The Roles of Rasd1 small G proteins and leptin in the activation of TRPC4 transient receptor potential channels

Jinhong Wie^{1,#}, Byung Joo Kim^{2,#}, Jongyun Myeong¹, Kotdaji Ha¹, Seung Joo Jeong¹, Dongki Yang³, Euiyong Kim⁴, Ju-Hong Jeon¹, and Insuk So^{1,*}

¹Department of Physiology; Seoul National University College of Medicine; Seoul, Republic of Korea; ²Division of Longevity and Biofunctional Medicine; Pusan National University School of Korean Medicine; Yangsan, Republic of Korea; ³Department of Physiology; College of Medicine; Gachon University; Incheon, Republic of Korea; ⁴Department of Physiology; College of Medicine; Inje University; Busan, Republic of Korea

[#]These authors contributed equally to this work

TRPC4 is important regulators of electrical excitability in gastrointestinal myocytes, pancreatic β -cells and neurons. Much is known regarding the assembly and function of these channels including TRPC1 as a homotetramer or a heteromultimer and the roles that their interacting proteins play in controlling these events. Further, they are one of the best-studied targets of G protein-coupled receptors and growth factors in general and G $\alpha_{i/o}$ and G α_q protein coupled receptor or epidermal growth factor and leptin in particular. However, our understanding of the roles of small G proteins and leptin on TRPC4 channels is still rudimentary. We discuss potential roles for Rasd1 small G protein and leptin in channel activation in addition to their known role in cellular signaling.

Introduction

Canonical transient receptor potential 4 (TRPC4) channels are calcium-permeable, nonselective cation channels that are widely distributed in mammalian cells. TRPC4 channels are activated by $G\alpha_{q/11}$ -phospholipase C (PLC) pathway or directly activated by $G\alpha_{i/o}$ proteins.¹⁻³ Rasd1 is a protein that is encoded by the *RASD1* gene. It is also known as Dexras1 (Dexamethasoneinduced Ras-related protein 1) or Ags1 (Activators of G-protein signaling 1). It belongs to the Ras superfamily of small GTPase.⁴ Our focus here will be on the roles of small G proteins Rasd1 in the activation of TRPC4 channels. Also, the roles of glucocorticoids and leptin via TRPC4 in insulin secretin will be discussed.

TRPC4 Channels

The TRPC subfamily consists of 7 proteins designated as TRPC1 to 7, which can be further divided into 4 subgroups

© Jinhong Wie, Byung Joo Kim, Jongyun Myeong, Kotdaji Ha, Seung Joo Jeong, Dongki Yang, Euiyong Kim, Ju-Hong Jeon, and Insuk So

*Correspondence to: Insuk So; Email: insuk@snu.ac.kr

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based on their sequence homology and functional similarities: (1) TRPC1, (2) TRPC4 and TRPC5, (3) TRPC3, TRPC6 and TRPC7, and (4) TRPC2.^{5,6} The activation of TRPC channels involves $G\alpha_{q/11}$ proteins and PLC. Molecules downstream of PLC, such as IP3 and DAG, and PIP2 hydrolysis have been suggested as activators and as an activation mechanism for TRPC channels, respectively.² TRPC4 channel consist of 6 transmembrane domains, intracellularly located amino and carboxyl termini and a putative pore-forming region.⁷ Within the TRPC family, TRPC4 and TRPC5 represent a structurally distinct subgroup that is characterized by its ability to form homo- and heteromultimeric channels with each other as well as with TRPC1.⁸ TRPC4 α and TRPC4 β are the most abundantly expressed and functionally characterized. TRPC4B channel lacks a domain of 84 amino acids in the C-terminal region containing a putative binding site for calmodulin (CaM) binding and inositol 1, 4, 5-triphosphate (IP₃) receptors. The deleted regions of TRPC4 is also responsible for the channel activity by PI(4, 5)P₂.⁷ TRPC4 has the 4 short amino acid sequence motifs (M1-M4) that show typical features of TRP. M1 motif is discovered upstream of TM1 and is conserved in this form throughout the TRPC subfamily. M2 motif located within the cytosolic loop between TM4 and TM5. M3 motif is part of TM6 of all TRPCs, and TRPVs. M4 motif is a highly conserved in this form throughout the TRPC family. This channel contains a coiled-coil domain, 4 ankyrin-like repeats, and predicted multimerization domain in the N-terminus as potential protein-protein interaction motifs.

TRPC4 is expressed in diverse organs and cell types including the dendrites and soma of various types of neurons, smooth muscles, the cardiovascular system, including endothelial and cardiac cells, skeletal muscle cells, the myometrium, the kidney, and immune cells. TRPC4 channels differ extremely in their permeability and pharmacological modulation, other biophysical properties, and mode of activation depending on the cellular environment. Activation of TRPC4 channels by agonists induced Ca^{2+} entry directly or indirectly via depolarization and activation of voltage-gated Ca^{2+} channels. TRPC4 channel was reported that this channel, as well as phospholipase C β_1 and β_2 interact with the first PDZ domain of NHERF, regulatory factor of the Na⁺/H⁺ exchanger.⁹ The C-terminal PDZ (Dlg, Zo-1, PSD-95) motif of TRPC4 has been implicated in the control of channels surface expression and localization.¹⁰ The PDZ deletion

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motif (T-R-L) of TRPC4 dramatically reduced the plasma membrane levels of the channel, which led the authors to suggest that the interaction between hTRPC4 and NHERF is required for the retention and stabilization of hTRPC4 channels in the cell membrane.¹⁰ Caveolin-1 (Cav-1) is associated with dynamic protein complex consisting of TRPC4, TRPC1 and IP₃Rs. The loss of Cav-1 blunted the localization of TRPC4 and agonist-stimulated complex formation.¹¹ TRPC4 channel has been proposed to be activated by $G\alpha_{\alpha}$ /phospholipase C signaling, release of intracellular Ca²⁺stores, or vesicular translocation to the membrane. Gag-PLC signaling mediated activation of PLC increased IP₃ that binds to the IP₃ receptor located on the endoplasmic reticulum and released intracellular Ca^{2+,12} A conformational change of TRPC channel bound to IP3 receptors, which bind to the C-terminal end of the TRPC channels via the IP₃/calmodulin receptor binding domain.¹² Among TRP channels, TRPC4 channels are unique in that they activated by trivalent cations Gd³⁺ and La³⁺ independently of GPCR. Previously, we reported that $G\alpha_i$ proteins are potent activators of the TRPC4 β channel. Gai2 protein strongly increased TRPC4B current. Mutation and modeling demonstrated that the K715 and R716 regions are important for the interaction between $G\alpha_{i2}$ and TRPC4 β channel. Thus, we suggest that the regions 700-720 containing the K715 and R716 regions are important for the interaction between $G\alpha_{i2}$ and TRPC4 β channel. $G\beta\gamma$ subunits are not obligatory for the action of TRPC4β channels.

The diversity of TRPC4 channels interfere with the development of specific agonists or antagonists. Recently, ML204 was identified as a blocker of both recombinant and endogenous TRPC4 channel that lacks activity on most voltage-gated channels and other TRPs.¹⁴ Selectivity for block of TRPC4 channels was examined in fluorescent and electrophysiological experiments against closely related TRPC channels and more distantly related TRPV, TRPA and TRPM channels, and against non-TRP ion channels.¹⁴ ML204 afforded good selectivity (fold19-) against TRPC6 channels and more modest selectivity against TRPC3 and TRPC5 (fold9-) channels. Little or no block of TRPV, TRPA, TRPM or voltage-gated ion channels was observed. ML204 exhibited properties useful for a variety of in vitro investigations.¹⁴ ML204 decreased the increase in intracellular Ca²⁺ mediated by TRPC4β after activation of μ -opioid receptors.

Recently, receptor-operated or/and store-operated mechanisms contribute to the activation of TRPC4 channels is especially important for the activation mechanism of epidermal growth factor (EGF) on TRPC4. TRPC4 mediated Ca²⁺ signaling pathway evoked by EGF was identified specifically in subconfluent, proliferating clusters of human microvascular endothelial cells.¹⁵ There are direct/indirect mechanisms that EGFR activates TRPC. Direct mechanism is that 2 tyrosine residues of the C-terminus in human TRPC4, Tyr-959 and Tyr-972 were phosphorylated following EGF receptor (EGFR) stimulation of COS-7 cells.¹⁶ This phosphorylation was mediated by Src family tyrosine kinases (STKs), with Fyn appearing to be the dominant kinase. There are indirect mechanisms to modulate TRPC. EGF activates Ras/Raf/Mek/Erk pathway. Erk1/2 in turn activates TRPC4. ¹⁷ On the contrary, EGF activates cPLA2/COX-2/

Prostaglandin/AC/cAMP/PKA pathway. PKA activation in turn has a negative feedback effect on EGF-induced stimulation of the MAPK cascade at the level of Raf-1 in the ERK limb of this superfamily. PKA activated by accumulation of the cAMP inhibits the Raf-1 activation. ¹⁷ We also showed that PKA mediate inhibition of TRPC.¹⁸ Similar study regarding leptin and glucose regulation of TRPC4 was recently published.¹⁹ In this study, leptin is shown to activate TRPC4 channel and Ca²⁺ influx through TRPC4 activates AMPK that triggers translocation of ATPdependent K⁺ channel (KATP) to the plasma membrane to hyperpolarize the resting membrane potential in pancreatic β -cell. Leptin activates TRPC4 in POMC, ^{20,21,22} and kisseptinsecreting hypothalamic neuron.²³ Through POMC, leptin increases energy expenditure without any effect on food intake and decreases insulin secretion. However, the exact mechanism how leptin activate TRPC4 at the molecular level, via tyrosine phosphorylation of TRPC4 like EGF or JAK2-PI3K-PIP3-PLCgamma1 pathway, needs more experiments. It seems that each growth factor activates each TRPC channel. PDGF activates TRPC6,²⁴ BDNF TRPC3,^{25,26} and VEGF TRPC5.²⁷

The physiological role of these channels was recently established, demonstrating that TRPC4 and TRPC6 are the molecular candidates for non-selective cation channels activated by the muscarinic receptor stimulation (mIcat) in visceral smooth muscle cells. *mIcat* mediates the physiological action of acetylcholine in evoking smooth muscle contraction.²⁸ Studies in knock-out showed that TRPC4 plays and important role in vascular physiology. TRPC4^{-/-} mice have markedly reduced store- and receptor-induced endothelial Ca²⁺ entry and impaired endotheliumdependent vasorelaxation. Knockout of TRPC4 strongly reduced acetylcholine-activated non-selective cation currents in visceral smooth muscle cells that are involved in the regulation of gastric motility. Airway tracheal and bronchial smooth muscles express TRPC4 and TRPC5 that are recognized as possible candidates in the pathogenesis of asthma and chronic obstructive pulmonary disease (COPD).⁵

Recently, single-nucleotide polymorphisms (SNPs) in the TRPC4 gene are associated with human diseases. For example, the TRPC4^{1957V} mutation of SNP in the TRPC4 gene is associated with a reduced risk of myocardial infarction. This mutation leads to an increase in receptor-operated cation currents and Ca²⁺ entry. This mechanism underlying protection is an improved endothelial function, but this needs more investigation.²⁹ In other study, photoparoxysmal responses (PPR) characterized by abnormal visual sensitivity of the brain to photic stimulation which is generally related to idiopathic generalized epilepsies (IGE) are related to various SNPs in the TRPC4 gene.³⁰ This relation was not significant after corresponding corrections, this trend toward relation of TRPC4 variants and PPR/IGE is of especial interest and deserves further investigation since it could be shown that TRPC4 is involved in numerous functions related with epilepsy.⁷

Small G Protein Rasd1

Rasd1 was first discovered as a dexamethasone inducible monomeric Ras protein in AtT-20 mouse corticotroph cells in the year 1998³¹ and is expressed at high concentrations in brain

and at lower concentrations in heart, liver, kidney, skeletal muscle, pancreas, and placenta.³²⁻³⁴ Expression of Rasd1 is upregulated by steroid hormones—glucocorticoid, dexamethasone, and β -estradiol. In pancreatic β -cell, dexamethasone decreased insulin secretion while mRNA level of Rasd1 was shown to be increased.³⁵ The upregulation of Rasd1 expression by glucocorticoids was due to glucocorticoid response element (GRE) in the 3'- flanking region of the human Rasd1 gene.³⁶ Rasd1 has all of the conserved domains of the Ras superfamily required for guanine nucleotide binding, hydrolysis, and effector interaction. The full-length cDNA of Rasd1 predicts a 280-aminoacid protein with a calculated molecular mass of 31,700Da.³¹

Rasd1 selectively activates the G α_i /G α_o -protein signaling pathway and appears to act as a guanine nucleotide exchange factor for G α_i , somewhat mimicking GPCR.^{32,37} Also, it has been reported that Rasd1 may have a dual role in modulating the activation of AC2 (adenylyl cyclase 2) signaling by concurrently blocking PKC (protein kinase C) and G $_{\beta\gamma}$ activity—2 proteins that function as activators of AC2. Rasd1 acts to negatively regulate PKC δ through an isoprenylation-dependent mechanism.³⁸ As Rasd1 can also regulate G $_{\beta\gamma}$ signaling,³⁹⁻⁴¹ it may be that Rasd1 interferes with multiple inputs to AC2 that function in an additive or synergistic manner for maximal AC2 activity.

Although their physiological functions remain to be fully elucidated, Rasd1 has been implicated in the photic and nonphotic responsiveness of the circadian clock,⁴² in tonic inhibitor of atrial natriuretic factor (ANF) secretion and a modulator of hormone secretion in volume overload condition of heart by inhibiting protein regulation of ANF release.⁴³ Also Rasd1 has a function to regulate neurotransmitter mediated behaviors⁴⁴ and cancers.⁴⁵

The functional relationship between TRPC4 and small GTPase Rasd1

In previous paper, we reported that $G\alpha_i$ proteins are potent activators of mTRPC4B channels.^{1,13} Also, Rasd1 is functionally classified as a guanine nucleotide exchange factor (GEF) for $G\alpha_i$ proteins⁴⁶ and facilitates guanosine diphosphate (GDP) to GTP exchange of $G\alpha_i$ proteins and triggers it to be functionally active until intrinsic GTPase activity of $G\alpha_i$ protein turns off the signal by hydrolyzing GTP to GDP. Therefore, since the function of Rasd1 as a GEF of $G\alpha_i$ may resemble the activators of $G\alpha_i$ proteins, we expressed Rasd1 and mTRPC4B in HEK293 cells and detected the effect of Rasd1 on the activity of mTRPC4 β . Rasd1^{S33V}, Rasd1^{G81A} and Rasd1^{A178V} (a constitutively active form of Rasd1) induced large mTRPC4B current. In contrast, expressing dominant-negative form of Rasd1, Rasd1^{G31V}, did not activate mTRPC4B current. In addition, to exclude the possibility that Rasd1 activates other TRP channels such as mTRPC4 α , hTRPC5, or mTRPC6 since amino acid sequence homology shows more than 40% similarity among those channels,⁴⁷ we tested whether Rasd1 activates these channels. However, Rasd1 could not activate mTRPC4a and mTRPC6 channels and reduced hTRPC5 currents. Also, we tested various Ras proteins other than Rasd1 to check the specificity of mTRPC4 β to Rasd1. Among various Ras proteins, Rasd1 was the only Ras protein to increase mTRPC4 β . Therefore, mTRPC4 β is the unique channel

activated by Rasd1 and Rasd1 is a unique TRP channel activator among various Ras proteins in Ras family.

The sequence homology analysis demonstrate that Rasd1 is most closely related to members of the Ras superfamily of SMWG proteins, with 55% amino acid homology to Rap2B, 36% homology to Rit1, and 50% homology to the prototypical Ras protein, H-Ras.^{48,49}. Recently, several nucleotide sequences predicting proteins with high degrees of homology to human Rasd1 (Genebank No. AF172846) have been reported, including mouse Rasd1 (Genebank No. AF009246), and Rat Rasd1(Genebank No.BC099136), which share 98% homology with human Rasd1. Human Rasd1 is located at chromosome 2q42. The most closely associated homologs to Rasd1 are human Rasd2 (Genebank No. BC013419), and mouse Rasd2 (Genebank No. BC026377), which share 62% homology with Rasd1 (Fig. 1). Ras superfamily proteins such as Rasd1, Rasd2, Rit1, H-Ras and Rap2B have highly conserved GTP binding pocket ($\Sigma 1-\Sigma 4$) domains and an effector loop which participates in proteinprotein interactions with other signaling molecules and is necessary for full biological activity.^{48,50,51,52}. The presumed structure of the Rasd1 contains several characteristic Ras superfamily motifs including the phosphate/magnesium binding regions GXXXXGK(S/T) (P-loop) (Σ 1), DXXG (Σ 2), and the guanine base binding loops NKXD (Σ 3) and EXSAK (Σ 4).^{51,52,54} The motif regions G-1 to G-3 which are characteristic of GTPases are present in Rasd1.^{51,52}. The C terminus possesses a typical CAAX motif,^{51,52,54} an important biochemical feature of a Ras superfamily. CAAX motif undergoes enzymatic posttranslational farnesylayion or prenylation, which related to its subcellular localization by promoting the translocation of the Rasd1 to the plasma membrane.^{51,54}. We tested whether Rasd2 and Rit1 can activate TRPC4 without any activator like rasd1 (Fig. 2). Since Rasd1 proteins were tagged with cyan fluorescent protein (pECFP) proteins at the C-terminus, pECFP was transfected as a negative control for Rasd1. As shown in Fig. 2B, pECFP alone did not affect mTRPC4 β current (1.39 ± 0.24 pA/pF, n = 20). Rasd2^{S40V}, a constitutively active form of Rasd2, induced large mTRPC4 β current (Fig. 2; 11.1 ± 2.23 pA/pF; n = 10). A mutation of serine at position 40 to valine (S40V) made Rasd2 unable to hydrolyse GTP to GDP, hence holding it constitutively active. This constitutively active Rasd2^{S40V} induced strong morphological change.⁵⁵ Rit1Q79L, a constitutively active form of Rit1, induced large mTRPC4 β current (Fig. 2; 9.86 ± 4.82 pA/ pF; n = 7). A mutation of glutamine at position 79 to leucine (Q79L) made Rit1 unable to hydrolyse GTP to GDP, hence holding it constitutively active. This constitutively active Rit1^{Q79L} induced strong morphological change.⁵⁵.

Next, to characterize the activation mechanism of $G\alpha_i$ proteins between Rasd1 and mTRPC4 β channels, we used pertussis toxin (PTX). PTX has been widely used as a reagent to characterize the involvement of heterotrimeric G proteins in signaling. This toxin catalyzes the ADP ribosylation of α subunits of $G\alpha_i$ family, and this modification prevents the occurrence of the receptor-G protein interaction.²⁸ Because the population of functional $G\alpha_i$ is strongly reduced by PTX, we examined whether PTX antagonizes the action of Rasd1. Rasd1-activated



Figure 1. Sequence alignment of several forms of Rasd1, Rasd2, and Rit1 proteins. The following sequences were aligned by using multiple sequence alignment by Florence Corpet: human Rasd1 (Genebank No. AF172846), mouse Rasd1 (Genebank No. AF009246), rat Rasd1 (Genebank No. BC099136), human Rasd2 (Genebank No. BC013419), mouse Rasd2 (Genebank No. BC026377), rat Rasd2 (Genebank No.AF134409), human Rit1 (Genebank No.U71203), mouse Rit1 (Genebank No. U712505), human H-Ras (Genebank No. AF493916), mouse H-Ras (Genebank No. AY373386), rat H-Ras (Genebank No.M13011), human Rap2B (Genebank No. AF493915), mouse Rap2B (Genebank No. BC032168), and rat Rap2B (Genebank No.AF386786). Regions defining the GTP-binding and hydrolysis domain (Σ 1- Σ 4) are boxed and annotated with their identifying consensus sequences. Consensus abbreviations: B, basic residue; J, polar residue; O, hydrophobic residue; X, any residue. The three conserved GDP/GTP binding motifs (G1-G3), and phosphate-magnesium binding motifs (PM1-PM3) are boxed. The red colors below the sequence indicate amino acid identity, whereas the blue color indicates similarity.

mTRPC4 β currents from Rasd1^{S33V}expressing cells at -60 mV were significantly reduced when PTX was pretreated. However, when GTP_yS was added to internal solution as a positive control for the activation, PTX could not reduce GTP_yS activated mTRPC4 β currents from Rasd1^{S33V}expressing cells. From these results, we reasoned that Rasd1 requires certain population of functional $G\alpha_i$ proteins to activate mTRPC4B channels. Muscarinic acetylcholine receptor type 2 (M_2R) , for example, is strong mTRPC4B channel activator, and functional $G\alpha_i$ proteins are quintessential for its action onto mTRPC4B channels. In a sense, if Rasd1 and M2R both exist, the competitive action for functional $G\alpha_i$ protein would be expected. Therefore, we expressed both Rasd1 and M₂R in HEK293 cells and tested how the mTRPC4B channel current results. Without Rasd1, M2R strongly activated mTRPC4B channels in response to extracellular carbachol (100 µM). When Rasd1 was co-expressed, however, M2R-activated mTRPC4B current was significantly reduced which clearly demonstrates competitive action between Rasd1 and M2R. Therefore, activation signaling of Rasd1 to mTRPC4 β involves $G\alpha_i$ protein, and certain population of functional $G\alpha_i$ protein is essential for activation of mTRPC4 β by Rasd1.

There are several types of $G\alpha_i$ proteins in HEK293 cells in which the level of expression decreases following order $G\alpha_{i2} > G\alpha_{i3} > G\alpha_{i1}$.⁵⁶ To verify the subtype of $G\alpha_i$ proteins, first, we tested the effect of the subtype of wild-type (WT) $G\alpha_i$ proteins on mTRPC4 β currents. WT $G\alpha_{i2}$ or $G\alpha_{i3}$ protein itself increased the mTRPC4B currents in HEK293 cells without activator GTP γ S. However, G α_{i1} did not increase mTRPC4B currents without Rasd1^{S33V}. mTRPC4β currents were increased by activating endogenous $G\alpha_i$ proteins in HEK cells via expressed Rasd1^{S33V}. Therefore, we co-expressed mTRPC4 β channel, Rasd1^{S33 \hat{V}}, and WT G α_i protein subtypes to test further effect of expressed WT Gai protein subtypes based on the endogenous $G\alpha_i$

protein subtypes in HEK cells on mTRPC4 β currents by Rasd1^{S33V}. mTRPC4 β currents by Rasd1^{S33V} with $G\alpha_{i1}$ protein, $G\alpha_{i2}$ protein or $G\alpha_{i3}$ protein showed no difference compared with that by Rasd1^{S33V} protein alone, or expression of WT $G\alpha_{i2}$ or $G\alpha_{i3}$ proteins alone. Because Rasd1 fully activated endogenous Gai proteins in HEK cells, mTRPCβ currents may not be further increased by co-expressing $G\alpha_i$ proteins with Rasd1^{S33V}. However, Rasd1^{S33V}-activated mTRPC4β currents were increased dramatically by coexpressing $G\alpha_{i1}$ protein with Rasd1^{S33V} compared with that by expression of WT $G\alpha_{i1}$ protein alone. These result suggested that either Rasd1 activates $G\alpha_{i1}$ protein more potently or Rasd1 is expressed at the low level in HEK cells. Next, we tested the effect of dominant negative Gai proteins on Rasd1 $^{\rm S33V}\text{-}activated mTRPC4\beta cur$ rents. Dominant negative Gail G202T and $G\alpha_{i3}^{G202T}$ significantly reduced



Figure 2. Rasd2 and Rit1 activates TRPC4 channel. (**A**) All panels indicate I-V relationship of currents measured from HEK293 cells expressing mouse TRPC4 β -ECFP channel and constitutively active Rasd2^{540V} or Rit1^{Q79L} proteins. We studied TRPC4 channel activity with Cs⁺-rich extracellular solution (Cs) since TRPC4 has greater permeability to Cs⁺ than Na⁺. Rasd2^{540V} and Rit1^{Q79L} proteins activated TRPC4 channel without GTP γ S. (**B**) Summarized current density measured above. Rasd2^{540V} or Rit1^{Q79L} proteins induced TRPC4 current increase. The comparison between pECFP and Rasd2^{540V} or Rit1^{Q79L} was carried out with Student's t-test. Statistical significance was denoted by an asterisk (*) at *P*< 0.05.

Rasd1-activated mTRPC4 β currents, but $G\alpha_{i2}^{G203T}$ did not affect. Therefore, $G\alpha_{i1}$ and $G\alpha_{i3}$ proteins are crucial for activation of mTRPC4 β by Rasd1 with more potent effect on $G\alpha_{i2}$ protein.

Since Rasd1 is GEF of $G\alpha_i$ proteins, it removes GDP from α subunit and changes GDP to GTP. This exchange dislocates α subunit from betagamma subunits (G $\beta\gamma$). Although classical

pathway of heterotrimeric G protein describes α subunit mostly, the release of $G\beta\gamma$ subunit is equally important. Therefore, we tested the involvement of $G\beta\gamma$ subunits in Rasd1 activation of mTRPC4 β . When $G\beta_1$ and $G\gamma_2$ proteins or $G\beta_1$ and $G\gamma_7$ proteins were coexpressed with mTRPC4B channels, neither of them was able to activate mTRPC4β channels. Also, co-expression of β-adrenergic receptor kinase (β ARK) that scavenges free G $\beta\gamma$ proteins did not reduce Rasd1-activated mTRPC4β currents. Furthermore, $G\beta^{W99A}$ (holds G proteins as a heterotrimer⁵⁵ making G proteins to be insensitive to activating signaling) and $G\beta^{I80A}$ (keeps $G\beta\gamma$ dimers in free form⁵⁵) mutant did not induce significant difference in Rasd1 activation of mTRPC4 β . From these results, $G\beta\gamma$ signaling can be excluded from possible signaling candidates of Rasd1. In previous study, we experimented about the relation between various small G protein (Ras, Rho, Rab proteins) and TRPC4 ion channel. As results, we found that Rasd1 activated the largest TRPC4 ion channels. In addition, both Rasd2 and Rit also activated TRPC4 with a lesser degree. It is known that Rasd2 can bind G $\beta\gamma$ and G α_i and Rit can also bind G α_o .⁵⁷ Since TRPC4 can be activated by G α_i and G α_o , TRPC4 could activated by not only Rasd1 and Rasd2 acting G α_i but also



Figure 3. $G\alpha_i$ and TRPC4 β interact with Rasd1. Rasd1-CFP and $G\alpha_{i1}$ was co-transfected in HEK293 cells. Immunoprecipitation was performed with $G\alpha_{i1}$ followed by immunoblotting with GFP antibody. Total proteins of Rasd1-CFP and $G\alpha_{i1}$ were detected using GFP and $G\alpha_{i1}$ antibody. (**B**) FRET efficiency was measured in HEK293 cells Rasd1-CFP, $G\alpha_{i2}$ WT-YFP, $G\alpha_{i2}$ QL-YFP, Gq WT-YFP, Gq QL-YFP, YFP-mTRPC4 β , mTRPC4 β -CFP, YFP-mTRPC5, YFP-G β G γ , and G β YFP-G γ . Among these proteins, Rasd1 bound $G\alpha_{i2}$ WT, $G\alpha_{i2}$ QL, and Gq WT. QL: Constitutively active form Q205L to impair GTPase activity of $G\alpha_{i2}$. Rit acting on $G\alpha_o$. Taken together, Rasd1 selectively activated TRPC4 channels, and it was the only Ras protein among Ras protein family that can activate TRPC4 channels. For this to occur, it was found that certain population of functional $G\alpha_{i1}$ and $G\alpha_{i3}$ proteins are essential.



Together, the results indicate that activation of $G\alpha_i$ subunits by Rasd1 is the primary mechanism for activating TRPC4. This raised the question of whether activation of the channels requires direct interaction of Rasd1 with the $G\alpha_i$ subunits. We tested whether Rasd1 and $G\alpha_{i1}$ interacts directly (Fig. 3A). To characterize the association between Rasd1 with $G\alpha_{i1}$ in vivo, HEK cells were transfected with Rasd1-CFP and $G\alpha_{i1}$, and their association was analyzed by co-immunoprecipitation. Immunoprecipitation of $G\alpha_i$ pulled down Rasd1-CFP (Fig. 3A). We also used FRET method (Fig. 3B). Key prerequisites for our experiments of interaction of Rasd1 and G proteins were strategic attachment of CFP or YFP to G protein. To study G protein subunit interactions in cell, CFP or YFP were inserted into the aBCloop within the α -helical domain of the G α_i , a domain that has been used previously to insert various sequences into Ga-subunits.⁵⁸ Ga_{i2}(WT)-YFP was well targeted to the plasma membrane. Rasd1-CFP showed the greatest FRET efficiency with $G\alpha_{i2}$ (WT)-YFP (Fig. 3B). There was no significant FRET of Rasd1 with TRPC4, TRPC5, or G $\beta\gamma$. Whether G α_{i1} (WT)-YFP or $G\alpha_{i3}(WT)$ -YFP has more FRET efficiency than $G\alpha_{i2}(WT)$ -YFP or not needs further studies.

The roles of glucocorticoids and leptin via TRPC4 in insulin secretion

Dexamethasone (a synthetic glucocorticoid) is a widely used immunosuppressant and insulin secretion regulator. In a previous research, Kemppainen et al.³¹ suggested that dexamethasone increased the expression of Rasd1 mRNA in murine AtT-20 corticotroph cells. Therefore, we experimented whether dexamethasone increases mTRPC4 β current by activating Rasd1. mTRPC4 β current was significantly increased with dexamethasone. With these results on hands, we tested whether Rasd1 activation by dexamethasone is sufficient to generate TRPC4-like current in INS-1 cells, since both Rasd1 and TRPC4 are detectable in pancreatic β -cells.

100 nM dexamethasone was sufficient to generate cationic current in INS-1 cells, and the I-V curve resembled that of TRPC4, suggesting that dexamethasone activates TRPC4 in INS-1 cells. Also, inhibition of Rasd1 with small interfering RNA (siRNA) blunted dexamethasone-induced TRPC4 current in INS-1 cells, indicating that dexamethasone activates TRPC4 via Rasd1 in INS-1 cells. Though Rasd1 is functionally classified as a guanine nucleotide exchange factor (GEF) for G α_i proteins, Lellis-Santos et al.³⁵ suggested that dexamethasone induced a rapid and sustained increase in *Rasd1* mRNA expression in MIN6 cells (3.5- and fold15- compared with control after

Figure 4. Schematic diagram for the signaling pathway involved in dexamethasone-induced TRPC4 channel activity. Rasd1 is classified as a guanine nucleotide exchange factor (GEF) for G α_i proteins. Dexamethasone enhanced Rasd1 that promotes GDP to GTP exchange in G α_i proteins and triggers it to be functionally active until the intrinsic GTPase activity of G α_i proteins turns off the signal by hydrolyzing GTP to GDP. G α_i -GTP protein activates TRPC4 β channel. Leptin binds to its receptor (LRB) to activate Jak2, which phosphorylates IRS proteins and activates PI3 kinase. PI3 kinase activates PLC γ 1 to increase TRPC4 channel activity.

24 and 48 h of dexamethasone treatment). We also showed that dexamethasone increased protein expression level of Rasd1. Therefore, we think that dexamethasone triggers TRPC4-like cationic current in INS-1 cells at least via increasing protein expression level of Rasd1. Whether dexamethasone enhances the GEF activity of Rasd1 or not needs more experiments. The finding of TRPC4 current in pancreatic B-cell in response to dexamethasone was interesting since similar study regarding leptin and glucose regulation of TRPC4 was recently published.¹⁹ Leptin was first discovered in 1994 by Zang et al.⁵⁹ and it is the most important and widely studied hormones in the control of energy balance.⁶⁰ Leptin is a 16 kDa protein mostly secreted from adipose tissue which has a critical role regulating body weight and energy homeostasis.^{61,62} Leptin signals via the leptin receptors (LepRs), which exists as several isoforms. These receptors are expressed in the brain and in peripheral tissues.⁶³ In the hypothalamus leptin acts as an anorexigenic hormone regulating the melanocortine/neuropeptide Y system to reduce food intake, increase energy expenditure, and decrease body weight.⁶⁴ However, circulating leptin levels are increased in obese humans,⁶⁵ suggesting that obesity may be either a result or a cause of leptin resistance.⁶⁶⁻⁶⁸ Leptin is an important regulator of the immune system. Leptin-deficient and leptin receptor-deficient mice exhibit thymic atrophy and are immunodeficient.⁶⁹ Leptin is also considered a proinflammatory adipokine because of the actions it exerts in several cells of the immune system, including monocytes/macrophages, dendritic cells, neutrophils, eosinophils, basophils, natural killer cells, and lymphocytes.⁷⁰⁻⁷³ Qiu et al.⁷⁴ suggested that a Janus 2 tyrosine kinase (Jak 2)-dependent pathway via stimulation of PI3 kinase and PLCgamma1 activated TRPC channels. Therefore, we think that leptin through a Jak2-PI3 kinase–PLCgamma pathway may activate TRPC4β channels

in INS-1 cells. In this study, leptin is shown to activate TRPC4 channel and Ca^{2+} influx through TRPC4 activates AMPK that triggers translocation of ATP-dependent K⁺ channel (KATP) to the plasma membrane to hyperpolarize the resting membrane potential in pancreatic β -cell. Another study reported that dexamethasone increased the protein expression of Rasd1 in pancreatic β -cells and diminished insulin secretion.³⁵ These studies indicated that Rasd1 and TRPC4 could be common molecular candidates in controlling insulin secretion in β -cells (**Fig. 4**).

Summary

We showed 1) that Rasd1 small G protein activates TRPC4 channel via GEF for $G\alpha_i$ proteins. Two) Related small G proteins Rasd2 and Rit1 also activated TRPC4 channels. Three)

Rasd1 interacts with $G\alpha_i$ protein rather than TRPC4 or $G\beta\gamma$. Four) In pancreatic β cell line, INS-1, Rasd1 induced TRPC4-like current.

In the previous studies, we reported that $G\alpha_{i/\alpha}$ proteins are essential for the activation of TRPC4.^{1,2,13}. Related to Ga; protein, Rasd1 and homologs also activated TRPC4 by inducing GTP exchange for $G\alpha_i$ proteins.⁷⁵ Interestingly, PTX did not inhibit Rasd1-induced current in the presence of intracellular GTPyS.75 In the beginning, we thought that $G\alpha_i$ proteins are really essential for the activation of TRPC4. However, we could not obtain the clear evidence showing the mutant responding to GTP γ S but not G α_i proteins. Any mutant that did not respond to $G\alpha_{i2}$ protein also did not respond to $GTP\gamma S$.¹ On the contrary, α -spectrin domain (730-758 amino acids) deleting mutant showed the response to $G\alpha_i$ protein but not to GTPyS, suggesting that $G\alpha_i$ protein-dependent pathway might be different from more general activation pathways by GTP_γS including Ga_i protein pathways. We need TRPC4 mutants which respond to GTPyS but not to $G\alpha_i$ proteins for clear distinction between $G\alpha_i$ protein pathway and GTPyS dependent pathway or to find other activation pathways independent from $G\alpha_i$ protein pathways. It will also be important to test whether TRPC4 currents can be recorded simultaneously with intracellular calcium concentration to see the relationship of TRPC4 with calcium itself. Finally, as was done for GIRK, we should study further the binding site for intracellular calcium itself, the effect of polyamine and Mg²⁺ on TRPC4 in the insideout patch mode, the roles of PtdIns(4,5)P2 and PKC in the desensitization and the role of G proteins in channel assembly and trafficking in addition to their known role in cellular signaling.





Clinically, the effect of glucocorticoid is important in care of hospitalized diabetes mellitus (DM) patients.⁷⁶ It is known that long-term prescription of dexamethasone could lead to insulin resistance and decrease in insulin secretion.⁷⁶ If the patient has been treated with dexamethasone as an anti-inflammatory drug or immunosuppressant, diabetes mellitus as his/her underlying disease could occur as a paradoxical effect of the drug. The relationship among Rasd1, TRPC4, and insulin secretion might suggest new therapeutic agent for this particular clinical situation. However, the exact mechanism how leptin activates TRPC4 at the molecular level, via tyrosine phosphorylation of TRPC4 like EGF or JAK2-PI₃K-PIP₃-PLCgamma1 pathway, needs more experiments.

It is well known that insulin secretion from pancreatic β-cells is increased by parasympathetic nervous system. When we tried to activate TRPC4 channel with carbachol, however, carbachol did not activate TRPC4 current in INS-1 cells (Fig. 5) although muscarinic M3 receptor is expressed in pancreatic β-cells. There must be some missing links between muscarinic stimulation and insulin secretion. Miguel et al.⁷⁸ showed that acetylcholine (ACh) acts on different receptor subtypes producing both a stimulatory (M1, M3) and an inhibitory (M2, M4) action on insulin release. Gautam et al.78 also showed that mutant mice selectively lacking the M3 muscarinic acetylcholine receptor subtype in pancreatic β -cells display impaired glucose tolerance and greatly reduced insulin release. In contrast, transgenic mice selectively overexpressing M3 receptors in pancreatic β -cells show a profound increase in glucose tolerance and insulin release. Moreover, these mutant mice are resistant to diet-induced glucose

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intolerance and hyperglycemia. Gautam et al.⁷⁹ suggested that β-cell M3 muscarinic receptors play a key role in maintaining proper insulin release and glucose homeostasis. On the other hand, Miguel et al.⁸⁰ showed an important functional cooperation between the cholinergic neurotransmitter ACh and the incretin hormone GLP-1 on insulin secretion mediated through the M3 muscarinic receptor subtype. However, the insulinotropic action of ACh was associated with a paradoxical inhibitory effect on GLP-1 stimulated cAMP production, achieved through a novel PTX- and pirenzepine-sensitive M1 muscarinic receptor activated pathway. These results suggest that intracellular Ca²⁺ and cAMP are important for the insulinotropic action of ACh. Considering the role of TRPC4 in M2 and M3 muscarinic receptor signaling, there is still the possibility for TRPC4 to be involved in the insulinotropic action of ACh in native pancreatic β-cells.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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