

Sup. Table 4: Correlation between maternal stress during infancy and increased DNA methylation in adolescence in boys only (all CpG sites with FDR<20%). Sites with greater than 5% differential methylation are highlighted in grey.

Gene	CpG Site	Slope	Range	Differential Methylation	Spearman ρ	FDR
RASA2	cg20247102	0.043	0.045	0.025	0.581	0
MGC4562	cg15198335	0.015	0.021	0.009	0.536	0
PKN1	cg19103609	0.131	0.178	0.075	0.501	0

Note. FDR = false discovery rate (%). Slope = slope of least squared line. Range = range of average beta.