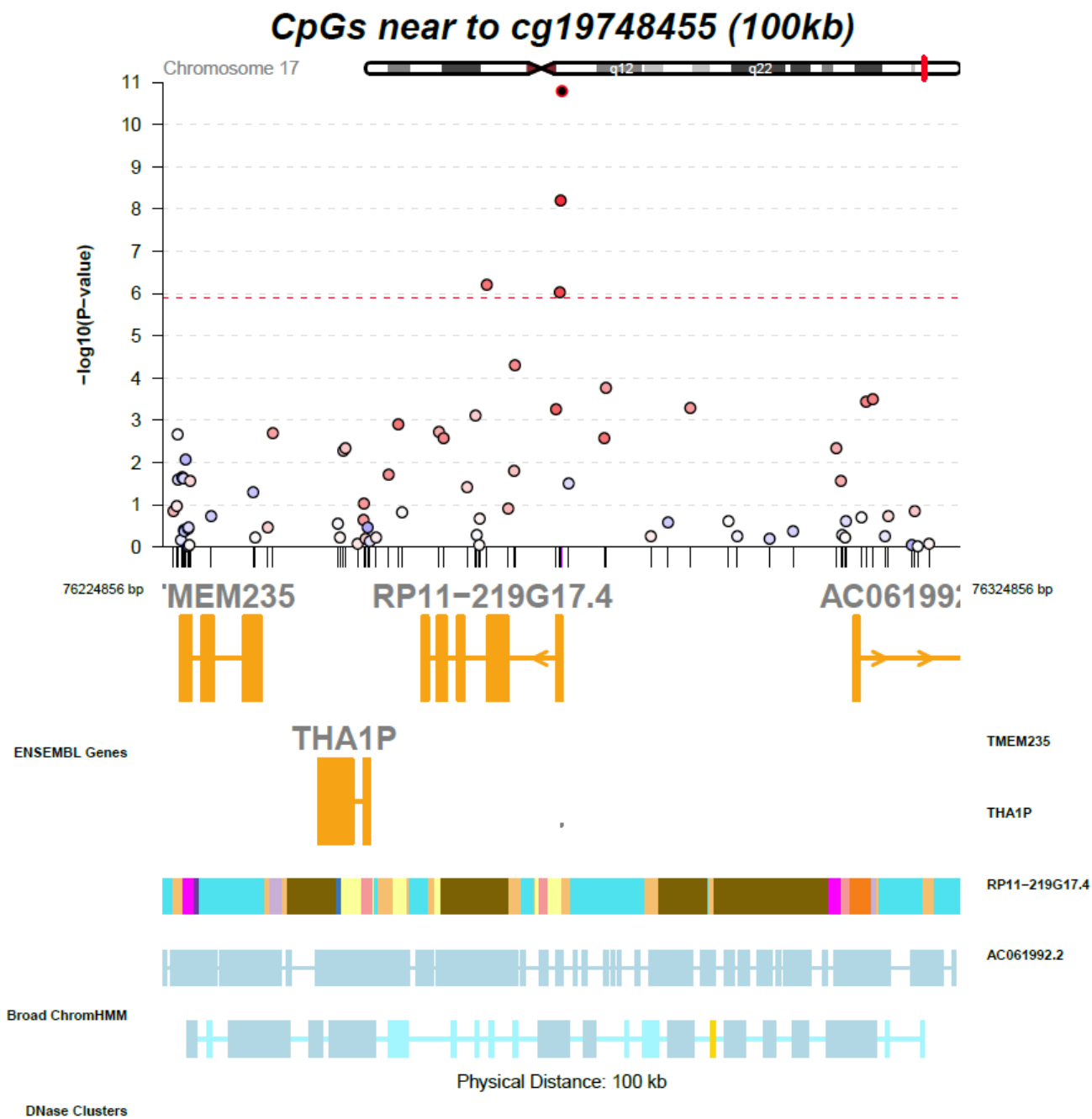
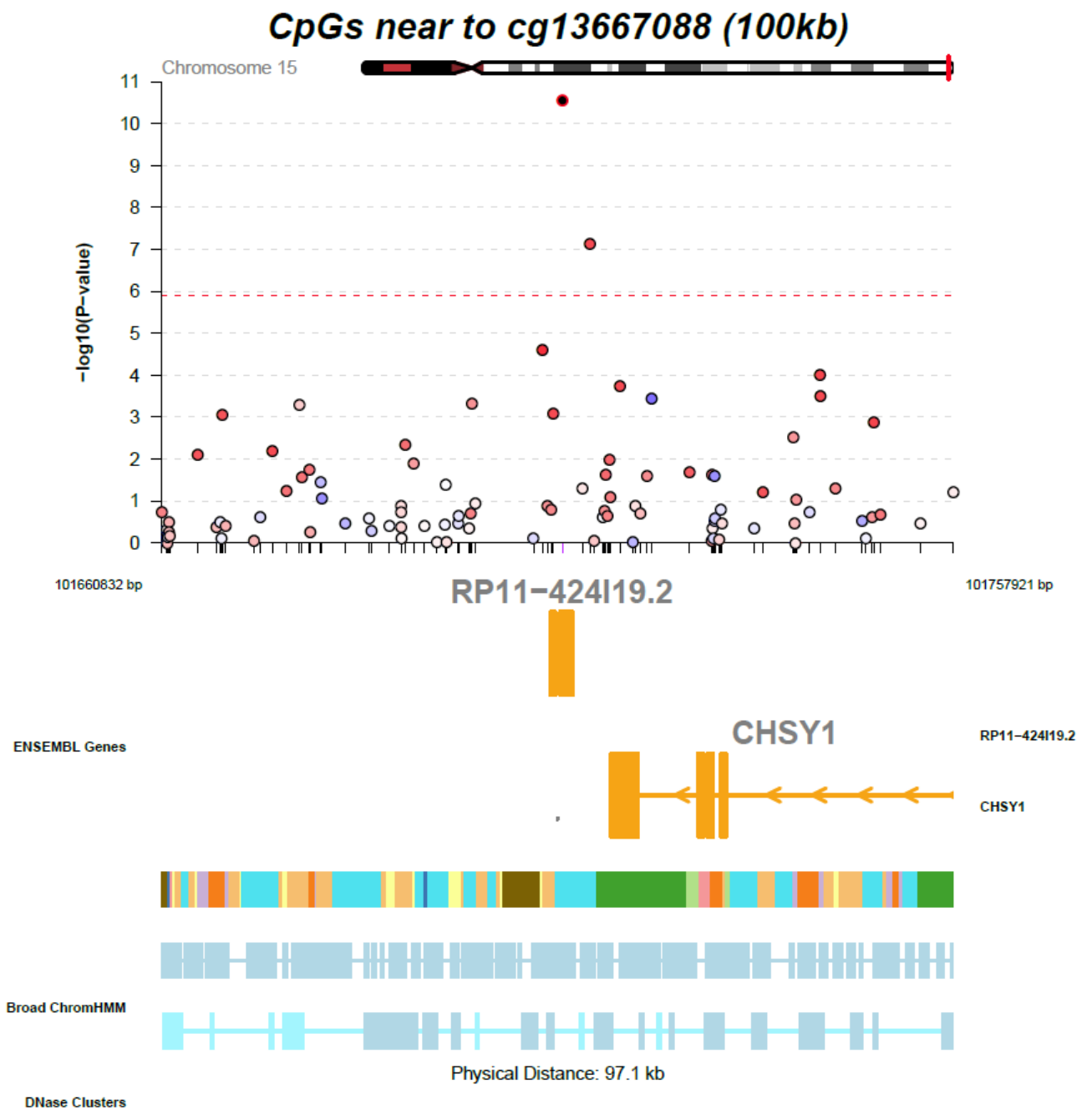


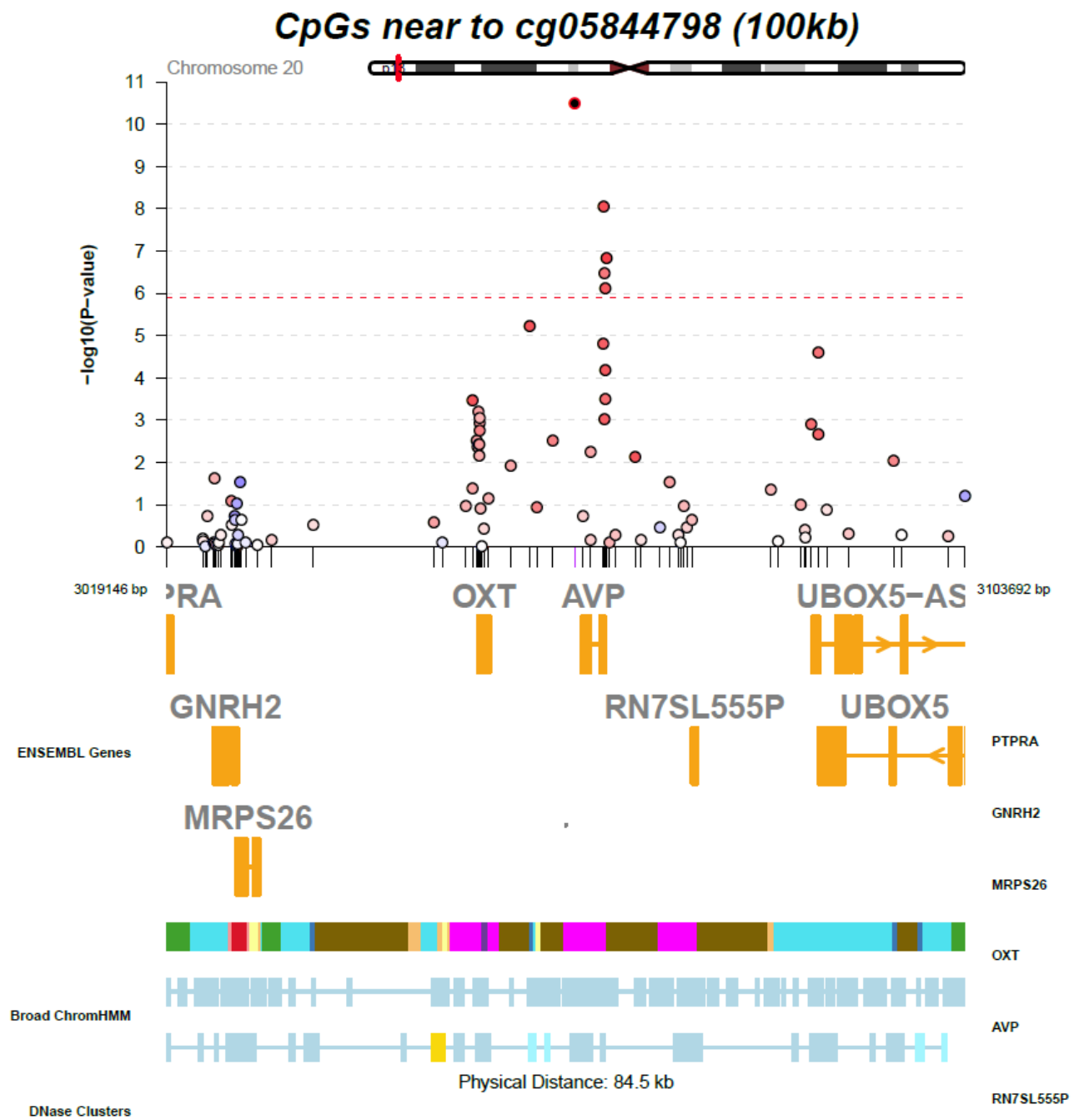
A) cg19748455 (*LOC100996291*)



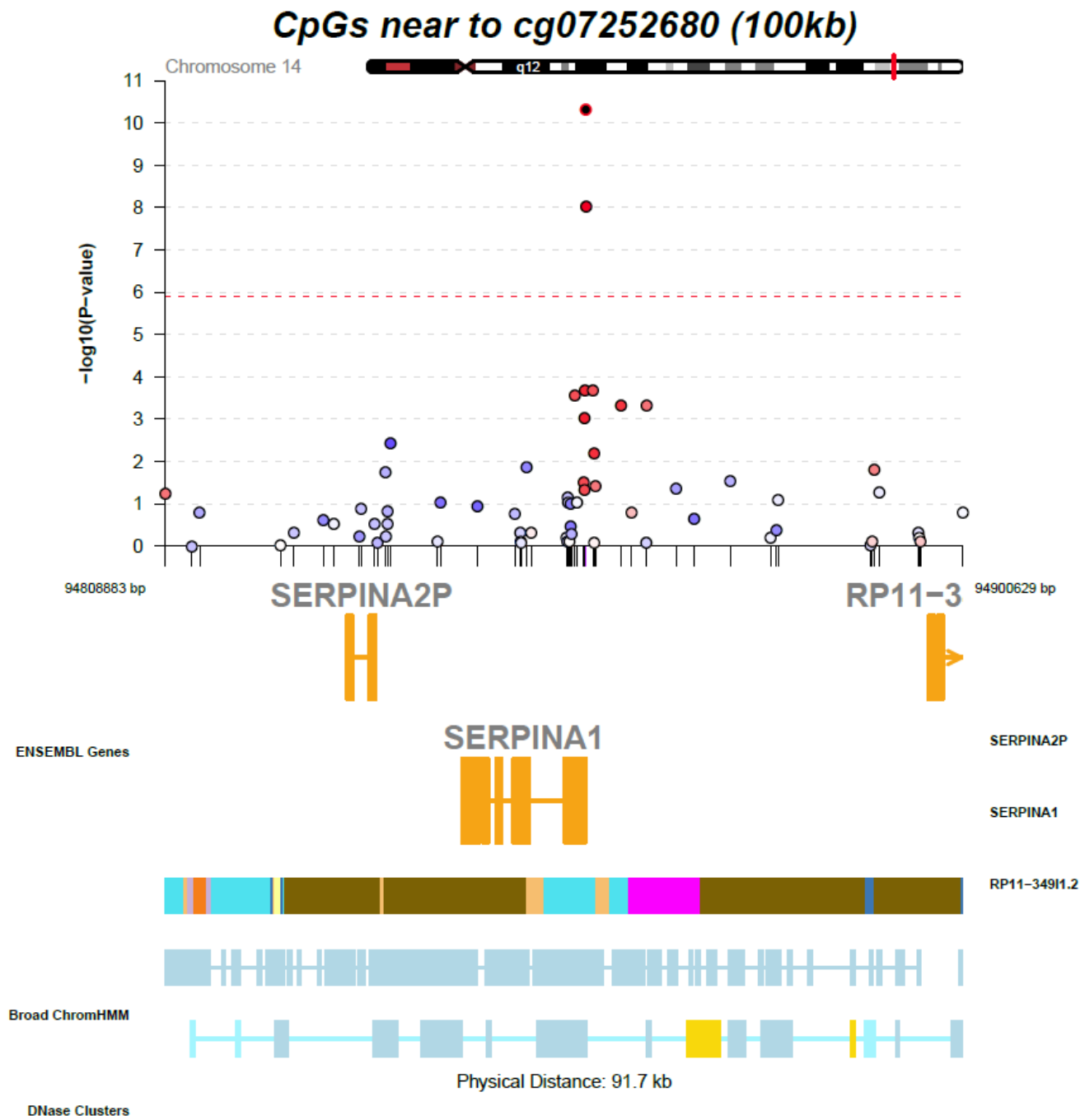
B) cg13667088



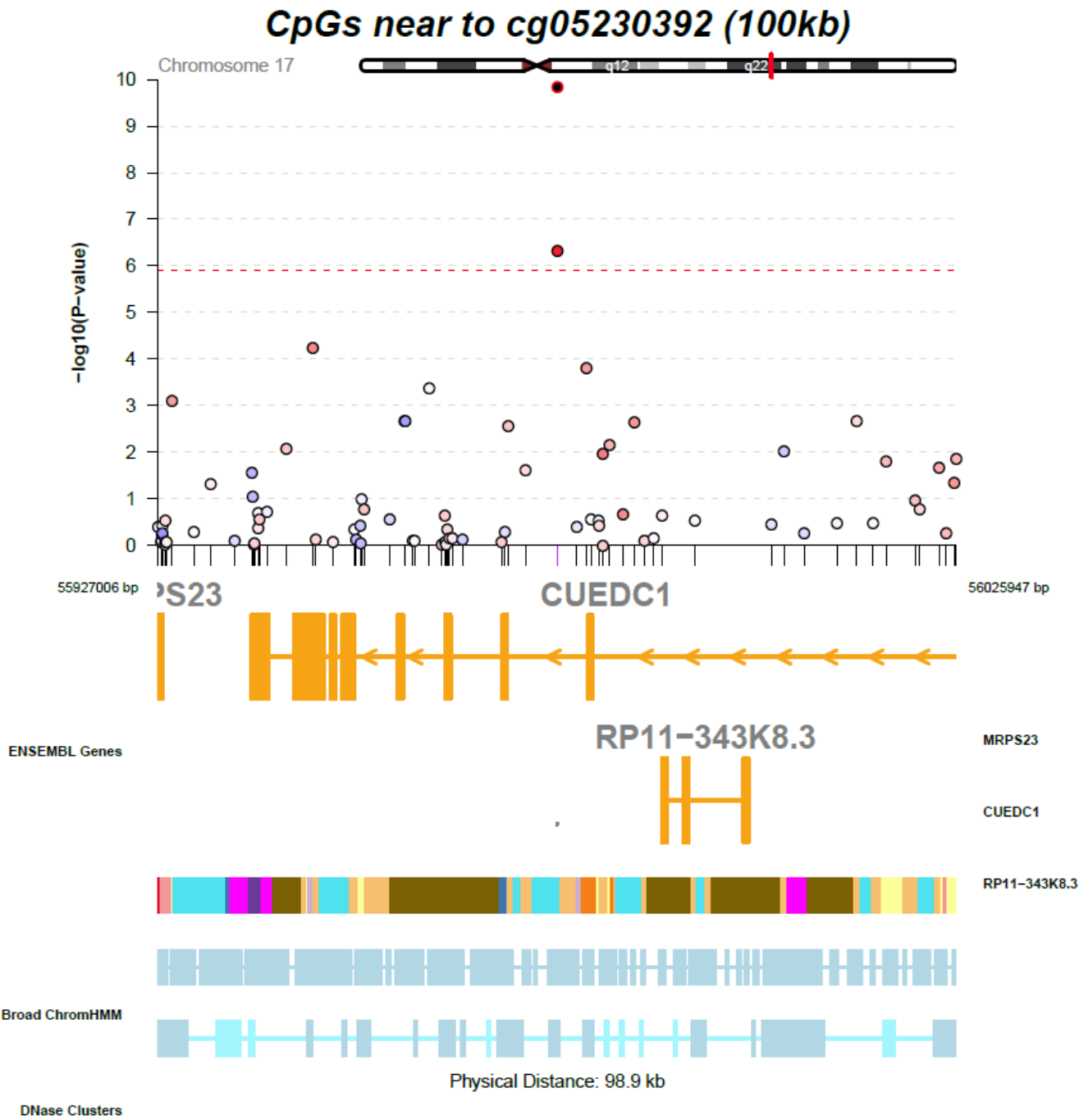
C) cg05844798



D) cg07252680 (*SERPINA1*)



E) cg05230392 (*CUEDC1*)



F) cg14047370 (*HECW1*)

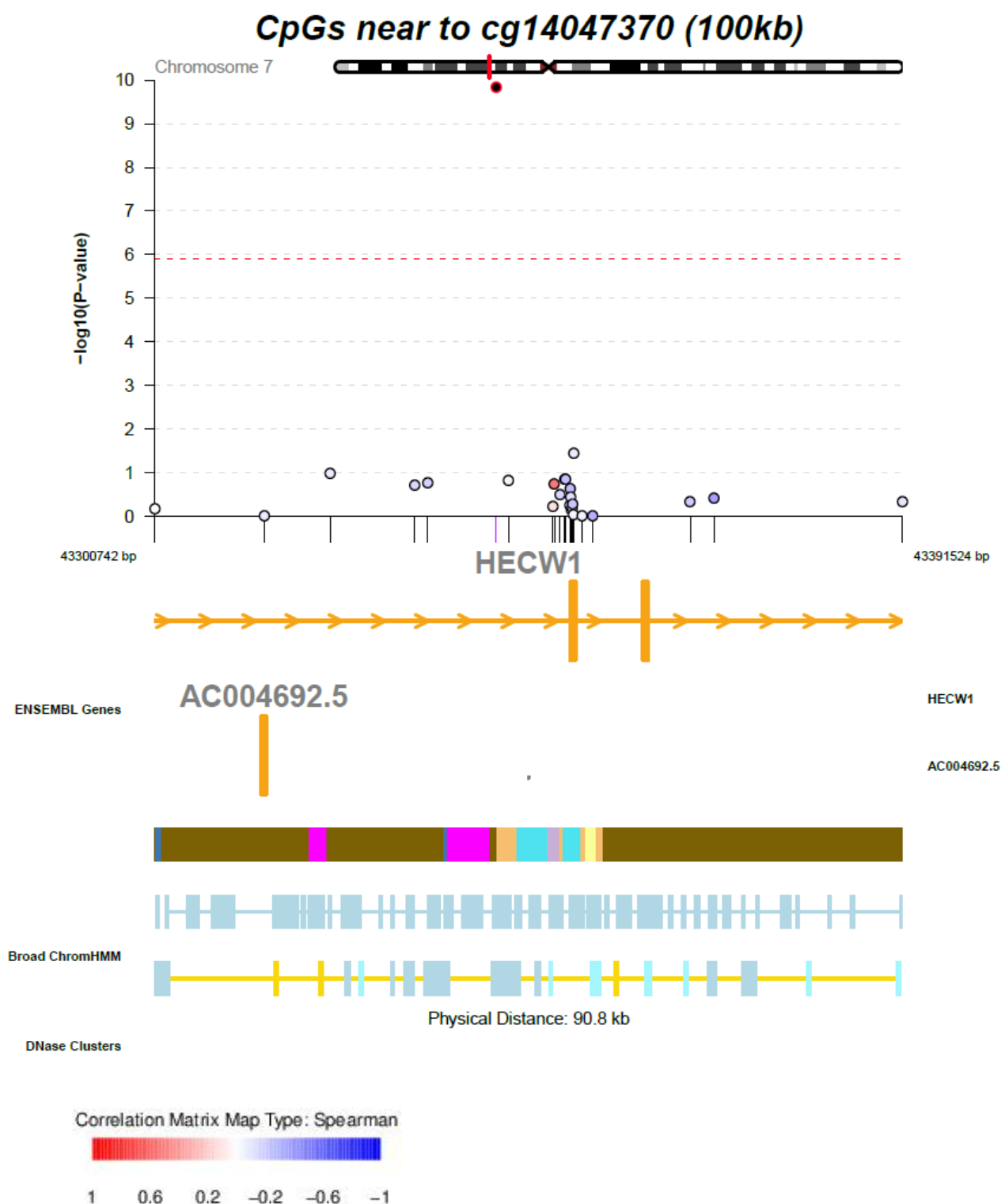


Figure S2. Regional plots for the genes containing top six FDR-significant EWAS CpG sites (circles): A) cg19748455 (*LOC100996291*), B) cg13667088 (*cg24172570*), C) cg05844798 (*cg08973675*), D) cg07252680 (*SERPINA1*), E) cg05230392 (*CUEDC1*), and F) cg14047370 (*HECW1*). $-\log_{10}(p\text{ values})$ from EWAS analysis, CpGs indicated by dots, color coded based on pairwise correlation with the index CpG.