**Partial alignment from Janecka et al. 2008 *Current Biology***

Dr. Janecka (Janecka Genomics)

**Data file: ColugoMtDNAnoHSAv2.fas**

These are concatenated mtDNA sequences used in the analysis. The file corresponds to the sequence matrix used to make the phylogenetic tree in Figure S1 of the Supplementary Materials file. If you use this alignment please site Janecka et al. (2007).

Positions 1-371 are 12S and positions 372-748 are CYTB.

**Reference**

Janecka JE, Lim NT-L, Baba M, Izawa M, Boeadi, Murphy WJ (2008) Evidence for multiple species of Sunda colugo. *Current Biology* 18:R1001–R1002.