

Erratum to: Epigenetic and genetic variation in *GATA5* is associated with gastric disease risk

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Unfortunately, the online published article has formatting error in Table 5, that complicates interpretation.

The correct Table 5 is given in the following page.

The online version of the original article can be found under doi:[10.1007/s00439-016-1687-1](https://doi.org/10.1007/s00439-016-1687-1).

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Table 5 Variables affecting gastric histopathology scores in the methylation group

Independent variable(s)	β	95 % CI	<i>p</i> value
Age	0.026	(0.006, 0.045)	0.009
Region of origin	0.40	(0.10, 0.69)	0.008
Dominant rs6587239	-0.45	(-0.76, -0.15)	0.004
<i>GATA5</i> methylation	0.028	(0.012, 0.045)	0.001
Sex (male)	-0.28	(-0.58, 0.023)	0.070
<i>vacA</i> s	-0.59	(-0.96, -0.22)	0.002
Age	0.022	(0.005, 0.040)	0.013
Region of origin	0.18	(0.093, 0.46)	0.194
Dominant rs6587239	-1.20	(-2.07, -0.33)	0.007
<i>GATA</i> methylation	-0.012	(-0.044, 0.020)	0.463
<i>vacA</i> s2	-0.73	(-1.08, -0.39)	4.28×10^{-5}
Interaction of Dom rs6587239 * <i>GATA5</i> methylation	0.040	(0.003, 0.076)	0.033

n = 189

The top half of the table shows independent covariates in univariate tests

The lower half shows covariates in a single regression model