



Supplemental Figure 12. (A) Histograms depicting the distribution of fractional methylation across all promoters in each of the 34 tissue types assessed. Each distribution is fit by a 3-component Gaussian mixture model to distinguish subpopulations of lowly methylated (LMP; red), intermediate methylated (IMP; green) and highly methylated (HMP; blue) promoters. (B) Boxplot of the fraction of MethExp promoters that belong to the HMP category across 34 tissue types.