

Table S7. Differentially methylated fragments in neutrophils and placenta.

Criteria	R\geq6, S\geq6*
Significant fragments ($1.55 \times 10^{-6**}$ and > 0.25 diff)	6767
CpG sites within these	38307
Mean CpGs per fragment	5.66
Placenta hypo	6146
Mean CpGs per fragment (hypo)	5.56
Placenta hyper	621
Mean CpGs per fragment (hyper)	6.67
% of DMFs that are hypo meth in placenta	90.8

* R = number of neutrophil samples fulfilling coverage criteria and S = number of placenta samples fulfilling coverage criteria.

** Significance level = 0.05, adjusted with multiple test correction by the Bonferroni method.