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| --- | --- | --- | --- | --- | --- |
| Table S4. Binary logistic regression model controlled for age and sex in methylation analysis. | | | | | |
| Characteristic | *β* | S.E. | Wald | df | *P* - value |
| Age | 0.011 | 0.015 | 0.582 | 1 | 0.445 |
| Sex | -0.829 | 0.371 | 5.006 | 1 | **0.025** |
| Methylation |  |  |  |  |  |
| CpG1 | -0.018 | 0.022 | 0.709 | 1 | 0.400 |
| CpG2 | 0.064 | 0.018 | 12.609 | 1 | **0.000** |
| CpG3 | -0.004 | 0.041 | 0.010 | 1 | 0.921 |
| Genotype |  |  | 0.349 | 2 | 0.840 |
| Val/Val | 0.190 | 0.664 | 0.082 | 1 | 0.774 |
| Val/Met | 0.331 | 0.657 | 0.255 | 1 | 0.614 |
| Constant | -2.074 | 1.266 | 2.686 | 1 | 0.101 |