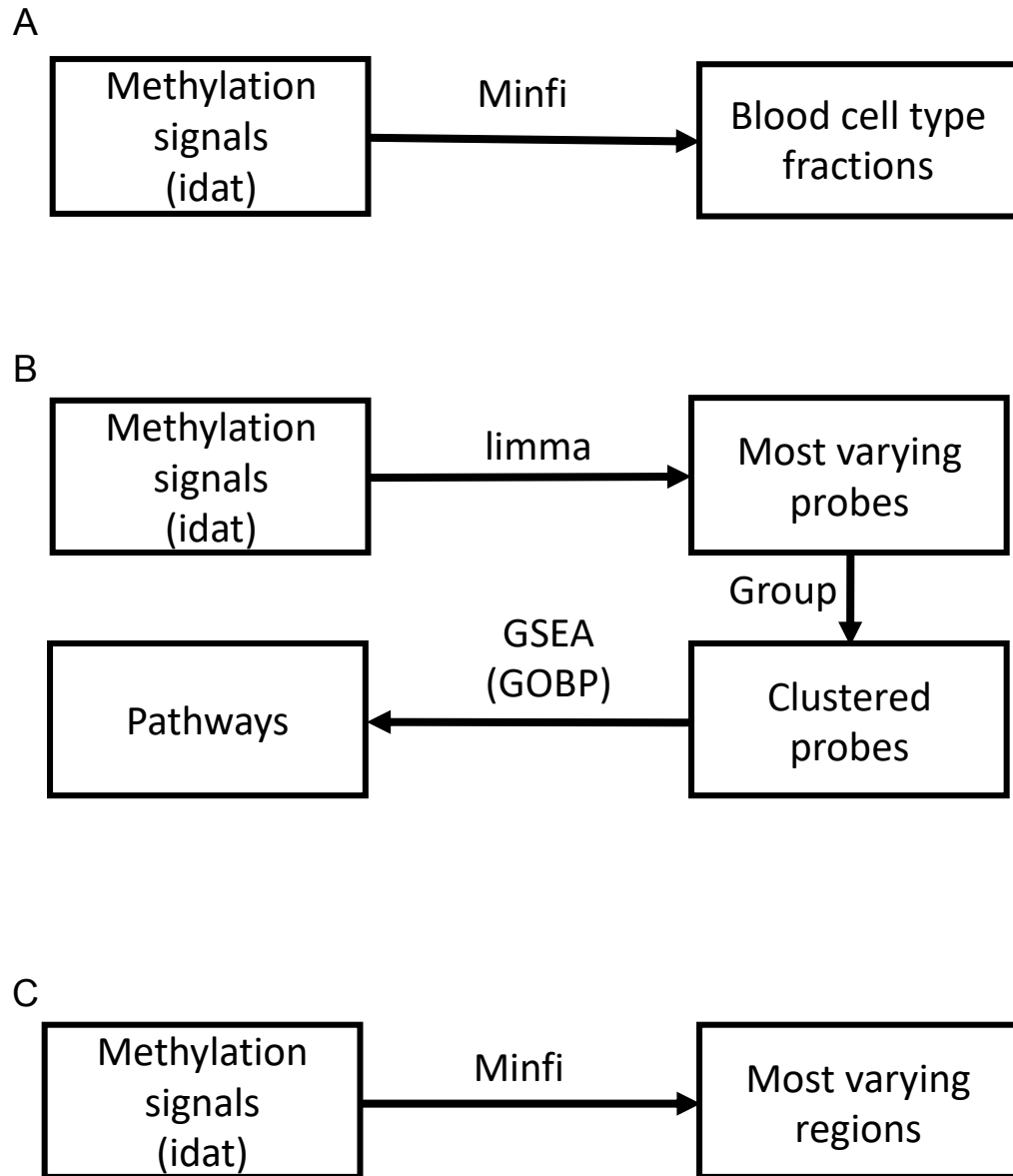




Supplementary Figure 3. *Blood DNA methylation bioinformatics pipeline.* **(A)** Blood cell fraction estimation by Minfi's function estimateCellCounts. **(B)** Detection of differentially methylated sites by limma, detection of methylation patterns by grouping sites for similar methylation changes over time, and pathway analysis by GSEA. **(C)** Detection of differential methylation regions by Minfi's function bumphunter.