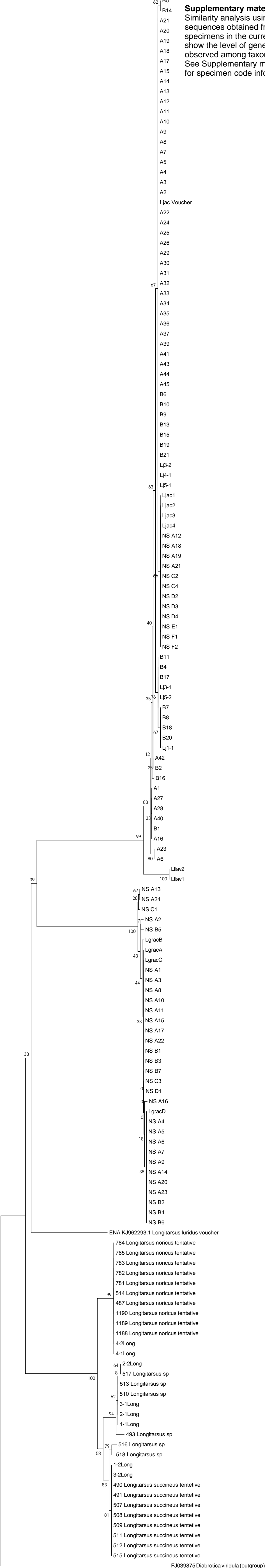


Supplementary material, Figure S1.
 Similarity analysis using all *CO1* gene sequences obtained from *Longitarsus* specimens in the current study. Results show the level of genetic variation observed among taxonomic groups. See Supplementary material, Table S1 for specimen code information.



0.02