

CORRECTION

Correction: The GATA Factor *elt-1* Regulates *C. elegans* Developmental Timing by Promoting Expression of the *let-7* Family MicroRNAs

The PLOS Genetics Staff

There are errors in [Table 1](#). The trans-heterozygous genotypes under ‘*elt-1* genotype’ are not displayed in the standard way; *ku491*+ should be *ku491*/+, *ok1002*+ should be *ok1002*/+ and *ku491ok1002* should be *ku491*/*ok1002*. The correct table is displayed underneath.



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Table 1. *elt-1* mutants have an L4-stage bursting vulva and defective alae formation and *elt-1(ku491)* is a partial loss-of-function allele.

Strain	RNAi	L4 Bursting Vulva			L4 Molt Alae			Young Adult Seam Cells		
		%	n	Absent (%)	Gapped (%)	Present (%)	n	SCM	Std Dev	n
wild-type	Empty vector ^a	0.0	245	0.0	0.0	100.0	27	16.0	0.4	27
<i>daf-12(rh61rh411)</i>	Empty vector ^a	0.0	183	0.0	0.0	100.0	24	18.1	1.6	24
<i>elt-1(ku491)</i>	Empty vector ^a	3.6	169	88.4	3.8	7.6	26	12.9	2.2	15
<i>elt-1(ku491);daf-12(rh61rh411)</i>	Empty vector ^a	55.1	198	88.8	0.0	11.1	18	39.7	9.8	29
wild-type	<i>elt-1</i>	48.7	117	100.0	0.0	0.0	21	0.5	0.8	21
<i>daf-12(rh61rh411)</i>	<i>elt-1</i>	56.7	90	90.0	5.0	5.0	20	0.2	0.4	20
<i>elt-1(ku491)</i>	<i>elt-1</i>	76.3	114	100.0	0.0	0.0	27	1.3	1.2	27
<i>elt-1(ku491);daf-12(rh61rh411)</i>	<i>elt-1</i>	94.8	231	96.4	3.6	0.0	28	2.0	2.2	28
<i>elt-1 genotype^b</i>		<i>daf-12 genotype</i>								
<i>ku491/+</i>	Wild-type	0.0	60	0.0	0.0	100.0	19	15.9	0.3	18
<i>ku491/+</i>	<i>rh61rh411</i>	0.0	57	0.0	0.0	100.0	18	18.3	1.8	18
<i>ok1002/+</i>	Wild-type	0.0	92	0.0	0.0	100.0	18	16.0	0.3	18
<i>ok1002/+</i>	<i>rh61rh411</i>	0.0	78	0.0	0.0	100.0	17	18.2	1.9	17
<i>ku491/ok1002</i>	Wild-type	1.9	52	84.0	16.0	0.0	25	9.4 ^c	2.0	25
<i>ku491/ok1002</i>	<i>rh61rh411</i>	65.9	41	100.0	0.0	0.0	20	18.2 ^d	6.6	20

^aPhenotypes on empty-vector control RNAi were similar to the standard *E. coli* strain OP50.

^bAn allele of *elt-1(ku491)* linked to mutations in *unc-24* and *dpy-20* was used for these strains.

^cFor *elt-1(ku491)* animals vs *elt-1(ku491)-over-elt-1(null)* animals at the young adult stage, the p-value for the comparison of seam-cell numbers is 0.0340

^dFor *elt-1(ku491);daf-12(rh61rh411)* animals vs *elt-1(ku491)-over-elt-1(null);daf-12(rh61rh411)* animals at the young adult stage, the p-value for the comparison of seam-cell numbers is < 0.0001.

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Reference

1. Cohen ML, Kim S, Morita K, Kim SH, Han M (2015) The GATA Factor *elt-1* Regulates *C. elegans* Developmental Timing by Promoting Expression of the *let-7* Family MicroRNAs. PLoS Genet 11(3): e1005099. doi: [10.1371/journal.pgen.1005099](https://doi.org/10.1371/journal.pgen.1005099) PMID: [25816370](https://pubmed.ncbi.nlm.nih.gov/25816370/)