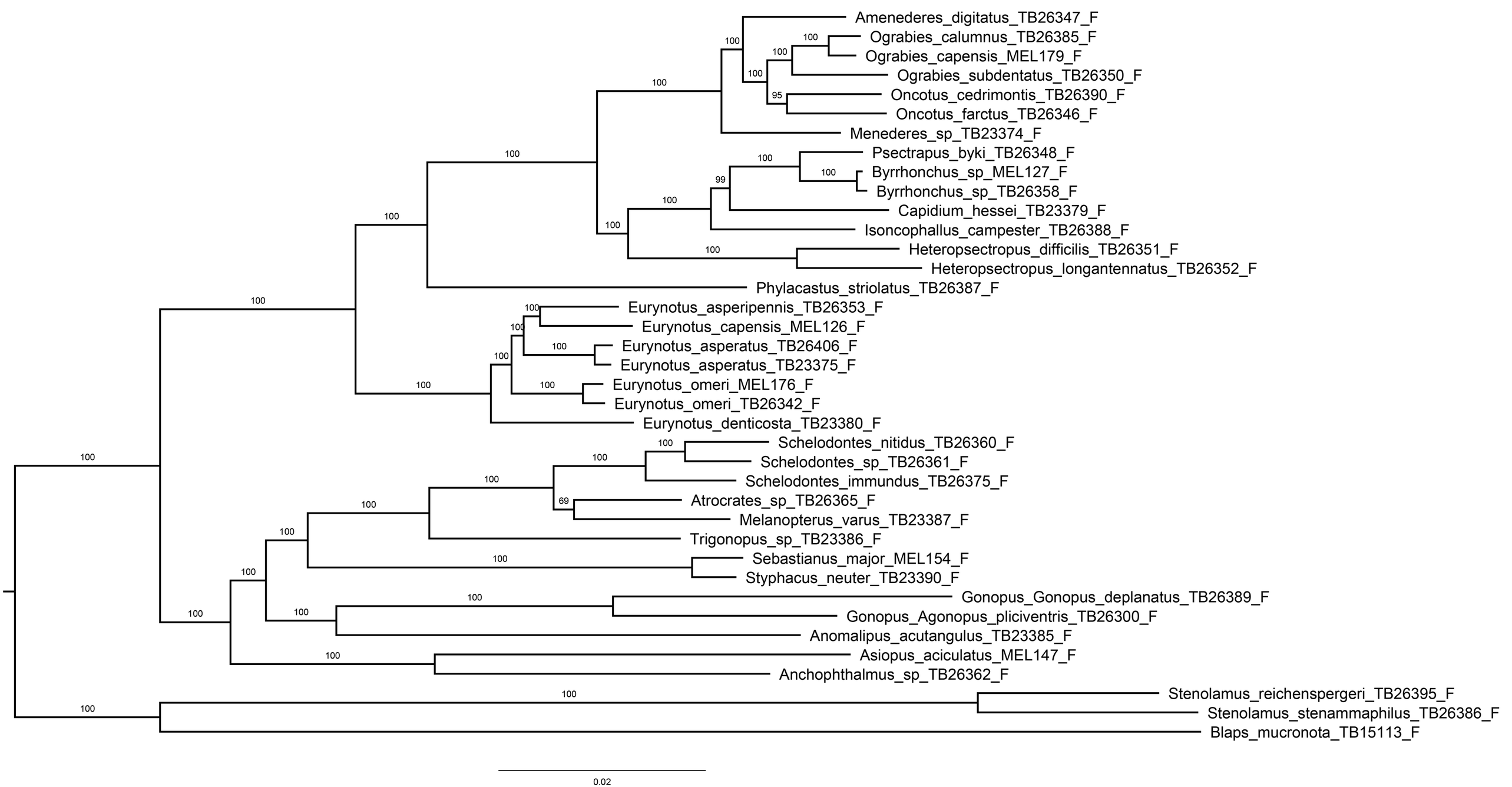

Phylogeny of the tribe Platynotini inferred from a targeted enrichment dataset. Only fresh, ethanol-preserved specimens were included. The tree shown is the Majority Rule consensus from a IQ-TREE2 analysis of the concatenated matrix comprising 499 loci and 155,172 aligned amino acid positions. Ultrafast bootstrap values (IQ-TREE 2) are indicated below above.