This Dryad deposition includes the following files:

(1) Sequence alignment for MrBayes analysis of mitochondrial DNA
(2) Input files for *X. clivii* and *X. largeni* for MIMAR analysis
(3) Input files for *X. clivii* and *X. largeni* for IMa2 analysis, including a smaller input file for *X. clivii* that has loci with evidence of non-neutral evolution removed
(4) Input files for Structure analysis, including a smaller file that as individuals with lots of missing data removed
(5) Input files for Structurama analysis