|  |
| --- |
| 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  |