



Figure S5. GC-content variation among taxa in phylogenomic datasets for birds showing the distribution of differences between the minimum and maximum GC-content for each locus (ΔGC_{inf}). Values that fell outside the interquartile range by >1.5 -fold were designated outliers. We limited GC-content calculations to parsimony-informative sites from regions with ≥ 50 parsimony-informative sites. The identity of two outliers in the EB2 data is noted next to the relevant data points. We used the EB2 taxon set for the EB2 data and all of the birds analyzed by Prum et al. (2015).