

Supplementary Figure 1. Regions in the *Trachelipus rathkei* genome assembly matching the *Wolbachia* cytoplasmic incompatibility genes *cifA* and *cifB*. Black bars on top represent a contig in the *T. rathkei* assembly. Lines underneath represent BLAST hits against *cifA* and *cifB* from different *Wolbachia* genomes; numbers to the left and right indicate sequence coordinates (amino acids) of the endpoints of the matching regions of the *cifA* and *cifB* genes in the wCon genome (which has multiple copies of each gene).

