Table S5. TELIS analysis relating maternal disadvantage to bioinformatic indications of transcription factor activity in newborn cord blood.

Transcription Factor	Transfac TFBM	Mean Fold-Difference	Standard Error	p value
EVI1	V$EVI1_04	0.27	0.07	0.0001
STAT3	V$STAT3_01	0.32	0.1	0.004
NF-κB	V$NFKAPPAB_01	0.55	0.1	0.004
TAL1	V$TAL1BETAE47_01	0.63	0.12	0.024
AP1	V$AP1_Q6	0.75	0.07	0.004
C/EBP-β	V$CEBPB_02	0.64	0.12	0.029
EGR1	V$EGR1_01	2.19	0.79	0.012
EGR3	V$EGR3_01	2.01	0.74	0.029
GATA1	V$GATA_C	1.29	0.14	0.033
ETS1	V$CETS1P54_01	1.43	0.22	0.014
cMyb	V$VMYB_02	1.99	0.32	0.0001

Note. TELIS quantities the prevalance of transcription factor binding motifs in promoters of differentially expressed genes.
For details, see Cole SW et al. Bioinformatics. 2005;21 (6):803-810.