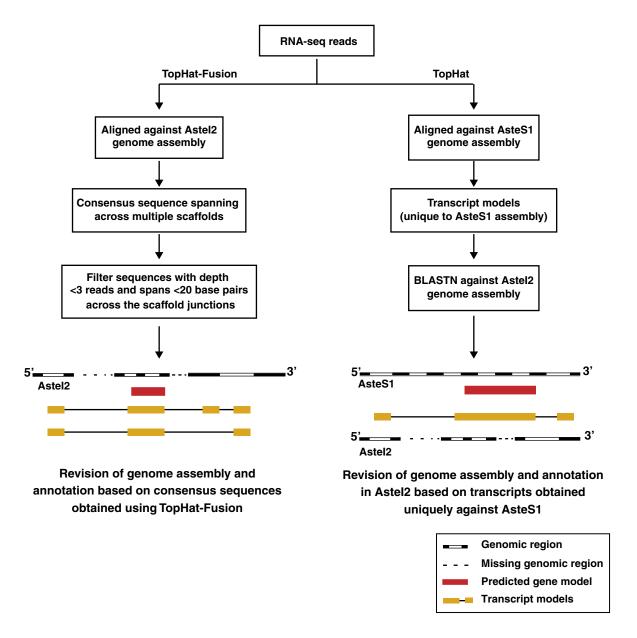
Supplemental Figure S1



Supplemental Figure S1. Improvement in genome annotation and assembly based on RNA-seq data. Transcripts spanning to multiple scaffolds were obtained using two independent approaches, which led to improvement in genome assembly and annotation in Astel2 and AsteS1 genome assemblies.