



Figure S1 Tissue-specific RE expression patterns

Hierarchally-clustered heatmap of RE-specific probes significantly regulated between B6 tissues for three independent experiments using the Mouse Genome 430 v2 microarray platform ($p < 0.001$ by ANOVA comparing tissues and eliminating experiment). Data are obtained from E-GEOD-1986, -9954, and -10246, which are identified with numbers 1, 2 and 3, respectively.