

Figure S2. Gene tree derived from STACEY analysis for BEAST-2 using the *trnT-L-F* marker sequences. PP values ≥ 0.90 on the branches.

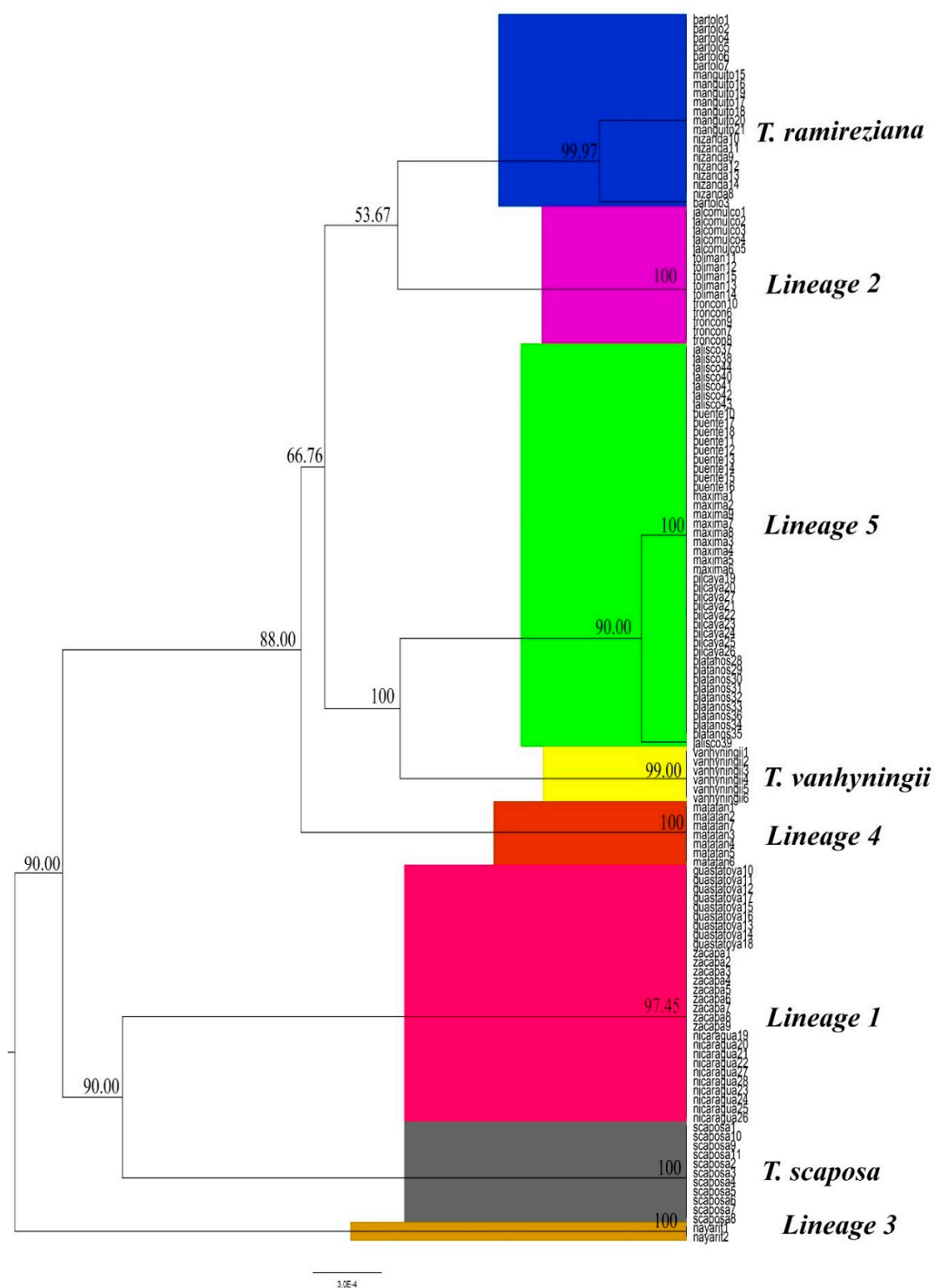


Figure S3. Gene tree derived from STACEY analysis for BEAST-2 using the *rps16-trnQ* marker sequences. PP values ≥ 0.90 on the branches.

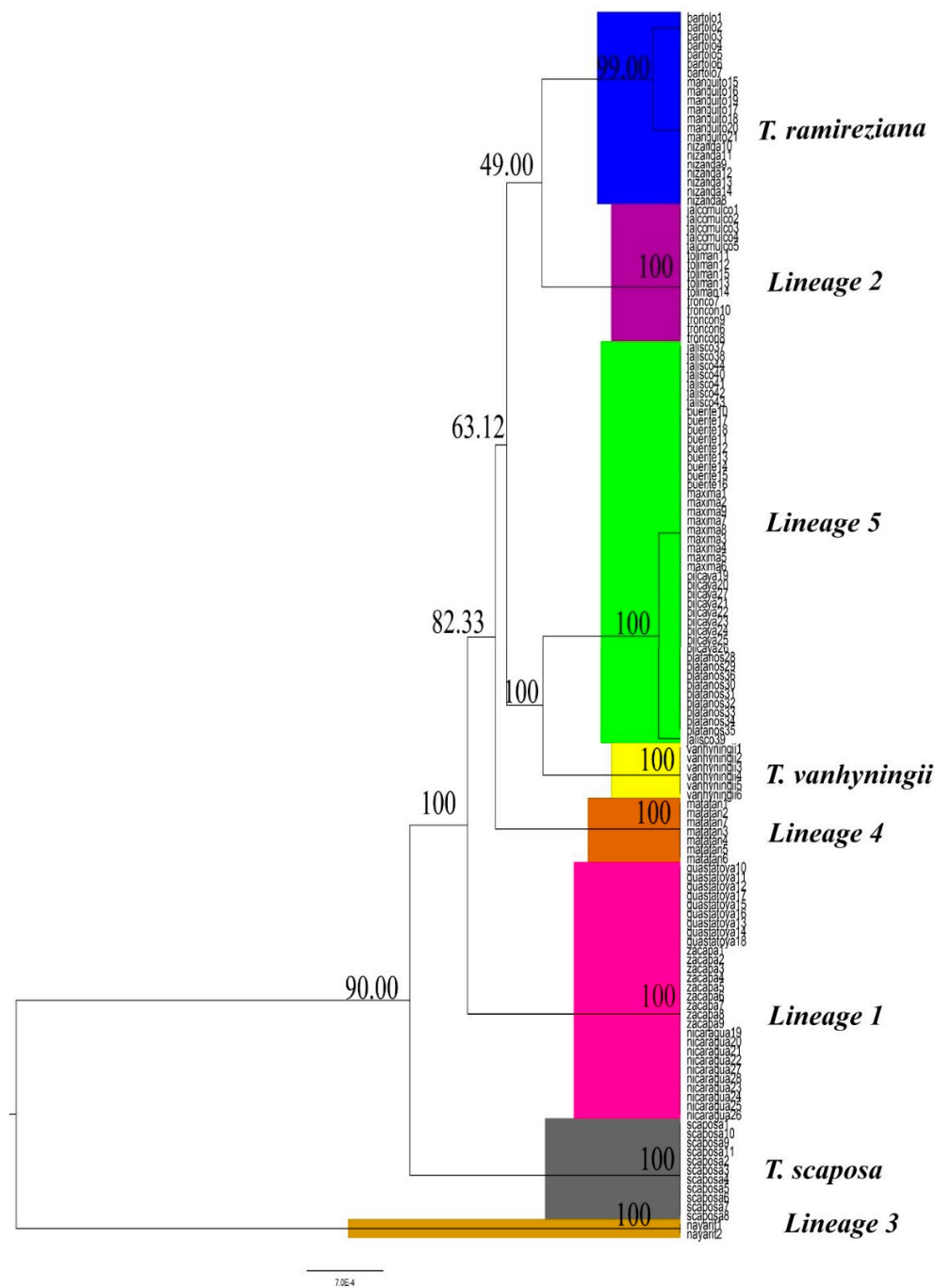


Figure S4. Gene tree derived from STACEY analysis for BEAST-2 using the *rpl16-rps3* marker sequences. PP values ≥ 0.90 on the branches.