SUPPLEMENTARY INFORMATION

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

Je-Keun Rhee[1], Jin-Hee Lee[2], Hae Kyung Yang[3], Tae-Min Kim[1]*, Kun-Ho Yoon[1,2,3]**

[1]Department of Medical Informatics, College of Medicine, The Catholic University of Korea, Seoul 06591, Korea, [2]Catholic Institute of U-Healthcare, Institute of Biomedical Industry, The Catholic University of Korea, Seoul 06591, Korea, [3]Department of Internal Medicine, College of Medicine, The Catholic University of Korea, Seoul 06591, Korea

Supplementary Table 1. Expression comparison of genes with significant enrichment of
differentially methylated CpGs

Symbol	p-value	FDR
C21orf56	0.030503	0.957686
ZNF154	0.040895	0.957686
SDK1	0.23544	0.965175
KIAA0146	0.289321	0.965318
SKIV2L	0.094446	0.957686
GPR125	0.819418	0.99262
SORBS2	0.885967	0.997319
C14orf70	0.816179	0.992498
POLR3E	0.765737	0.989988
CTBP2	0.080374	0.957686
DLGAP2	0.224745	0.962153
CAPS2	0.588693	0.986149
GIMAP1	0.140035	0.957686
RNF213	0.547384	0.982494
MND1	0.954998	0.999836
SRM	0.070763	0.957686
TGM6	0.506865	0.977098
WDR27	0.124784	0.957686

FDR, false discovery rate.