

Bulletin of Insectology Supplemental Material (supplement 2)

Title: **Distribution and genetic variability of Staphylinidae across a gradient of anthropogenically influenced insular landscapes**

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Figure S5. Neighbor Joining tree for Staphylinidae species in subfamilies and tribes of the four informal groups of Staphylinidae *sensu* Lawrence and Newton (1995).

Figure S5. Neighbor Joining tree for Staphylinidae species in subfamilies and tribes of the four informal groups of Staphylinidae *sensu* Lawrence and Newton (1995), found in the Azores and worldwide, for which COI sequences were generated (1265 bp) or available in Genbank, respectively. Staphylinidae groups colored as follows: Tachyporine (Green), Omaliine (Red), Oxytelina (Black) and Staphylinina (Brown). Outgroup was colored in Brown. Taxa are followed by their Genbank accession number, tribe or subfamily (*) when tribe was not available in Genbank. COI sequences for taxa in bold were generated in this study. Island of origin for each species is specified as follows: SMG – Sao Miguel, TER – Terceira, SMR – Santa Maria, FLO – Flores and PIC – Pico.

