

Fig S9

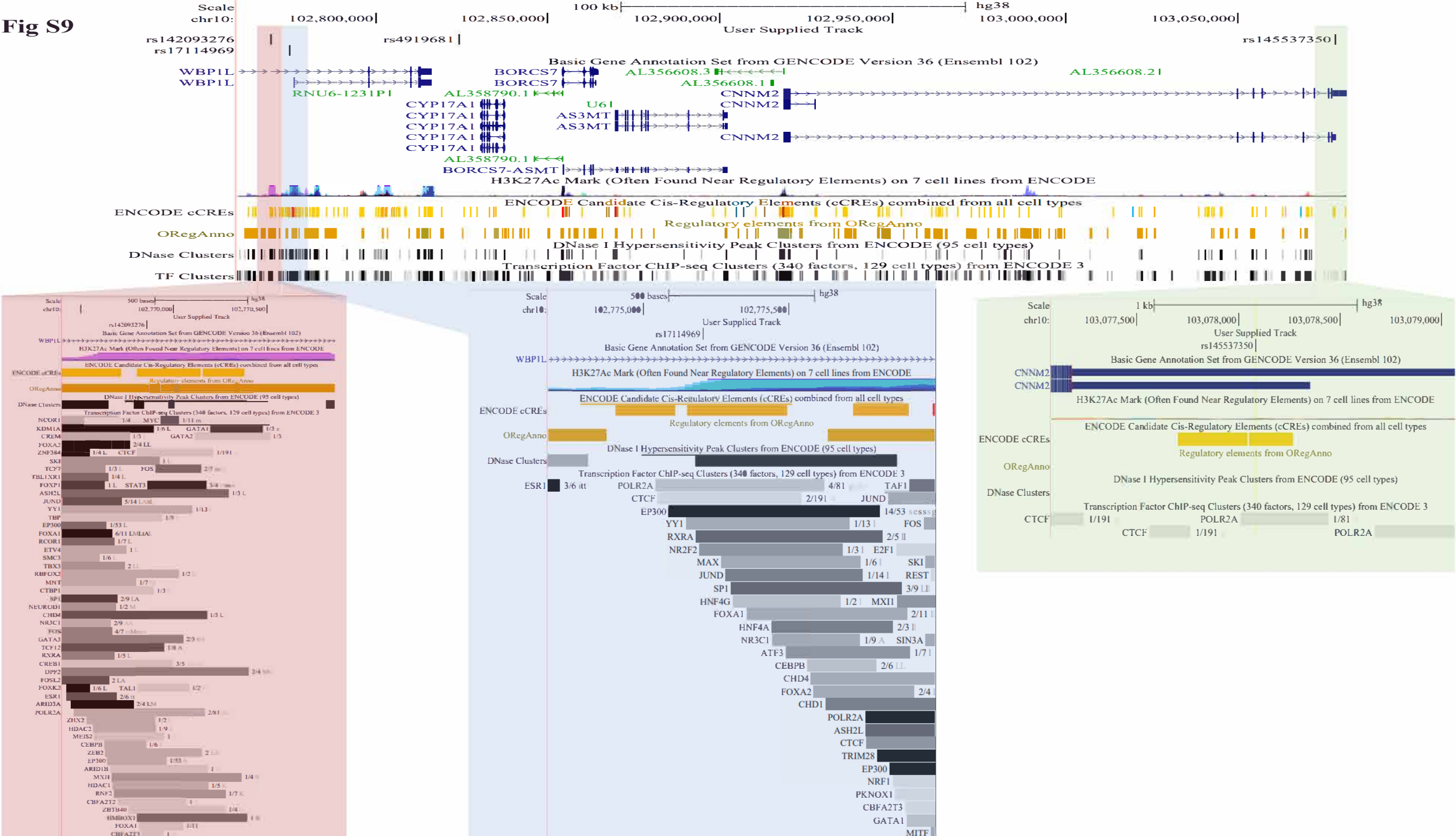


Fig S9. Genomic annotations of HEALS confidence set 1.

Results of in silico-functional analysis reveals overlap with genomic annotations for candidate causal SNPs from HEALS confidence set 1, corresponding to the primary association signal (lead SNP rs145537350). Multiple SNPs in the confidence set overlap genomic features including candidate cis-regulatory elements and transcription factor binding sites. Highlighted panels show details of candidate causal SNPs overlapping these features.