

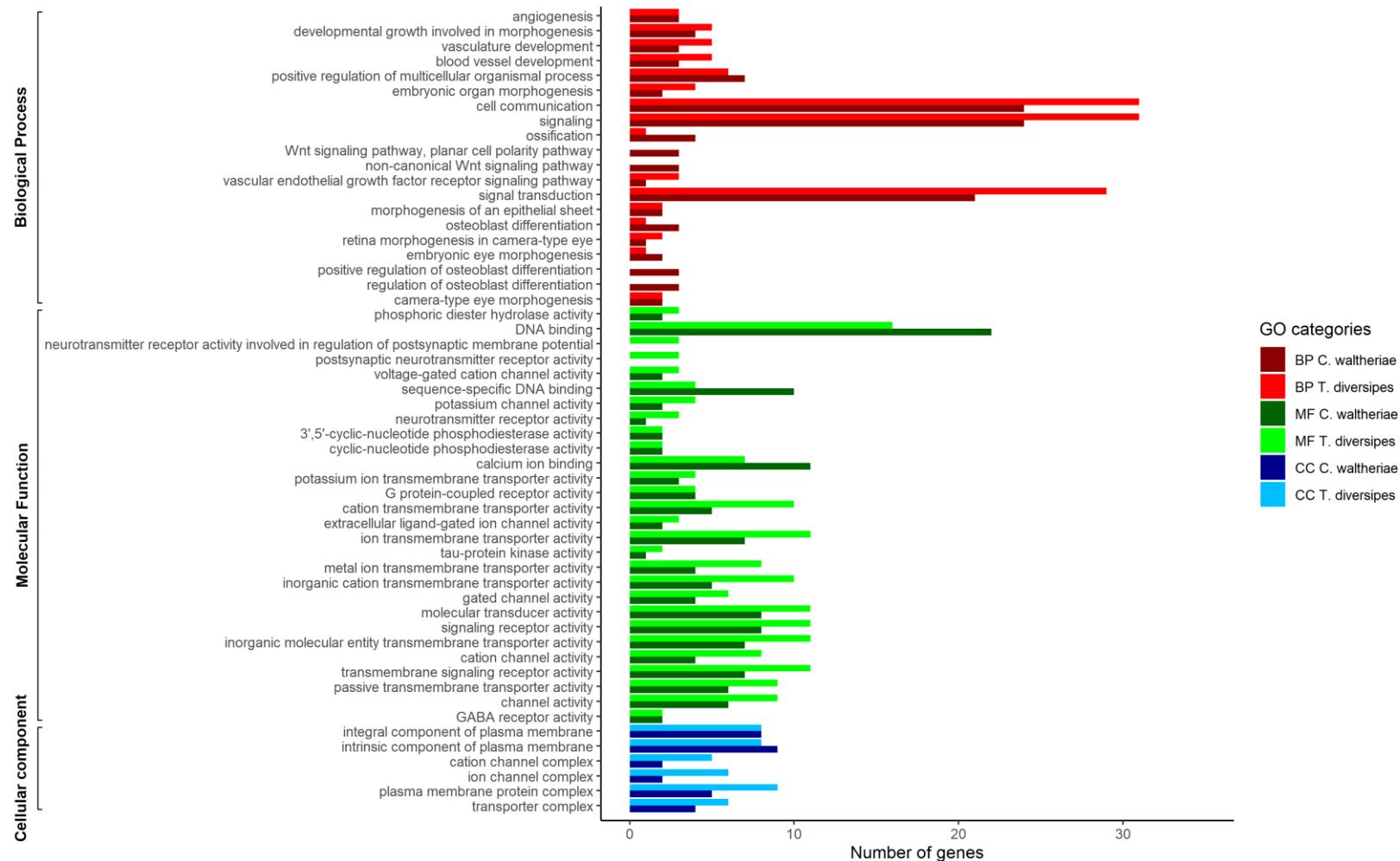
# Decoding bee cleptoparasitism through comparative transcriptomics of *Coelioxoides waltheriae* and its host *Tetrapedia diversipes*

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**Figure 2.** Gene ontology (GO) functional classification of *Coelioxoides waltheriae* (darker) and *Tetrapedia diversipes* (lighter) differently expressed orthologs. The GO functional annotations were classified into three main categories: Cellular Component (blue), Molecular Function (green) and Biological Process (red).