



PUBLISHER CORRECTION

Publisher Correction: Single-cell transcriptomic analysis of somatosensory neurons uncovers temporal development of neuropathic pain

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We apologize for the mistake in the Supplementary information Fig. S7 which was incorrectly presented as Supplementary information Fig. S9 online, the correct Fig. S7 is shown as below.

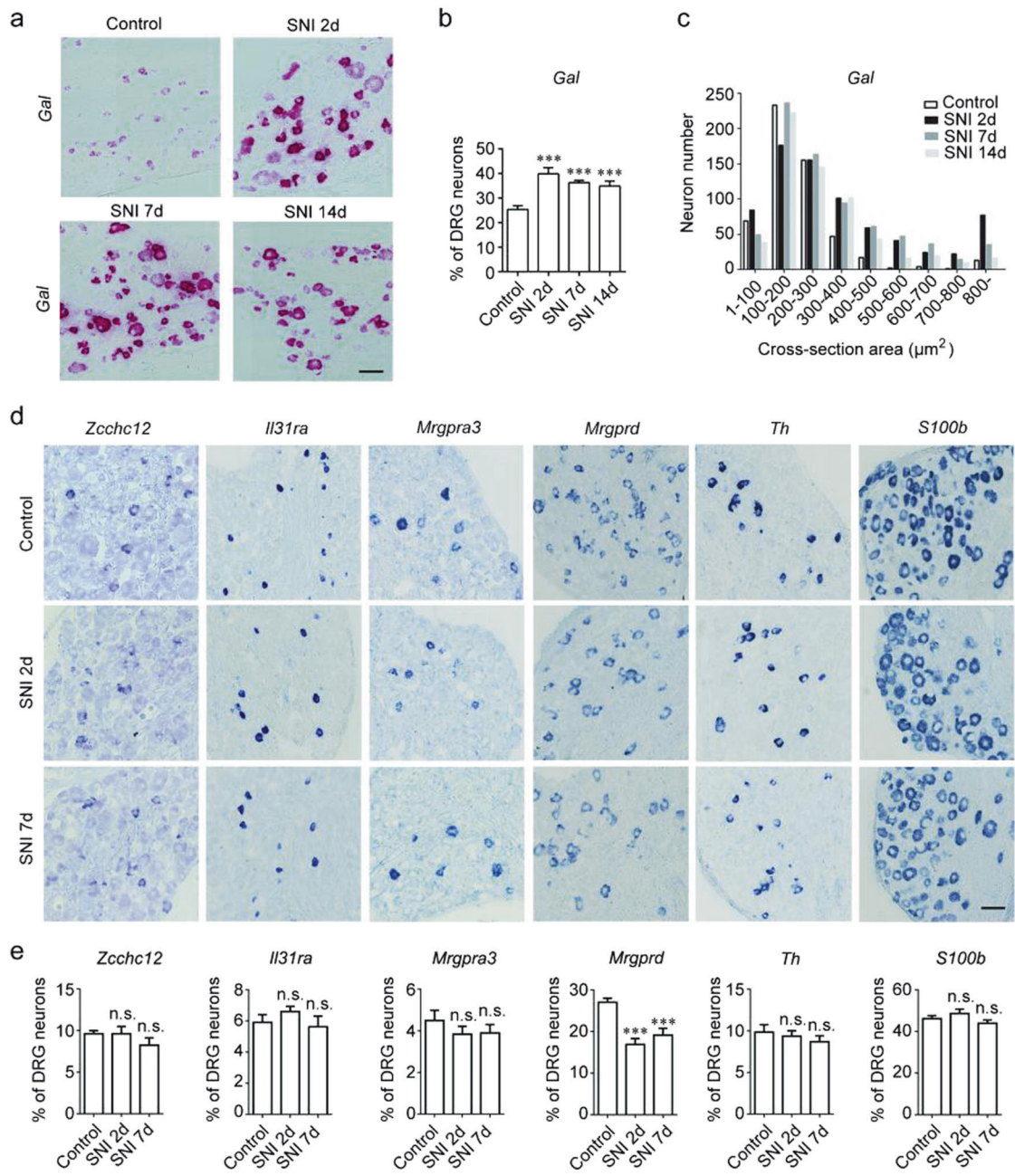


Fig. S7. Expression patterns of the marker genes for each neuron cluster after SNI. a, b RNA scope result shows *Gal* is upregulated after SNI. Scale bar, 50 μm . **c** Histogram showing the size changes of DRG neurons expressing *Gal* before and after SNI. *Gal* was highly expressed in small, medium, and large-diameter

neurons after SNI. **d, e** ISH shows the percentage of *Mrgprd*⁺ neurons was decreased after SNI, but no significant changes in other neuron clusters (labeled by *Zchc12*, *Il31ra*, *Mrgpra3*, *Th*, and *S100b*). Scale bar, 50 μm . The data are shown as mean \pm SEM. *** $P < 0.001$, n.s. no significant vs the control group.