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Transcription Factor *Shox2* Is Required for Proper Development of TrkB-Expressing Mechanosensory Neurons

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Abstract

Dorsal root ganglia (DRG) contain somatosensory neurons of diverse sensory modalities. Among these different types of sensory neurons, the molecular mechanisms that regulate the development and specification of touch neurons are least well understood. We took a candidate-approach and searched for transcription factors that are expressed in subsets of DRG neurons, and found that the transcription factor *Shox2* is expressed in subpopulations of TrkB- and Ret-expressing neurons at neonatal stages. Since TrkB is a known marker selectively expressed in touch sensory neurons, we decided to examine the function of *Shox2* in specifying TrkB-positive DRG neurons. Conditional deletion of *Shox2* in neural crest cells (which give rise to all DRG neurons) caused a 60~65% reduction in the number of TrkB-expressing neurons. It also resulted in an increase in co-expression of TrkC in Ret-positive sensory neuron. Deletion of *Shox2* in differentiating DRG neurons at later time points caused only a moderate reduction in TrkB expression. Overexpression of *Shox2* in all neural crest cells resulted in a small increase in the number of TrkB-expressing neurons. Finally, *Shox2*-deletion also caused reduced touch sensory axonal innervation to layers III/IV of the spinal cord. Taken together, our findings identify *Shox2* as an essential but not sufficient component of the transcription programs required in neural progenitor cells for the proper specification of subsets of TrkB-expressing touch/mechanosensory neurons.

Introduction

Somatosensory neurons located in dorsal root ganglia (DRG) consist of many different types that detect diverse modalities of sensory stimuli. All DRG neurons are originated from neural crest cells (NCCs) (Chai et al., 2000; Ma et al., 1999; Szeder et al., 2003). We are interested in identifying molecular mechanisms that enable NCCs to differentiation into touch/mechanosensory neurons. Different DRG neurons have unique molecular compositions. Receptors for neurotrophic factors are among the best characterized markers for sensory neurons. It has been shown that TrkA and Ret receptors are mainly expressed in nociceptive and thermal sensory neurons (Chen et al., 2006; Kramer et al., 2006; Luo et al., 2007). TrkC is expressed primarily in proprioceptive neurons innervating the skeletal muscles (Hasegawa and Wang, 2008; Hippenmeyer et al., 2005; Inoue et al., 2007; Sedy et al., 2006). TrkB is expressed by a subpopulation of cutaneous low-threshold touch neurons

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(Gonzalez-Martinez et al., 2004; Perez-Pinera et al., 2008; Shimizu et al., 2007). An early born population of Ret-expressing neurons develops into rapid-adapting mechanosensory neurons (Bourane et al., 2009; Luo et al., 2009). Signaling through neurotrophic receptors is important for neuron survival, axon growth, innervation of central and peripheral targets, and proper differentiation into specialized and modality-specific sensors (da Silva and Wang, 2010; Marmigere and Ernfors, 2007).

Significant progresses have been made in elucidating the transcriptional programs specifying nociceptive and proprioceptive neurons. For example, the transcription factor Runx1 is essential for differentiation and diversification of nociceptive neurons into peptidergic and non-peptidergic lineages (Chen et al., 2006; Inoue et al., 2008; Kramer et al., 2006; Marmigere and Ernfors, 2007), whereas Runx3 and Er81 are important for the specification of TrkC-expressing proprioceptive sensory neurons (Hippenmeyer et al., 2005; Inoue et al., 2007; Kramer et al., 2006; Levanon et al., 2001). Recently, MafA was shown to be involved in the development of Ret-positive rapid-adapting mechanoreceptors (Bourane et al., 2009). However, the transcription factors that enable progenitor cells to differentiate into TrkB-expressing mechanosensory neurons remain unclear.

We searched for transcription factors expressed in a subset of DRG neurons and found the gene encoding Shox2 (Short Stature Homeobox 2), a homeobox transcription factor, is dynamically expressed during DRG development. The mouse *Shox2* gene displays 99% and 73% similarity to human *Shox2* and *Shox*, respectively. Mutations in human *Shox* cause short stature and Leri-Weill dyschondrosteosis (Binder, 2011; Marchini et al., 2007). Mice only have the *Shox2* gene. Thus, it appears that mouse Shox2 assumes the functions of both human Shox and Shox2. *Shox2* mutant mice show defects in bone, heart, and palate development (Blaschke et al., 2007; Cobb et al., 2006; Yu et al., 2005). However, the role of Shox2 in neuronal development has not been examined. Here we performed loss- and gain-of-function analyses to determine the role of Shox2 in DRG development. We discovered that Shox2 is important for the development of TrkB-expressing mechanosensory neurons.

Materials and Methods

Mice

Shox2^{fllox/fllox} (Cobb et al., 2006), *Wnt1-Cre* (Danielian et al., 1998), *Advillin^{Cre/+}* (Zhou et al., 2010), *Isl1^{Cre/+}* (Srinivas et al., 2001; Yang et al., 2006) and *Advillin^{PLAP/+}* (Hasegawa et al., 2007) mice are all previously described. *Rosa^{CAG-STOP-Shox2/+}* mouse was generated by inserting a cassette of “chicken β Actin promoter-LoxP-neo-polyA-LoxP-Shox2-polyA” into the *Rosa26* locus via homologous recombination. Genotyping of the *Rosa^{CAG-STOP-Shox2/+}* mice was performed by PCR. PCR primers were designed as follows: *Rosa/01*, 5-CACTTGCTCTCCCAAAGTCG-3; *Rosa/02*, 5-TAGTCTAACTCGCGACACTG-3; *CAG/02*, 5-GTTATGTAACGCGGAACTCC-3. The wild-type allele produces a 560 bp fragment with *Rosa/01* and *Rosa/02* primers, whereas the knock-in allele results in a 300 bp fragment with *Rosa/01* and *CAG/02* primers. Furthermore, primers were designed to specifically detect the Shox2 cDNA in the *Rosa^{CAG-STOP-Shox2}* allele: *RShox2/01*, 5-GTGTCCCCTGAACTGAAGGA-3; and *RShox2/02*, 5-GCCTGAACCTGAAAGGACAA-3. The knock-in allele produces a 400 bp fragment using the *RShox2/01* and *RShox2/02* primers. All experiments were conducted according to protocols approved by The Duke University Institutional Animal Care and Use Committee.

In situ hybridization—The mouse cDNA fragments of the neurotrophic receptors and Shox2 were amplified by PCR with the antisense primers containing the T7 promoter sequence. *In vitro* transcription was then performed from the PCR-amplified template using

T7 RNA polymerase (Roche, Indianapolis, IN) with Digoxigenin-UTP (Roche) for the synthesis of the antisense probes. *In situ* hybridization was performed according to standard methods (Hodge et al., 2007). Fluorescent two-color *in situ* hybridization was performed according to standard methods (Hasegawa and Wang, 2008).

Immunostaining—Immunostaining was performed according to standard methods (Hodge et al., 2007). Antibody used are: Alexa Fluor 488-conjugated IB4 (Molecular Probes/Invitrogen, Eugene, OR), Anti-Caspase 3 (active) (1:1000; R&D Systems, Minneapolis, MN), Anti-CGRP antibody (1:2000; Chemicon/Invitrogen), Anti-PGP9.5 antibody (UltraClone, Isle of Wight, UK), Anti-vGluT1 antibody (1:1000; Millipore, Billerica, MA), Alexa 488-labeled anti-rabbit IgG (1:400; Invitrogen), Alexa 488-labeled anti-guinea pig IgG (1:400; Invitrogen), and Cy3-labeled anti-rabbit IgG (1:400, Jackson Labs, Bar Harbor, MA).

Alkaline phosphatase staining—Alkaline phosphatase staining was performed according to standard methods (Hasegawa et al., 2007).

Quantification methods—For every developmental stage, at least 3 embryos/pups from 2 to 3 different litters were analyzed. *In situ* (or immuno) signal positive DRG neurons were counted from randomly selected sections. For counting TrkA or Ret positive neurons, N=20 randomly selected sections from each animal are counted; for counting Shox2, TrkC, TrkB or activated caspase-3 positive neurons, N=40~50 randomly selected sections from each animal are counted; for counting MafA, Runx3, or parvalbumin positive neurons, N=30~35 randomly selected sections from each animal are counted. On each section, the area of the DRG was measured using MetaMorph software. Number of cells per unit area is then calculated, and averaged over all embryos/animals. *p* values were calculated using Student's *t* test. PLAP staining intensity (from 50 randomly selected sections of animals of 2 different litters) and vGluT1 staining intensities and areas (from 60 randomly selected sections of animals of 2 different litters) were measured using MetaMorph software and were set to artificial units. *p* values were calculated with Student's *t* test.

Results

***Shox2* expression pattern in the developing mouse DRG**

In situ hybridization was carried out to examine the expression of *Shox2* in the developing mouse DRG. At early stages beginning at E10.5 (embryonic day 10.5), *Shox2* is expressed throughout the ganglion (Figure 1A). By E14.5, its expression begins to down-regulate in the majority of sensory neurons (Figure 1A). This down regulation continues after E14.5 and by E18.5 only a small population of DRG neurons retain stable expression of *Shox2*, which persist into adulthood (Figure 1A and data not shown). The dynamic expression pattern of *Shox2* led us to hypothesize that *Shox2* is involved in the specification and development of subtypes of DRG neurons. To test this, we carried out fluorescent two-color *in situ* hybridization experiments to examine the potential co-expression of *Shox2* with *Trks* or *Ret* receptors (which are markers for different somatosensory lineages). We observed that at P0 (postnatal day 0), *Shox2* co-localizes with subsets of *TrkB* or *Ret*-expressing neurons and is absent from the *TrkA*- and *TrkC*-expressing cells (Figure 1B). Upon quantification, we found that 65% of *Shox2*-positive cells co-localize with 65% of *TrkB*-neurons, and the remaining 35% of *Shox2*-expressing cells co-localizes with a small number of *Ret*-positive DRG neurons. Since *TrkB* has been shown to be a marker for subsets of mechanosensory neurons, and *Ret* is expressed in some of the rapid-adapting mechanosensory neurons in addition to nociceptive neurons, the co-expression results suggest that *Shox2* may be involved in the differentiation of NCCs into mechanosensory neurons.

Significant reduction in the number of *TrkB*-expressing DRG neurons in *Wnt1-Cre; Shox2^{flox/flox}* mouse

To examine the function of *Shox2* in DRG neuron development, we used *Wnt1-Cre* (Danielian et al., 1998) to conditionally delete the *Shox2* gene in all neural crest derived cells including DRG neurons. We crossed *Wnt1-Cre* mice with *Shox2^{flox/flox}* mice (Cobb et al., 2006; Danielian et al., 1998) to obtain conditional mutant *Wnt1-Cre; Shox2^{flox/flox}* (as well as control *Wnt1-Cre; Shox2^{flox/+}*) embryos and neonates. Figure 2A shows the absence of *Shox2* expression in DRG from *Shox2*-mutant embryo at E10.5. In our hands, all *Shox2* mutant mice die within two days after birth due to an anterior cleft palate defect (data not shown), which limited our characterization of the role of *Shox2* to embryonic and neonatal stages.

Using in situ hybridization, we examined the expression of *Trks* and *Ret* receptors in developing DRG at different stages in the mutant and control embryos. There is no observable difference in the number of *TrkA* or *Ret* positive cells at all time points examined, suggesting that *Shox2* does not play a major role in the development of nociceptive lineage (Figure 3C–D). In contrast, we observed a significant decrease in the number of *TrkB*-expressing cells in the *Shox2*-deleted DRG compared with controls at all time points examined (Figure 2B–C). On average, there is a 60~65% loss of *TrkB*-expressing DRG neurons at stages after E16.5 (Figure 2C), which is consistent with the fact that 65% *TrkB*-neurons express *Shox2*. Finally, the proprioceptive lineage marker *TrkC* showed a mild increase in the *Shox2*-deleted DRG at perinatal stages starting after E18.5 (Figure 3A–B). The detailed method used for quantifying the numbers of different types of DRG neurons in these and subsequent results is described in Materials and Method.

Previous studies have shown that the transcription factors *MafA* and *Runx3* are involved in the development of the *Ret*-positive mechanosensory neurons and *TrkC*-positive proprioceptive neurons, respectively (Bourane et al., 2009; Inoue et al., 2007; Kramer et al., 2006). We used in situ hybridization to examine the expression of both these transcription factors in mutant and control mice. We found that there was no difference in the number of DRG neurons expressing either *MafA* or *Runx3* in the *Shox2*-deleted versus control DRG at P0 or P2 (Figure 3E–G). The result suggests that *Shox2* do not regulate *MafA* or *Runx3* expression, and by extension, probably do not play a major role in the development/specification of *Ret*-positive rapid adapting mechanosensory neurons or *TrkC*-expressing proprioceptive neurons.

Loss of *TrkB*-expressing DRG neurons in *Wnt1-Cre; Shox2^{flox/flox}* mouse is not caused by elevated apoptosis

We next investigated whether apoptosis could account for the observed loss of *TrkB*-expressing neurons in *Wnt1-Cre; Shox2^{flox/flox}* DRG. We used anti-activated caspase-3 antibody to detect cell death, and found no statistically significant difference in the number of apoptotic cells in *Shox2*-deleted versus control DRG at all time points examined (Figure 4A–B). Note that both *Shox2*-deficient and control DRG show increased numbers of caspase-3 positive cells at E14.5 (Figure 4B), a time point of natural occurring cell death of developing sensory neurons as shown previously (Raff et al., 1993; White et al., 1998). Thus, the loss of *TrkB*-expression in *Shox2* mutant mice is not likely due to apoptosis of sensory neurons.

TrkC co-expression in subsets of *TrkB*- or *Ret*-positive sensory neurons in *Wnt1-Cre; Shox2^{flox/flox}* mouse

It is known that during early DRG neurogenesis, the transient population of *TrkC/TrkB* double positive progenitor neurons later differentiate into *TrkB*-single positive or *Ret*-single

positive mechanosensory neurons, or *TrkC*-single positive proprioceptive neurons (Kramer et al., 2006; Marmigere and Ernfors, 2007). However, there appears to be a small percent of sensory neurons that maintain co-expression of *TrkB/TrkC* or *Ret/TrkC*. We thus examined the co-expression of *TrkC* in the remaining *TrkB*-, as well as in *Ret*-expressing DRG neurons in *Shox2* mutant mice.

We first performed fluorescent two-color in situ hybridization to detect *TrkB* and *TrkC* mRNA simultaneously. In control DRG at E14.5, about 33% of *TrkB* positive neurons also express *TrkC*. By E16.5, less than 10% of *TrkB*-cells still express *TrkC* in wild type DRG. This number is further reduced during postnatal development (Figure 5A). In *Wnt1-Cre; Shox2^{fllox/fllox}* DRG, there is an apparent increase in the relative percentage of *TrkB/TrkC* double positive cells. However, when we quantified the actual average numbers of *TrkB/TrkC* double positive DRG neurons per unit area, there is no statistically significant difference between control and *Shox2*-mutant DRG (Figure 5B). This result suggests that normally in wildtype mouse, *TrkB/TrkC* double positive mechanosensory neurons belong to the 35% *TrkB*-positive but *Shox2*-negative populations, and thus their number is unaffected by *Shox2*-deletion.

Since *Shox2* is also expressed in a subset of *Ret*-positive neurons (Figure 1B), we also examined co-expression of *TrkC* in *Ret*-expressing sensory neurons. Again, using two-color in situ hybridization, we found that there are a small percentage of *Ret*-positive cells also expressing *TrkC* at perinatal stages in both control and *Shox2*-mutant DRG. Interestingly, the average number of *Ret/TrkC* double positive neurons per unit area is increased in the mutant (Figure 5C–D). This result is consistent with the observed increase in the total number of *TrkC*-expressing DRG neurons, and suggests that normally, the function of *Shox2* in *Ret*-positive neurons may be to suppress *TrkC* expression.

To examine whether any of the *Ret/TrkC* double positive cells in *Shox2*-deleted DRG differentiate toward a proprioceptive neuron fate, we carried out two-color in situ hybridization to detect *Ret* and *Parvalbumin (PV)* simultaneously. Parvalbumin is a known marker for *TrkC* expressing proprioceptive sensory neurons (Arber et al., 2000). *Ret/PV* double positive cells are rarely seen in either control or *Shox2*-mutant DRG, and there is no statistically significant difference in the total number of *PV* positive neurons between control and mutant mice (Figure 5E and data not shown). This result suggests that the increased number of *Ret/TrkC* double positive neurons in *Shox2*-mutant DRG do not appear to be proprioceptive neurons. To determine whether any of the *Ret/TrkC* double-positive neurons belong to the late-born *Ret*-positive nociceptive neurons, we performed two-color in situ using *MrgD* and *TrkC* probes. *MrgD* is expressed exclusively in *Ret/Runx1*-expressing non-peptidergic nociceptive neurons (Liu et al., 2008). We did not find any *MrgD/TrkC* double positive neurons in either control or *Shox2*-mutant mice (Figure 5F). Thus, by exclusion, the increased *Ret/TrkC* double positive neurons in *Shox2*-mutant are mechanosensory neurons.

Taken together, *Shox2*-deletion resulted in a significant loss of *TrkB* expression and a mild increase of *TrkC* co-expression in small number of *Ret*-positive mechanosensory neurons. These data suggest that *Shox2* is essential for the proper expression of neurotrophic receptors during the differentiation of mechanosensory neurons.

Developmental time-dependent requirement of *Shox2* for proper *TrkB* expression in DRG neurons

Previous studies have found time-dependent roles of certain transcription factors (ETS genes, *Islet1*) in DRG neuron development (Hippenmeyer et al., 2005; Sun et al., 2008). We therefore wanted to determine the time window when *Shox2* is required for the development

of a subset of *TrkB*-expressing sensory neurons. We used *Avil*^{Cre/+} (Zhou et al., 2010) to delete *Shox2* gene at later stages by generating *Avil*^{Cre/+}; *Shox2*^{flox/flox} mice. *Advillin* is a gene whose expression is largely restricted to peripheral sensory neurons. *Advillin* is weakly expressed in a few DRG neurons at E12.5 and reaches peak expression at E16.5 (da Silva et al., 2011; Hasegawa et al., 2007; Zhou et al., 2010). *Avil*^{Cre/+} mediated deletion of *Shox2* is completed at E18.5 (Figure 6A).

The *Avil*^{Cre/+}; *Shox2*^{flox/flox} mice are viable, fertile and appear normal. Using in situ hybridization, we found that the number of *TrkB* positive cells is only moderately decreased in these mutants compared to the controls (Figure 6B–C). Furthermore, there is no difference in the numbers of *TrkC*, *TrkA* or *Ret* positive neurons between *Avil*^{Cre/+}; *Shox2*^{flox/flox} and the control DRG (Figure 6D–F). Thus, deleting *Shox2* at stages after E12.5 resulted in a milder loss of *TrkB* expression and no effect on *TrkC* expression, suggesting *Shox2* is required primarily at the very early stages of mechanosensory neuron differentiation to ensure normal Trk receptor expression.

Effects of induced constitutive expression of *Shox2* in all neural crest derived cells

To gain further insight into the functions of *Shox2*, we asked whether over-expressing *Shox2* in all DRG neurons would have a dominant effect on *Trk* receptor expression. To achieve this, we created a knock-in mouse that allows Cre-dependent over-expression of *Shox2*. Briefly, a CAG promoter followed by loxP-STOP-loxP cassette, followed by *Shox2* cDNA and polyA was inserted into *Rosa26* locus (*Rosa*^{CAG-STOP-*Shox2*}). The knock-in mice were crossed to *Wnt1-Cre* to obtain *Wnt1-Cre; Rosa*^{CAG-STOP-*Shox2*/+} mice. In situ hybridization confirmed the over-expression of *Shox2* mRNA in all DRG cells (Figure 7A). *Wnt1-Cre; Rosa*^{CAG-STOP-*Shox2*/+} mice die immediately at birth, due to a completely cleft palate (data not shown). We examined the expression of *Trks* and *Ret* in these mice at E18.5 and P0 stages. Upon quantification, we found a 20% increase in the number of *TrkB* positive neurons in DRG from *Shox2*-overexpression embryos (Figure 7B–C). However, no apparent changes in *TrkA*, *TrkC* and *Ret* expression were detected in these mice compared to the controls (Figure 7D–F). Thus, although *Shox2* is necessary for inducing and/or maintaining *TrkB* expression in subsets of mechanosensory neurons, it is not sufficient to induce *TrkB* or suppress *TrkC* expression in all DRG neurons.

Defects in mechanosensory neuron central innervations in *Shox2*-deficient mice

Finally, we examined the consequences of loss of *Shox2* on the peripheral and central axonal projections of mechanosensory DRG neurons. To visualize the axonal projections, we utilized the *Avil*^{PLAP/+} mice in which human placenta alkaline phosphatase (PLAP) is inserted into the *Advillin* locus (Hasegawa et al., 2007). We crossed *Wnt1-Cre; Shox2*^{flox/+} males to *Avil*^{PLAP/+}; *Shox2*^{flox/flox} females to generate *Avil*^{PLAP/+}; *Wnt1-Cre; Shox2*^{flox/flox} mutant mice and their littermate controls. Since no mutant mice survive past P2, we examined axonal projection at E18.5, P0 and P2. At these stages, the peripheral axons have reached their targets, but have not yet fully differentiated into specialized sensory endings (Albuerne et al., 2000; Hasegawa et al., 2007), thereby preventing us from definitively determining the exact morphological subtypes of neurons that are affected by *Shox2* deletion. Using PLAP staining and anti-PGP9.5 staining, we did not detect any apparent differences in the general peripheral sensory projections into the hairy or the glabrous skin (data not shown).

However, PLAP staining on spinal cord sections from control or *Avil*^{PLAP/+}; *Wnt1-Cre; Shox2*^{flox/flox} mutant mice revealed that the *Shox2*-deleted DRG neurons showed reduced central axonal innervation to layers III/IV of the spinal cord compared to the control (Figure 8). Note that layer III/IV stained strongly for PLAP in the control (arrow in Figure 8A), but

the corresponding region in the *Wnt1-Cre; Shox2^{flox/flox}* mutant spinal cord stained much weaker (Figure 8A). On average, there is a 10% reduction in the PLAP-staining intensity in *Shox2*-mutant ($p < 0.001$). Since layers III/IV receive inputs from mechanosensory neurons, including *TrkB* and *Ret*-expressing touch neurons, this observation suggests that *Shox2* deficiency causes central innervation defects in subsets of mechanosensory neurons. Note that the reduced PLAP staining in layer III/IV was not due to changes in the expression of *Advillin* locus as PLAP staining in the DRG cell bodies and peripheral axons were equally intense, and in situ hybridization showed similar level of *Advillin* expression in both control and *Shox2*-deleted DRG (data not shown).

To confirm the central axon innervation defects with an independent method, we also used *Isl1^{cre/+}* (Srinivas et al., 2001) to conditionally delete *Shox2*. Islet1 (*Isl1*) is expressed in all sensory neurons beginning at E10, thus in these mice *Shox2* is deleted at early stages of DRG neuron differentiation. In *Isl1^{cre/+}; Shox2^{flox/flox}* mice, we observed 53% reduction in the number of *TrkB* expressing cells, only slightly less than what we observed in *Wnt1-Cre; Shox2^{flox/flox}* mice ($p < 0.001$, data not shown). We used anti-vGluT1 staining to specifically visualize mechanosensory afferent termini (whereas *Avil^{PLAP/+}* labels all axons including nociceptive afferents). The average vGluT1 staining intensity in layer III/IV showed an 11% reduction in *Isl1^{cre/+}; Shox2^{flox/flox}* dorsal spinal cord compared to that in controls (Figure 8B, $p < 0.002$). As a control, the vGLUT staining signals in DRG neuronal cell bodies were indistinguishable between control and *Shox2*-mutant mouse (insets in Figure 8B). In addition, the average area covered by vGluT1 positive mechanosensory axons in dorsal horn spinal cord is also reduced by about 20% in *Isl1^{cre/+}; Shox2^{flox/flox}* mice (Figure 8B, $p < 0.001$). Not surprisingly, nociceptive innervation to the dorsal horn as revealed by anti-CGRP and IB4 staining (to visualize peptidergic and nonpeptidergic nociceptive central axonal projections, respectively) was unchanged in *Shox2*-mutant mice (Figure 8C). Taken together, our study uncovered an important requirement for the transcription factor *Shox2* for the proper development, differentiation and central innervation of subset of *TrkB*-expressing mouse mechanosensory neurons.

Discussion

In the mammalian peripheral somatosensory system, mechanosensory and proprioceptive lineages arise from the same progenitor populations, through the first wave of neurogenesis from the precursor NCCs that migrate into the site of the future DRG at E9 (Fode et al., 1998; Ma et al., 1999). The initially *TrkC/TrkB* double positive progenitor neurons eventually differentiate into *TrkC* single positive proprioceptive neurons, as well as *TrkB* or *Ret* single positive touch sensory neurons, although a very small number of mechanosensory neurons are *TrkB/TrkC* or *Ret/TrkC* double positive. Previously, the molecular mechanisms regulating the specification of *TrkB*-expressing mechanosensory neuron lineage are unknown. In this study, we discovered that *Shox2* is an essential, although not sufficient, component required for the proper development of a subpopulation of *TrkB* positive DRG neurons.

Wnt1-Cre mediated deletion of *Shox2* in NCCs result in a 60–65% reduction in the number of *TrkB*-expressing DRG neurons at stages after E16.5. *Shox2*-deletion also caused a small increase in *TrkC/Ret* double positive neurons at later stages. These findings suggest that *Shox2* is important for ensuring *TrkB* expression and may contribute to *TrkC* repression in *Ret*-expressing touch sensory neurons, although at present we do not yet know whether such effects of *Shox2* is direct or indirect.

Later deletion of *Shox2* using *Avil^{Cre/+}* resulted in only a moderate decrease in *TrkB* positive neurons than that observed in *Wnt1-Cre* mediated deletion mice, suggesting *Shox2*

is required in progenitor and early born neurons to promote their differentiation into TrkB-expressing touch neurons. The onset of Advillin expression is at E12.5 and reaches a maximum at E16.5. It is likely that only those neurons that express Cre at early stages (E12.5) are affected and lose TrkB expression, whereas those that express Cre after E12.5 are not affected by *Shox2* deletion. Perhaps that once the stable high-level expression of TrkB is established, *Shox2* is no longer needed for the maintenance of TrkB expression. In our gain-of-function studies, overexpression of *Shox2* in all NCC progenitor cells only mildly increased the number of *TrkB*-positive cells, but had no effects on other *Trks* or *Ret* receptor expressions, including *TrkC*. Taken together, *Shox2* is necessary for inducing or maintaining TrkB expression in a subpopulation of mechanosensory neurons, but itself alone is not sufficient to induce TrkB expression in non-mechanosensory neurons. A model summarizing previous and current finding related to proprioceptive and mechanosensory neuron specification is shown in Figure 9.

At present, we do not know the downstream targets of *Shox2*, nor do we know whether *Shox2* only regulates Trk receptor expression or whether it also regulates other genes involved in other aspects of the touch/mechanosensory neuronal development. In other sensory lineages, the transcription factors Runx3 and Runx1 control the whole gene expression programs relevant to proprioceptive or nociceptive sensory neurons development and differentiation, respectively (Chen et al., 2006; Inoue et al., 2008; Kramer et al., 2006). On the other hand, the transcription factor Klf7 is required only for TrkA gene expression by binding to an enhancer element in the TrkA promoter. The loss of Klf7 leads to increased apoptosis of nociceptive sensory neurons. However, Klf7 did not appear to regulate other aspects of the differentiation program of TrkA-positive neurons (Lei et al., 2005). Future work is needed to determine the transcription targets of *Shox2*.

In *Shox2*-deficient mice, the central afferent innervations from mechanosensory neurons to layers III/IV in the spinal cord are reduced. Again, this could be a direct consequence of failed differentiation of a subset of TrkB-expressing touch neurons, or a secondary effect due to the loss of TrkB expression. Unlike the wealth of marker genes known for proprioceptive or nociceptive neurons, very limited molecular markers are known to specifically label TrkB-expressing mechanosensory neurons. We, therefore, could not examine other molecular aspects of mechanosensory neuron development and differentiation in *Shox2* mutant mice. Nonetheless, *Shox2* is the first transcription factor discovered that affects TrkB expression in mechanosensory DRG neurons.

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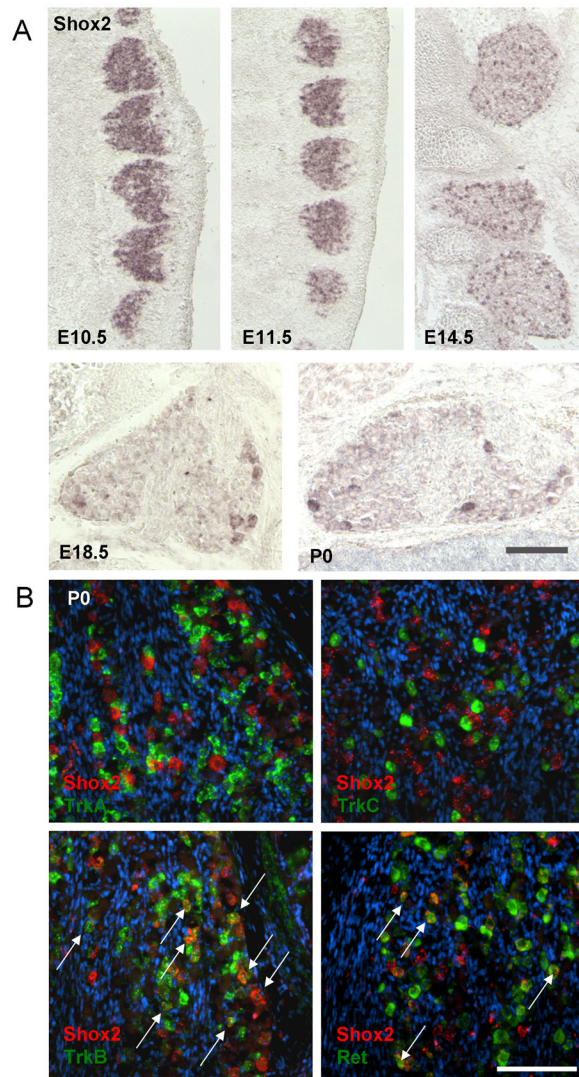


Figure 1. *Shox2* expression in the developing DRG

A. In situ hybridization experiments show that at E10.5 and E11.5, *Shox2* is expressed throughout the entire DRG; at E14.5, *Shox2* expression begins to down-regulate in subsets of DRG neurons; and by E18.5 and P0, *Shox2* is only present in a small number of DRG neurons. **B.** Two color in situ hybridization with probes for *Shox2* (red) and different receptors (*TrkA*, *TrkB*, *TrkC* and *Ret*) (green). Arrows indicate cell bodies in the ganglia showing co-localization of *Shox2* with either *TrkB* or *Ret*. Note that not all *TrkB* or *Ret* positive cell bodies have *Shox2* expression at P0. Scale bar = 100um

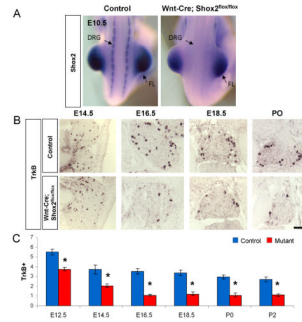


Figure 2. Reductions in the number of *TrkB*-expressing DRG neurons in *Wnt1-Cre*; *Shox2*^{flox/flox} embryos

A. In situ hybridization reveals the absence of *Shox2* expression in DRGs of *Wnt1-Cre*; *Shox2*^{flox/flox} mice at E10.5. **B.** Representative images of *TrkB* expression in control and *Shox2*-deleted DRG at E14.5, E16.5, E18.5 and P0. **C.** Quantifications of the numbers of *TrkB*-expressing DRG neurons per unit area in control and *Wnt1-Cre*; *Shox2*^{flox/flox} embryos. **p*<0.001. Error bars represent ±SEM. Scale bar = 100µm.

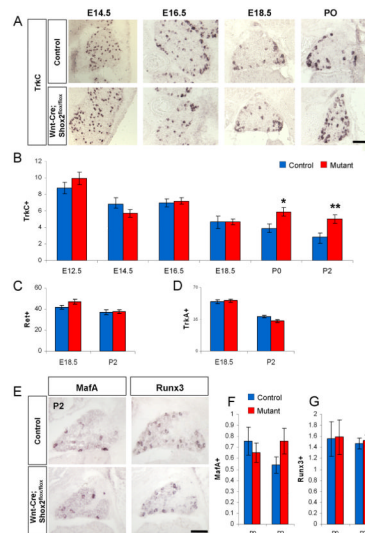


Figure 3. Increases in the number of *TrkC*-expressing DRG neurons in *Wnt1-Cre; Shox2^{flox/flox}* mice at perinatal stages

A. Representative images of *TrkC* expression in control and *Shox2*-deleted DRG at E14.5, E16.5, E18.5 and P0. **B.** Quantifications of the numbers of *TrkC*-expressing DRG neurons per unit area in control and *Wnt1-Cre; Shox2^{flox/flox}* embryos at 6 different stages. Note that there is a increase in the number of *TrkC* expressing neurons in the mutant at P0 (* $p < 0.005$) and P2 (** $p < 0.001$). **C and D.** Quantification of the number of Ret and TrkA expressing neurons in control and *Wnt1-Cre; Shox2^{flox/flox}* mice at E18.5 and P2. Error bars represent \pm SEM. Scale bar = 100 μ m. **E.** Representative images of *MafA* and *Runx3* expression in control and *Shox2*-deleted DRGs at P2. **F and G.** Quantifications show no significant differences in the numbers of cells expressing *MafA* or *Runx3* per unit area between the control and *Shox2*-deleted DRG at P0 and P2.

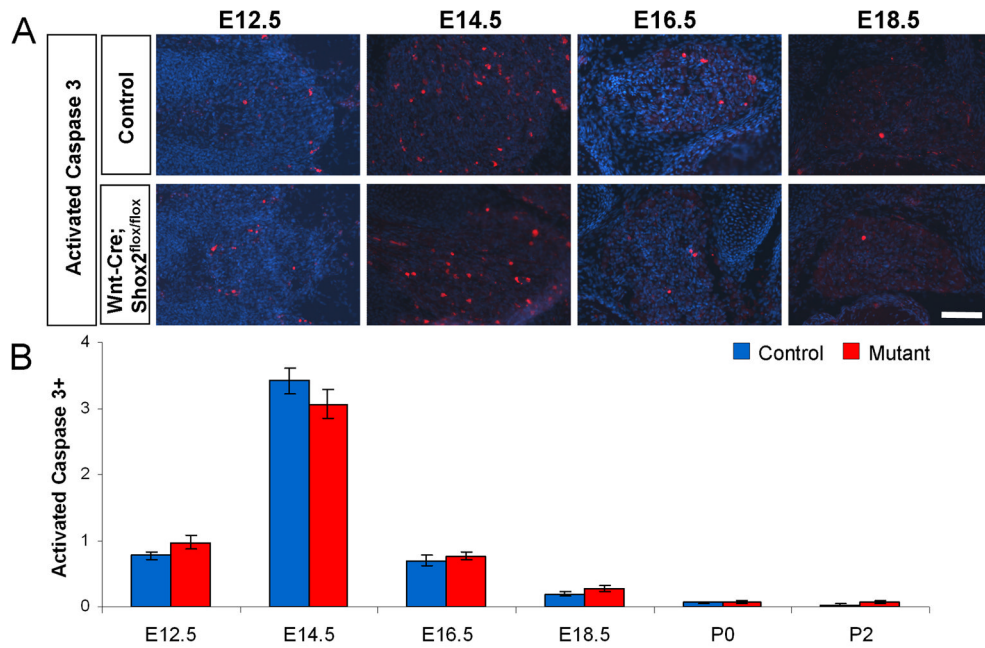


Figure 4. Apoptosis is normal in *Wnt1-Cre; Shox2^{flox/flox}* DRG

A. Representative images of immunostaining with anti-activated-caspase-3 in control and *Shox2*-deleted DRG at E12.5, E14.5, E16.5, and E18.5. **B.** Quantifications show no significant differences in the amount of apoptotic cells between control and *Shox2*-deleted DRG. Note that at E14.5, a time point of natural occurring cell death, there is an increase in activated-caspase-3 positive cells in both control and mutant DRG. Error bars represent \pm SEM. Scale bar = 100 μ m.

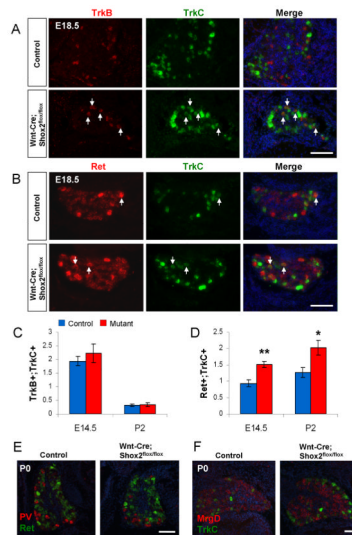


Figure 5. Increased co-expression of *Ret* and *TrkC* in DRG neurons in *Wnt1-Cre; Shox2^{flox/flox}* mice

A. Representative results of two-color in situ hybridization using *TrkB* (red) and *TrkC* (green) probes at E18.5. Arrows indicate neurons in the DRG that are positive for both receptors. **B.** Representative results of two-color in situ hybridization using *Ret* (red) and *TrkC* (green) probes at E18.5. Arrows indicate cell bodies in the DRG that are positive for both receptors. **C.** Quantification of the average number (per unit area) of *TrkB/TrkC* double positive neurons revealed no significant difference between the control and *Shox2*-deleted mice. **D.** Quantification of the average number (per unit area) of *Ret/TrkC* double positive neurons showed a significant increase in the *Shox2*-deleted versus control DRG. **E.** Representative images of two color in situ hybridization results with *Parvalbumin* (red) and *Ret* (green) probes at P0. **F.** Representative images of two color in situ hybridization results with *MrgD* (red) and *TrkC* (green) probes at P0. * $p < 0.01$ and ** $p < 0.001$. Scale bar = 100 μ m.

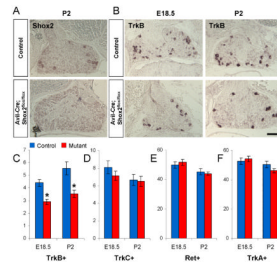


Figure 6. Mild reduction in the number of *TrkB*-expressing cells in *Avil^{Cre/+}; Shox2^{lox/lox}* mice
A. In situ hybridization confirms the loss of *Shox2* expression in the DRG from *Avil^{Cre/+}; Shox2^{lox/lox}* mice. **B.** Representative images of *TrkB* expression in control and *Avil^{Cre/+}; Shox2^{lox/lox}* mice at E18.5 and P2. **C–F.** Quantifications of the numbers of *TrkB* (C), *TrkC* (D), *Ret* (E), and *TrkA* (F) expressing DRG neurons per unit area in control and *Avil^{Cre/+}; Shox2^{lox/lox}* mice at E18.5 and P2. **p*<0.001. Error bars represent ±SEM. Scale bar = 100 μm.

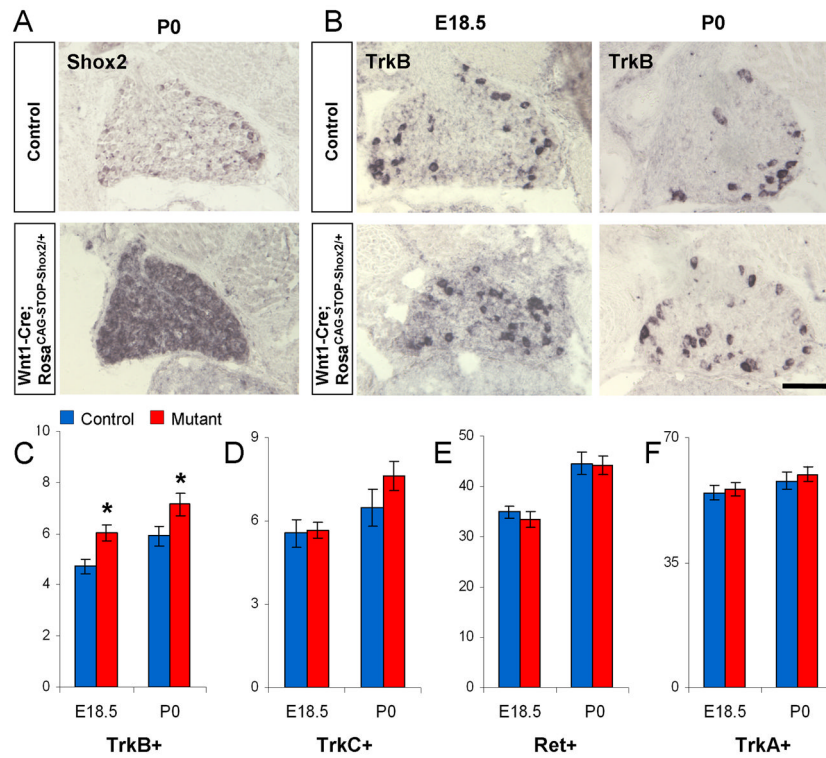


Figure 7. Over-expression of *Shox2* in all sensory neurons results in a mild increase in the number of *TrkB*-positive cell
A. Representative in situ hybridization images confirming the overexpression of *Shox2* in all DRG neurons in *Wnt1-Cre; Rosa^{CAG-STOP-Shox2}* mice. **B.** Representative images of *TrkB* in situ hybridization results in control and *Shox2* overexpression mice. **C–F.** Quantifications of the numbers of *TrkB* (**C**), *TrkC* (**D**), *Ret* (**E**), and *TrkA* (**F**) expressing DRG neurons per unit area in control and *Shox2* overexpression mice at E18.5 and P0. * $p < 0.05$. Error bars represent \pm SEM. Scale bar = 100 μ m.

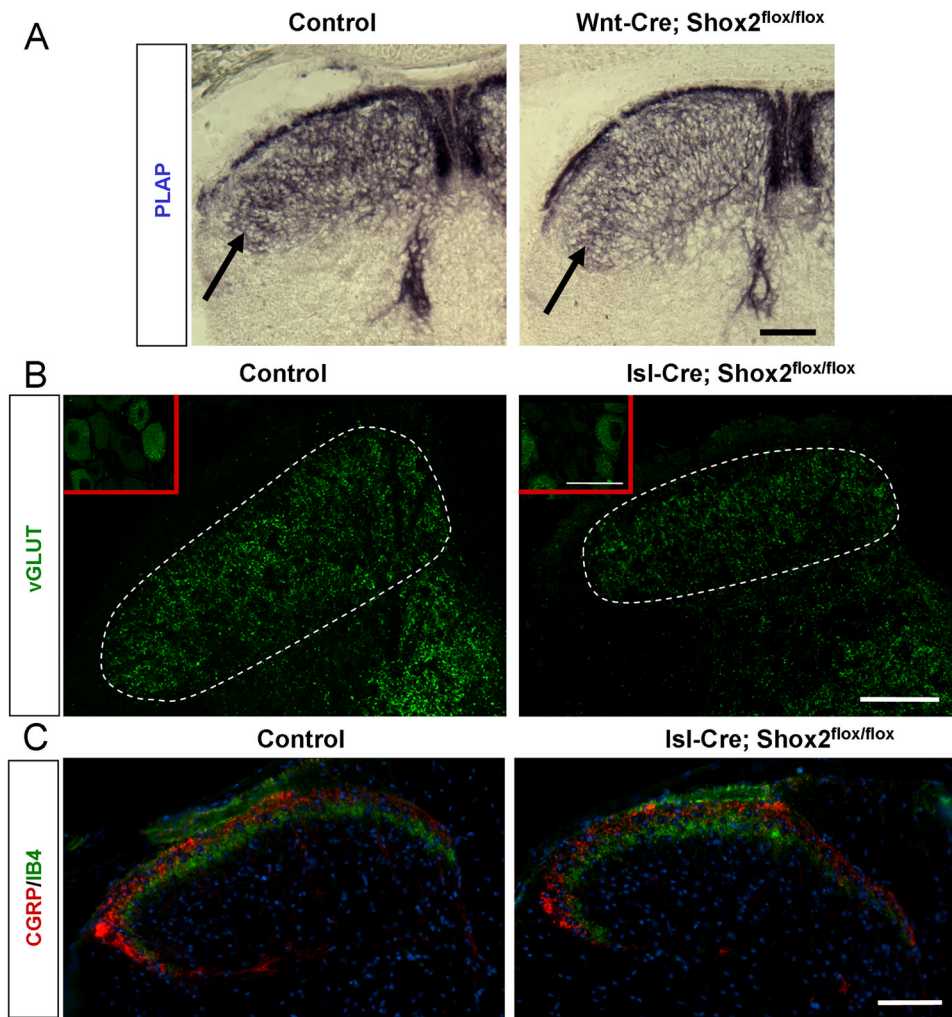


Figure 8. Reduced mechanosensory central innervation in the spinal cord in *Shox2*-deleted mice
A. Representative images of sensory afferent projections in the spinal cord from control (*Wnt1-Cre; Shox2^{flox/+}; Avil^{PLAP/+}*) and *Shox2*-deleted mice (*Wnt1-Cre; Shox2^{flox/flox}; Avil^{PLAP/+}*) as revealed by PLAP staining at P0. Arrows point to a densely stained band in layer III/IV in control that is only moderately stained in the mutant. **B.** Representative images of immunofluorescence staining with anti-vGluT1 in the spinal cord of control (*Isl1-Cre; Shox2^{flox/+}*) and *Shox2*-deleted (*Isl1-Cre; Shox2^{flox/flox}*) mice. Inset shows the anti v-GluT1 immunofluorescence signal in the DRG of control and *Shox2*-deleted mice. **C.** Representative images of immunofluorescence staining with anti-CGRP (red) and anti-IB4 (green) in the spinal cord of control and *Shox2*-deleted mice. Blue is DAPI. Scale bar in A and C is 100 μ m. Scale bar in B is 50 μ m.

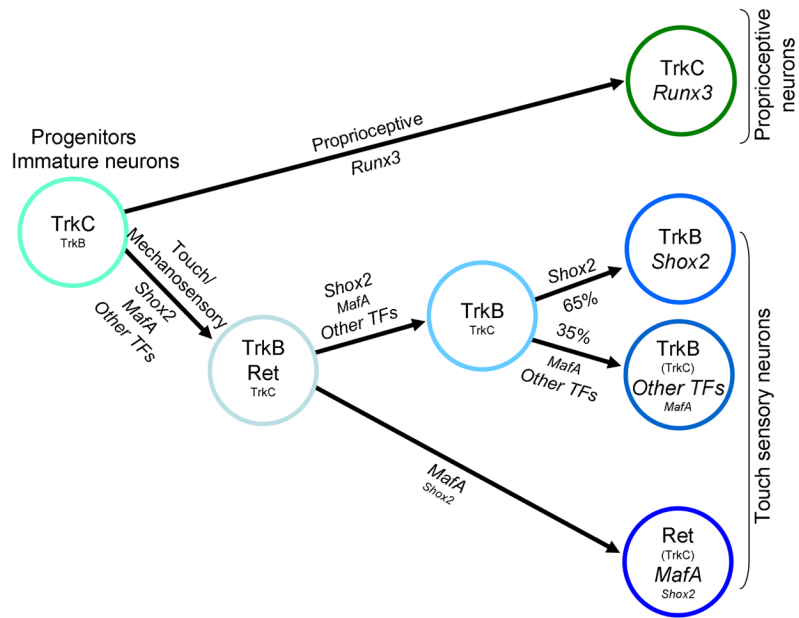


Figure 9. A model for the development and diversification of proprioceptive and touch sensory neurons
 Schematic model shows the transcription factors involved in the progressive specification of TrkB/TrkC double positive progenitor/immature neurons into proprioceptive and different types of touch sensory neurons.