

Supporting Information

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SI Methods

Tissue Fixation. Adult rats, mice, and *Oprd1* exon 1-deleted mice (Jackson Lab) were anesthetized and perfused with 20 mL of warm saline, followed by a warm mixture of 4% paraformaldehyde and 0.1% picric acid in 0.1 M phosphate buffer (pH 7.4) and then the same but ice-cold fixative (50 mL). The lumbar segments of the spinal cord and DRGs were dissected out and postfixed in the same fixative for 90 min at 4 °C.

Gene Transfection in DRG Neurons. Dissociated DRG neurons were transfected with pCMV-Myc-DOR1 or DOR1-EGFP in pEGFP-N3 by electroporation with Nucleofector II (Amaxa) using Neuron Nucleofector Solution and program O-003 and were then plated on coverslips in DMEM containing 10% (vol/vol) FBS. After 6 h, neurons were cultured in the medium containing 50% (vol/vol) DMEM, 50% (vol/vol) F12, 1% N2, and 100 ng/mL NGF for small neurons or 50 ng/mL NT3 for large neurons for 48–72 h.

Whole-Cell Patch-Clamp Recording. The composition of the pipette solution was 120 mM CsCl, 1 mM MgCl₂, 10 mM Hepes, 10 mM EGTA, 4 mM Mg-ATP, and 0.3 mM Na-GTP (pH 7.2, 300 mOsm). Cells were kept in Tyrode's solution containing 135 mM NaCl, 2.5 mM KCl, 4 mM CaCl₂, 2 mM MgCl₂, 10 mM D-glucose, and 10 mM Hepes (pH 7.4, 300 mOsm) until the whole-cell configuration was achieved. The external solution was then exchanged with the Ba²⁺-containing solution, referred to as Ba-solution, containing 5 mM BaCl₂, 150 mM N-methyl-D-glucamine, 2 mM MgCl₂, 10 mM D-glucose, and 10 mM Hepes (pH 7.4, 300 mOsm), while holding the cell at -70 mV. The drug stock solutions were diluted in Ba-solution before use. Effects of agonists on barium currents flowing through Ca²⁺ channels were measured on current waveforms activated by 100- or 200-ms command pulses to 0 mV from a holding potential of -70 mV and recorded with a MutiClamp-700B amplifier (Molecular Devices). The data were quantified off-line using custom-made software by measuring the peak current amplitude of the depolarizing test pulses.

Whole-Cell Recording in Spinal Cord Slice. Transverse slices (~600 μm) of L4 and L5 spinal cord segments with an attached dorsal root from adult rats were prepared with a vibrating microslicer and perfused in oxygen-bubbled Krebs' solution (117 mM NaCl, 3.6 mM KCl, 2.5 mM CaCl₂, 1.2 mM MgCl₂, 1.2 mM NaH₂PO₄, 25 mM NaHCO₃, and 11 mM D-glucose) for blind whole-cell recording (1). Resistance of patch pipettes was typically 4–10 MΩ. The pipette solution contained 135 mM K-gluconate, 0.5 mM CaCl₂, 2 mM MgCl₂, 5 mM KCl, 5 mM EGTA, 5 mM Hepes, and 5 mM D-glucose. Currents were filtered at 2 kHz and digitized at 5 kHz using the Axopatch 200B (Molecular Devices) and were analyzed by pCLAMP8.5 (Molecular Devices) at a holding potential of -70 mV. Amplitude of mEPSCs was analyzed with an Axograph (Molecular Devices). Dorsal root stimulation, which was sufficient to recruit C-fibers, was delivered with a suction electrode linked to a constant-current stimulator (Electronic Stimulator; Nihon Kohpen). Afferent C-fibers were identified by their estimated evoked EPSC latency at 0.9 ± 0.5 m/s (range: 0.3–1.5 m/s).

Electric Stimulation, Cell Surface Biotinylation, and Immunoblotting. Neurons from L4 and L5 DRGs were cultured in the medium containing an equal volume of DMEM and F12 medium with 1% N2 for 3 d. Cells were stimulated by 10-Hz electric stimulation with platinum wires contacting the culture medium. Each stimulation pulse (0.5 ms, 8–10 V) was sufficient to elicit action potential. Cells were stimulated for 5 min. Surface protein biotinylation was performed after 5-min electrical stimulation by incubating cultured neurons with Sulfo-NHS-LC-Biotin (0.5 mg/mL; Pierce) for 60 min at 4 °C. Cells were then processed for precipitation of biotinylated proteins with ImmunoPure Immobilized Streptavidin (Pierce) and immunoblotting with DOR antibodies (1:3,000; Neuromics) and actin (1:10,000; Santa Cruz). The intensity of DOR bands was normalized to actin.

1. Nakatsuka T, Park JS, Kumamoto E, Tamaki T, Yoshimura M (1999) Plastic changes in sensory inputs to rat substantia gelatinosa neurons following peripheral inflammation. *Pain* 82:39–47.

Table S1. Primers for PCR experiments and probes for in situ hybridization

No.	Gene	Primer name	Primer sequence	Product length, bp
1	<i>OPRD1</i> (NM_013622)	OPRD1-F	5'-GTGCAAGGCTGTGCTCTCCATTG-3'	770
		OPRD1-R	5'-GTCGGGTAGGTCAGGCGGCAGCGCCACCG-3'	
2	<i>OPRM1</i> (NM_001039652)	OPRM1-F	5'-ATGGACAGCAGCGCCGGCCC-3'	1,317
		OPRM1-R	5'-TCACCTGCCAAGCTGGCCTTC-3'	
3	<i>PPT-A</i> (NM_009311)	PPT-A-F	5'-ATGAAAATCCTCGTGG-3'	393
		PPT-A-R	5'-TTATTTACGTCTTCTTC-3'	
4	<i>GAPDH</i> (NM_008084)	GAPDH-F	5'-TGCTGAGTATGTCGTGGAG-3'	555
		GAPDH-R	5'-TAGCCCAAGATGCCCTTCAGT-3'	
5	<i>OPRD1</i> (NM_013622)	OPRD1-F	5'-ATGGAGCTGGTCCCTCTG-3'	
		OPRD1-F	5'-TCCAGAGCGCCAAGTACTTG-3'	
		OPRD1-R	5'-TCAGGCGGCAGCGCCACCG-3'	
6	<i>OPRD1</i> (NM_013622)	OPRD1-F	5'-GTGCAAGGCTGTGCTCTCCATTG-3'	770
		OPRD1-R	5'-GTCGGGTAGGTCAGGCGGCAGCGCCACCG-3'	
7	<i>OPRD1</i> (NM012617)	OPRD1-F	5'-GTGCAAGGCTGTGCTCTCCATTG-3'	770
		OPRD1-R	5'-GCCGGGTAGGTCAGGCGGCAGCGCCACCG-3'	
8	<i>OPRD1</i> (NM_013622)	OPRD1-F1	5'-TTCTGGGCAACGTGCTCGTC-3'	510
		OPRD1-R1	5'-CATAGCACACCGTGATGATG-3'	
		OPRD1-F2	5'-TGTTGGCATCGTCCGGTAC-3'	
9	<i>OPRM1</i> (NM_001039652)	OPRD1-R2	5'-TGAAGCCAAGACCCAGATGC-3'	320
		OPRM1-F1	5'-GTATCTTCACCCCTCTGCACC-3'	
		OPRM1-R1	5'-AGGCAATGCAGAAGTGCCAG-3'	
10	<i>PPT-A</i> (NM_009311)	OPRM1-F2	5'-AGGCCCTGGATTTCCGTACC-3'	272
		OPRM1-R2	5'-CATGCGGACACTCTTGAGTC-3'	
		PPT-A-F1	5'-ATGAAAATCCTCGTGGCCGT-3'	
11	β -Actin (X03672)	PPT-A-R1	5'-CTTTCGTAGTTCTGCATCGC-3'	380
		PPT-A-F2	5'-CCACTCAACTGTTTGACAGAG-3'	
		PPT-A-R2	5'-GCTCTTTTGCCCATAGTCC-3'	
11	β -Actin (X03672)	β -Actin-F1	5'-GCCAACCGTGAAAAGATGAC-3'	556
		β -Actin-R1	5'-GCACTGTGTTGGCATAGAGG-3'	
		β -Actin-F2	5'-GGCTGTGCTGTCCCTGTATG-3'	
		β -Actin-R2	5'-GGCTGTGCTGTCCCTGTATG-3'	

Nucleotide sequence of PCR primers is listed with reference to corresponding GenBank accession numbers. Primers 1–4 were used in PCR of three subsets of DRG neurons, primer 5 was used in PCR of *Oprd1* exon 1-deleted mice, primers 6 and 7 were used for probes in situ hybridization, and primers 8–11 were used in single-cell PCR. F: forward; R: reverse. F1 and R1 and F2 and R2 primer pairs were used in the first- and second-round single-cell PCR assays. For each primer pair, forward and reverse primers are located in different exons to avoid the amplification of genomic DNA.