

Figure S3. A Bayesian phylogenetic tree of *Rickettsia* sequences based on 16S rDNA sequences (GenBank accession MN621120 - MN621139) with *Orientia tsutsugamuschi*, *Wolbachia- E. formosa*, *Wolbachia- N. vitripennis* as outgroups. Bayesian posterior probability values are given above branches.

