Supplementary Figure S1:

**Figure S1.** (A). GO functional classification of DEGs between CLso infected and uninfected midguts of B. trigonica. (B). Pathway enrichment statistics of DEGs between CLso infected and uninfected midguts of B. trigonica. Accession numbers: The transcriptome de novo assembly generated in this study is available at the DDBJ/EMBL/GenBank database under the BioProject ID: PRJNA561752. Accession number of the raw reads are SRR10019128 and the transcriptome assembly is GHUW01000000.