# Increased Na<sub>v</sub>I.7 expression in the dorsal root ganglion contributes to pain hypersensitivity after plantar incision in rats

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### Abstract

Postoperative pain remains a complex problem that is difficult to manage in the clinical context, seriously affecting rehabilitation and the quality of life of patients after surgery. Nociceptors, of which the cell bodies are located in the dorsal root ganglion, are crucial for initiating and conducting the pain signal. The peripheral voltage-gated sodium channels, including Na, 1.7, which is mainly expressed in the dorsal root ganglion, are key to understanding the mechanism underlying postoperative pain. Na, 1.7, in particular, of which mutations in the encoding gene (SCN9A) can determine whether pain occurs, has aroused most attention. Previous studies have shown that Nav I.7 in dorsal root ganglion is critical for the development of inflammatory pain and some neuropathic pain. However, the expression of NavI.7 in the dorsal root ganglion after surgery and its role in postoperative pain hypersensitivity remain unclear. Therefore, in this study, in order to gain a better understanding of the role of dorsal root ganglion NavI.7 in pain hypersensitivity following operation, we dynamically examined the pain-related behavior and expression of NavI.7 in L4–L6 dorsal root ganglion before and after plantar incision in rats (an acute postoperative pain model). After plantar incision, the mechanical and thermal pain threshold decreased significantly, the cumulative pain score was increased significantly, meanwhile quantitative polymerase chain reaction and Western blotting results showed that expression of Nav I.7 in L4-L6 dorsal root ganglion was enhanced significantly. After pretreatment using SCN9A-RNAi-LV delivered via an intrathecal tube, immunohistochemistry showed that increased expression of Nav1.7 in L4–L6 dorsal root ganglion after plantar incision was inhibited, as also confirmed by quantitative polymerase chain reaction and Western blotting. Moreover, pain hypersensitivity was alleviated. These results suggested that Na, 1.7 of L4–L6 dorsal root ganglion plays an important role in the development of pain hypersensitivity after plantar incision.

### **Keywords**

Dorsal root ganglion, Nav1.7, pain hypersensitivity, plantar incision, postoperative pain

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# Introduction

Postoperative pain is the most common form of acute pain. In 2015, the annual global number of surgeries exceeded 310 millions.<sup>1</sup> Although analgesic technologies have continually improved (e.g., multimodal analgesia, advanced analgesia, etc.), 39%-50% of surgical patients experience moderate to severe acute postoperative pain, which seriously affects the rehabilitation of these patients.<sup>2,3</sup> Thus, there are tens of millions of patients who suffer from moderate to severe pain every year worldwide. Therefore, effective treatment of

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Xianwei Zhang, Department of Anesthesiology, Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology, No. 1095 Jie-Fang Road, Wuhan 430030, Hubei, People's Republic of China. Email: ourpain@163.com

Creative Commons Non Commercial CC BY-NC: This article is distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 License (http://www.creativecommons.org/licenses/by-nc/4.0/) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (https://us. sagepub.com/en-us/nam/open-access-at-sage). postoperative pain remains an urgent and key problem in clinical practice.

There is strong evidence that peripheral voltage-gated sodium channels, including Na<sub>v</sub>1.7, which is primarily expressed in the dorsal root ganglion (DRG), are associated with initiating and conducting pain signal.<sup>4,5</sup> In tetrodotoxin-sensitive sodium particular, channel  $Na_v 1.7$ , which is able to amplify sub-threshold excitability and regulate neuronal excitability as a threshold channel, has recently emerged as a potential analgesics target because mutation of the gene encoding Nav1.7 (SCN9A) can determine whether pain is present or not.<sup>6-9</sup> In humans, gain-of-function mutations result in pain hypersensitivity (primary erythermalgia (PE),<sup>10</sup> paroxysmal extreme pain disorder,<sup>11</sup> and small-fiber neuropathy<sup>12</sup>), while loss-of-function mutations cause congenital insensitivity to pain.<sup>13</sup> In addition, previous studies of conditional knockout mice and multiple rodent models of inflammatory and neuropathic pain have shown that  $Na_v 1.7$  expressed in the DRG is crucial for generating and conducting pain signals, for example, inflammatory pain induced by formalin, carrageenan, nerve growth factor (NGF), or complete Freund's adjuvant and the neuropathic pain induced by chronic constriction injury.<sup>6,14,15</sup> Preoperative or perioperative use of nonselective sodium channel blockers (e.g., lidocaine) in patients can relieve postoperative pain sensitivity and reduce postoperative opioid consumption.<sup>16,17</sup> Furthermore, our previous study confirmed that genetic polymorphisms in SCN9A might affect postoperative pain sensitivity in surgical patients.<sup>18,19</sup> These findings suggested that Nav1.7 may play an important role in postoperative pain. However, its role in the development of postoperative pain remains unclear.

To explore the role of Na<sub>v</sub>1.7 in L4–L6 DRG in pain hypersensitivity after plantar incision, a rodent model of postoperative pain, we observed pain-related behavior and dynamically detected the expression of Na<sub>v</sub>1.7 in L4–L6 DRG before and after plantar incision in rats. Then, Na<sub>v</sub>1.7 in L4–L6 DRG was knocked down using *SCN9A*-RNAi-LV, administered via an intrathecal tube, and the effect on pain-related behavior after plantar incision was observed.

# Materials and methods

## Animals and groups

The study was approved by Animal Care and Use Committee of Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology. The protocols were conducted according to the ethical guidelines of the International Association for Pain Research. Male Sprague–Dawley rats, weighing 200 - 220 g, were provided by the Animal

Center of Hubei Province, China. The rats were kept under controlled conditions, at  $23 \pm 1^{\circ}$ C, with a 12-h light/dark schedule and had *ad libitum* access to food and water.

To detect Na<sub>v</sub>1.7 expression in the DRG of rats after plantar incision, the rats (N = 36) were randomly divided into two groups: control group (n = 6) and skin-fasciamuscle incision group (n = 30). The rats in the skinfascia-muscle incision group were further divided into five groups: 2 h (n = 6, rats were sacrificed at 2 h), 24 h (n = 6, rats were sacrificed at 24 h), 48 h (n = 6, rats were sacrificed at 48 h), 72 h (n = 6, rats were sacrificed at 72 h), and 120 h (n = 6, rats were sacrificed at 120 h). Before 0 h and at 2 h, 24 h, 48 h, 72 h, and 120 h following surgery, the rats underwent behavior testing (withdrawal responses, thermal hyperalgesia, and cumulative pain score). Six rats from each of the incision and the control groups were used for quantitative polymerase chain reaction (Q-PCR) (n = 3 per group) and Western blotting analysis (n=3 per group) after painrelated behavior.

To explore whether knockdown of  $SCN9A/Na_v 1.7$  in DRG is effective in attenuating pain hypersensitivity after plantar incision, the rats (N=48) were divided into four groups: a control group (n=12), an incision group (n=12), SCN9A-RNAi-LV + incision group (n=12), and a negative control lentivirus (NC-LV) + incision group (n=12). Before 0 h and at 2 h, 24 h, and 48 h following surgery, the rats were used for behavior testing. At 2 h following surgery, nine rats of each group were used for quantitative PCR (Q-PCR) (n=3 per group), Western blotting analysis (n=3 per group), and immunohistochemistry (n=3 per group); the other rats (three rats of each group) were used for behavior testing until 48 h after surgery.

# Construction of SCN9A-RNAi-LV and microinjection of virus via intrathecal catheter

For the knockdown experiment with *SCN9A*-RNAi-LV, the sequences of rat *SCN9A* siRNA lentivirus (*SCN9A*-RNAi-LV) were designed and synthesized using the following target sequences: 5'-ATCGAAGAA GCTAAACAGAAA-3'. The same vector without *SCN9A* siRNA was used as the NC-LV. The lentivirus vectors were constructed by Shanghai Genechem Co., Ltd. (Shanghai, China). The titer of the virus was  $5 \times 10^8$  TU/ml and stored at  $-80^{\circ}$ C.

According to previously described methods,<sup>20,21</sup> intrathecal catheters (PE-10, Smiths Medical, England) were constructed, and 3 days before operation, 10  $\mu$ l of *SCN9A*-RNAi-LV or NC-LV was administered intra-thecally, followed by a 10- $\mu$ l saline flush.

### Establishment of incisional pain in rats

According to previously reported methods,<sup>22</sup> the rats received general anesthesia with 2.5% isoflurane. Before surgery, each rat received 30,000 IU penicillin by intramuscular injection. Then, the left hindpaw plantar was sterilized with iodophor. A 1-cm longitudinal incision was made through the skin and underlying fascia (starting 0.5 cm from the proximal heel and extending toward the toes). The flexor digitorum brevis muscles of the plantar were exposed and incised longitudinally (leaving the muscle origin and insertion intact). After hemostasis by compression, the skin was sutured using 5–0 silk. The incision wound site was covered with penicillin ointment.

### Measurement of pain behaviors

All behavior tests were performed under quiet conditions in the morning.

Measurement of mechanical hyperalgesia. The rats were habituated in plastic cages  $(10 \times 10 \times 19 \text{ cm})$ , with a grid floor, for 30 min. Mechanical allodynia was measured using electronic von Frey filaments according the method of Vivancos et al.,<sup>23,24</sup> the tip of the filament was applied perpendicularly to the mid-plantar with gradually increasing pressure; the pressure of the stimulus was recorded automatically when the paw was withdrawn. The tests were repeated until three similar measurements (within 10 g) were obtained; the test-free period was 10 min. The average of three measurements was considered as the mechanical withdrawal threshold.

Pain scoring based on weight bearing. According to a previously described method,<sup>22</sup> the pain score was obtained based on the position of the operated paw: score = 0, the operated foot completely touched the mesh; score = 1, the wound touched the mesh and the foot distorted; and score = 2, the operated foot was kept completely off the mesh.

Measurement of thermal hyperalgesia. The rats were allowed to habituate in the test equipment for 30 min, after which thermal hyperalgesia was performed according to method of Hargreaves et al.<sup>25</sup> A radiant light was focused on the hindpaw, and the response latency was automatically recorded when the rat lifted or licked its hindpaw, the average of three repeated measurements was considered the withdrawal latency (5-min intervals between measurements). To avoid tissue damage, a 20-s cutoff time was established.

# Expression of Nav1.7 in the DRG

Quantitative PCR. Total RNA was extracted from the left L4-L5 DRG of the rats using Trizol (Invitrogen, Carlsbad, CA, USA) and 2 µg of total RNA was reverse-transcribed into cDNA using RevertAid Reverse Transcriptase (Thermo Scientific, Waltham, MA, USA) according to the manufacturer's instructions. Quantitative RT-PCR was performed using PCR system (ABI 7900/Viia, Foster City, CA, USA). SYBR Green was used to detect amplification of Nav1.7; reaction conditions were 95°C for 1 min (initial denaturation), followed by 40 cycles each consisting of 95°C for 30 s (denaturation) and at 60°C for 30 s (annealing extension). The primer sequences were as follows. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH): 5'-ACAGCAACAGGGTGGTGGAC-3' (forward) and 5'-TTTGAGGGTGCAGCGAACTT-3' (reverse), Na<sub>v</sub>1.7: 5'-CAGCCGCAGATAGCCG TCGT-3' (forward) and 5'-GGGCGTCCGCAAAGT CAGAG-3' (reverse). The data of Na<sub>v</sub>1.7 were normalized to the internal reference (GAPDH). The relative level of Nav1.7 mRNA was calculated using the comparative cycle threshold (CT) method  $(2^{-\Delta\Delta CT})$ . The samples were quantified in triplicate.

Western blotting. The DRGs were dissected from the left L4–L5 DRG of the rats and the total protein was RIPA lysis buffer extracted using (Beyotime Biotechnology, Jiangsu, China). The samples were separated by electrophoresis by 8% sodium dodecyl sulfatepolyacrylamide gel electrophoresis and then transfected to a 0.45-µm nitrocellulose membrane (Merck Millipore, Billerica, MA, USA), which were probed with primary antibodies (Nav1.7 rabbit polyclonal antibodies (1:200, Alomone Labs, Jerusalem, Israel) or GAPDH rabbit polyclonal antibody (1:1000, Abcam, Cambridge, MA, USA)) overnight at 4°C. Then, the nitrocellulose membranes were incubated with a secondary antibody for 2 h at room temperature. Positive signals were quantified using ImageJ software.

*Immunohistochemistry.* Under deep anesthesia with isoflurane, the rats were perfused intracardially with normal saline and subsequently fixed with 4% paraformaldehyde. Left L4–L5 DRGs were collected and postfixed in 4% paraformaldehyde for 4 h at room temperature. After dehydration, DRG sections (4  $\mu$ m) were prepared and incubated in 5% bovine serum albumin for 30 min at room temperature and then incubated with Na<sub>v</sub>1.7 polyclonal antibodies (1:200) overnight at 4°C, followed by secondary antibody (1:500, Life Technologies, Carlsbad, CA, USA) for 90 min at 37°C. Images were captured with a Leica microscope (Welzar, Germany).

#### Statistical analysis

All data were described as means  $\pm$  SD. Pain measurement data were analyzed by SPSS 19.0. Statistical analysis of Q-PCR, Western blotting, and immunostaining intensity was performed using GraphPad software. The statistically significant difference between groups was analyzed using one-way analysis of variance with Tukey's post hoc test. For all analyses, P < 0.05 was considered as statistically significant difference.

# Results

# Pain-related behavior in rats following plantar incision

Pain-related behaviors were measured before and at 2 h, 24 h, 48 h, 72 h, and 120 h after plantar incision. Compared with the control group, the mechanical and thermal pain threshold decreased significantly from 2 h to 72 h after plantar incision (P < 0.05) and showed a minimum value at 2 h after the operation (Figure 1(a) and (b)). Cumulative pain scores were increased significantly as compared with the control group (P < 0.05) and reached a maximum value at 2 h after the operation (Figure 1(c)). These findings indicated that the model of plantar incision-related postoperative pain was established successfully. Subsequently, incision-related pain behaviors started to return to the preoperative value; although the pain threshold remained lower than that in the control group, no statistically significant differences (P > 0.05) were found between the two groups at 120 h after the operation (Figure 1(a) to (c)).

# Dynamic expression of $Na_v I.7$ in the DRG after plantar incision

To analyze the expression of  $Na_v 1.7$  in the DRG during pain hypersensitivity after plantar incision, the expression of  $Na_v 1.7$  mRNA and  $Na_v 1.7$  proteins in L4–L6 DRG of rats was detected by Q-PCR and Western blotting, respectively, before and at 2 h, 24 h, 48 h, 72 h, and 120 h after plantar incision. The results revealed that both  $Na_v 1.7$  mRNA and  $Na_v 1.7$  proteins in L4–L6 DRG significantly increased from 2 h to 72 h postoperatively and reached a maximum at 2 h after plantar incision. Subsequently, the expression began to return to preincision levels (Figure 2(a) to (c) and Figure S1).

# $Na_v I.7$ expression in the DRG after plantar incision in rats pretreated with SCN9A-RNAi-LV

To detect whether *SCN9A*-RNAi-LV can knockdown  $Na_v1.7$  expression, mRNA of  $Na_v1.7$ ,  $Na_v1.8$ , and  $Na_v1.9$  were examined by Q-PCR. We found that *SCN9A*-RNAi-LV significantly reduced mRNA levels of  $Na_v1.7$  (72.6% as compared to the NC-LV group) (Figure 3(a)), while the mRNA levels of  $Na_v1.8$  and  $Na_v1.9$  were not affected (Figure 3(c) and (d)).  $Na_v1.7$  protein levels were also reduced significantly, by 60.8%, by this treatment (Figure 3(b)). These results showed that *SCN9A*-RNAi-LV down-regulated  $Na_v1.7$  expression efficiently and specifically.

As the expression of Na<sub>v</sub>1.7 in L4–L5 DRG peaked at 2 h after plantar incision, when the postoperative pain sensation was most significant, we investigated Na<sub>v</sub>1.7 expression in the DRG at 2 h after plantar incision in rats pretreated with *SCN9A*-RNAi-LV. As shown in Figure 4(d) and Figure S2, compared with the NC-LV pretreatment group (c), the enhanced Na<sub>v</sub>1.7 expression in L4–L6 DRG was inhibited in the *SCN9A*-RNAi-LV group (d) after plantar incision (b).

# Effect of SCN9A-RNAi-LV pretreatment in in incision-related pain behavior

Pretreatment with *SCN9A*-RNA-LV significantly inhibited the enhanced expression of Na<sub>v</sub>1.7 in L4–L6 DRG after plantar incision (Figure 5(a) to (c)). As compared with the NC-LV + incision group, the mechanical pain threshold was significantly increased by 108% (10.2  $\pm$  1.5 vs. 4.9  $\pm$  0.9), 88% (13.6  $\pm$  1.8 vs. 7.2  $\pm$  2.5), and



**Figure 1.** Pain-related behavior in rats following plantar incision. (a) Mechanical hypersensitivity. (b) Thermal hypersensitivity. (c) Cumulative pain score. \*P < 0.05 compared to the control group.



**Figure 2.** Dynamic expression of Na<sub>v</sub>1.7 in L4–L6 DRG of rats after plantar incision. (a) Quantitative analysis of Na<sub>v</sub>1.7 mRNA expression in the DRG after plantar incision. (b) The expression of Na<sub>v</sub>1.7 in the DRG as analyzed by Western blotting. (c) Quantitative analysis of Na<sub>v</sub>1.7 protein expression in the DRG after plantar incision. \*P < 0.05 compared with controls. GAPDH: glyceraldehyde-3-phosphate dehydrogenase.



**Figure 3.** The efficient and special of SCN9A-RNAi-LV. (a) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.7 mRNA in DRG. (b) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.7 protein in the DRG. (c) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.8 mRNA in the DRG. (d) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (e) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) The influence of SCN9A-RNAi-LV on Na<sub>v</sub>1.9 mRNA in the DRG. (f) Th

91% ( $14.3 \pm 1.6$  vs.  $7.5 \pm 0.7$ ) at 2 h, 24 h, and 48 h, respectively (Figure 5(a)). In the pretreated group, the thermal pain threshold was significantly increased by 46% ( $13.0 \pm 3.1$  vs.  $8.9 \pm 0.5$ ), 50% ( $15.9 \pm 5.3$  vs.  $10.6 \pm 3.2$ ), and 27% ( $15.7 \pm 2.3$  vs.  $12.4 \pm 3.6$ ) at 2 h, 24 h and 48 h, respectively (Figure 5(b)). The cumulative pain score in the pretreated group also significantly declined by 43% ( $12.5 \pm 0.6$  vs.  $22.0 \pm 2.2$ ), 56% ( $9.3 \pm 1.5$  vs.  $21.3 \pm 2.9$ ), 47% ( $6.8 \pm 1.0$  vs.  $12.8 \pm 1.7$ ) at 2 h, 24 h and 48 h, respectively (Figure 5(c)).

#### Discussion

The rat model of plantar incision-related pain is a reliable model for simulating postoperative pain in humans.<sup>22</sup> It can be used to facilitate understanding of the mechanism underlying postoperative pain and to explore new treatments for postoperative pain. In the present study, after plantar incision, the rats developed pain sensitization; the mechanical and thermal pain thresholds were significantly decreased and cumulative



**Figure 4.** Nav 1.7 expression in the DRG after plantar incision in rats pretreated with *SCN9A*-RNAi-LV. (a) The influence of *SCN9A*-RNAi-LV on Nav 1.7 mRNA in the DRG after plantar incision. (\*P < 0.05 compared with incision group.). (b) The influence of *SCN9A*-RNAi-LV on Nav 1.7 proteins in the DRG after plantar incision. (c) Quantitative analysis of Nav 1.7 protein expression in the DRG after plantar incision. (\*P < 0.05 compared with incision group.). (b) The influence of *SCN9A*-RNAi-LV on Nav 1.7 proteins in the DRG after plantar incision. (c) Quantitative analysis of Nav 1.7 protein expression in the DRG after plantar incision. (\*P < 0.05 compared with incision group.). (d) Representative immunohistochemical images showed the Nav 1.7 expression in DRG of naïve control group (a'), incision group (b'), NC-LV + incision group (c'), and SCN9A-RNAi-LV + incision group (d'). Nav 1.7-positive neurons exhibited a brown color. The naïve control group is the section of DRG from naïve rat, in which anti-Nav 1.7 antibodies were pre-incubated with Nav 1.7. (e) Quantification of Nav 1.7 labelling intensity in normalized to naïve control values. The increased expression of Nav 1.7 in L4–L6 DRG after plantar incision was inhibited by *SCN9A*-RNAi-LV. \*P < 0.05 compared with incision group. GAPDH: glyceraldehyde-3-phosphate dehydrogenase; NC-LV: negative control lentivirus.

pain scores were significantly increased (from 2 h to 72 h after plantar incision). The mechanical and thermal pain thresholds were lowest and the cumulative pain scores were highest at 2 h after the operation and then subsequently recovered gradually. Na<sub>v</sub>1.7 expression in L4–L6 DRG increased significantly and reached a peak at 2 h after plantar incision and then recovered from 72 h after plantar incision. However, when the rats were

pretreated with SCN9A-RNAi-LV to inhibit the enhancement of Na<sub>v</sub>1.7 expression in the DRG, the pain hypersensitivity after plantar incision was alleviated.

Voltage-gated sodium channel  $Na_v 1.7$  plays an important role in the generation and transmission of pain signals.<sup>26</sup> Electrophysiological studies confirmed that  $Na_v 1.7$  is able to amplify sub-threshold excitability



**Figure 5.** The effect of SCN9A-RNAi-LV pretreatment in incision-related pain behavior. (a) Mechanical hypersensitivity. (b) Pain score. (c) Thermal hypersensitivity. \*P < 0.05 compared with the incision group. NC-LV: negative control lentivirus.

and regulate neuronal excitability.<sup>4,27</sup> The increased  $Na_v 1.7$  expression in DRG suggests hyperexcitability of DRG neurons and significant stimulate synaptic transmission following plantar incision, which may contribute to hyperalgesia after plantar incision. These findings demonstrate that the increased  $Na_v 1.7$  in the DRG is very important for postoperative pain hypersensitivity after plantar incision.

This is in agreement with greatly increased Nav1.7 expression and pain hypersensitivity in preclinical studies from rodent model of inflammatory pain and peripheral neuropathic pain.<sup>6,14,28,29</sup> A previous study has shown that the expression of Nav1.7 is regulated by inflammatory mediators, such as NGF and prostaglandin  $E_2$  via the protein kinase B/Akt pathway.<sup>30</sup> Pronociceptive factors (e.g., NGF) are increased after plantar incision.<sup>31</sup> These findings dovetail well with previous results and show that NGF signaling may be important in the up-regulation of Nav1.7 in postoperative pain.

In addition, in clinical context, postoperative pain is categorized as at-rest pain and movement-induced pain. In previous studies, the mechanical and the thermal pain thresholds were quantified by the von Frey filaments method and thermal radiation, respectively, these pain sensations are similar to pain evoked by coughing or movement. However, in this study, we also used a cumulative pain score, which reflects a kind of nonevoked pain and is based on the position chosen by the rats after the incision, and which is similar to surgical limb protection behavior of patients, to assist in the analysis of mechanical sensitivity.<sup>16,19</sup>

In summary, this study confirms that  $Na_v 1.7$  in DRG plays an important role in a preclinical rodent model of acute postoperative pain and suggests that  $Na_v 1.7$ -targeted analgesics currently under development may be effective in patients with acute postoperative pain.

#### **Author Contributions**

Xianwei Zhang conceived the study and supervised project, Xianwei Zhang and Jiaoli Sun designed the experiment, analyzed the data, and wrote the manuscript. Ningbo Li and Guangyou Duan were responsible for establishment of incisional pain in rats. Yi Liu and Shanna Guo conducted the measurement of pain behaviors. Cong Wang and Changmao Zhu carried out Q-PCR, Western blotting, and immunohistochemistry analysis. All authors approved the version to be published.

#### **Declaration of Conflicting Interests**

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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#### **Supplemental Material**

Supplementary material is available for this article online.

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