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| Table S2 Comparisons of *COMT* methylation levels of exon1 between cases and controls. | | | | |
| CpG locus  (%) | Controls  (Mean ± S.D.) | SCZ  (Mean ± S.D.) | *t* value | *P -* value |
| CpGa | 2.95±2.96 | 2.03±2.87 | -3.741 | 0.000 |
| CpGb | 2.31±3.42 | 1.71±2.27 | -0.545 | 0.586 |
| CpGc | 1.92±2.71 | 0.98±1.37 | -3.048 | 0.002 |
| CpGd | 3.86±8.90 | 2.23±2.63 | -1.283 | 0.200 |
| CpGe | 1.76±4.65 | 1.31±2.48 | -1.257 | 0.209 |
| CpGf | 3.58±9.24 | 2.23±4.66 | -1.070 | 0.284 |
| CpGg | 1.01±2.17 | 0.58±0.85 | -0.090 | 0.928 |