Erratum: Extending Phylogenetic Regression Models for Comparing Within-Species Patterns Across the Tree of Life

06 November, 2026: Post-production of the manuscript, a coding error was discovered in a downstream function for assembling components of manova statistics, which affected the Table of results for the empirical example. The corrected table is below:

Table 1: MANOVA table of variance components from E-PGLS. All variance components are obtained while conditioning the individual-level data on the phylogeny as described in the paper.

	Wilks	LR	Z	pval
Species	0.470	731.296	6.143	0.001
Sex	0.895	107.875	2.760	0.005
Population	0.022	3676.763	27.420	0.001
Species:Sex	0.421	838.337	10.581	0.001
Full.Model	0.005	5102.098	24.553	0.001