

Original Article

Increased store-operated Ca^{2+} entry mediated by *GNB5* and *STIM1*

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ABSTRACT Recent human genetic studies have shown that *Gβ5* is related to various clinical symptoms, such as sinus bradycardia, cognitive disability, and attention deficit hyperactivity disorder. Although the calcium signaling cascade is closely associated with a heterotrimeric G-protein, the function of *Gβ5* in calcium signaling and its relevance to clinical symptoms remain unknown. In this study, we investigated the *in vitro* changes of store-operated calcium entry (SOCE) with exogenous expression of *Gβ5*. The cells expressing *Gβ5* had enhanced SOCE after depletion of calcium ion inside the endoplasmic reticulum. *Gβ5* also augmented Stim1- and Orai1-dependent SOCE. An *Orai1* loss-of-function mutant did not show inhibition of *Gβ5*-induced SOCE, and a *STIM1-ERM* truncation mutant showed no enhancement of SOCE. These results suggested a novel role of *GNB5* and Stim1, and provided insight into the regulatory mechanism of SOCE.

INTRODUCTION

Before the discovery of direct activation and interaction of *Gβγ* with the potassium channel in the heart [1], the G_α subunit was considered the only effector, and the *Gβγ* complex was considered a signal inhibitor molecule. However, after the discovery of several biological effects of *Gβγ*, many studies have attempted to understand its functions. It is currently known that the *Gβγ* complex has its own properties and functional roles in cellular signaling pathways [2,3]. *Gβγ* has specific tissue expression patterns, and specific coupling between the GPCRs [3]. For example, a study showed that *Gβ1γ1* assisted *Gat* binding to rhodopsin, but *Gβ1γ2* did not [4]. *Gα13β1γ3* complexes are coupled with the angiotensin receptor (*AT1_A*) in rat portal vein myocytes, and enhance cytosolic calcium ion transport [5]. These previous studies as well as others suggest that each component of the *Gβγ* subunit complex has its own unique role in specific tissues.

Interestingly, recent genetic studies have reported that mutations of *GNB5* (the gene encoding *Gβ5*) were associated with the

symptoms of sinus bradycardia, cognitive disability [6], and attention deficit hyperactivity disorder (ADHD) [7,8] in the human. Moreover, a mouse model that lacked *Gβ5* showed hyperactivity [9].

GNB5 related symptom related with calcium signaling. Calcium signaling is closely related with cardiac and neural function. In particular, store-operated calcium entry (SOCE) plays important roles in the heart [10-12]. SOCE allows replenishment of sarcoplasmic reticulum calcium in the cardiomyocytes; therefore, defects in SOCE lead to some cardiac diseases [12,13]. In neurons, calcium signaling is also important. Spontaneously hypertensive rats, as an animal model of ADHD [14], have a dysfunction of calcium signaling [15,16] and lower brain Ca^{2+} -ATPase activity [17].

Consequently, previous studies have implied a relationship of the *GNB5* gene and SOCE. However, there are no studies describing this relationship, and therefore we investigated the regulatory mechanism of *GNB5*-induced effects on SOCE.



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METHODS

cDNA and transfection

The genes, *GNBI* (NCBI accession number: NM_002074), *GNB5* (NCBI accession number: NM_006578), *GNG11* (NCBI accession number: NM_004126), and *GNG13* (NCBI accession number: NM_016541), were cloned into the mammalian expression vectors, pEGFP-N1 and p3XFLAG-CMV14. The mutant Orai1^{R91W} [18,19] and Δ ERM-Stim1^{D76A} [20,21] were also used. All the samples were incubated for 24 h after transfection using Lipofectamine 2,000 (Invitrogen, Carlsbad, CA, USA).

Cell culture

HEK293T cells were maintained in high-glucose Dulbecco's Modified Eagle's Medium with 10% fetal bovine serum and 1 \times antibiotic-antimycotic reagent (Gibco, Gaithersburg, MD, USA) in 95% CO₂, 5% O₂, at 37°C. Subculturing was done with trypsin/EDTA when the cells reached confluency. The cells used in the calcium imaging experiments were cultured on 24 mm \times 24 mm glass slides.

Measurement of intracellular calcium

The cells were cultured on 24 mm \times 24 mm glass slides at a density of 1.5 \times 10⁶ cells/ml for 48 h and then treated with 4 μ M Fura-2/AM and 0.05% pluronic acid F-127 for 30 min in physiological salt solution at room temperature. Fura-2/AM fluorescence was measured at an excitation wavelength of 340/380 nm, and an emission wavelength of 510 nm (expressed as the ratio=F340/F380), using an imaging system (Molecular Devices, Sunnyvale, CA, USA). The emitted fluorescence was monitored using a CCD camera (CoolSNAP, Tucson, AZ, USA) attached to an inverted microscope. Fluorescence images were obtained at 2-s intervals. All data were analyzed using MetaFluor software (Molecular

Devices, Downingtown, PA, USA). Intracellular calcium is shown as representative traces. The data represent the mean \pm S.E.M. Δ Plateau was defined as the value that difference of the Fura-2/AM ratio between the injection point of 2 mM Ca²⁺ solution and highest value.

Statistical analysis

The statistical significance of differences between groups was determined using the paired Student's *t*-test. In all cases, *p* values<0.05 were considered significant.

RESULTS

GNB5 expression increases SOCE

To investigate the role of the G β and G γ subunits in SOCE, we overexpressed *GNBI*, *GNB5*, *GNG11*, and *GNG12* in HEK293T cells. After the expression of each protein subunit was confirmed, we measured the amount of SOCE (Fig. 1A). To maximize SOCE, we used 25 μ M cyclopiazonic acid (CPA), and induced calcium entry using 2 mM Ca²⁺ physiological salt solution. SOCE was increased after exogenous overexpression of the *GNB5* gene (Fig. 1B), suggesting that G β 5 protein affects the SOCE in HEK293T cells.

GNB5 enhances Stim1-Orai1 dependent SOCE

Because Stim1-Orai1 complexes are known as ubiquitous store-operated Ca²⁺ channels [22], we investigate SOCE after co-expression of Stim1, Orai1, and G β 5. SOCE was significantly increased when all the components were expressed (Fig. 2B). With G β 5 expression, the SOCE was approximately 2.5- fold higher than that with Stim1 or Orai1 expression. SOCE was also enhanced in Stim1-G β 5 expressing cells and in Orai1-G β 5 expressing cells.

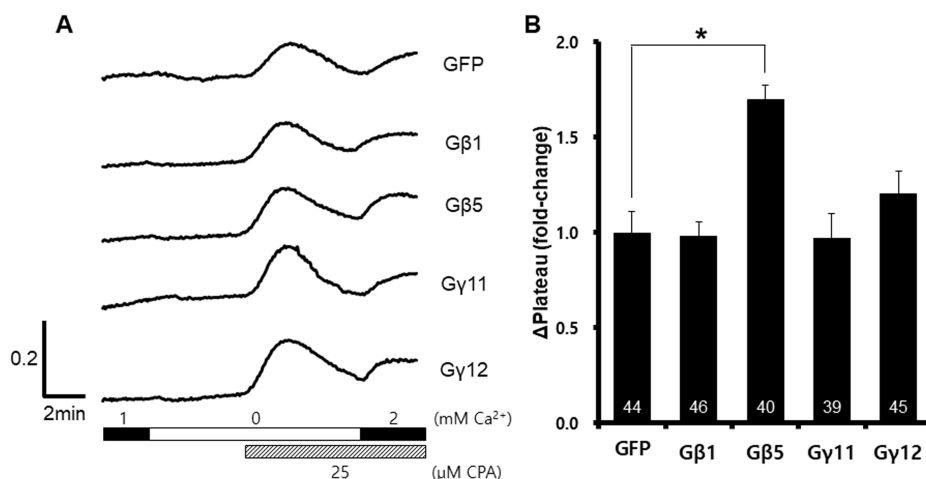


Fig. 1. Enhanced store-operated Ca²⁺ entry induced by overexpression of *GNB5*. After depletion of ER Ca²⁺ stores using cyclopiazonic acid and EGTA, Ca²⁺ entry was measured by Fura2-AM fluorescent signal ratios. A representative trace (A) summarizes the data (B), and the number inside the column is the total cell number from multiple independent experiments (n=3-4). Data were expressed as the mean \pm SEM. **p*<0.05 compared with GFP.

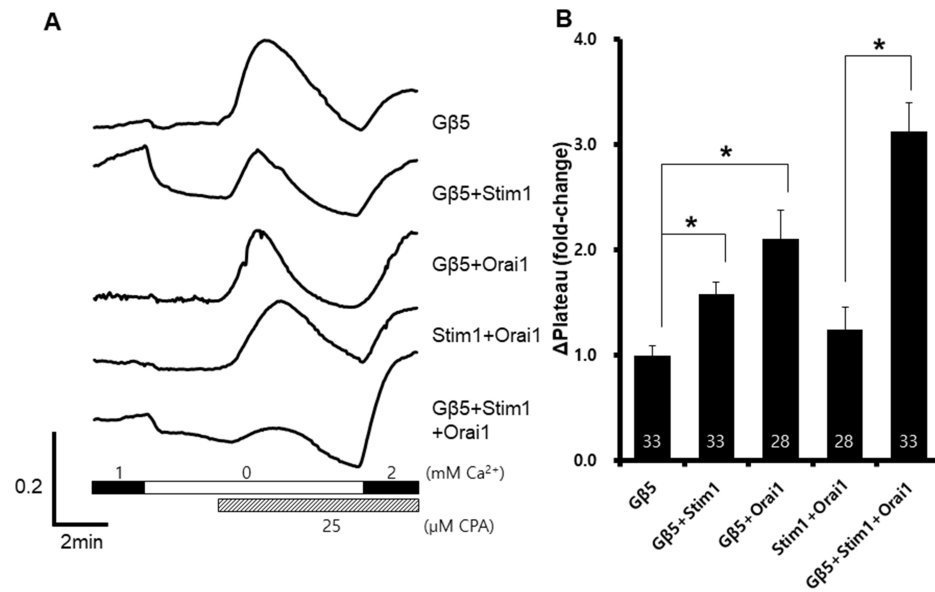


Fig. 2. Increased Stim1-Orai1 dependent store-operated Ca²⁺ entry after GNB5 overexpression. Gβ5, Stim1, and Orai1 were co-expressed in HEK293T cells. A representative trace (A) summarizes the data (B), and the number inside the column is the total cell number from multiple independent experiments (n=3-4). Data were expressed as the mean±SEM. *p<0.05 compared with control.

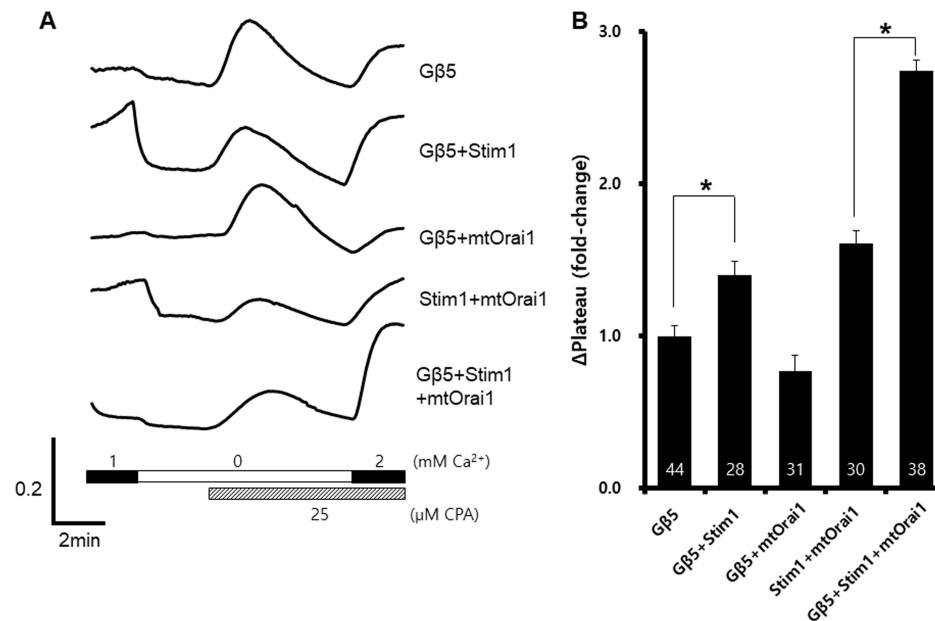


Fig. 3. Effects of a loss-of-functional Orai1 mutant on store-operated Ca²⁺ entry. Wild-type Gβ5, wild-type Stim1, and Orai1^{R91W} were co-expressed in HEK293T cells. A representative trace (A) summarizes the data (B), and the number inside the column is the total cell number from multiple independent experiments (n=3-4). Data were expressed as the mean±SEM. *p<0.05 compared with control.

These results suggested that Gβ5 interacts with Orai1 or Stim1, or both.

Mutant Orai1 has no effect on the GNB5-induced enhancement of SOCE

To confirm the target of Gβ5, we co-expressed Orai1^{R91W}, a loss-of-function mutant of Orai1, with Gβ5. Although mutant Orai1 was expressed, SOCE was still enhanced by Gβ5 expression (Fig. 3B). This might be a result of the presence of endogenous Orai1 and its participation in SOCE.

Stim1-ΔERM abolishes the augmentation of SOCE induced by GNB5 gene expression

ΔERM-Stim1^{D76A} is a dominant-negative form of *STIM1* (DN-STIM1). When we expressed DN-STIM1 with *GNB5*, the enhancement of SOCE was diminished (Fig. 4B). This result supported the hypothesis that *GNB5* may interact with *STIM1*.

DISCUSSION

GPCR signaling is a well-established pathway by which intracellular signals interact with extracellular signals [23]. Coupling

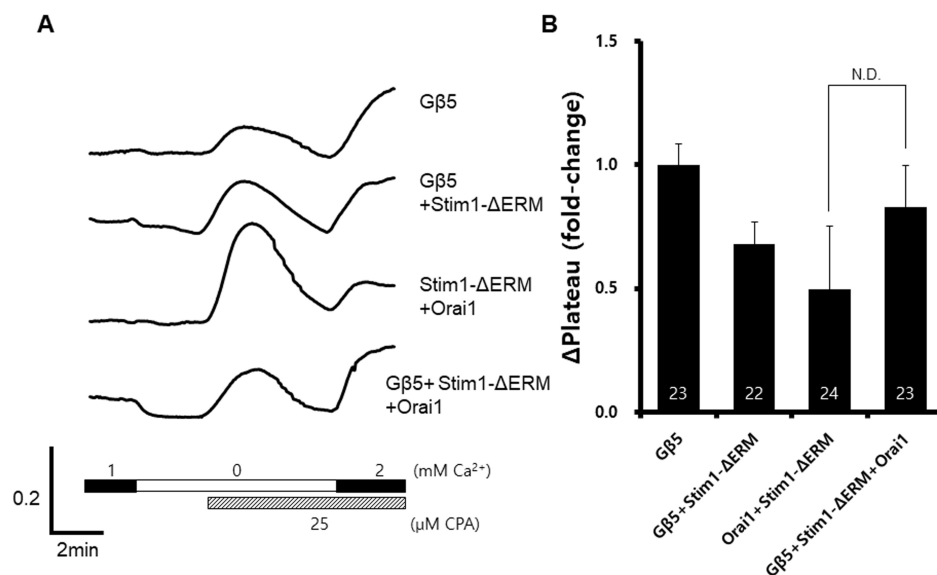


Fig. 4. Diminished enhancement of the store-operated Ca^{2+} entry induced by $\text{G}\beta 5$ and the co-expression of a dominant-negative Stim1. Wild-type $\text{G}\beta 5$, $\Delta\text{ERM-Stim1}^{\text{D76A}}$, and wild-type Orai1 were co-expressed in HEK293T cells. A representative trace (A) summarizes the data (B), and the number inside the column is the total cell number from multiple independent experiments ($n=3-4$). N. D.=not determined.

between phospholipase $\text{C}\beta$ ($\text{PLC}\beta$) induces inositol-1, 4, 5-triphosphate (IP_3) generation and subsequent cytosolic Ca^{2+} ion cascades [24]. Specific $\text{G}\beta\gamma$ subunit complexes activate $\text{PLC}\beta$ and cause its translocation to the cytosol, to subsequently interact with the IP_3 receptor in the endoplasmic reticulum (ER), and induce changes in Ca^{2+} conductance [25,26]. Evidence also suggests a direct activation of IP_3 by $\text{G}\beta\gamma$ complexes [27]. However, the role of $\text{G}\beta\gamma$ complexes in SOCE remains unknown. From our present results, we concluded that $\text{G}\beta 5$ expression enhanced SOCE (Fig. 1B).

The mechanism by which $\text{G}\beta 5$ increased SOCE remains unknown. Stim1-Orai1 dependent SOCE was significantly increased by the expression of $\text{G}\beta 5$ (Fig. 2B). The replacement of wild-type Orai1 with a loss-of-function Orai1 mutant, Orai1^{R91W}, did not alter SOCE. However, a loss-of-function Stim1 mutant, $\Delta\text{ERM-Stim1}^{\text{D76A}}$, inhibited the SOCE induced by $\text{G}\beta 5$. These results suggested that *GNB5* gene expression may affect Stim1 action, however, whether this is a direct or indirect relationship is unclear. Further studies are needed to evaluate the interactions of $\text{G}\beta 5$ with Stim1.

The Stim1 protein has an EF-hand domain for calcium sensing at the ER lumen [28]. The cytosolic portion of Stim1 begins with an ERM domain that starts with a coiled-coil domain for oligomerization and oligomer-induced recruitment of Orai1 protein to the plasma membrane [19]. In the ERM domain of the Stim1 protein, the Stim1 Orai1 activating region (SOAR) performs key functions in the activation of Orai1 [19,29]. In our study, $\Delta\text{ERM-Stim1}^{\text{D76A}}$ inhibited the SOCE induced by $\text{G}\beta 5$. $\text{G}\beta 5$ also has α -helical extension of 20 amino acids forms a coiled-coil domain [30]. This suggests that the coiled-coil domain may enhance the recruitment of the Orai1 protein.

Many Stim1-modulating molecules have been identified. For example, SARAF and TEM66 bind to the Stim1 SOAR domain to cause reduced Orai1 activity [31]. STIMATE and TEM110 in-

teract with Stim1 to promote its conformational switch changes [32]. Cracm1, calcium related activating current modulator 1, is a plasma membrane protein that is essential for SOCE [33]. The R7 RGS protein, regulator of G-protein signaling protein, mediates $\text{G}\beta 5$ neuronal signaling [8,9]. Similarly, *GNB5* may act by modulating molecules related to *STIM1*.

The TRPC pathway is another important pathway of SOCE control in mammalian cells [19,29,34]. In a mouse model that lacks *Trpc3*, reduced SOCE in pancreatic acinar cells and reduced amylase secretion were observed [35]. Stim1 SOAR domains are also important for activation of *Trpc3* [19]. Previous studies indicated that $\text{G}\beta 5$ interacts with Stim1, and its expression promotes *Trpc3*-dependent SOCE. Studies of SOCE in $\text{G}\beta 5$ -*Trpc3* expressing cells, compared with cells expressing *Trpc3* alone, have not been reported. Therefore, the role of $\text{G}\beta 5$ and Stim1 should be further studied.

The mechanism and function of the $\text{G}\beta\gamma$ complexes still remain to be determined. $\text{G}\beta\gamma$ complexes are potential therapeutic targets, therefore, our findings may contribute to the development of novel approaches for cures of diseases.

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CONFLICTS OF INTEREST

The authors declare no conflicts of interest.

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