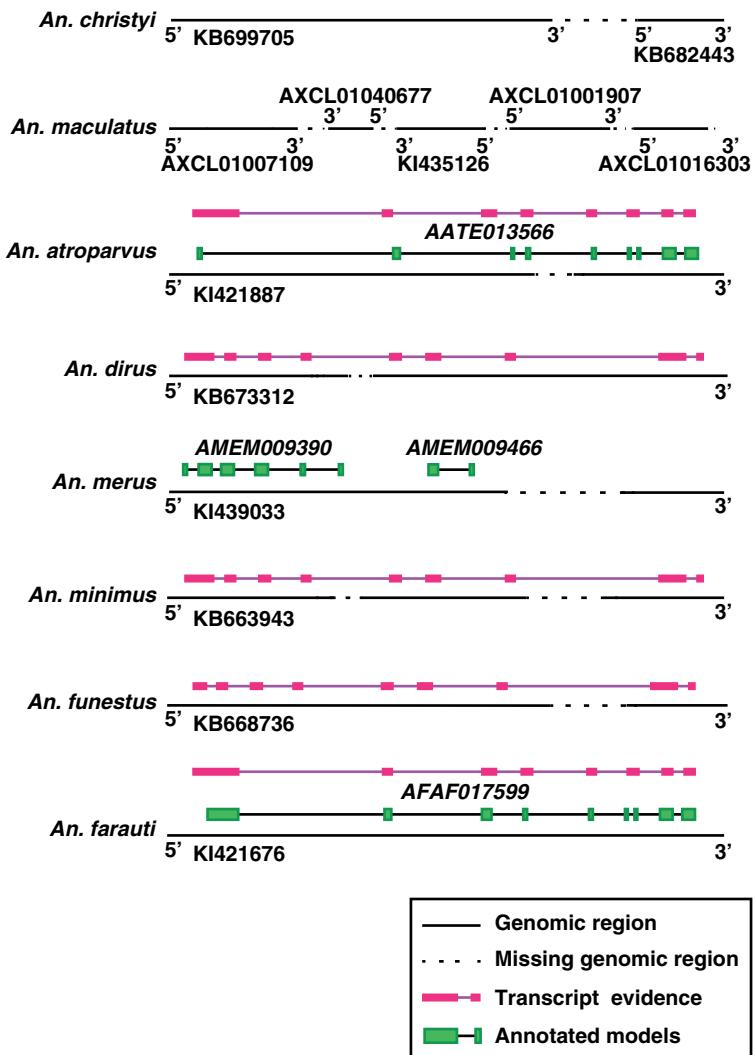
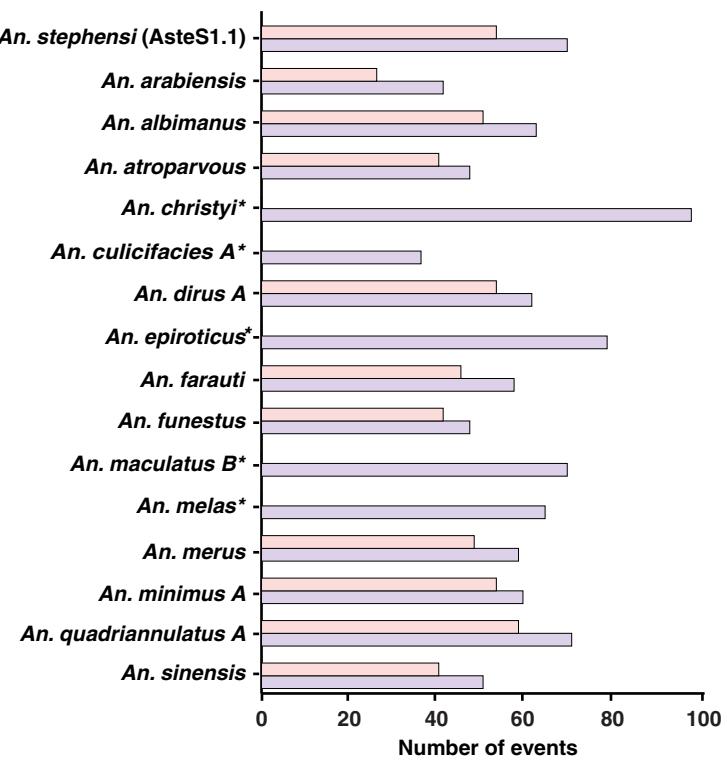


## Supplemental Figure S4

**A**



**B**



- Number of missed genome annotations
- Missed genome annotations with RNA-Seq evidence
- \* Species with no RNA-Seq data

**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.