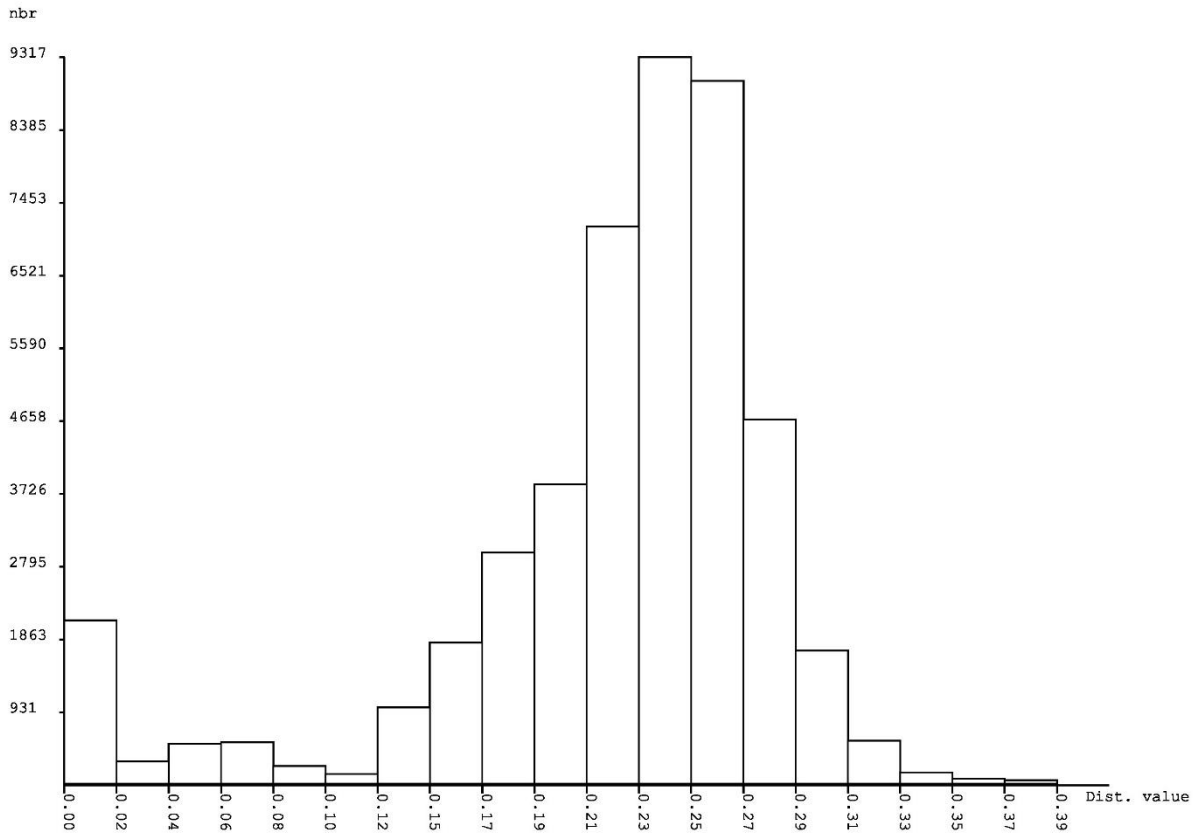


Figure S1. Distribution of genetic distances indicating two modes separated by a barcode gap ranging from 0.02 to 0.12. Species delimitation analysis using the ABGD method.