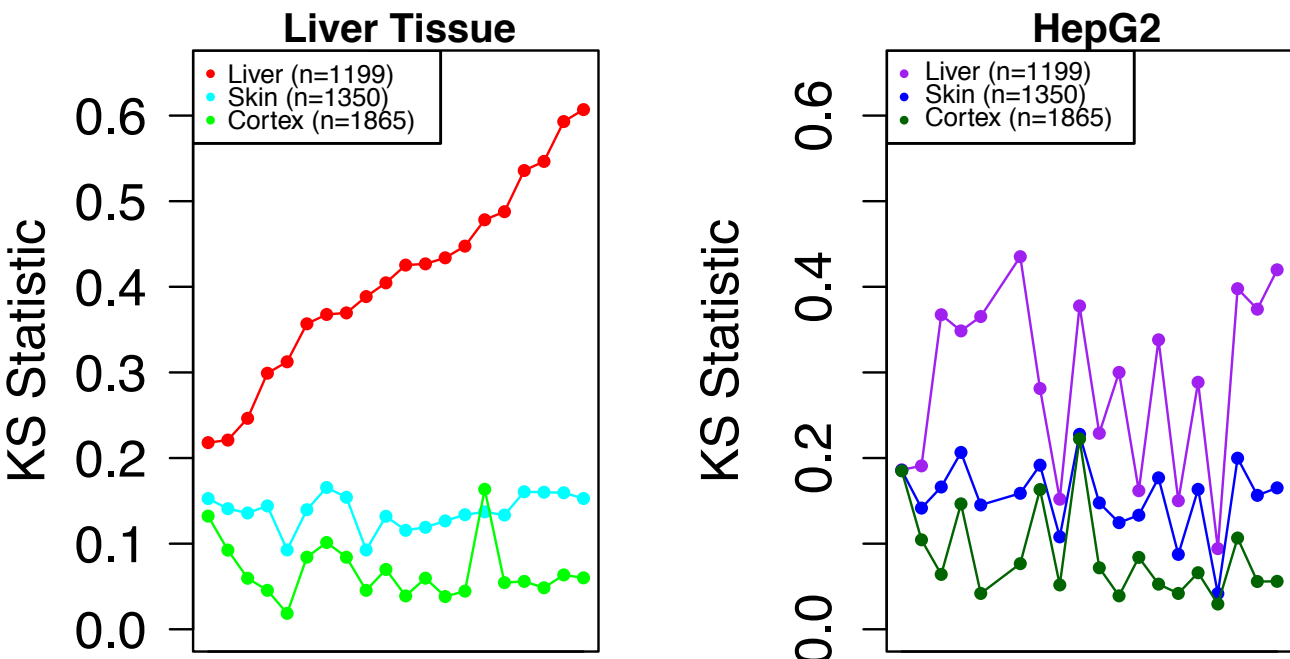
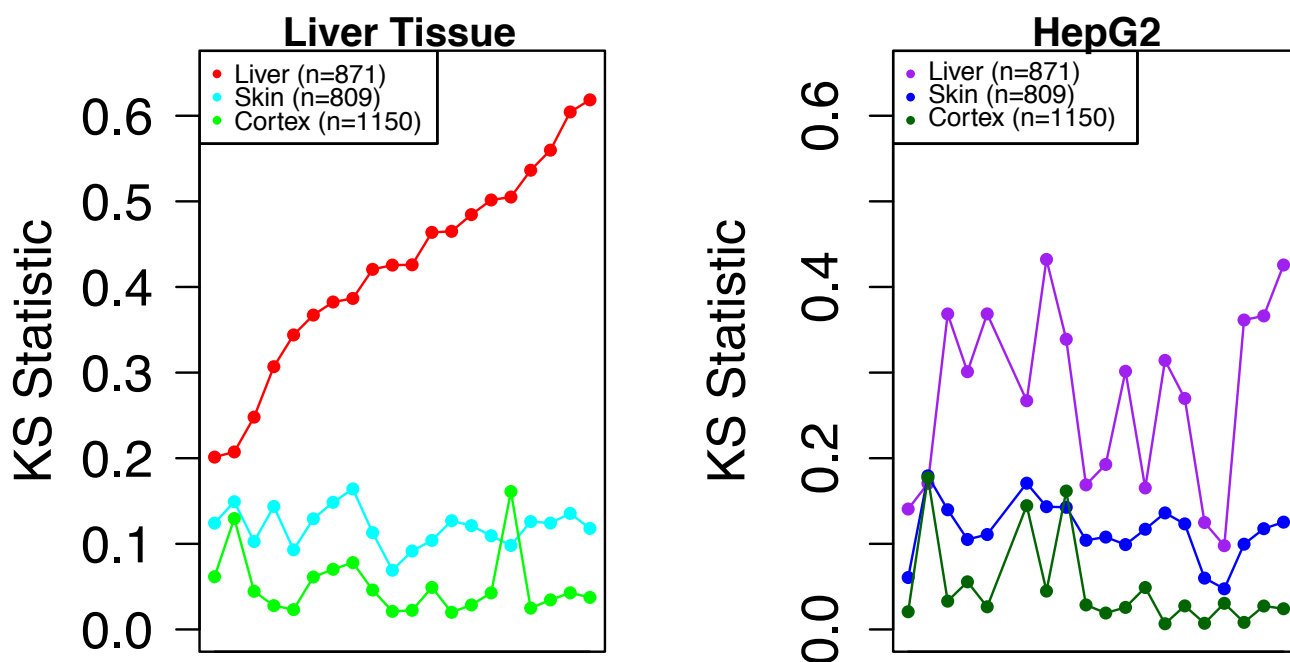


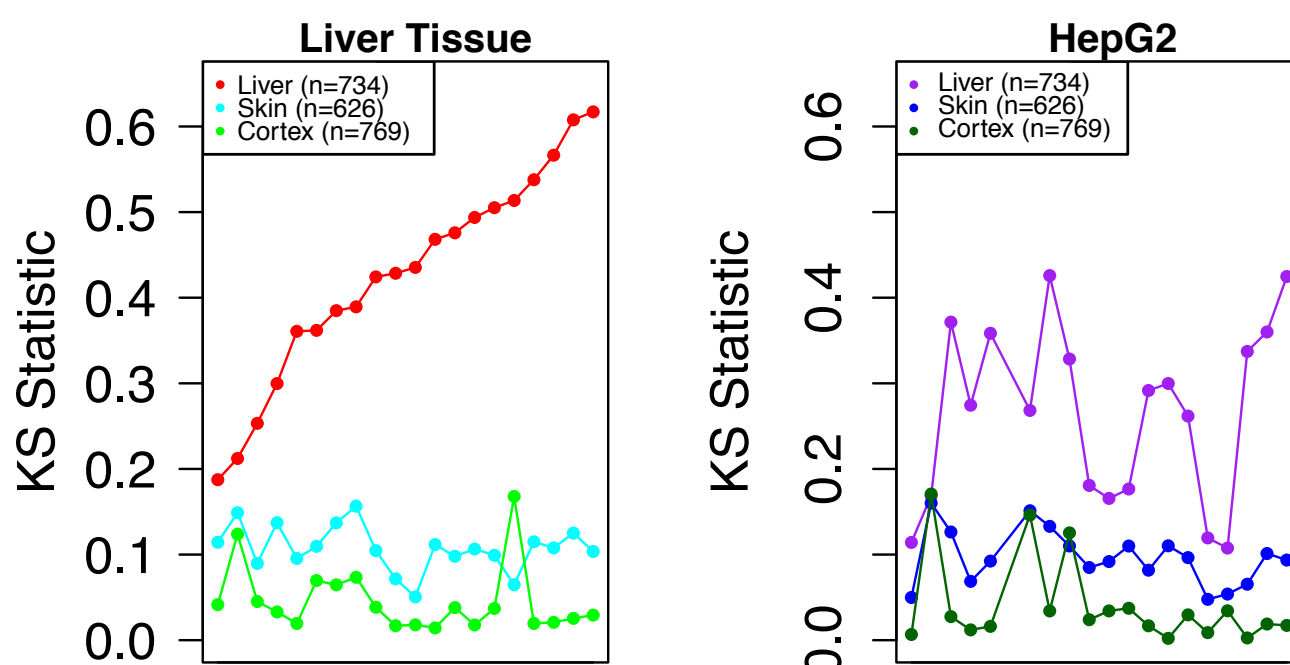
2-Fold Enrichment Threshold



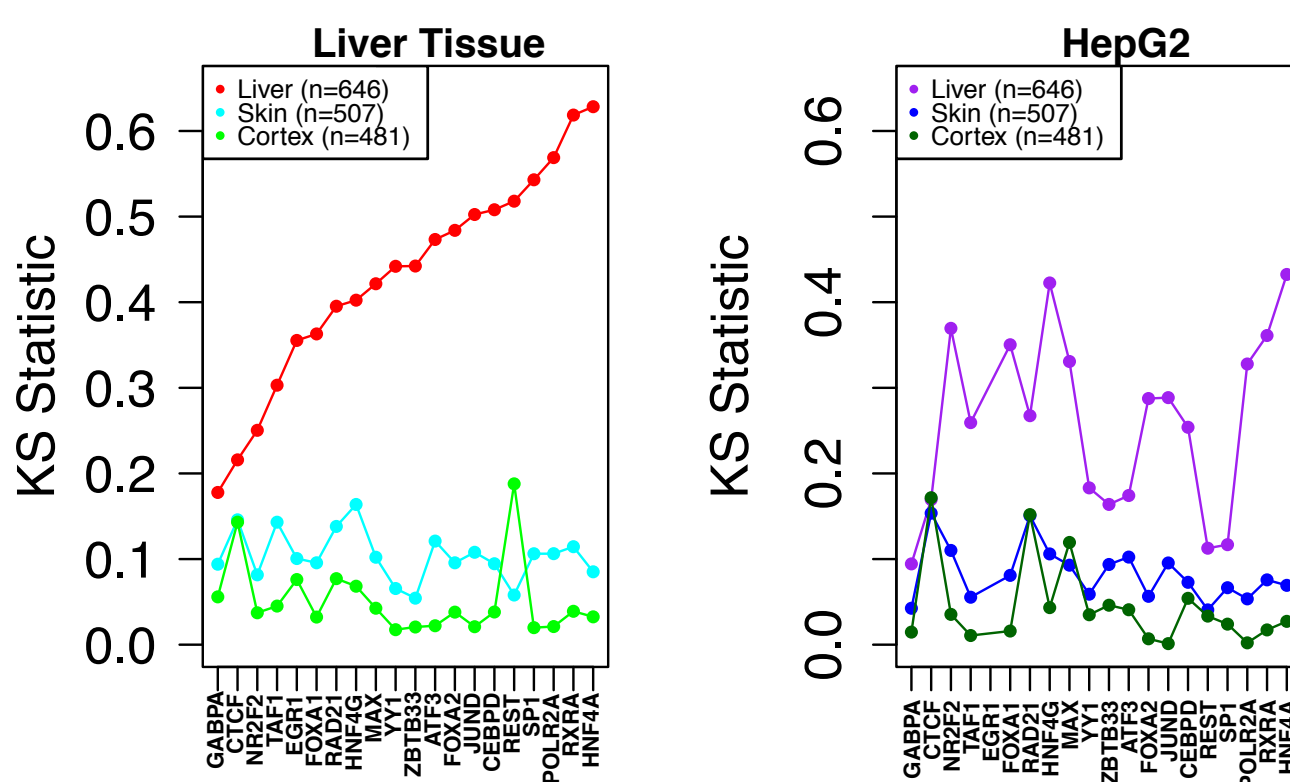
3-Fold Enrichment Threshold



4-Fold Enrichment Threshold



5-Fold Enrichment Threshold



Supplemental_Fig_S12. Enrichment for proximal binding to liver specific expression analysis, as shown in Figure 3C-D, over a range of fold enrichment threshold cutoffs. From top to bottom, tissue specific transcripts were defined as those with a mean RPKM of 2 in a given tissue of interest and 2, 3, 4, and 5-fold increased mean expression in the tissue of interest relative to the mean expression across all other tissues. Dots represent KS-statistic of enrichment for proximal binding of each DAP to liver (red/purple), skin (blue) and cortex (green) -specific genes in adult liver tissue (left) and HepG2 cell line (right).