

Table S1. TUNEL analysis for third instar Garland cells

Genotype	Imaged nuclei (sample)	TUNEL positive	
		Number of positive signals	%
<i>sns-GCN-nGFP, sns-GCN-Gal4/+</i>	181 (4)	6	3.31
<i>sns-GCN-nGFP, sns-GCN-Gal4/UAS-sns-IR(1)</i>	180 (4)	70	38.90
<i>sns-GCN-nGFP, sns-GCN-Gal4/UAS-sns-IR(2)</i>	93 (4)	38	40.86
<i>sns-GCN-nGFP, sns-GCN-Gal4/UAS-duf-IR(1)</i>	102 (3)	52	50.90
<i>sns-GCN-nGFP, sns-GCN-Gal4/UAS-duf-IR(2)</i>	109 (4)	18	16.51
<i>sns^{4.3}, Act5C-Gal4/sns^{4.3}, UAS-2xEGFP</i>	83 (4)	36	43.37
<i>sns-GCN-nGFP, sns-GCN-Gal4/UAS-pyd-IR</i>	144 (4)	14	9.72
<i>sns-Gal4/sns^{zf1.4}, UAS-sns[C-HA]</i>	66 (3)	0	0
<i>sns^{zf1.4}, sns-Gal4/sns^{zf1.4}, UAS-sns[C-HA]</i>	101 (4)	22	21.80
<i>sns-Gal4/sns^{zf1.4}, UAS-sns[mlg4]</i>	64 (4)	18	28.13
<i>sns^{zf1.4}, sns-Gal4/sns^{zf1.4}, UAS-sns[mlg4]</i>	34 (3)	16	47.06
<i>sns-Gal4/sns^{zf1.4}, UAS-sns[mlg6]</i>	92 (3)	0	0
<i>sns^{zf1.4}, sns-Gal4/sns^{zf1.4}, UAS-sns[mlg6]</i>	25 (3)	17	68.00