



Figure S8 GABP α and CREB1 binding to SNPs in genomic regions co-bound by CREB1 and GABP α , in different cell lines. (A-B) Enrichment of consensus and each 1-bp variation of ETS motif (CCGGAAGT) in GABP α ChIP-seq peaks (A) that do not overlap with CREB1 ChIP-seq peaks (GABP α - CREB1), and (B) that do overlap with CREB1 ChIP-seq peaks (GABP α + CREB1), in h1HESC cells. (C) Histogram of the ratio of motif enrichment +/- CREB1 to the consensus and 1-bp variations of the ETS motif (CCGGAAGT) in h1HESC cells. (D-L) same as in A-C, but for: (D-F) GM12878 cells, (G-I) HepG2 cells, and (J-L) K562 cells.