

Table S2. *Post hoc* χ^2 comparisons of DNA methylation band numbers between the four treatment groups sampled as 4th and 5th instar larvae

Treatment	χ^2	d.f.	P-value
4th instar larvae			
Control and <i>IRS</i> ⁻	2.78	1	0.095*
Control and <i>TOR</i> ⁻	6.62	1	0.010*
Control and <i>IRS/TOR</i>	5.82	1	0.016*
<i>IRS</i> ⁻ and <i>IRS/TOR</i>	0.58	1	0.448 (n.s.)
<i>TOR</i> ⁻ and <i>IRS/TOR</i>	.03	1	0.872 (n.s.)
<i>IRS</i> ⁻ and <i>TOR</i> ⁻	0.85	1	0.357 (n.s.)
5th instar larvae			
Control and <i>IRS</i> ⁻	3.76	1	0.053*
Control and <i>TOR</i> ⁻	9.41	1	0.002**
Control and <i>IRS/TOR</i>	1.12	1	0.289 (n.s.)
<i>IRS</i> ⁻ and <i>IRS/TOR</i>	0.78	1	0.378 (n.s.)
<i>TOR</i> ⁻ and <i>IRS/TOR</i>	4.14	1	0.042*
<i>IRS</i> ⁻ and <i>TOR</i> ⁻	1.38	1	0.239 (n.s.)

Asterisks indicate significant differences between groups (* $P < 0.05$, ** $P < 0.005$).