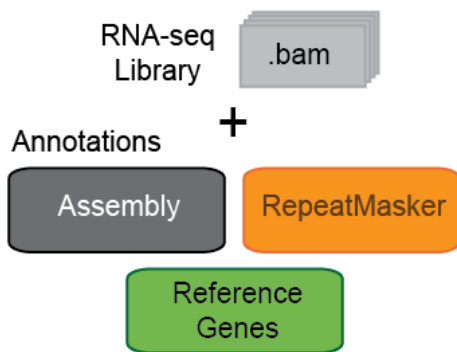
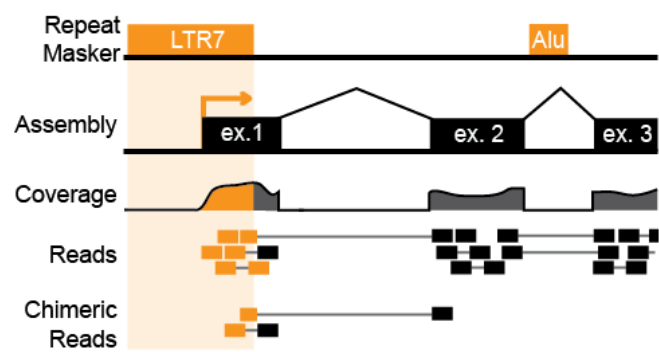


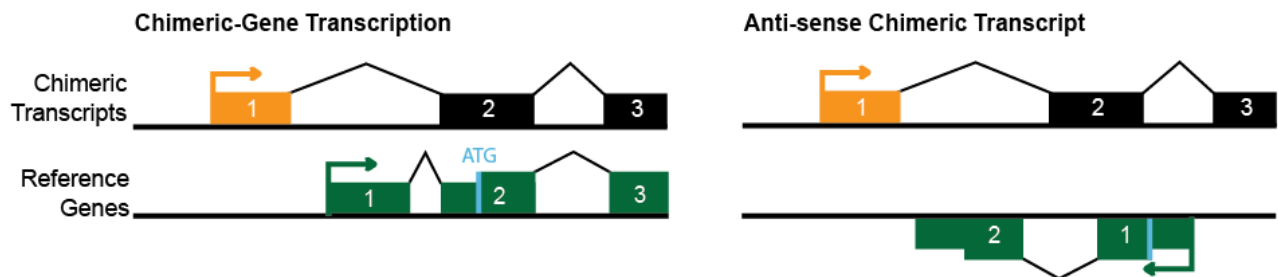
A) i. Input



ii. Detect & Classify TE-Derived Transcripts



iii. Intersect TE-Derived Transcripts to Reference Genes



B)

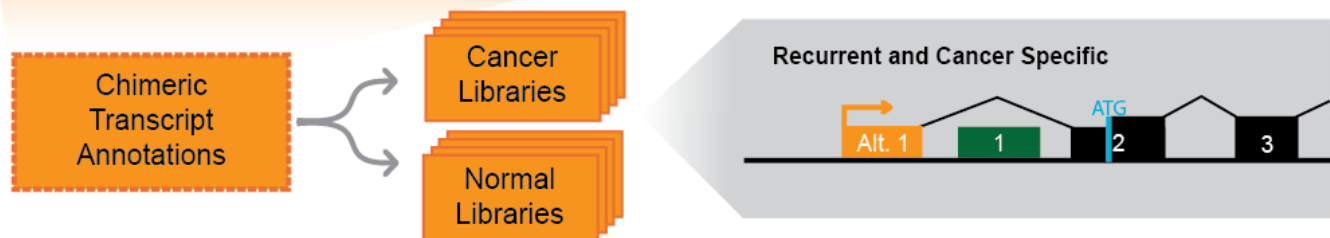


Figure 1: Identification of Chimeric Transcripts Workflow

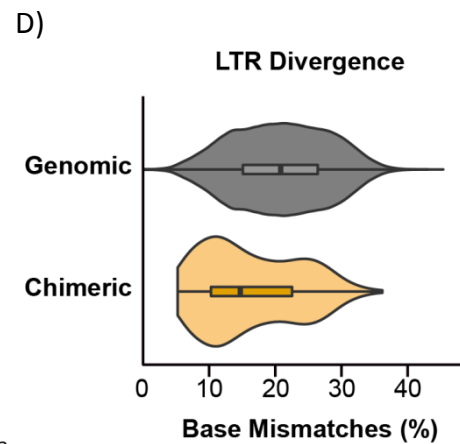
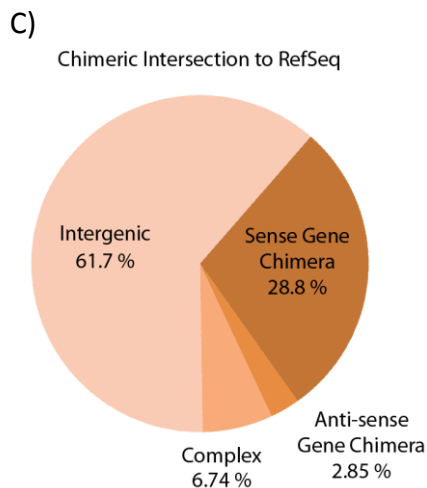
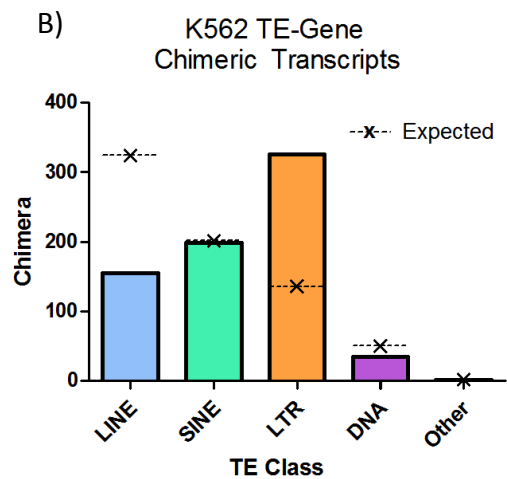
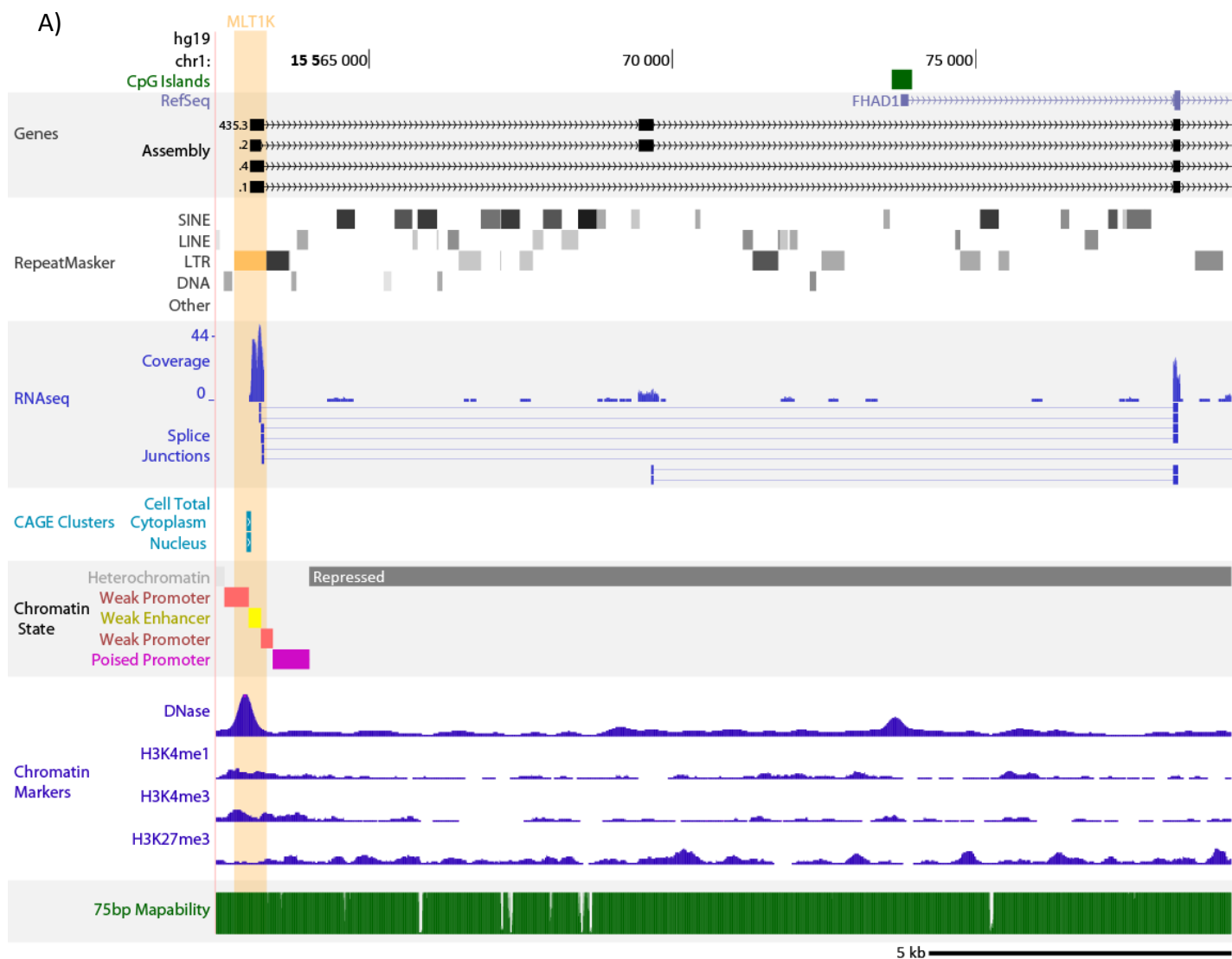


Figure 2: Chimeric Transcription in the K562 ENCODE training data

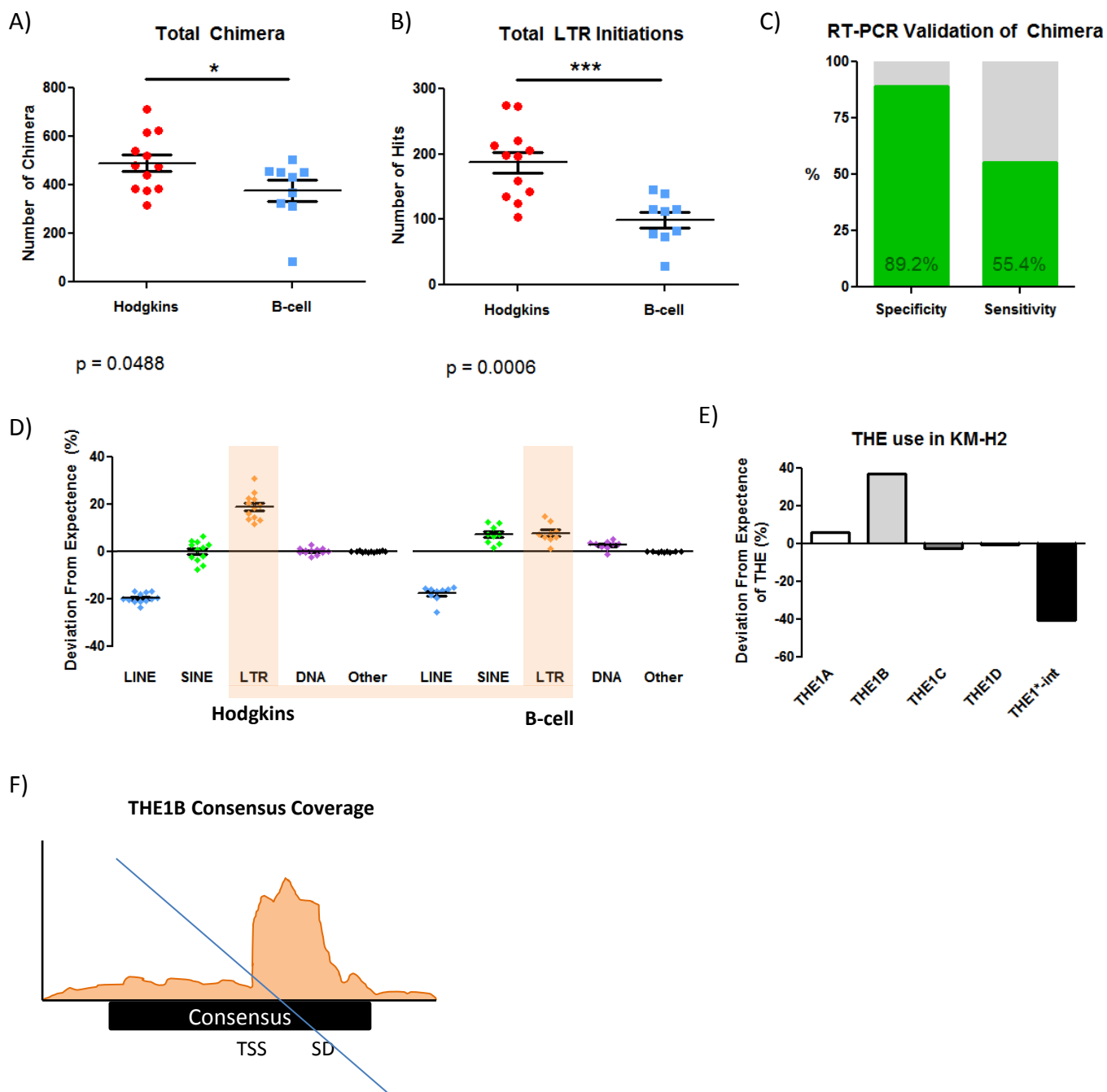


Figure 3: Chimeric Transcription in the Hodgkin's Lymphoma and B-cell Transcriptome

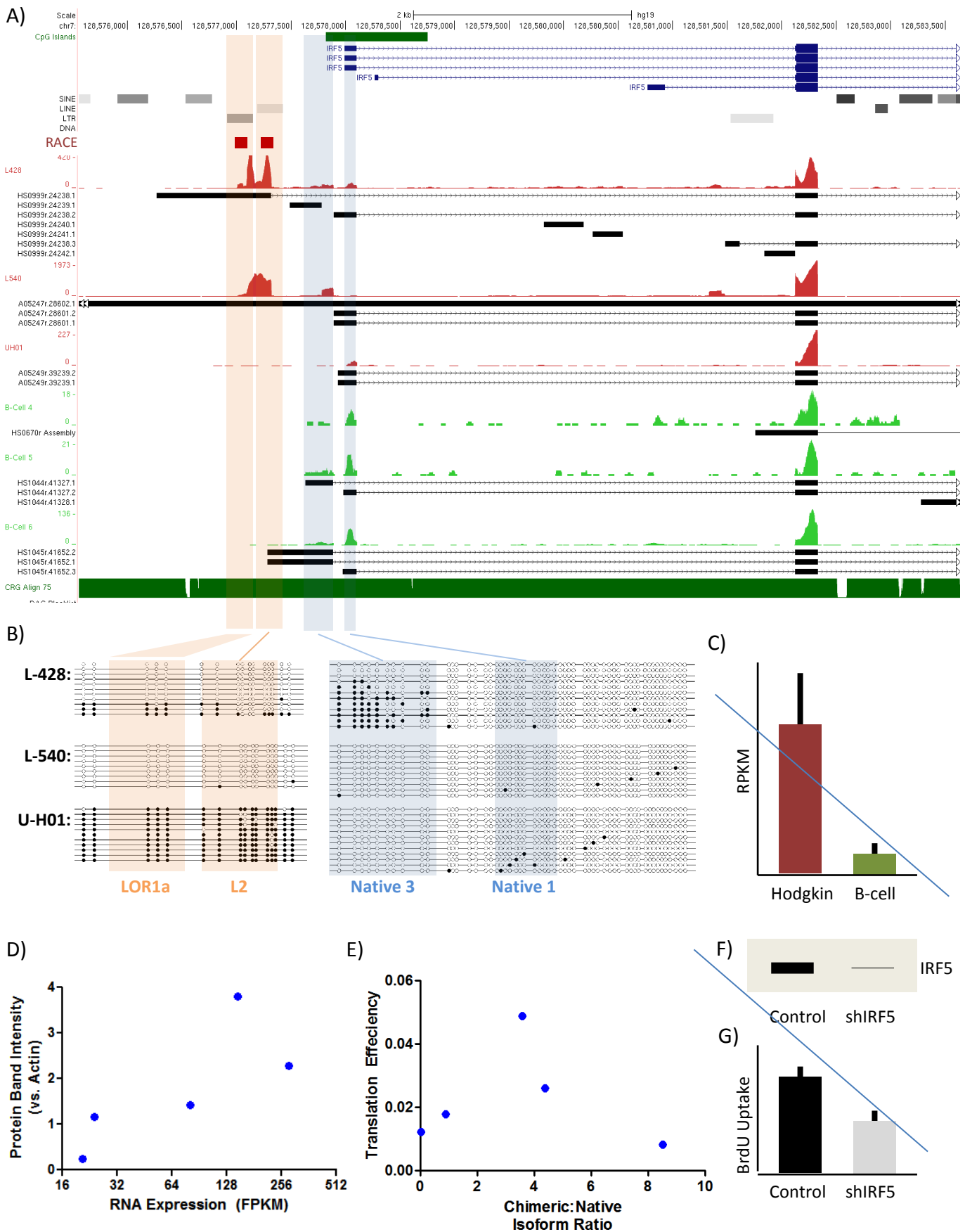


Figure 4: Chimeric Transcription of *IRF5* in Hodgkin's Lymphoma

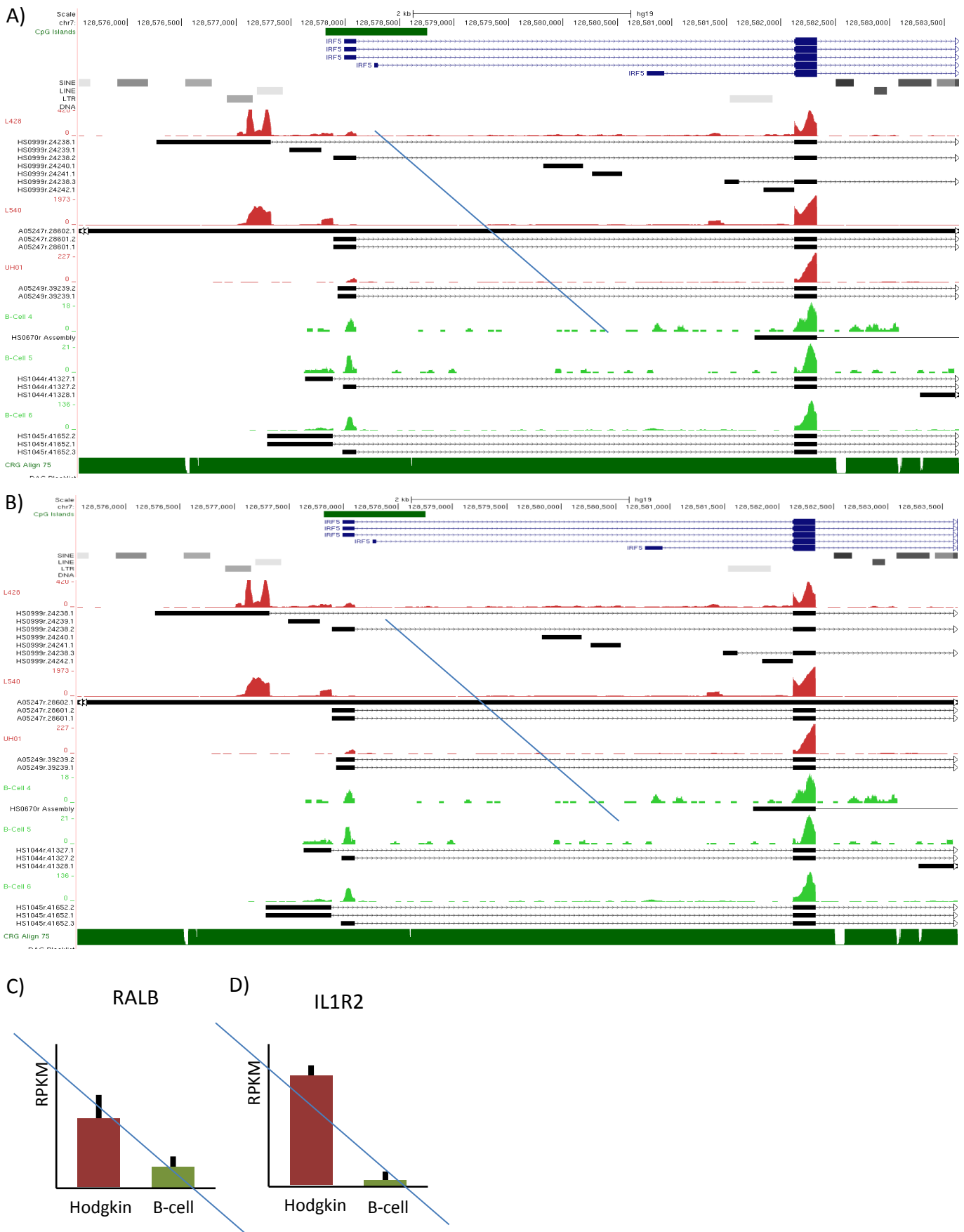


Figure 5: Chimeric Transcription of *RALB* and *IL1R2* in Hodgkin's Lymphoma

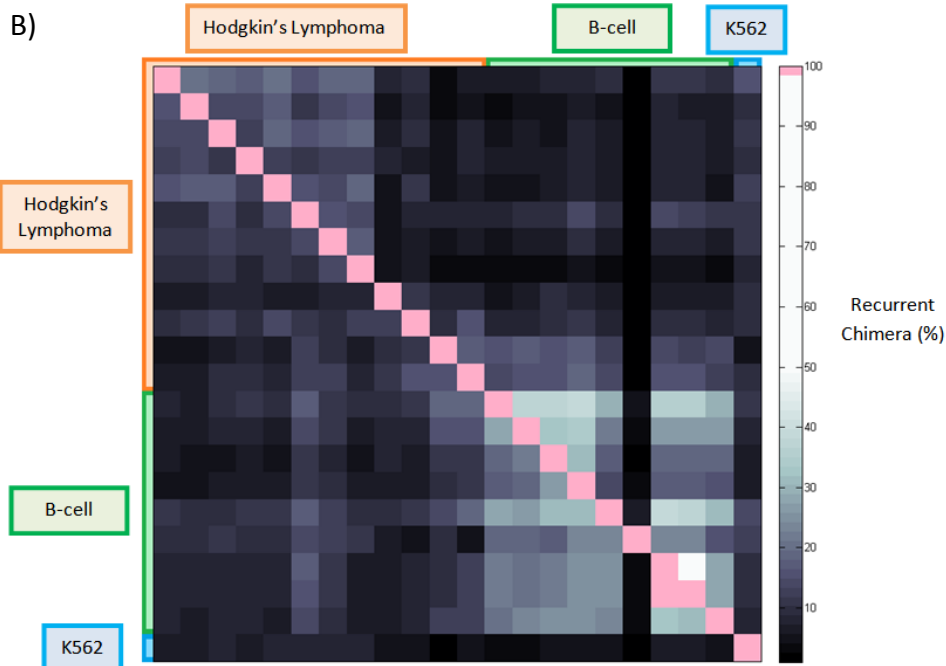
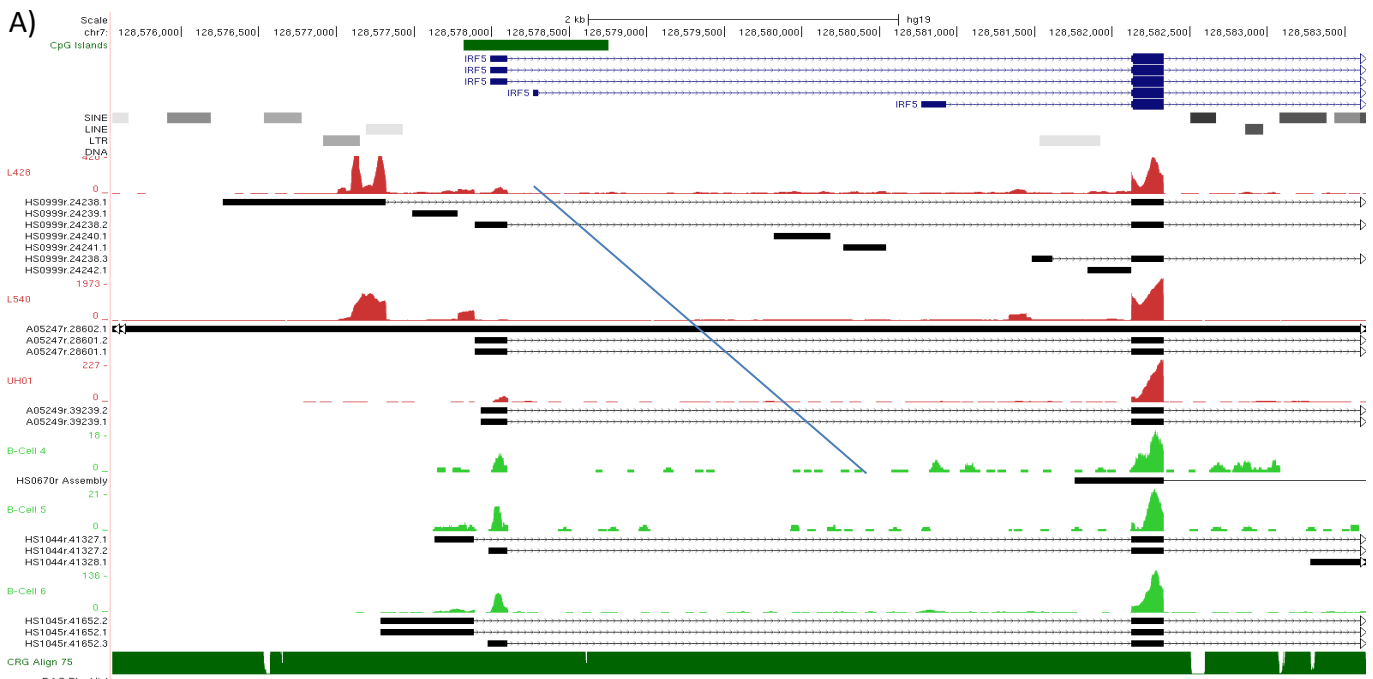


Figure 6: Chimeric Transcription of *HBE* and Chimera Recurrence in Hodgkin's Lymphoma