



Supplemental Figure 3. Phylogenetically corrected linear regression analysis of the relationship between (A) total TE, (B) *Copia*, (C) *Gypsy* element abundance and genome size for all species, including outliers. PIC=phylogenetically independent contrasts. Asterisks (*) next to r^2 values denotes significant p-values at the $\alpha = 0.1$ level.