

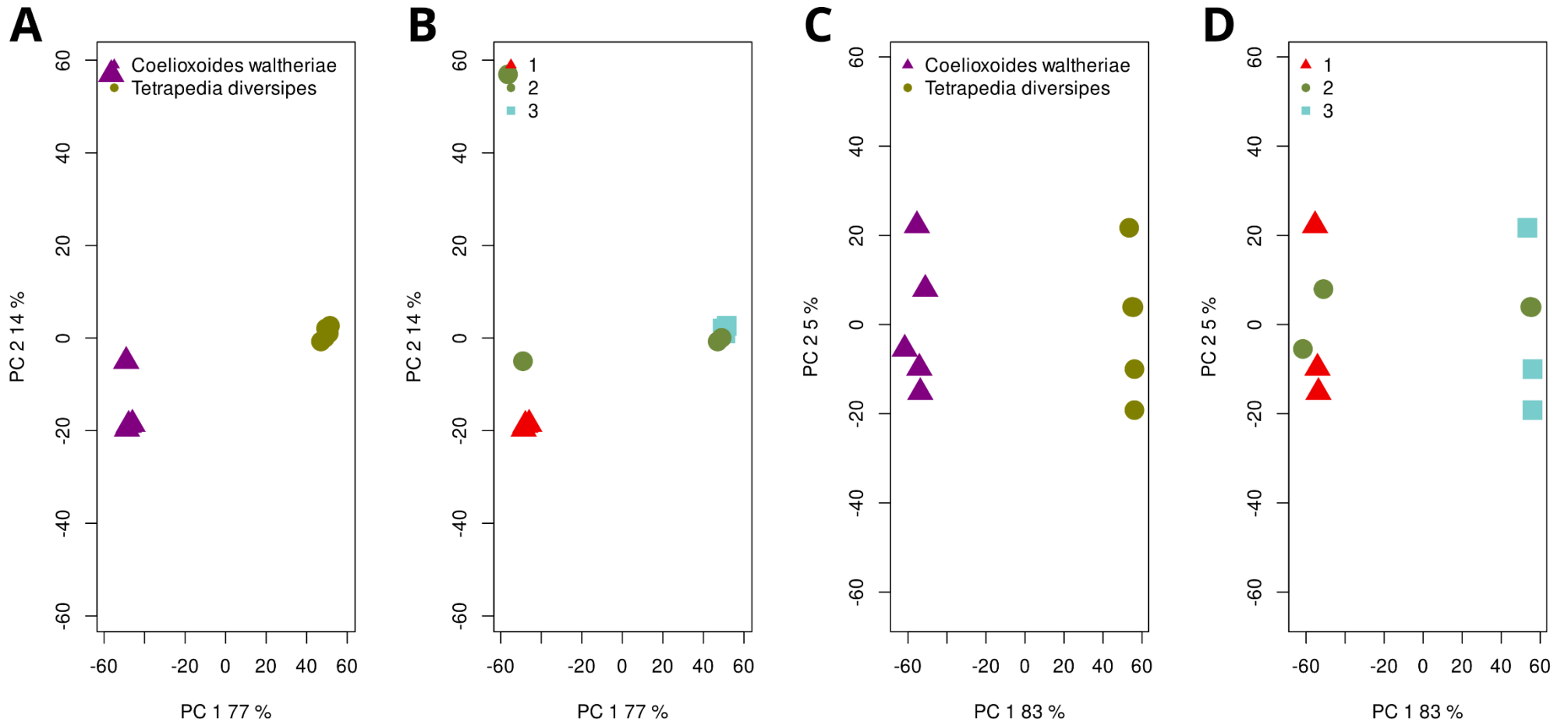
# Decoding bee cleptoparasitism through comparative transcriptomics of *Coelioxoides waltheriae* and its host *Tetrapedia diversipes*

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**Figure 1.** Principal component analysis (PCA) plots showing samples of *Coelioxoides waltheriae* (purple triangles) and *Tetrapedia diversipes* (olive circles) based on gene expression profile, before (A) and after (C) batch correction on raw data. The samples from each batch (batch 1: red triangles; 2: green circles; 3: blue squares) are represented in the same way, before (B) and after (D) correction. The X- and Y-axes represent the first and second principal components (PC1 and PC2) and the associated percentage of variation.