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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  |