

Identification of the Gene Repertoire of the IMD Pathway and Expression of Antimicrobial Peptide Genes

in Several Tissues and Hemolymph of the Cockroach *Blattella germanica*

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Supplementary Figure S1

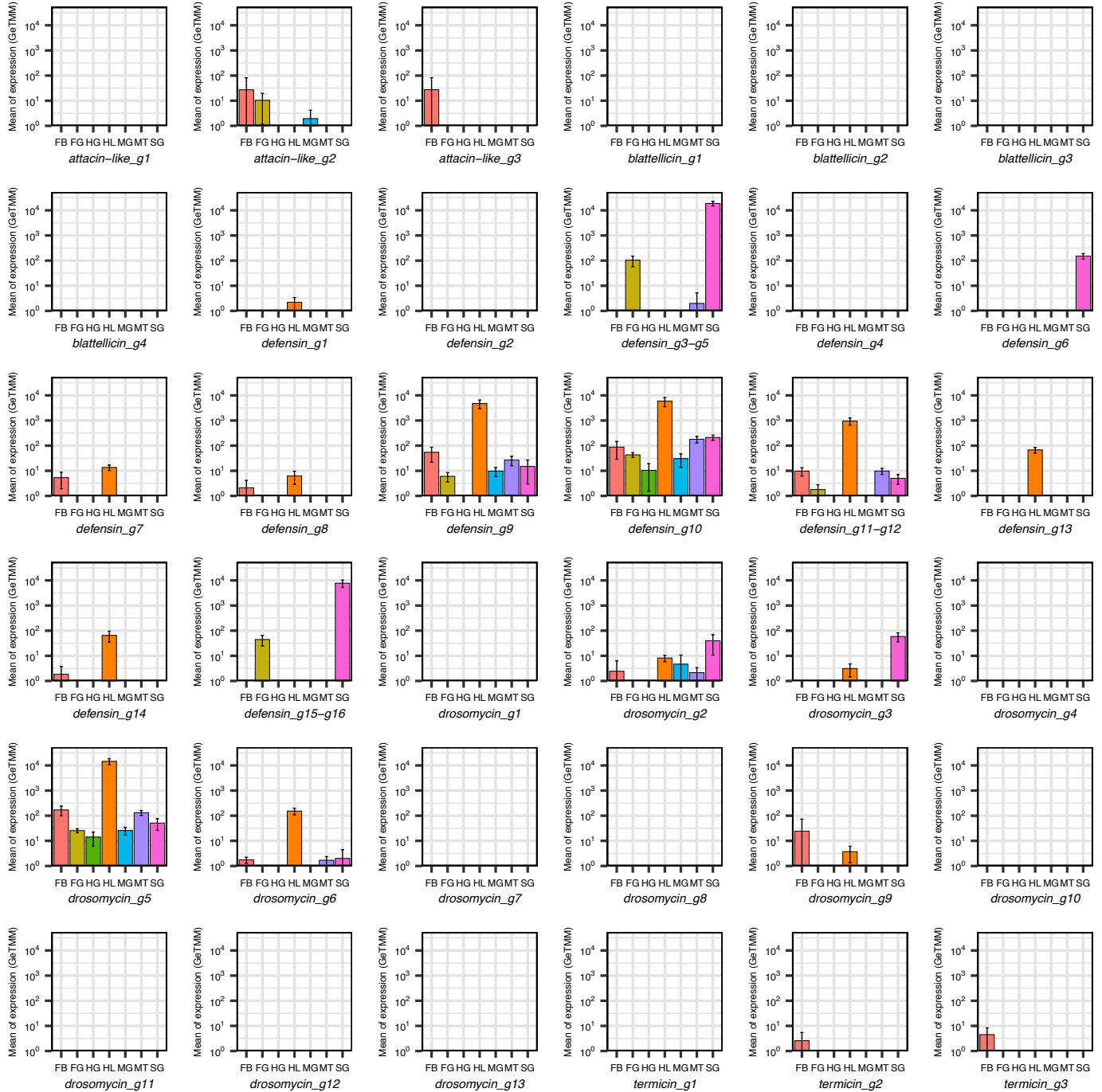


Figure S1. Expression patterns of each AMP gene in the analyzed sample types of *B. germanica*. Means and standard deviations of normalized expression (GeTMM) of AMP genes are shown. Abbreviations: FB (fat body), FG (foregut), HG (hindgut), HL (hemolymph), MG (midgut), MT (Malpighian tubules) and SG (salivary glands). Graph bars are standard deviations. Y-axis is in log scale. The lower ends of the error bars are not shown below 10^0 .