

**Supplementary Figure 4.** Tissue-specific expression patterns of Cry receptor encoding genes in different tissues of *S. litura* fourth-instar larvae. Fold change in expression of a candidate gene in different tissues (fat body (FB), foregut (FG), midgut (MG), hindgut (HG) and Malpighian tubules (MT)) was quantified in relation to the gene's expression in head (H) tissue (value set at 1). Significant differential expression is indicated by different letters ( $P < 0.01$ , Tukey's HSD test). *S. litura* *GAPDH* gene was used as the internal reference. Each bar represents the mean fold change value  $\pm$  SE of qPCR runs in five biological and three technical replicates.

