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Supplemental information

Loss of *Prdm12* during development, but not in mature nociceptors, causes defects in pain sensation

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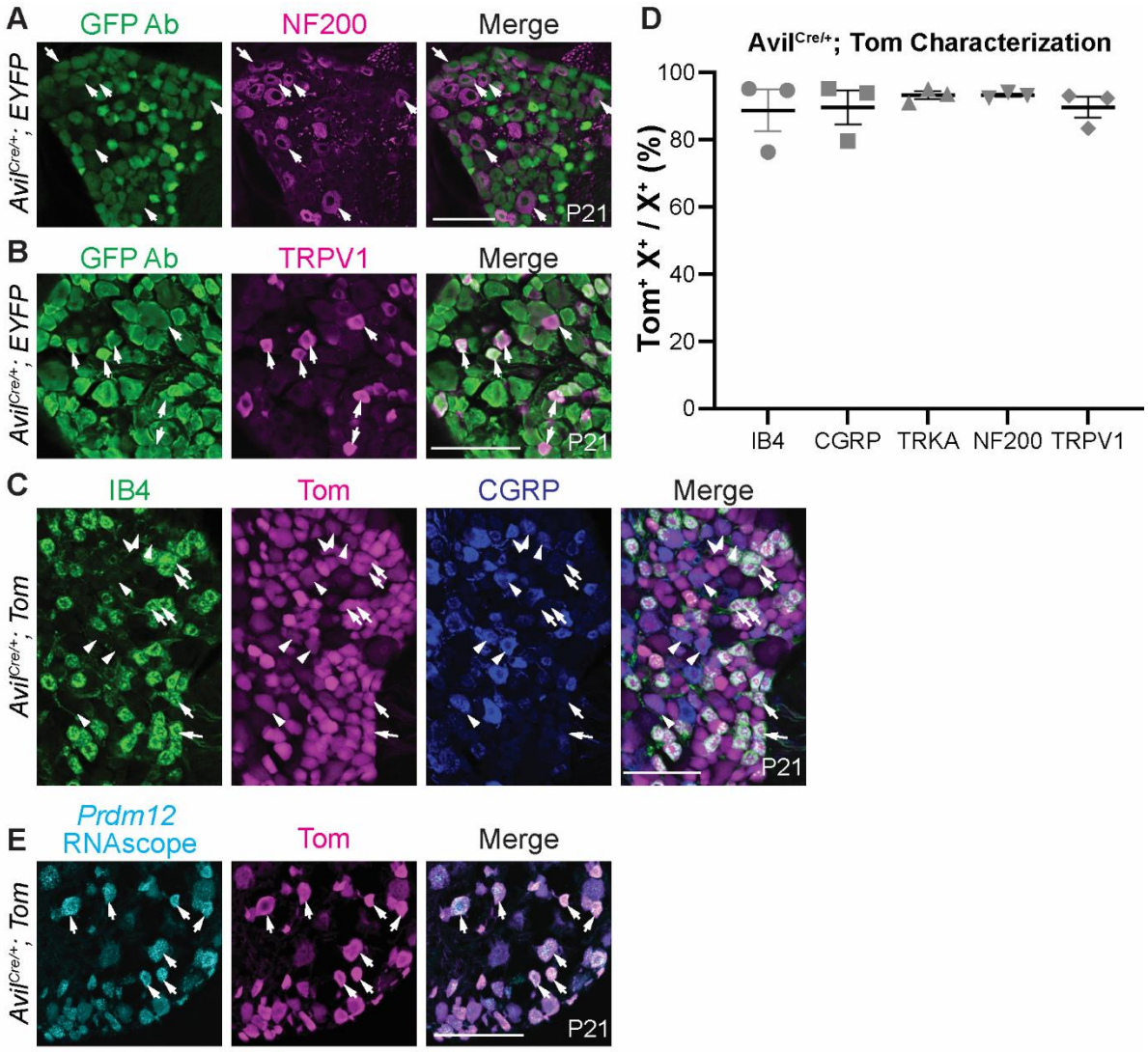

Supplemental Figure 1. Characterization of a sensory neuron specific CRE line. Related to Fig. 3. *Avil*$^{Cre/+}$ crossed to a CRE-dependent reporter identifies DRG neurons expressing CRE recombinase. (A-C) *Avil*$^{Cre/+}$-lineage neurons colocalize with myelinated DRG neurons (A, NF200^{+}, arrows) unmyelinated TRPV1^{+} neurons (B, arrows), and nonpeptidergic (C, IB4^{+}, arrows) and peptidergic (C, CGRP^{+}, arrowheads) C-fibers. (D) Quantification of Tom^{+} neurons as a percent of neurons expressing other markers in the *Avil*$^{Cre/+}$ lineage. Counts are the average of three L2-5 DRG sections per animal in n=3 mice. (E) The *Avil*$^{Cre/+}$-lineage also colocalizes with *Prdm12* mRNA expression (arrows, RNAscope). Scale bars 100 μm.

Table S3. Sequences for qPCR and genotyping primers, and the *Prdm12* exon V probe.

Related to the STAR Methods.

Sequence Type	Name	Sequence	Reference
qPCR Primers	Prdm12 exon 1 F	TCAAAACAGGGCTGAAGGCT	Desiderio et al. 2019
	Prdm12 exon 1 R	TTGCGCCATCGGCCATATAA	
	Prdm12 exon 5 F	CTTGGGGAGGGCAATGCTTA	Desiderio et al. 2019
	Prdm12 exon 5 R	TCCAATCACAGCCAAGGGTC	
	Ntrk1 F	AGAGTGGCCTCCGCTTTGT	Bartesaghi et al. 2019
	Ntrk1 R	CGCATTGGAGGACAGATTCA	
	Calca F	TTTGAGGTCAATCTTGGAAAGCA	Harvard PrimerBank
	Calca R	CTGAGCAGTGACACTAGAGCC	
	Scn10a F	ACCGACAATCAGAGCGAGGAG	Laedermann et al. 2014
	Scn10a R	ACAGACTAGAAATGGACAGAATCACC	
	Mrgprd F	TTTTCAGTGACATTCCTCGCC	Harvard PrimerBank
	Mrgprd R	GCACATAGACACAGAAGGGAGA	
	Trpv1 F	CCGGCTTTTTGGGAAGGGT	Harvard PrimerBank
	Trpv1 R	GAGACAGGTAGGTCCATCCAC	
	Trpm3 F	CAAGATGCCTGCCGTTTTTCT	Harvard PrimerBank
	Trpm3 R	GTCTAATTGGCAACTACCCCAAA	
	Trpa1 F	GTCCAGGGCGTTGTCTATCG	Harvard PrimerBank
	Trpa1 R	CGTGATGCAGAGGACAGAGAT	
	Trpv2 F	TGCTGAGGTGAACAAAGGAAAG	Harvard PrimerBank
	Trpv2 R	TCAAACCGATTTGGGTCCTGT	
	Trpm8 F	ACAGACGTGTCCTACAGTGAC	Harvard PrimerBank
	Trpm8 R	GCTCTGGGCATAACCACACTT	
	Sst	ACCGGGAAACAGGAACTGG	Harvard PrimerBank
	Sst	TTGCTGGGTTCGAGTTGGC	
	Nefh F	AGCCTGCACTACTCGCTGA	Harvard PrimerBank
	Nefh R	GGCCGTTGCTTAGGGTGTC	
	Ntrk2 F	CTGGGGCTTATGCCTGCTG	Harvard PrimerBank
	Ntrk2 R	AGGCTCAGTACACCAAATCCTA	
	Ntrk3 F	CTGAGTGCTACAATCTAAGCCC	Harvard PrimerBank
	Ntrk3 R	CACACCCCATAGAACTTGACAAT	
	Pvalb F	ATCAAGAAGGCGATAGGAGCC	Harvard PrimerBank
	Pvalb R	GGCCAGAAGCGTCTTTGTT	
	Atf3 F	AGCTGAGATTCGCCATCCAGAA	Laedermann et al. 2014
	Atf3 R	CTCGCCGCCTCCTTTTCCT	
	Kcnma1 F	AAGACGAAGGAGGCCCAGAA	Laumet et al. 2015
	Kcnma1 R	TCAGTGTCTGGGCGGATATC	
	Hprt F	GTCCTGTGGCCATCTGCCTAG	

	Hprt R	TGGGGACGCAGCAACTGACA	Harvard PrimerBank
Genotyping Primers	Prdm12dlg	GGTGCAAGGTCCAAATGTCT	Chen et al. 2020
	Prdm12drg/Null R	CACACTGCAGCTGGCTAAAC	
	Prdm12 Null F	TTCGGATGAGATGGCTAAGG	
	Cre F	GGACATGTTCAGGGATCGCCAGGCG	This lab
	Cre R	GCATAACCAGTGAAACAGCATTGCTG	
	iCre F	AGATGCCAGGACATCAGGAACCTG	The Jackson Laboratory
	iCre R	ATCAGCCACACCAGACACAGAGATC	
ISH Probe	Prdm Exon V	GCAGGGCCGCGTCTCCACGCGGACATC TCTGTCCTAGATGACAGGAGAGGGGGAA CAGACGATCTCCAAAGAGAGCTGAGTAG TTGGGAGTGGAAGAGAGAGCTACCTCCT TGGCCCAGAGTGGGCTTGGGGAGGGCA ATGCTTATCTCTGCCTCCTGGGAATGGT GTCTTGATGTGTCACGAGTGGAGGGAAT GACGTTGACCCTTGGCTGTGATTGGAGC CCTAAAGTGATTAGCACAGACCGGGAGA GACCTGATCATTCGGGGGGCCCTGCTTT CCTTCAGGCTACTTTCCCTAGCCTCCAG CTCCCAGGGAAGGACAAGACGACAATAT ATTTTTTGAAGTGTGCTGGGCAGATTGA GAGTGGAGTCACAAGAGACATTTACCAC AGCCCGTGGGCAAGACCCTGACCCTTTT GGGGCTTCAGTTTCCCCATCTATAGAAC GCAGTAATTAGATCAGTGCTCCCTCTCC TAAATCTC	This paper