

Supplementary Material to “The shared performance of different COI regions increases the niche breadth detectable from bat guano eDNA”

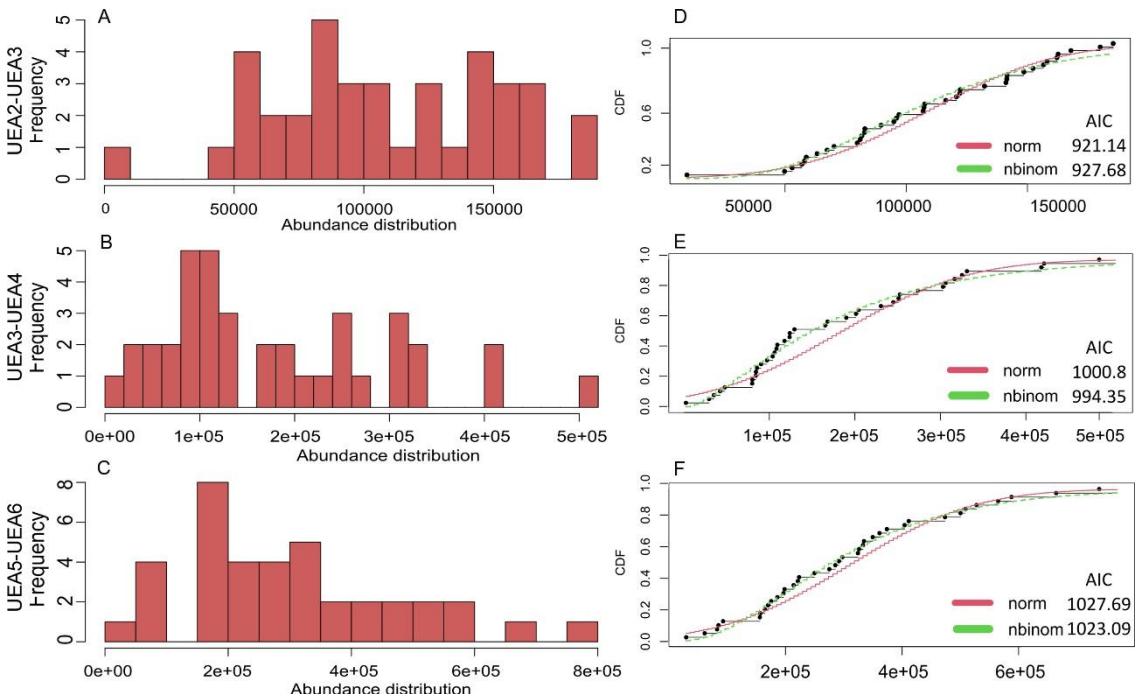


Figure S1 - Total read abundance distribution for UEA2-UEA3, UEA3-UEA4, and UEA5-UEA6.