99 \pm 0.56; hSK2, 6.28 ± 0.91 ; hSK3, 6.33 ± 0.51), whereas the enhancement of the hIK current level was more prominent (15.56 \pm 1.96-fold stimulation, P < 0.001, one-way ANOVA). In contrast, GW542573X was a more potent modulator of hSK1 than of hSK2, hSK3, or hIK channels at all concentrations tested. GW542573X induced no effects at the hSK2, hSK3, or hIK current level at 0.1 and 1 μ M, but hSK2 and hSK3 currents were increased at 10 μ M GW542573X to a similar degree that was higher than the response obtained for the hIK current (P < 0.001, one-way ANOVA). Although we have not done an exhaustive analysis of activity across other channel and receptor types, GW542573X produced <10% inhibition at 30 μ M versus representative voltage-gated Na⁺ (Na_v1.3), Ca²⁺ (Ca₂2.2), and K⁺ channels (K₂1.3), as well as exemplar ligand-gated ion channels (P2X2/3, nicotinic α 7). Thus, GW542573X is an SK1-selective activator.

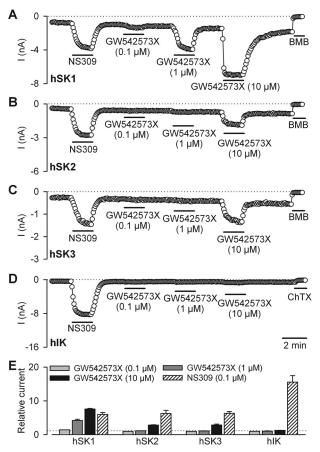


Fig. 2. GW542573X is an SK1-selective activator. Whole-cell currents recorded from HEK293 cells stably expressing hSK1 (A), hSK2 (B), hSK3 (C), or hIK (D). The pipette [Ca²+] was buffered at 0.3 μ M. The voltage protocol was as described in the legend to Fig. 1, and current recorded at -75 mV is plotted as a function of time. NS309 (0.1 μ M) or GW542573X (0.1, 1, or 10 μ M) were added to the bath solution as indicated by the bars. The experiments were terminated by the addition of 300 μ M BMB (SK experiments) or 30 nM ChTX (IK experiment). E, average currents (mean \pm S.E.M.) in the presence of GW542573X (0.1, 1, or 10 μ M) or NS309 (0.1 μ M). Currents in the presence of compound are shown relative to the SK/IK current level just before compound addition (indicated by the dashed line; n=4-5).

We next investigated whether GW542573X acted as a classic positive modulator of hSK1. The hitherto known positive modulators increase the apparent Ca2+ sensitivity of SK channels, a property reflected in a leftward shift in the Ca²⁺response curve (Pedarzani et al., 2001; Hougaard et al., 2007). The effect of GW542573X on the Ca²⁺-response relationship of hSK1 is elucidated in Fig. 3A using inside-out patches. The patch was exposed first to 0.01 µM Ca²⁺ to determine the background current level and subsequently to $10~\mu\mathrm{M~Ca^{2+}}$ to define the maximal current. After re-exposure to $0.01~\mu\mathrm{M~Ca^{2+}}$, the $[\mathrm{Ca^{2+}}]_{\mathrm{i}}$ in the intracellular solution was increased in a stepwise manner with GW542573X present as shown by the bar. Note the clear activation at 0.01 μ M Ca²⁺ and the drop in current upon compound washout at 10 μM Ca²⁺. In Fig. 3B, the effect of 10 μM GW542573X on the Ca²⁺-response relationship of hSK1 is presented. GW542573X induced a leftward shift in the apparent Ca²⁺dependence, reducing the EC_{50} value for Ca^{2+} from 410 \pm 20 (n = 9) to 240 ± 10 nM (n = 5, P < 0.01, Student's t test). In addition, GW542573X also increased the maximal degree of hSK1 channel activation, as evidenced by an increased current level at saturating Ca^{2+} concentration ($I/I_{max} > 1$). It is noteworthy that a small $(4.0 \pm 0.8\%)$ of the maximal current, n = 5) but reproducible and significant increase in current was also observed at very low Ca²⁺ concentrations. In contrast to GW542573X, NS309 only induces a leftward shift in the Ca²⁺-response relationship, reducing the EC₅₀ value for Ca^{2+} to 140 \pm 20 nM (n=3) without affecting the current level at the very low or saturating Ca²⁺ concentrations.

The complex effects of GW542573X on the Ca²⁺-response

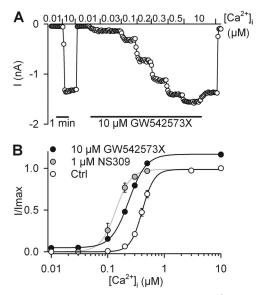


Fig. 3. GW542573X exerts complex effects on the Ca²⁺ concentration-response relationship of SK1. A, hSK1 current (inside-out patch) measured at $-75~\mathrm{mV}$ and depicted as a function of time. The [Ca²⁺] in the bath/intracellular solution was increased from 0.01 to 10 $\mu\mathrm{M}$ in a step-wise manner as indicated on the abscissa, and 10 $\mu\mathrm{M}$ GW542573X was present as shown by the bar. B, Ca²⁺-response curves, obtained from experiments similar to the one depicted in A either in the absence (Ctrl) or presence of 10 $\mu\mathrm{M}$ GW542573X or 1 $\mu\mathrm{M}$ NS309. Currents from individual patches were normalized with respect to the effect of 10 $\mu\mathrm{M}$ Ca²⁺. Data points are the mean \pm S.E.M. of nine experiments under control conditions, five experiments in the presence of GW542573X, and three in the presence of NS309. The solid lines are the fit of data to the Hill equation. Note that in the presence of GW542573X, the efficacy with respect to 10 $\mu\mathrm{M}$ Ca²⁺ without compound was 117% and that a small increase in current was induced by the compound at the very low [Ca²⁺].

relationship of hSK1 prompted a more detailed study of GW542573X at different [Ca²⁺]_i and a comparison with the hSK1 current enhancement induced by the more classic positive modulator NS309. Figure 4A shows the time course of an experiment in which 10 µM GW542573X was tested at nominally 0, 0.2, or 10 μ M intracellular Ca²⁺, and Fig. 4B shows the corresponding I-V relationships. GW542573X induced a small but distinct increase in the current level in the absence of Ca²⁺, and the I-V relationship shifted from being linear (reflecting nonspecific leak current) to exhibiting the characteristic inward rectification typical for SK channels recorded under these conditions. At the intermediate Ca²⁺ concentration, GW542573X produced a prominent stimulation of the current [from 5 (Ctrl) to 60% of full Ca²⁺ activation], and at the saturating Ca²⁺ concentration, GW542573X enhanced the current by approximately 15%. Both at the intermediate and at the high Ca2+ concentrations, the current stimulated by GW542573X maintained the typical inward rectification. Figure 4, C and D, show results from a similar experiment conducted with 1 µM NS309. Unlike GW542573X, NS309 was unable to stimulate the current at 0 or 10 μM Ca²⁺, whereas a substantial potentiation of the current was observed at $0.2 \mu M Ca^{2+}$ [from 8 (Ctrl) to 90% of full activation].

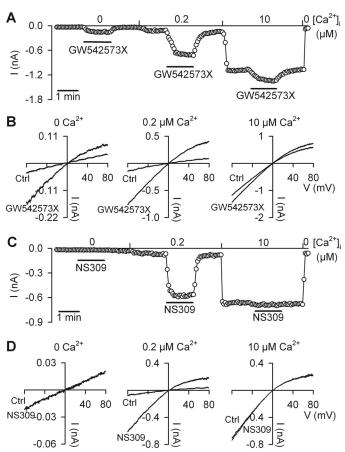


Fig. 4. GW542573X is not just a positive modulator but also an SK channel opener. A, hSK1 current at -75 mV depicted as a function of time. Current was measured from an inside-out patch obtained from an HEK293 cell stably expressing hSK1. The patch was exposed to 10 μ M GW542573X either in the complete absence of Ca²+ (0) or at 0.2 or 10 μ M intracellular Ca²+. B, I-V relationships at 0, 0.2, or 10 μ M Ca²+ in the absence (Ctrl) or presence of 10 μ M GW542573X as indicated. C and D, similar experiment as in A and B but with NS309 (1 μ M) instead of GW542573X (n=3–5).

The rather surprising observation that GW542573X is not just a positive modulator, but is, in fact, also able to activate hSK1 Ca²⁺-independently led us to investigate whether the GW542573X-activated hSK1 channel, like the Ca2+-activated, was sensitive to the SK channel blocker apamin and to positive modulation by NS309. Figure 5A shows the time course of a whole-cell experiment with 0 added Ca²⁺ and 10 mM EGTA in the pipette solution. The whole-cell configuration was chosen for this purpose because apamin is unable to block SK channels when applied from the intracellular side of the membrane, and GW542573X-induced activation at 0 Ca²⁺ only amounts to approximately 4% of the maximal current, which is considered too low for reliable assessment of additional pharmacological modulation. The letters "a" to "d" indicate the time points at which the I-V relationships shown in Fig. 5B were obtained. First, the cell was exposed to 10 μ M NS309 (a) that did not induce any increase in control current (b), suggesting that under these experimental conditions, [Ca²⁺]; was too low (well below 10 nM; Hougaard et al., 2007) to sustain NS309-mediated hSK1 activation. A subsequent application of 10 µM GW542573X (c) resulted in a robust stimulation of a current (1.6 \pm 0.3 nA, n = 11), which was sensitive to inhibition by apamin (200 nM, d). We then tested whether the GW542573X-activated channel was sensitive to positive modulation (Fig. 5, C and D). Although 10 μM NS309 (d) alone was unable to activate hSK1 channels in the absence of added Ca^{2+} , it could further enhance (by 0.7 \pm 0.2 nA, n = 6) (c) the hSK1 current activated by GW542573X (b), suggesting a synergistic or additive action of GW542573X and NS309 at the hSK1 channel level. Similar effects were observed with a classic SK channel modulator from another chemical class (a riluzole analog; data not shown).

Based on experiments with IK/SK2 chimeras, Pedarzani et al. (2001) showed that 1-EBIO mediated its stimulating effect via the C-terminal of the channels. Using hSK1-hSK3 chimeras, we have reported previously that amino acids within and just distal to the CaMBD is responsible for the SK3/2-selective channel modulation by CyPPA (Hougaard et al., 2008). A similar approach was adopted to identify the channel regions important for the selectivity of GW542573X

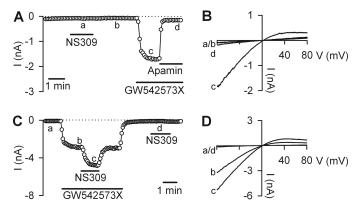


Fig. 5. The SK current activation by GW542573X in the absence of Ca²+ is sensitive to positive modulation and block. A and C, whole-cell hSK1 current at -75 mV depicted as a function of time. The [Ca²+] in the pipette solution was lower than 1 nM (no added Ca²+ and 10 mM EGTA). NS309 (10 μM), GW542573X (10 μM), and the SK channel blocker apamin (200 nM) were added to the perfusate as shown by the bars. B and D, whole-cell I-V relationships measured at the time points indicated by letters in A and C. Data are from single experiments and are representative of five to six independent observations.

for SK1 over SK3. We focused on the SK subtype selectivity rather than on the more pronounced SK1-IK selectivity. based on the assumption that chimeras from the nearly identical SK channels would most likely reflect genuine SK channel pharmacological properties. However, preliminary experiments were conducted with SK1-IK chimeras (including C-terminal switch; data not shown), yielding confirmative information to the more refined analysis outlined below. All chimeras were screened using whole-cell experiments on transiently transfected HEK293 cells. NS309 (30 nM) was applied at the beginning of the experiments to ensure that the chimeras had retained normal sensitivity to positive modulation. After washout of NS309, 200 nM GW542573X was applied, because this concentration induced a potentiation of SK1 channels of similar amplitude as 30 nM NS309, whereas having no effect on SK3. The overall topology of the chimeras and the results from this screening are presented in Fig. 6A; furthermore, an alignment showing the exact point mutations is presented in Fig. 6B.

Initially, two hSK1-hSK3 chimeras were generated, addressing the roles of the N-terminal plus the transmembrane regions versus the C-terminal harboring the CaMBD. Chimera 1 represents an hSK3 channel with an hSK1 C-terminal, and whole-cell experiments showed that insertion of the hSK1 C-terminal consistently failed to enable the modulation by GW542573X. Introduction of an SK3 C-terminal into hSK1 as for chimera 2 did not eliminate the effect of GW542573X. Chimera 1 and 2 were characterized further, and Fig. 7, A and B, show that whereas 1 µM GW542573X induced an increase in current mediated by chimera 2, 10 μ M GW542573X was required to modulate chimera 1, demonstrating that the SK1-selective modulation by GW542573X is dependent on the N-terminal and/or the transmembrane regions. Comparison of Fig. 7B with Fig. 2C shows furthermore that the response of chimera 2 to GW542573X was not different from that of wild-type SK1, whereas chimera 1 was hSK3-like in response to GW542573X.

To address specifically the role of the N-terminal domain and the intracellular loop between S2 and S3, chimera 3 was generated, and GW542573X induced an increase in current similar to that on hSK1. Furthermore, testing chimera 4, which is SK1 only in S3 to S6, elucidated that determinants of SK1-selective modulation are located in transmembrane regions surrounding the pore. Five regions within S3 to S5 were point-mutated (chimeras 5-9), walking from S3 to S5, exchanging only hSK1 amino acids being different from both hSK2 and hSK3 to their equivalent hSK3 amino acids. Chimeras 5 to 8 were all hSK1-like in their responses to GW542573X. However, chimera 9 exhibited an increase of only 1.09 ± 0.04 , indicating relative insensitivity to GW542573X. The only difference between chimera 3 and chimera 9 is the introduction of a serine residue in position 293; thus, this amino acid seems to play a critical role for the sensitivity toward GW542573X.

A leucine residue is present in both hSK2 and hSK3 at the equivalent position of the hSK1S293. This leucine residue (Leu476 in hSK3) was mutated to a serine residue in hSK3 resulting in hSK3 $_{\rm L476S}$ (chimera 10), being a GW542573X-sensitive channel. Chimera 10 was further investigated in whole-cell experiments (Fig. 7C) in a protocol similar to that used for chimera 1 and 2. It is evident that this single amino acid substitution in hSK3 fully mimicked the potency of

GW542573X on chimera 2 and hence on hSK1 (compare with Figs. 7B and 2A, respectively). Figure 7D summarizes the concentration responses of GW542573X on chimeras 1, 2, and 10 (hSK3 $_{\rm L476S}$) bracketed by the concentration responses on the hSK1 and hSK3 wild types.

Finally, Fig. 8A shows concentration-response relationships of GW542573X performed on inside-out patches with hSK3 and chimera 10 (hSK3_{L476S}). Whereas GW542573X barely stimulated the hSK3 current in concentrations up to $50 \mu M$, substantial activation of hSK3_{L476S} was seen. Fitting these data to the Hill equation yielded EC₅₀ values of 9.3 \pm 1.4 μ M (n=5), which is not significantly different from that estimated for the hSK1 (8.2 \pm 0.8 μ M, n = 6, dashed curve) and is in good agreement with the whole-cell data shown in Fig. 7. The only difference between $hSK3_{L476S}$ and hSK1 was an apparent slightly lower efficacy of GW542573X on $hSK3_{L476S}$ (50 ± 7%, n = 5) compared with hSK1 (76 ± 5%, n=6). Finally, to investigate whether Ser293 also accounted for the new MOA of GW542573X, we tested the ability of the compound to activate $hSK3_{L476S}$ in the absence of intracellular Ca²⁺. As seen from the whole-cell data in Fig. 8B, GW542573X is, in contrast to NS309, able to activate hSK3_{L476S} Ca²⁺-independently, similar to what is observed for hSK1 wild-type channels.

Thus, we conclude that the serine residue at position 293 in

hSK1 essentially accounts for the selective activation of SK1 and most likely also for the novel MOA described here for GW542573X. This shows that Ser293 positioned in S5 at a position closely opposed to the pore region constitutes an important site for modulation of the gating process of SK channels.

Discussion

In this article, we present a new chemical substance, GW542573X, characterized as a selective activator of human SK channels. The compound is structurally unrelated to previously described compounds in this pharmacological category: 1-EBIO, DC-EBIO, riluzole, SKA-31, and NS309 have limited SK subtype selectivity and higher affinity for IK channels (IK > SK1 = SK2 = SK3); and the selective SK3/ SK2 modulator CyPPA (SK3 > SK2 \gg SK1 = IK). GW542573X exhibits a novel selectivity profile being selective for SK1 over SK2/SK3 and essentially inactive on IK (SK1 > SK2 = SK3 > IK). From a selectivity point of view, GW542573X is therefore complementary to CyPPA, and these compounds (or more potent congeners) have the potential to be useful tools for investigations aimed at defining the physiological roles of the SK channel subtypes, including the poorly defined function of SK1. However, because the rat SK1

Channel	NS309 30 nM	GW542573X 200 nM		GW542573X current increase normalized to NS309			
	Fold curre	ent increase	n	0.6	8.0	1.0	1.2
SK1	1.99 <u>+</u> 0.15	1.67 <u>+</u> 0.15	5		Ι Φ Ι		
SK3	1.70 <u>+</u> 0.10	1.00	3	i e			
Chimera 1	1.93 <u>+</u> 0.13	1.09 <u>+</u> 0.04	6	Н			
Chimera 2	1.65 <u>+</u> 0.08	1.47 <u>+</u> 0.06	6		Н		
Chimera 3	1.66 <u>+</u> 0.06	1.69 <u>+</u> 0.09	3				
Chimera 4	1.78 <u>+</u> 0.31	1.69 <u>+</u> 0.32	3		 	_	l
Chimera 5	2.03 <u>+</u> 0.07	1.71 <u>+</u> 0.19	4		—		
Chimera 6	1.69 <u>+</u> 0.20	1.34 <u>+</u> 0.06	4		H		
Chimera 7	1.58 <u>+</u> 0.12	1.28 <u>+</u> 0.02	5		H		
Chimera 8	1.61 <u>+</u> 0.10	1.43 <u>+</u> 0.18	3		H •	1	
Chimera 9	1.76 <u>+</u> 0.08	1.09 <u>+</u> 0.04	6	H			
Chimera 10	1.73 <u>+</u> 0.05	1.50 <u>+</u> 0.13	6		H		

Fig. 6. Identification of the amino acid important for functional effects of GW542573X on hSK1 via effects on chimera and point-mutated channels. A. effects of NS309 and GW542573X on wild-type hSK1 and hSK3 and various channel chimeras. Regions shown in gray indicate hSK1 domains, and regions shown in black are from hSK3. -Fold increase in whole-cell current in the presence of 30 nM NS309 and in the presence 200 nM GW542573X is given for each channel as mean ± S.E.M. GW542573X current increase normalized to NS309 is calculated as (-fold increase GW542573X)/(-fold increase NS309) for each cell and is shown graphically as mean ± S.E.M. Vertical lines represents values for wt hSK1 (dashed) and wt hSK3 (solid). B, amino acid alignment of hSK1 and hSK3. S3, S4, and S5 are indicated as black lines above the alignment, and mutated amino acids are indicated by stars for chimeras 5 to 9.

В

isoform does not express in cell lines, possible species differences in the pharmacology of SK channels remain a caveat for such studies.

It is significant that the MOA of GW542573X also is unique. Previously published compounds all induce a leftward shift of the [Ca²⁺]_i-response curves without activating at low [Ca²⁺], at which the channel open state probability is 0, or augmenting at high $[Ca^{2+}]_i$, at which the channel is maximally activated (open state probability ≈ 80%; Hirschberg et al., 1998). In contrast, GW542573X causes a (minor) leftward shift in the $[Ca^{2+}]_i$ response curve for SK1 but additionally causes a partial activation in the absence of [Ca²⁺]_i and furthermore increases the current level achievable at saturating [Ca²⁺]_i. GW542573X is thus able to partially uncouple SK1 channel activation from the physiological ligand. Because the activation occurs swiftly upon application to completely closed channels, we hypothesize that the molecular mechanism relates to destabilization of a closed state rather than to stabilization of an open state. The additive effect observed at a $[Ca^{2+}]_i$ too low for NS309 to activate SK1 by itself probably represents the polar opposite effect (open channel stabilization), thereby pinpointing another fundamental difference between GW542573X and the hitherto known positive modulators.

The unusual selectivity profile and MOA is independent of the CaMBD in the C terminus, which is essential for the positive modulation by 1-EBIO (Pedarzani et al., 2001) and CyPPA (Hougaard et al., 2008). Rather, both the selectivity profile and the special MOA are tightly linked to a single amino acid, Ser293 located in S5 at the level of the inner pore vestibule just below the ion-selectivity filter. This conclusion was obtained from the hSK3/hSK1 chimera approach using

the nondiscriminating NS309 as a control for preserved "normal SK activator pharmacology" of the various chimeras, thereby gradually honing in on domains important for the selectivity of GW542573X. Eventually, the point mutation L476S in hSK3 (the position equivalent to Ser293 in hSK1) was identified as the substitution that can restore the functional potency of GW542573X to the same extent as observed for hSK1. It is important to note, however, that although this point mutation fully accounts for the selectivity pattern among the SK channels (hSK2 express Leu321), additional determinants must participate in the even more pronounced selectivity against hIK (which express the equivalent Leu215). Because the Ser293 in hSK1 is conserved in the rat isoform, it is anticipated that GW542573X will be preferentially active on rSK1 over rSK2, rSK3, and rIK as well.

These experiments show for the first time that gating properties of SK channels can be influenced by interaction with amino acids in transmembrane regions close to the inner pore vestibule and the selectivity filter. The results may be interesting, not only from a pharmacological perspective defining a new site of action, but also in relation to the ongoing debate on the nature and position of the natural gating process of SK/IK channels: Upon binding of Ca²⁺ at the N-lobes of the attached CaM, a conformational change in the CaM/CaMBD (Schumacher et al., 2001) induces an intramolecular gating cascade that is transmitted by the CaMBD-S6 linker via S6 to a structure in or close to the selectivity filter, which constitutes the physical gate. This conclusion is based on a series of articles discussing SK2 channels, in which substituted cysteine accessibility scans were performed from the proximal CaM/CaMBD-S6 linker along the S6 deep into the pore. The main conclusion reached

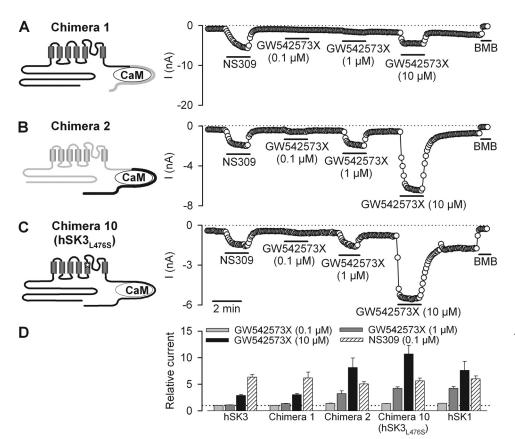


Fig. 7. The effect of GW542573X is mainly conferred by parts upstream the C terminus. A to C, whole-cell currents recorded from HEK293 cells expressing chimera 1 (A), chimera 2 (B), or chimera 10 (hSK3 $_{\rm L476S},\,$ C). Currents were recorded at -75 mV and plotted as a function of time. NS309 (0.1 μ M) or GW542573X $(0.1, 1, or 10 \mu M)$ was added to the bath solution as indicated by the bars. The $[Ca^{2+}]_i$ was 0.3 μ M, and the experiments were terminated by the addition of 300 μM BMB. D, current (mean \pm S.E.M.) in the presence of GW542573X (0.1, 1, or 10 μ M) or NS309 (0.1 μ M). For illustrative purposes, data on SK1 and SK3, previously presented in Fig. 2, are included. Currents in the presence of compound are shown relative to the SK current level just before compound addition (indicated by the dashed line) (n = 4-5).

from these experiments (Bruening-Wright et al., 2002, 2007) is that the equivalent position to the classic S6 "bundlecrossing" at the intracellular end of $K_{\rm v}$ channels does not exclude even bulky reducing agents like MTSEA+ from acting at "deep pore" positions like A384C in SK2 channels, even when the channel is closed by low [Ca²⁺];. On the other hand Ba²⁺, a K⁺ analog that specifically binds at the selectivity filter and blocks the current of K+ through SK channels only in the open state, prevents MTSEA⁺ modification at Ala384, indicating that the gate and the selectivity filter may be functionally connected. A similar experimental strategy was also adopted for the IK channel (Klein et al., 2007; Garneau et al., 2009), clearly revealing that Val275, which is equivalent to Ala384 in SK2, is accessible in the closed state for reduction by Ag⁺, a K⁺-sized ion. It is noteworthy that this study also showed that mutations at specific hydrophobic amino acids (A279G, V282G in IK, and V391G in SK2) in the pore region lead to constitutively active channels being insensitive toward [Ca²⁺]_i. This finding suggests hydrophobic interactions of S6 as being essential for the stabilization of the gate in the closed conformation and, furthermore, that transmission of the CaM-Ca²⁺ gating signal to the gate most likely involves an exposure of the hydrophobic Val282 to

Assuming that the actions of GW542573X shown in the present study are influencing the physiological "gating machinery" of SK channels (although twisting it away from normal physiological function), it is tempting to assume that the compound interacts with gating structures residing deep in the inner pore vestibule, possibly close to the selectivity filter. Therefore, these data may be an independent pharmacological support of the concept of deep-pore gating in SK channels. Although the above-mentioned attempts to localize

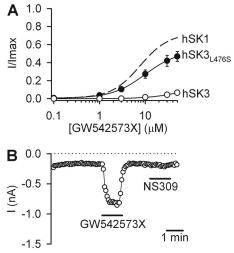


Fig. 8. Exchanging leucine in position 476 of hSK3 with the corresponding serine in hSK1 makes the channel SK1-like in the response to GW542573X. A, concentration-response relationship of GW542573X on hSK3_L476S and hSK3 channels measured on inside-out patches. Currents were normalized with respect to the effect of $10~\mu M$ ${\rm Ca^{2^+}}$, and data points represent the mean \pm S.E.M. of five experiments, for both hSK3 and hSK3_L476S channels. The solid lines are the fit of the averaged data to the Hill equation. For comparison, the data obtained on hSK1 (from Fig. 1) is shown as a dashed line. B, whole-cell current from a cell transfected with hSK3_L476S channels depicted as a function of time. The [Ca^{2^+}] in the pipette solution was lower than 1 nM (no added Ca^{2^+} and 10 mM EGTA). GW542573X (10 μM) and NS309 (10 μM) were added to the perfusate as shown by bars.

the gate by cysteine scanning has exclusively been directed at the most likely "transduction pathway" between the CaMBD and the gate—the S6—the present study pinpoints amino acids in S5 as being close enough to central gating mechanisms to influence the process. S5 amino acids in this region are not lining the pore vestibule in the potassium channel structures solved so far and are not depicted as such in SK/IK homology models. Hence it is unlikely that Ser293 acts as a "gating particle" per se, whereas it may well serve a stabilizing function for the closed channel, possibly by interhelical S5-S6 interactions. We cannot, however, distinguish between Ser273 being an important determinant of natural gating of SK1 or if it rather constitutes a favorable coordination site for GW542573X, which then by itself interferes with critical amino acid interactions important for stabilization of the closed conformation.

Conclusion and Perspective

Classic IK/SK channel openers (1-EBIO, CyPPA, and probably SKA-31 and NS309 also), act via the CaM/CaMBD at the C terminus. These compounds exclusively activate SK channels by increasing the apparent affinity for Ca²⁺ and can be formally categorized as positive allosteric modulators. In contrast, GW542573X acts via Ser293 in S5, possibly at a site near the physical gate, thereby influencing gating processes seemingly intercalated between Ca2+-binding and channel opening. GW542573X partially "bypasses" the physiological activating signal and is therefore best described as a partial agonist of hSK1 channels. In our view, GW542573X represents primarily an in vitro tool for research aimed at elucidating the gating process in SK channels. It may also delineate a route toward the synthesis of SK1-selective agents with higher potency and stability that may also be useful in the establishment of the-still obscure-physiological/pathological relevance of SK1 in the brain.

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