Functional and evolutionary insights from the genomes of three parasitoid Nasonia species

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Table S50: Hymenoptera-specific orthologs. For each orthologous group (OG), the number of paralogs in the 2 hymenopterans *A. mellifera* (*A. mel.*) and *N. vitripennis* (*N. vit.*) is shown together with BLASTP E-values of the bee protein to *N. vit.*, *D. melanogaster* (*D. mel.*), and *T. castaneum* (*T. cas.*). Some bee proteins have a significant BLASTP hit with ESTs of a third hymenopteran species, *S. invicta* (fireant).

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