

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

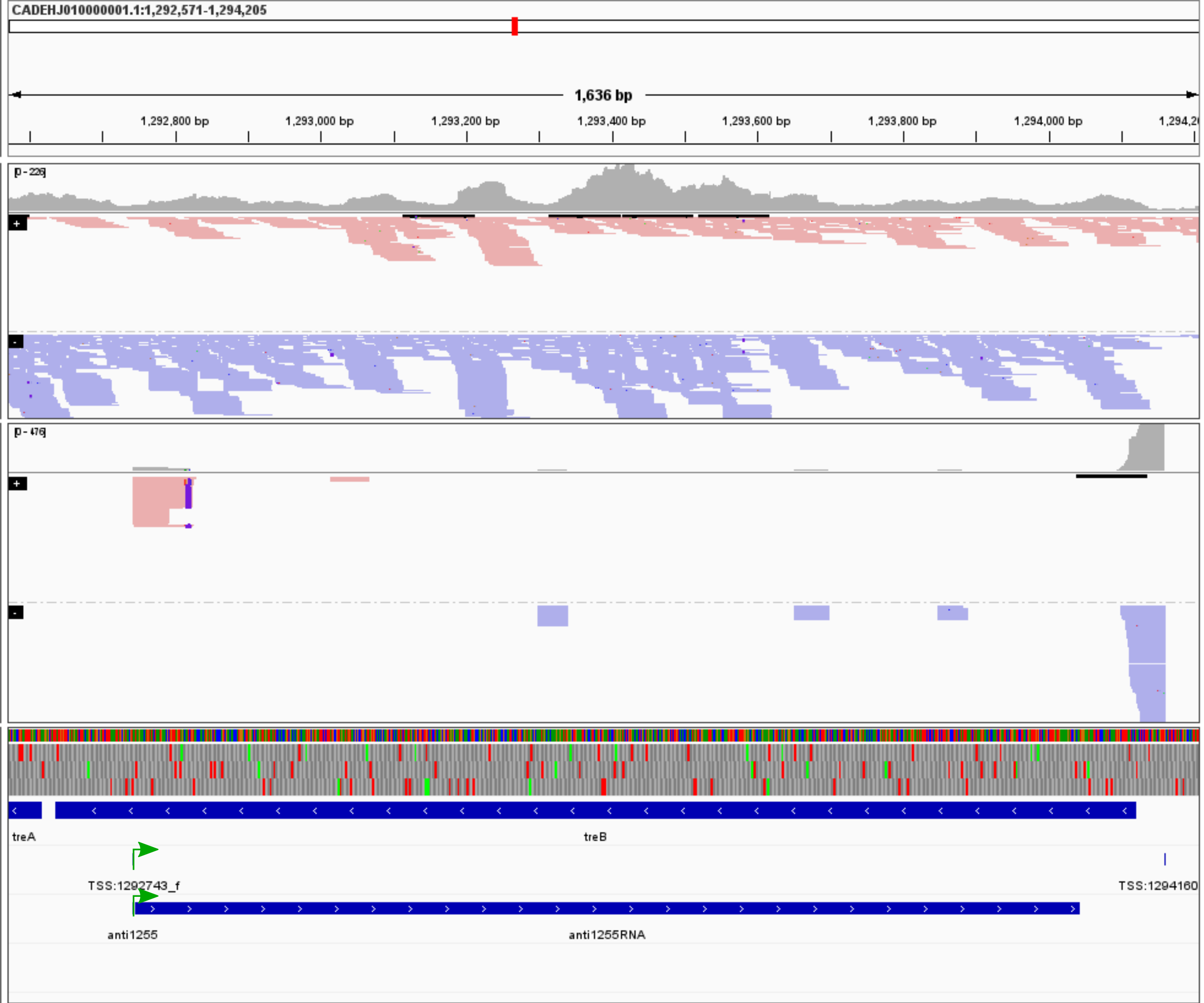
Combined Cap-seq

Annotation

RO15 TSSs

EGD-e lifted TSSs
and asRNAs

Annogestic sRNA
prediction



B)

Combined RNA-seq

Combined Cap-seq

Annotation

RO15 TSSs

EGD-e lifted TSSs
and asRNAs
Annogesic sRNA
prediction

