

Supplemental Table S6. Genes that are not sex-biased, yet down-regulated by Rsl in females.

Gene	Accession Number	Expression Ratio (average)			
		F-wt / M-wt	F- <i>rsl</i> / F-wt	F-R1 / F- <i>rsl</i>	F-R2 / F- <i>rsl</i>
<i>Cap2</i>	AV261931	1.03	4.55	0.73	1.00
<i>Alas1</i>	BM021574	0.87	2.33	0.66	0.52
<i>Fancc</i>	BB522905	0.96	2.17	0.60	0.67
<i>Zfp259</i>	AK016551	0.83	2.04	0.78	0.71
<i>Col5a2</i>	AV229424	1.03	2.04	0.66	1.08
<i>Phr1</i>	BM238052	0.90	1.69	0.51	0.70
<i>Gda</i>	AW911807	0.88	1.56	0.61	0.80
<i>Fzd8</i>	BB086994	1.06	1.54	0.75	0.90

Shown are the 8 non-sexually dimorphic genes that are increased in expression in *rsl* females ($\Delta \geq 1.50$ fold) and down-regulated in Rsl1 (R1) and Rsl2 (R2) transgenic females relative to wt.