

Supplementary Figure legends:

Figure S1: A model for linker cell death.

Based on ¹⁶.

Figure S2: Histone methyltransferase genes not relevant for linker cell death.

Error bars, SEM. Numbers within bars, no. animals.

Figure S3: NHR-67 protein binds sequences upstream of the *let-70* gene.

let-70 gene structure is shown. NHR-67 enrichment (\log_2) over background is shown in blue.

Supplementary Table S1. CPF components are not required for linker cell death

Genotype^a	% LC survival	n^b
Wild type + empty vector	3 ± 2	122
<i>ztf-1</i> (RNAi)	4 ± 2	90
<i>nrd-1</i> (RNAi)	6 ± 2	104
<i>B0035.11</i> (RNAi)	6 ± 2	100
<i>cpf-2</i> (RNAi)	0 ± 0	82
<i>pcf-11</i> (RNAi)	2 ± 2	52
<i>nab-3</i> (RNAi)	4 ± 2	90
<i>clpf-1</i> (RNAi)	7 ± 2	82
<i>pap-1</i> (RNAi)	2 ± 2	51

^aAll animals contained *rrf-3(pk1426)*; *him-8(e1489)* mutations and a *lag-2::gfp (qIs56)* reporter gene to identify the linker cell. ^bNumber of animals scored.