

CORRECTION

Correction: Genome-Wide Analyses Reveal a Role for Peptide Hormones in Planarian Germline Development

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The authors would like to clarify the figure presentation of Figure 7. A corrected legend for <u>Figure 7</u> is provided here.



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Figure 7. Some prohormone genes are expressed differentially in the CNS of sexual and asexual planarians. Comparison of the ventral expression of (A) *ppl-1* or (B) *npy-8* between asexual, immature sexual hatchlings, and mature sexual animals. Whole animal images were obtained using dark-field microscopy. Enlarged views were captured at higher magnification using Rottermann Contrast imaging. Magnified views do not necessarily represent the same focal plane or same individual shown in the whole-animal image. Red dashed boxes indicate the general body regions from which these images were obtained. (C) Dorsal expression of *npy-8* in immature sexual hatchlings (left) and mature sexual animals (right). (D) Transparency rendering showing expression of *npy-8* in a cell in close proximity to testes lobes. Inset shows higher magnification of *npy-8*-expressing cell. (E) Northern blot comparing expression of *npy-8* in asexual animals (J. Stary and P. Newmark, unpublished observations) and is shown as a loading control. Scale bars: (A–C) 300 µm; (D) 10 µm.

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Reference

 Collins JJ III, Hou X, Romanova EV, Lambrus BG, Miller CM, Saberi A, et al. (2010) Genome-Wide Analyses Reveal a Role for Peptide Hormones in Planarian Germline Development. PLoS Biol 8(10): e1000509. doi: <u>10.1371/journal.pbio.1000509</u> PMID: <u>20967238</u>