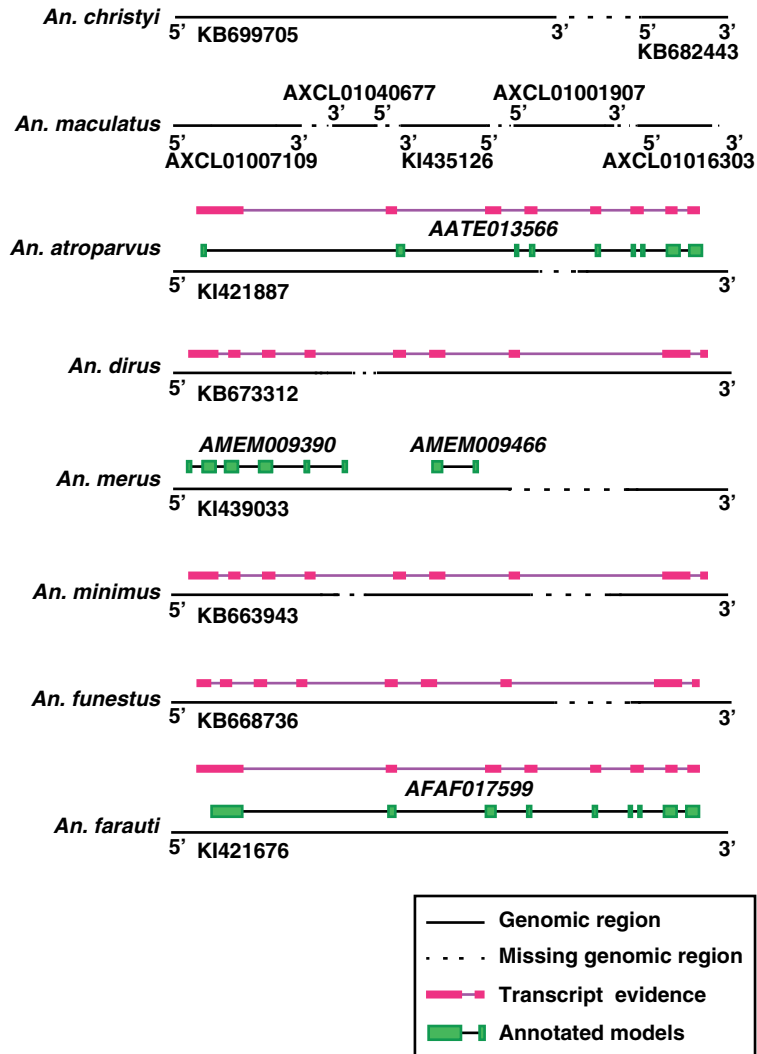
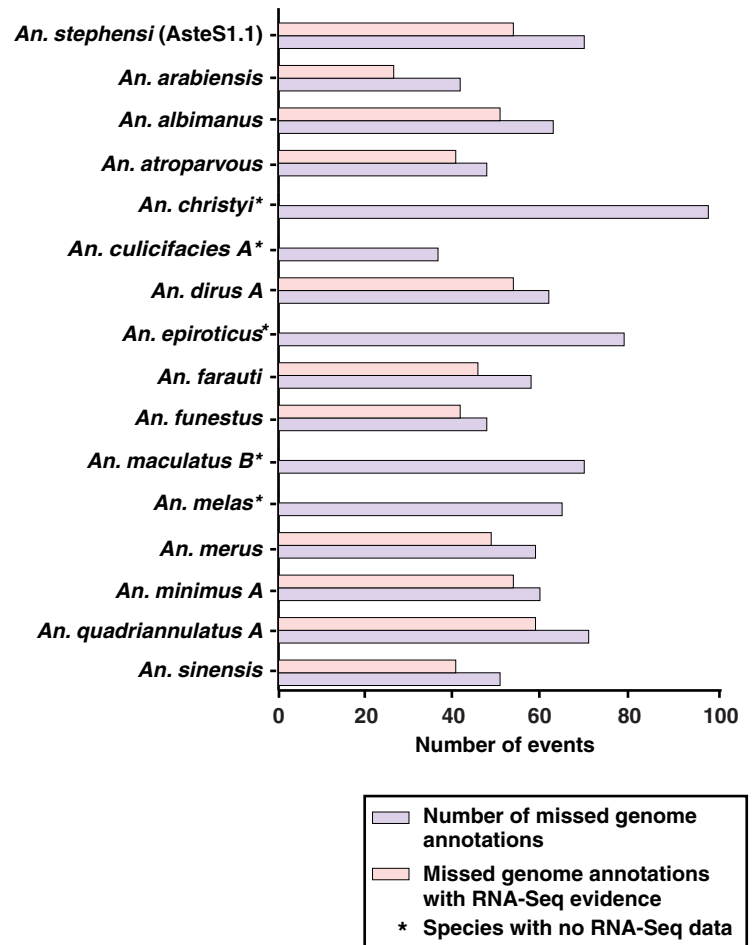


A



B



Supplemental Figure S4. Identification of missed and revised genome annotation in 16 Anopheline species. **A.** shows representation of comparison of one such example against rest of the genomes. **B.** Most of the missed genome annotations identified across 15 Anopheline genomes were supported by RNA-seq evidence from the same species.