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G γ 7 proteins contribute to coupling of nociceptin/orphanin FQ peptide (NOP) opioid receptors and voltage-gated Ca²⁺ channels in rat stellate ganglion neurons

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Abstract

The nociceptin/orphanin FQ peptide (NOP) opioid receptors regulate neurotransmitter release via inhibition of voltage-gated Ca²⁺ channels (Ca_v2.2) in sympathetic and sensory neurons. Stimulation of NOP receptors by its endogenous agonist, nociception (Noc), leads to membrane-delimited, voltage-dependent (VD) block of Ca_v2.2 channel currents mediated by G $\beta\gamma$ protein subunits. Previously we reported that the pertussis toxin-sensitive G α i1 and G β 2/ β 4 isoforms mediate the functional coupling of NOP opioid receptors with Ca_v channels in rat stellate ganglion (SG) sympathetic neurons. In the present report we extended our studies by identifying the G γ subunit that forms the heterotrimer within this signaling pathway. Small interference RNA (or siRNA) was employed to silence the expression of the natively expressed G γ subunits. Initial PCR assays indicated that SG neurons expressed seven G γ subunits. Silencing G γ 3 subunits did not alter signaling between NOP receptors and Ca²⁺ channels. However, after G γ 7 isoforms were silenced, the Noc-mediated inhibition of Ca_v channels was significantly decreased when compared to SG neurons transfected with scrambled siRNA. We observed that G γ 10 and G γ 11 mRNA levels increased 2.5- and 2.7-fold, respectively, after G γ 7 subunits were silenced. However, this compensatory increase in mRNA expression did not appear to fully rescue the NOP receptor coupling efficiency. Additionally, both G γ 2 and G γ 5 levels increased 50 and 75%, respectively, while G γ 3 and G γ 4 expression levels remained relatively unchanged. Taken together, our findings suggest that the G α i1/G β 2(β 4)/G γ 7 heterotrimeric G protein complex determines the NOP receptor-mediated modulation of Ca_v channels in SG neurons.

Keywords

G protein subunits; opioid receptors; Ca²⁺ channels; G protein-coupled receptors

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Conflict of Interest There is no conflict of interest to report.

1. Introduction

Nociceptin/orphanin FQ peptide (NOP) opioid receptors are G protein-coupled receptors (GPCR) that are activated by the endogenous ligand, nociceptin (Noc). The NOP receptor system has been demonstrated to regulate processes including cardiovascular, pain, stress, renal function, learning and memory [1–2]. Recent studies also have shown that this opioid receptor plays an important role in sepsis and inflammatory conditions [3–4]. NOP receptor activation leads to voltage-gated Ca^{2+} channel (Ca_V) inhibition, G protein inwardly rectifying K^+ (GIRK) channel activation and negative coupling to adenylyl cyclase [1–2]. $\text{Ca}_V2.2$ channels, the major Ca^{2+} ion carriers in SG neurons [5], have been shown to interact with NOP receptors, which influences their trafficking to the cell surface and internalization processes [6–7].

SG neurons, which innervate cardiac muscle and exert chronotropic and inotropic effects, express NOP opioid receptors [8–9]. Noc mediates its effects by coupling NOP receptors to members of the pertussis toxin (PTX)-sensitive $\text{G}_{\alpha i/o}$ family of heterotrimeric G proteins [1]. Previously, we reported that the Noc-mediated modulation of Ca_V currents occurred primarily via PTX-sensitive $\text{G}_{\alpha i1}$ protein subunits in rat SG neurons [10]. Subsequently, employing small interference RNA (siRNA) approach, both $\text{G}\beta 2$ and $\text{G}\beta 4$ subunits were shown to mediate the coupling of NOP receptors with Ca_V channels [11].

$\text{G}\gamma$ subunits are crucial for efficient attachment of the $\text{G}\beta\gamma$ complex to the cell membrane. There are twelve known $\text{G}\gamma$ subunits [12] and recent studies have indicated that the interaction between GPCR and effector proteins can be influenced by $\text{G}\gamma$ proteins [13–16]. In the present study, we investigated whether specific $\text{G}\gamma$ subunits influenced coupling specificity of NOP opioid receptors with Ca_V channels in rat SG neurons.

2. Materials and Methods

2.1. Rat SG tissue isolation

The experiments performed on animals were approved by the Penn State College Medicine Institutional Animal Care and Use Committee. Male Sprague-Dawley rats were anesthetized with CO_2 and then rapidly decapitated with a laboratory guillotine. After the SG tissue was removed, it was desheathed in ice-cold Hanks' balanced salt solution (Sigma-Aldrich, St. Louis, MO). Following removal of the connective tissue, multiple parallel slits (~ 1 mm apart) were made perpendicular to the long axis. The SG tissue was then placed in an RNase- and DNase-free 2 ml microcentrifuge tube containing optimum minimal essential medium (O-MEM, Thermo-Fisher Scientific, Grand Island, NY) supplemented with 2 mM 2,3-butanedione monoxime (BDM, Sigma-Aldrich) and kept on ice until ready for siRNA transfection.

2.2. Small interference RNA (siRNA) transfection

SG neurons were transfected with siRNA employing both electroporation and lipofection as previously described [11]. $\text{G}\gamma 7$ subunit siRNA sequences designed in this study were chosen based on an eight point criteria with a scoring system (10 = best, 1 = worse) [17]. The sequences were obtained with a macro written by Dr. Stephen R. Ikeda (National Institutes

of Health/NIAAA) on Igor Pro 6.37 (WaveMetrics, Inc., Lake Oswego, OR). The target siRNA oligonucleotide sequences for rat G γ 7, score in parenthesis, were as follows: sense, 5'–UGA UGU CAG GUA CUA ACA AUU-3' (8) corresponding to nucleotide positions 2–20. The antisense sequence was 5'–UUG UUA GUA CCU GAC AUC AUU-3'. The control SG tissue was transfected with scrambled siRNA sequences (both from Thermo-Fisher Scientific).

Immediately after dissection, the SG tissues were preserved in ice-cold Opti-MEM and BDM prior to transfer to the electroporation solution. The lipofection solutions were prepared first and contained Opti-MEM, scrambled siRNA or G γ 7 siRNA (both at 1.5 μ M), BDM (2 mM) and 10 μ l Lipofectamine 2000 (Thermo-Fisher Scientific) in a final volume of 1 ml per group. Thereafter, each lipofection solution was transferred to a 22 mm dish (12-well plate) and stored in a 37°C humidified incubator (95% air-5% CO $_2$) until ready for use. The electroporation solutions were prepared by mixing R solution (provided with NEON electroporator kit, Thermo-Fisher Scientific) with either scrambled siRNA (1.5 μ M) or G γ 7 (1.5 μ M) siRNA and BDM (2 mM). The ganglia were then transferred into their respective solution for 15–20 min at room temperature. After the incubation period in the electroporation solution, the ganglia were drawn up into a 100 μ l electroporator tip and electroporated using the NEON electroporator (Thermo-Fisher Scientific), 3 pulses, each 1000 V and for 20 msec.

Immediately after electroporation the ganglia were placed in the 22 mm dish containing the appropriate lipofection solution and incubated for 5 hr in the humidified incubator. Following the 5 hr incubation, the ganglia were rinsed 3 times with MEM (minimum essential medium, Thermo Fisher Scientific) and placed back in the incubator. The transfection procedure was repeated 48 and 96 hr later.

2.3. SG tissue dispersion

Following the transfection period, the SG tissue was enzymatically dissociated as previously described [11]. Briefly, the tissue was incubated in a shaking water bath (35°C) for 60 min in Earle's balanced salt solution containing 0.6 mg/ml collagenase Type D (Roche Diagnostics, Indianapolis, IN), 0.4 mg/ml trypsin (Worthington Biochemicals, Freehold, NJ) and 0.1 mg/ml DNase (Sigma-Aldrich). Afterwards, the cells were dissociated in the flask by vigorous shaking and the neuron suspension was centrifuged twice for 6 min at 53 \times g and resuspended in MEM supplemented with 10% fetal bovine serum, 1% penicillin-streptomycin and 1% glutamine. The dispersed neurons were plated onto 35 mm poly-L-lysine-coated dishes and stored in a humidified incubator.

2.4. Electrophysiology and data analysis

SG Ca $^{2+}$ channel currents were recorded at room temperature employing the whole-cell variant of the patch-clamp technique. Recording pipettes were pulled on a micropipette puller (P-97, Sutter Instrument, Novato, CA) and fire polished on a microforge. Ca $^{2+}$ channel currents were acquired with a patch-clamp amplifier (Axopatch 200B, Molecular Devices, Sunnyvale, CA), analog filtered at 2 kHz (–3 dB; 4-pole low-pass Bessel filter) and digitized at 10 kHz with an A/D converter board (ITC-18, HEKA Instruments Inc., Bellmore, NY)

employing custom-designed software (S5, Dr. Stephen R. Ikeda, NIH/NIAAA) on a PowerMacG4 computer (Apple Inc., Cupertino, CA). The series resistance and membrane capacitance were electronically compensated (80–85%). Ca² currents were evoked with the triple-pulse voltage protocol [18]. This protocol is routinely employed to study the agonist-mediated voltage-dependent inhibition of Ca² channel currents. The protocol consists of a test pulse to +10 mV (prepulse) followed by a large depolarizing conditioning test pulse to +80 mV, a brief return to –80 mV, and followed by a test pulse to +10 mV (postpulse). The peak Ca² current amplitude was measured isochronally 10 msec after the initiation of the prepulse and postpulse. Data and statistical analyses were performed with Igor Pro 6.37 and Prism 6.0 (GraphPad Software, Inc., San Diego, CA) software packages with $P < 0.05$ considered statistically significant. Current traces and graphs were created with both Igor Pro 6.37 and Graphic (Autodesk, San Rafael, CA) software packages. The data shown are mean \pm SE.

2.5. Solutions and drugs

The internal pipette solution contained (in mM): 80 *N*-methyl-D-glucamine, 20 tetraethylammonium hydroxide (TEA-OH), 11 EGTA, 10 HEPES, 1 CaCl₂, 4 Mg-ATP, 0.3 Na₂GTP, 20 CsCl, 40 CsOH and 14 tris creatine phosphate. Methanesulfonic acid was used to adjust the final pH to 7.2 and the osmolality ranged from 293–303 mOsmol/kg. The external solution consisted of (in mM): 145 TEA-OH, 140 methanesulfonic acid, 10 HEPES, 15 glucose, 10 CaCl₂, and 0.0003 tetrodotoxin. The pH was adjusted to 7.4 with TEA-OH and the osmolality ranged from 316–320 mOsmol/kg. A stock solution of Noc (Tocris Cookson, Ellisville, MO) was prepared in water and diluted in the external solution to its final concentration.

2.6. Quantitative real-time-PCR (QRT-PCR)

In this set of experiments, RNA from SG tissue was isolated with the Nucleospin RNA/Protein Kit (Macherey-Nagel, Bethlehem, PA). RNA concentrations were determined with the Qubit@2.0 fluorometer (Thermo Fisher Scientific) and stored at –80°C until ready for use. For QRT-PCR assays, equal quantities of total RNA were used to synthesize cDNA using the High Capacity cDNA RT Kit (Thermo Fisher Scientific). For the QRT-PCR reactions, we employed the TaqMan Gene Expression Assay (Thermo Fisher Scientific) specific for rat G γ and GAPDH with equal cDNA quantities for each group. The assays were then run on a QuantStudio 12K Flex Real-Time PCR system (Thermo Fisher Scientific) and analyzed using the comparative *C_t* method. The results were normalized to internal GAPDH mRNA controls.

3. Results

To determine which of the known G γ subunits couple NOP receptors to Ca_v, we first focused on the G γ 3 isoform, which has been reported to be involved in mu opioid receptor signaling cascades associated with rewarding of high fat diets in female mice [19]. Thus, G γ 3 subunits were silenced in SG neurons. QRT-PCR assays of SG tissue were performed 72, 96 and 120 hr post-siRNA transfection. When compared to scrambled siRNA-transfected SG tissue, G γ 3 mRNA levels were 76, 92 and 91% lower 72, 96, and 120 hr post-

transfection, respectively. However, when the NOP receptor-mediated inhibition of Ca_V was examined, the modulation of the Ca^{2+} currents was indistinguishable in scrambled and $\text{G}\gamma 3$ siRNA-transfected groups. For instance, exposure to Noc (1 μM) resulted in a mean Ca^{2+} current block of $61.1 \pm 4.2\%$ ($n=9$) and $58.2 \pm 6.8\%$ ($n=6$) in SG neurons transfected with scrambled and $\text{G}\gamma 3$ siRNA, respectively, 120 hr post-transfection.

Thereafter, $\text{G}\gamma 7$ proteins were targeted and their silencing in SG neurons was optimized. Figure 1 shows time courses of Ca^{2+} current modulation by Noc in SG neurons 96 hr post-transfection with either scrambled siRNA (Fig. 1A) or $\text{G}\gamma 7$ siRNA (Fig. 1B). Pre- and post-pulse Ca^{2+} currents were evoked every 10 sec with the triple pulse protocol shown in Fig. 1A (top right) and the current traces are shown to the right. The protocol consists of two identical test pulses to +10 mV (holding potential of -80) separated by a large depolarizing conditioning pulse to +80 mV. Ca^{2+} current amplitude was measured isochronally 10 msec after the initiation of the prepulse. In the control neuron, it can be observed that prior to Noc (1 μM) application the prepulse current (trace 1) was fast and reached a plateau within 5 msec. During Noc application, the prepulse current was blocked maximally (~ 50%) within 20 sec. It can also be observed that the prepulse current (trace 3) exhibited kinetic 'slowing', characteristic of the voltage-dependent (VD) modulation of Ca^{2+} currents [20]. Following a recovery period, the neuron was exposed to NE (10 μM) and the prepulse current was also blocked 50% (traces 5 and 7). This $\alpha 2$ -adrenergic receptor agonist was employed as a positive control for stimulation of G protein signaling. The time course in Figure 1B shows the modulation of Ca_V currents of an SG neuron transfected with $\text{G}\gamma 7$ siRNA in the presence of Noc or NE. Similar to the scrambled siRNA-transfected neuron, application of Noc led to a 50% block of Ca^{2+} currents. However, it can be observed that the cell exhibited a delay (> 60 sec) in reaching maximal inhibition (trace 3) when compared to the control SG neuron. The onset of the steady-state Noc-mediated Ca^{2+} current inhibition was fit to a single exponential function. The mean tau values for control and $\text{G}\gamma 7$ siRNA-transfected cells were 30.4 ± 4.4 and 46.5 ± 8.4 sec ($P = 0.16$, NS, unpaired t -test), respectively. On the other hand, a delay was not observed following NE exposure (trace 7) when compared to the control neuron. The summary plot in Figure 1C illustrates that silencing $\text{G}\gamma 7$ did not significantly alter the coupling pathways between NOP receptors and Ca_V channels.

In the next set of experiments, we examined the coupling in SG neurons 120 hr post-siRNA transfection. Figures 2A and 2B are time courses of Ca^{2+} current inhibition for neurons transfected with scrambled and $\text{G}\gamma 7$ siRNA, respectively. Figure 2A shows that the modulation of Ca^{2+} currents by Noc or NE were similar to control the group described for Fig. 1A. Further, the Ca^{2+} currents were blocked in a VD manner. Exposure of the $\text{G}\gamma 7$ siRNA-transfected neuron to Noc (Fig. 2B), on the other hand, again resulted in a delay in reaching maximal Ca^{2+} current inhibition as well as an attenuation of the current block. The mean tau value for the onset of the steady-state Noc-mediated Ca^{2+} current inhibition in control neurons was 21.0 ± 3.1 sec, while it was 37.5 ± 9.6 sec for $\text{G}\gamma 7$ siRNA-transfected cells ($P=0.14$, NS, unpaired t -test). Unlike Noc, the NE-mediated modulation of Ca_V currents was unaffected by $\text{G}\gamma 7$ silencing. The summary dot plot in Figure 2C shows that $\text{G}\gamma 7$ siRNA transfection significantly ($P < 0.05$) decreased the Noc-mediated Ca^{2+} modulation in SG neurons. These results suggest that with this approach, a period of 120 hr was required to decrease $\text{G}\gamma 7$ expression levels and alter NOP receptor coupling.

The lack of specific rat G γ 7 protein antibodies precluded us from performing Western blotting assays on siRNA-transfected tissue to measure protein levels. However, QRT-PCR assays on scrambled and G γ 7 siRNA-transfected SG tissue were performed 72, 96 and 120 hr post-siRNA transfection. Figure 3A shows that G γ 7 mRNA expression levels in SG tissue decreased 80% 72 hr post-transfection and greater than 98% 96–120 hr following siRNA transfection. In a second set of experiments, QRT-PCR assays were performed in SG tissue to examine the effect of G γ 7 siRNA transfection on mRNA expression levels of other G γ protein subunits 120 hr post-siRNA transfection. Of the six G γ proteins tested, G γ 10 and G γ 11 mRNA levels exhibited the highest increased changes of 2.5- and 2.7-fold, respectively. G γ 2 and G γ 5 increased by 50 to 75% while G γ 3 and G γ 4 levels remained relatively unchanged.

4. Discussion

In the present study we examined whether G γ subunits influence coupling specificity between NOP receptors and Ca $_v$ channels. We identified previously that G β 2, G β 4 and PTX-sensitive G α subunits play a crucial role in the Noc-mediated modulation of Ca $_v$ currents in SG neurons [10–11]. The approach employed in our experiments was to silence specific G γ subunits employing siRNA assays. Since twelve G γ proteins have been described and few studies have explored G γ coupling specificity, we employed a judicious approach. G γ 1 and G γ 8 subunits, for instance, exhibit restricted tissue distribution and, thus, were not studied [12, 21]. Our initial focus was the G γ 3 isoform that has been previously linked to mu opioid receptor signaling mechanisms in response to a high-fed diet [19]. Our results showed that silencing G γ 3 did not overtly alter the coupling between NOP receptors and Ca $^{2+}$ channels, which suggests this isoform does not play a crucial role in this signaling pathway.

The sequence homology of G γ 7 [12, 21], coupled with its functional interaction with G γ 3 subunits [22], prompted us to next probe the role of this isoform on the NOP receptor signaling pathway. After the G γ 7 proteins were silenced, the Noc-mediated block of Ca $^{2+}$ currents was significantly decreased, indicating it is an important component in this signaling process. That G γ subunits can influence coupling has been shown with other GPCR. Robishaw and colleagues, for instance, reported that G γ 7 subunits were critical components of the G protein heterotrimer employed by adenosine A $_{2A}$ receptors in mice striatum [14], dopamine (D1) receptors [15] and β 1 adrenergic receptors [16]. Others have found that G γ 11 subunits exhibited preferential coupling to adenosine A $_1$ and 5-HT $_{1A}$ receptors [13].

Similar to our previous study [10], a 120 hr period was necessary to effectively silence G γ 7 proteins and alter coupling specificity between NOP receptors and Ca $_v$ channels. On the other hand, we also reported that a 72 hr period was required to silence G β 2 subunits to significantly uncouple NOP receptor signaling in SG neurons [11]. However, unlike G γ 7 and G α i1, silencing G β 2 led to an upregulation of G β 4 mRNA and protein such that the Noc-mediated Ca $^{2+}$ current inhibition was restored. In the present study, QRT-PCR results showed upregulation of some G γ isoforms following G γ 7 siRNA transfection, but lacked a compensation of the NOP coupling pathway.

The mRNA expression profile of non-targeted G γ subunits indicated that the G γ 7 protein levels influence the expression of G γ 10 and G γ 11 subunits, and to lesser extent that of G γ 2 and G γ 5. The upregulation of these G γ proteins suggests that there is a compensatory mechanism for a G γ 7-dependent signaling pathway, but independent of Ca $_v$ channel modulation. It is also interesting to note that we did not observe decreases of G γ expression levels in light of a recent study which reported a series of experiments designed to silence eight G α and G β isoforms in HeLa cells [23]. In that report, for instance, G α i1 expression levels significantly increased while G β 2 and G β 4 levels decreased significantly following G α s silencing. Further, when G β 1- β 4 subunits were silenced, the expression levels of all G α subunits tested significantly decreased. The effect of silencing either G α or G β isoform on G γ subunits was not determined [23].

In summary, the findings of the present study show that G γ 7 subunits are crucial elements for NOP receptor coupling to Ca $_v$ channels, while the G γ 3 does not appear to be involved in this signaling pathway. In addition, as G γ 7 mRNA levels decreased 96 hr post-transfection, and presumably protein, there was a slowing of the onset of the Noc-mediated Ca $^{2+}$ current inhibition when compared to scrambled siRNA-transfected cells. The modulation of Ca $^{2+}$ currents was significantly decreased 120 hr post-siRNA transfection when compared to the control group. Following G γ 7 silencing, the mRNA levels of other G γ proteins (G γ 2, G γ 5, G γ 10 and G γ 11) increased, yet the coupling between Ca $_v$ channels and NOP receptors was diminished. The current results, in combination with our previous reports, indicate that NOP receptors employ G α i1, G β 2/G β 4, and G γ 7 proteins to modulate Ca $_v$ channel function (Fig. 4). Thus, the G $\alpha\beta\gamma$ heterotrimer highly impacts the NOP receptor signal transduction pathway in SG neurons. This striking specificity could be exploited therapeutically in conditions, such as sepsis, where high Noc levels are associated with increased morbidity and/or mortality (see [4] for review). Targeting these G protein isoforms would suppress the NOP signaling pathway, preserve modulation of Ca $_v$ channels by other GPCR and markedly limit side effects.

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Highlights

- Examined G γ coupling specificity of NOP receptors and Ca $_V$ in rat sympathetic neurons.
- Silencing G γ 7 isoform significantly decreased the Noc-mediated Ca $_V$ modulation.
- G γ 7 siRNA-transfected cells exhibited increased G γ 2, G γ 5, G γ 10 and G γ 11 mRNA levels.
- G α i1/G β 2(β 4)/G γ 7 heterotrimer couples mediate coupling of NOP receptors and Ca $_V$.

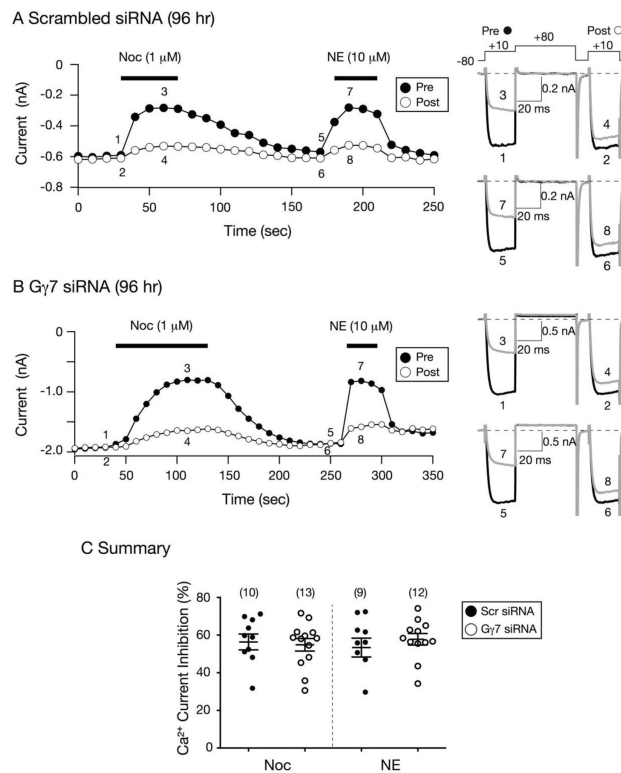


Figure 1. Effect of G γ 7 knockdown on the Noc- and NE-mediated Ca^{2+} current inhibition 96 hr post siRNA transfection

A and B) Time courses of peak Ca^{2+} current amplitude inhibition for pre- (●) and postpulse (○) acquired from the sequential application of Noc (1 μ M) and NE (10 μ M) in neurons transfected with control scrambled (A) and G γ 7 (B) siRNA, respectively. Superimposed Ca^{2+} current traces evoked with the 'triple-pulse' voltage protocol (shown on top of 1A, right) in the absence (1 and 2; 5 and 6, black) or presence (3 and 4; 7 and 8, grey) of Noc and NE, respectively, are shown to the right. Currents were evoked every 10 sec. The filled bars indicate the application of agonists. C) Summary dot plot of mean (\pm SE) Ca^{2+} current inhibition produced by application of Noc or NE in neurons transfected with control scrambled or G γ 7 siRNA. Inhibition was determined from the Ca^{2+} current amplitude measured isochronally at 10 msec into the prepulse (+10 mV) in the absence or presence of Noc or NE. Numbers in parenthesis indicate the number of neurons tested.

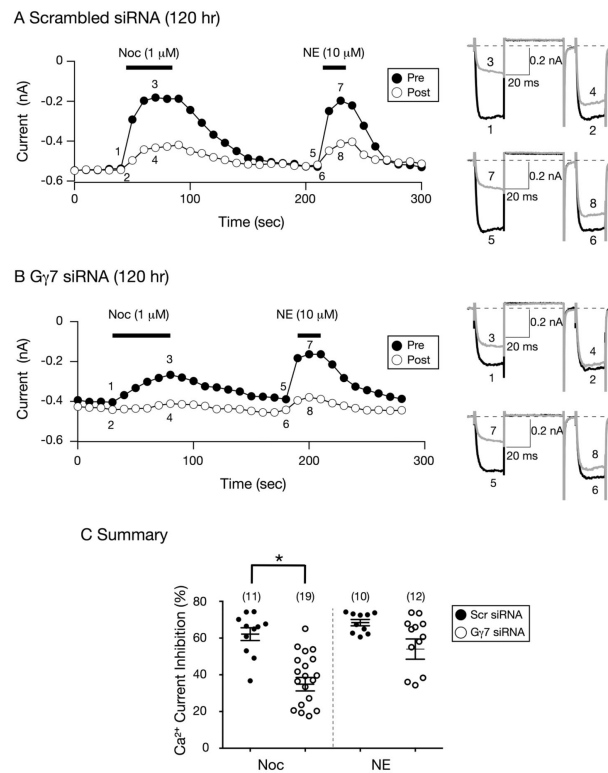


Figure 2. Effect of G γ 7 silencing 120 hr post siRNA transfection on the Noc- and NE-mediated Ca²⁺ current inhibition in SG neurons

A and B) Time courses of peak Ca²⁺ current amplitude inhibition for pre- (●) and postpulse (○) acquired from the sequential application of Noc (1 μ M) and NE (10 μ M) in neurons transfected with control scrambled (A) and G γ 7 (B) siRNA, respectively. Superimposed Ca²⁺ current traces (shown to the right) evoked with the 'triple-pulse' voltage protocol (shown on top of 1A) in the absence (1 and 2; 5 and 6, black) or presence (3 and 4; 7 and 8, grey) of either NE or Noc. Currents were evoked every 10 sec. The filled bars indicate the application of agonists. C) Summary graph of mean (\pm SE) Ca²⁺ current inhibition produced by application of Noc or NE in neurons transfected with scrambled or G γ 7 siRNA. Inhibition was determined from the Ca²⁺ current amplitude measured isochronally at 10 msec into the prepulse (+10 mV) in the absence or presence of Noc or NE. Numbers in parenthesis indicate the number of neurons tested. * P < 0.05 compared to neurons transfected with scrambled siRNA, Student's t test.

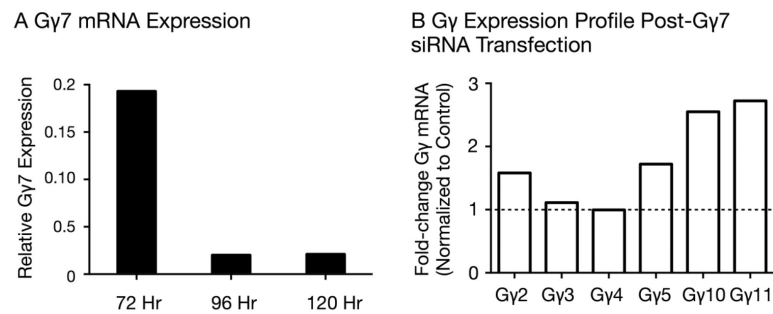


Figure 3. Quantitative assessment of G γ subunit mRNA expression by QRT-PCR in SG tissue 72–120 hr post-G γ 7 siRNA transfection

A) QRT-PCR analysis showing G γ 7 mRNA expression levels in SG tissue transfected with either scrambled siRNA or G γ 7 siRNA 72, 96 and 120 hr post-transfection. For each experimental group, 1–2 rats were employed. B) Relative G γ mRNA expression levels 120 hr post-transfection in SG tissue following G γ 7 subunit silencing. QRT-PCR was carried out with total RNA from scrambled and G γ 7 siRNA-transfected SG tissue. The fold-differences were calculated with the Ct values for each probe and corrected for GAPDH expression levels in each sample. The corrected expression level of each G γ subunit was then normalized to its corresponding value obtained from SG tissue transfected with scrambled siRNA.

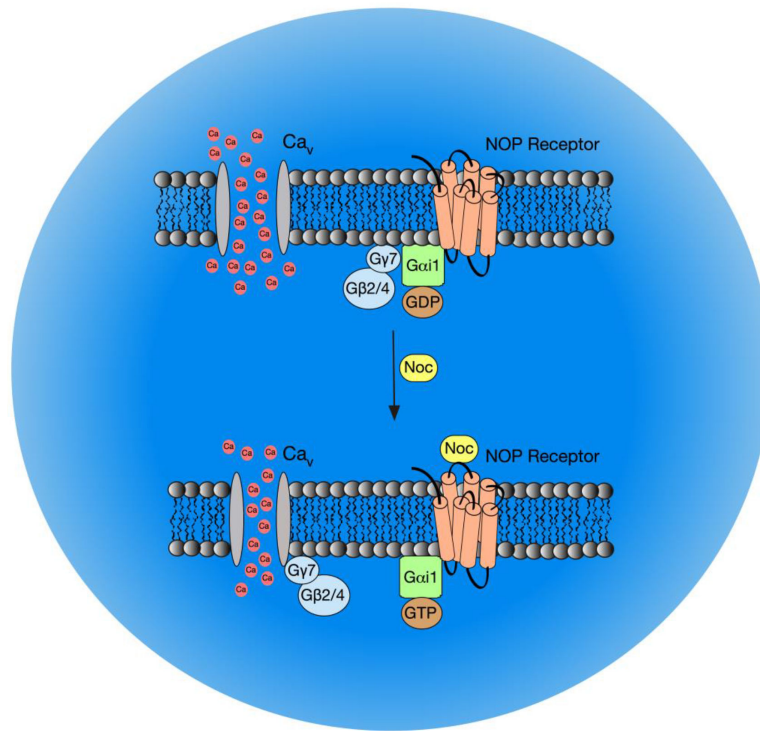


Figure 4. Diagram summarizing the NOP receptor signaling pathway that mediates Ca_v inhibition following NOP receptor stimulation by Noc in rat SG neurons
 Prior to Noc exposure, NOP receptors are coupled to the Gai1/β2(β4)/γ7 heterotrimer. Upon activation by Noc, free Gβ2(β4)γ7 dimers bind to Ca_v channels leading to VD inhibition of Ca²⁺ currents.