

**Figure S13. Covariation in DNA methylation between whole blood and individual blood cell-types is higher in the subset of DNAm sites classed as having ‘intermediate’ levels of DNAm or being highly ‘variable’.** Shown are boxplots of variance explained in whole blood for each cell type separately where DNAm sites are split by mean DNA methylation level (x-axis, left panels) and variability (right panels).

