

Figure S4. Histogram of the number of sample types in which each DMP is differentially methylated compared to whole blood. Taking all sites identified as having a significantly different level of DNA methylation compared to whole blood in at least one sample type ($n = 611,070$, ANOVA $P < 9 \times 10^{-8}$) we counted the number each of individual sample types characterized by differential DNAm ($P < 0.05$).

