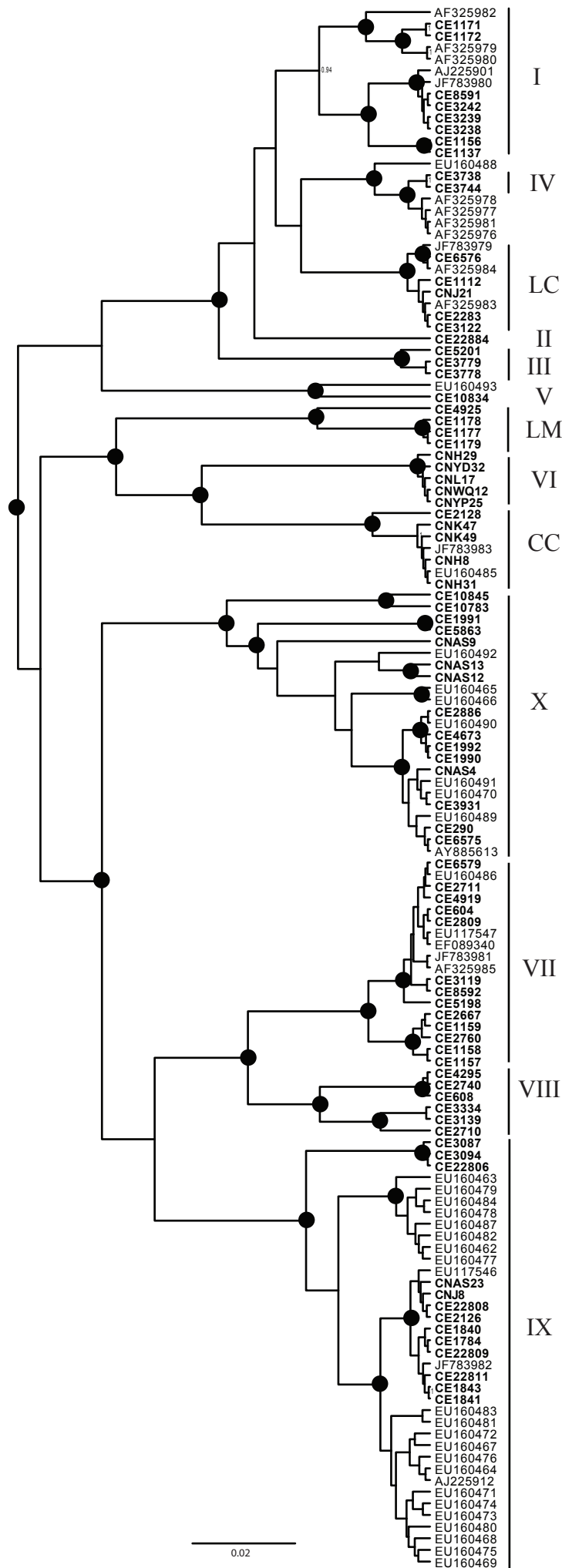




Supplementary Fig. S3

Maximum clade credibility tree for mitochondrial 16S dataset, with nodal support given as Bayesian posterior probability (PP): black dots at nodes denote $PP \geq 0.99$. Note that data include several sequences (in regular font) downloaded from GenBank, see Table S1 for details.

Species labelled I-IV and VI-X are identified as members of the *L. hoffmeisteri* complex, CC as "*L. clapedianus-cervix*", LM as *L. maumeensis*, and LC as *L. clapedianus*.