

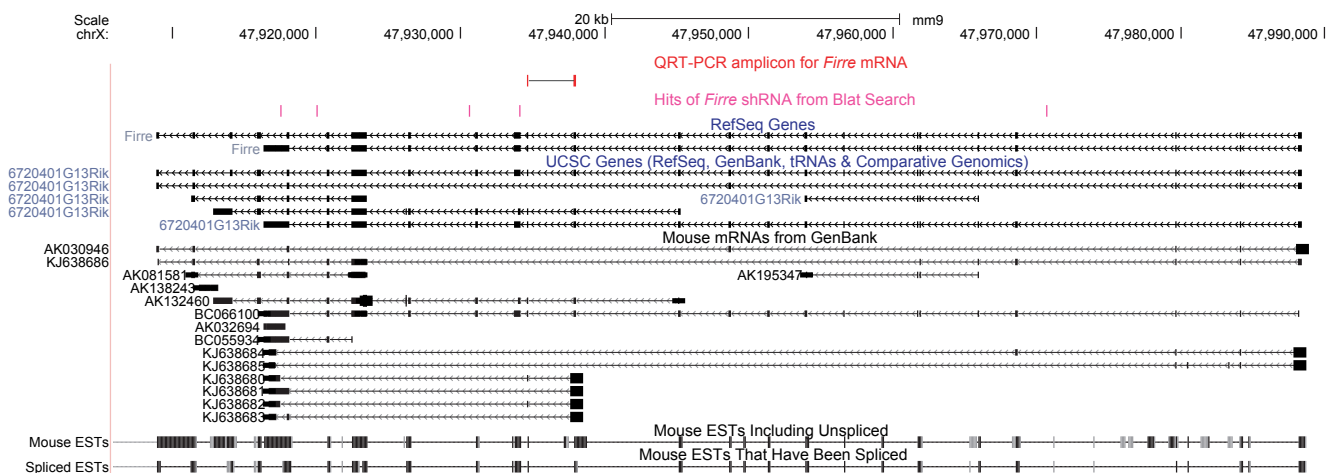


Figure S1. Location of qRT-PCR amplicon and of shRNA hits in the *Firre* gene viewed in the UCSC genome browser. The *Firre* qRT-PCR amplicon spans an intron and targets most transcripts. The shRNAs hit five regions in *Firre*, two in exons and three in introns. There is no other hit in the mouse genome from Blat search.