



**S6 Fig. Gene expression of gestational duration- and SPTB-annotated genes across 30 tissue types in GTEx v8.** The histograms show MAGMA gene set enrichment analysis for gestational duration (above) and SPTB (below). The heatmap features gene expression depicted as averaged expression value per tissue type, with hierarchal clustering for both genes and tissues. Across genes and tissues, the cells filled in red represent higher expression compared to the cells filled in blue.