

Table S1. Genetic interaction of *ot101*

Genotype	ASEL fate in ASEL	ASEL fate in ASER	<i>n</i>
Wild type	100%	0%	>100
<i>ot101</i>	17%	0%	31
<i>ls[die-1r]</i>	99%	85%	62
<i>ls[die-1r]; ot101</i>	100%	59%*	85
<i>ls[ceh-36::lsy-6]</i>	99%	89%	54
<i>ls[ceh-36::lsy-6]; ot101</i>	76%	28%	79

Marker used: *lim-6::gfp*.

*Expression of marker dimmer in ASER than in ASEL in 40% of these animals.