

01 **Chapter 5**
02 **Mechanosensory Transduction in the Nematode**
03 ***Caenorhabditis elegans***
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07 **Nikos Kourtis and Nektarios Tavernarakis**
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13 **Abstract** Mechanotransduction, the process of converting a mechanical stimulus
14 into a biological signal, appeared very early in the evolution and underlies a plethora
15 of fundamental biological processes such as osmosensation, touch, hearing, bal-
16 ance and proprioception. Mechanosensory transduction has been studied extensively
17 in simple animal models such as the nematode *Caenorhabditis elegans* and the
18 fruit fly *Drosophila melanogaster*. Genetic and physiological studies have revealed
19 that specialized macromolecular complexes, encompassing mechanically gated ion
20 channels, play a critical role in the conversion of mechanical energy into cellu-
21 lar response. Members of two large ion channel families, the degenerin/epithelial
22 sodium channels (DEG/ENaC) and the transient receptor potential ion channels
23 (TRP), have emerged as candidate mechanosensitive channels. Several auxiliary
24 proteins associate with the core mechanosensitive channels to form the mechan-
25 otransducing apparatus in specialized mechanosensory cells. *C. elegans* displays a
26 variety of mechanosensory behaviours. In this chapter, we survey the mechanisms
27 of mechanosensory transduction in *C. elegans*. The exceptional amenability of this
28 simple metazoan to genetic and molecular manipulations has facilitated the dissec-
29 tion of the mechanotransduction process to unprecedented detail.
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31 **Key words:** Degenerin · Ion channels · Proprioception · Touch receptor neurons ·
32 TRP channels
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40 **5.1 Introduction**
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42 Animals receive and process information about their surroundings through
43 specialized sensory cells. Ubiquitous mechanical stimuli permeate the environment
44 of every living cell and every organism. The process by which cells convert mechan-
45 ical energy into electrical or chemical signals is called *mechanotransduction*. Be-
46 cause mechanical force is everywhere, mechanosensation probably represents one

01 of the oldest sensory transduction pathways that evolved in living organisms, from
02 bacteria to humans (Blount and Moe, 1999; French, 1992; Gillespie and Walker,
03 2001; Sackin, 1995) The capacity to respond and adjust to mechanical inputs plays
04 a pivotal role in numerous fundamental physiological phenomena such as the per-
05 ception of sound and gravity, which underlie our senses of hearing and balance
06 (Garcia-Anoveros and Corey, 1997; Hackney and Furness, 1995; Kellenberger and
07 Schild, 2002). Touch sensation and proprioception (the coordinated movement of
08 our body parts) are additional manifestations of responsiveness to mechanical stim-
09 ulation (Garcia-Anoveros and Corey, 1996; Tavernarakis and Driscoll, 1997; Tav-
10 ernarakis et al., 1997; Welsh et al., 2002). Moreover, mechanotransduction is equally
11 critical for the stretch-activated reflexes of vascular epithelia and smooth muscle,
12 and in the regulation of systemic fluid homeostasis and blood pressure (barore-
13 ception; Garcia-Anoveros and Corey, 1996; Lee and Huang, 2000; Tavernarakis
14 and Driscoll, 2001a; Tavi et al., 2001; Welsh et al., 2002). Mechanotransduction
15 is also important for the prevention of polyspermy during fertilization, cell volume
16 and shape regulation, cell locomotion and tissue development and morphogenesis
17 (Ingber, 1997; Ko and McCulloch, 2001; Rossier et al., 1994). In plants, mechan-
18 otransduction is the basis of gravitaxis and turgor control (Lynch et al., 1998;
19 Pickard and Ding, 1993). In protists (*Paramecium*, *Stentor*) mechanotransduction
20 underlies gravikinesis (the swimming against the gravity vector in order to avoid
21 sedimentation; Block et al., 1999; Gebauer et al., 1999; Hemmersbach et al., 2001;
22 Marino et al., 2001).

23 All living organisms have developed highly specialized structures that are re-
24 ceptive to mechanical forces originating either from the surrounding environment
25 or from within the organism itself. Mechanotransducers are among the most elab-
26 orate and efficient, such structures, which are responsible for sensory awareness,
27 for example, those facilitating touch, balance proprioception and hearing (Garcia-
28 Anoveros and Corey, 1997; Gillespie and Walker, 2001; Hackney and Furness,
29 1995; Tavernarakis and Driscoll, 2001a). The mechanisms underlying the capability
30 of living cells to receive and act in response to mechanical inputs are among the most
31 anciently, implemented during evolution. Proteins with mechanosensitive properties
32 are ubiquitously present in eubacteria, archaea and eukarya, and are postulated to
33 have been an essential part of the physiology of the Last Universal Ancestor (Kloda
34 and Martinac, 2001; Koch, 1994; Koprowski and Kubalski, 2001; Martinac, 2001).
35 The first mechanosensitive processes may have evolved as backup mechanisms for
36 cell protection, e.g. to reduce intracellular pressure and membrane tension during
37 osmotic swelling. Subsequent organismal diversification and specialization resulted
38 in variable requirements for mechanotransduction in different organisms (Norris
39 et al., 1996). Hence, evolutionary pressure has shaped a large repertoire of mechan-
40 otransducers, optimized for a great assortment of tasks that range from maintenance
41 of intracellular osmotic balance and pressure, to our impressive ability of hearing
42 and discriminating sounds, and reading Braille code with our fingertips (Gillespie
43 and Walker, 2001; Hamill and Martinac, 2001).

44 In this chapter, we describe *C. elegans* mechanosensory behaviours and survey
45 the genes implicated in mechanosensory perception. We also discuss the relevant
46 mechanisms underlying mechanotransduction in the nematode.

5.2 *C. elegans*: Background Information

Caenorhabditis elegans is a small (about 1mm length), hermaphroditic, soil-dwelling nematode. The size and simple dietary demands permit easy and cheap cultivation of the animal in the lab. The worm completes a reproductive life cycle in 2.5 days at 25°C progressing from a fertilized embryo through four larva stages, to become an egg-laying adult which lives for about 2–3 weeks (Brenner, 1974). Under non-favourable conditions such as starvation, high temperature or overcrowding, larvae may enter an alternative life stage, called dauer larva, a very resistant larval form that survives for months (Golden and Riddle, 1984). The simple body plan, the transparent egg and cuticle, and the nearly invariant developmental plan of this nematode has facilitated exceptionally detailed developmental and anatomical characterization of the animal (Ward et al., 1975; White et al., 1986; www.wormatlas.org). The complete sequence of somatic cell divisions from the fertilized egg to the 959-cell adult hermaphrodite has been described (Sulston and Horvitz, 1977). The *C. elegans* nervous system consists of 302 neurons of 118 types that interconnect in a stable and reproducible manner to create a variety of neural circuits (White et al., 1976; White et al., 1986). Individual neurons can be ablated through laser microsurgery and genetic manipulation, and neuronal pathways responsible for different behaviours have been characterized.

C. elegans is especially amenable to both forward and reverse genetic analysis. Mutagenized parents segregate homozygous F2 mutant progeny without any requirement for genetic crossing. Self-fertilization of heterozygotes leads to genetically homogeneous populations, while crossing with males that appear at a small percentage in the population facilitates genetic manipulations. Rapid and precise genetic mapping can be achieved, by taking advantage of a dense single nucleotide polymorphism map (Koch et al., 2000; Wicks et al., 2001). A physical map of the *C. elegans* genome, consisting of overlapping cosmid and YAC clones covering most of the six chromosomes, has been constructed to facilitate cloning of genes that have been positioned on the genetic map (Coulson et al., 1988; Waterston and Sulston, 1995). Double-stranded RNA mediated interference (dsRNAi) enables probable loss-of-function phenotypes to be readily evaluated (Fire et al., 1998; Tavernarakis et al., 2000). Transgenic animals carrying the gene of interest can be easily constructed by microinjecting the transgene into the gonad of hermaphrodite animals (Mello and Fire, 1995).

C. elegans displays a variety of behaviours, including mechanosensitive behaviours. The broad range of genetic and molecular tools available in the worm facilitates thorough and multifaceted investigation of the pathways that govern these behaviours.

5.3 *C. elegans* Mechanosensory Behaviours

Many behaviours displayed by *C. elegans* are direct manifestations of mechanosensitivity, making it exceptionally attractive for investigating mechanotransduction (Bargmann and Kaplan, 1998; Baumeister and Ge, 2002; Chalfie, 1993; Chalfie

01 and Sulston, 1981; Chalfie et al., 1985; Driscoll and Tavernarakis, 1997; Herman,
02 1996; Kaplan and Horvitz, 1993; Mah and Rankin, 1992; O'Hagan and Chalfie,
03 2006; Rankin, 1991; Syntichaki and Tavernarakis, 2004; Tavernarakis and Driscoll,
04 1997; Wicks and Rankin, 1995; Wolinsky and Way, 1990). The best characterized
05 such behaviour is the response to a gentle mechanical stimulus delivered trans-
06 versely along the body of the animal, typically by means of an eyelash hair at-
07 tached onto a toothpick (the 'gentle body touch response'; (Chalfie and Sulston,
08 1981; Chalfie et al., 1985; Herman, 1996; Tavernarakis and Driscoll, 1997)). We
09 discuss studies elucidating the molecular mechanisms of this touch response in the
10 following section. Other mechanosensory responses are the generation and mainte-
11 nance of the characteristic coordinated sinusoidal pattern of locomotion (analogous
12 to proprioception; (Li et al., 2006; Tavernarakis and Driscoll, 1997; Tavernarakis
13 et al., 1997); see below), and the nose touch response, which can be further catego-
14 rized into the head-on collision response and the head withdrawal response (Driscoll
15 and Kaplan, 1996; Kaplan and Horvitz, 1993).

16 When animals collide with an obstacle in a nose-on fashion during the course of
17 normal locomotion they respond by reversing their direction of movement
18 (Bargmann and Kaplan, 1998; Colbert et al., 1997). This response is independent
19 of touch receptor neurons, needed for gentle touch (Chalfie and Sulston, 1981).
20 Three classes of mechanosensory neurons, ASH, FLP, and OLQ, mediate this avoid-
21 ance response (Bargmann and Kaplan, 1998; Herman, 1996; Kaplan and Horvitz,
22 1993; Wicks and Rankin, 1995). Each of these sensory neurons accounts for a
23 part of the normal response, which is quantitative with normal animals respond-
24 ing about 90% of the time. Laser ablation and genetic studies have demonstrated
25 that each sensory neuron contributes to the overall responsiveness as follows: ASH,
26 45%; FLP, 29%; and OLQ, 5%. The remaining 10% of the responses are medi-
27 ated by the ALM and AVM neurons, which sense anterior body touch (Driscoll
28 and Kaplan, 1996; Kaplan and Horvitz, 1993). It is unclear what distinguishes the
29 function of the three nose touch neurons. One attractive possibility is that these
30 cells differ in their sensitivities and that the intensities of nose touch stimuli vary
31 according to the violence of the collision. If this were the case, it would be ex-
32 pected that the most sensitive neuron (ASH) would account for the majority of
33 responses while less sensitive neurons (FLP and OLQ) would account for the re-
34 mainder. In addition to their mechanosensory properties, the ASH neurons are part
35 of a chemosensory organ, the amphid sensilla, with their sensory endings exposed to
36 the external environment (Perkins et al., 1986; Ward et al., 1975). The ASH neurons
37 serve chemosensory and osmosensory functions, mediating avoidance of osmotic
38 repellents (Hart et al., 1999; Hart et al., 1995). Several classes of chemosensory
39 neurons respond to multiple chemical stimuli in *C. elegans*. However, ASH is unique
40 among them in responding to such divergent stimuli. In this respect, ASH neu-
41 rons are similar to vertebrate neurons that sense painful stimuli, which are called
42 nociceptors. For their multi-sensory capabilities, the ASH neurons have been cat-
43 egorized as polymodal sensory neurons (Driscoll and Kaplan, 1996; Kaplan and
44 Horvitz, 1993).

45 In addition to DEG/ENaC proteins, another major family of channel proteins
46 implicated in sensory mechanotransduction is the transient receptor potential (TRP)

01 ion channels. TRPs are non-specific cation-permeable channels that are present
02 in diverse species ranging from yeast, flies, and worms to humans (Kahn-Kirby
03 and Bargmann, 2006). All TRP channels appear to form tetrameric assemblies and
04 include six predicted transmembrane domains and a variable number of ankyrin
05 motifs, which are suggested to mediate protein-protein interactions. Members of
06 individual subfamilies may bear several other domains, such as coiled-coil motifs,
07 protein kinase domains, transmembrane segments, and TRP domains
08 (Montell, 2005).

09 In *C. elegans*, the TRPV (vallinoid TRP) subfamily genes *osm-9* and *ocr-2*
10 are required for the aversive responses of ASH neurons to various noxious stimuli,
11 including high osmolarity and noxious chemicals (Bargmann et al., 1990;
12 Hilliard et al., 2005; Troemel et al., 1995). The OSM-9 and OCR-2 proteins localize
13 to the sensory cilia of ASH, suggesting a direct role in sensory transduction
14 (Colbert et al., 1997). Additionally, OSM9::GFP is expressed in FLP, OLQ
15 and PVD mechanosensory neurons. Several genetic studies suggest that the function
16 of the putative OSM-9/OCR-2 ion channel is regulated by G protein signalling and
17 specific polyunsaturated fatty acids (PUFAs), which act upstream of
18 OSM-9/OCR-2 to modulate nociceptive responses in ASH neurons, including the
19 mechanosensory nose touch avoidance behaviour (Kahn-Kirby et al., 2004; Roayaie
20 et al., 1998). Expression of a mammalian TRPV4 protein in ASH can rescue
21 defects in osmotic and nose-touch avoidance in *osm-9* mutants (Liedtke et al.,
22 2003).

23 *C. elegans* also distinguishes textural differences in its substrate. When worms
24 enter a bacterial lawn they slow their movement, a behaviour known as basal
25 slowing. This response is indeed mechanosensory since worms entering a lawn
26 of Sephadex beads instead of bacteria, slow similarly (Sawin et al., 2000). This
27 behaviour depends on the CEP, ADE and PDE dopaminergic neurons which have
28 ciliated sensory endings, embedded in the cuticle (Perkins et al., 1986; Ward et al.,
29 1975; White et al., 1986). Laser ablation experiments confirmed that these neurons
30 are required for response to small particles (Sawin et al., 2000). In support of this
31 model, dopaminergic neurons transduce an inhibitory signal to the motor circuit
32 (White et al., 1986).

33 The complicated male-mating behaviour of *C. elegans* is probably based on
34 chemosensory and mechanosensory cues (Liu and Sternberg, 1995). Males have 87
35 additional neurons, comparing with the hermaphrodite many of which are ciliated
36 and considered to be sensory (Sulston et al., 1980). Two genes, *lov-1* and *pkd-2*,
37 needed for male mating have been implicated in mechanical signalling. *lov-1* and
38 *pkd-2*. LOV-1 and PDK-2 are the nematode homologs of mammalian PDK-1 and
39 PDK-2 TRPP ion channels respectively (Corey, 2003). Interestingly, mutations in
40 the mammalian PDK-1 and PDK-2 cause autosomal dominant polycystic kidney
41 disease (ADPKD). PDK-1 and PDK-2 form a Ca^{2+} -permeable ion channel which
42 is mechanically activated by fluid flow in certain epithelial cells (Nauli et al., 2003).

43 An additional mechanosensitive behaviour in *C. elegans* is the tap withdrawal
44 reflex, where animals retreat in response to a tap on the culture plate (Chiba and
45 Rankin, 1990; Rankin et al., 2000; Wicks and Rankin, 1997). Worms respond to a
46 diffuse mechanical stimulus (a tap to the side of the dish they are resting on) by

01 either accelerating forward movement or by initiating backward movement (Chiba
02 and Rankin, 1990; Rankin et al., 2000). Given that the stimulus is not spatially
03 coherent and that the animal's response is variable, it was proposed that the tap
04 response reflects the simultaneous activation of the anterior and posterior touch
05 cells. The behavioural outcome is likely determined by the integration of these two
06 antagonistic circuits.

07 Mechanotransduction appears to also play a regulatory role in processes such
08 as egg laying, feeding, defecation, and maintenance of the pseudocoelomic body
09 cavity pressure (Avery, 1993; Du and Chalfie, 2001; Liu and Thomas, 1994; Liu and
10 Sternberg, 1995; Thomas et al., 1990; Wolinsky and Way, 1990). These behaviours
11 add to the large repertoire of mechanosensitive phenomena, amenable to genetic and
12 molecular dissection in the nematode (Bargmann and Kaplan, 1998; Syntichaki and
13 Tavernarakis, 2004; Tavernarakis and Driscoll, 2001b).

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16 **5.4 The Gentle Body Touch Transduction System**

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19 In its natural habitat, the soil, *C. elegans* encounters a large number of mechanical
20 stimuli. While crawling on surfaces of soil particles, the worm receives external
21 forces generated by bumping on soil materials and other animals. The laboratory
22 assay for the gentle body touch response involves a mild stroke of the animal with
23 an eyelash hair attached to a toothpick, transversely to the anterior-posterior body
24 axis (Chalfie and Sulston, 1981; Chalfie et al., 1985; Syntichaki and Tavernarakis,
25 2004). When no response is observed, animals are prodded with a thin platinum
26 wire to confirm that they are touch insensitive rather than paralyzed (gentle-touch
27 insensitive animals typically still respond to a strong stimulus-the harsh touch re-
28 sponse) (Chalfie et al., 1985; Chalfie and Wolinsky, 1990; Driscoll and Kaplan,
29 1996; Wolinsky and Way, 1990). Depending on the part of the body touched, an-
30 imals will either accelerate or initiate forward movement (when stimulated at the
31 posterior or the tail), or reverse and move backwards (when stimulated at the ante-
32 rior part of the body). Hermaphrodite, male, juvenile (except L1), and dauer animals
33 respond identically to touch. The response is adaptive: repetitive stimulation leads
34 to short periods of insensitivity (Mah and Rankin, 1992; Rankin et al., 2000; Wicks
35 and Rankin, 1995).

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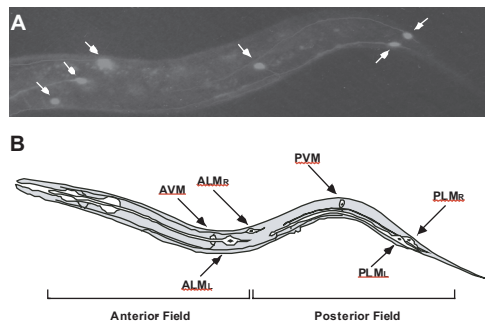
38 **5.4.1 The Sensory Cells**

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41 The touch reflex of the mature animal involves six touch receptor neurons, 5 pairs
42 of interneurons and 69 motorneurons (Chalfie, 1993; Chalfie et al., 1985). The six
43 touch receptor neurons share common ultrastructural features and express many of
44 the same genes (Chalfie and Sulston, 1981; Chalfie and Thomson, 1979; Chalfie
45 and Thomson, 1982). The six touch receptor neurons were originally designated as
46 the microtubule cells because of distinctive bundles of 15-protofilament (pf; tubulin

01 dimmer filaments) microtubules that fill their processes (ALML/R: anterior lateral
 02 microtubule cell, left/right; AVM: anterior ventral microtubule cell; PLML/R; pos-
 03 terior lateral microtubule cell, left/right; and PVM: posterior ventral microtubule
 04 cell; (Chalfie, 1993; Chalfie et al., 1986; Chalfie et al., 1985; Chalfie and Thomson,
 05 1979; Chalfie and Thomson, 1982)). Two fields of touch sensitivity, anterior and
 06 posterior are defined by the arrangement of the six touch receptor neurons along
 07 the body axis (Fig. 5.1; Chalfie et al., 1985; Tavernarakis and Driscoll, 1997).
 08 All six cells have anteriorly directed processes and, except for PVM, an anterior
 09 branch. The processes of touch neurons are localized in the hypodermis, just be-
 10 neath the cuticle, an ideal position for sensing external stimuli and vibrations. All
 11 six cells are dispensable for the viability of the organism. Apart from insensitiv-
 12 ity to gentle body touch, laser ablation of all six neurons does not result in any
 13 additional adverse effects (Chalfie, 1995; Chalfie et al., 1985; Tavernarakis and
 14 Driscoll, 1997).

15 Laser microsurgery established that PLML and PLMR are required for response
 16 to a touch to the tail. If either is present, tail touch sensitivity is observed. When
 17 both are ablated, animals are completely insensitive to gentle touch stimuli ad-
 18 ministered to the posterior (Chalfie et al., 1985; Chalfie et al., 1983; Kitamura
 19 et al., 2001). Either ALML or ALMR can mediate a response to a mechanical
 20 stimulus delivered to the anterior part of the body. AVM, which is added into the
 21 touch circuitry postembryonically, can mediate a weak response to some stimu-
 22 li but not all. In animals in which both ALM cells are killed, partial touch
 23 sensitivity returns 35–40 hours after hatching, which is attributable to AVM being
 24 generated. PVM alone does not produce a touch response, but its synaptic
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 38 **Fig. 5.1** The *C. elegans* touch receptor neurons. (A) Visualization of touch receptors. Worms
 39 are expressing the Green Fluorescence Protein (GFP) under the control of the *mec-4* promoter,
 40 which is active only in the six-touch receptor neurons. Arrows indicate the cell bodies of the
 41 neurons. (B) Schematic diagram, showing the position of the six touch receptor neurons in the
 42 body of the adult worm. The two fields of touch sensitivity are defined by the arrangement of touch
 43 receptor neurons along the body axis. The ALMs and AVM mediate the response to touch over the
 44 anterior field whereas PLMs mediate the response to touch over the posterior field. PVM does not
 45 mediate touch response by itself (Chalfie, 1995; Chalfie, 1997; Chalfie et al., 1985; Tavernarakis
 46 and Driscoll, 1997). Reproduced from (Voglis and Tavernarakis, 2005) with copyright permission
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01 pattern implicates it in the touch behaviour (Chalfie, 1993; Chalfie and Au, 1989;
02 Chalfie et al., 1985).

03 Bundles of darkly staining large diameter microtubules distinguish the touch
04 receptor neurons (Chalfie and Thomson, 1979; White et al., 1976). Cross bridges be-
05 tween microtubules of a bundle are observed in micrographs obtained with electron
06 microscopy, and may increase the structural integrity of the bundle. These micro-
07 tubules are unique to the nematode touch receptor neurons and contain
08 15-protofilament microtubules, a unique feature of these six cells (Chalfie and
09 Thomson, 1982; Tavernarakis and Driscoll, 1997). In most eukaryotic cells, α - and
10 β -tubulin co-assemble into 13-protofilament microtubules, whereas the vast major-
11 ity of microtubules in *C. elegans* cells have 11-protofilament (Chalfie and Thomson,
12 1982; White et al., 1986). In normal touch receptors, 11-protofilament
13 microtubules typical of most other cells in this nematode are occasionally observed.
14 If the 15-protofilament microtubules are eliminated by mutation, the number of
15 11-protofilament microtubules in the touch cell processes increases (Chalfie, 1993;
16 Chalfie et al., 1986; Chalfie and Thomson, 1979; Fukushige et al., 1999; Savage
17 et al., 1989). Individual microtubules are 10–20 μm , but overlap and create bundles,
18 filling in that way the process of the neuron which is 400–500 μm long (Chalfie
19 and Thomson, 1979). The distal ends of the microtubules are apposed to the plasma
20 membrane and are associated with structures of a diameter up to twice that of the
21 microtubules (Chalfie and Thomson, 1979).

22 Touch receptors also are uniquely surrounded by an osmiophillic extracellular
23 material referred to as the mantle. The amount of mantle varies along the length of
24 the process (Chalfie and Sulston, 1981). The mantle is needed for the attachment
25 of the touch receptor process to the body wall, bringing it to an ideal position for
26 detection of external mechanical stimuli.

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32 **5.4.2 The *C. elegans* Mechanosensory Apparatus**

33 To identify molecules dedicated to touch transduction, Martin Chalfie and colleagues
34 mounted a forward genetics approach to isolate gentle body touch-insensitive ne-
35 matode mutants (Chalfie and Au, 1989; Chalfie et al., 1986; Chalfie and Sulston,
36 1981; Du and Chalfie, 2001; Gu et al., 1996; Tavernarakis and Driscoll, 1997).
37 Briefly, populations of wild type, touch sensitive animals were mutagenized and
38 touch insensitive individuals were sought among their descendants by stroking with
39 an eyelash hair and prodding with a platinum wire (Chalfie, 1997). During the course
40 of this very tedious screening process, over 417 mutations in 17 different genes,
41 randomly distributed in all six chromosomes of *C. elegans* were isolated (Driscoll
42 and Kaplan, 1996; Tavernarakis and Driscoll, 1997). By design, the screen yields
43 mutations in genes that are fairly specific for normal gentle body touch perception.
44 For example, gene mutations with pleotropic effects that result in lethality or unco-
45 ordinated and paralyzed phenotypes would have been missed. In addition to being
46 touch insensitive *mec* mutants tend to be lethargic when grown normally in the pres-
ence of amble food (Driscoll and Kaplan, 1996). Reduced spontaneous movement

01 is probably due to their inability to sense micro vibrations in their environment,
02 interaction with external objects or stretch produced by the locomotory movements
03 themselves. However, when starved or during mating they move as well as wild type.
04 The 17 genes isolated are designated as the *mec* genes for their ‘mechanosensory
05 abnormal’ phenotype. Corroborating the high specificity of the screen, while most
06 of the alleles generated cause complete touch insensitivity, only few other ab-
07 normalities accompany the mutants (Driscoll and Kaplan, 1996). Depending on
08 their role and point of action, *mec* genes can be loosely classified into three main
09 categories. First, the regulatory/specification genes which control the expression
10 touch receptor neuron specific genes or modify the activity of the mechanotrans-
11 ducer complex; second, the *mec* genes encoding core structural components of the
12 mechanosensitive ion channel; and third the genes encoding peripheral, associated
13 proteins.

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16 **5.4.2.1 Genes Needed for the Development of Touch Receptor Neurons**

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18 The UNC-86 and MEC-3 transcription factors are essential for proper development
19 and differentiation of the six touch receptor neurons (Chalfie and Sulston, 1981;
20 Duggan et al., 1998; Way and Chalfie, 1988; Way and Chalfie, 1989; Xue et al.,
21 1992). UNC-86 is a POU domain protein which is required in several distinct neu-
22 roblast lineages for daughter cells to become different from their mothers (Finney
23 and Ruvkun, 1990; Finney et al., 1988). UNC-86 activates the expression of *mec-3*
24 which encodes a LIM-type homeodomain protein needed for the differentiation of
25 the six touch receptor cells (Way and Chalfie, 1989). In mutants lacking *mec-3*
26 activity, the touch receptors express none of their unique differentiated features
27 and appear to be transformed to other types of neurons (Way and Chalfie, 1988).
28 UNC-86 and MEC-3 bind cooperatively as a heterodimer to the *mec-3* promoter
29 as well as to the promoters of other genes required for the function of touch cells
30 (Duggan et al., 1998; Way and Chalfie, 1988; Way and Chalfie, 1989). *unc-86* and
31 *mec-3* genes are also expressed in the PVD and FLP neurons which also function
32 as mechanoreceptors but they do not express the same genes as the touch receptor
33 neurons (Way and Chalfie, 1989).

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36 **5.4.2.2 Genes Needed for Function of Touch Receptor Neurons**

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38 Four *mec* genes can be classified in the category of core structural components of
39 the putative mechanosensory ion channel in touch receptor neurons, *mec-2*, *mec-4*,
40 *mec-6* and *mec-10*. MEC-4 and MEC-10 form the core ion channel, while MEC-
41 2 and MEC-6 physically interact with the channel subunits to shape and modu-
42 late their gating properties. Animals bearing loss-of-function mutations in *mec-4*
43 or *mec-10* are touch-insensitive despite the fact that in these mutant backgrounds
44 the touch receptor neurons develop normally and exhibit no apparent defects in ul-
45 tra structure (Driscoll and Chalfie, 1991; Huang and Chalfie, 1994; Tavernarakis
46 and Driscoll, 1997). *mec-4* and *mec-10* encode homologous proteins related to

01 subunits of the multimeric amiloride sensitive Na^+ channel which mediates Na^+
02 re-absorption in vertebrate kidney, intestine and lung epithelia (the ENaC channel;
03 (Kellenberger and Schild, 2002; Rossier et al., 1994)). In addition to being involved
04 in mechanotransduction, MEC-4, MEC-10 and several other related nematode pro-
05 teins have a second, unusual property: specific amino acid substitutions result in
06 aberrant channels that induce the swelling and subsequent necrotic death of the
07 cells in which they are expressed (Driscoll, 1996; Driscoll and Chalfie, 1991; Hall
08 et al., 1997; Harbinder et al., 1997). This pathological property is the reason that
09 this family of proteins was originally called degenerins (Chalfie et al., 1993; Chalfie
10 and Wolinsky, 1990; Tavernarakis and Driscoll, 2001a; Tavi et al., 2001). *C. elegans*
11 degenerins, together with their mammalian relatives, the ENaCs, comprise the large
12 DEG/ENaC family of ion channels (Fig. 5.2). The relationship of these channel
13 subunits to subunits of an amiloride-sensitive ENaCs is intriguing because amiloride
14 is a general inhibitor of mechanosensitive ion channels (Alvarez de la Rosa et al.,
15 2000; Hamill et al., 1992; Hamill and McBride, 1996; Hoyer et al., 1997; Lane et al.,
16 1991; Rossier et al., 1994; Voilley et al., 1997).

17 *mec-4* is expressed solely in the six touch receptor neurons, while *mec-10* in ad-
18 dition to the six touch receptor neurons, is expressed in two other neuron pairs that
19 may mediate stretch-sensitive responses (FLPs and PVDs; (Driscoll and Chalfie,
20 1991; Driscoll and Tavernarakis, 1997; Huang and Chalfie, 1994; Tavernarakis and
21 Driscoll, 1997)). Interestingly, a MEC-4::GFP fusion localizes in distinct puncta
22 along the processes of the touch receptor neurons (Fig. 5.3). Such punctuate lo-
23 calization may reflect the distribution of mechanotransducing complexes on the
24 axon of the touch receptor neuron. MEC-4 and MEC-10 co-localize exclusively in
25 mechanosensitive neurons, where they may co-assemble into a mechanically-gated
26 ion channel. Optical imaging studies using genetically encoded Ca^{2+} indicators
27 (cameleons), which monitor intracellular Ca^{2+} changes in response to gentle body
28 touch support this hypothesis (Suzuki et al., 2003). More definitive proof is provided
29 by recent electrophysiological, whole-cell voltage-clamp recordings of mechanore-
30 ceptor currents (MRCs) from the PLM touch receptors *in vivo* (O'Hagan et al.,
31 2005). The mechanoreceptor currents are extremely rapid; they turned on within
32 a millisecond of force application and quickly decreased in amplitude, character-
33 istic of adaptation. MRCs recorded from PLM neurons are probably representing
34 transduction by direct activation of the channel and not through second messengers.
35 MRCs are absent in MEC-4, MEC-2 and MEC-6 null mutants, whereas mechanore-
36 ceptor current was only partially decreased when testing mutations in non-conserved
37 regions of the proteins (O'Hagan et al., 2005). In addition, MRCs are reduced but not
38 eliminated in worms that carry a null mutation in *mec-7* and *mec-12* genes, affect-
39 ing α and β -tubulin which forms the 15-protofilament microtubule bundles. These
40 results question the absolute requirement of microtubules for touch transduction.

41 MEC-2 is an additional component of the channel complex. *mec-2* encodes a
42 481-amino acid protein and is expressed in the touch receptor neurons and in a
43 few additional neurons in the nerve ring region (Du and Chalfie, 2001; Gu et al.,
44 1996; Huang et al., 1995). MEC-2 features three candidate protein-protein interac-
45 tion domains (Fig. 5.4). First, part of the amino-terminal domain (situated in part
46 between AA 42–118) is needed for the proper localization of a *mec-2/lacZ* fusion

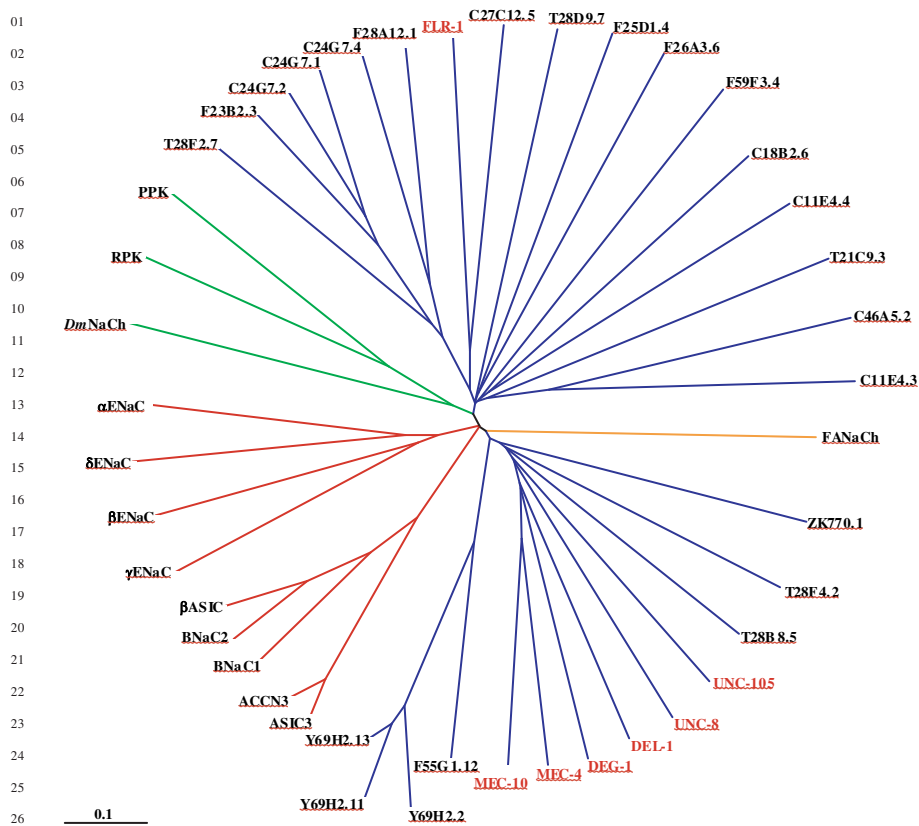


Fig. 5.2 Phylogenetic relationships among DEG/ENaC proteins. Nematode degenerins are shown with blue lines. The current degenerin content of the complete nematode genome is included. The seven genetically characterized (DEG-1, DEL-1, FLR-1, MEC-4, MEC-10, UNC-8 and UNC-105) are shown in red. Representative DEG/ENaC proteins from a variety of organisms, ranging from snails to humans, are also included (mammalian: red lines; fly: green lines; snail: orange line). The scale bar denotes relative evolutionary distance equal to 0.1 nucleotide substitutions per site (Sadoshima et al., 1992). Reproduced from (Voglis and Tavernarakis, 2005) with copyright permission of the Academia Publishing House Ltd

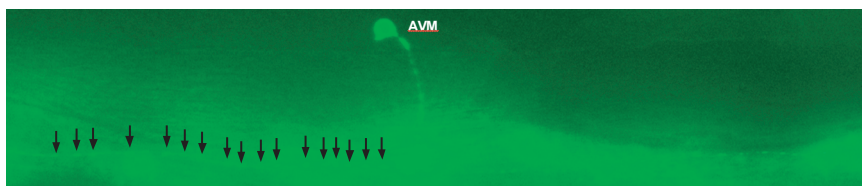


Fig. 5.3 Punctuate localization of a putative mechanosensitive ion channel subunit. Image of an AVM touch receptor neuron expressing a GFP-tagged MEC-4 protein. Fluorescence is unevenly distributed along the process of the neuron in distinct puncta, which may represent the location of the mechanotransducing apparatus. Reproduced from (Voglis and Tavernarakis, 2005) with copyright permission of the Academia Publishing House Ltd

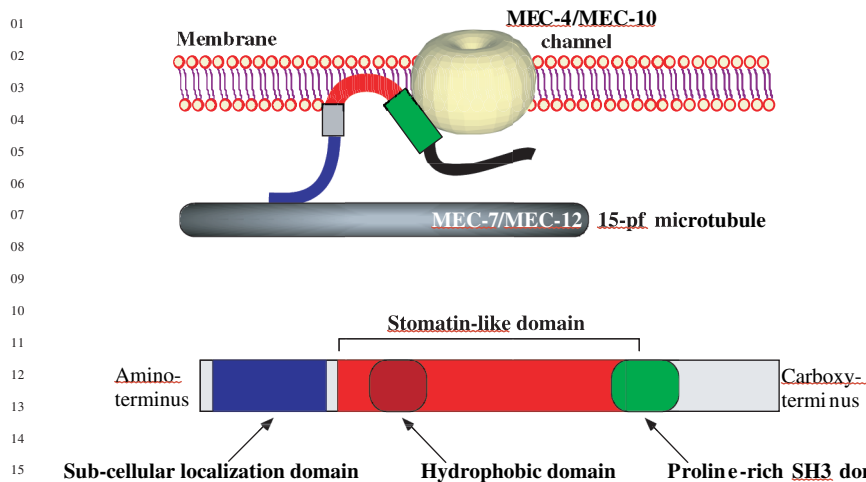
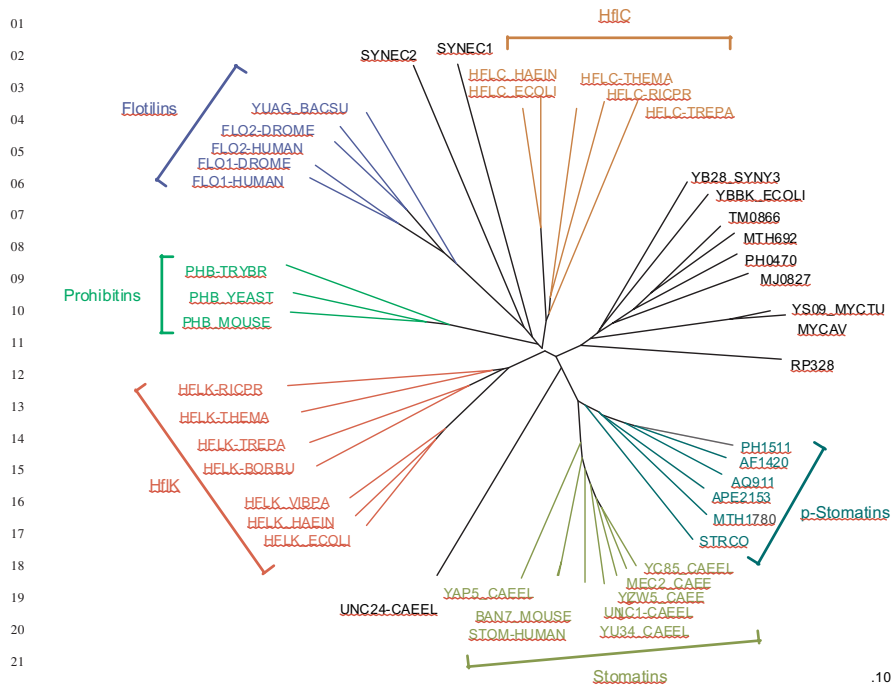


Fig. 5.4 Schematic representation and topology of the MEC-2 protein. Conserved domains as well as hydrophobic regions are highlighted. Putative interactions with the degenerin channel and the cytoskeleton are indicated (Goodman et al., 2002). Reproduced from (Voglis and Tavernarakis, 2005) with copyright permission of the Academia Publishing House Ltd

protein to the touch receptor process. Second, the carboxy-terminal domain includes a proline-rich region that is similar to SH3-binding domains. Third, the central region (AA 114–363) encompasses an SPFH domain with a membrane-associated hydrophobic part (AA 114–141) and a cytoplasmic hydrophilic part that together exhibit 65% identity to the human red blood cell protein stomatin (Huang et al., 1995; Tavernarakis et al., 1999). The SPFH domain is the common denominator of stomatins, prohibitins, flotilins and bacterial *HflK/C* proteins, all of which are membrane associated regulators (Fig. 5.5; Tavernarakis et al., 1999). Stomatin, also known as band 7.2b protein, is a membrane-associated protein originally identified as a component of human red cells (Delaunay et al., 1999; Sedensky et al., 2001; Snyers et al., 1998; Stewart, 1997; Stewart et al., 1993). In humans, stomatin is missing from erythrocyte membranes in autosomal dominant hemolytic disease overhydrated hereditary stomatocytosis, despite an apparent normal stomatin gene. Many of the 54 mutant *mec-2* alleles have dominant effects and exhibit a complex pattern of inter-allelic complementation (Chalfie and Sulston, 1981; Gu et al., 1996), indicating that MEC-2 protein molecules form higher order complexes. However, there is also genetic data suggesting that MEC-2 interacts with the specialized touch cell microtubules encoded by *mec-7* and *mec-12* (α -tubulin and β -tubulin respectively; (Gu et al., 1996; Huang et al., 1995)). Normally, a *mec-2::lacZ* fusion protein is distributed along the touch receptor axon (Huang et al., 1995). The axonal distribution of a MEC-2::*lacZ* fusion protein is mildly disrupted in a *mec-7* null or *mec-12* strong loss-of-function background, implying that the 15-protofilament microtubules are not essential for the localization of MEC-2 to the neuronal process. However, two specific *mec-12* missense alleles interfere dramatically with localization of MEC-2 fusion proteins, restricting the fusion proteins to the cell body (Huang et al., 1995).



23 **Fig. 5.5** Phylogenetic relations among SPFH domain proteins. A dendrogram showing distance relationships among most of the stomatin protein super-family members (the complete ClustalW generated alignment on which the dendrogram was based is available at <http://www.imbb.forth.gr/worms/worms/alignment.gif>). The dendrogram was constructed with the neighbor-joining method based on pairwise distance estimates of the expected number of amino acid replacements per site (0.10 in the scale bar), and visualized by TreeTool. Protein sub-families are denoted in different colours (Tavernarakis et al., 1999)

31 MEC-2 colocalizes with MEC-4 in the six touch receptor neurons and is distributed
32 along neuronal processes in punctuate pattern (Zhang et al., 2004). This is consistent
33 with the co-immunoprecipitation of the two proteins in *Xenopus* oocytes (Goodman
34 et al., 2002). The stomatin-like domain of MEC-2 interacts specifically with the
35 N-terminus cytoplasmic region of MEC-4 (Zhang et al., 2004). Punctuate expression
36 of MEC-2 is disrupted in the *mec-4(u253)*, *mec-6(u450)* and *mec-10(u20)* loss-of-
37 function mutants indicating that the MEC-2 subcellular localization depends on the
38 other partners of the mechanosensory complex (Zhang et al., 2004). These genetic
39 studies, which do not by themselves prove a direct interaction, have recently been
40 complemented by elegant heterologous expression experiments in *Xenopus* oocytes
41 that support physical interaction between MEC-2 and the channel subunits MEC-4
42 and MEC-10 (Goodman et al., 2002). Reconstitution of channel activity in *Xenopus*
43 oocytes revealed that MEC-2 regulates the activity of the MEC-4/MEC-10 channel,
44 providing the first direct support for the hypothesis that stomatin-like proteins in-
45 teract with and regulate ion channels (Goodman et al., 2002; Stewart et al., 1993).
46 This interaction appears to dramatically potentiate the conductivity of the channel in

01 oocytes. Co-expression of MEC-2 with the hyperactive MEC-4(d) and MEC-10(d)
02 derivatives in *Xenopus* oocytes resulted in about 40-fold increase in the amplitude
03 of amiloride-sensitive ionic currents, and this amplification allowed currents to be
04 detected even with wild-type MEC-4 and MEC-10 proteins (Goodman et al., 2002;
05 Stewart et al., 1993). Visualization of tagged MEC-4(d) and MEC-10(d) in live
06 oocytes demonstrated that MEC-2 does not increase the number of MEC-4(d)/MEC-
07 10(d) channels that reach the plasma membrane, and probably acts by regulating
08 their activity. In *mec-2(u37)* loss-of-function mutants the touch-evoked currents are
09 abolished confirming that MEC-2 is one of the major components needed for the
10 proper function of the MEC-4/MEC-10 ion channel (O'Hagan et al., 2005).

11 A second stomatin-like protein, UNC-24 appears to be part of the channel com-
12 plex. In addition to the stomatin domain, UNC-24 has a lipid transfer domain
13 (Barnes et al., 1996; Sedensky et al., 2001), probably important for the mem-
14 brane localization of the membrane channel complex. The *unc-24* gene is expressed
15 in the touch neurons, while the UNC-24 protein appears in puncta that colocal-
16 ize with MEC-4 and MEC-2. Mutation of *unc-24* enhances the Mec phenotype
17 caused by *mec-4* and *mec-6* temperature sensitive alleles (Zhang et al., 2004).
18 UNC-24 appears to interact with MEC-2 and MEC-4 through its stomatin-like
19 domain.

20 *mec-6* encodes a 377-amino acid protein and is expressed in muscle cells, neu-
21 rons and other tissues (Chelur et al., 2002). Recessive *mec-6* mutations disrupt
22 touch sensitivity but do not cause detectable changes in touch cell ultrastructure
23 (Chalfie and Sulston, 1981; Tavernarakis and Driscoll, 1997). *mec-6* alleles have
24 the interesting property that they completely block *mec-4(d)* and *mec-4(A673V)*-
25 induced touch cell degeneration (Harbinder et al., 1997; Huang and Chalfie, 1994;
26 Tavernarakis and Driscoll, 1997). MEC-6 encodes a protein with limited similarity
27 to Paraoxonases/Arylesterases that physically interacts with MEC-4 and MEC-10
28 (Chelur et al., 2002). MEC-6 has a short cytoplasmic N-terminus, a single trans-
29 membrane domain, and a large extracellular C-terminus. How exactly MEC-6 acts
30 to influence MEC-4/MEC-10 channel activity is unknown. Nevertheless, it appears
31 that *mec-6* mutations do not affect *mec-4* transcription, although they do cause
32 full-length MEC-4::LacZ or MEC-4::GFP reporter fusion chimeras to be rapidly
33 degraded (N. T. unpublished observations; (Chelur et al., 2002)). Thus, working
34 hypotheses concerning the function of MEC-6 focus on two possibilities. First,
35 MEC-6 is another subunit needed for channel function or assembly, or second, it
36 mediates localization or post-translational modification essential for MEC-4 and
37 MEC-10 activity/stability. It should be noted that MEC-6 function is not exclu-
38 sively related to the MEC-4/MEC-10 touch receptor channel. *mec-6* mutations also
39 suppress the deleterious consequences of neurodegeneration-inducing mutations in
40 other *C. elegans* degenerins including *deg-1*, *unc-8* and partly *unc-105* ((Chalfie and
41 Wolinsky, 1990; Liu et al., 1996; Shreffler et al., 1995; Tavernarakis et al., 1997);
42 N.T unpublished observations). *mec-6* loss-of-function mutations affect localization
43 of the MEC-4 channel and disrupt touch evoked membrane currents (Chelur et al.,
44 2002).

45 Although the exact stoichiometry of the components of the mechanotransducer
46 channel complex is not known, genetic data suggest that several proteins are

01 present in multiple copies and in various combinations. Two subgroups of *mec*
02 genes encoding peripheral components required for mechanotransduction in the
03 touch receptor neurons can be defined, those encoding intracellular (*mec-7*, *mec-*
04 *12*) and those encoding extracellular (*mec-1*, *mec-5*, *mec-9*) proteins (Driscoll and
05 Kaplan, 1996; Driscoll and Tavernarakis, 1997; Tavernarakis and Driscoll, 1997).
06 As described previously, the touch receptor processes are filled with bundled 15-
07 protofilament microtubules. Mutations in two genes, *mec-7* and *mec-12*, disrupt the
08 formation of these microtubules (Chalfie and Au, 1989; Chalfie and Sulston, 1981;
09 Fukushige et al., 1999; Hamelin et al., 1992; Savage et al., 1989; Savage et al.,
10 1994). *mec-7* encodes a β -tubulin expressed at high levels in the touch receptor
11 neurons (Hamelin et al., 1992; Savage et al., 1989; Savage et al., 1994). MEC-7
12 is highly conserved—apart from the carboxy-terminal domain that is characteris-
13 tically highly variable; only 7 amino acids differ from other β -tubulins. *mec-12*
14 encodes an α -tubulin expressed at high levels in the touch receptor neurons but
15 also expressed in several other neurons that do not assemble 15-protofilament mi-
16 crotubules (Fukushige et al., 1999). Thus, the presence of the MEC-12 tubulin is
17 not sufficient to nucleate assembly of the touch-cell specific microtubules. As is the
18 case for *mec-7*, many *mec-12* mutations are semi-dominant or dominant and are
19 likely to disrupt subunit interactions or protofilament assembly (Gu et al., 1996).
20 Recent data suggest that tethering of the channel complex to the microtubules is not
21 essential for transduction, as mechanoreceptor currents are reduced but not elim-
22 inated in *mec-7* β -tubulin mutants (O'Hagan et al., 2005) and *mec-7* and *mec-12*
23 null mutations do not prevent the formation of channel puncta (Emtage et al., 2004;
24 Zhang et al., 2004).

25 In *mec-1* mutants, touch cells generally lack the mantle and associated periodic
26 specializations of the overlying cuticle (Chalfie and Sulston, 1981; Gu et al., 1996;
27 Savage et al., 1994). *mec-1* is expressed in touch receptor neurons, other lateral
28 neurons and intestinal muscles. It encodes a likely secreted protein with multiple
29 Kunitz-type serine protease inhibitor and EGF domains. The Kunitz and EGF do-
30 mains are likely to be protein interaction domains. The C terminus of MEC-1 is
31 needed for touch sensitivity, while the N terminus mediates the attachment of the
32 touch neuron processes to the hypodermis (Emtage et al., 2004). MEC-1 is local-
33 ized along the touch receptor processes in a punctuate manner and colocalizes with
34 MEC-5 and the MEC-4/MEC-10 mechanosensory channel complex (Emtage et al.,
35 2004).

36 *mec-5* mutations disrupt the extracellular matrix in a subtle manner; the mantle
37 in a wild-type animal can be stained with peanut lectin, whereas the mantle in *mec-5*
38 mutants cannot (Chalfie and Sulston, 1981; Du et al., 1996; Gillespie and Walker,
39 2001). *mec-5* encodes a novel collagen type that is secreted by hypodermal cells
40 (Du et al., 1996). The central portion of the *mec-5* protein is made up of Pro-rich
41 Gly-X-Y repeats. *mec-5* mutations, many of which are temperature-sensitive, cluster
42 toward the carboxy terminus of the protein and affect these repeats. Genetic
43 interactions suggest that *mec-5* influences MEC-4/MEC-10 channel function (for
44 example, *mec-4* and *mec-10* mutations can enhance the *mec-5(ts)* mutant phenotype;
45 (Gu et al., 1996)). Thus, a specialized collagen could interact with the touch receptor
46 channel, perhaps acting to provide gating tension.

01 *mec-9* mutations do not alter mantle ultrastructure in a detectable manner, despite
02 the fact that *mec-9* encodes a protein that appears to be secreted from the touch
03 receptor neurons (Chalfie and Sulston, 1981; Du et al., 1996). The *mec-9* gene
04 generates two transcripts, the larger of which encodes an 834 amino acid protein
05 (MEC-9L) that is expressed only by the touch receptors (Chalfie and Sulston, 1981;
06 Du et al., 1996). Akin to MEC-1, the predicted MEC-9L protein contains several do-
07 mains related to the Kunitz-type serine protease inhibitor domain, the Ca²⁺-binding
08 EGF repeat, the non-Ca²⁺-binding EGF repeat and a glutamic acid-rich domain
09 (Chalfie and Sulston, 1981; Du et al., 1996). Single amino acid substitutions that
10 disrupt MEC-9 function affect the two Ca²⁺-binding EGF repeats, the sixth EGF re-
11 peat and the third Kunitz-type domain, thus implicating these regions as important in
12 MEC-9 function (Chalfie and Sulston, 1981; Du et al., 1996). How MEC-9 is needed
13 for touch cell activity is not clear, but it is interesting that MEC-9 appears specialized
14 for protein interactions and that agrin, a protein that acts to localize acetylcholine
15 receptors, has a domain structure that appears similarly specialized (agrin features
16 multiple EGF and Kazal-type serine protease inhibitor repeats; (Rupp et al., 1992a;
17 Rupp et al., 1992b; Rupp et al., 1991)). *mec-9* mutations are dominant enhancers of
18 a *mec-5(ts)* allele, suggesting that these proteins might interact in the unique mantle
19 extracellular matrix outside the touch receptor neuron (Du et al., 1996; Gu et al.,
20 1996).

21 22 **5.5 Proprioception** 23 24

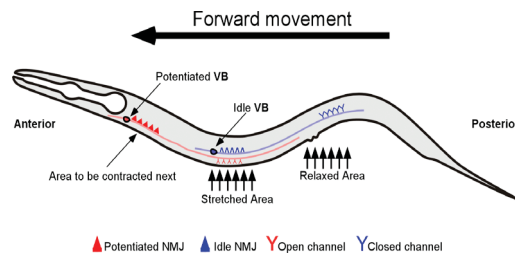
25 *C. elegans* senses forces arising within the body itself during movement, a phe-
26 nomenon called proprioception. Animal locomotion results from alternate contrac-
27 tion and relaxation of dorsal and ventral body wall muscles, which generates a
28 canonical sinusoidal pattern of movement (White et al., 1986; Wolinsky and Way,
29 1990). The arrangement of the body wall muscles and their synaptic inputs re-
30 stricts locomotion to dorsal and ventral turns of the body. The body wall muscles
31 are organized into two dorsal and two ventral rows. Each row consists of 23 or
32 24 diploid mononucleate muscle cells arranged in an interleaved pattern (Francis
33 and Waterston, 1991; Moerman et al., 1996; Waterston et al., 1980; Williams and
34 Waterston, 1994). Distinct classes of motoneurons control dorsal and ventral body
35 muscles. To generate the sinusoidal pattern of movement, the contraction of the
36 dorsal and ventral body muscles must be out of phase. For example, to turn the
37 body dorsally, the dorsal muscles contract, while the opposing ventral muscles re-
38 lax. Interactions between excitatory and inhibitory motoneurons produce a pat-
39 tern of alternating dorsal and ventral contractions (Francis and Waterston, 1991;
40 Hresko et al., 1994; Tavernarakis and Driscoll, 1997). Relatively little is known
41 about how the sinusoidal wave is propagated along the body axis. Adjacent muscle
42 cells are electrically coupled via gap junctions, which could couple excitation of ad-
43 jacent body muscles. Alternatively, ventral cord motoneurons could promote wave
44 propagation since gap junctions connect adjacent motoneurons of a given class
45 (Chalfie et al., 1985; White et al., 1976; White et al., 1986). A third possibility is
46 that motoneurons could themselves act as stretch receptors so that contraction of

01 body muscles could regulate adjacent motorneuron activities, thereby propagating
02 the wave (Tavernarakis and Driscoll, 1997; Tavernarakis and Driscoll, 2001b; Tav-
03 ernarakis et al., 1997). Numerous mutations disrupt normal sinusoidal locomotion
04 in *C. elegans*, resulting in animals with movement defects ranging from total paral-
05 ysis, to severe uncoordination, to subtle and almost imperceptible irregularities in
06 movement (Tavernarakis and Driscoll, 1997; Tavernarakis et al., 1997). Unusual,
07 semi-dominant (sd), gain-of-function mutations in the gene *unc-8* induce transient
08 neuronal swelling of embryonically derived motorneurons as well as some neurons
09 in the head and tail ganglia, and severe uncoordination (Park and Horvitz, 1986b;
10 Shreffler et al., 1995; Tavernarakis et al., 1997). Swelling is absent at hatching and
11 peaks in severity late in L1 and L2. *unc-8* encodes a degenerin expressed in sev-
12 eral motor neuron classes, in some interneurons and in nose touch sensory neurons.
13 Interestingly, semi-dominant *unc-8* alleles alter an amino acid in the region hypoth-
14 esized to be an extracellular channel-closing domain defined in studies of *deg-1*
15 and *mec-4* degenerins. The genetics of *unc-8* are further similar to those of *mec-4*
16 and *mec-10*; specific *unc-8* alleles can suppress or enhance *unc-8(sd)* mutations
17 in trans, suggesting that UNC-8::UNC-8 interactions occur (Shreffler et al., 1995;
18 Tavernarakis et al., 1997). Another degenerin family member, *del-1* (degenerin-like)
19 is co-expressed in a subset of neurons that express *unc-8* (the VA and VB motor
20 neurons) and is likely to assemble into a channel complex with UNC-8 in these
21 cells (Tavernarakis et al., 1997). The UNC-8 and DEL-1 proteins include all do-
22 mains characteristic of degenerin family members and are likely to adopt similar
23 transmembrane topologies (amino and carboxy termini situated inside the cell and
24 a large extracellular domain that includes three cysteine-rich regions). Neither de-
25 generin has any primary sequence features that are markedly different from other *C.*
26 *elegans* family members although one somewhat atypical feature of UNC-8 is that it
27 has a relatively long C-terminal domain that shares some primary sequence homol-
28 ogy with the extended C-terminus of another degenerin implicated in locomotion,
29 UNC-105 (Liu et al., 1996; Park and Horvitz, 1986a).

30 The exact function of the UNC-8 degenerin channel in motorneurons was elu-
31 cidated through genetic approaches. *unc-8* null mutants have a subtle locomotion
32 defect; they inscribe a path in an *E. coli* lawn that is markedly reduced in both
33 wavelength and amplitude as compared to wild type (Tavernarakis et al., 1997).
34 This phenotype indicates that the UNC-8 degenerin channel functions to modulate
35 the locomotory trajectory of the animal. How does the UNC-8 motor neuron channel
36 influence locomotion? As mentioned earlier, one highly interesting morphological
37 feature of some motorneurons (in particular, the VA and VB motorneurons that
38 co-express *unc-8* and *del-1*) is that their processes include extended regions that
39 do not participate in neuromuscular junctions or neuronal synapses. These “undif-
40 ferentiated” process regions have been hypothesized to be stretch-sensitive (White
41 et al., 1986). Given the morphological features of certain motor neurons and the
42 sequence similarity of UNC-8 and DEL-1 to the candidate mechanically-gated chan-
43 nels MEC-4 and MEC-10, we have proposed that these subunits co-assemble into
44 a stretch-sensitive channel that might be localized to the undifferentiated regions
45 of the motor neuron process (Tavernarakis et al., 1997). When activated by the
46 localized body stretch that occurs during locomotion, this motor neuron channel

01 potentiates signaling at the neuromuscular junction, which is situated at a distance
 02 from the site of stretch stimulus (Fig. 5.6). The stretch signal enhances motorneuron
 03 excitation of muscle, increasing the strength and duration of the pending muscle
 04 contraction and directing a full size body turn. In the absence of the stretch acti-
 05 vation, the body wave and locomotion still occur, but with significantly reduced
 06 amplitude because the potentiating stretch signal is not transmitted. This model
 07 bears similarity to the chain reflex mechanism of movement pattern generation.
 08 However it does not exclude a central oscillator that would be responsible for the
 09 rhythmic locomotion. Instead, we suggest that the output of such an oscillator is
 10 further enhanced and modulated by stretch sensitive motorneurons (Tavernarakis
 11 and Driscoll, 1997; Tavernarakis et al., 1997). One important corollary of the *unc-8*
 12 mutant studies is that the UNC-8 channel does not appear to be essential for motor
 13 neuron function. If this were the case, animals lacking the *unc-8* gene would be
 14 severely paralyzed. This observation strengthens the argument that degenerin chan-
 15 nels function directly in mechanotransduction rather than merely serving to maintain
 16 the osmotic environment so that other channels can function.

17 Muscle cells may also play part in the coordination of locomotion by sensing
 18 their own extent of stretch. Mutations in the muscle degenerin *unc-105* cause muscle
 19 hypercontraction (Garcia-Anoveros et al., 1998). The muscle hyper-contraction phe-
 20 notype caused by dominant *unc-105* mutations can be suppressed by mutations near
 21 the carboxy-terminus of *let-2*, a gene that encodes the $\alpha 2$ chain of type-IV collagen
 22 found in the basement membrane between muscle cells and the hypodermis (Liu
 23 et al., 1996). It is tempting to speculate that LET-2 normally carries gating tension
 24 to the UNC-105 channel, when the muscle is stretched, thus providing regulatory
 25 feedback for muscle contraction. However, results from *in vivo* electrophysiology
 26 suggest that UNC-105 is not involved in the formation of a muscle stretch receptor
 27 complex (Jospin et al., 2004).



38 **Fig. 5.6** A model for UNC-8 involvement in stretch-regulated control of locomotion. Schematic
 39 diagram of potentiated and inactive VB class motor neurons. Neuro-muscular junctions (signi-
 40 fied by triangles) are made near the cell body (Tavernarakis et al., 1997; White et al., 1986).
 41 Mechanically-activated channels postulated to include UNC-8 (and, possibly in VB motor neurons,
 42 DEL-1) subunits (signified by Y figures) are hypothesized to be concentrated at the synapse-free,
 43 undifferentiated ends of the VB neuron. Mechanically-gated channels could potentiate local exci-
 44 tation of muscle. Body stretch is postulated to activate mechanically-gated channels which potentiate
 45 the motor neuron signal that excites a specific muscle field. Sequential activation of motor neurons
 46 that are distributed along the ventral nerve cord and signal non-overlapping groups of muscles, am-
 plifies and propagates the sinusoidal body wave (NMJ: neuromuscular junction). Reproduced from
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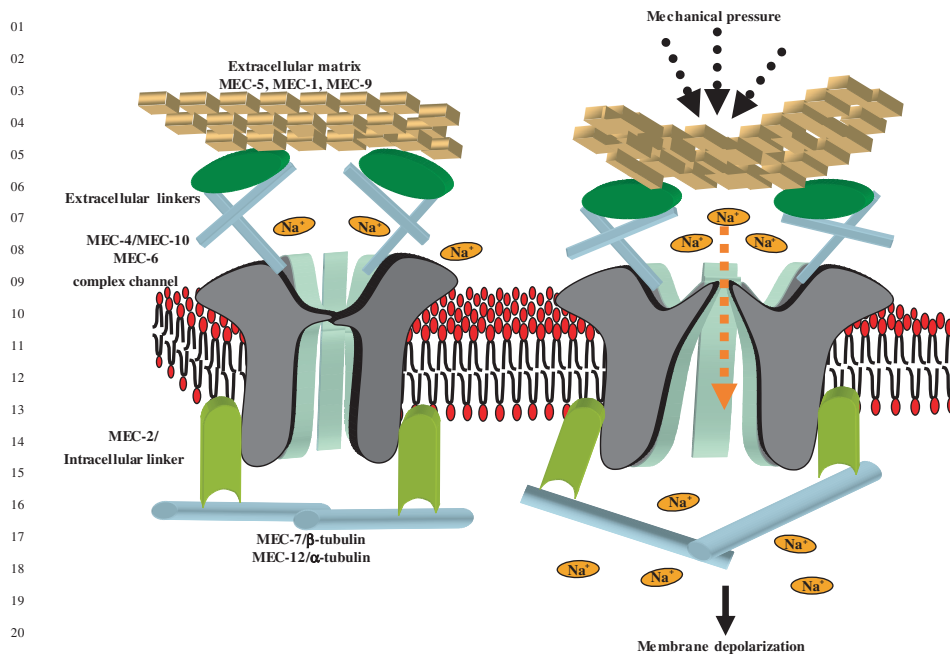
01 In addition to DEG/ENaC ion channels, a transient receptor potential (TRP)
02 channel has recently been found to be critical for proprioception in *C. elegans*.
03 TRP cation channels are present in all eukaryotes, from yeast to mammals and can
04 be divided into seven subfamilies based on sequence similarities (Montell, 2005).
05 TRP channels are linked to many physiological processes ranging from temperature
06 sensation to mechanosensation and osmosensation (Caterina et al., 1999; Caterina
07 et al., 1997; Colbert et al., 1997; Peier et al., 2002; Tobin et al., 2002). The *trp-4*
08 gene is orthologous to the *Drosophila* *nompC* channel which is critical for hair cell
09 mechanotransduction (Walker et al., 2000) and to the zebrafish *nompC* which is
10 required for the function of auditory hair cells (Sidi et al., 2003). *C. elegans* *trp-4*
11 mutants generate abnormal locomotion waves, with exaggerated body bands and
12 larger than normal wave amplitudes (Li et al., 2006). TRP-4 acts in a single in-
13 terneuron, called DVA and TRP-4 channels may act as stretch receptors in these
14 neurons to provide sensory feedback to the locomotor control circuit. Thus, DVA
15 appears to be a primary proprioceptor neuron which modulates the shape of the
16 locomotion sinusoidal wave.

17
18

19 **5.6 Mechanotransduction in Other Organisms**

20
21

22 Genetic studies of sensory mechanotransduction, which were initiated in *C. elegans*
23 and are now also being carried out in *Drosophila* and in mammals, have converged
24 to reveal a limited set of underlying mechanisms (Eberl, 1999; Gillespie and Walker,
25 2001; Harteneck et al., 2000; Kellenberger and Schild, 2002). For example, the
26 model proposed for mechanotransduction in the touch receptor neurons (Fig. 5.7)
27 and motoneurons of *C. elegans* shares the same underlying principle and features of
28 the proposed gating mechanism of mechanosensory ion channels in *Drosophila* sensory
29 bristles and the channels that respond to auditory stimuli in the hair cells of the
30 vertebrate inner ear (Fettiplace and Fuchs, 1999; Gillespie and Walker, 2001; Hamill
31 and McBride, 1996; Hudspeth, 1989; Hudspeth et al., 2000; Jaramillo and Hudspeth,
32 1991; Pickles and Corey, 1992; Pickles et al., 1991; Tavernarakis and Driscoll, 1997;
33 Weinbaum et al., 2001). Hair cells have bundles of a few hundred stereocilia on their
34 apical surface, which mediate sensory transduction. Stereocilia are connected at their
35 distal ends to neighboring stereocilia by filaments called tip links. The integrity of
36 the tip links is essential for channel opening and the mechanosensitive channels
37 appear to be situated at the ends of the stereocilia, near the connecting tip links. Di-
38 rectional deflection of the stereocilia relative to each other introduces tension on the
39 tip links, which is proposed to open the mechanosensitive hair cell channels directly.
40 This remarkable convergence of independent studies in distant species, strongly sug-
41 gests that different mechanotransducers in different systems have evolved to strictly
42 adhere to the same set of principles. Besides the DEG/ENaC family of ion channels,
43 members of the TRP family are involved in mechanosensation in different organ-
44 isms. (Alvarez de la Rosa et al., 2000; Duggan et al., 1998; Minke and Cook, 2002;
45 Montell, 2001; Tavernarakis and Driscoll, 2001a; Welsh et al., 2002). Experiments
46 in *Drosophila* revealed the involvement of TRP-like channel genes in the function of



22 **Fig. 5.7** A model for a mechanotransducing complex in *C. elegans* touch receptor neurons. The
 23 extracellular matrix contains MEC-5, MEC-9 and MEC-1; the sensory transduction channel is
 24 formed by MEC-4, MEC-10 and possibly MEC-6; MEC-2 enhances channel activity and tethers
 25 the channel to specialized microtubules containing MEC-7/ β -tubulin and MEC-12/ α -tubulin.
 26 Recent findings suggest that mechanotransduction can occur even when the 15-protofilament
 27 microtubules are missing in *mec-7* mutants, questioning in that way the necessity of tethering of
 28 the complex to microtubules (O'Hagan et al., 2005). In the absence of mechanical stimulation,
 29 the channel is closed and the sensory neuron is idle. Application of mechanical force to the body of
 30 the animal results in distortion of a network of interacting molecules that opens the degenerins
 31 channel. Na^+ influx depolarizes the neuron, initiating a cascade that leads to the integration of the
 32 stimulus

33 mechanosensory bristles (*nompC*; (Walker et al., 2000)), auditory receptors (*iav* and
 34 *nan*; (Gong et al., 2004)), and nociceptors (*painless*; (Tracey et al., 2003)). Another
 35 member of the TRP protein family, the TRPA1 channel has been identified as a
 36 candidate mechanotransducing channel in the mouse (Corey et al., 2004). *In situ*
 37 hybridization revealed that the TRPA1 channel is expressed in the cochlea organ of
 38 Corti, which contains the auditory hair cells. Additional colocalization experiments
 39 link TRPA1 to mechanosensation: TRPA1 is expressed together with two accessory
 40 proteins of the mechanosensory apparatus, myosin 1c and cadherin 23, at the tips of
 41 stereocilia throughout the kinocilium and in the pericuticular zone. Whole-cell
 42 patch clamp recording of inner hair cells in mice show that the transduction current
 43 produced is significantly reduced in the absence of TRPA1, indicating that this
 44 channel is a component of the mechanosensitive transduction channel of vertebrate
 45 hair cells (Corey et al., 2004). Additionally, two TRP proteins, a NompC-like protein
 46 and TRPA1 are required for hair cell function in zebrafish (Corey et al., 2004;
 Sidi et al., 2003).

01 Despite enormous progress on the illumination of vertebrate mechanosensory
02 cell biology achieved in recent years, there is still a striking gap between the bio-
03 physical information that has accumulated and our understanding of the molecular
04 aspects of mechanosensation. Sophisticated experiments in mice and humans re-
05 vealed many genes involved in the development and function of the mammalian
06 cochlea and have cumulated in the formulation of the gating-spring model for hair
07 cell mechanotransduction (Gillespie, 1995; Gillespie and Walker, 2001). However,
08 many pieces of the mechanotransducing apparatus puzzle are still missing. Work
09 in lower vertebrates such as birds, amphibians and fish has also contributed sig-
10 nificantly in complementing and extending the studies with mammals. In these
11 animals mechanosensory structures are often much easier to access, follow and
12 monitor providing large potential for investigating the molecular basis of audi-
13 tory transduction (Ashmore, 1998; Smotherman and Narins, 2000). An increas-
14 ing amount of evidence suggests that some mammalian DEG/ENaC proteins may
15 play a role in mechanosensation similarly to their nematode counterparts. In mam-
16 mals, there are strong indications that ENaC subunits may be components of
17 the baroreceptor mechanotransducer, one of the most potent regulators of arte-
18 rial pressure and neurohumoral control of the circulation (Drummond et al., 1998;
19 Drummond et al., 2001). Members of the ASIC (acid sensing ion channel) sub-
20 group of the DEG/ENaC family have been implicated in mechanotransduction in
21 mammals. BNC1 (brain Na⁺ channel; also known as MDEG, BNaC1, ASIC2;
22 (Garcia-Anoveros and Corey, 1997; Price et al., 1996; Waldmann et al., 1996; Wald-
23 mann and Lazdunski, 1998)) has emerged as promising candidate for a mechanosen-
24 sitive channel. In BNC1 null mice touch receptor neurons of the skin produce fewer
25 action potentials than in wild type animals over a comparable range of stimuli (Price
26 et al., 2000).

29 5.7 Conclusions

32 Studies in *C. elegans* have contributed critical insights into the cellular and molecu-
33 lar mechanisms of mechanotransduction (Fig. 5.7; Chalfie, 1997; Syntichaki and
34 Tavernarakis, 2004; Tavernarakis and Driscoll, 1997; Tavernarakis and Driscoll,
35 2001b). Recently developed powerful methodologies, such as direct electrophysio-
36 logical recordings from neurons and imaging of genetically encoded calcium
37 sensors provide the unique opportunity to investigate the properties of the mechan-
38 otransduction apparatus in the context of live, behaving animals.

39 Although our understanding of metazoan mechanosensation has been advanced
40 significantly, open questions still remain. While specific DEG/ENaC and TRP ion
41 channels have been directly implicated in the process of mechanotransduction the
42 *C. elegans* genome encodes many more members of these ion channel proteins. The
43 role of these proteins in mechanotransduction remains to be elucidated.

44 An additional major question that remains to be addressed is whether the mam-
45 malian counterparts of the *C. elegans* degenerins play specialized roles in
46 mechanical signalling in humans. A significant step toward addressing this ques-

tion has been accomplished with the demonstration that BNC1 is involved in mechanosensory signalling in the skin as we have described above. Even though the candidacy of BNC1 for being in the core of a mechanotransducing complex was greatly boosted by these results, a demanding critic would argue that albeit very strong, it still remains just a candidacy. The potential role of BNC1 as part of the core mechanotransducing channel can still only be inferred from these experiments and is not directly proven. It is still possible that BNC1 forms or participates in an auxiliary channel that facilitates the function of the actual mechanotransducing channel. A BNC1 knockout does not completely eliminate the responses to mechanical stimuli (Price et al., 2000). The incomplete nature of the BNC1 deficiency effects indicates that even if BNC1 is indeed part of the core mechanosensory channel, it most likely is not the only critical one. Alternatively, there might be more than one, different mechanotransducing complexes within one neuron, with different properties and composition. The above arguments however, are by no means confined to BNC1. On the same basis, MEC-4/MEC-10 and UNC-8/DEL-1 in *C. elegans* as well as PPK in *Drosophila* might not be parts of the real mechanotransducer but only auxiliary ion channels. The recent identification of another strong candidate mechanosensory channel, the *Drosophila* NompC, adds to the list of candidate mechanosensitive ion channels (Walker et al., 2000). Evidence implicating NompC in mechanotransduction is especially convincing given the supporting electrophysiological analysis that is feasible in this system, and the availability of mutants with altered properties and intermediate effects (Walker et al., 2000). Therefore, NompC homologues in other organisms, including humans, emerge putative mechanosensitive ion channels. Even in this case however, there are caveats; the absence of NompC does not completely eliminate mechanosensitive currents in *Drosophila* hair bristles. Furthermore, the identities and properties of force-generating tethers of NompC in mechanotransducing complexes will need to be determined. Another issue that needs to be addressed is the potential interplay between DEG/ENaC and NompC channels in mechanosensory cells before a clear understanding of mechanotransduction can be achieved.

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