

Description of Additional Supplementary Files

Supplementary Data 1:

Differentially expressed genes in LTbR- deficient HSCs relative to LTbR proficient HSCs
FACS-purified *Ltbr*^{-/-} and Ly5.1 LT/ST-HSCS from chimeras six weeks after primary transplantation (n=6, pooled to 3 biological replicates) were used for RNA-seq analysis. Supplementary data 1 includes all 227 differentially expressed genes.