



Supplemental Figure S13. Enrichment of CRs localized in regulatory regions for RBP binding sites. The heatmaps show the fold enrichment (left) and the corresponding BH adjusted p -values (right) for the enrichment of CRs located in enhancers, 3' extensions, 5' extensions and upstream antisense of mRNA/lncRNA TSSs for RBP binding sites. The considered regulatory regions were defined by experimentally determined but unannotated transcript boundaries (CAGE TSSs and poly(A) sites).