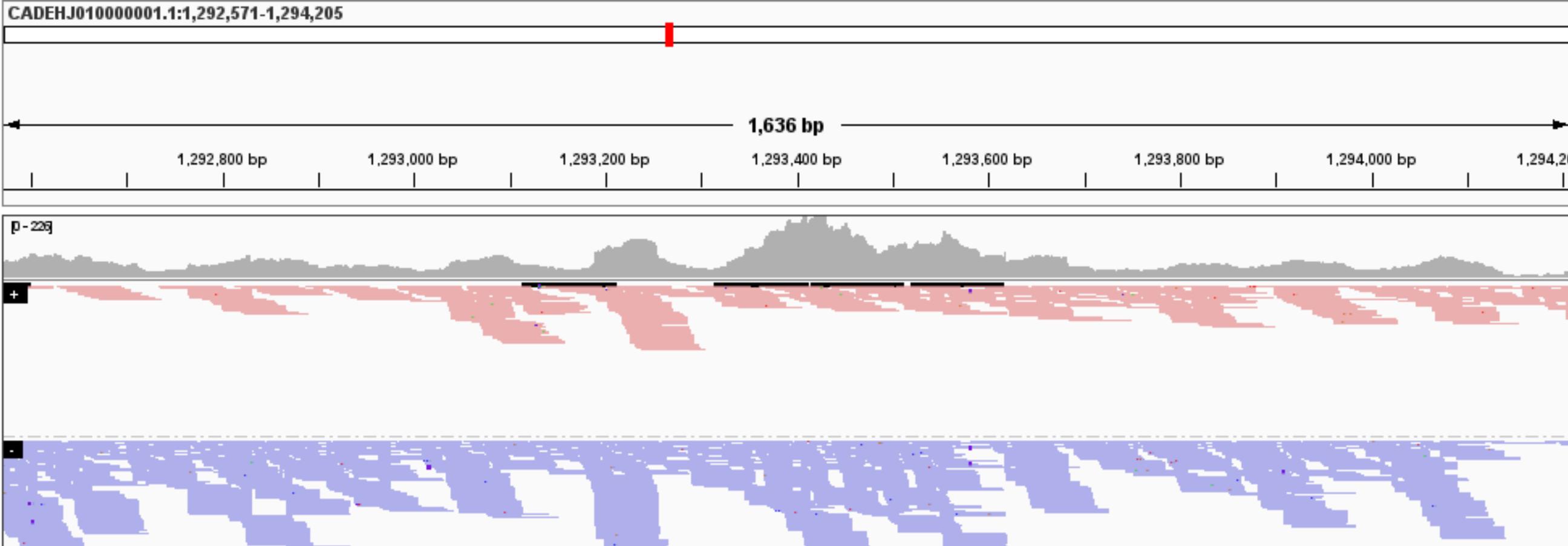
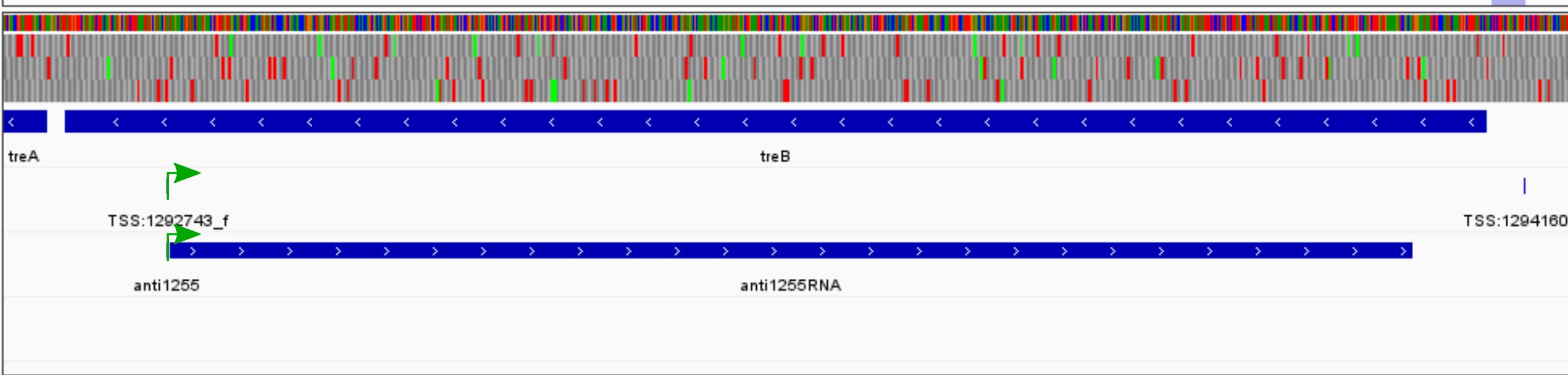
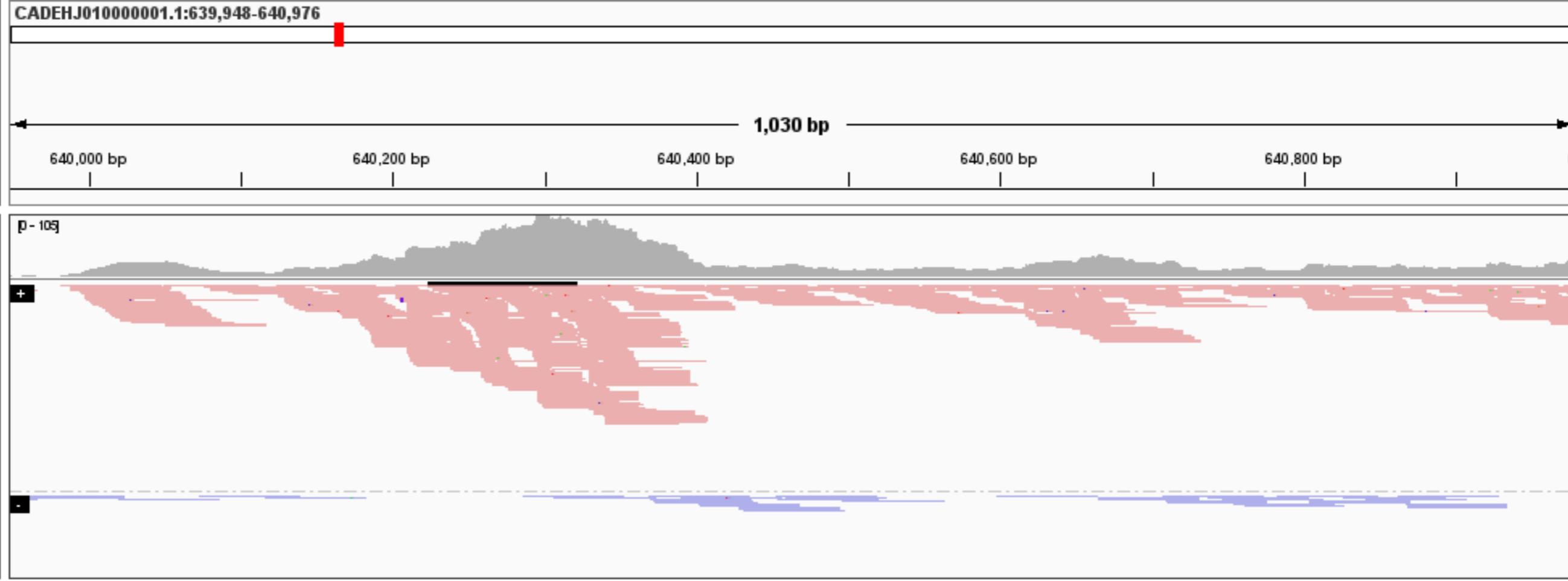


**Figure S3. Visualization of antisense TSS prediction sites verified previously in EGD-e: A) anti1255 and B) anti0605.** Figure shows two rows of reads data, in descending order: combined RNA-seq Illumina reads and combined cap-seq reads. Red lines are forward reads, and blue lines are reverse reads. Reads are sorted by strand. Below reads, annotation rows are shown in order of Genbank gene annotation, ANNOgesic TSS prediction in RO15, EGD-e TSSs lifted to RO15 genome, and ANNOgesic sRNA prediction. Green arrow represents TSSs.

**A)****Combined RNA-seq****Combined Cap-seq****Annotation****RO15 TSSs****EGD-e lifted TSSs  
and asRNAs****Annogesic sRNA  
prediction**

**B)**

Combined RNA-seq



Combined Cap-seq



Annotation

RO15 TSSs

EGD-e lifted TSSs  
and asRNAs

Annogesic sRNA  
prediction

