

SUPPLEMENTARY INFORMATION

DNA Methylation Profiles of Blood Cells Are Distinct between Early-Onset Obese and Control Individuals

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Supplementary Table 2. Differentially expressed genes

Symbol	p-value	FDR
<i>GNG7</i>	5.43E-05	0.913233955
<i>LOC400890</i>	0.00010231	0.913233955
<i>RTN4</i>	0.000123827	0.913233955
<i>HDAC2</i>	0.000151315	0.913233955
<i>REEP5</i>	0.000164331	0.913233955
<i>MIR30A</i>	0.000236918	0.913233955
<i>TMOD1</i>	0.000237557	0.913233955
<i>XPR1</i>	0.000252695	0.913233955
<i>COL4A1</i>	0.000261597	0.913233955
<i>MSX2</i>	0.000294059	0.923903924
<i>UTP23</i>	0.000345272	0.957686345
<i>BCL9</i>	0.000440861	0.957686345
<i>MAP1LC3C</i>	0.000461436	0.957686345
<i>LOC644682</i>	0.000461498	0.957686345
<i>BTF3L4</i>	0.000476407	0.957686345
<i>MTPN</i>	0.00050862	0.957686345
<i>LOC646278</i>	0.000588438	0.957686345
<i>PTPN12</i>	0.000598659	0.957686345
<i>LOC284023</i>	0.000614055	0.957686345
<i>LOC729510</i>	0.000618711	0.957686345
<i>HIST1H3G</i>	0.000654751	0.957686345
<i>TRIM69</i>	0.000700616	0.957686345
<i>STON1</i>	0.000764487	0.957686345
<i>LOC652579</i>	0.000808139	0.957686345
<i>DTL</i>	0.000815974	0.957686345
<i>LOC652127</i>	0.000847992	0.957686345
<i>LOC647850</i>	0.000904861	0.957686345
<i>DYRK1A</i>	0.000942617	0.957686345
<i>DECR2</i>	0.000947844	0.957686345
<i>LOC100132964</i>	0.000992485	0.957686345

FDR, false discovery rate.