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4	Touch-induced Mechanical Strain in Somatosensory Neurons is Independent of Extracellular
5	Matrix Mutations in C. elegans
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20	Running Head: Mechanical strain in sensory neurons
21	Abbreviations:
22	TRN: touch receptor neuron
23	ECM: extracellular matrix

24 Abstract

Cutaneous mechanosensory neurons are activated by mechanical loads applied to the 25 26 skin, and these stimuli are proposed to generate mechanical strain within sensory neurons. Using a microfluidic device to deliver controlled stimuli to intact animals and large, immobile, and 27 28 fluorescent protein-tagged mitochondria as fiducial markers in the touch receptor neurons 29 (TRNs), we visualized and measured touch-induced mechanical strain in C. elegans worms. At steady-state, touch stimuli sufficient to activate TRNs induce an average strain of 3.1% at the 30 31 center of the actuator and this strain decays to near zero at the edges of the actuator. We also 32 measured strain in animals carrying mutations affecting links between the extracellular matrix 33 (ECM) and the TRNs but could not detect any differences in touch-induced mechanical strain between wild-type and mutant animals. Collectively, these results demonstrate that touching the 34 35 skin induces local mechanical strain in intact animals and suggest that a fully intact ECM is not essential for transmitting mechanical strain from the skin to cutaneous mechanosensory neurons. 36 37

38 Introduction

Touch and proprioception are essential to the daily lives of all animals, including 39 humans. Classical examples of mechanotransduction, both of these senses depend on activation 40 of mechano-electrical transduction (MeT) channels arrayed within somatosensory neurons (Katta 41 42 et al., 2015) and are thought to activate following physical deformation of sensory cells during 43 touch and movement. Consistent with this inference, neurons innervating stretch receptor organs 44 in vertebrates (Bewick and Banks, 2015) and arthropods (Suslak and Jarman, 2015) are activated 45 by experimentally applied mechanical strain (stretch). High-speed volumetric imaging reveals that proprioceptors deform in crawling Drosophila larvae (He et al., 2019; Vaadia et al., 2019) 46 47 and that physical deformations are correlated with calcium transients. Similarly, movement 48 induces localized calcium transients in the tertiary dendrites of the C. elegans PVD 49 proprioceptors that depend on expression of a DEG/ENaC/ASIC protein (Tao et al., 2019). These 50 studies reinforce the idea that physical deformation of sensory neurons during movement is 51 critical for activation of native MeT channels, but do not address whether or not similar deformations occur in response to touch or measure the extent of mechanical strain that occurs 52 53 within the sensory neurons themselves.

54 Using C. elegans touch receptor neurons (TRNs), we investigated the quantitative relationship between touch and physical deformation of sensory neurons in intact living animals. 55 56 The ready availability of transgenic animals expressing TRN-specific markers and their 57 transparent body make C. elegans an especially useful animal for investigating touch-evoked 58 sensory neuron deformation. Adult animals have six TRNs, consisting of two bilaterally symmetric pairs of touch receptor neurons (ALM and PLM) and two neurons that run along the 59 ventral midline (AVM and PVM) (Goodman, 2006). These six neurons extend long, unusually 60 61 straight sensory neurites (Krieg et al., 2017) that are embedded in epidermal cells and have a 62 distinctive, electron-dense extracellular matrix or ECM as well as hemidesmosome structures 63 attaching the TRNs to the cuticle (Chalfie and Sulston, 1981; Chalfie and Thomson, 1979).

All of the TRNs express the MEC-4 channel, which is required for touch-evoked calcium
transients (Suzuki et al., 2003) and touch-evoked MeT currents (O'Hagan et al., 2005). The
MEC-4 channels localize to puncta arrayed along the entire length of wild-type TRN sensory

neurites (Chelur et al., 2002; Cueva et al., 2007; Emtage et al., 2004; Katta et al., 2019), but are
disrupted in ECM mutants (Emtage et al., 2004).

69 Touch-evoked behavior (Petzold et al., 2013) and MeT channel activation (Eastwood et al., 2015) depend on body indentation rather than the force applied. Slow stimuli fail to activate 70 71 MeT currents and their size increases with stimulus frequency, indicating that activation of MeT 72 channels in their native environment depends on tissue viscoelasticity (Eastwood et al., 2015; 73 Katta et al., 2019; Sanzeni et al., 2019). Although slow, movement-induced physical 74 deformations of the worm's body and its neurons are too slow to activate MeT currents in the 75 TRNs (Eastwood et al., 2015; Katta et al., 2019), these undulatory movements produce 76 mechanical strains of up to 40% (Krieg et al., 2017; Krieg et al., 2014), indicating that TRNs can withstand significant mesoscale extension and compression in situ. The ability to withstand 77 78 movement-induced strain is shared by mammalian nerves that experience up to 30% strain 79 during limb movement (Phillips et al., 2004).

80 Here, we sought to determine whether or not body indentations sufficient to evoke calcium transients in TRNs also generate local strain. To achieve this goal, we visualized steady-81 82 state, touch-induced strain in TRNs in living animals restrained within a microfluidic stimulation device equipped with the ability to deliver mechanical stimuli (Fehlauer et al., 2018; Nekimken 83 et al., 2017a). Our approach benefits from large, immobile mitochondria distributed within TRN 84 85 sensory neurites (Sure et al., 2018), exploits the ability to tag these mitochondria with a red 86 fluorescent protein (Zheng et al., 2014), and borrows analytic principles from traction force 87 microscopy (Ribeiro et al., 2016). Although mitochondria have been used to evaluate neuronal mechanics in culture (O'Toole et al., 2015), we believe this study is the first to make use of 88 mitochondria as mechanical fiducial markers in living animals. We show that body indentation 89 90 increases local steady-state mechanical strain in the TRNs in a manner that is robust to mutations 91 known to affect attachment of the TRN to the extracellular matrix and to epidermal cells.

92 **Results**

93 Mechanical stimulation in a microfluidic device

To directly observe touch-induced deformation of the TRNs, we sought to develop a 94 95 method that combines the delivery of controlled mechanical stimuli and three-dimensional optical imaging. To reach this goal, we applied mechanical stimuli to adult worms confined in a 96 microfluidic device (Nekimken et al., 2017, Figure 1A). Our device has pneumatic actuators that 97 98 consist of a thin flexible wall that separates a channel filled with air from a worm in the trap channel. When air pressure in the actuator channel is increased using a pressure controller, the 99 thin wall expands like a balloon, deforming the trapped worm and generating indentations 100 101 sufficient to activate the TRNs (Nekimken et al., 2017a). Other devices that deliver mechanical stimuli to restrained (Cho et al., 2018) or moving (McClanahan et al., 2017) worms in 102 103 microfluidic chambers have been reported, but they use larger actuator channels and are thus not 104 well-matched to our goal of investigating local deformation of the TRNs.

105 Our device is designed to increase the probability that animals will enter the trap with 106 either their left or right sides in contact with the bottom of the chamber and their ventral and 107 dorsal sides near the actuator (Nekimken et al., 2017a). Consistent with this expectation and the 108 position of the AVM and ALM neurons within the worm's body, previous experiments using this 109 device resulted in a higher frequency of AVM activation compared to ALM (Nekimken et al., 110 2017a). However, the rotational orientation of each worm was variable. We took advantage of this variation to select animals oriented such that ALM (rather than AVM) was near the actuator 111 112 and used only ALM neurons for this study. In all cases, this variation in worm positioning leads to variation in the size of the effective stimulus. To ensure that all animals received a deforming 113 114 stimulus, we limited our analysis to animals whose TRNs were deformed enough to observe 115 visually during the experiment.

116

117 Mitochondria as fiducial markers for inferring strain

We used fluorescent protein-tagged mitochondria as fiducial markers to measure strain,
an approach that uses particle tracking with large (~1µm), immobile mitochondria in the TRNs
(Sure et al., 2018) serving as natural fiducial markers. We imaged mitochondria position before

and after mechanical stimulation (Figures 1B and 1C). Next, we used the displacement between 121 122 adjacent mitochondria to quantify one-dimensional mechanical strain along the long axis of the 123 TRN (Figure 1D) according to: $\varepsilon = \Delta L/L_0$, where L_0 is the resting, undeformed length of an 124 object, and ΔL is the change in length of the object when deformed. For clarity, we refer to this 125 as longitudinal strain. As shown in Figure 1, mitochondria adjacent to the actuator moved more 126 than those anterior and posterior to the actuator (Figures 1B and 1C) and the inferred strain was 127 greatest near the center of the actuator (Figure 1D). This method enables direct observation of 128 touch-evoked strain in C. elegans TRNs in a single dimension aligned with the TRN's longest 129 dimension.

130 We note that this method is not suitable for detecting shear strain or bending strain. To 131 derive these measurements, we would need to detect angular changes (shear) or relative position 132 (bending) of the mitochondria within the TRNs. Given that the diameter of each mitochondrion 133 is only ~100 nm and the TRN is diameter is not much larger than this (200-300 nm, see Figure 3C, Cueva et al., 2007), such movements cannot be resolved because these objects are similar in 134 135 size to the estimated lateral resolution of 254 nm and axial resolution of 632 nm of our imaging system. What about strain in directions orthogonal to the long axis of the TRN dendrite? 136 137 Although our image stacks contain three-dimensional position data (Figures 2A and 2B), the 138 initial distance between adjacent pairs of mitochondria in the x and z directions is small 139 compared to the resulting displacement. Thus, measurement resolutions lead to large uncertainties in the strain calculation for these dimensions. Both limitations can be attributed to 140 141 the confinement of the mitochondria within the narrow caliber of the TRN and the fact that the TRNs lie primarily within a single focal plane. Thus, the size of mitochondria, the geometry of 142 143 the TRNs and the optical resolution of the spinning-disk confocal limit this measurement to one-144 dimensional strain along the y-dimension that traces the main axis of the TRN.

145

146 Local indentation induces local mechanical strain in TRNs

147 Next, we estimated the spatially averaged distribution of touch-induced strain in the
148 TRN. The mitochondria are sparsely distributed in the TRN and strain is a pairwise measurement
149 between adjacent mitochondria, so our strain measurements result in a series of discontinuous
150 step functions for a single trial rather than a smooth curve (Figure 1D). Each segment of the plot

indicates the average strain between two markers, so the maximum local strain on each interval 151 152 may be larger than our measured strain. The average distance between mitochondria in control 153 *jsIs1073* animals was 26 µm (min: 5 µm; max: 64 µm) and we were able to analyze an average 154 of 8.9 intervals per trial (Table S1). To obtain averages across trials, we plotted deformation 155 against position in the longitudinal or y direction, fit this profile with a Gaussian function and 156 defined the position of the maximum as the center of the actuator and y=0. (see dotted line in Figure 1C). In this coordinate system, positions anterior to y=0 are negative and positions 157 158 posterior to y=0 are positive. Finally, we determined the average strain at each longitudinal position along the neurite across actuation trials (see Figure 2C). Note that the average strain is 159 160 both larger and noisier for anterior (negative) positions. This is likely to reflect the fact that the 161 head is less constrained in the microfluidic channel than other parts of the body and suggests that 162 displacement and strain on the anterior side of the actuator include both touch- and movementinduced mechanical strain on the neuron. 163

164 As expected, strain at the center of the actuator (y=0) increases with TRN deformation (Figure 2D), but the dependence on TRN deformation was weak. The observed, but modest 165 166 variation in TRN deformation arises from two aspects of our method. First, because the actuator 167 is fixed to the surrounding material of the device on all sides, its center is the location of 168 maximum deformation. Second, although most animals are trapped with either their left or right 169 side in contact with the coverslip at the bottom of the device, some are rotated along their anterior-posterior axis such that the position of the imaged TRN varies with respect to the 170 171 actuator (Figure 2A). Thus, the average strain induced between the pair of mitochondria at the 172 center of the actuator was 0.031 ± 0.005 (mean \pm SEM, n = 61 trials, N=15 animals) in transgenic 173 *jsIs1073* TRN neurons. Although additional studies exploring a wider range of deformations will 174 be needed to determine the nature relationship between TRN deformation and strain, to our 175 knowledge, these are the first *in vivo* measurements of touch-induced longitudinal strain in touch 176 receptor neurons, and they link local body indentation to local cellular deformation.

177

178 Do tagged mitochondria affect TRN function?

We used classical touch assays to address this question (Chalfie et al., 2014; Nekimken et
al., 2017b) by measuring touch sensitivity in wild-type (N2) and transgenic worms carrying the

181 transgenic mitochondria marker, *jsIs1073*. In blinded assays of three independent cohorts of 25

animals for each genotype, wild-type and *jsIs1073* animals had touch response rates of 0.869 and

- 183 0.865, respectively. The difference between the means was 0.00667 [95% CI (-0.0347, 0.048)].
- 184 Thus, the *jsIs1073* transgene does not decrease touch sensitivity.
- 185

186 Selected ECM mutants and their touch sensation and anatomical phenotypes

187 Having measured touch-induced mechanical strain in control TRNs, next we measured longitudinal strain in four existing ECM mutants: him-4(e1267), mec-1(e1738), mec-1(e1066), 188 189 and mec-5(u440). One of these mutants, him-4(e1267), has a partial defect in touch-sensitivity, and the others are touch-insensitive (Du et al., 1996; Emtage et al., 2004; Vogel and Hedgecock, 190 191 2001). We reproduced these previous results using a ten-touch assay (Table 1). Whereas him-4 192 mutants retain MEC-4 puncta that are grossly wild-type, TRN neurites in mec-1 and mec-5 193 mutants lack prominent MEC-4 puncta (Emtage et al., 2004). In him-4 and in mec-1(e1738) 194 mutants analyzed here, the ALM and PLM neurons are displaced from their normal body 195 position near the lateral midlines and are not properly embedded in the epidermis (Emtage et al., 196 2004; Vogel and Hedgecock, 2001). This effect is inferred to arise from a defective TRN-ECM 197 attachment (Vogel and Hedgecock, 2001). Consistent with this idea, the electron-dense ECM is not detected in mec-1 mutants (Chalfie and Sulston, 1981). Analyzing these four mutants enables 198 199 us to evaluate the relationship, if any, between touch-induced longitudinal strain and three other 200 phenotypes: behavioral responses to touch, the distribution of MEC-4 puncta, and TRN-ECM attachment. 201

202 The *him-4* gene encodes hemicentin, a conserved ECM protein that is rich in Ig and EGF domains and is expressed by body wall muscles (Vogel and Hedgecock, 2001). The him-4 locus 203 204 is >30 kb, which complicated determination of the molecular defect encoded by e1267 using classical sequencing methods. We used whole-genome sequencing to circumvent this limitation 205 206 and found that e1267 encodes a single base indel in the third intron of the him-4 gene, 207 introducing a shift in the reading frame of the encoded protein (Figure S1). A second polymorphism was detected in the intron following the 48th exon. Thus, the e1267 allele is a null 208 209 allele and animals carrying this mutant are likely to lack the HIM-4 protein. The mec-1 gene 210 encodes a large secreted protein rich in kunitz-like domains, is expressed by the TRNs, and many

211 alleles of this gene were found in forward genetic screens for touch-insensitive animals (Chalfie 212 and Au, 1989; Chalfie and Sulston, 1981). The e1066 allele is a null and animals carrying this 213 allele are likely to lack the MEC-1 protein, whereas the *e1738* allele encodes a premature stop 214 codon and is proposed to express a truncated MEC-1 protein. Both mec-1 mutants are touch 215 insensitive, but only e1066 lacks proper TRN-ECM attachments (Emtage et al., 2004). mec-5 216 encodes an atypical collagen and, unlike *mec-1*, it is not made by the TRNs (Du et al., 1996). 217 Like other ECM mutants we analyzed here, the *u440* allele is null and *u440* mutants do not express the MEC-5 protein. The TRN neurites in mec-5 mutants are positioned near the lateral 218 219 midlines, but are less straight than they are in wild-type animals, meandering such that some 220 segments of the neurite are close to the muscle and other parts closer to the lateral midline 221 (Emtage et al., 2004).

222

223 Touch-induced longitudinal strain in TRNs with ECM mutants

224 To analyze transmission of mechanical energy from the skin in animals lacking proper TRN-ECM attachment, we introduced the *jsIs1073* transgene into *him-4* and *mec-1* mutants and 225 226 applied mechanical stimuli to animals trapped in our pneumatic microfluidic device. Touch-227 induced longitudinal mechanical strain in him-4(e1267) TRNs was indistinguishable from that observed in control animals (Figure 3A, Table 2). Touch-induced longitudinal strain was 228 229 likewise similar to control in both *mec-1* mutants (Figure 3B) and in *mec-5* mutants (Figure 3C). Figure 3D shows that the average longitudinal strain profiles for control, him-4, mec-1, and 230 231 *mec-5* mutants are similar across the entire 200 µm segment of the TRN neurite analyzed here. 232 Values for strain measured at the actuator center (y=0) had similar mean values and distribution 233 for all genotypes tested (Figure 4A) and estimation graphics (Ho et al., 2019) indicate that the 234 mean values for all genotypes are not different than control values (Figure 4B). Except for mec-5 235 mutants, the number of mitochondria available for tracking and the average distance between 236 mitochondria was similar in control transgenic and mutant animals (Table S1). mec-5 mutants 237 had fewer, more widely spaced mitochondria than control animals, an effect would be expected to impair the spatial resolution of longitudinal strain measurements. Collectively, these findings 238 239 suggest that neither severe nor mild defects in TRN-ECM attachment play a significant role in the generation of touch-induced longitudinal mechanical strain at steady-state. Future studies 240

and techniques will be needed to resolve dynamic changes in local strain that are expected to 241 242 occur on the millisecond timescale of MEC-4-dependent MeT channel activation.

243

244 Discussion

245 Using mitochondria as natural fiducial markers in C. elegans TRNs, we showed for the first time that body indentation sufficient to activate the TRN causes an increase in longitudinal 246 strain. The magnitude of this strain increases with TRN displacement. In wild-type animals, the 247 248 touch-induced strain closest to the point of maximum indentation was roughly 3% (Figure 3, 249 Table 2). This change in strain is unlikely to damage these sensory neurons, since they are 250 subjected to mechanical strain on the order of 40% during locomotion (Krieg et al., 2017; Krieg 251 et al., 2014). Such global changes in strain are too slow to activate the TRNs (Eastwood et al., 252 2015; Katta et al., 2019).

253 The 3% strain we measured may be a lower bound on the true value for longitudinal 254 strain. Simulations predicted an elongated strain field with a peak value of 12% for longitudinal 255 strain (Sanzeni et al., 2019). Unlike our measurements, which used a wide $(50\mu m)$, flexible 256 PDMS membrane to indent the worm's body and have limited spatial resolution, the simulations 257 employed a stiff spherical bead (10µm diameter) to deliver touch stimuli and a continuous 258 deformation function. The simulations (Sanzeni et al., 2019) predict strains are present in the 259 other two dimensions (along the direction of the stimulus and tangential to the circumference of 260 the worm at the point of the stimulus). Due to the nature of our measurement, however, we were 261 only able to measure one-dimensional strain along the length of the TRN. Nevertheless, these 262 measurements of touch-induced strain can be incorporated into future models of C. elegans touch 263 sensation to improve understanding of the mechanical state of the TRN and mechano-sensitive 264 ion channel complex upon touch stimulation.

265

266

Origins of mechanical coupling between the skin surface and somatosensory neurons

We tested the idea that strain transmission would depend on TRN-ECM attachment, be 267 268 correlated with impaired touch sensation, and be independent of the proper distribution of 269 MEC-4 channels. To our surprise, we found that touch-induced longitudinal mechanical strain

was similar in control TRNs and all ECM mutants tested here, including the *him-4(e1267)* and *mec-1(e1066)* mutants exhibiting severe defects in TRN-ECM attachments. Thus, touch-induced
longitudinal strain we observed in the ALM neurons is independent of their attachment to other
tissues or the expression of ECM proteins MEC-1 and MEC-5, at least at steady state.

274 This finding implies that mechanical coupling between the skin and sensory neurons 275 persists in mutants with ECM defects. Friction is one alternative source of mechanical coupling 276 between TRNs and surrounding tissues. In this scenario, all of the worm's tissues are compacted 277 together by a high hydrostatic pressure and this would elevate friction between tissues. Further compression applied to the outside of the worm during touch stimulation could lead to stiffening 278 279 of the worm's body that might be caused by internal structures jamming together (Gilpin et al., 280 2015). When we and others immobilize worms in microfluidic devices, we fabricate channels 281 small enough to apply gentle compressive forces that the worm cannot overcome. As a result, the 282 worm is compressed on all sides except the nose and tail when in the microfluidic trap, 283 potentially jamming the TRNs against surrounding tissues and further increasing friction. 284 Another possibility is that the generation of touch-induced local mechanical strain is dominated 285 by the cytoskeleton. In support of this idea, mutations that disrupt the expression of MEC-12 α -286 tubulin and MEC-7 β-tubulin decrease mechanoreceptor currents and increase stimulus amplitude needed to activate these currents (Bounoutas et al., 2009; O'Hagan, 2005). Additional 287 288 experimental work and methods with improved spatial and temporal resolution will be needed to 289 differentiate among these possibilities.

290

291 Spatial and temporal resolution of mechanical strain transmission

292 The spatial resolution of our strain measurements is limited by the average distance 293 between adjacent, immobile mitochondria, which was 26µm (Table S1). Thus, this method may 294 lack sufficient spatial resolution to detect micro- or nanoscale variations in mechanical strain 295 transmission at steady state. The spatial resolution of strain measurements could be improved by 296 using more closely spaced fiducial markers. In principle, MEC-4 channels (with an average 297 spacing of 2-3 µm) tagged with a very bright fluorescent protein would provide a ten-fold 298 increase in resolution. Independent of potential improvements in spatial resolution, the present 299 approach is limited to steady-state measurements because our volume imaging rate is low (15

seconds per stack). In this regard, it is important to note that wild-type *C. elegans* TRNs are
preferentially activated mostly by high-velocity stimuli (Eastwood et al., 2015; Katta et al., 2019;
Nekimken et al., 2017a; Suzuki et al., 2003). Temporal resolution could be improved using other
imaging techniques enabling rapid acquisition of imaging volumes. If volumes could be acquired
at 20 Hz or faster, it would possible to observe mechanical strain in a 10 Hz buzz stimulus that
we previously used to activate the TRNs (Nekimken et al., 2017a).

306 *Conclusion*

307 We performed the first *in vivo* measurements of touch-induced mechanical strain in C. elegans TRNs. We used mechanically stable mitochondria in the TRNs as natural fiducial 308 309 markers to observe deformation of the TRN and found that local touch stimuli applied in a 310 microfluidic device induces local strain in the TRN. Defects in the ECM surrounding the TRN did not alter the steady-state mechanical strain in the TRN, suggesting that explicit attachments 311 are not necessary for deformation of the TRN and that the bulk properties of tissues are sufficient 312 313 to sustain significant mechanical energy transfer from the skin surface to the embedded neurons. In light of the temporal limitations of our measurements, however, we cannot exclude the 314 315 possibility that TRN-ECM attachments contribute to dynamic aspects of mechanical strain 316 transmission. Collectively, these findings provide an empirical basis for the idea that mechanical stimuli applied to the skin stretch embedded sensory neurons that may be shared by the sensory 317 318 neurons that innervate the skin and other tissue in mammals. 319

320 Methods and Materials

- 321
- 322 Nematode strains
- 323 For all experiments measuring mechanical strain, we used animals carrying *jsIs1073*
- 324 [mec-7p::TagRFP-mito::CBunc-119], a transgene that drives expression of TagRFP in the
- mitochondria in the TRNs (Zheng et al., 2014), alone or together with ECM mutants (Table 3).
- We relied on visualization of the *jsIs1073* transgene and behavioral phenotypes (for *mec* genes)
- 327 or anatomical defects (for *him-4*) to perform genetic crosses. We used gene sequencing to verify
- 328 the presence of ECM mutants in all strains created for this study. The primer pairs we used to
- amplify the relevant segments of each gene are:
- 330 *mec-1(e1066)*: Forward–catettccacgccgcaaagtc, Reverse—aatcctctctgccctcatgttcc
- 331 *mec-1(e1738)*: Forward—tcacagtcagacgtgcctcg, Reverse—cattgcctcacaccaacttccac
- 332 mec-5(u444): Forward—cagaatactatgtacgtaacttgggatc; Reverse—ctcatgggtacgcaaatgatactc
- *him-4(e1267)*: Forward—tttcgtgatgactgtggtgactgtgg; Reverse—ttaaagtcaacagcaccgtgacc
- 334

335 Immobilization and mechanical stimulation with a pneumatic microfluidics device

336 To provide a repeatable mechanical stimulus that is compatible with imaging the mitochondria of 337 the TRN, we used a microfluidic device made of the transparent elastomer PDMS for 338 simultaneous mechanical stimulation and imaging of *C. elegans* (Nekimken et al., 2017a). Using a pressure controller (Elveflow OB1), we applied 300 kPa of pressure to one of the device's 339 340 actuators to create a mechanical stimulus. We fabricated and operated devices designed for use 341 with young adult worms as described in our previous work (Fehlauer et al., 2018; Nekimken et 342 al., 2017a). Because worms immobilized in this trap are uniformly and partially deformed by the 343 channel, we refer to this as the rest configuration.

344

345 Sample preparation and inclusion criteria

We performed all experiments using young adult animals that were synchronized by
hypochlorite treatment (Stiernagle, 2006) and cultured for three days on NGM agar plates seeded
with OP50 bacteria for food. We transferred animals from the agar plates with a platinum wire

pick to a drop of imaging medium (see below) in a small Petri dish. After using the pick to push 349 350 away large particles that might clog the microfluidic device, we aspirated worms into 351 polyethylene tubing (PE50, Intramedic[™] brand, Becton-Dickson) with a syringe and connected 352 the tubing to the device's inlet. For each trial, we pushed a worm into the trap channel using the syringe and then evaluated whether to use this worm for an experiment based on our inclusion 353 354 criteria. We included animals that fit in the microfluidic trap with minimal movement and could be ejected through the narrow opening at the head of the trap. When performing experiments 355 with him-4 mutants, we chose animals whose gross morphology was wild-type and avoided 356 357 animals whose intestines were everted. In all cases, we only acquired images from animals 358 whose ALM neurons were oriented adjacent to the mechanical actuation channels.

359 To improve image quality by reducing reflections generated by the walls of the 360 microfluidic trap, we designed a non-toxic imaging medium with a refractive index similar to 361 PDMS (1.4). The imaging medium was a 70%:30% (vol:vol) mixture of physiological saline 362 and iodixanol (OptiprepTM, Sigma-Aldrich), a non-toxic density-gradient medium (Boothe et al., 363 2017). We used the same physiological saline as that used for electrophysiological recordings 364 from C. elegans neurons (O'Hagan et al., 2005), which contains (in mM): NaCl (145), KCl (5), 365 MgCl₂ (5), CaCl₂ (1), and Na-HEPES (10), adjusted to a pH of 7.2 with NaOH. The imaging 366 medium has an osmolality of 300-325 mOsm (Fiske Micro-Osmometer Model 210), so it does 367 not cause large osmotic shocks to C. elegans. By contrast with other fluids we tested (e.g. 368 glycerol, halocarbon oil), this medium has a viscosity that appeared to be similar to that of 369 physiological saline.

370

371 *Image acquisition*

372 Although the mechanical stimulus is mostly in the horizontal direction, there is enough

deformation in the z-direction to move the neuron out of plane during stimulation in some cases.

374 In initial experiments with a traditional epifluorescence microscope, the mitochondria often

375 moved out of focus during stimulation due to movement in the z-direction. To account for this

problem, we used a spinning disk confocal microscope (Nikon TiE, Yokogawa CSU-X1,

377 40x/NA 1.4 oil objective, and Photometrics Prime95B sCMOS camera) to acquire z-stacks,

378 which provided adequate resolution in the z-direction.

For each trial, we acquired eleven (11) z-stacks containing the neuron of interest, with 300 kPa of pressure applied during the even-numbered stacks, and 0 kPa applied during the oddnumbered stacks. We started acquisition of a stack every 15 seconds, although the time to acquire each given stack was approximately 12 seconds. The exact time varied depending on the height of the z-stack, which was manually set to accommodate observed motion in the z direction during the test actuation. During the short delay between acquisition of stacks, we toggled the applied pressure.

386

387 Image analysis

388 To detect the mitochondria, we used a particle-tracking algorithm implemented in Python (Allan et al., 2018), based on work by Crocker and Grier (Crocker and Grier, 1996). Briefly, the 389 390 algorithm involves applying a spatial band pass filter, finding peaks, refining the position of 391 peaks by finding their center of mass, and linking particles across timepoints into trajectories. In 392 some stacks, the algorithm failed to detect the mitochondria because they were too close to the 393 top or bottom of the stack, were not bright enough, overlapped with the mitochondria of another 394 TRN, or were blurred due to the motion of the worm. We discarded all stacks subsequent to a 395 stack where the image processing failed, because the strain measurement requires comparison to 396 a previous stack. Additionally, not all of the stacks from each trial were usable, since the 397 fluorophores bleached over time. As a result, not all actuation events yielded strain 398 measurements (see Supplementary Figure S2).

399 Due to variability across trials, we manually selected a region of interest around the TRN 400 and tuned parameters in the particle tracking algorithm. Primarily, we changed the minmass 401 threshold, which filters out particles where the sum of pixel values within the boundaries of the 402 particle is below the chosen threshold, and the search radius for the linking step, which specifies 403 how far a particle can travel between timepoints and still be identified as the same particle. Less 404 frequently, we changed the cutoff size for the bandpass filter or the brightness percentile 405 threshold, which sets a minimum value for the brightest pixel in a particle as a percentile of the brightness of pixels in the image. We tuned these parameters until the particles found and linked 406 407 included only particles along the location of the neuron and not autofluorescent spots. 408 Supplemental Figure S3 shows the range of parameters we used.

409

410 *Touch assays*

To test the touch-sensitivity of the mutants used in our strain transmission experiments, we performed touch assays by lightly stroking an eyebrow hair across the body of a worm and scoring its behavioral response (Goodman, 2006). For each session of touch assays, we tested 25 worms from a plate, performing 10 touches per worm. We performed the touch assays blinded with respect to genotype. For each touch event, we counted a response consisting of reversing direction or speeding up to move away from the stimulus as a positive response.

417

418 *Whole-genome sequencing of* him-4(e1267)

419 Our whole-genome sequencing protocol involved four sub-protocols: 1) DNA extraction, 2) 420 sequencing library preparation, 3) sequencing, and 4) analysis. We isolated DNA from CB1267 421 him-4(e1267) X animals with a phenol chloroform isoamyl alcohol (PCI) extraction. First, we 422 washed worms off a mostly starved plate using M9 buffer, rinsed the animals twice in M9 buffer, 423 resuspended them in EN buffer (0.1M NaCl and 20 mM EDTA), removed the supernatant, and 424 flash-froze the sample in liquid nitrogen. Next, we added 450 µL of worm lysis buffer (0.1 M TRIS pH 8.5, 0.1 M NaCl, 50 mM EDTA, and 1% SDS) and 40 µL of proteinase K (10 mg/ml) 425 to 50 µL of frozen worms and incubated at 62°C for 45 minutes, vortexing occasionally. Then, 426 427 we performed the PCI extraction in a phase lock gel tube (VWR) by adding 500 µL of PCI to the sample, vortexing, spinning for 5 minutes at 10,000 rpm, and then collecting the upper phase. 428 429 We repeated the PCI extraction step, and then extracted twice using chloroform. We precipitated 430 the DNA by adding 40 µL of 5M sodium acetate and 1 mL of ethanol, spinning for 5 minutes at 431 10,000 rpm, removing the supernatant, washing in 70% ethanol, and resuspending in 50 μ L of TE buffer at pH 7.4. 432

We created a sequencing library according to manufacturer instructions (Nextera DNA Library Prep Kit, Illumina). Briefly, this includes tagmentation of the DNA using the Illumina Tagment DNA Buffer and Enzyme, clean-up of the tagmented DNA using a Zymo DNA Clean and Concentrator Kit, PCR to add the library indices to the tagmented DNA, and PCR cleanup by gel extraction using a Qiagen MinElute Kit. We did a quality control step to confirm that the average length of DNA fragments was in the expected range of 300-500 bp using the Agilent

Bioanalyzer at the Stanford Protein and Nucleic Acid Biotechnology Facility. Sequencing wascompleted on an Illumina NextSeq sequencer in the Stanford Functional Genomics Facility.

We analyzed the sequencing data using the computing cluster of the Stanford Center for Genomics and Personalized Medicine. Briefly, we mapped reads using Bowtie2, used Picard to sort reads, mark duplicates, and prepare read groups, then used GATK to select high quality

444 SNPs and INDELs, and SnpEff to annotate the results.

445

446 *Code availability*

447 Two kinds of data were analyzed using custom code: image analysis (Python,

448 <u>https://github.com/anekimken/SSN_ImageAnalysis</u>) and whole-genome sequence analysis

449 available (scripts, <u>https://github.com/wormsenseLab/whole_genome_sequencing</u>).

450

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- 580
- 581

582 Figure Legends

583

584 Figure 1: Touch-induced mechanical strain measured using mitochondria as fiducial

- 585 markers in *C. elegans* TRNs *in vivo*.
- 586 A) Brightfield image of worm in the microfluidic device. Scale bar 20 µm. B) Maximum
- 587 projection image of TRN mitochondria before (magenta) and during (green) mechanical
- stimulus. Mitochondria that did not move appreciably appear white due to the overlap of
- 589 magenta and green. Image intensity and contrast was adjusted to improve visualization. Scale bar
- 590 20 μm. C) Displacement in the direction of actuation before (magenta) and during (green)
- stimulation. The smooth line (green) is a Gaussian fit used to infer the center of actuator. D)
- 592 Distribution of touch-induced strain in the TRN for a single actuation trial.

594 Figure 2: Indentation induces local longitudinal strain in *C. elegans* TRNs.

- 595 A) Three-dimensional diagram the positioning of the worm in the microfluidics trap, the animal-
- 596 centric coordinate system used to characterize strain, and the consequences of trapping animals
- 597 in different orientations. TRN is red, and muscles are included in brown as visual aid. B) Touch-
- induced displacement in three dimensions. C) Touch-induced strain, anterior is to the left and
- posterior is to the right. y=0 is the center of the actuator. n=61 trials from N=15 worms. D)
- 600 Mechanical strain at y=0 as a function of maximum neuron displacement. The line is a linear fit
- to the data: $\varepsilon = 0.0061 \text{ x} 0.0067$, where ε is the strain and x is the maximum displacement. The
- 602 shaded area indicates 95% confidence intervals of the fit. $R^2 = 0.04$.

603

604

605

Figure 3: Touch-induced mechanical strain profiles are similar in control animals and ECM mutants.

- A) Spatially averaged strain in control ALM neurons with normal TRN attachment (15 animals)
- 610 and *him-4(e1267)* with attachment defects (16 animals). The data for control animals are the
- 611 same data as Figure 2. B) Spatially averaged strain in *mec-1* mutants. 14 independent animals for
- 612 mec-1(1066) and 16 animals for mec-1(e1738). C) Spatially averaged strain in mec-5(u444)
- 613 mutants (17 animals). D) Overlay of all spatially averaged strain profiles. Smooth lines are the
- averages across all trials and shaded areas show the error (95% confidence intervals). One ALM
- 615 neuron tested in each animal.

617 Figure 4: Strain at the center of the actuator is similar in control animals and ECM

618 mutants

- A) Strain at the center of the actuator. Each point is the result of a single trial. The vertical lines
- 620 next to the swarm plots indicate the median and quartiles of the data. Data collected from 15, 16,
- 621 14, 16, and 17 animals (left to right). B) Difference in the mean strain between control and each
- 622 mutant with a bootstrapped resampled distribution of the data. Estimation plots generated using
- 623 the DABEST plotting package (Ho et al., 2019).

624

625 Table 1: Touch sensitivity of ECM mutants.

626

Genotype	Response	Mean difference from	95% confidence interval of mean
	probability	control	difference
control	0.742		
him-4(e1267)	0.527	-0.215	(-0.275, -0.163)
mec-1(e1066)	0.083	-0.659	(-0.698, -0.619)
mec-1(e1738)	0.237	-0.505	(-0.551, -0.461)
mec-5(u444)	0.208	-0.534	(-0.580, -0.487)

627 All mutants we tested were less sensitive to touch than control animals; effect size was smallest

628 for *him-4(e1267)* and largest for *mec-1(e1066)*. All strains tested contained the transgene

jsIs1073 that tags mitochondria in the TRNs with RFP. *N*=100 worms for each genotype, tested

630 in four independent cohorts of 25 blinded to genotype.

631

633

634 Table 2: Touch-evoked longitudinal strain in the TRNs as a function of genotype.

635

Genotype	Mean strain \pm	Mean	95% confidence	stimulation	animals
	SEM	difference	interval of mean	trials	
		from control	difference		
control	0.031 ± 0.005			61	15
him-4(e1267)	0.024 ± 0.005	-0.007	(-0.020, 0.007)	61	16
mec-1(e1066)	0.034 ± 0.007	0.003	(-0.012 0.022)	49	14
mec-1(e1738)	0.035 ± 0.005	0.004	(-0.009 0.018)	68	16
mec-5(u444)	0.026 ± 0.009	-0.005	(-0.028 0.013)	44	17

636 There was no detectable difference in the magnitude of the touch-evoked strain in control and

637 ECM mutant animals.

639 Table 3: *C. elegans* strains

Strain Name	Genotype	Source	
N2 (Bristol)	wild-type	CGC	
NM3573	jsIs1073	(Zheng et al., 2014)	
GN885	jsIs1073;him-4(e1267)	this study	
GN886	jsIs1073;mec-1(e1066)	this study	
GN887	jsIs1073;mec-1(e1738)	this study	
GN906	jsIs1073;mec-5(u444)	this study	

649 Figure S1: *him-4(e1267)* is likely to be a null allele. A) Map of *him-4* with *e1267* allele

annotated. B) Close-up of insertion in sequence as indicated by dotted lines in Panel A. C)

651 Sequences of both Indels found by sequencing. The first indel is likely to cause the null

652 phenotype because it causes a frameshift in an early exon, whereas the second indel is in a later

653 intron.

654

Figure S2: Quality control for image (z) stacks as a function of genotype. Eleven (11) image

656 stacks were collected from each TRN analyzed for touch-induced strain. This plot shows the

number of stacks that passed quality for each experiment. Note: The final stack was not used as

it was acquired in the resting configuration and could not be compared to a subsequent stack

659 collected in the indented configuration.

660

Figure S3: We tuned parameters of the particle tracking algorithm to account for variabilityacross images.

664 Table S1: Average number of detected mitochondria and distance between adjacent

665 detected mitochondria as a function of genotype.

666

Number of mitochondria							
Genotype	Mean number ± SEM	Mean difference from control	95% confidence interval of mean difference	Minimum	Maximum		
Control	8.9 ± 0.6			5	15		
him-4(e1267)	9.1 ± 0.5	0.2	(-1.5, 1.6)	6	13		
mec-1(e1066)	8.8 ± 0.5	-0.1	(-1.7, 1.5)	6	13		
mec-1(e1738)	8.9 ± 0.5	0.01	(-1.6, 1.5)	6	15		
mec-5(u444)	7.2 ± 0.5	-1.7 (-3.3, -0.3)		5	12		
Distance between adjacent mitochondria							
Genotype	Mean distance ± SEM (µm)	Mean difference from control (µm)	95% confidence interval of mean difference (μm)	Minimum (µm)	Maximum (µm)		
Control	26.0 ± 0.4			5.4	63.7		
him-4(e1267)	27.3±0.4	1.4	(-2.0, 4.8)	5.0	76.1		
mec-1(e1066)	27.0±0.5	1.0	(-2.6, 4.7)	5.3	65.1		
mec-1(e1738)	26.9±0.4	0.9	(-2.4, 4.3)	6.0	64.1		
<i>mec-5(u444)</i>	31.1±0.6	5.2	(1.2, 9.3)	5.5	84.1		

667

668 The number of detected mitochondria and distance between adjacent mitochondria in the ALM

669 neurons was similar in control and the ECM mutants, except for mec-5(u444) which had fewer

670 detected mitochondria and larger distances between adjacent mitochondria, on average. Number

671 of animals is the same as Table 2.

672







