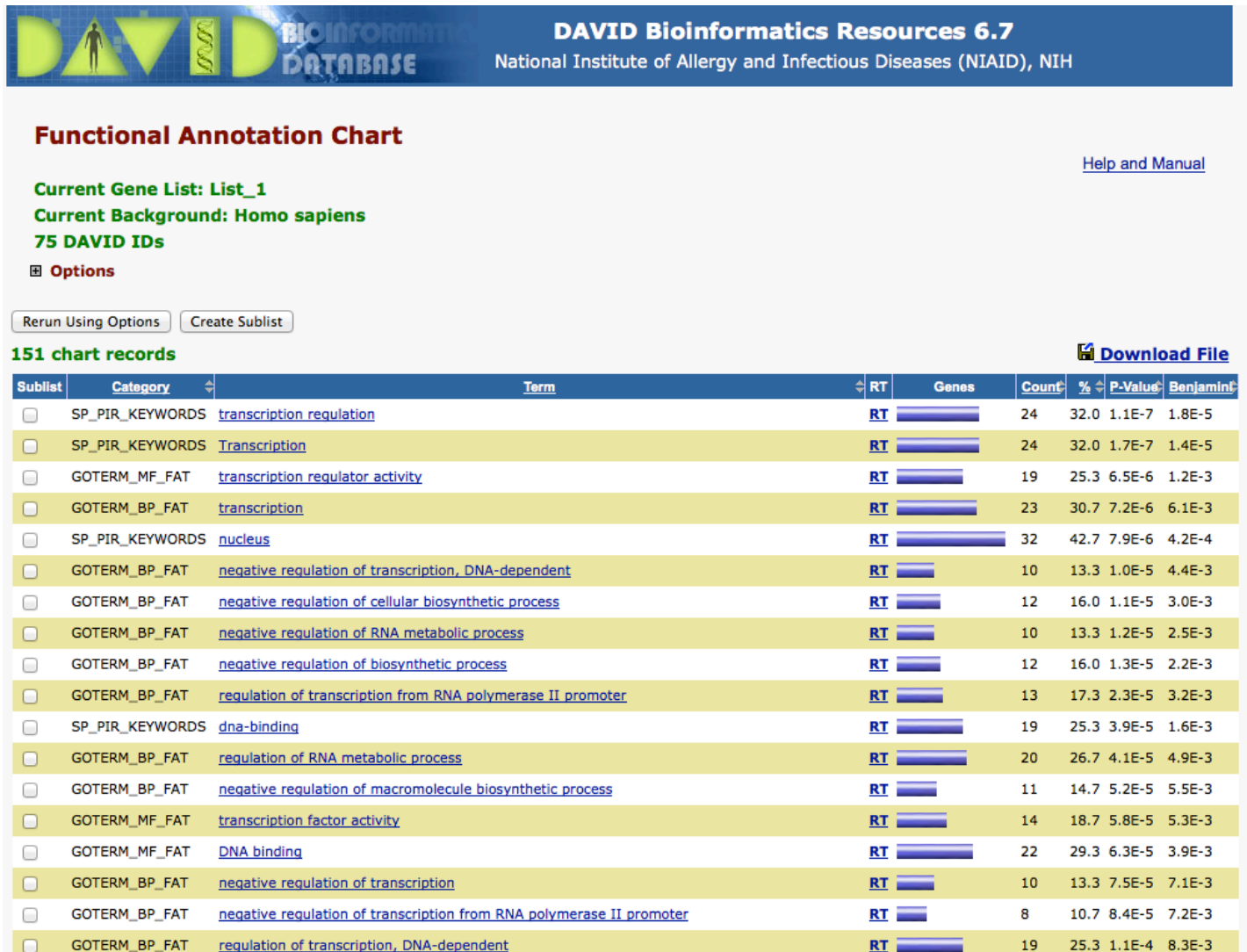
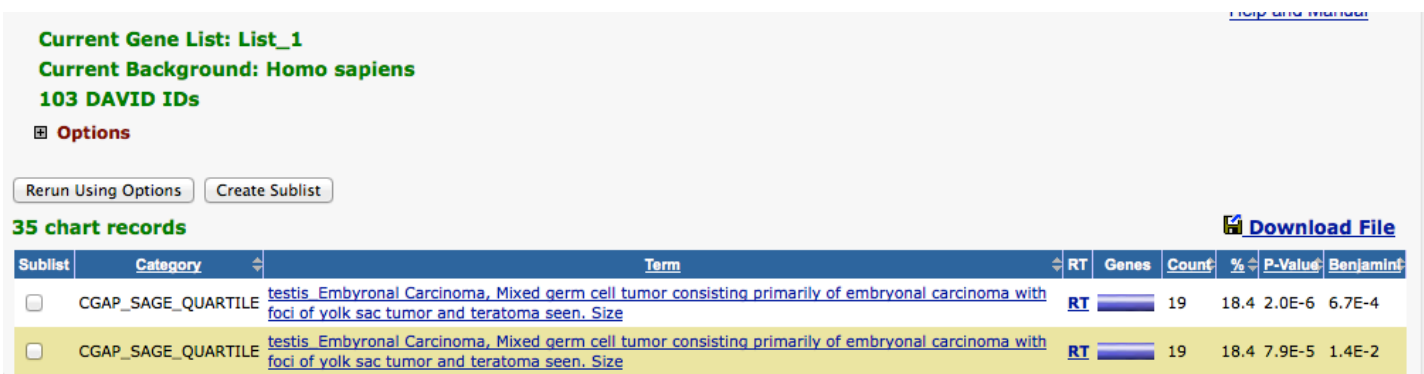


A EC/ESC/iPSC co-hypomethylation module



B EC-unique hypomethylation module



Supplemental Fig. S10. **A:** Functional annotation of the EC-ESC/iPSC co-hypomethylation module of 141 CpG targets mapped to 76 genes revealed significant association with homeobox and cell fate commitment genes (Table S2). **B:** Targets uniquely erased in EC relative to pluripotent references (161 targets mapped to 56 genes) are significantly enriched for genes with upregulated transcription in EC (Table S2).