# Voltage- and temperature-dependent activation of TRPV3 channels is potentiated by receptor-mediated PI(4,5)P<sub>2</sub> hydrolysis

Julia F. Doerner,<sup>1</sup> Hanns Hatt,<sup>1</sup> and I. Scott Ramsey<sup>2</sup>

<sup>1</sup>Department of Cell Physiology, Ruhr University Bochum, 44801 Bochum, Germany

<sup>2</sup>Department of Physiology and Biophysics, Virginia Commonwealth University School of Medicine, Richmond, VA 23298

TRPV3 is a thermosensitive channel that is robustly expressed in skin keratinocytes and activated by innocuous thermal heating, membrane depolarization, and chemical agonists such as 2-aminoethyoxy diphenylborinate, carvacrol, and camphor. TRPV3 modulates sensory thermotransduction, hair growth, and susceptibility to dermatitis in rodents, but the molecular mechanisms responsible for controlling TRPV3 channel activity in keratinocytes remain elusive. We show here that receptor-mediated breakdown of the membrane lipid phosphatidylinositol (4,5) bisphosphate  $(PI(4,5)P_2)$  regulates the activity of both native TRPV3 channels in primary human skin keratinocytes and expressed TRPV3 in a HEK-293-derived cell line stably expressing muscarinic M<sub>1</sub>-type acetylcholine receptors. Stimulation of PI(4,5)P<sub>2</sub> hydrolysis or pharmacological inhibition of PI 4 kinase to block PI(4,5)P<sub>2</sub> synthesis potentiates TRPV3 currents by causing a negative shift in the voltage dependence of channel opening, increasing the proportion of voltage-independent current and causing thermal activation to occur at cooler temperatures. The activity of single TRPV3 channels in excised patches is potentiated by  $PI(4,5)P_2$  depletion and selectively decreased by PI(4,5)P<sub>2</sub> compared with related phosphatidylinositol phosphates. Neutralizing mutations of basic residues in the TRP domain abrogate the effect of  $PI(4,5)P_2$  on channel function, suggesting that  $PI(4,5)P_2$  directly interacts with a specific protein motif to reduce TRPV3 channel open probability.  $PI(4,5)P_2$ -dependent modulation of TRPV3 activity represents an attractive mechanism for acute regulation of keratinocyte signaling cascades that control cell proliferation and the release of autocrine and paracrine factors.

## INTRODUCTION

Transient receptor potential (TRP) channels have emerged as cellular sensors for a variety of environmental and endogenous stimuli (Clapham, 2003). Synthetic and naturally occurring organic compounds (capsaicin, menthol, and 2-aminoethyoxy diphenylborinate [2-APB]), membrane lipids such as phosphatidylinositol phosphates (PIPs) and products of lipid metabolism (diacylglycerol [DAG], eicosanoids, and arachidonate), inorganic ions (Ca<sup>2+</sup>, Mg<sup>2+</sup>, and H<sup>+</sup>), and transmembrane voltage and temperature changes modulate or directly activate a variety of TRP channels (Ramsey et al., 2006). A subset of "polymodal" TRP channels including TRPV1, TRPV3, TRPA1, TRPM4, and TRPM8 respond to multiple distinct activators, suggesting that they are likely to function as important integrators of convergent cellular stimuli. Depending on the cell- and tissue-specific contexts of their expression, polymodal TRP channels are poised to perform a variety of physiological functions in health and disease (Nilius, 2007).

TRPV3 (human TRP channel, vanilloid family, subtype 3; NCBI Protein database accession no. AF514998) is a Ca<sup>2+</sup>-permeable cation channel that is activated by moderate thermal heating  $(\geq 32^{\circ}C)$  in the range where warmth is sensed (Peier et al., 2002; Smith et al., 2002; Xu et al., 2002). Organic chemicals such as 2-APB and structurally related compounds, plant-derived compounds (camphor, carvacrol, and thymol), and unsaturated fatty acids also activate or potentiate TRPV3 activation (Peier et al., 2002; Smith et al., 2002; Xu et al., 2002, 2005, 2006; Chung et al., 2004a; Hu et al., 2006; Phelps et al., 2010). TRPV3 also displays an unusual sensitization to repeated activation that was attributed to modulation of channel activity by Ca<sup>2+</sup> and calmodulin (Xiao et al., 2008a; Phelps et al., 2010). TRPV3 protein and mRNA are highly expressed in epidermal keratinocytes, and TRPV3 currents were reported in primary rodent keratinocytes and immortalized 308 cells (Peier et al., 2002; Smith et al., 2002; Xu et al., 2002, 2006; Chung et al., 2003, 2004b; Hu et al., 2006; Cheng et al., 2010). Targeted deletion of TRPV3 in mice abolishes TRPV3 currents in keratinocytes, causes a

Correspondence to I. Scott Ramsey: isramsey@vcu.edu

J.F. Doerner's present address is Dept. of Cardiology, Children's Hospital Boston, and Dept. of Neurobiology, Harvard Medical School, Boston, MA 02115.

Abbreviations used in this paper: 2-APB, 2-aminoethyoxy diphenylborinate; CCh, carbamylcholine; DAG, diacylglycerol; OAG, 1-oleoyl-2-acetyl*sn*-glycerol; PAO, phenylarsine oxide; PI(4,5)P<sub>2</sub>, phosphatidylinositol (4,5) bisphosphate; PIP, phosphatidylinositol phosphate; PI-PLC, phosphatidylinositol-specific PLC; TRP, transient receptor potential; VIF, voltage-independent fraction; WM, wortmannin; WT, wild type.

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wavy hair/curly whisker phenotype and erythroderma, and alters thermal selection, indicating that the channel is necessary for normal hair development, epidermal barrier formation, and thermosensation (Mogrich et al., 2005; Cheng et al., 2010). In addition, the release of ATP and transforming growth factor  $\alpha$  (TGF- $\alpha$ ) elicited by TRPV3 agonists and thermal warming of primary keratinocytes is attenuated in TRPV3-null mice (Mandadi et al., 2009; Cheng et al., 2010). Keratinocyte secretion of chemical messengers and growth factors thus transmits important information to proliferating and differentiating keratinocyte as well as sensory neurons (Lee and Caterina, 2005; Lumpkin and Caterina, 2007; Mandadi et al., 2009; Cheng et al., 2010). Interestingly, spontaneous autosomal dominant mutations in TRPV3 that result in constitutive channel activity cause hairlessness and increase susceptibility to dermatitis and inflammatory skin lesions in rodents (Asakawa et al., 2006; Xiao et al., 2008b; Imura et al., 2009; Yoshioka et al., 2009). TRPV3-mediated Ca<sup>2+</sup> influx and depolarization may therefore represent an important link in signaling cascades that control cellular activation, the release of autocrine and paracrine factors, hair morphogenesis, and cell proliferation in keratinocytes.

Studies performed using heterologous expression systems showed that TRPV3 currents are also strongly potentiated by stimulation of G<sub>q/11</sub>-coupled receptors in a PLCβ-dependent fashion (Xu et al., 2006). Activation of PLC-coupled growth factor receptors also enhances TRPV3-dependent Ca<sup>2+</sup> influx (Cheng et al., 2010). The precise mechanism of TRPV3 potentiation by PLCcoupled receptors is unknown, but the response is inhibited by U73122, suggesting that stimulation of PLC activity may relieve a tonic inhibitory influence on channel opening that limits the amount of TRPV3 current under resting conditions (Xu et al., 2006; Cheng et al., 2010). PLCB mediates hydrolysis of plasma membrane phosphatidylinositol (4,5) bisphosphate ( $PI(4,5)P_2$ ), which is known to modulate the activity of a plethora of ion channels and transporters in the plasma membrane, including several members of the TRP channel family (Gamper and Shapiro, 2007; Rosenhouse-Dantsker and Logothetis, 2007; Nilius et al., 2008; Suh and Hille 2008; Rohács, 2009). We therefore hypothesized that TRPV3 channel activity might be negatively regulated by resting levels of  $PI(4,5)P_2$ , such that  $G_{q/11}$ -coupled receptor activation would result in an increase in TRPV3 conductance.

Elucidating the mechanisms by which PIPs regulate the function of ion channels and transporters is an area of active investigation, but the details of protein–PIP interactions are not well understood. Putative PI(4,5)P<sub>2</sub>binding motifs identified in various integral membrane proteins commonly contain several basic amino acid residues that are predicted to form electrostatic interactions with negatively charged phosphates on the inositol ring, but lipid contacts with noncharged residues are likely to contribute to PIP specificity; to date, no consensus  $PI(4,5)P_2$ -binding sequence has emerged (Rosenhouse-Dantsker and Logothetis, 2007; Suh and Hille, 2008). In TRPV1, TRPV5, TRPM5, and TRPM8, modulation of channel activity by  $PI(4,5)P_2$  is sensitive to mutations of residues located in the TRP domain, an  $\sim$ 25-residue motif that is conserved in most TRP channel subfamilies (Nilius et al., 2008; Rohács, 2009). Within the TRP domain is a six-amino acid TRP box motif that typically contains at least two basic residues interspersed with hydrophobic amino acids and is predicted to form an amphipathic helix (Montell, 2001; Ramsey et al., 2006; Brauchi et al., 2007). A nearby poly basic motif, which is found C-terminal to the TRP domain, has also been implicated in  $PI(4,5)P_2$  binding to TRPV1 (Ufret-Vincenty et al., 2011). The functional consequences of  $PI(4,5)P_2$  binding range from inhibition (Estacion et al., 2001; Otsuguro et al., 2008) to bidirectional modulation (Chuang et al., 2001; Stein et al., 2006; Lukacs et al., 2007) to activation (Runnels et al., 2002; Liu and Liman, 2003; Liu and Qin, 2005; Rohács et al., 2005; Nilius et al., 2006; Karashima et al., 2008; Mercado et al., 2010; Ufret-Vincenty et al., 2011).

We show here that the activation of G<sub>q/11</sub>-coupled receptors alters the voltage dependence of activation and potentiates the amplitude of current carried by endogenous TRPV3 channels in primary human keratinocytes. The activation of M<sub>1</sub> receptors in HM1 cells (HM1 is a HEK-293 cell clone that stably expresses the G<sub>q/11</sub>-coupled human muscarinic acetylcholine receptor type 1 [M<sub>1</sub>]) (Peralta et al., 1988) additionally sensitizes recombinant TRPV3 channels to activation by thermal heating and increases the potency of the channel agonist 2-APB. Pharmacological depletion of membrane  $PI(4,5)P_2$  mimics the effect of G<sub>q/11</sub>-coupled receptor stimulation. In excised patches, the open probability of TRPV3 singlechannel currents is reduced by a water-soluble  $PI(4,5)P_2$ analogue. Finally, we identify two basic residues in the TRP domain that are required for  $PI(4,5)P_2$ -dependent modulation of TRPV3. The majority of our data are well explained by a mechanism that involves tonic suppression of TRPV3 channel activity by  $PI(4,5)P_2$  and rapid relief by receptor-mediated activation of PLCβ.

### MATERIALS AND METHODS

### DNA constructs, cells, and transfection

cDNA encoding human TRPV3 (Xu et al., 2002) was subcloned into pIRES-EGFP (Invitrogen) as an EcoRV/NotI fragment for expression in mammalian cells. TRPV3 R696A and K705A mutagenesis was performed using overlap extension PCR with synthetic complementary oligonucleotides (Ho et al., 1989). All cDNA mutations were verified by DNA sequencing. HM1 cells (Peralta et al., 1988) were maintained in DMEM supplemented with 10% fetal bovine serum, 100 µg/ml penicillin/streptomycin (Invitrogen), and 0.5 mg/ml G-418 (A.G. Scientific, Inc.) at 37°C, 5% CO<sub>2</sub>. Cells were transfected using the calcium phosphate precipitation method

(Zufall et al., 1993). Transfected cells were trypsinized and plated onto glass coverslips and used for electrophysiological recordings 24–48 h after transfection. Transfected cells were identified by GFP fluorescence before electrophysiological recording.

Freshly collected human skin was processed as described by Jacobsen et al. (2005). Human keratinocytes were provided by the BG University Hospital Bergmannsheil (Ruhr University, Bochum, Germany) and maintained in complete keratinocyte medium (DMEM/Ham's F12 [3:1; Invitrogen], 10% FBS, 1% penicillin/streptomycin [Invitrogen], 5 µg/ml human insulin [Sigma-Aldrich], 24.3 µg/ml adenine [Sigma-Aldrich], 1 µM isoproterenol [Sigma-Aldrich], 0.8 µg/ml hydrocortisone [Sigma-Aldrich], 1.36 ng/ml triiodothyronine [Sigma-Aldrich], and 20 ng/ml human EGF [Sigma-Aldrich]) at 37°C, 5% CO<sub>2</sub>, for four passages. Keratinocytes were trypsinized and plated onto glass coverslips for electrophysiological recordings 24–48 h later.

#### Electrophysiology

The standard bath solution used for whole cell recordings contained (in mM): 139 NaCl, 5 KCl, 10 HEPES, 10 glucose, 2 MgCl<sub>2</sub>, and 1.8 BaCl<sub>2</sub>, pH 7.4 with NaOH. The pipette solution used to acquire 2-APB concentration-response curves and heat-evoked currents contained (in mM): 120 cesium methanesulfonate (CsMeSO<sub>3</sub>), 10 EGTA, 10 HEPES, 2 MgCl<sub>2</sub>, and 3.15 CaCl<sub>2</sub> (free  $[Ca^{2+}] \approx 50$  nM), pH 7.3 with CsOH. For experiments involving muscarinic and purinergic receptor activation, the pipette solution contained (in mM): 120 CsMeSO<sub>3</sub>, 10 1,2-Bis(2aminophenoxy)ethane-N,N,N',N'-tetraacetic acid (BAPTA), 10 HEPES, 2 CaCl<sub>2</sub> (free  $[Ca^{2+}] \approx 50 \text{ nM}$ ), 0.3 Na-GTP, 2 Mg-ATP, and 8 NaCl, pH 7.3 with CsOH. Ca<sup>2+</sup>-free pipette solution contained (in mM): 120 CsMeSO<sub>3</sub>, 10 BAPTA, 10 HEPES, 0.3 Na-GTP, 2 Mg-ATP, and 12 NaCl (free  $[Ca^{2+}] < 10 \text{ nM}$ ), pH 7.3 with CsOH. The development of nonspecific leak currents during thermal heating was assessed by superfusing an Na<sup>+</sup>-free bath solution (in mM: 150 NMDG, 10 HEPES, and 10 glucose, pH 7.4 with HCl) at the peak of the temperature ramp. Cells displaying measurable inward currents in the presence of NMDGCl were omitted from analysis. The pipette and bath solution used for excised insideout patch recordings was symmetrical (in mM): 145 NaCl, 5 CsCl<sub>2</sub>, 20 HEPES, and 1 EGTA, pH 7.4 with NaOH. Inside-out patches were washed by bath superfusion after excision and reagents were added by perfusion ( $\sim 1 \text{ min}$ ), followed by incubation in a still bath chamber (total time of exposure to the indicated reagents is indicated by horizontal bars in the figures); subsequent washout was accomplished by restarting the superfusion. For cell-attached recordings, NaCl in the bath solution was replaced by KCl. Free calcium concentrations were calculated using MaxChelator.

Unless otherwise indicated, recordings were performed at room temperature (21–24°C). In some experiments, bath solution was regulated using a bipolar temperature controller and an in-line heater/cooler with continuous superfusion (Warner Instruments). The temperature of the superfusate was recorded using a thermistor placed in close proximity to the cell.

Membrane voltage was controlled and currents were recorded using an amplifier (EPC7 Plus; HEKA) interfaced with a PC computer running Pulse and PulseFit software (HEKA). Whole cell current recordings were acquired at 5–10 kHz and low-pass filtered at 2 kHz. Capacitance correction and series resistance compensation (30–50%) were routinely used. Excised patch recordings were acquired at 10 kHz and low-pass filtered at 5 kHz. Currents were further digitally filtered offline for display purposes. For both wild-type (WT) TRPV3 and TRP domain mutants, currents were activated by 2-APB at 0.3 times its respective [ $EC_{50}$ ] to minimize irreversible entry of channels into the I2 conducting state and to control for changes in 2-APB potency observed in TRPV3 mutants.

#### Data analysis

Data were analyzed using Pulse and PulseFit (HEKA), OriginPro (OriginLab), IgorPro (WaveMetrics), Clampfit 9 (Axon Instruments), and Excel (Microsoft). Data are expressed as mean ± SEM unless otherwise noted. Statistical comparisons were made using Student's unpaired t test. Concentration-response relations were fitted with a Hill equation of the form  $I = I_{max} ([X]^n / EC_{50}^n + [X]^n)$ , where I is current, [X] is ligand concentration, EC<sub>50</sub> is halfmaximal current, and n is the Hill coefficient. In some cases, n was constrained during the curve fitting. The thermal coefficient (Q<sub>10</sub>) for temperature-dependent activation of TRPV3 currents was calculated from the slope of the steep phase of log(I) versus 1,000/T (1/°K) plots, determined by linear regression fits (y = A + B \* X, where B is the slope and A is the intercept). Peak tail current amplitude  $\left(I_{\text{TAIL}}\right)$  was calculated by extrapolating the time-decaying current to the instant of membrane voltage change (t = 0) using mono-exponential fits. The I<sub>TAIL</sub>-V relation for each cell was fitted to a Boltzmann function (I = I2 + (I1-I2)/(1 + I2)) $\exp((V-V_0)/dV))$ , where I is current, I1 is current minimum, I2 is current maximum, and V is membrane voltage). Where the data were not well fit to a complete Boltzmann function (i.e., in the absence of 2-APB), we empirically determined an apparent threshold for voltage-dependent activation (V<sub>THR</sub>) from the inspection of tail currents elicited by voltage steps in +20-mV increments, as described previously (Musset et al., 2008). The most negative voltage that generated I<sub>TAIL</sub>, which was unambiguously larger than the background current, was designated as V<sub>THR</sub>. Single-channel amplitudes were estimated by Gaussian fits to all-points histograms and direct inspection of single-channel records. Open probability (NP<sub>OPEN</sub>) was computed by the equation NP<sub>OPEN</sub> =  $T_{OPEN}/T_{OPEN} + T_{CLOSED}$ , where N is channel number,  $P_{OPEN}$  is open probability, T<sub>CLOSED</sub> is total closed time, and T<sub>OPEN</sub> is total open time, using Clampfit 9.2 (Axon Instruments). We selected patches with low basal channel activity for excised patch recordings.

### Chemicals

2-APB, camphor, carvacrol, probenecid, allyl isothiocyanate, menthol, carbamylcholine (CCh), wortmannin (WM), phenylarsine oxide (PAO), BAPTA, EGTA, NMDG, and capsaicin were from Sigma-Aldrich. Ruthenium red was purchased from EMD. ATP was from Roche, purified bacterial phosphatidylinositol-specific PLC (PI-PLC) was obtained from Invitrogen, and diC8-PIPs and anti-PI(4,5)P<sub>2</sub> mouse monoclonal antiserum (mAb) were from Echelon Biosciences.

### Online supplemental material

Fig. S1 shows current responses in primary human keratinocytes to various TRP channel agonists and compares the single-channel I-V relations of endogenous keratinocyte and expressed TRPV3 channels. Fig. S2 illustrates changes in the ligand, voltage, and temperature sensitivity that result from R696A and K705A mutations. Fig. S3 shows that R696A and K705A display reduced sensitivity to modulation of membrane PI(4,5)P<sub>2</sub> levels in cell-free membrane patches. The online supplemental material is available at http://www.jgp.org/cgi/content/full/jgp.200910388/DC1.

# RESULTS

# Endogenous TRPV3 currents in primary human keratinocytes are potentiated by $G_{q/11}$ -coupled purinergic receptor activation

We measured membrane currents elicited by voltage steps in primary cultures of human epidermal keratinocytes using the whole cell voltage clamp mode and a Cs<sup>+</sup>-based pipette solution that blocks most outward K<sup>+</sup> channel current but permits measurement of outward conductances mediated by nonselective cation channels. Superfusion of the nonselective TRPV1-3 channel agonist 2-APB (30 µM) typically activated small voltage-dependent currents with steady-state outward rectification that reversed near 0 mV (Fig. 1, A and B). Carvacrol and camphor, but not capsaicin, probenecid, allyl isothiocyanate, or menthol, elicited currents with a similar steady-state I-V relation (Fig. S1), indicating that the observed currents were most likely a result of activation of TRPV3 and not TRPV1, TRPV2, TRPA1, or TRPM8 channels, respectively (Hu et al., 2004; Ständer et al., 2004; Bang et al., 2007; Facer et al., 2007; Anand et al., 2008). TRPV4 channels, which are also expressed in keratinocytes, are insensitive to 2-APB (Chung et al., 2004a). Ruthenium red, a voltage-dependent blocker of expressed TRPV3 and TRPV4 (Peier et al., 2002; Smith et al., 2002; Voets et al., 2002; Xu et al., 2002; Chung et al., 2004a), eliminated only the inward component of 2-APB-activated current, as expected (Fig. S1). The slope conductance of 2-APB-activated single-channel currents measured in inside-out membrane patches excised from primary keratinocytes (184.6  $\pm$  2.1 pS) was



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close to that measured in patches from HM1 cells transfected with human TRPV3 cDNA ( $206.3 \pm 1.6$  pS; Fig. S1) under identical recording conditions. Collectively, these data indicate that the membrane currents measured under our experimental conditions in primary human keratinocytes are attributable to the expression of functional TRPV3 channels.

TRPV3 currents evoked by channel agonists such as 2-APB are potentiated by  $G_{q/11}$ -coupled receptor stimulation in HEK-293 cells (Xu et al., 2006), raising the question of whether native TRPV3 channels might be similarly modulated. We found that the amplitude of 2-APB-activated TRPV3 whole cell current in keratinocytes increased substantially (5.6  $\pm$  1.6- and 6.3  $\pm$ 1.9-fold at +80 and -80 mV, respectively) after exposure to the purinergic P2Y receptor agonist ATP (Fig. 1, A, B, and E). P2Y stimulation also altered the voltage dependence of native TRPV3 currents, such that half-maximal activation (V<sub>0.5</sub>) was shifted  $-61.7 \pm 17.8$  mV after exposure to ATP (Fig. 1, C, D, and F). In addition, the nondecaying portion of current increased substantially after ATP exposure (Fig. 1 C), and Boltzmann fits to the data clearly saturated at negative voltages, allowing us to

> Figure 1. Endogenous TRPV3 currents in primary human keratinocytes are potentiated by Gq/11-coupled purinergic receptor activation. (A) Representative currents elicited by voltage steps to +80 mV (filled circles) or -80 mV (open circles) in the presence of 30 µM 2-APB (black bars) before and after exposure to 100 µM ATP (open bar). (B) Representative steadystate I-V relations taken at time points indicated in A: a, basal (circles); b, 30 µM 2-APB (squares); c, 30 µM 2-APB plus 100 µM ATP (triangles). (C) Representative currents (bottom) evoked by steps to the indicated voltage (top) at times a, b, andc indicated in A. Dashed line indicates the zero current level. (D) ITAIL-V relation for voltage-dependent currents elicited in the presence of 2-APB before (squares) and after (triangles) exposure to ATP. Data are from the experiment shown in A, and lines represent fits to a Boltzmann function. Note that the midpoint of the Boltzmann fit (V<sub>0.5</sub>) shifts negatively and I<sub>TAIL</sub> at negative voltages (VIF) increases after ATP exposure. (E) Average change in step current at +80 mV (open bars) and -80 mV (shaded bars) after exposure to ATP (left) or vehicle control (right). (F) Average change in V<sub>0.5</sub> before and after ATP (open bar) or vehicle control (hatched bar) exposure. (G) Average VIF determined from the saturation of Boltzmann fits to I<sub>TAIL</sub> at negative voltages before and after exposure to ATP (open bar) or vehicle control (hatched bar). For E-G, data represent means  $\pm$  SEM from n = 4 cells.



calculate the change in the voltage-independent fraction (VIF) of TRPV3. Voltage-independent conducting modes have been previously reported for several TRP channels, including TRPV3 (Chung et al., 2005). VIF increased 26.7  $\pm$  5.1% after ATP exposure (Fig. 1, D and G), indicating that G<sub>q</sub>-coupled receptor activation not

Figure 2. Potentiation of recombinant expressed TRPV3 currents by G<sub>q/11</sub>-coupled muscarinic receptor activation. Whole cell currents were recorded from HM1 cells expressing WT TRPV3 under conditions that support ( $\sim 50$  nM of free  $[Ca^{2+}]_i$ ) or prevent (free [Ca<sup>2+</sup>]<sub>i</sub>, <10 nM) PLCβ activity. (A) Representative current recordings elicited by voltage steps to +100 and -100 mV (upper panel, top) in the presence of  $10 \,\mu M$ 2-APB before and after exposure to 100 µM CCh in the presence (lower panel) or absence (upper panel) of [Ca<sup>2+</sup>]. Dashed lines indicate the zero current level. (B) I<sub>TAIL</sub>-V relations for 2-APB-activated currents before (squares) and after (triangles) exposure to CCh in the presence  $(\sim 50 \text{ nM} [\text{Ca}^{2+}]_i; \text{ filled symbols}) \text{ or absence}$ (<10 nM [Ca<sup>2+</sup>]<sub>i</sub>; open symbols) of free Ca<sup>2+</sup> in the pipette solution. Lines indicate fits to a Boltzmann function. (C) Average fold change in step current at +80 mV (white bars) and -80 mV (gray bars) after exposure to CCh in the presence (open bars) or absence (hatched bars) of  $[Ca^{2+}]_i$ . The average increase in TRPV3 current was  $21.4 \pm 4.5$ -fold at +80 mV (P = 0.0025) and  $31.1 \pm 7.9$ -fold at -80 mV (P = 0.0091). (D) Average change in  $V_{0.5}$  before and after CCh in the presence (open bar) or absence (hatched bar) of  $[Ca^{2+}]_i$  (P = 0.014). (E) Average VIF determined from Boltzmann fits before and after exposure to CCh in the presence (open bars) or absence (hatched bars) of  $[Ca^{2+}]_i$  (P < 0.001,  $CCh + Ca^{2+}$  pre vs. post; P = 0.003,  $CCh + Ca^{2+}$ post vs. CCh – Ca<sup>2+</sup> post). For C–E, data represent means  $\pm$  SEM from n = 5-8 cells.

only modulates the voltage dependence of TRPV3 channels but also causes a significant fraction of the current to remain constitutively open. The activation of purinergic P2Y receptors therefore appears sufficient to increase the amplitude of TRPV3 currents in keratinocytes by causing a shift in voltage-dependent activation toward

 TABLE I

 Effects of  $M_1$  receptor stimulation and PI 4 kinase inhibition on TRPV3 channel activity

 CCh (+Ca<sup>2+</sup><sub>i</sub>)

 CCh (+Ca<sup>2+</sup><sub>i</sub>)

Treatment:			$CCh (+Ca^{2+}_{i})$	)		$CCh (-Ca^{2+})$	i)		PAO	
DNA:		WT	R696A	K705A	WT	R696A	K705A	WT	R696A	K705A
Fold change	Mean	21.4	9.7	$3.6^{\mathrm{a}}$	2.9	4.0	$1.5^{\mathrm{a}}$	19.6	$3.3^{\mathrm{a}}$	$1.5^{a}$
	SEM	1.6	3.3	0.3	0.4	0.6	0.1	5.6	1.0	0.1
$\Delta V_{0.5}$ (mV)	Mean	-117.3	$-45.8^{a}$	$-64.4^{a}$	-51.0	-35.1	-22.9	-109.9	$-30.1^{a}$	$-22.5^{a}$
	SEM	19.4	10.8	7.8	10.0	8.6	7.8	23.9	7.6	5.3
$\Delta \text{VIF}(\%)$	Mean	23.1	21.3	$8.3^{a}$	5.5	24.0	$1.3^{a}$	19.1	12.2	$2.3^{\mathrm{a}}$
	SEM	3.5	5.2	3.0	1.2	8.6	1.2	3.4	3.8	1.0
	n	8	6	7	7	4	6	8	5	8

Effects of  $M_1$  stimulation (CCh) or PI 4 kinase inhibition (PAO) on current amplitude and voltage dependence for WT TRPV3, R696A, and K705A channels expressed in HM1 cells recorded with either 50 nM [Ca<sup>2+</sup><sub>i</sub>] (+) or <10 nM [Ca<sup>2+</sup><sub>i</sub>] (-) to control PLCβ activity. Currents were recorded in the presence of 10 µM (WT), 3 µM (R696A), or 6 µM (K705A) 2-APB at +80 mV. Final drug concentrations: 100 µM CCh and 30 µM PAO. <sup>a</sup>P < 0.05 comparing WT TRPV3 and R696A or K705A, respectively.

negative membrane potentials and increasing the size of TRPV3 current that flows at voltages where channels are normally closed.

To examine the mechanism of G<sub>q/11</sub>-coupled receptor potentiation of TRPV3 currents, we used the HM1 cell (Peralta et al., 1988) heterologous overexpression system that permits expression of recombinant TRPV3 channels together with recombinant human M1 ACh receptors. In HM1 cells, the M1 agonist CCh caused a larger potentiation of TRPV3 current amplitude (21.4and 31.1-fold at +80 and -80 mV, respectively; Fig. 2, A and C, and Table I) than that seen in keratinocytes. The increased TRPV3 current amplitude and larger magnitude of G<sub>q/11</sub>-coupled receptor responses in HM1 cells facilitated our analysis of changes in TRPV3 channel properties. As in keratinocytes, G<sub>q/11</sub>-coupled receptor stimulation caused a large negative shift in  $V_{0.5}$  (-117.3 ± 19.4 mV; Fig. 2, B and D, and Table I) and an increase in VIF (23.1 ± 3.5%; Fig. 2, B and E, and Table I).

To test whether  $M_1$  modulation of TRPV3 required PLC $\beta$  activity, we omitted added Ca<sup>2+</sup> from the BAPTA-

buffered pipette solution, thereby reducing  $[Ca^{2+}]_i$ from ~50 to <10 nM (refer to Materials and methods). Lowering  $[Ca^{2+}]_i$  to a similar extent was previously shown to inhibit PLC $\beta$  activity and abrogate M<sub>1</sub>-dependent modulation of KCNQ2/3 channels expressed in HEK-293 cells (Horowitz et al., 2005). Blocking PLC $\beta$  by decreasing  $[Ca^{2+}]_i$  largely abolished the CCh-induced increase in TRPV3 current amplitude, negative shift in V<sub>0.5</sub>, and increase in VIF, but the effects of CCh were largely reversible on washout (Fig. 2 and Table I; not depicted). Activation of PLC $\beta$  by G<sub>q/11</sub>-coupled receptor is therefore sufficient to rapidly amplify TRPV3 currents in both native and heterologous cellular contexts.

# Stimulation of $\rm G_{q/11}\mbox{-}coupled$ receptors shifts TRPV3 agonist, voltage, and temperature sensitivity

In addition to modulating voltage-dependent activation of TRPV3,  $M_1$  stimulation altered the potency for TRPV3 channel activation by the chemical agonist 2-APB, as indicated by an approximately fourfold increase in 2-APB potency after CCh exposure (Fig. 3 A). To determine

> Figure 3. Stimulation of G<sub>q/11</sub>-coupled receptors shifts TRPV3 agonist, voltage, and temperature sensitivity. (A) Concentration-response relation of 2-APB-activated currents recorded at +80 mV after exposure to either CCh (filled circles) or vehicle control (open circles). Solid lines represent fits to the Hill equation (CCh, EC\_{50} = 6.7  $\pm$ 0.8  $\mu$ M; vehicle control, EC<sub>50</sub> = 28.6 ± 2.1  $\mu$ M; n = 5 each). (B) Representative I<sub>TAIL</sub>-V relation of tail currents (measured at -100 mV) elicited by steps to the indicated voltages before (open circles) and after (filled circles) exposure to CCh. Apparent V<sub>THR</sub> shifted from >+200 mV before to  $+168.6 \pm 9.6$  mV (n = 7) after CCh exposure. (C) Representative currents at +80 mV (filled circles) or -80 mV (open circles) during repeated heating (17-40°C) in the absence or presence of CCh (open bar). The bath was superfused with NMDGCl (black bar) to check for inward leakage current after heating. (D) Representative temperature-current relations for heat-activated TRPV3 currents recorded in the presence of 100 µM CCh (filled circles) or vehicle control (open circles). Thermal coefficient (Q<sub>10</sub>) values were determined from linear fits to each of two distinct phases of temperature-dependent activation. (E) Average Q10 values for temperaturedependent activation at +80 mV before and after exposure to either CCh (open bars) or vehicle control (hatched bars). (F) Apparent thermal threshold (T<sub>h</sub>) for TRPV3 activation at +80 mV before and after exposure to CCh (filled circles; P<0.001) or vehicle control (open circles). Open squares represent means ± SEM. For E and F, data represent means  $\pm$  SEM from n = 3-7 cells.



whether Gq/11-coupled receptor modulation of TRPV3 activity requires channels to be simultaneously activated by the chemical agonist 2-APB, we measured voltageand heat-activated currents before and after M1 stimulation in cells that were naive to 2-APB exposure (Fig. 3). Before CCh, TRPV3 tail currents were not readily detectable (up to +200 mV); after CCh, we observed voltagedependent activation of TRPV3 at  $\sim$ +170 mV (Fig. 3 B and Table II). M<sub>1</sub> activation caused the apparent threshold for thermal activation  $(T_h)$  to shift  $-10^{\circ}C$  (Fig. 3, C, D, and F), but the thermal coefficient  $(Q_{10})$  was changed only slightly by CCh exposure (Fig. 3, D and E), suggesting that G<sub>q/11</sub>-coupled receptor stimulation selectively affects the set point for thermal activation but not the fundamental mechanism of temperature sensing. Control recordings revealed only small and statistically insignificant changes in T<sub>h</sub> and Q<sub>10</sub> (Fig. 3, D-F). To rule out the possibility that heating induced a nonspecific leakage current, we superfused cells with NMDG at the peak of heat activation (Fig. 3 C). As expected, inward TRPV3 current was blocked, whereas outward currents were either unaffected or slightly potentiated (Xu et al., 2002; Chung et al., 2005). The sensitivity of TRPV3 channels to voltage, heating, and the chemical agonist 2-APB is therefore subject to modulation by receptor-catalyzed activation of PLCB and increases the likelihood that channels will conduct current at typical skin temperature and resting membrane potential.

## PI(4,5)P2 depletion potentiates 2-APB-activated **TRPV3** currents

Given that TRPV3 activity is positively modulated by  $G_{\alpha/11}$ -coupled receptors and related TRP channels are regulated by  $PI(4,5)P_2$ , we wondered whether a PLC $\beta$ catalyzed decrease in membrane  $PI(4,5)P_2$  levels could account for the changes in TRPV3 properties that we observed. We therefore measured 2-APB-activated current before and after treating cells with drugs that block either PI 3 kinase (10 nM WM) or PI 4 kinase (30 µM PAO or 10 µM WM) (Nakanishi et al., 1995; Wiedemann et al., 1996; Várnai and Balla, 1998; Suh and Hille, 2007, 2008). PAO treatment resulted in a strong potentiation of TRPV3 current (19.6  $\pm$  5.6–fold) that was similar to the effect of M<sub>1</sub> stimulation (Fig. 4, A and C, and Table I). Like CCh, PAO caused  $V_{0.5}$  to shift  $-109.9 \pm 23.9$  mV and VIF to increase  $19.1 \pm 3.4\%$  (Fig. 4, B, D, and E, and Table I). At 10  $\mu$ M, the effect of WM on V<sub>0.5</sub> and VIF  $(\Delta V_{0.5} = -98.4 \pm 18.6 \text{ mV}; \Delta \text{VIF} = 19.7 \pm 3.8\%; n = 5)$  was similar to PAO, but the average current potentiation by 10  $\mu$ M WM (9.6 ± 2.2– and 13.2 ± 2.8–fold increase at +80 and -80 mV, respectively) was less robust and failed to reach statistical significance (+80 mV, P = 0.06; -80 mV, P = 0.27; by Student's nonpaired t test). Although the data suggest that 30 µM PAO and 10 µM WM similarly affect TRPV3 activity by inhibiting PI 4 kinase, the effects of the two drugs at the concentrations tested are not identical. At a lower concentration (10 nM), the effect of WM was similar to vehicle control (Fig. 4, A-E), suggesting that PI 3 kinase activity is unlikely to participate in Gq/11-coupled receptor modulation of TRPV3 under our experimental conditions.

Combining PI 4 kinase inhibition (PAO) and CCh simulation resulted in greater current potentiation, a further negative shift in  $V_{0.5}$ , and a larger increase in VIF than was seen with either agent alone (Fig. 4). One possible explanation is that inhibiting  $PI(4,5)P_2$  synthesis while simultaneously stimulating its hydrolysis more profoundly or persistently decreases the amount of plasma membrane  $PI(4,5)P_2$ , resulting in a larger effect on TRPV3 channels. Alternatively, the combined effect of CCh and PAO may

	Effects of $M_1$ receptor stimulation on WT and mutant TRPV3 channel properties									
		WT (before CCh)	WT (after CCh)	R696A	K705A					
	Mean	28.6	$6.7^{a}$	11.9 <sup>a</sup>	$14.6^{a}$					
2-APB EC <sub>50</sub> (µM)	SEM	2.1	0.8	1.0	2.1					
	n	5	5	7	5					
V <sub>THR</sub> (mV)	Mean	>200	168.6	165.7	178.2					
	SEM		9.6	11.3	5.7					
	n	7	7	7	11					
T <sub>h</sub> (°C)	Mean	28.4	$18.3^{a}$	$22.9^{a}$	$23.5^{a}$					
	SEM	1.9	0.2	1.2	0.6					
	n	6	6	7	6					
Arrhenius slope	Mean	7.2	10.3	5.6	8.3					
	SEM	0.9	1.1	0.5	0.4					
	n	6	6	6	6					

TABLE II

Effects of G<sub>q/11</sub> activation and TRP domain mutations on 2-APB potency (EC<sub>50</sub>), apparent threshold for voltage-dependent activation (V<sub>THR</sub>), apparent threshold for temperature-dependent activation (T<sub>h</sub>), and Arrhenius slope of temperature-dependent activation for WT TRPV3, R696A, and K705A channels expressed in HM1 cells at +80 mV (except for  $V_{THR}$ ).

<sup>a</sup>P < 0.05 compared to WT TRPV3 before CCh by Student's non-paired t test.

indicate that other signaling molecules downstream of  $M_1$  activation, such as DAG, modulate TRPV3 activity. Collectively, our data suggest that depleting  $PI(4,5)P_2$  is at least partially sufficient to modulate both voltage-dependent and voltage-independent activation of TRPV3 channels.

# TRPV3 channel activity in excised patches is stimulated by $PI(4,5)P_2$ depletion

To more directly determine whether  $PI(4,5)P_2$  is necessary for controlling basal TRPV3 channel activity, we treated excised inside-out patches with agents that are known to decrease membrane  $PI(4,5)P_2$  levels (Suh and Hille, 2008). Spontaneous TRPV3 channel open probability (NP<sub>OPEN</sub>) after patch excision was typically low (refer to Materials and methods), but the addition of the  $PI(4,5)P_2$ -scavenging mAb (Zhang et al., 2003; Albert et al., 2008; Suh and Hille, 2008) caused NP<sub>OPEN</sub> to increase (NP<sub>OPEN</sub> = 0.005 ± 0.002 before mAb; NP<sub>OPEN</sub> = 0.64 ± 0.23 after mAb; Fig. 5, A–C). In contrast, control recordings showed only a negligible increase in NP<sub>OPEN</sub>

over the same 8-min incubation period (initial  $NP_{OPEN}$  =  $0.015 \pm 0.007$ ; final NP<sub>OPEN</sub> =  $0.085 \pm 0.055$ ; Fig. 5, B and C). Purified bacterial PI-PLC (Sui et al., 1998; Suh and Hille, 2008) essentially mimicked the effect of mAb, causing TRPV3 channel activity to persistently increase  $(NP_{OPEN} = 0.021 \pm 0.006 \text{ before PI-PLC}; NP_{OPEN} = 0.89 \pm$ 0.24 after PI-PLC; Fig. 5, B and C). In contrast to its effect on TRPC6 channels (Hofmann et al., 1999), the DAG analogue 1-oleoyl-2-acetyl-sn-glycerol (OAG; 100 µM) did not significantly increase TRPV3 channel activity  $(NP_{OPEN} = 0.025 \pm 0.021 \text{ before OAG; } NP_{OPEN} = 0.069 \pm$ 0.031 after OAG; Fig. 5, B and C), suggesting that DAG production alone is not sufficient to potentiate TRPV3. Our results therefore suggest that depleting  $PI(4,5)P_2$ increases the open probability of TRPV3 channels by sensitizing the channel to activation by other stimuli (i.e., thermal heating, membrane depolarization, and chemical agonists).  $PI(4,5)P_2$  thus appears necessary to maintain tonic inhibition of TRPV3 channel activity under typical resting cellular conditions.



**Figure 4.**  $PI(4,5)P_2$  depletion potentiates 2-APB–activated TRPV3 currents. HM1 cells expressing WT TRPV3 were treated for 2 min (CCh) or 5 min (PAO and WM) with the indicated drug, and currents were elicited in the presence of 10 µM 2-APB. (A) Representative currents resulting from voltage steps to +100 and -100 mV after treatment with the indicated drugs. (B) Representative I<sub>TAIL</sub>–V relations from the same cells as current traces shown in A. (C) Average fold change in step current at +80 mV (open bars) and -80 mV (gray bars) elicited by treatment with the indicated drugs or vehicle control. (D and E) Average changes in  $V_{0.5}$  (D) and VIF (E) under the indicated conditions. In D and E, bar shading represents CCh alone (open), PAO (diagonally hatched), CCh plus PAO (gray), WM (crosshatched), or vehicle control (black). Final drug concentrations in A–E: CCh, 100 µM; PAO, 30 µM; WM, 10 nM. For C–E, data represent means ± SEM from n = 5-8 cells. See Table I for a summary of the data.

To determine whether  $PI(4,5)P_2$  was sufficient to decrease TRPV3 channel NP<sub>OPEN</sub>, we applied a synthetic water-soluble analogue, diC8-PI(4,5)P2 (dioctanoyl  $PI(4,5)P_2$  or D-myo-phosphatidylinositol 4,5-bisphosphate C8) to the intracellular side of excised inside-out patches that had been previously treated with mAb or PI-PLC. Between 3 and 60 µM, diC8-PI(4,5)P<sub>2</sub> decreased PI-PLC-stimulated TRPV3 channel activity in a concentration-dependent fashion that was well fit to the Hill equation (IC<sub>50</sub> =  $10.0 \pm 4.5 \mu$ M; n = 5 different patches; Fig. 6, A and D). Similar results were seen when TRPV3 channels were activated by  $PI(4,5)P_2$  depletion with mAb (Fig. 6 C). TRPV3 activity rapidly returned after diC8-PI(4,5)P2 washout in four of six patches, and the addition of either vehicle control or other diC8-phosphatidylinositol polyphosphates ( $PI(3,4,5)P_3$  or  $PI(3,5)P_2$ ) or monophosphates (PI(3)P, PI(4)P, or PI(5)P) at 30 µM failed to significantly decrease NP<sub>OPEN</sub> (Fig. 6, B and E).

Although low concentrations (0.1–1  $\mu$ M) of diC8-PI(4,5)P<sub>2</sub> sometimes appeared to stimulate TRPV3

activity relative to NP<sub>OPEN</sub> measured after washout of PI(4,5)P<sub>2</sub>-depleting agents (Fig. 6, A and D), this effect was inconsistent, and similar time-dependent changes in NP<sub>OPEN</sub> after PI-PLC washout were also seen in some vehicle-treated patches (Fig. 6 B). A simple explanation for the data from excised patches is that TRPV3 activity is selectively inhibited by PI(4,5)P<sub>2</sub> in cell-free membrane patches.

### Effects of mutating putative $PI(4,5)P_2$ -binding residues in the TRP domain of TRPV3

To identify specific residues that are required for PLC $\beta$ and PI(4,5)P<sub>2</sub>-dependent modulation of TRPV3, we mutated two conserved basic residues in the TRP domain (R696A and K705A; Fig. 7 A) that had been previously identified as potential PI(4,5)P<sub>2</sub>-interacting residues in TRPV1 and TRPV5 (Rohács et al., 2005; Brauchi et al., 2007). Both R696A and K705A expressed robust 2-APB–activated and voltage-dependent whole cell currents, but M<sub>1</sub>-dependent channel modulation of



**Figure 5.** TRPV3 channel activity in excised patches is stimulated by  $PI(4,5)P_2$  depletion. Inside-out patches pulled from HM1 cells expressing WT TRPV3 were exposed to either anti- $PI(4,5)P_2$  mAb or PI-PLC for 5 min, or OAG for 2 min ( $V_m = +80$  mV). (A) A representative current record (left) showing the potentiating effect of mAb (1:200 dilution) on TRPV3 channel activity. Open boxes before (*a*) and after (*b*) reagent addition indicate time segments used for determination of NP<sub>OPEN</sub>. Expanded current traces (right) during the time period shown in box *a* or *b*. The level of current associated with the indicated number of open TRPV3 channels is shown on the left. (B) Selected all-points histograms showing TRPV3 channel activity in patches before and after treatment with vehicle control, mAb (1:200 dilution; data from A), 100 µM OAG, or 0.5 U/ml PI-PLC. (C) Mean calculated NP<sub>OPEN</sub> values before and after reagent addition in patches treated with vehicle control (*n* = 7), mAb (*n* = 7), OAG (*n* = 11), or PI-PLC (*n* = 29). Error bars represent SEM; \*, P < 0.05; \*\*, P < 0.001 versus control after vehicle addition by Student's nonpaired *t* test.

the mutant channels was attenuated compared with WT TRPV3 (Fig. 7 and Table I). Stimulation of  $M_1$  by CCh caused 2-APB-activated current to increase only 9.7-fold and 3.6-fold in cells expressing R696A or K705A, respectively (Fig. 7, B, D, G, and I, and Table I), which was significantly less than the 21.4-fold potentiation seen for WT TRPV3 (Fig. 2). The CCh-induced negative shift in V<sub>0.5</sub> observed for WT TRPV3 was also significantly attenuated in both R696A and K705A (Fig. 7, C, E, H, and J, and Table I). Despite their similarity, some functional differences between R696A and K705A were noted. Whereas M<sub>1</sub> stimulation increased VIF of K705A channels only 8.3% in the presence of 50 nM  $[Ca^{2+}]_i$  (compared with 23.1% for WT TRPV3), VIF increased 21.3% for R696A (Fig. 7 and Table I). Unlike WT TRPV3 and K705A, the increase in VIF for R696A appeared to be independent of  $[Ca^{2+}]_i$  (Table I). The effects of neutralizing mutations at Arg696 and Lys705 support the hypothesis that TRP domain residues are required for the robust potentiation of TRPV3 channel activity resulting from receptor-mediated  $PI(4,5)P_2$  hydrolysis or pharmacological depletion of  $PI(4,5)P_2$ .

# R696 and K705 contribute to ligand, voltage, and temperature sensitivity in TRPV3

Because  $M_1$  stimulation altered the sensitivity of WT TRPV3 to activation by 2-APB, voltage, and temperature (Fig. 3), and the R696A and K705A mutations were both much less sensitive to  $G_{q/11}$ -dependent modulation (Fig. 7 and Table I), we reasoned that both mutations might also affect 2-APB potency, voltage-dependent activation, and apparent thermal threshold. As summarized in Table II, 2-APB activated R696A and K705A currents approximately twofold more potently than it did WT TRPV3. V<sub>THR</sub> measured in the absence of CCh was also more negative for both R696A and K705A than it was for WT TRPV3 (Table II and Fig. S2). In addition, T<sub>h</sub> for R696A and K705A was shifted toward cooler temperatures compared with WT TRPV3 (Table II and Fig. S2). Interestingly, V<sub>THR</sub>,



**Figure 6.**  $PI(4,5)P_2$  decreases TRPV3 single-channel open probability. Inside-out patches excised from HM1 cells expressing WT TRPV3 (V<sub>m</sub> = +80 mV) were treated with 0.5 U/ml PI-PLC (open bar), followed by water-soluble diC8-phosphatidylinositol phosphate analogues. (A and B) Representative current records showing the [diC8-PI(4,5)P\_2]-dependent decrease in TRPV3 channel activity (A) and lack of similar effect in vehicle-treated patches (B). Filled bars in B represent serial dilutions of vehicle that mimic those used to achieve the indicated [diC8-PI(4,5)P\_2] shown in A. (C) A representative patch recording in which TRPV3 activity was stimulated by mAb and reversibly inhibited by 30 µM diC8-PI(4,5)P\_2. (D) The average diC8-PI(4,5)P\_2 concentration–response relation (data represent means ± SEM; n = 5 patches). Solid line represents a Hill fit to the data for 1–60 µM diC8-PI(4,5)P\_2 (IC<sub>50</sub> = 10.0 ± 4.5 µM). (E) Average change in NP<sub>OPEN</sub> induced by the addition of the indicated diC8-phosphatidylinositol phosphates. Data represent means ± SEM from n = 3-6 patches.





**Figure 7.** Effects of  $G_{q/11}$ -coupled receptor activation are attenuated in TRP domain mutants. (A) Sequence alignment of TRP domains from TRPV1, TRPV3, and TRPV5. Numbers indicate starting position in the amino acid sequence of human TRPV1 (NCBI Protein database accession no. NM\_080705), human TRPV3 (AF514998.1), and human TRPV5 (NM\_019841). Residues that are identical to TRPV3 are shown in gray, and basic residues previously identified as being important for PI(4,5)P<sub>2</sub>-dependent modulation of TRPV1 and TRPV5 that are conserved in TRPV3 are shown within open boxes. Asterisks denote positions of TRPV3 mutations reported here. Whole cell currents were recorded from

2-APB EC<sub>50</sub>, and T<sub>h</sub> values for R696A and K705A under resting conditions (i.e., without M<sub>1</sub> activation) were similar to those found for WT TRPV3 after G<sub>q/11</sub>-dependent receptor stimulation (Table II), suggesting that these basic residues in the TRP domain contribute to the modulation of voltage, temperature, and chemical ligand sensitivity in TRPV3.

# TRP domain–mutant channel activity is unaffected by Pl(4,5)P<sub>2</sub> depletion

To determine whether the R696A and K705A mutations affected the sensitivity of TRPV3 to  $PI(4,5)P_9$ , we measured TRPV3 single-channel currents in excised patches treated with  $PI(4,5)P_2$ -depleting agents and diC8-PI(4,5)P<sub>2</sub>. Spontaneous (basal) channel activity during the first 2 min after patch excision was sevenfold and 23-fold higher for R696A (NP<sub>OPEN</sub> = 0.13  $\pm$ 0.06; Fig. 8, A–C) and K705A (NP<sub>OPEN</sub> =  $0.45 \pm 0.10$ ; Fig. 8, D-F), respectively, than for WT TRPV3 (basal  $NP_{OPEN} = 0.02 \pm 0.004$ ; Fig. 5). The single-channel conductances of R696A (175.5  $\pm$  2.6 pS) and K705A  $(178.0 \pm 3.2 \text{ pS})$  were similar to WT TRPV3  $(180.8 \pm$ 2.4 pS). The addition of either mAb or PI-PLC failed to cause a significant increase in NP<sub>OPEN</sub> for either mutant compared with vehicle controls (Figs. 8 and S3). Likewise, R696A and K705A channel activity was unaffected by the addition of 30 µM diC8-PI(4,5)P<sub>9</sub> (Figs. 8, B and E, and S3), which exhibited a nearly maximal inhibitory effect on WT TRPV3 NPOPEN (Fig. 6). Collectively, our findings indicate that TRPV3 channel activity is negatively regulated by resting  $PI(4,5)P_2$ levels in the plasma membrane, and that basic residues in the TRP domain are essential for the effects of  $PI(4,5)P_2$  on TRPV3.

HM1 cells transfected with TRPV3 R696A (B-F) or K705A (G-K). M<sub>1</sub> was activated by bath application of 100 µM CCh in the presence  $(CCh + Ca^{2+})$  or absence  $(CCh - Ca^{2+})$  of buffered free  $[Ca^{2+}]_i$ , as described in Fig. 2. Cells were incubated with 30  $\mu$ M PAO for 5 min between the first and second 2-APB applications to inhibit PI 4 kinase activity. The 2-APB concentrations used for R696A and K705A were 3 and 6 µM, respectively. (B and G) Representative current records (bottom) elicited by steps to the indicated voltages (top) recorded under the following conditions:  $CCh + Ca^{2+}$  (black circles),  $CCh - Ca^{2+}$  (gray squares), and PAO (gray triangles). The zero current level is indicated by a solid line. (C and H) Representative ITAIL-V relations from the cells shown in B and G under the following conditions:  $CCh + Ca^{2+}$  (circles),  $CCh - Ca^{2+}$  (squares), and PAO (triangles). Lines represent fits to a Boltzmann function. (D and I) Average fold change in current amplitude at +80 mV (open bars) and -80 mV (black bars) elicited by treatment under the indicated conditions. (E and J) Average change in V<sub>0.5</sub> ( $\Delta V_{0.5} = \Delta V_{0.5 POST} - \Delta V_{0.5 PRE}$ ) measured in cells treated under the indicated conditions. (F and K) Average change in VIF (VIF<sub>POST</sub> - VIF<sub>PRE</sub>) before and after treatment under the following conditions:  $CCh + Ca^{2+}$  (open bars),  $CCh - Ca^{2+}$ (diagonally hatched bars), and PAO (crosshatched bars). For D-F and I–K, data represent means  $\pm$  SEM from n = 5-7 cells.

### DISCUSSION

Keratinocytes reside in close proximity to sensory nerve endings and leukocytes in skin, and bidirectional communication between different cell types is mediated by a variety of paracrine signaling molecules including G protein–coupled receptor agonists ATP, serotonin, histamine, prostaglandin E2, and leukotriene B4, and cytokines such as IL-1 $\alpha$ , IL-6, IL-8, IL-17, IFN- $\gamma$ , and TNF (Maurer et al., 1997; Corsini et al., 1998; Giustizieri et al., 2004; Koizumi et al., 2004; Lumpkin and Caterina, 2007; Huang et al., 2008; Zhu et al., 2009). Temperature-sensitive TRPV3 channels are highly expressed in keratinocytes, and thermal or mechanical stimulation of keratinocytes causes ATP release, which might cause autocrine stimulation of purinergic P2Y receptors (Peier et al., 2002; Smith et al., 2002; Xu et al., 2002; Koizumi et al., 2004; Inoue et al., 2007). ATP, histamine, and serotonin are known to generate a transient rise in  $[Ca^{2+}]_i$  via receptor-mediated PLC $\beta$  activation and to stimulate keratinocyte proliferation, which is required for wound healing (Maurer et al., 1997; Lee et al., 2001; Giustizieri et al., 2004; Braun et al., 2006; Greig et al., 2008). TRPV3 current is potentiated by



Figure 8. The TRP domain mutants R696A and K705A are insensitive to changes in PI(4,5)P<sub>2</sub>. Inside-out patches were excised from HM1 cells expressing TRPV3 R696A or K705A, and voltage was clamped at +80 mV. (A and D) Representative current records from inside-out patches exposed to mAb (1:200 dilution; black bars, left). Open boxes indicate the time segments used for analysis of singlechannel properties (a, before exposure to mAb; b, after exposure to mAb). Expanded current traces (right) during the time period shown in box a or b. (B and E) All-points histograms of raw data shown in A and D and similar recordings (see Fig. S3) in nontreated controls or patches treated with 0.5 U/ml PI-PLC and 30  $\mu$ M diC8-PI(4,5)P<sub>2</sub>. (C and F) Average NP<sub>OPEN</sub> in patches expressing R696A or K705A either before (open bars) or after (black bars) the addition of mAb or PI-PLC, or in nontreated control recordings. Data represent means  $\pm$  SEM from n = 3-5 patches.

 $G_{q/11}$ -coupled receptors in a heterologous overexpression system (Xu et al., 2006), suggesting that native TRPV3 channels could be subject to a similar regulatory mechanism in keratinocytes.

We show here that primary human keratinocytes express voltage- and agonist-evoked currents with biophysical and pharmacological properties that are strikingly similar to expressed TRPV3 channels. Extracellular ATP potentiated agonist-evoked TRPV3 current amplitude, shifted voltage-dependent activation toward negative potentials, and increased the proportion of voltageindependent current in both primary keratinocytes and HM1 cells, indicating that native and expressed TRPV3 channels are stimulated by similar mechanisms. The increased susceptibility to dermatitis and inflammatory skin lesions in hairless rodents bearing constitutively active TRPV3 channel mutations demonstrates that elevated TRPV3 activity is sufficient to dramatically alter skin health (Asakawa et al., 2006; Xiao et al., 2008b; Imura et al., 2009; Yoshioka et al., 2009). TRPV3 channels regulate growth factor receptor signaling and are required for TGF- $\alpha$  release (Cheng et al., 2010) and may also control the release of paracrine messengers (i.e., ATP, prostaglandin E2) from keratinocytes (Huang et al., 2008; Mandadi et al., 2009).

Consistent with previous reports describing activation mechanisms in TRPV1 and TRPM8 (Prescott and Julius, 2003; Rohács et al., 2005; Brauchi et al., 2007; Voets et al., 2007), we also found that the voltage, temperature, and agonist sensitivity of TRPV3 are coordinately modulated by G<sub>q/11</sub>-coupled receptor stimulation. Both receptor-mediated PLCB activation and pharmacological depletion of  $PI(4,5)P_2$  were sufficient to cause a large negative shift in the voltage dependence of TRPV3 activation and increase the proportion of voltage-independent current. M<sub>1</sub> stimulation also increased the potency for 2-APB to activate TRPV3 current and shifted thermal activation toward cooler temperatures without significantly affecting Q<sub>10</sub>. The inhibition of PI 4 kinase (but not PI 3 kinase) largely mimicked the M<sub>1</sub> effect, suggesting that CCh modulation of TRPV3 is primarily attributable to receptor-catalyzed hydrolysis of  $PI(4,5)P_2$ . Buffering [Ca<sup>2+</sup>]<sub>i</sub> below 10 nM was previously demonstrated to prevent PLCB-dependent modulation of KCNQ2/3 channels (Horowitz et al., 2005), and this maneuver also significantly attenuated the CCh-dependent effects on TRPV3 measured here.

DAG that is generated by PLC $\beta$  (and the PI-PLC used in excised patch recordings) may contribute to the increases in TRPV3 channel activity observed in some of our experiments. PI(4,5)P<sub>2</sub> and DAG were shown to coordinately regulate mammalian TRPC6 (Hofmann et al., 1999; Gudermann et al., 2004; Albert et al., 2008) and *Drosophila melanogaster* TRPL (Estacion et al., 2001) channel activities. However, a DAG analogue (OAG) failed to significantly elevate NP<sub>OPEN</sub> in excised patches

containing TRPV3 channels. In vitro depletion of  $PI(4,5)P_2$  therefore appears to be sufficient to increase TRPV3 channel activity, but our data do not rule out the possibility that DAG (or other signaling molecules) generated by G<sub>a</sub>-coupled receptor stimulation could also stimulate TRPV3 under different experimental conditions. Indeed, the synergistic effect of M<sub>1</sub> activation in cells pretreated with the PI 4 kinase inhibitor PAO is consistent with the possibility that another factor, possibly DAG, could increase TRPV3 current after  $PI(4,5)P_2$ becomes depleted. Additional studies to elucidate other potentially important mechanisms of TRPV3 regulation are required. However, lowering  $[PI(4,5)P_2]$  by the addition of mAb to inside-out patches was sufficient to increase TRPV3 channel activity, indicating that PLCB activity is not a prerequisite for TRPV3 sensitization. We therefore conclude that TRPV3 channel activity is subject to negative regulation by resting levels of  $PI(4,5)P_2$ in plasma membrane, such that the activation of  $G_{q/11}$ coupled receptors leads to a rapid and dramatic increase in channel activity.

The large unitary conductance and high Ca<sup>2+</sup> permeability of TRPV3 channels suggest that under physiological conditions, even a moderate elevation in TRPV3 channel opening is likely to have profound consequences for voltage- and Ca2+-dependent cellular processes. Indeed, expression of constitutively active mutant TRPV3 channels or strong activation of related TRPV1 channels by capsaicin causes cell death both in vitro and in vivo (Sugimoto et al., 1998; Jin et al., 2005; Xiao et al., 2008b). Regulation of TRPV3 channel activity by PLC-coupled receptor-catalyzed  $PI(4,5)P_2$  hydrolysis represents a plausible mechanism for maintaining strict control of basal channel activity under resting conditions while also allowing for rapid and reversible amplification of TRPV3 responses. TRPV3-mediated Ca<sup>2+</sup> influx and depolarization thus allows for temporal and spatial control of signaling cascades that may control release of factors that are known to modulate inflammation, hair growth, keratinocyte differentiation, wound healing, and thermosensation (Braun et al., 2006; Lumpkin and Caterina, 2007; Greig et al., 2008; Cheng et al., 2010).

Changes in membrane  $PI(4,5)P_2$  levels are known to modulate the activity of several TRP channels (Estacion et al., 2001; Runnels et al., 2002; Liu and Liman, 2003; Liu and Qin, 2005; Rohács et al., 2005; Nilius et al., 2006, 2008; Karashima et al., 2008; Otsuguro et al., 2008; Rohács, 2009; Mercado et al., 2010; Ufret-Vincenty et al., 2011). In a subset of TRP channels, basic residues in the TRP domain were proposed to mediate  $PI(4,5)P_2$ binding (Nilius et al., 2008; Rohács, 2009). We found that TRPV3 channels are also sensitive to  $PI(4,5)P_2$  in both whole cell and excised patch recording configurations and show here that neutralizing mutations of TRP domain residues R696 and K705 largely abrogate the effects of techniques designed to reduce  $PI(4,5)P_2$ levels in both whole cell and excised patch configurations. In addition, V<sub>THR</sub>, T<sub>h</sub>, and 2-APB potency values in R696A and K705A mimicked those seen for WT TRPV3 after M<sub>1</sub> activation, indicating that both residues are required for normal modulation of TRPV3 downstream of PLCB. Elevated WT TRPV3 channel activity resulting from  $PI(4,5)P_2$  depletion was reversed by the addition of water-soluble diC8-PI(4,5)P<sub>2</sub> with an IC<sub>50</sub> of  $\sim 10 \mu$ M.  $PI(4,5)P_2$  therefore appears to modulate TRPV3 with potency comparable to other "high affinity" channels such as Kir2.1 (Huang et al., 1998; Kobrinsky et al., 2000). Despite exhibiting higher basal activities than WT TRPV3, both R696A and K705A mutants were insensitive to diC8-PI(4,5)P<sub>2</sub>, suggesting that  $PI(4,5)P_2$ could exert its effects on TRPV3 gating by directly interacting with these residues.

Electrostatic interactions between positively charged side chains of Arg and Lys residues and negatively charged phosphates on the inositol ring appear to be essential components of discrete PI(4,5)P<sub>2</sub>-binding sites in a variety of ion channels and transporters (Rosenhouse-Dantsker and Logothetis, 2007; Suh and Hille, 2008). R696 and K705 in TRPV3 are homologous to conserved basic residues that are thought to participate in  $PI(4,5)P_2$ interactions in several other TRP channels (Rohács et al., 2005; Brauchi et al., 2007). However, most of the tools commonly used to study ion channel interactions with PIPs do not directly measure lipid binding to functional channels under voltage control, and even excised membrane patches could include cytosolic enzymes that regulate lipid turnover and modulate channel activity. Although the effect of  $PI(4,5)P_2$  on TRPV3 activity may therefore be indirect and require interactions with channel-associated PI(4,5)P<sub>2</sub>-binding proteins (Suh and Hille, 2008), a recent report showed that the TRPV1interacting and  $PI(4,5)P_2$ -binding protein Pirt is not required for channel activation by  $PI(4,5)P_2$  (Ufret-Vincenty et al., 2011). Our data are consistent with the large body of experimental evidence indicating that PIP binding to basic residues directly regulates the activity of many ion channels and transporters (Rosenhouse-Dantsker and Logothetis, 2007; Suh and Hille, 2008; Ufret-Vincenty et al., 2011).

The rapid shift in TRPV3 voltage dependence that we observed after  $M_1$  stimulation in HM1 cells or P2Y stimulation in keratinocytes is reminiscent of the effect of prolonged heating or exposure to high concentrations of channel agonists such as 2-APB (Chung et al., 2005). Previous studies concluded that both TRPV1 and TRPV3 channels undergo an apparently irreversible transition from voltage-dependent ("I1") to voltage-independent ("I2") conducting modes. Entry into I2 mode was shown to be accompanied by changes in  $T_h$ , 2-APB potency, and ion selectivity (Chung et al., 2005, 2008; Chen et al., 2009). Although transition to I2 is reported to be

irreversible, we show that the activation of  $G_{q/11}$ -coupled receptors reversibly promotes occupancy of a voltageindependent TRPV3 gating mode that shares at least some properties with I2. Although detailed mechanistic explanations for agonist-, time-, and PI(4,5)P<sub>2</sub>dependent changes in TRPV3 gating require further investigation, studies on related TRP channels may provide some insight.

The voltage dependence of TRPM8, TRPA1, and TRPM4 is also subject to modulation by channel ligands, and under some circumstances, these channels also generate voltage-independent current (Zhang et al., 2005; Nilius et al., 2006; Matta and Ahern, 2007; Chen et al., 2009). Although the voltage-independent opening of TRP channels argues against an obligatory voltage-dependent transition in activation gating (Brauchi et al., 2007; Matta and Ahern, 2007; Nilius et al., 2007), it remains possible that voltage-independent current actually reflects a shift in voltage sensitivity to extremely negative potentials that are outside of the readily measurable range, using standard voltage clamp techniques (typically -200 to +200 mV). Nonetheless, G<sub>a/11</sub>coupled receptor modulation of TRPV3 leads to an increase in the probability that channels will generate current at the resting membrane potential, effectively abrogating the effect of hyperpolarizing voltage to induce channel closure.

The discovery of heat-insensitive but agonist- and voltage-sensitive TRPV3 pore domain mutants also demonstrates that voltage-dependent gating is separable from other modes of activation, at least in TRPV3 channels (Grandl et al., 2008). Residues in S4 and the S4–S5 linker of TRPM8 were reported to contribute to voltage sensitivity in TRPM8 (Voets et al., 2007), and point mutations of G573 in mouse TRPV3 that cause constitutive voltage-independent opening are also predicted to lie in the S4-S5 linker (Xiao et al., 2008b). The high resolution structure of a mammalian voltagedependent K<sup>+</sup> channel suggests that voltage-dependent movement of S4 gating is coupled to pore opening via interactions between residues in the S4-S5 linker and C-terminal end of S6 (Long et al., 2007). The TRP domain is located at the intracellular end of S6, potentially placing it in proximity to the S4-S5 linker if basic channel architecture is similar in TRP channels and voltagegated K<sup>+</sup> channels. Docking stimulations of a full atom-refined model of TRPV1 place the  $PI(4,5)P_2$  polar head in position to interact with positive charges located in the TRP domain and the S4-S5 linker. Residues homologous to R696 and K705 lie on opposite sides of a putative PI(4,5)P<sub>2</sub>-binding pocket and may differentially affect voltage sensing (Brauchi et al., 2007). Additional residues in the proximal C terminus could also be required for  $PI(4,5)P_2$  binding in TRP channels (Ufret-Vincenty et al., 2011). The molecular determinants of voltage sensitivity and  $PI(4,5)P_2$  binding may therefore lie in close proximity, perhaps explaining the typically close coupling between voltage and lipid sensitivity (Nilius et al., 2007), and perhaps offering some insight into the differences in voltage-dependent gating between R696A and K705A observed here.

Unlike prototypical voltage-gated K<sup>+</sup> channels, in which voltage sensing is obligately coupled to channel opening, TRP channels appear to share more functional homology with Ca2+-activated K+ (BK) channels wherein voltage dependence is modulated by a ligand,  $[Ca^{2+}]_i$  (Magleby, 2001; Horrigan and Aldrich, 2002; Bezanilla, 2008). A cogent and detailed understanding of gating mechanisms in polymodal TRPs and other channels activated by multiple stimuli will likely require a combination of high resolution protein structure, computer simulation, direct measurement of ligand-channel and lipid-channel interactions, and real-time electrophysiological analysis of channel function. Determining whether bound PI(4,5)P<sub>2</sub> directly participates in the formation or function of sensors for voltage, heat, and chemical ligands, or merely influences the function of other motifs that separately comprise sensors for various activation stimuli in TRPV3 and related channels, is a goal for future study.

In summary, we showed that native TRPV3 channels in primary human keratinocytes and recombinant TRPV3 channels expressed in HEK-293 cells are strongly modulated by stimulation of G<sub>q/11</sub>-coupled receptors. PLC $\beta$ -catalyzed hydrolysis of PI(4,5)P<sub>2</sub> accounts for most of the observed effects on TRPV3 channel activity: (a) a potentiation of current amplitude that is driven by a negative shift in the voltage dependence of opening and an increase in the fraction of channels that conduct in a voltage-independent mode; and (b) a decrease in the apparent threshold for thermal activation. Each of these effects will tend to increase the likelihood that TRPV3 channels will be open under physiological conditions. Two basic residues in the TRP domain are required for Gq/11-coupled receptor modulation of TRPV3 and could participate in  $PI(4,5)P_2$  binding. We hypothesize that PI(4,5)P<sub>2</sub>-bound TRPV3 channels open only rarely and that PLC activity or PI(4,5)P<sub>2</sub> depletion provides relief of tonic inhibition.  $PI(4,5)P_2$  thus appears well suited to regulate TRPV3 channel activity in vivo and thereby control Ca<sup>2+</sup>-dependent cellular signaling in keratinocytes or other cells expressing TRPV3.

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