

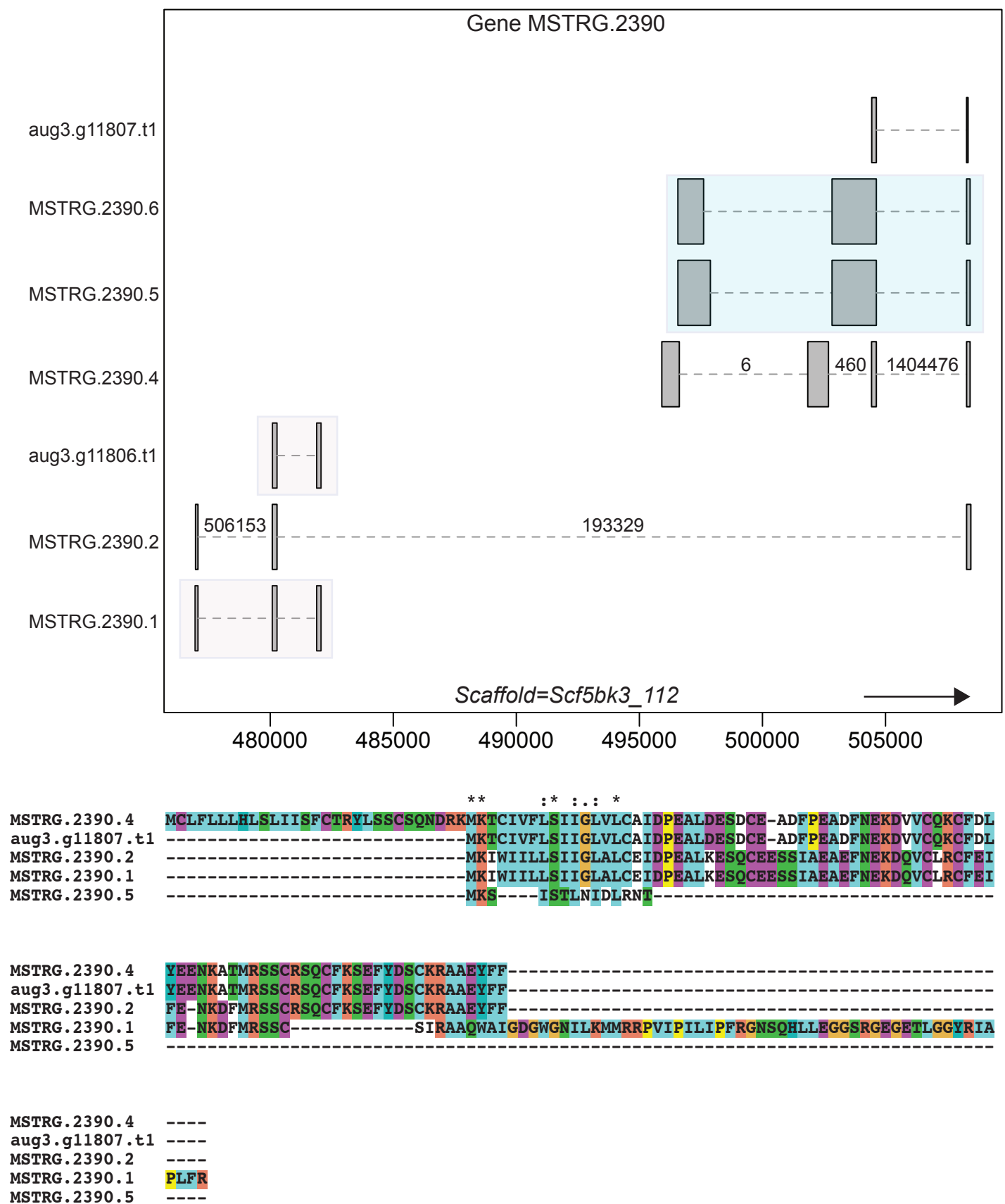


Figure S4. The exon-intron structure of predicted transcripts at gene MSTRG.2390 (no BLAST hit) is shown at top. The alignment of all distinct proteins predicted at this locus is shown at bottom. Transcripts producing identical proteins are indicated by shaded boxes of the same color. The arrow indicates inferred direction of transcription. Numbers over introns represent spliced read counts for novel junctions across all libraries.