

Figure S2

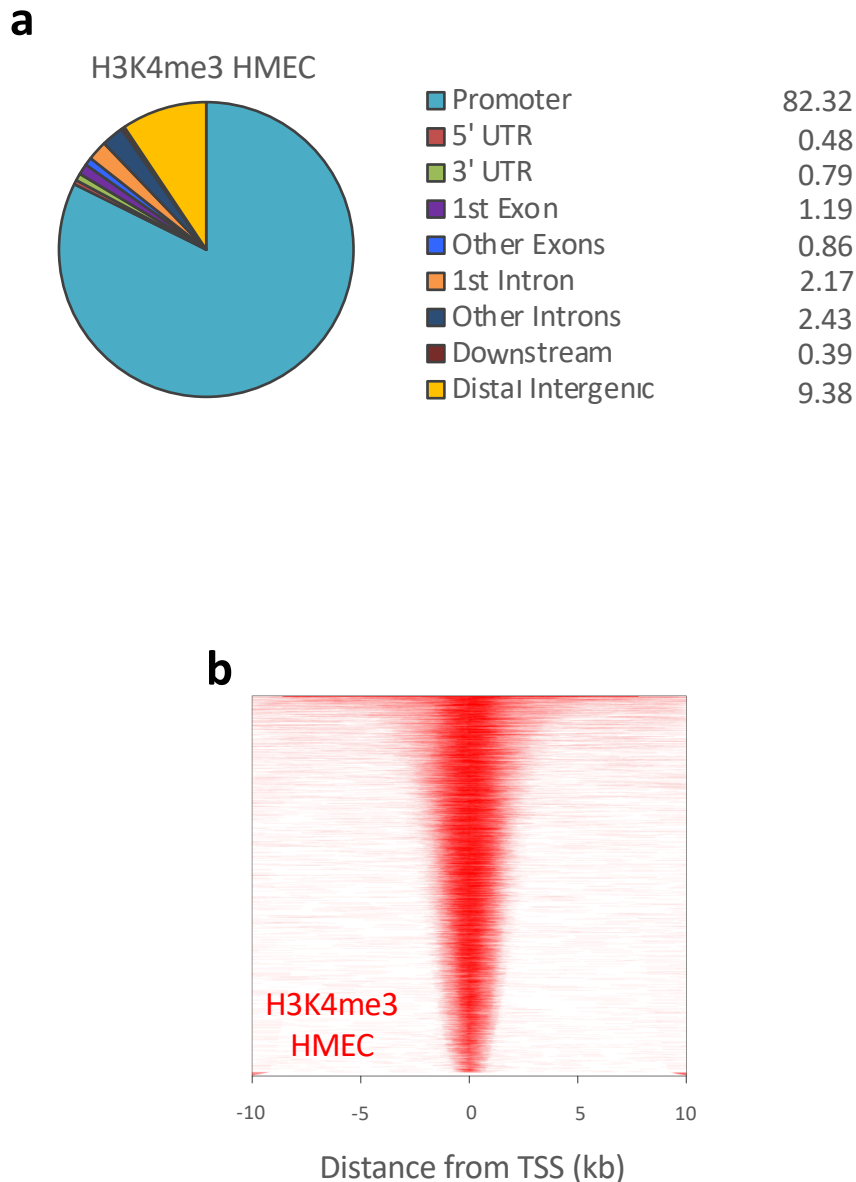


Figure S2. Analysis of H3K4me3 ChIP-Seq data from ENCODE project. H3K4me3 data from human mammary epidermal cells (HMEC) was taken from UCSC Genome Browser and analysed following the same pipeline used for standard PAT-ChIP and EPAT-ChIP (LRC) data sets. Pie charts depicting the distribution of peaks across genomic features with relative percentage values shown on the right (**a**). Heatmaps illustrating H3K4me3 peak densities from -10 Kb to +10 Kb relative to the TSS (**b**).