Figure S9. CAGE-enhancers expressed in ESCs and NESC. A) The graph shows the epigenetic signature of CAGE-enhancers in ESCs and NESC. Most CAGE-enhancers carried either a promoter (H3K4me3+/high/H3K4me1-/low) or an enhancer (H3K4me1+/high/H3K4me3-/low) signature. B and C) Analysis of putative TFBS enrichment within down- (B) and up- (C) regulated CAGE-enhancers, as determined by HOMER tool.