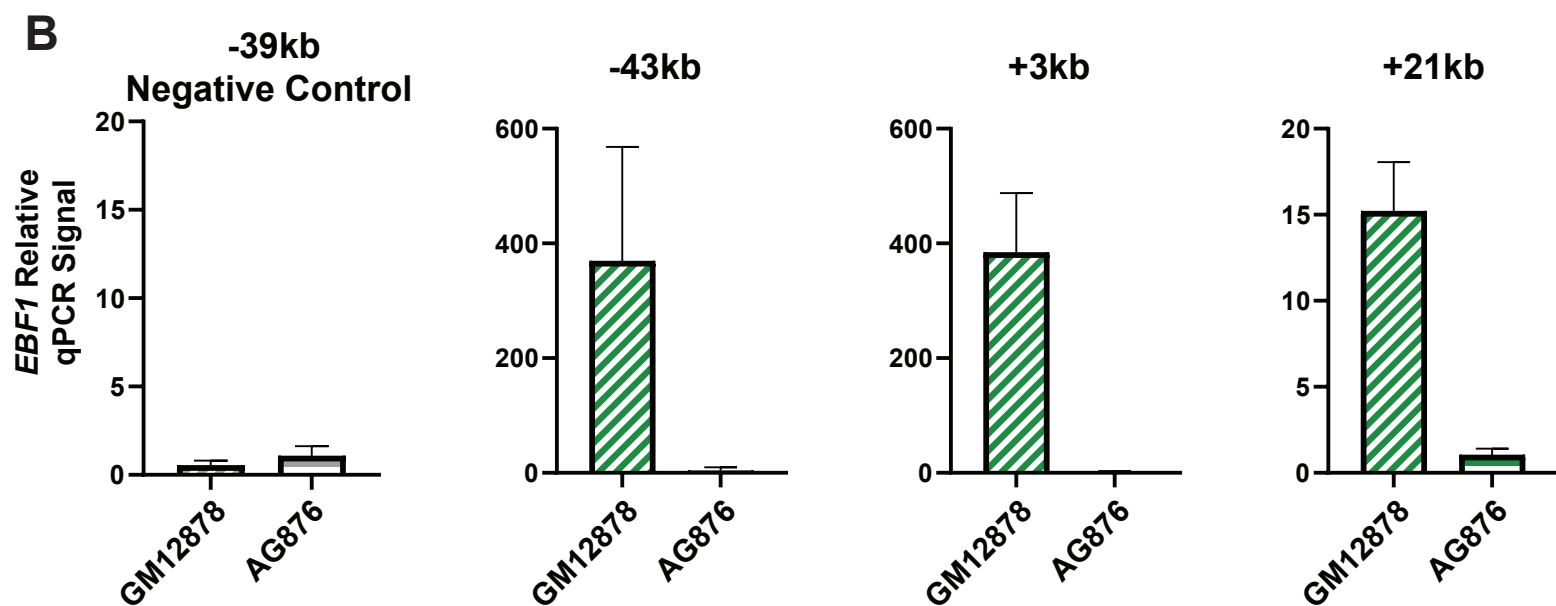
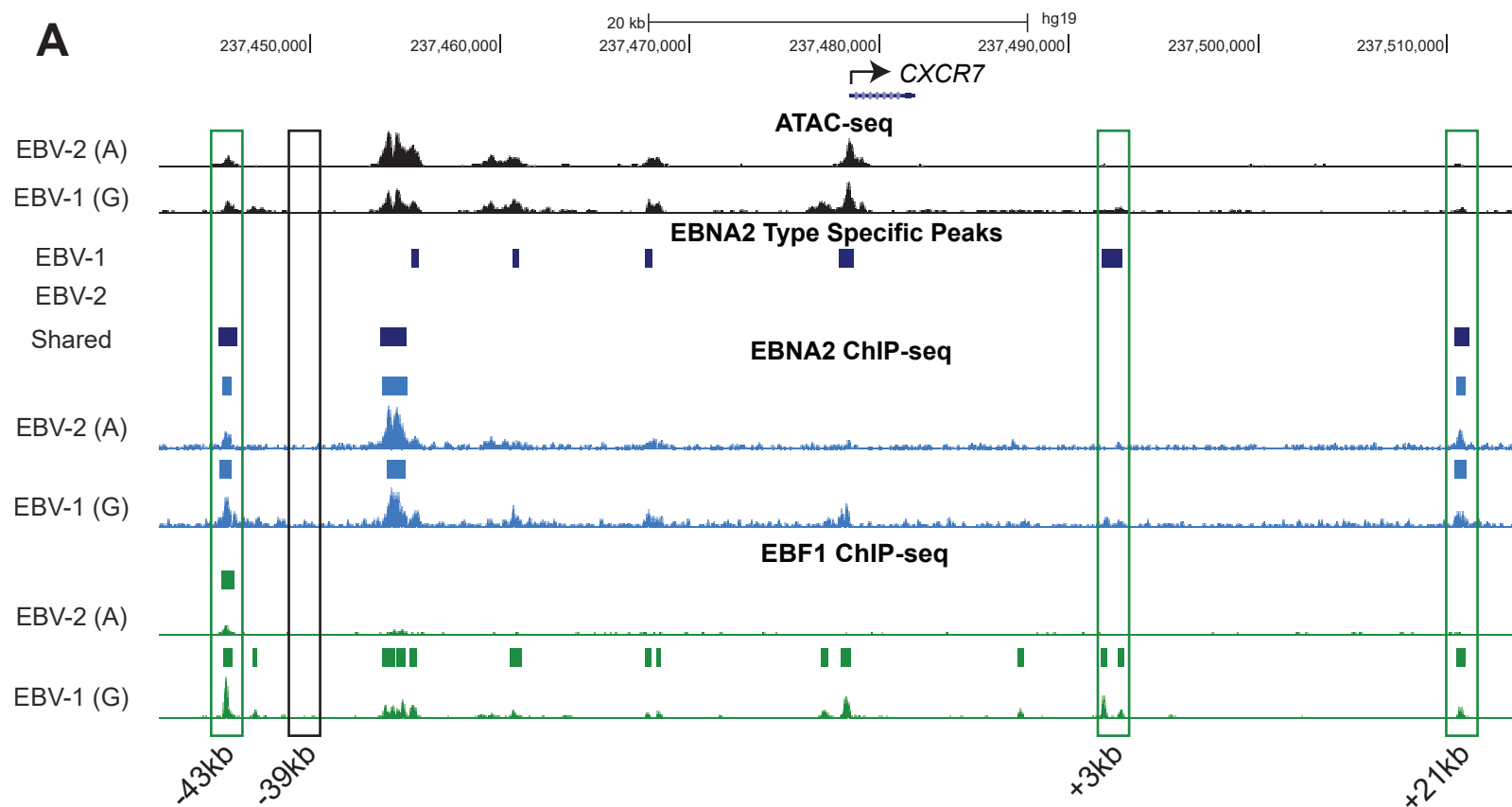




Additional File 11: Supplemental Figure 11. EBF1 ChIP-qPCR analysis of EBNA2 ChIP-seq peaks in the *CXCR7* locus. A) Genome Browser track showing chromatin accessibility and EBNA2 & EBF1 ChIP-seq signal at the *CXCR7* locus. Range: ATAC-seq (0 to 5); EBNA2 ChIP-seq (0 to 1.5); and EBF1 ChIP-seq (0 to 4). A = AG876; G = GM12878 B) Chromatin immunoprecipitation of EBF1 at the *CXCR7* locus (chr2:237,441,986-237,513,677) was normalized to the cycle threshold of the AG876 immunoprecipitation. The positions of each qPCR plot are relative to the transcriptional start site of *CXCR7*. -39kb was selected as a negative control. The mean of three technical replicates is plotted; error bars are SD.