Nav1.1 in mammalian sensory neurons is required for normal motor behaviors

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6 Cyrrus M. Espino¹⁺, Cheyanne M. Lewis¹⁺, Serena Ortiz², Miloni S. Dalal³, Kaylee M. Wells⁴,
 7 Darik A. O'Neil⁴, Katherine A. Wilkinson², and Theanne N. Griffith^{1,4*}

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¹Department of Physiology and Membrane Biology, University of California Davis, Davis, CA,
 95616

¹² ²Department of Biological Science, San José State University, San José, CA, 95192

¹³ ³Department of Pharmacology, Physiology, and Neuroscience, New Jersey Medical School-

14 Rutgers University, Newark, NJ, 07103

⁴Neurobiology course, Marine Biological Laboratory Woods Hole, MA, 02540

1617 *Corresponding Author: Theanne N. Griffith

18	 1275 Med Science Drive
19	Tupper Hall 4135
20	Davis, CA 95616
21	530.754.2780
22	tgriffith@ucdavis.edu

²⁴ [†]These authors contributed equally to this work.

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36 Abstract

37 The mammalian voltage-gated sodium channel (Na_V), Na_V1.1, has been well-studied in the central 38 nervous system; conversely, its contribution to peripheral sensory neuron function is more 39 enigmatic. Here, we report a new role for peripherally expressed Nav1.1 in murine motor 40 behaviors. RNAscope analysis found 100% of proprioceptors express Nav1.1 transcript, 41 consistent with in vitro patch clamp recordings showing this channel is required for repetitive firing 42 in proprioceptors. Notably, genetic deletion of Nav1.1 in all sensory neurons caused profound 43 motor coordination deficits in homozygous conditional knockout animals of both sexes, a 44 phenotype similar to conditional Piezo2-knockout animals. Movement deficits were also observed 45 in heterozygotes, demonstrating that Nav1.1 haploinsufficiency in sensory neurons leads to motor 46 deficiencies. This behavioral phenotype was not due to reduced proprioceptor numbers or 47 abnormal muscle spindle formation; however, we observed decreased proprioceptor innervation 48 of motor neurons in the spinal cord in conditional knockouts, indicating loss of Nav1.1 in sensory 49 neurons alters spinal cord circuitry. Ex vivo muscle afferent recordings also support the notion 50 that loss of Nav1.1 leads to aberrant proprioceptor function. Collectively, these data provide the 51 first evidence that Nav1.1 in mammalian sensory neurons is essential for motor coordination. 52 Importantly, human patients harboring Nav1.1 loss-of-function mutations often present with motor 53 delays and ataxia. Thus, our data suggest sensory neuron dysfunction may contribute to the 54 clinical manifestations and co-morbidities of neurological disorders in which Nav1.1 function is 55 compromised.

56

57 INTRODUCTION

58 Voltage gated sodium channels (Navs) are critical mediators of neuronal excitability and are 59 responsible for action potential generation and propagation (Ahern et al., 2016; Bean, 2007). In 60 the mammalian nervous system there are nine isoforms (Nav1.1-1.9), each with unique 61 biophysical properties, as well as distinguishing cellular expression and subcellular localization 62 patterns (Bennett et al., 2019; Catterall, 2017). Of these different subtypes, Nav1.1 is notable for 63 its role in brain disease (Escavo and Goldin, 2010; Mullev et al., 2005; Ogiwara et al., 2007). 64 Indeed, Scn1a, the gene that encodes Nav1.1, is referred to as a "super culprit" gene, with over 65 1,000 associated mutations that lead to abnormal brain function, resulting in brain disorders such 66 as epilepsy and migraine, as well as neurodivergent phenotypes, such as autism spectrum disorder (Ding et al., 2021; Lossin, 2009). Homozygous Nav1.1^{-/-} global knockout mice are ataxic 67 and die by P15, while heterozygous Nav1.1^{+/-} animals develop seizures and begin to die 68 69 sporadically starting at P21 (Yu et al., 2006). In addition to the central nervous system, Na $_{\rm V}$ 1.1 is 70 also expressed in the peripheral nervous system (Sharma et al., 2020; Usoskin et al., 2015); yet, 71 the prominent role this channel plays in brain function has left its physiological roles in sensory 72 neuron populations understudied.

73 Peripheral sensory neurons of the dorsal root and trigeminal ganglia (DRG and TG, 74 respectively) are tasked with encoding somatic sensations, such as touch, temperature, pain, and 75 proprioception, and are anatomically and functionally heterogenous (Kupari et al., 2021; Nguyen 76 et al., 2021; Oliver et al., 2021; Wu et al., 2021). Nav1.1 transcript and protein have been observed 77 primarily in myelinated mechanosensory DRG and TG neurons (Fukuoka et al., 2008; Ho and 78 O'Leary, 2011; Osteen et al., 2016). Indeed, subcutaneous injection of the Nav1.1 activator, 79 Hma1, into mouse hind paw causes non-inflammatory mechanical pain and spontaneous pain 80 behaviors (Osteen et al., 2016). Interestingly, pharmacological inhibition of Nav1.1 does not affect 81 mechanical thresholds in uninjured mice but does reduce mechanical pain in a spared-nerve

82 injury model (Salvatierra et al., 2018), suggesting Na $_{\rm V}$ 1.1 may have a more prominent role in 83 mechanical pain as opposed to normal touch sensing. Nav1.1 in TG neurons has also been 84 reported to mediate mechanical pain in an orbitofacial chronic constriction injury model (Pineda-85 Farias et al., 2021). In addition to somatosensory neurons, Nav1.1 is found in colon-innervating 86 vagal neurons, where it contributes to firing of colonic mechano-nociceptors and is upregulated 87 in a mouse model of chronic visceral hypersensitivity (Osteen et al., 2016; Salvatierra et al., 2018). 88 Lastly, Nav1.1 contributes to action potential firing in a subset of DRG neurons that express the 89 cold sensitive ion channel, transient receptor potential melastatin 8 (TRPM8), suggesting the 90 channel may also contribute to thermosensory transmission (Griffith et al., 2019). While most data 91 support a role for Na_V1.1 in pain, the limited number of studies that have investigated Na_V1.1 92 function in sensory neurons has left gaps in our knowledge regarding other potential roles this 93 channel may play in somatosensation.

94 Given the relatively under-explored role of $Na_V 1.1$ in the peripheral nervous system, we 95 set out to determine what other somatosensory modalities rely on Nav1.1 expression in sensory 96 neurons. Here, we show that 100% of proprioceptors express Nav1.1 mRNA. A functional role for 97 the channel in these cells was supported by both in vitro and ex-vivo electrophysiological 98 recordings from genetically identified proprioceptors and functionally identified muscle spindle 99 afferents, respectively. Importantly, mice lacking Nav1.1 in all sensory neurons display visible and 100 profound motor deficits and ataxic-like behavior, which were quantified in rotarod and open field 101 assays. These deficits may be due to a combination of aberrant electrical signaling and altered 102 spinal cord circuitry. Collectively, our data provide evidence for a new role for peripherally 103 expressed Na_V1.1 in proprioceptor function and motor coordination.

104

105 MATERIALS AND METHODS

106 *Key resources*. Table 1 contains a list of key resources and supplies used for this study.

Reagent or Resource	Source	Identifier
Deposited Data		
Source data	This paper	doi: 10.17632/kt23th75v9.2
Matlab scripts	This paper	https://github.com/doctheagrif/Curre nt-Clamp-Matlab-Code_O-Neil-DA
Antibodies		· –
Rabbit polyclonal anti-DsRed	Takara Bio	Catalogue #632496
Chicken polyclonal GFP	Abcam	Catalogue #ab13970
Chicken polyclonal NFH	Abcam	Catalogue #ab4680
Rabbit polyclonal CGRP	Immunostar	Catalogue #24112
Guinea pig polyclonal VGLUT1	Thomas Jessell Laboratory; HHMI Columbia University	Catalogue #CU1706, RRID:AB_2665455
Chicken polyclonal β 3-tubulin	Abcam	Catalogue #ab41489
Rabbit polyclonal β3-tubulin	Abcam	Catalogue #ab18207
Chemicals		
VECTASHIELD® Antifade Mounting Media with DAPI	Vector Laboratories	Catalogue #H-2000
Tissue-Tek OCT compound	Sakura	Catalogue #4583
Fluromount-G DAPI	SouthernBiotech	Catalogue #0100-20
Laminin	Sigma-Aldrich	Catalogue #L2020-1MG
Collagenase type P	Sigma-Aldrich	Catalogue #11213865001
TrypLE Express	ThermoFisher	Catalogue #12605-010
MEM	ThermoFisher	Catalogue #11095-080
Penicillin-streptomycin	ThermoFisher	Catalogue #15140-122
MEM vitamin solution	ThermoFisher	Catalogue #11120-052
B-27 supplement	ThermoFisher	Catalogue #17504-044
Horse serum, heat inactivated	ThermoFisher	Catalogue #26050-070
ICA 121431	Tocris Bioscience	Catalogue #5066/10
Critical Commercial Assays		
RNAscope Fluorescence Multiplex Kit	Advanced Cell Diagnostics	Catalogue #320851
Oligonucleotides		
Pvalb probe channel 1	Advanced Cell Diagnostics	Catalogue #421931
Scn1a probe channel 2	Advanced Cell Diagnostics	Catalogue #556181-C2
Runx3 probe channel 3	Advanced Cell Diagnostics	Catalogue #451271-C3
Experimental Models: Mouse Strain	IS	
Pirt ^{cre} Dr. Xinzhong Dong		
Rosa26 ^{Ai14}	Jackson Laboratories	Stock #007914

PVcre Na _V 1.1 ^{fl/fl} MrgprD ^{GFP}	Jackson Laboratories UC Davis MMRRC Zheng et al., 2019	Stock # 008069 Stock # 041829-UCD
Software and Algorithms		
pClamp 11.2 Software Suite	Molecular Devices	https://www.moleculardevices.com
Image J	Schneider et al. (2012)	https://imagej.nih.gov
Prism 9	Graphpad	https://www.graphpad.com
LabChart	ADInstruments	https://www.adinstruments.com/pro ducts/labchart
MATLAB	MathWorks	https://www.mathworks.com
Software, algorithm, custom (MATLAB)	This study	https://github.com/doctheagrif/Curre nt-Clamp-Matlab-Code O-Neil-DA



108 Animals. Pirt^{cre} mice were a kind gift from Dr. Xinzhong Dong (Johns Hopkins University). Rosa26^{Ai14} (stock #007914; (Madisen et al., 2010)) and PV^{cre} (stock # 008069) were obtained from 109 110 Jackson Laboratories. Nav1.1^{fl/fl} (stock # 041829-UCD) mice were purchased from the UC Davis 111 MMRRC. MrgprD mice were originally published in Zheng et al., 2019. Genotyping was 112 outsourced to Transnetyx. Animal use was conducted according to guidelines from the National 113 Institutes of Health's Guide for the Care and Use of Laboratory Animals and was approved by the 114 Institutional Animal Care and Use Committee of Rutgers University-Newark, UC Davis, and San 115 José State University (ex vivo muscle recordings). Mice were maintained on a 12 h light/dark 116 cycle, and food and water was provided ad libitum.

117

Rotarod. To assess motor coordination, a rotarod machine (IITC Life Sciences, Woodland Hills, CA) that has an accelerating rotating cylinder was used. Mice were acclimated to the behavior room for 2 h prior to testing. Mice were assayed on the rotarod for 3 consecutive days, with 3 trials per day and an intertrial interval of at least 15 min. The average of the three trials per day was used. The experimenter was blind to genotype.

123

Open field test. Mice were acclimated to the behavior room for 2 h prior to testing. The open field
 apparatus consisted of a black square sound attenuating box of dimensions 40.6 cm × 40.6 cm.

A camera suspended above the arena was connected to a computer running Ethovision XT software, which tracked animal movement and velocity. An animal was placed in the center of the arena and allowed to freely explore for a 10 min trial. The experimenter was blind to genotype.

130 *Tissue processing.* For spinal cord immunolabeling experiments, whole spinal columns from adult Pirt^{Cre}:Rosa26^{Ai14} and Pirt^{Cre}:Nav1.1^{fl/fl} animals were harvested on ice. For Tdtomato IHC. 131 132 spinal columns were fixed overnight at 4°C in 4% paraformaldehyde. For vesicular glutamate 133 transporter 1 (VGLUT1) and choline acetyltransferase (ChAT) co-labeling experiments, spinal 134 columns were fixed in 4% paraformaldehyde for 1 h on ice. Tissue was then placed in 30% 135 sucrose solution overnight at 4°C. Following cryoprotection, tissue was embedded in optimal 136 cutting temperature compound (OCT, Tissue-Tek® Sakura) and stored at -80°C until sectioning. 137 DRG were harvested from thoracic spinal levels and fixed in 4% formaldehyde for 15 min at 4°C 138 and were then incubated in 30% sucrose for 2-4 h at 4°C. DRG were embedded in OCT and 139 stored at -80°C until sectioning.

140

141 *Immunohistochemistry*. Immunohistochemistry of spinal cord cryostat sections (30µm) was 142 performed using the following primary antibodies: Rabbit anti-DsRed (1:3000, Takara Bio, 143 632496), guinea pig anti-VGLUT1 (1:8000, Zuckerman Institute, 1705), and rabbit anti-ChAT 144 (1:10,000, Zuckerman Institute, 1574). Secondary antibodies used were as follows: anti-rabbit 145 594 (1:1000, ThermoFisher, A32740), anti-guinea pig 488 (1:1000, ThermoFisher, A11073), and 146 anti-chicken 647 (ThermoFisher, A32733). Specimens were mounted with Fluoromount-G with 147 DAPI (SouthernBiotech, 0100-20). EDL muscles used in ex vivo muscle afferent recordings were 148 placed in ice cold 4% paraformaldehyde for 1 h followed by ice cold methanol for 15 min. Muscles 149 were incubated in blocking solution (0.3% PBS-T and 1% BSA) followed by incubation in primary 150 antibodies (guinea pig anti-VGLUT1 1:800 and chicken anti-NFH 1:300, ThermoFisher ab4680)

for 3-6 days at 4°C. After primary antibody treatment, tissue was washed in blocking solution and
treated with secondary antibody (anti-guinea pig 488 1:50 and anti-chicken 594 1:300, Invitrogen,
WA316328) for 2-3 days. Specimens were mounted with VECTASHIELD® with DAPI (H-2000,
Vector Laboratories). All specimens were imaged in three dimensions on a on a Zeiss LSM880
Airyscan confocal microscope. Images were analyzed using ImageJ software.

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157 *Multiplex* in situ *hybridization*. Fixed-frozen DRG tissue was cut in 25µm sections and placed 158 on electrostatically coated slides. Sections were processed for RNA in situ detection using a 159 modified version of the manufacturer's protocol ((Griffith et al., 2019), Advanced Cell Diagnostics) 160 and the following probes: Pvalb (421931- C1, mouse), Runx3 (451271-C3, mouse), Scn1a 161 (556181-C2, mouse). Following *in situ* hybridization, sections were incubated in blocking solution 162 (5% normal goat serum, 0.1% PBS-T) for 1 h at RT. Tissue was then incubated in primary 163 antibodies overnight at 4°C. The following antibodies were used: rabbit DsRed (1:3000, Takara 164 Bio, 632496), rabbit β 3-tubulin (1:3000, Abcam, ab18207), chicken β 3-tubulin (1:500, Abcam, 165 ab41489), rabbit CGRP (1:1000, Immunostar, 24112), chicken GFP (1:3000, Abcam, ab13970), 166 and chicken NFH (1:3000, Abcam, ab4680). Tissue was treated with the following secondary 167 antibodies for 45 min at RT: anti-rabbit 448 (1:1000, Invitrogen, A32731), 594 (1:1000, Invitrogen, 168 A11037) and 647 (1:1000, Invitrogen, A32733), anti-chicken 488 (1:1000, Invitrogen, A32931) 169 and 594 (1:1000, Invitrogen, A32740). Sections were washed and mounted with Fluoromount-G 170 with DAPI and imaged in three dimensions (2μ m axial steps) on an Olympus confocal (LV3000) 171 using a 40X 0.90 NA water objective lens. Images were auto-thresholded and analyzed using 172 ImageJ software.

173

174 **DRG culture preparation**. DRG were harvested from thoracic spinal levels of adult 175 PV^{cre};Rosa26^{Ai14} (6-16 weeks) mice of both sexes and transferred to Ca²⁺-free and Mg²⁺-free

176 HBSS solution (Invitrogen, 14170-112). Upon isolation, processes were trimmed, and ganglia 177 were transferred into collagenase (1.5 mg/mL; Type P, Sigma-Aldrich, 11213865001) in HBSS 178 for 20 min at 37°C followed by TrypLE Express (ThermoFisher, 12605-010) for 3 min with gentle 179 rotation. TrypLE was neutralized with 10% horse serum (heat-inactivated; Invitrogen, 26050-070) 180 and supplemented with culture media (MEM with L-glutamine, Phenol Red, without sodium 181 11095-080), containing 10,000 U/mL Penicillin-streptomycin pyruvate, ThermoFisher, 182 (ThermoFisher, 15140-122), MEM Vitamin Solution (Invitrogen, 11120-052), and B-27 183 supplement (ThermoFisher, 17504-044). Serum containing media was decanted and cells were 184 triturated using a fire-polished Pasteur pipette in the MEM culture media described above. Cells 185 were resuspended and triturated using a plastic pipette tip. Cells were plated on glass coverslips 186 that had been washed in 2M NaOH for at least 4 h, rinsed with 70% ethanol, UV-sterilized, and 187 treated with laminin (0.05 mg/mL, Sigma-Aldrich, L2020-1MG) for 1 hour prior to plating. Cells 188 were then incubated at 37°C in 5% CO₂. Cells were used for electrophysiology experiments 14-189 36 h post-plating.

190

191 In Vitro *Electrophysiology*. Whole-cell voltage-clamp recordings were made from TdTomato-192 expressing dissociated DRG neurons using patch pipettes pulled from Model P-1000 (Sutter 193 Instruments). Patch pipettes had a resistance of $3-6M\Omega$ when filled with an internal solution 194 containing the following (in mM): 140 CsF, 10 NaCl, 1.1 EGTA, .1 CaCl₂, 10 HEPES, and 2.5 195 MgATP, pH with CsOH to 7.2. Seals and whole-cell configuration were obtained in an external 196 solution containing the following (in mM): 145 NaCl, 5 KCl, 10 HEPES, 10 Glucose, 2 CaCl₂, 2 197 MgCl₂, pH 7.3 with NaOH, osmolarity ~320mOsm. Series resistance was compensated by 70-198 80%. To isolate whole-cell sodium currents during voltage clamp experiments, a modified external 199 solution was applied containing the following solution (in mM): 50 NaCl, 95 TEA-Cl, 10 HEPES, 200 2 BaCl₂, 13 glucose, 0.03 CdCl₂, pH 7.3 with NaOH, Osmolarity ~320mOsm. Voltage clamp

recordings were performed at room temperature and current clamp recordings were conducted
 at 37°C. Bath temperature was controlled and monitored using CL-100 (Warner Instruments).

203

204 **Ex vivo** electrophysiology. The effect of the loss of Nav1.1 on muscle spindle afferent firing 205 rates during muscle stretch and sinusoidal vibration was determined using an isolated muscle 206 nerve preparation. The extensor digitorum longus muscle and innervating peroneal branch of the 207 sciatic nerve were dissected from adult (2-4 month old) mice of both sexes. Muscles were held at 208 optimal length (Lo), or the length of the muscle that maximal force of twitch contraction occurred. 209 A series of nine 4 s ramp-and-hold stretches were given to 3 different stretch lengths repeated 3 210 times each (2.5%, 5%, and 7.5% Lo; ramp speed 40% Lo/s). A series of twelve 9 s sinusoidal 211 vibrations were given (25, 50, and 100 µm amplitude; 10, 25, 50, and 100 Hz frequency). A one-212 minute rest was given between each length change. Firing rates during a 10 s baseline before 213 stretch (resting discharge or RD) and the maximal firing rate during the ramp up phase of stretch 214 (dynamic peak or DP) were calculated for all animals. We determined whether the response to 215 static stretch was maintained consistently throughout the 4s stretch, as well as the coefficient of 216 variability of the interspike interval (ISI) during the plateau phase of stretch (CV = Std Dev/Mean 217 of ISI over the time period of 1.5-3.5s after end of ramp up). Average firing rate during the 9 s of 218 vibration and whether the unit could entrain in a 1:1 fashion to vibration was also determined. 219 Detailed methods can be found in Wilkinson et al., 2012.

220

221 **Data acquisition and analysis**. Currents and voltages were acquired using pClamp software 222 v11.2 (Molecular Devices). Recordings were obtained using an AxoPatch 200b patch-clamp 223 amplifier and a Digidata 1550B and filtered at 5 kHz and digitized at 10 kHz. Voltage-clamp and 224 current-clamp experiments were analyzed with Clampfit software v11.2 (Molecular Devices) and 225 custom MatLab Scripts. *Ex vivo* recordings were obtained using an A-M Systems Model 1800

extracellular amplifier with headstage and digitized using an ADInstruments PowerLab. Data was
analyzed using ADInstruments LabChart software using the Spike Histogram function.

228

Pharmacology. ICA 121431 was purchased from Tocris Bioscience. All other chemicals were
 from Sigma-Aldrich and Fisher Chemical.

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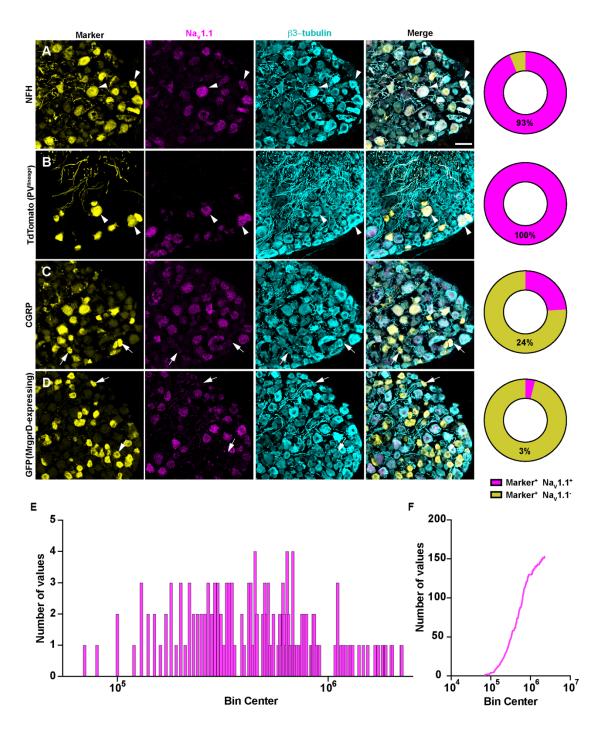
232 *Experimental design and statistical analysis.* Summary data are presented as mean ± SD from 233 *n* cells or afferents, or *N* animals. For quantitative analysis of *in situ* hybridization data, at least 3 234 biological replicates per condition were used and the investigator was blinded to genotype for 235 analysis. Statistical differences was determine using parametric tests for normally distributed data 236 and non-parametric tests for data that did not conform to Gaussian distributions or had different 237 variances. Statistical tests are listed in *Results* and/or figure legends. Statistical significance in 238 each case is denoted as follows: *p < 0.05, **p < 0.01, ***p < 0.001, and ****p < 0.0001. Statistical 239 tests and curve fits were performed using Prism 9.0 (GraphPad Software).

240

241 **Results**

242 Most studies have found Na_V1.1 expression primarily in myelinated sensory neurons that 243 transmit mechanical signals (Fukuoka et al., 2008; Ho and O'Leary, 2011; Wang et al., 2011). In 244 line with prior work, RNAscope analysis of DRG sections from adult mice show that 93% of 245 myelinated neurons, as determined by neurofilament heavy chain (NFH) labeling, expressed 246 Na_V1.1 transcripts (Fig 1A). RNA-sequencing datasets have consistently identified Na_V1.1 247 expression in proprioceptors; thus, we next analyzed Nav1.1 expression in genetically identified 248 proprioceptors and found 100% of those neurons were positive for Na_V1.1 message (**Fig 1B**), 249 consistent with the observation that $Na_V 1.1$ protein is expressed in muscle spindle afferents 250 (Carrasco et al., 2017), which innervate proprioceptive end organs found in skeletal muscle. This

contrasted with low expression of Na_V1.1 in both calcitonin gene related peptide (CGRP) expressing neurons, which represent peptidergic nociceptors, and non-peptidergic polymodal MrgprD-expressing nociceptors (24% and 3%, respectively, **Fig 1C-D**). The coefficient of variation for the integrated fluorescence density of Na_V1.1 transcripts from RNAscope experiments in proprioceptors was 76.5 (**Fig 1E-F**).



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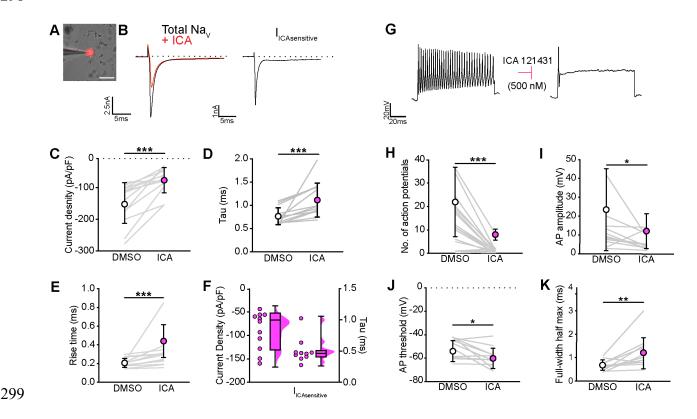
Figure 1. Na_v1.1 is ubiquitously expressed in genetically identified proprioceptors. A-D, Representative confocal images of cryoprotected adult DRG sections (25 μm) with quantifications indicating the percentage of Na_v1.1+ and Na_v1.1- neurons in each subpopulation. Images were acquired with a 40X, 0.9 NA water-immersion objective. Sections were hybridized using RNAscope with probes targeting Na_v1.1 (*Scn1a*, magenta) and stained with the following antibodies (yellow): (**A**) anti-neurofilament heavy (NFH, n=787) (**B**) anti-DsRed (TdTomato, n=143) (**C**) anti-calcitonin gene-related peptide (CGRP, n=877) (and (**D**) anti-GFP to label MrgprD+ neurons. C57BL/6, Parvalbumin^{Cre:}Rosa26^{Ai14}, and Mrgprd^{GFP} DRG we used (n=774). All sections were stained with anti-β3-tubulin (cyan). Scale bar 50μm. White arrowheads indicate Na_v1.1+ neurons while white arrows indicate Na_v1.1- neurons. Frequency (**E**) and cumulative (**F**) distribution plots of integrated fluorescence density of Na_v1.1 in TdTomato+ proprioceptors (n=153) **B**. n = cells.

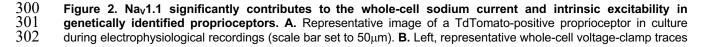
267

268	To determine if $Na_V 1.1$ message in proprioceptors is translated into functional surface
269	protein, we performed pharmacological in vitro patch clamp experiments from TdTomato+
270	neurons harvested from Parvalbium ^{Cre} ;Rosa26 ^{Ai14} (PV ^{Ai14}) adult mice (6-15 weeks, N = 10 mice),
271	which represent genetically identifiable proprioceptors harvested from thoracic spinal levels (de
272	Nooij et al., 2013). The Nav1.1 blocker, ICA 121431 (ICA, 500 nM), was used to block Nav1.1.
273	While ICA also blocks Na $_{\rm V}$ 1.3, those channels are not expressed in uninjured adult DRG neurons
274	(Chang et al., 2018; Felts et al., 1997; He et al., 2010; Usoskin et al., 2015; Waxman et al., 1994)
275	and RNA-sequencing has not found Na $_{ m V}$ 1.3 message in proprioceptors. In voltage clamp
276	experiments, ICA application reduced the whole-cell sodium current (I_{Na}) on average by 46%,
277	suggesting Nav1.1 is a dominant sodium channel isoform in these neurons (Fig 2A-B). I_{Na} density
278	in proprioceptors on average fell from -150 pA/pF to -80 pA/pF following wash on of ICA (Fig 2C).
279	I_{Na} had an average decay tau of 0.8 ms in proprioceptors, which was significantly slowed to 1.1
280	ms following application of ICA (Fig 2D). Blocking Na $_{\rm V}$ 1.1 also resulted in a moderate but
281	significant slowing of the sodium current rise time from 0.2 ms to 0.3 ms (Fig 2E). Quantification
282	of the ICA-sensitive component found an average decay tau of 0.5 ms and an average current
283	density of -89 pA/pF (Fig 2F). Of note, there was a wide distribution of current densities for the
284	ICA-sensitive component, ranging from ~-37 pA/pF to ~-168 pA/pF, suggesting some variability
285	in the contribution of Na $_{\rm V}$ 1.1 to proprioceptor excitability. As such, we next used current clamp
286	experiments to determine of effect of ICA on proprioceptor action potential firing (Fig 2G).

287 Analysis of current clamp data showed that pharmacological inhibition of Nav1.1 significantly 288 reduced the number of evoked action potentials in most genetically identified proprioceptors (Fig 289 **2H**); however, of the 20 cells recorded, 5 had low firing rates that were not further inhibited by 290 ICA, indicating in some proprioceptors subtypes other Navs mediate to action potential firing. 291 Action potential amplitude (Fig 2I, p = 0.0420) and action potential threshold (Fig 2J, p = 0.0186) 292 were also significantly reduced following ICA application. The reduction in action potential 293 threshold is consistent with loss of Na_V1.1, which has a V $\frac{1}{2}$ of activation of ~ -15 mV (Aman et 294 al., 2009). We also observed a significant increase in action potential full-width half-max following 295 ICA application (**Fig 2K**, p = 0.0068), in line with loss of the fast Na_V1.1-mediated current. To our 296 knowledge, these data are the first reported analysis of the I_{Na} in proprioceptors and identify 297 $Na_V 1.1$ as a major contributor.







303 elicited before (black) and after (red) application of ICA121431 (500nM). Right, the subtracted ICA-sensitive current 304 shown in black. C. Quantification of the reduction in whole-cell current density before (white) and after (magenta) ICA, 305 306 p = 0.0002, n=13 cells. D. Quantification of rate of current decay before and after ICA, p = 0.0002, n=13 cells. E. Quantification of whole-cell current rise time before and after ICA, p = 0.0002, n=13 cells. F. Left, current densities of 307 ICA-sensitive sodium currents (n=11), right, current decay taus of ICA-sensitive sodium currents (n=10 cells), G. 308 Representative whole-cell current clamp traces before (left) and after (right) application of ICA. H-K. Quantification of 309 number of action potentials in response to current injection (H, p = 0.0002, n=20 cells), action potential amplitude (I, p310 = 0.0420, n=20 cells), action potential threshold (J, p = 0.0186, n=20 cells), and full-width half max (K, p = 0.0068, n=20 311 cells). Grey lines represent paired observations, circles and lines represent means and standard deviations. White 312 circles, before ICA application. The Wilcoxon matched-pairs signed rank test was used to determine significance.

313

314 We next sought to determine if Na_V1.1 plays an *in vivo* role in motor behaviors. Na_V1.1 315 makes critical contributions to the excitability of parvalbumin-positive interneurons in the brain; 316 thus, we were precluded from using a PV^{cre} driver line to directly interrogate a role for Na_V1.1 in 317 proprioceptors, as loss of Nav1.1 in these interneurons produces an epilepsy phenotype that 318 complicates behavioral analyses in adult animals (Ogiwara et al., 2007). As such, we used a 319 Pirt^{Cre} driver to delete Na_V1.1 in all sensory neurons to determine the contribution of peripherally 320 expressed Nav1.1 to motor behaviors (Kim et al., 2008). Consistent with in vitro data, Pirt^{Cre};Na_V1.1^{fl/fl} (Na_V1.1^{cKO}) animals of both sexes displayed profound and visible motor 321 322 abnormalities. These abnormalities include ataxic-like tremors when suspended in the air 323 (Supplemental video 1), abnormal limb positioning (Supplemental videos 2-3), and paw clasping, which are absent in Nav1.1^{fl/fl} littermate controls and heterozygous animals 324 (Pirt^{Cre};Na_V1.1^{fl/+}, Na_V1.1^{Het}, respectively, **Fig 3A**). We first ran animals in the open field test for 325 326 ten minutes each to quantify spontaneous locomotor behaviors (**Fig 3B**). We found that Na_V1.1^{cKO} animals traveled significantly less (Fig 3C) and slower (Fig 3D) than Na_V1.1^{fl/fl} littermate controls 327 (p = 0.0077 and 0.0057, respectively). Surprisingly, $Na_V 1.1^{Het}$ mice also displayed motor 328 abnormalities in the open field test, performing similarly to Nav1.1^{cKO} animals (Fig 3B-D), 329 330 demonstrating Nav1.1 haploinsufficiency in sensory neurons for motor behaviors. No genotype-331 dependent differences were observed in the amount of time spent moving, suggesting gross 332 motor function was intact (Fig 3E). Additionally, the amount of time spent in the center of the open 333 field chamber was also independent of genotype (Fig 3F). We next used the rotarod assay to 334 investigate differences in motor coordination. Mice were assayed on three consecutive days and

latency-to-fall and RPM were quantified. Unlike in the open field assay, both Nav1.1^{fl/fl} and 335 Na_v1.1^{Het} mice performed at similar levels during the three-day period (**Fig G-H**). Conversely, 336 Nav1.1^{cKO} animals performed significantly worse. By day 3, on average they were only able to 337 maintain their position on the rotarod for 41 s, falling over 50% faster Nav1.1^{fl/fl} and Nav1.1^{Het} mice. 338 339 We did not observe any sex dependent differences in performance in the open field or rotarod tests (Supplementary Fig 1). We confirmed that our mouse model selectively targeted sensory 340 neurons by crossing a Pirt^{Cre} driver with a fluorescent reporter line (Pirt^{Cre};Rosa26^{Ai14}). We 341 342 observed little-to-no neuronal expression of TdTomato in both dorsal and ventral spinal cord 343 (Supplementary Fig 2). In contrast, DRG neurons somata and axons showed strong labeling. 344 Collectively, our behavioral data provide evidence for a new *in vivo* role of Nav1.1 in sensory 345 neurons in motor behaviors and coordination.

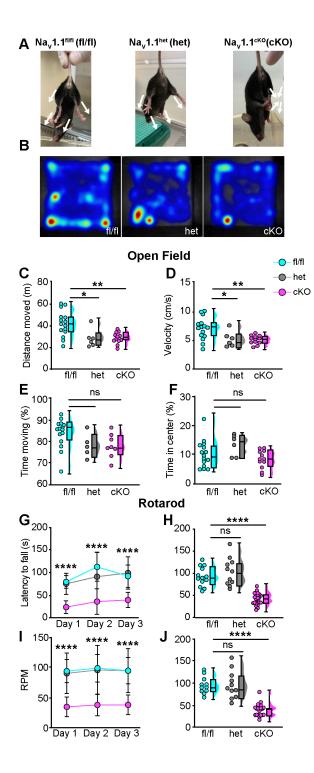
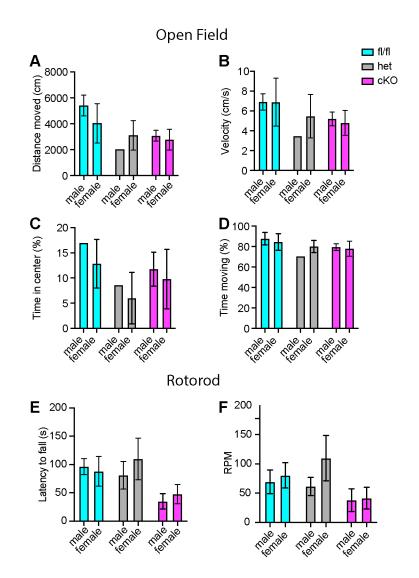


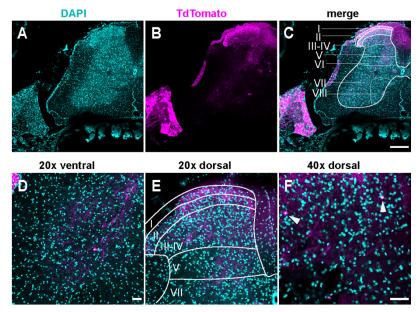


Figure 3. Loss of Nav1.1 in peripheral sensory causes deficits in motor behaviors. A. Representative images showing limb positions of adult Nav1.1^{fl/fl} (left), Nav1.1^{Het} (middle), and Nav1.1^{cKO} (right) mice. White arrows represent the direction of limbs. **B.** Representative heat maps from open field experiments between Nav1.1^{fl/fl} (left), Nav1.1^{Het} (middle), and Nav1.1^{cKO} (right) mice. White arrows represent (middle), and Nav1.1^{cKO} (right), Nav1.1^{Het} (left), Nav1.1^{Het} (middle), and Nav1.1^{cKO} (right). Open Field (**C-F**), Quantification of total distance traveled during a 10-minute open-field test between Nav1.1^{fl/fl} (cyan), Nav1.1^{Het} (grey), and Nav1.1^{cKO} (magenta) mice (**C**, Nav1.1^{Het} p = 0.0255, Nav1.1^{cKO}, p = 0.0077, compared to Nav1.1^{fl/fl}, average animal velocity (**D**, Nav1.1^{Het} p = 0.00311, Nav1.1^{cKO}, p = 0.0057, compared to Nav1.1^{fl/fl}), percent time moving (**E**, Nav1.1^{Het} p = 0.1362, Nav1.1^{cKO}, p = 0.0730, compared to Nav1.1^{fl/fl}), and percent time spent in center (**F**, Nav1.1^{Het} p = 0.2297, Nav1.1^{cKO}, p = 0.2494, compared to Nav1.1^{fl/fl}) during the test. Rotarod

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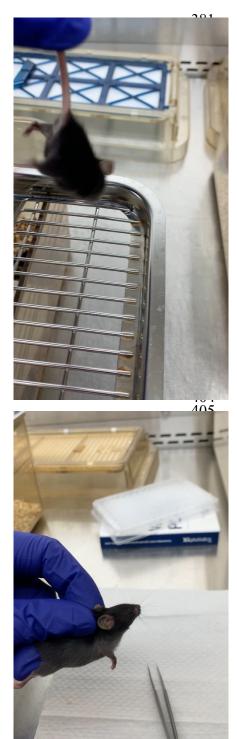
Supplementary Figure 1. Motor deficits in Nav1.1^{Het} and Nav1.1^{cKO} animals are not sex dependent. A-D.
 Quantification of sex-dependent parameters during a 10-minute open field trial. Total distance moved (A). Velocity (B).
 Percent time spent in center of box (C). Total percent time moving (D). Quantification of the sex-dependent differences in rotarod day-three trial latency to fall (E) and RPM (F).



- 368 369 Supplementary Figure 2. TdTomato expression is limited to sensory neurons. A-D, Images stained using immunohistochemistry with anti-DsRed (TdTomato) (A-C) Representative confocal images of spinal cord sections at 370 10X, 0.45 NA dry objective. Scale bar 200µm. (D) Image was taken at 20X, 0.8 NA dry objective of the ventral horn.
- 371 372 (E) Image was taken at 20X, 0.8 NA dry objective of the dorsal horn. Scale bar 50μm. (F) Image was taken at 40X, 1.3
 - NA oil-immersion objective. Arrows indicate TdTomato+ puncta. Scale bar 50μ m. (N=3, n=30 sections). 373



Supplemental Video 1. Uncoordinated movements in Nav1.1cko animals. A Nav1.1^{cko} mouse (left) shows abnormal and spastic movements when suspended in the air. These movements are absent in $Na_V 1.1^{fl/fl}$ mice (right).

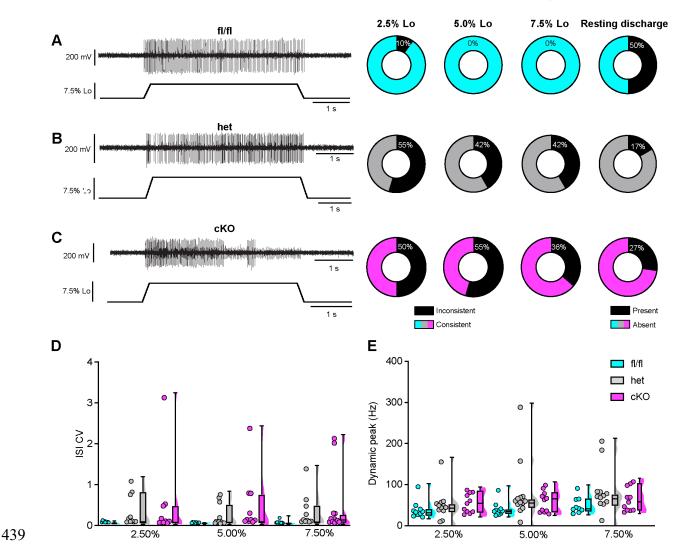


Supplemental Video 2. Abnormal limb position in Na_v1.1^{cko} animals. A Na_v1.1^{cko} mouse has uncoordinated leg movements and makes an abnormal rotation of its hind paw to grasp its tail while suspended in the air.

Supplemental Video 3. Abnormal paw position in Nav1.1^{cko} animals. A Nav1.1^{cko} mouse is scruffed and places hind paws with foot pads facing down. This contrasts with the normal paw positioning seen in the forepaws, in which foot pads are in the outward facing position.

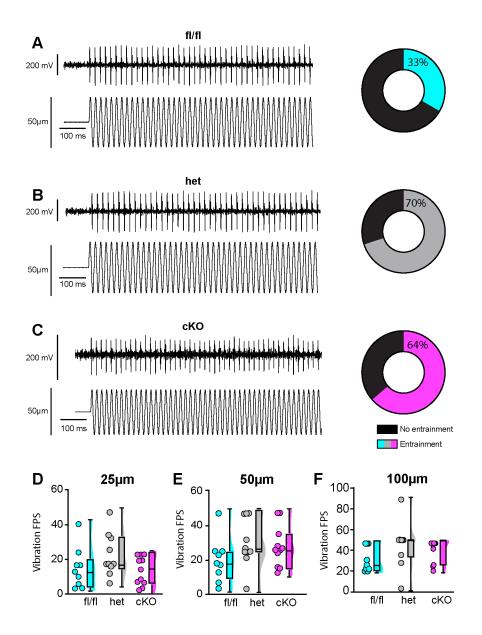
412 Pharmacological inhibition of Nav1.1 in proprioceptors in vitro significantly reduced intrinsic excitability. Thus, we next asked whether proprioceptor transmission was reduced in 413 414 Nav1.1^{cKO} animals. We recorded proprioceptor firing in an ex vivo preparation during ramp-andhold stretch and sinusoidal vibration. Afferents from both Na_V1.1^{Het} and Na_V1.1^{CKO} mice exhibited 415 416 impaired static stretch sensitivity as evidenced by a decreased likelihood of firing during rest as 417 compared to Na_V1.1^{fl/fl} mice as well as an inability to maintain firing throughout the entire 4s 418 stretch. Almost all afferents from Na_V1.1^{fl/fl} mice could fire consistently throughout the entire 4s 419 hold phase (Fig 4A), but loss of one or both copies of Na_V1.1 led to either firing only near the 420 beginning of stretch or inconsistent firing in a high percentage of afferents lacking Na_V1.1 (Fig 421 **4B,C**). We quantified this inconsistent firing by determining the coefficient of variation (CV) of the 422 interspike interval (ISI) during the plateau phase of stretch (1.5-3.5 s into the hold phase) across 423 different stretch lengths and found a significant effect of genotype, with the knockout afferents both having higher ISI CV than the Na_V1.1^{fl/fl} afferents (**Fig 4D**; 0.074 \pm 0.06, 0.313 \pm 0.456, .497 424 ±.831, at 7.5% Lo, Na_V1.1^{fl/fl}, Na_V1.1^{Het}, and Na_V1.1^{cKO} afferents, respectively, Two-way ANOVA. 425 426 p = 0.015). In contrast to the clear deficits in static sensitivity in afferents lacking Na_V1.1, dynamic 427 sensitivity was not significantly impaired. The maximum firing frequency during the ramp up phase 428 (Dynamic Peak) was independent of genotype, and even trended slightly higher in afferents 429 lacking Nav1.1 (Fig 4E; Two-way ANOVA, effect of genotype p=0.0633). We next examined the 430 requirement of Nav1.1 for proprioceptor afferent responses to sinusoidal vibration, which were 431 similarly unchanged or enhanced (Fig 5A-C, Tables 2-4). We characterized a unit as having 432 entrained to vibration if it fired at approximately the same time every cycle of the 9s vibration. In 433 most cases, afferents lacking Nav1.1 were similarly or more likely to entrain to vibration than Na_V1.1^{fl/fl} afferents (Fig 5C). Indeed, Na_V1.1^{cKO} afferents were able to maintain firing during the 434 435 entire 9 s sinusoidal vibration, in contrast to their inability to maintain consistent firing during static stretch. There were no significant differences in firing rate during vibration between Nav1.1^{fl/fl}. 436

437 Na_V1.1^{Het}, and Na_V1.1^{cKO} afferents. Thus, Na_V1.1 is required in proprioceptive afferents for normal



438 responses to static stimuli but is dispensable for afferent responsiveness to dynamic stimuli.

440 Figure 4. Loss of Nav1.1 reduces static muscle stretch sensitivity and reliability. A-C. Representative responses 441 to ramp-and-hold muscle stretch at 7.5% of Lo from Nav1.1^{fl/fl} (A), Nav1.1^{Het} (B) and Nav1.1^{cKO} (C) afferents. Afferents 442 from Na_V1.1^{Het} and Na_V1.1^{cKO} mice were more likely to show inconsistent firing during the hold phase of stretch. The 443 percentage of afferents from each genotype that were able to fire consistently for the entire duration of stretch at 2.5%, 444 445 5.0%, and 7.5% of Lo are shown in the pie charts next to the representative trace from their genotype (black indicates percentage with inconsistent firing). The final pie charts represent the proportion of afferents that exhibited resting 446 discharge at Lo for each genotype. D. Inconsistency in firing was quantified as the interspike interval coefficient of 447 variation (ISI CV) during the plateau stage of the hold phase of stretch (1.5-3.5s into stretch) for the 3 different 448 genotypes. A significant effect of genotype was observed (two-way mixed-design ANOVA, p=0.015) E. The highest 449 firing rate during the ramp up phase of stretch (dynamic peak), which is a measure of dynamic sensitivity. No significant 450 effect of genotype was observed (two-way mixed-design ANOVA, p=0.0633). Each dot represents one afferent 451 (Na_V1.1^{fl/fl}, n=10; Na_V1.1^{Het}, n=12; Na_V1.1^{cKO}, n=11).



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Figure 5. Loss of Na_v1.1 does not alter muscle spindle afferent response to vibratory muscle stretch. A-C. Representative traces from afferents that were able to entrain to a 50Hz, 100µm vibration as well as graphs with the percentage of all Na_v1.1^{fl/fl} (cyan; **A**), Na_v1.1^{Het} (gray; **B**), and Na_v1.1^{cKO} (magenta; **C**) afferents that could entrain to the vibration shown in **A-C**. Average firing frequency during a 9 s 50 Hz vibration shown for a 25 µm (**D**), 50 µm (**E**), and 100 µm (**F**) amplitude vibration. There was no significant effect of genotype on the firing frequency during vibration (25 µm, p = 0.2398, 50 µm, p = 0.2413, 100 µm, p = 0.1276). A one-way ANOVA was used to determine statistical significance in **D** and **E**. A Kruskal-Wallis test was used to determine statistical significance in **F**. Each dot represents one afferent (Na_v1.1^{fl/fl}, n=9; Na_v1.1^{Het}, n=10; Na_v1.1^{cKO}, n=11).

461 462

Table 2. Afferent entrainment to 25 µm amplitude vibration

Genotype	10 Hz	25 Hz	50 Hz	100 Hz
Na _V 1.1 ^{fl/fl}	33.33%	11.11%	0.00%	0.00%
Na _v 1.1 ^{het}	50.00%	40.00%	10.00%	10.00%

Na _V 1.1 ^{cKO}	27.27%	9.09%	0.00%	0.00%

Table 2. The percentage of muscle spindle afferents that entrained to a 25μm amplitude sinusoidal vibration.
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Table 3. Afferent entrainment to 50 μm amplitude vibration

Genotype	10 Hz	25 Hz	50 Hz	100 Hz
Na _V 1.1 ^{fl/fl}	88.89%	22.22%	11.11%	11.11%
Na _V 1.1 ^{het}	80.00%	70.00%	30.00%	10.00%
Na _V 1.1 ^{cKO}	45.45%	54.55%	18.18%	0.00%

Table 3. The percentage of muscle spindle afferents that entrained to a 50µm amplitude sinusoidal vibration.

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470 **Table 4. Afferent entrainment to 100 μm amplitude vibration**

Genotype	10 Hz	25 Hz	50 Hz	100 Hz
Na _V 1.1 ^{fl/fl}	100.00%	44.44%	33.33%	22.22%
Na _V 1.1 ^{het}	90.00%	90.00%	70.00%	40.00%
Na _v 1.1 ^{cKO}	63.64%	72.73%	63.64%	18.18%

471 472 Table 4. The percentage of muscle spindle afferents that entrained to a 100µm amplitude sinusoidal vibration.

473 Could reduced and inconsistent transmission of static stimuli in proprioceptors lead to a 474 concomitant activity-dependent reduction in the number of mature proprioceptors? To address this guestion, we performed RNAscope analysis of DRG sections from Nav1.1^{fl/fl}, Nav1.1^{Het}, and 475 476 Na_v1.1^{cKO} mice. We quantified the number of neurons per DRG section that were positive for both 477 Runx3 and parvalbumin transcript, the molecular signature of mature proprioceptors (Fig 6, Oliver 478 et al., 2021). We found no significant genotype-dependent differences in the number of proprioceptors in Na_V1.1^{Het} and Na_V1.1^{cKO} mice compared to Na_V1.1^{fl/fl} controls (p = 0.3824 and p 479 = 0.1665, respectively), indicating that the behavioral deficits observed in Na_V1.1^{cKO} mice are not 480 481 the result of a developmental loss of proprioceptors. We also analyzed muscle spindle 482 morphology to determine if aberrant sensory end organ development may contribute to the motor 483 abnormalities of Nav1.1^{cKO} mice. Similar to conditional Piezo2-knockout animals (Woo et al., 484 2015), no qualitative differences were observed between genotypes (Supplementary Fig 3).

- 485 Thus, abnormal proprioceptor development does not contribute to the overall phenotype of
- 486 Na_V1.1^{cKO} mice.

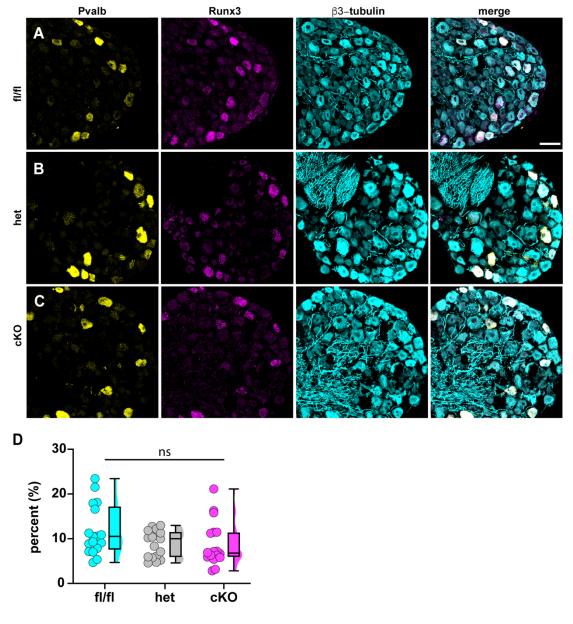
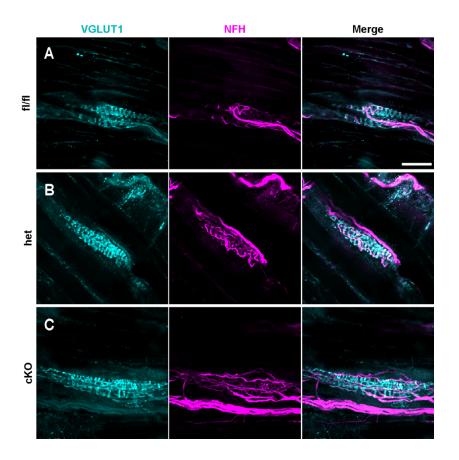


Figure 6. Loss of Na_v1.1 in sensory neurons does not affect proprioceptor development. Representative images of Na_v1.1^{f//f}(**A**), Na_v1.1^{het} (**B**), and Na_v1.1^{cKO} (**C**) adult DRG neuron sections (25 μm). Images were acquired with a 400, 0.9 NA water-immersion objective. Sections were hybridized with probes targeting parvalbumin (*Pvalb*, yellow) and *Runx3* (magenta) and immunostained with anti-β3-tubulin (cyan). (**D**) Quantification of the percentage of Pvalb+/Runx3+ neurons in each genotype. Each dot represents one DRG section. A Kruskal-Wallis test with Dunn's post hoc comparison was used to determine statistical significance (p = 0.1971, Na_v1.1^{f/fl} n=17, Na_v1.1^{Het} n=17, Na_v1.1^{cKO} n=18)). Scale bar 50μm.

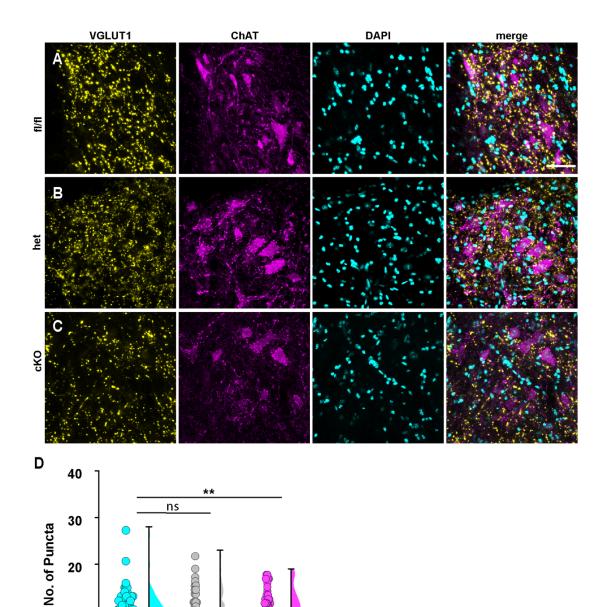
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503 We next asked whether reduced proprioceptor signaling decreased synaptic connectivity 504 between proprioceptive axons and motor neurons in the spinal cord. Spinal cord sections were harvested from Na_V1.1^{fl/fl}, Na_V1.1^{Het}, and Na_V1.1^{cKO} mice and stained with antibodies against 505 506 vesicular glutamate transporter 1 (VGLUT1) and choline acetyltransferase (ChAT), to label 507 proprioceptor axons and α - motor neurons in the ventral horn, respectively. Our analysis found a significant decrease in VGLUT+ puncta per ChAT+ motor neuron in Nav1.1^{cKO} mice compared to 508 $Na_V 1.1^{fl/fl}$ littermate controls (**Fig 7**, p = 0.0031). $Na_V 1.1^{Het}$ mice showed no significant alterations 509 510 in connectivity between VGLUT1+ proprioceptive terminals and ChAT+ motor neuron compared to $Na_V 1.1^{fl/fl}$ littermate controls (p = 0.7695). Thus, in addition to decreased proprioceptive 511

- 512 transmission, Na_v1.1^{cKO} mice also have reduced proprioceptor innervation of motor neurons,
- 513 suggesting their behavioral deficits may also be due in part to aberrant signaling in the spinal
- 514 cord.
- 515





10

0

fl/fl

het



cKO

525 **Discussion**

526 The critical role for Nav1.1 in various brain disorders has overshadowed the potential 527 contributions of this channel in peripheral signaling. The results presented in this study are the 528 first to provide functional evidence that Nav1.1 in peripheral sensory neurons is required for 529 normal motor behaviors. We found that mice lacking Nav1.1 in sensory neurons exhibit visible 530 motor deficits and ataxic-like behaviors, which were quantified in the rotarod and open field 531 behavioral assays. We propose that this aberrant motor behavior is largely attributed to loss of 532 Na_V1.1 in proprioceptors. Indeed, RNAscope analysis showed expression of Na_V1.1 mRNA in 533 100% of parvalbumin-positive DRG neurons. In line with this finding, our functional in vitro patch-534 clamp experiments show pharmacological inhibition of Nav1.1 in proprioceptors is sufficient to 535 significantly attenuate action potential firing, likely due to Nav1.1 carrying nearly half of the I_{Na} in these cells. At the afferent level Nav1.1^{cKO} and Nav1.1^{Het} animals show clear deficits in static 536 537 stretch sensitivity, but not dynamic sensitivity, and could even entrain to vibrations as fast as 100 538 Hz, suggesting that Na $_{\rm V}$ 1.1 in proprioceptors is dispensable for encoding dynamic stimuli. Finally, 539 we found that loss of Nav1.1 in sensory neurons had no effect on proprioceptor development or 540 muscle spindle morphology; however, we did observe a significant reduction in proprioceptive 541 afferent innervation of ChAT+ motor neurons in the spinal cord. Thus, the observed motor deficits 542 are likely due to a combination of reduced static sensitivity of proprioceptor afferents and a 543 concomitant activity-dependent loss of motor neuron innervation.

Our model proposes that $Na_V 1.1$ is tasked with maintaining consistent firing in proprioceptors during static muscle stretch for normal motor behaviors, whereby activation of the mechanotransduction channel Piezo2 initiates electrical signaling, which in turn activates a complement of tetrodotoxin-sensitive Na_V channels (Carrasco et al., 2017; Florez-Paz et al., 2016;

548 Woo et al., 2015, Fig 8). During dynamic or vibratory stimuli, Piezo2, and likely a combination of 549 other molecular mediators, including Nav1.6 and Nav1.7, are sufficient to elicit normal electrical 550 activity. Conversely, during prolonged muscle stretch when Piezo2 channels presumably 551 inactivate, Na_V1.1 is required for regular and reliable firing. In the absence of functional Na_V1.1 552 channels, inconsistent proprioceptor firing leads to a decrease in synaptic connectivity in the 553 spinal cord between proprioceptors and motor neurons, likely exacerbating motor deficits. While 554 other signaling molecules and channels, such as vesicle-released glutamate (Bewick et al., 2005; 555 Than et al., 2021), and mechanosensitive ASIC channels (Lin et al., 2016) and ENaC channels 556 (Bewick and Banks, 2015), also contribute to mammalian proprioception, our data suggests 557 $Na_{V}1.1$ is a critical for muscle spindle mechanotransduction, given the overt behavioral deficits 558 observed in Nav1.1^{cKO} mice.

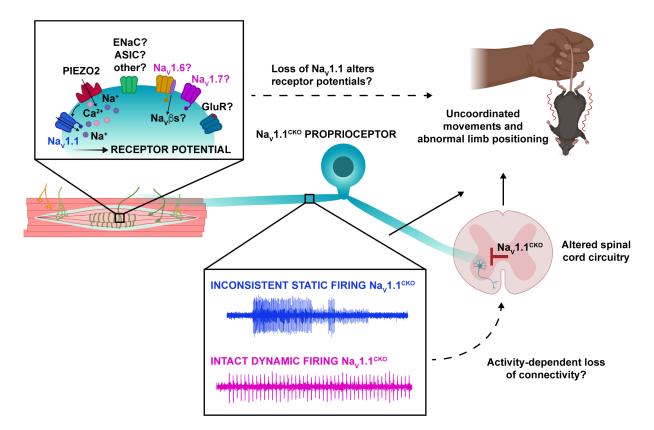


Figure 8. Proposed model of the role of Na_v1.1 in proprioception. Upon muscle static stretch, various channels activate, including Piezo2 (red) which results in an influx of calcium and sodium ions, causing a depolarization that activates Na_v1.1 (dark blue). Na_v1.1 activation drives reliable repetitive firing of proprioceptors during static stretch for normal motor behavior. Loss of Na_v1.1 in sensory neurons results in inconsistent static firing at the afferent level while

564 maintaining dynamic firing. It is possible that a combination of Piezo2, Na_V1.6 (yellow), Na_V1.7 (pink), and/or other 565 channels, such as glutamate receptors, Asic channels, and ENaC, mediate dynamic firing. Unreliable signaling may 566 alter spinal cord circuitry through an activity-dependent loss of synaptic connectivity, resulting in uncoordinated 567 movements and abnormal limb positioning.

568

569 To date, our knowledge of the functional contributions of Nav1.1 in the PNS is limited. 570 Most studies have identified roles for this channel in mechanical pain signaling in DRG, TG, and 571 vagal sensory neurons. Intraplantar pharmacological activation of Nav1.1 induces spontaneous 572 pain behaviors and mechanical pain, which is absent in mice lacking Nav1.1 in small- and 573 medium-diameter sensory neurons (Osteen et al., 2016). Inhibition of Nav1.1 prevented the 574 development of mechanical pain in several preclinical models, including spared nerve injury 575 (Salvatierra et al., 2018b), an irritable bowel syndrome mouse model (Salvatierra et al., 2018), 576 and infraorbital nerve chronic constriction injury (Pineda-Farias et al., 2021). Additionally, blocking 577 Na_v1.1 channels inhibited firing in TRPM8-expressing neurons *in vitro*, suggesting a potential role 578 for this channel in thermosensation (Griffith et al., 2019). No prior studies, however, have reported 579 a role for Na_V1.1 in proprioceptor function or motor behaviors.

580 The biophysical properties of $Na_V 1.1$ within the mammalian muscle spindle may help drive 581 static stretch induced action potential firing in proprioceptors. Indeed, the loss of consistent firing we observed during static stretch in Na_V1.1^{cKO} animals is functionally similar to loss of Na_V1.1 in 582 583 other brain cell types. Loss of a single copy of $Na_V 1.1$ is sufficient to attenuate sustained action 584 potential firing in parvalbumin-positive hippocampal interneurons (Ogiwara et al., 2007; Yu et al., 585 2006) and cerebellar Purkinje neurons (Yu et al., 2006). Mechanistically, Na $_{\rm V}$ 1.1 might contribute 586 to sustained firing in two distinct ways. First, Nav1.1 has been associated with persistent sodium 587 current ($I_{Na}P$) and resurgent sodium current ($I_{Na}R$), both of which promote repetitive firing in a wide 588 variety of cell types in the CNS and PNS (Barbosa et al., 2015; Kalume et al., 2007; Khalig et al., 589 2003). Second, Nav1.1 recovers rapidly from fast inactivation compared to other channel 590 subtypes (Herzog et al., 2003; Patel et al., 2015) and has been shown to be refractory to entry 591 into slow inactivation in TRPM8-expressing DRG neurons (Griffith et al., 2019). Thus, it is possible

592 that proprioceptors rely on these features of $Na_V 1.1$ for reliable and consistent firing in response 593 to static stretch.

594 Loss of Nav1.1 notably impacted proprioceptor afferent static sensitivity during ramp-and-595 hold stretch, but not dynamic sensitivity as measured by entrainment to sinusoidal vibrations using ex vivo muscle-nerve recordings. Afferents from Nav1.1^{cKO} animals were more likely to have 596 597 failures and thus were mostly unable to fire consistently throughout the 4 s of stretch, which was 598 accompanied by a higher coefficient of variability in the ISI. This is consistent with Na_V1.1 playing 599 a critical role in transmitting static stretch information to the central nervous system. Interestingly, 600 however, dynamic sensitivity in these afferents appears to be enhanced, or at least unimpaired. Though not statistically significant, Na_V1.1^{cKO} afferents were more likely to entrain throughout the 601 entire 9 s vibration compared to control mice. Therefore, Nav1.1^{cKO} afferents don't have a 602 603 generalized inability to maintain high frequency firing, but a more specific deficit in static 604 sensitivity. Na_V1.1 has been localized to muscle spindle afferent endings (Carrasco et al., 2017) 605 and has been hypothesized to help amplify receptor current. Our results support a model whereby 606 current via Piezo2 and potentially other mechanically sensitive ion channels at the start of stretch 607 produces a sufficient receptor potential to generate firing at the heminode, but that amplification 608 of the receptor potential by Na_V1.1 is necessary to maintain firing during held stretch. A similar 609 deficit in static but not dynamic sensitivity was seen following loss of synaptic-like vesicle released 610 glutamate from afferent endings (Than et al., 2021); however in those afferents firing only 611 occurred at the beginning of stretch and patchy firing was never observed. This may indicate that 612 glutamate plays a more general role maintaining excitability, whereas Nav1.1 is required for 613 reliable action potential generation at heminodes during static stimuli. Alternatively, or in addition, 614 Na_v1.1 expressed along the axon could be essential for sustained static firing. A detailed 615 examination of Nav1.1 subcellular localization along proprioceptor afferents could shed light on 616 how this channel contributes to signal propagation. The observed trend towards increased

dynamic sensitivity, especially in Na_V1.1^{Het} animals, could suggest the upregulation of other Na_V subtypes or other molecules as a compensatory mechanism to counteract the loss of Na_V1.1. Future studies using temporally controlled deletion of Na_V1.1 in sensory neurons could tease this apart. Nevertheless, as static sensitivity is still very much impaired in Na_V1.1^{cKO} afferents, Nav1.1 may play a potentially unique role in maintaining afferent firing during the sustained stretch.

Although though the response patterns of Nav1.1^{Het} and Nav1.1^{cKO} afferents were similar, 622 behavioral deficits were much more striking in Nav1.1^{cKO} mice (**Fig 4**). One potential explanation 623 624 is the significant decrease in Group Ia proprioceptor synapses (VGLUT1+) on α -motor neurons observed in Na_V1.1^{cKO} animals but not in Na_V1.1^{Het} mice. Nerve injury can permanently reduce 625 626 VGLUT1 synapses on motor neurons (Alvarez et al., 2011), suggesting they may be particularly 627 sensitive to changes in excitability. Given the similar deficits in static stretch sensitivity between these two genotypes, however, it is unclear why Nav1.1^{Het} mice maintain normal synaptic 628 connectivity between proprioceptive afferents and α -motor neurons. Na_V1.1^{Het} afferents trended 629 630 toward having increased dynamic sensitivity, making it is possible that this is sufficient to protect 631 from an activity-dependent loss of motor neuron innervation. Another explanation is a presynaptic 632 role of Na_V1.1 in proprioceptive terminals that is unveiled when both copies of Na_V1.1 are lost. 633 For example, loss of presynaptic Na $_{\rm V}$ 1.7 channels in the spinal cord reduced glutamate release 634 from nociceptive afferents onto dorsal horn neurons (MacDonald et al., 2021). If a similar 635 mechanism is at play for Nav1.1 in proprioceptors, reduced release from afferent terminals could 636 lead to homeostatic changes in connectivity with post-synaptic motor neurons. Future studies are 637 required to test this possibility.

We observed effects of both pharmacological inhibition and genetic deletion of Na_v1.1 in *in vitro* and *ex vivo* electrophysiological experiments, respectively (**Figs 2-3, 5**); however, in our mouse model Na_v1.1 is deleted in all sensory neurons. Thus, we cannot rule out that loss of Na_v1.1 in other mechanosenory neuron populations contributes to the motor deficits observed.

642 Deletion of Nav1.1 in small- and medium-diameter DRG neurons did not produce visible motor 643 deficits (Osteen et al., 2016), indicating sensory neuron populations in those categories are not 644 involved. We did observe Nav1.1 transcripts in the vast majority of myelinated DRG neurons (a 645 combination of large- and medium-diameter DRG neurons), consistent with its presence in 646 different subclasses of tactile sensory neurons (Zheng et al., 2019). However, the severe motor phenotype of Na_V1.1^{cKO} mice precludes mechanical threshold analysis using von Frey or tactile 647 648 sensitivity using tape test. Notably, baseline mechanical thresholds were unchanged following 649 intraplantar injection of a selective $Na_V 1.1$ inhibitor (Salvatierra et al., 2018). This suggests that 650 while $Na_V 1.1$ mRNA is expressed in most tactile sensory neurons, functional protein may only be 651 upregulated in these populations during pathological states.

652 Despite this limitation, one noteworthy and intriguing finding from our study was the 653 haploinsufficiency of Nav1.1 in sensory neurons for normal motor behavior in the open field test. Na_V1.1^{Het} mice had an identical phenotype to Na_V1.1^{cKO} mice, moving more slowly and less than 654 655 controls, despite not having the more severe and visible motor coordination deficits. Indeed, their performance on the rotarod was identical to that of $Na_V 1.1^{fl/fl}$ controls. The gene encoding $Na_V 1.1$, 656 657 Scn1a, is a super culprit gene with over one thousand associated disease-causing mutations, 658 most of which are linked to different forms of epilepsy. Many epilepsy patients with hemizygous 659 Na_V1.1 loss-of-function display ataxia and motor delays and deficiencies (Claes et al., 2001; 660 Fujiwara et al., 2003), which has traditionally been attributed to loss of Na_v1.1 function in the 661 brain, namely the cerebellum (Kalume et al., 2007). Our findings suggest that some of the clinical 662 manifestations associated with epilepsy are not solely due to $Na_V 1.1$ loss-of-function in the brain, 663 but also may manifest in part as a result from unreliable coding by peripheral proprioceptors.

Data presented in this study provide new evidence of a role for peripherally expressed Na_V1.1 in motor coordination. We show that Na_V1.1 is ubiquitously expressed in mammalian proprioceptors, contributes to proprioceptor excitability *in vitro* and *ex vivo*, and is required in

sensory neurons for normal spinal cord motor circuit development. Collectively, this works
identifies a new role for Na_v1.1 in peripheral sensory neuron physiology and motor behaviors.

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670 **AUTHOR CONTRIBUTIONS**

CME performed immunohistochemistry and voltage-clamp experiments. CML performed RNAscope experiments. SO performed *ex vivo* electrophysiology experiments. MSD performed behavioral experiments. KMW and DAO performed current clamp experiments. CME, CML, SO, KMW, and DAO conducted data analysis. TNG and KAW assisted with data analysis and experimental design. CME and CML made the figures. TNG wrote the first draft of the manuscript. CME, CML, and KAW assisted with writing and editing the manuscript. All authors approved the manuscript. TNG, CME, and KAW acquired funding, and TNG and KAW supervised the project.

679 DATA AVAILABILITY

680 All data generated or analyzed during this study are included in the manuscript and supporting 681 file; Source Data files have been uploaded to Mendeley for all figures (doi: 682 10.17632/kt23th75v9.2). GitHub Code has been uploaded to 683 (https://github.com/doctheagrif/Current-Clamp-Matlab-Code O-Neil-DA). A key resources table 684 with specific organism and reagent information has been included in the methods section.

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