



**Supplemental Figure S4.** (A) The proportion of tissue-specific (orange) splicing events and exons and those testable through eQTL (purple). Splicing *cis*-eQTLs in autoimmune disease genes in CD4+ (pink), CD8+ (blue), and CD19+ lymphocytes (green) are summarized as (B) SNPs shared between tissues, (C) SNPs shared between tissues where the associated feature is detected in all three tissues.