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**Title: Shaping the microbial landscape: Parasitoid-driven modifications of *Bactrocera dorsalis* microbiota**

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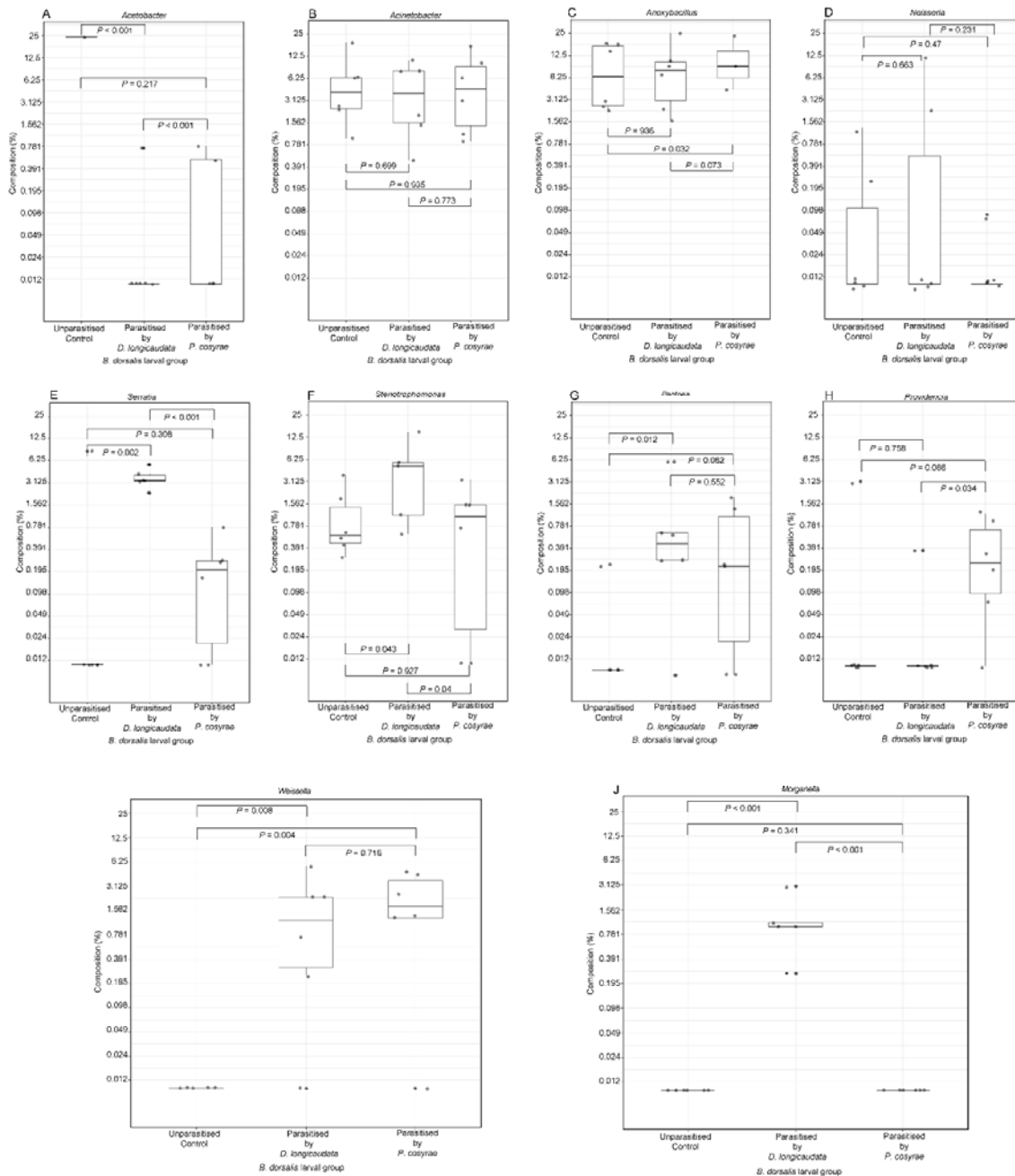
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Supplementary figure 1: Relative abundances of selected bacterial genera across the different *Bactrocera dorsalis* larval groups (Control, parasitized by *Diachasmimorpha longicaudata* and those parasitized by *Psytalia cosyrae*). Comparison across groups for A) *Acetobacter*, B) *Acinetobacter* c) *Anoxybacillus*, D) *Neisseria*, E) *Serratia*, F) *Stenotrophomonas*, G) *Pantoea*, H) *Providencia* I) *Weissella* and J) *Morganella* . Comparisons between groups are significant when  $P < 0.05$ .