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Supplemental Information

Innate Immunity in the *C. elegans* Intestine

Is Programmed by a Neuronal Regulator

of AWC Olfactory Neuron Development

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Figure S1. Loss-of-function mutations in *olrn-1* cause constitutive immune activation, Related to Figure 1. qRT-PCR data of *irg-4* (A) and *irg-5* (B) in wild-type, *olrn-1(ums9)*, *olrn-1(ums11)* and *olrn-1(ky626)* is presented. Data are the average of three independent replicates, each normalized to the control gene *snb-1* with error bars representing SEM, and are presented as the value relative to the average expression from all replicates of the indicated gene in wild-type animals. * equals p<0.05 by one-way ANOVA for the indicated comparison. **p=0.10 by one-way ANOVA and p<0.05 by two-tailed *t*-test. C. Data from a nanoString analysis of *irg-4* and *irg-5* expression, presented as the fold change of gene expression in *olrn-1(ums9)* mutants versus wild-type animals. Data are the average of two independent replicates for *olrn-1(ums9)* and one sample for wild-type, each normalized to three control genes (*snb-1, ama-1* and *act-1*) with error bars representing SEM. D. The lifespan of *olrn-1(ums9)* and *olrn-1(ky626)* mutant animals is shown relative to wild-type animals. Sample sizes, mean lifespan and p-values for each of two trials is shown in Table S2.



Figure S2. *olrn-1* **suppresses the p38 MAPK PMK-1 innate immune pathway, Related to Figure 2.** Venn diagrams show the overlap of genes induced in *olrn-1(ums9)* mutants with targets of **(A)** the bZIP transcription factor *zip-2*, **(B)** the G protein-coupled receptor *fshr-1*, and **(C)** the FOXO transcription factor *daf-16*. In A, B and C, the hypergeometric p value for the overlap between these datasets was not significant (n.s.).







Figure S4. Expression of *olrn-1* in chemosensory neurons is sufficient to regulate innate immunity in the intestinal epithelium, Related to Figure 5. qRT-PCR data of *irg-4* (A) and *irg-5* (B) in wild-type, *nsy-4(ky616)* and *nsy-5(ky634)* animals is presented. Data are the average of three independent replicates, each normalized to a control gene with error bars representing SEM, and are presented as the value relative to the average expression from all replicates of the indicated gene in wild-type animals. "n.s." equals not significant for the p value and * equals p<0.05 by one-way ANOVA for the indicated comparison.



Figure S5. Neuronal *olrn-1* regulates p38 MAPK PMK-1-dependent immune effector expression during nematode development, Related to Figure 6. A. Venn diagrams show the overlap of genes induced in *olrn-1(ums9)* mutants with genes that are upregulated at each larval stage in wild-type animals compared to wild-type adult animals. The hypergeometric p value for the overlap of each dataset is given in the figure. "n.s." equals not significant. qRT-PCR data of *irg-4* (B) and *irg-5* (C) in wild-type animals at the second or third larval stage (L2/L3), and the young adult (YA) stage are shown. Data are the average of four independent replicates, each normalized to a control gene with error bars representing SEM, and are presented relative to the average expression from all replicates of the indicated gene in animals at the young adult stage. * equals p<0.05 by one-way ANOVA for the indicated comparison.