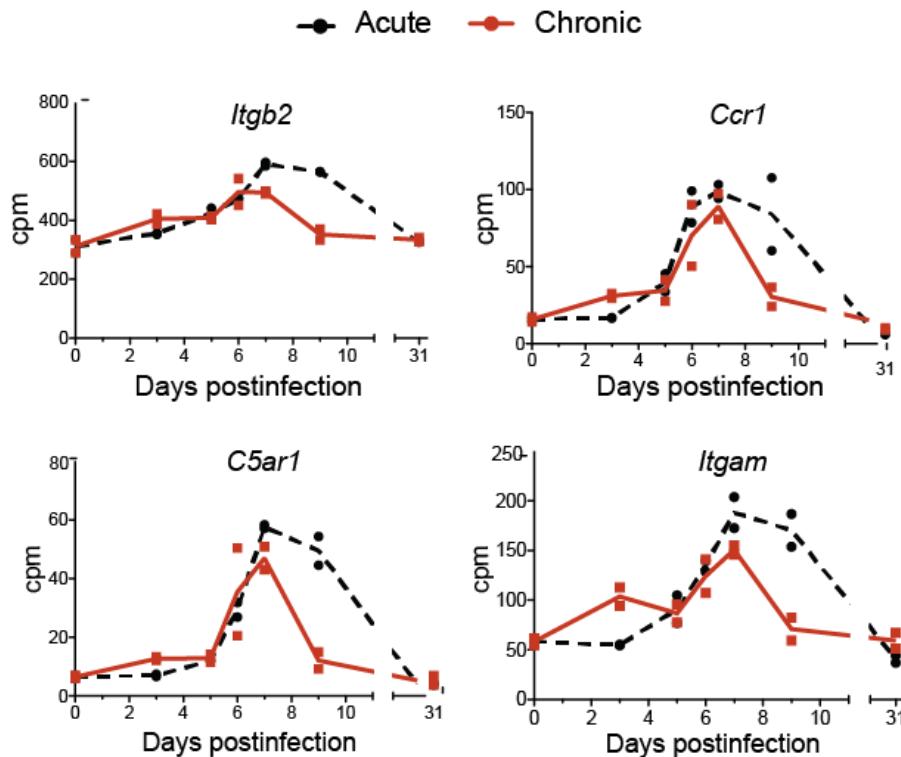


Supplemental Fig. S6. Validation of RNAseq-derived gene expression by qPCR.

(A) Normalized gene expression (cpm) kinetics obtained from RNAseq analysis of spleens from acute or chronic infected mice. (B) Quantitative PCR of the leukocyte chemotaxis related genes *Itgb2*, *Ccr1*, *C5ar1* and *Itgam* was performed at the indicated days postinfection from spleens of acute or chronic infected mice. Relative gene expression levels were normalized to *Gapdh*. Data are shown as mean \pm SEM from 5 to 6 mice. Significant differences were determined by an unpaired two-tailed *t*-test. ns, non significant; * $p \leq 0.05$; ** $p \leq 0.01$.

A**B**