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Corrigendum: Comparative analysis of the complete mitochondrial genomes of three *Zeugodacus* species (Insecta: Tephritidae: Dacinae) and their phylogenetic relationships with other congeners. Arthropod Systematics & Phylogeny 81: 747–759. doi: 10.3897/asp.81.e105025

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The article entitled "Comparative analysis of the complete mitochondrial genomes of three *Zeugodacus* species (Insecta: Tephritidae: Dacinae) and their phylogenetic relationships with other congeners", published in Arthropod Systematics & Phylogeny (DOI: 10.3897/asp.81.e105025), requires two corrections that do not affect the scientific content or integrity of the article. Figures 2 and 3, as presented in this corrigendum, now reflect our original intent.

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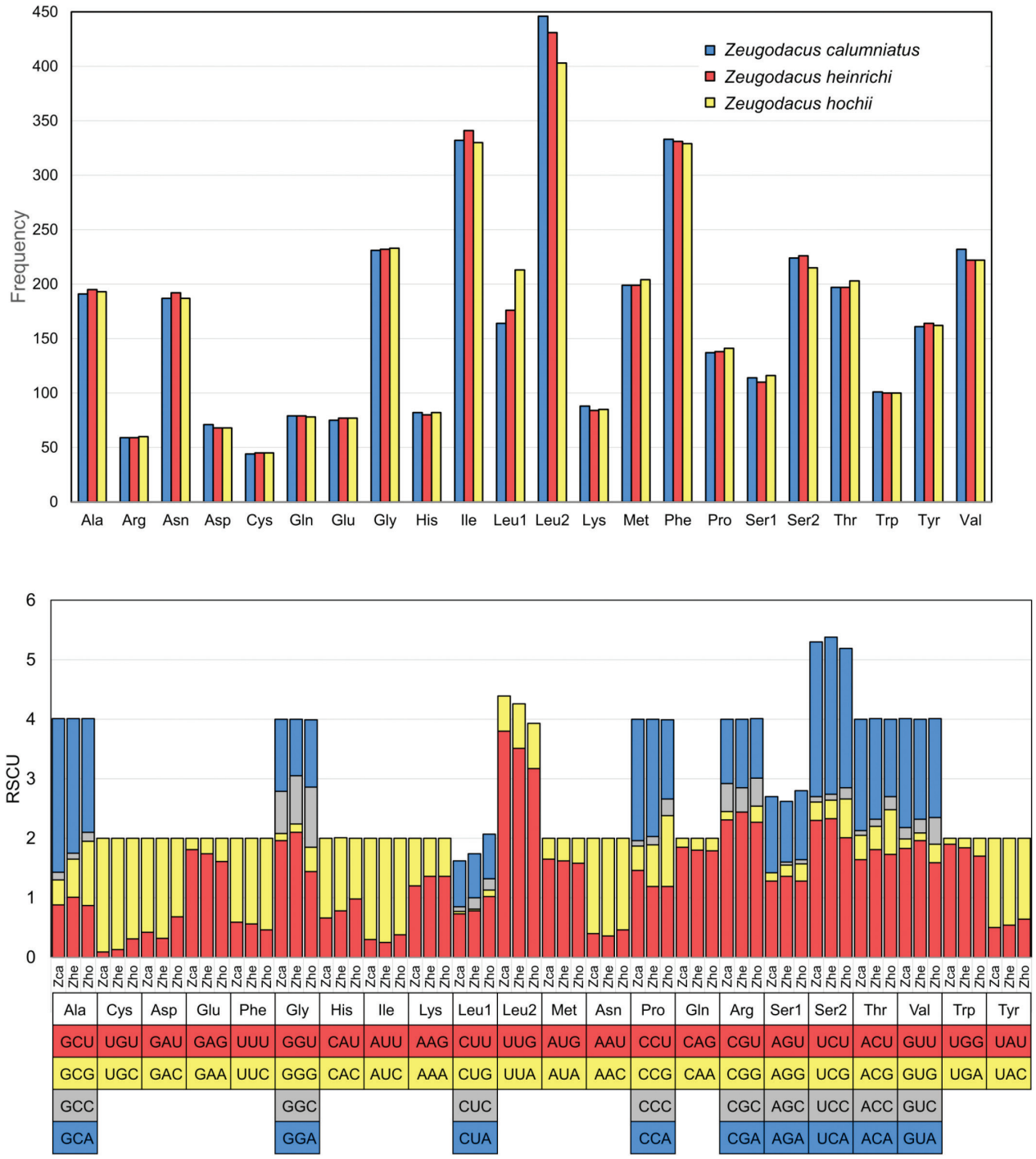

Figure 2. Amino acid frequency (Top) and relative synonymous codon usage (Bottom) of PCGs in the *Zeugodacus* mitogenomes generated using MEGA X (https://www.megasofware.net/). Zca, *Zeugodacus calumniatus*; Zhe, *Zeugodacus heinrichi*; Zho, *Zeugodacus hochii*.

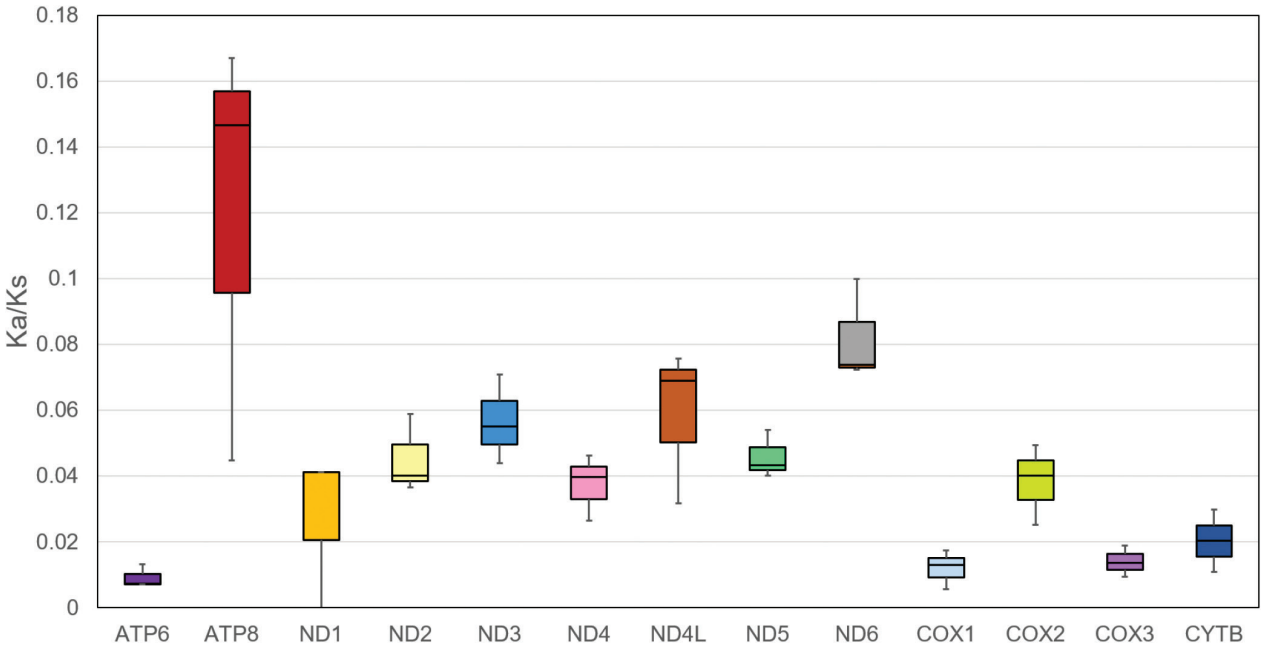

Figure 3. Box plot for pairwise divergence of Ka/Ks ratio (mean ± SD, and range) for 13 protein-coding genes (PCGs) of three *Zeugodacus* mitogenomes (*Z. calumniatus*, *Z. heinrichi*, *Z. hochii*) generated using DnaSP6.0. (http://www.ub.edu/dnasp).