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CORRECTION

Correction: Biologically anchored knowledge expansion approach uncovers *KLF4* as a novel insulin signaling regulator

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S2 Table is incomplete. The bottom part of S2 Table is missing. The full S2 Table can be viewed below.

Supporting information

S2 Table. L_0 and L_1 genes. L_0 represents genes that were differentially expressed between DW16 and DC16 adipocytes. L_1 represents genes in L_0 for which expression profiles significantly correlated with expression of insulin signaling pathway genes (L_{path}) in adipocytes using data for all four conditions DC8, DW8, DC16 and DW16 (marked L_1 in table).
(PDF)

Reference

1. Muthiah A, Angulo MS, Walker NN, Keller SR, Lee JK (2018) Biologically anchored knowledge expansion approach uncovers *KLF4* as a novel insulin signaling regulator. PLoS ONE 13(9): e0204100. https://doi.org/10.1371/journal.pone.0204100 PMID: 30240435

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