



**Supplemental Figure 5.** Phylogenetically corrected linear regression analysis of the relationship between (A) total TE, (B) *Copia*, (C) *Gypsy* element abundance and genome size for 20% sampling effort of genome wide coverage. PIC=phylogenetically independent contrasts. Asterisks (\*) next to  $r^2$  values denotes significant p-values at the  $\alpha = 0.1$  level.