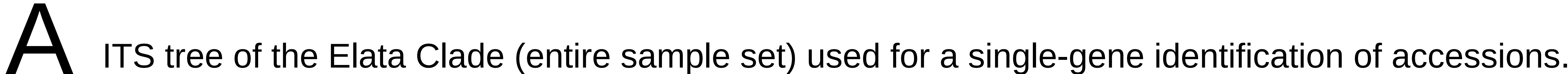
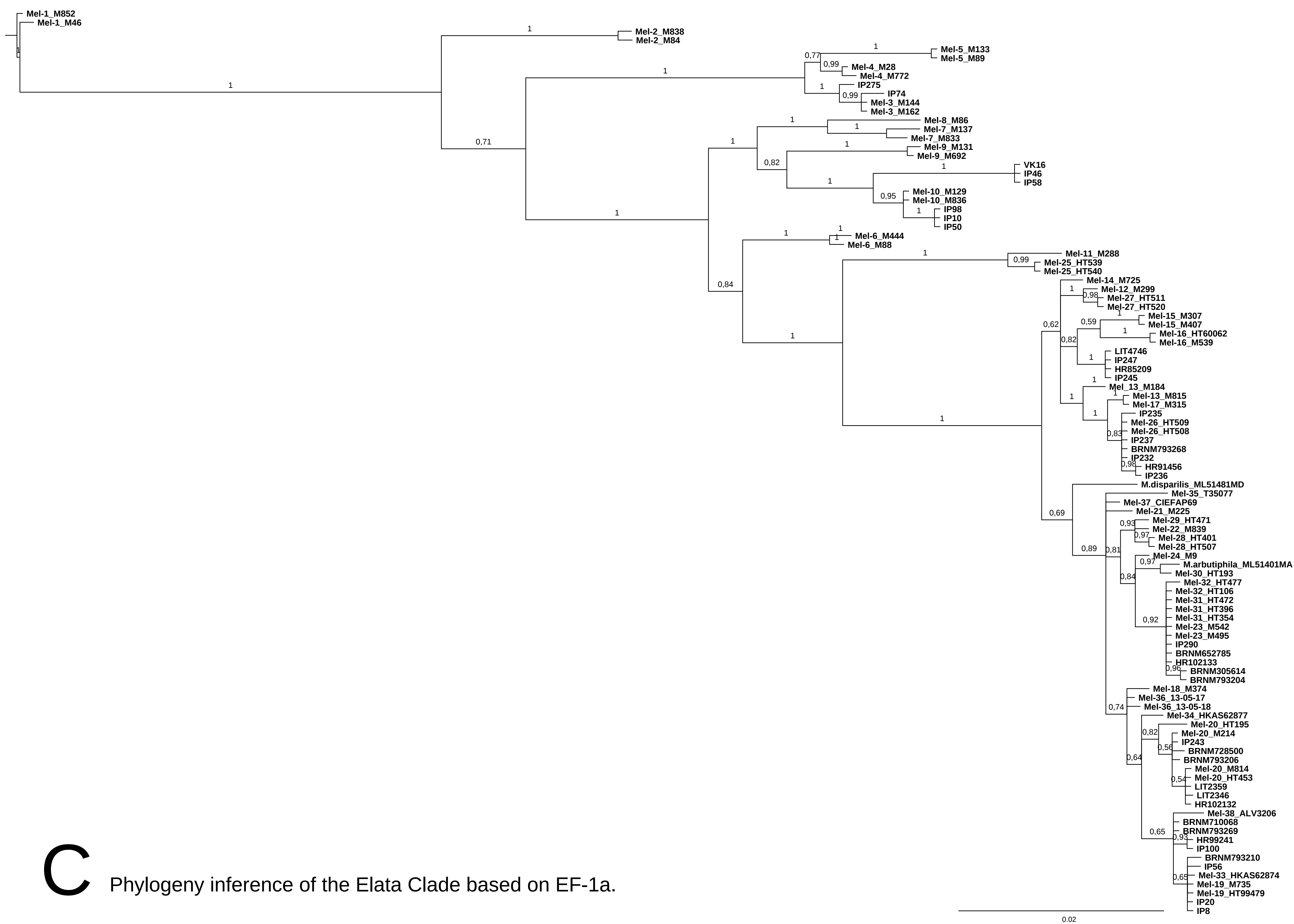


Petrželová and Sochor: How useful is the current species recognition concept for determination of true morels? Insights from the Czech Republic

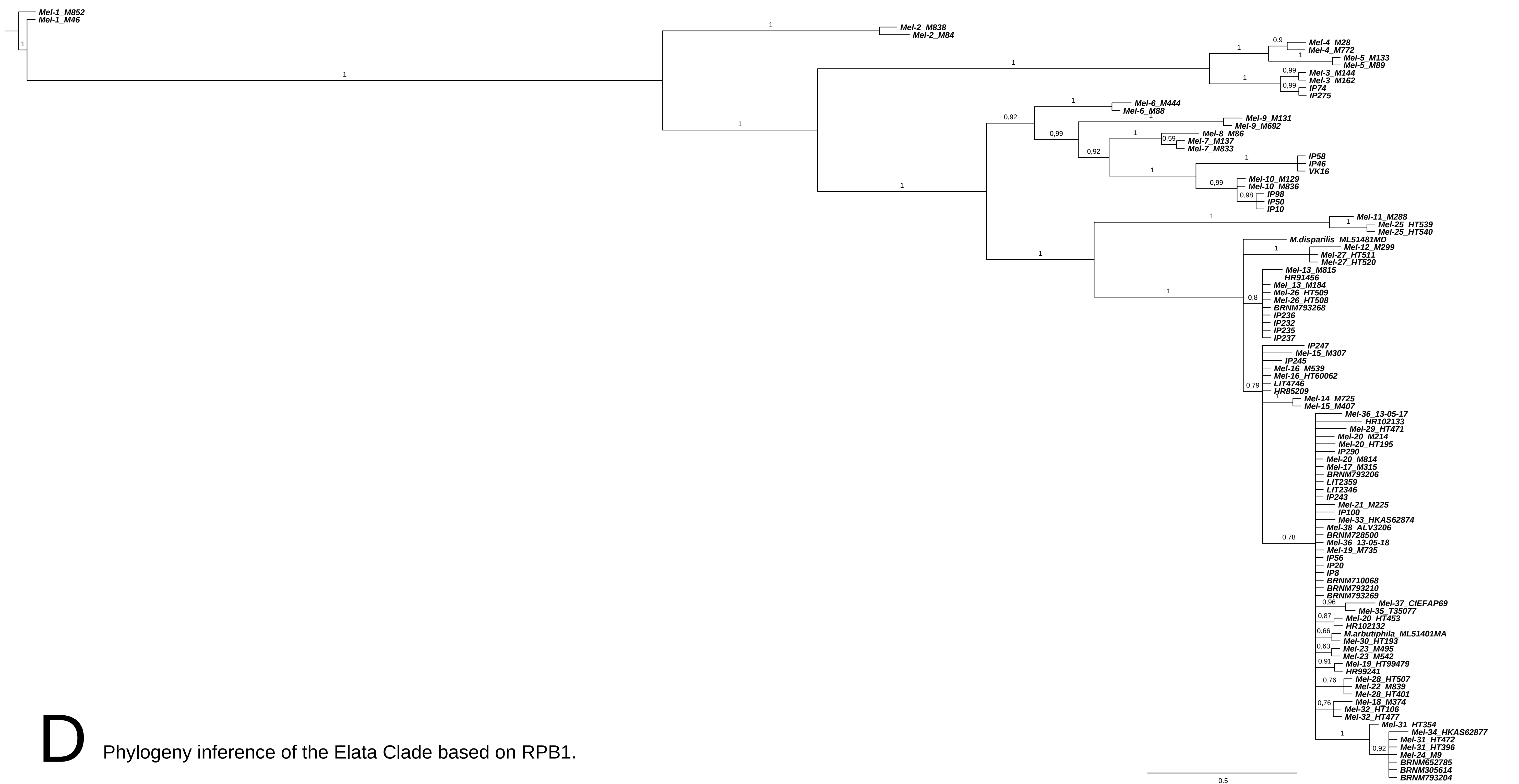
SUPPLEMENTARY FIGURES 1: Single-gene bayesian trees; posterior probabilities shown above branches. Content of this file:

- A – ITS tree of the Elata Clade (entire sample set) used for a single-gene identification of accessions.
- B – ITS tree of the Esculenta Clade (entire sample set) used for a single-gene identification of accessions.
- C – Phylogeny inference of the Elata Clade based on *EF-1 α* .
- D – Phylogeny inference of the Elata Clade based on *RPB1*.
- E – Phylogeny inference of the Elata Clade based on *RPB2*.
- F – Phylogeny inference of the Elata Clade based on LSU.
- G – Phylogeny inference of the Elata Clade based on ITS (multi-gene sample set).
- H – Phylogeny inference of the Esculenta Clade based on *EF-1 α* .
- I – Phylogeny inference of the Esculenta Clade based on *RPB1*.
- J – Phylogeny inference of the Esculenta Clade based on *RPB2*.
- K – Phylogeny inference of the Esculenta Clade based on LSU.
- L – Phylogeny inference of the Esculenta Clade based on ITS (multi-gene sample set).

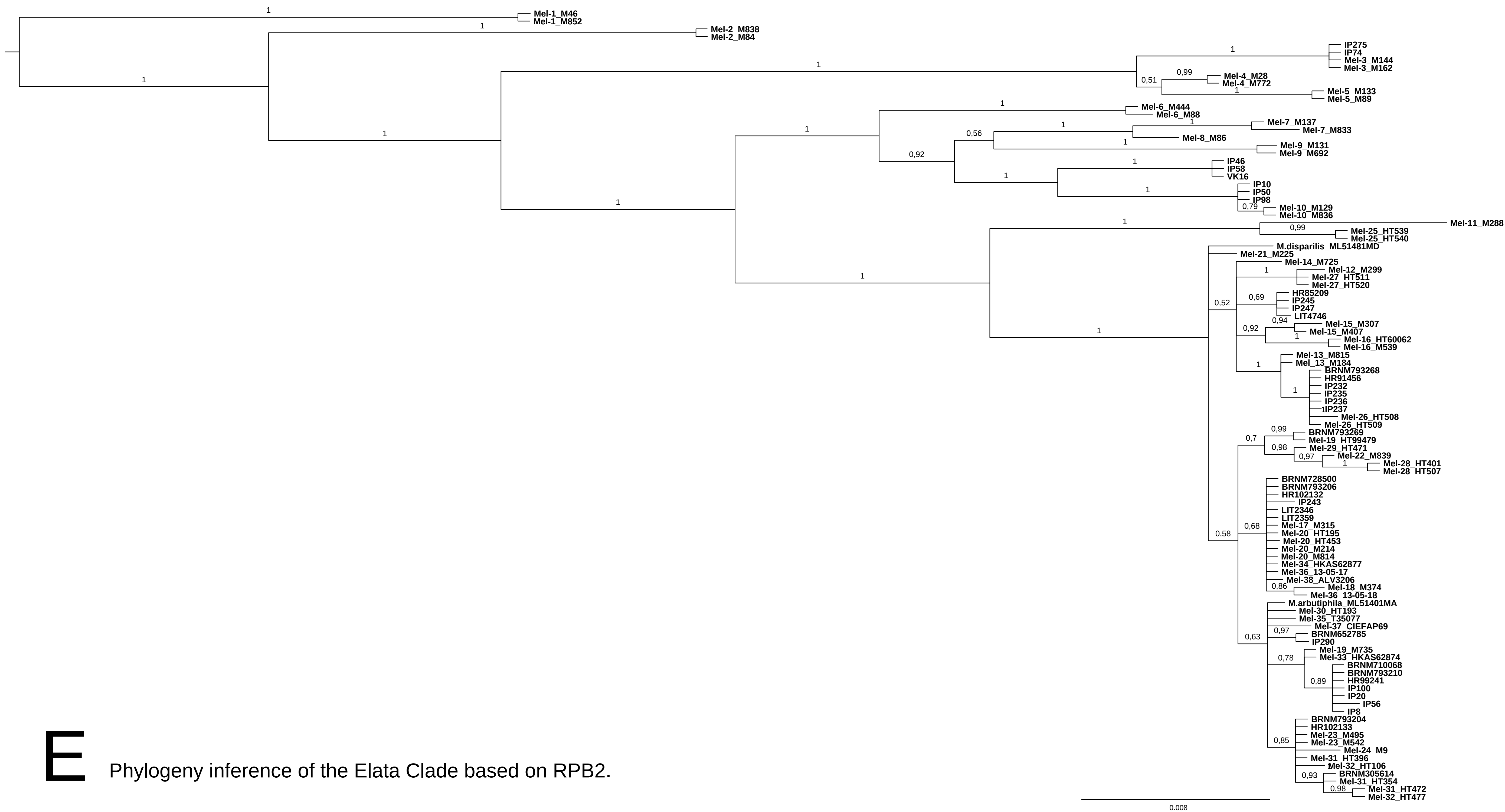


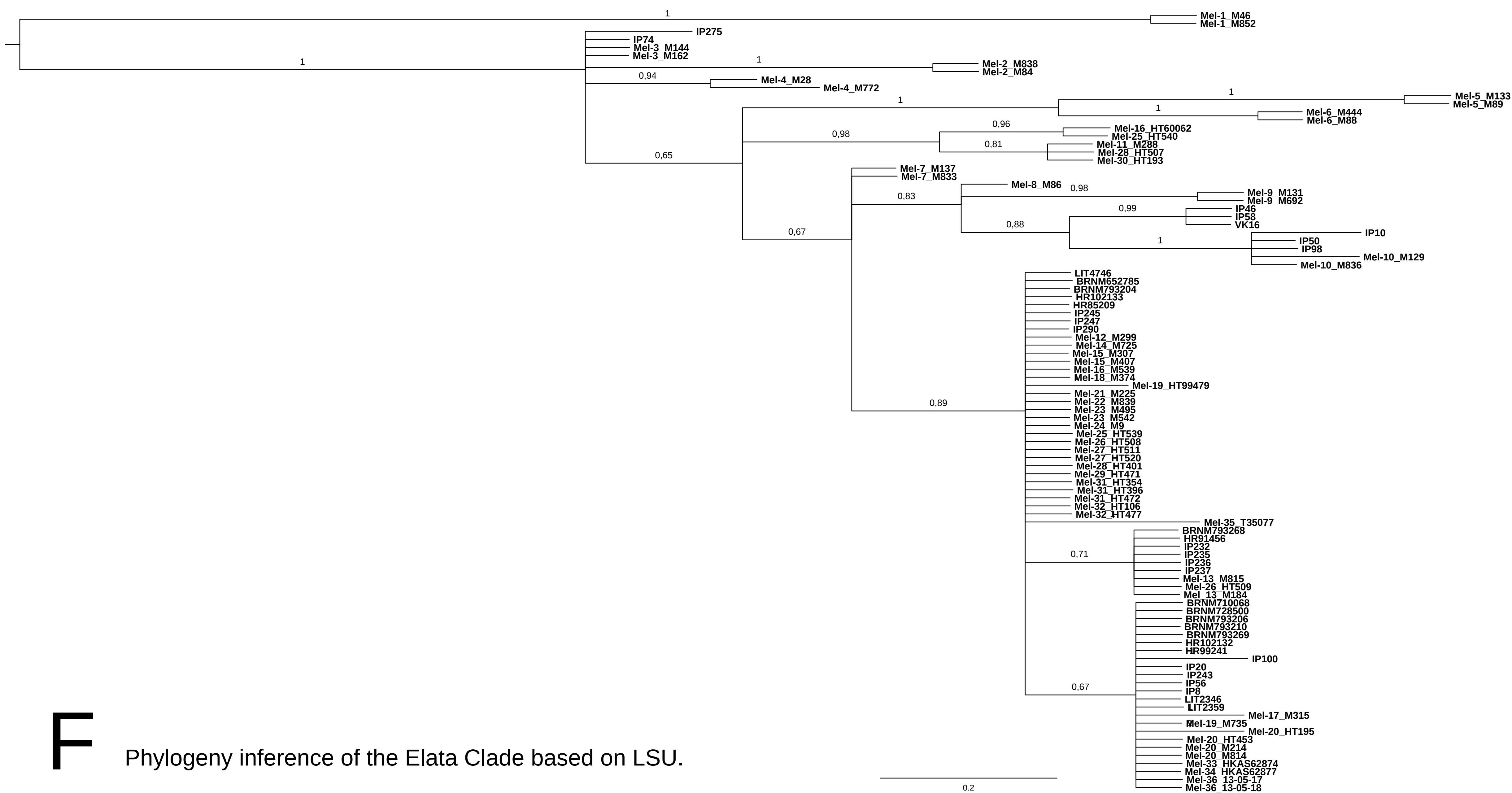


C Phylogeny inference of the Elata Clade based on EF-1a.



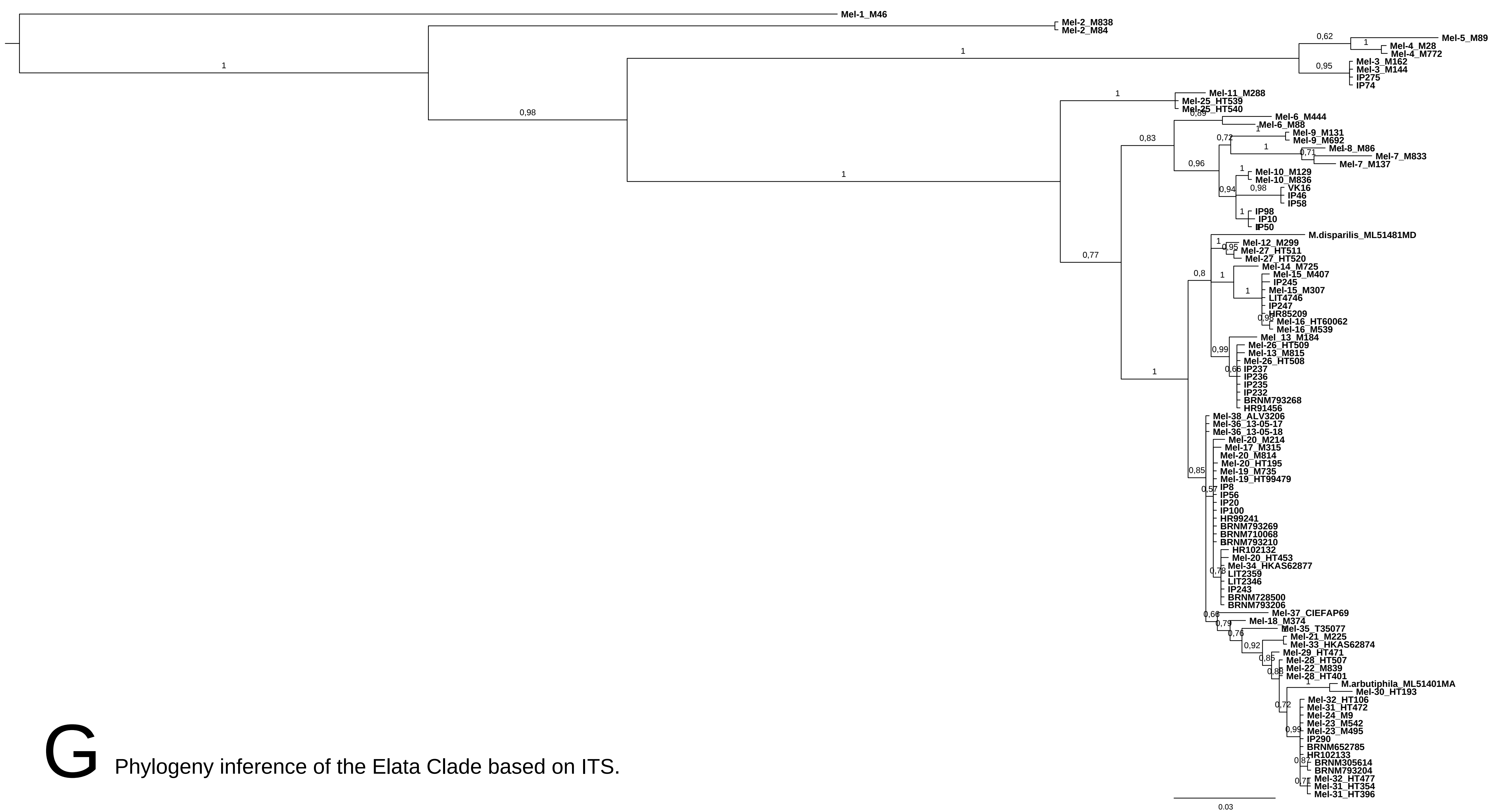
D Phylogeny inference of the Elata Clade based on RPB1.





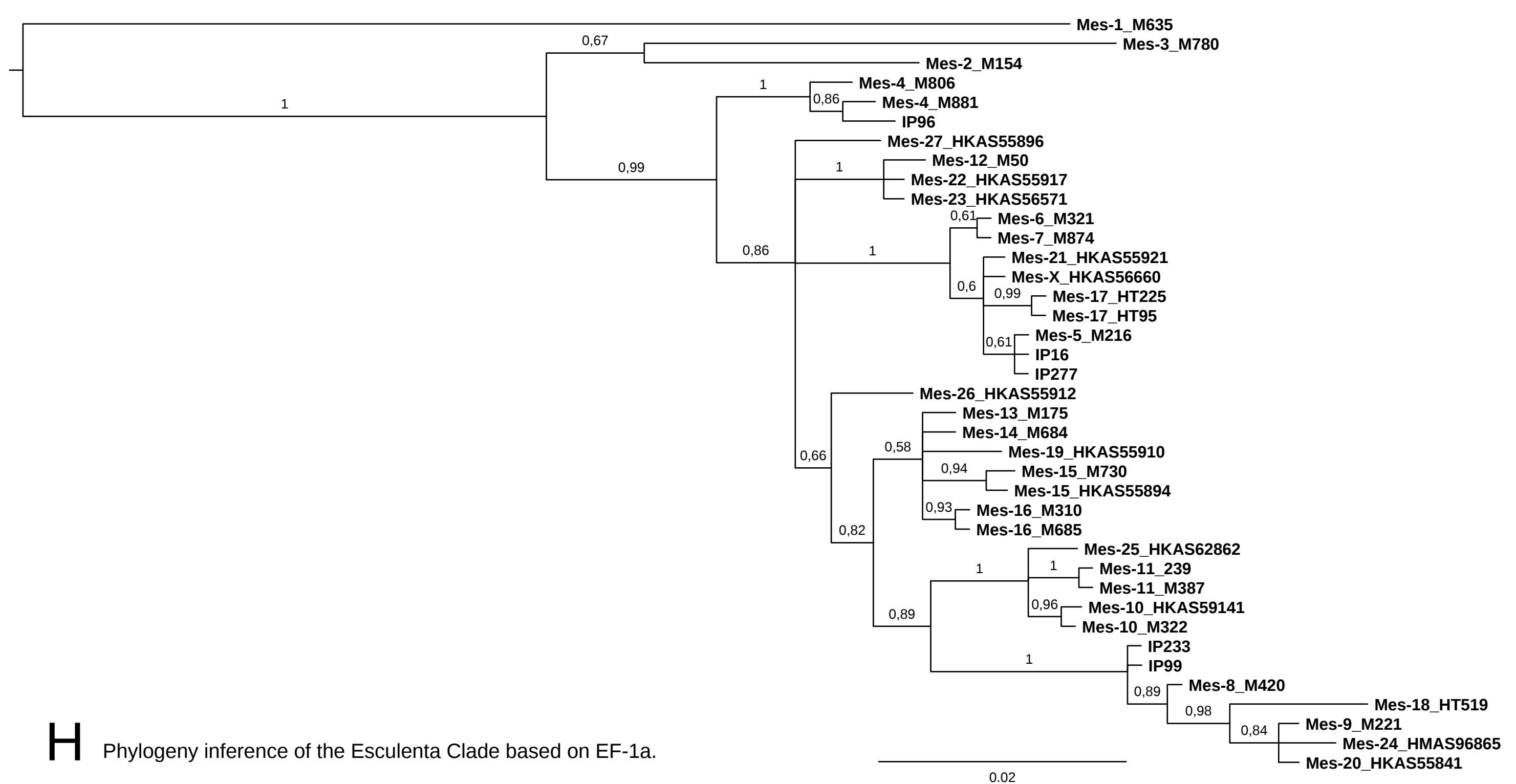
F

Phylogeny inference of the Elata Clade based on LSU.



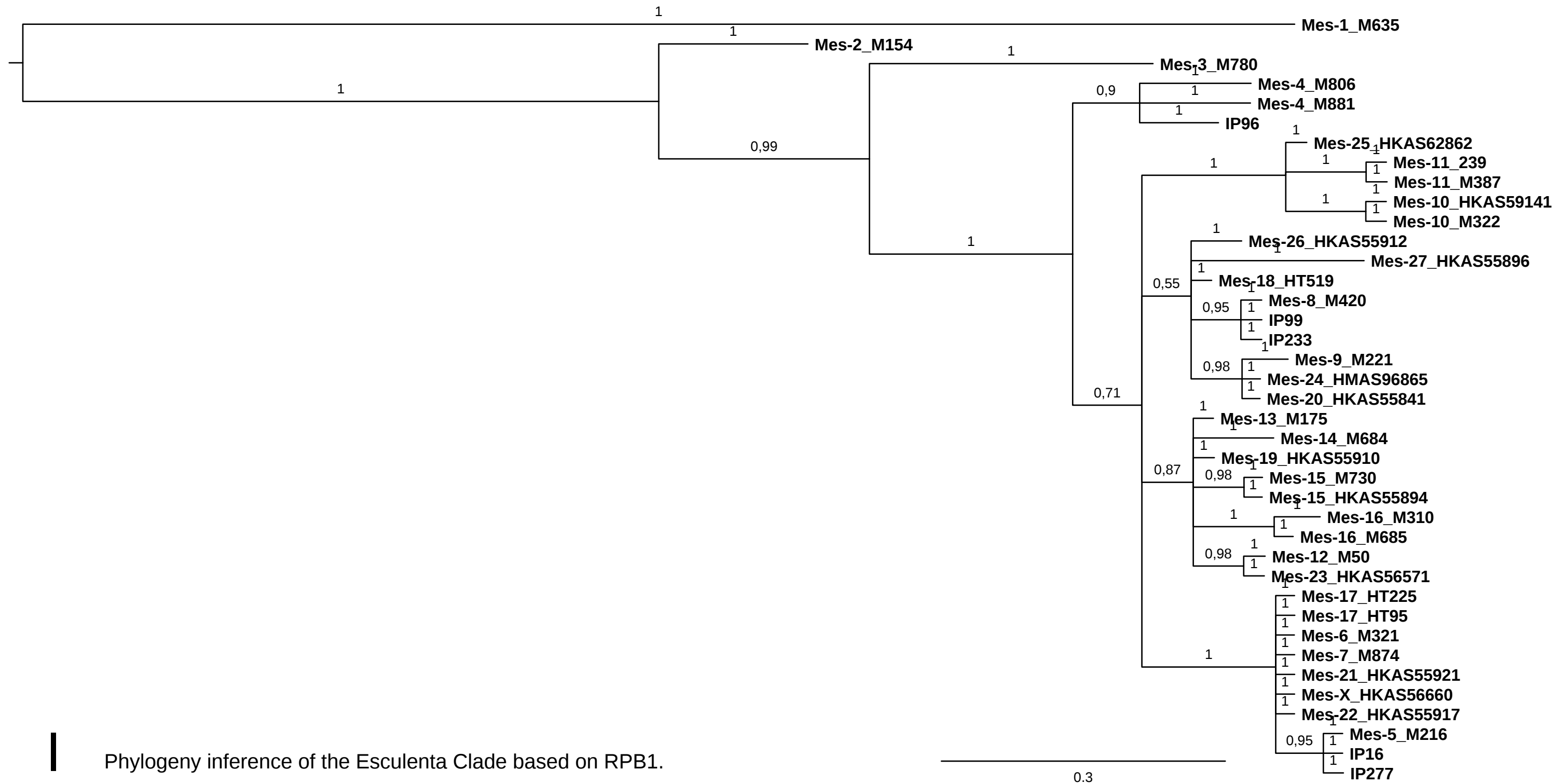
G

Phylogeny inference of the Elata Clade based on ITS.

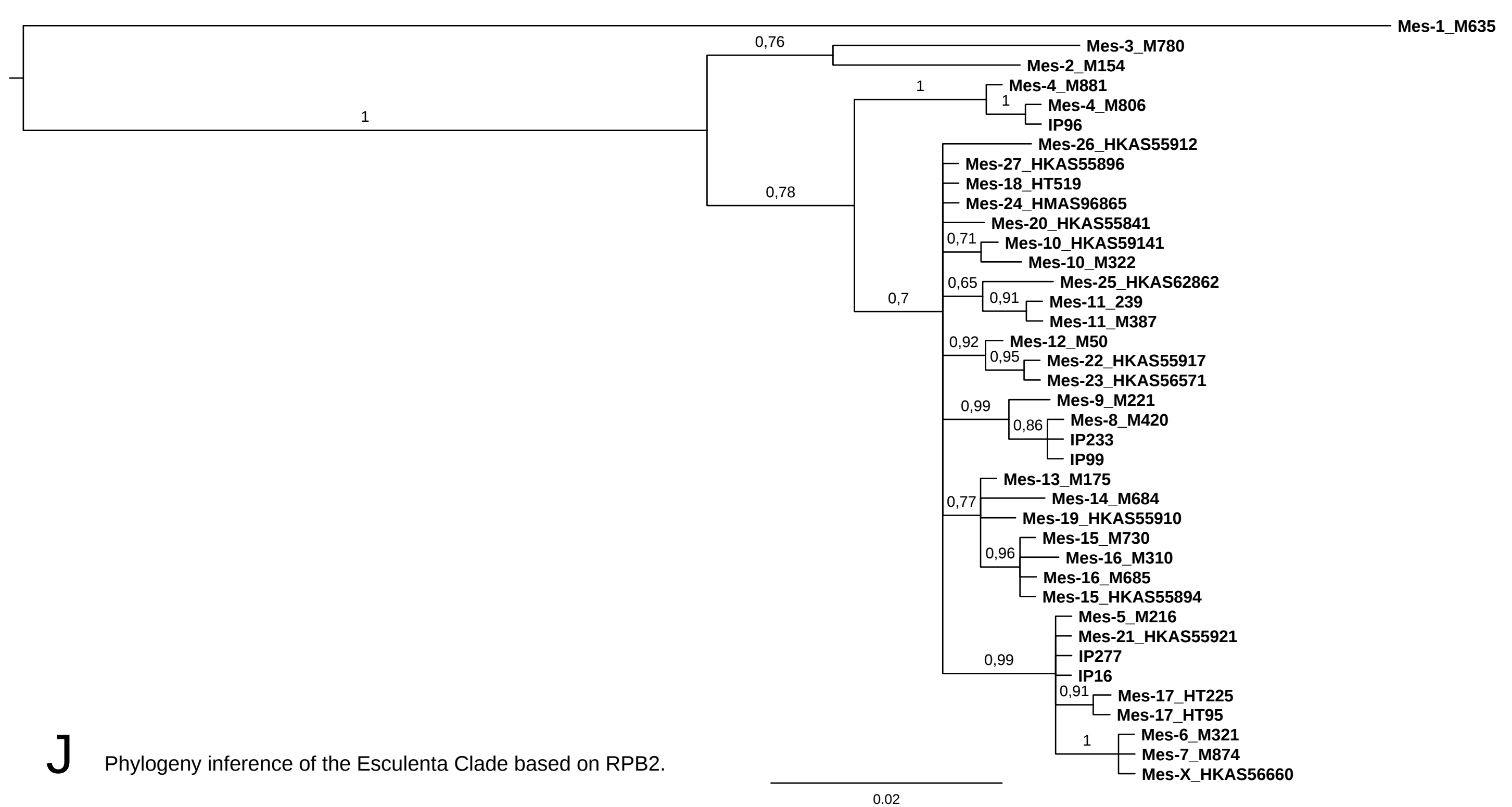


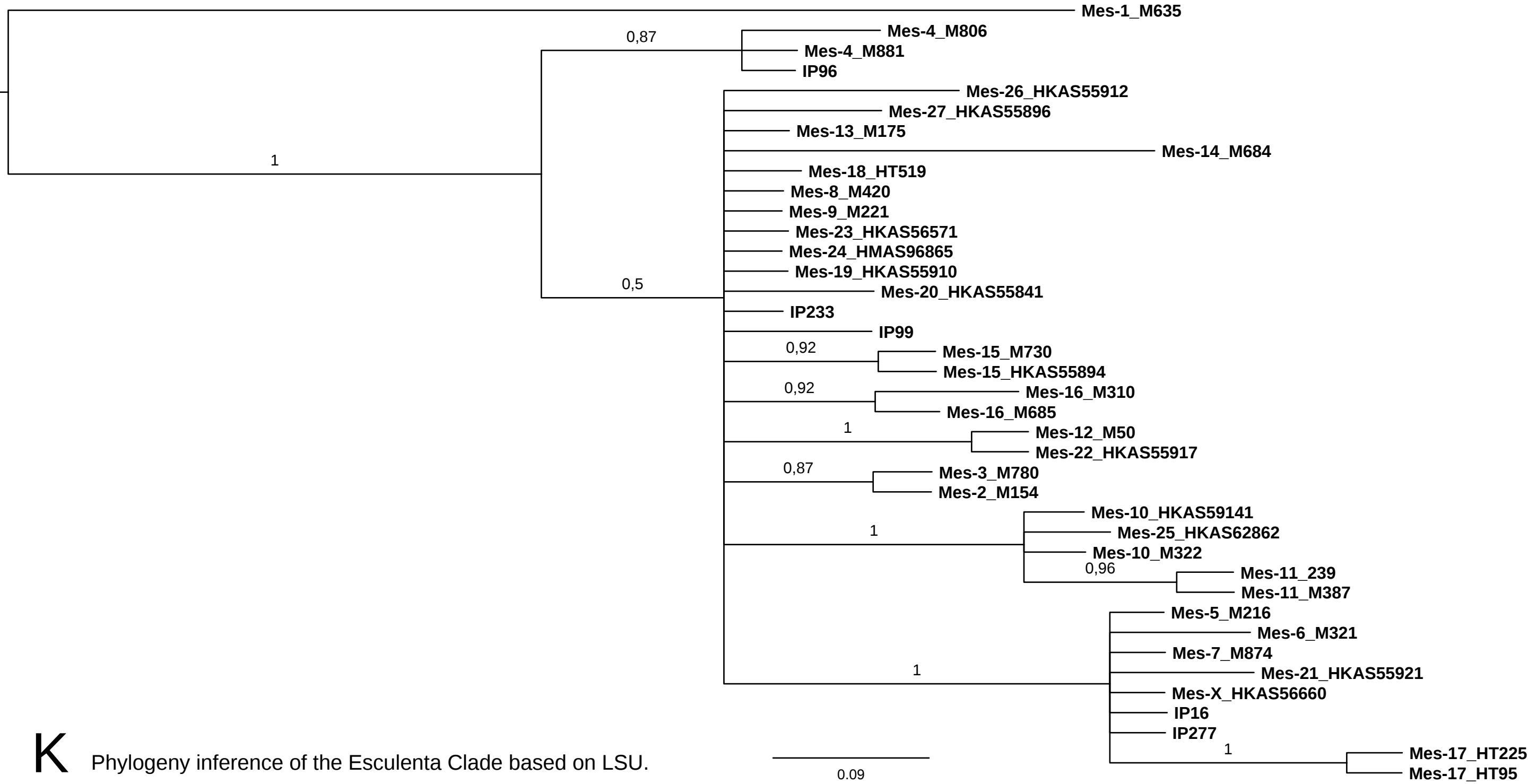
H

Phylogeny inference of the Esculenta Clade based on EF-1a.



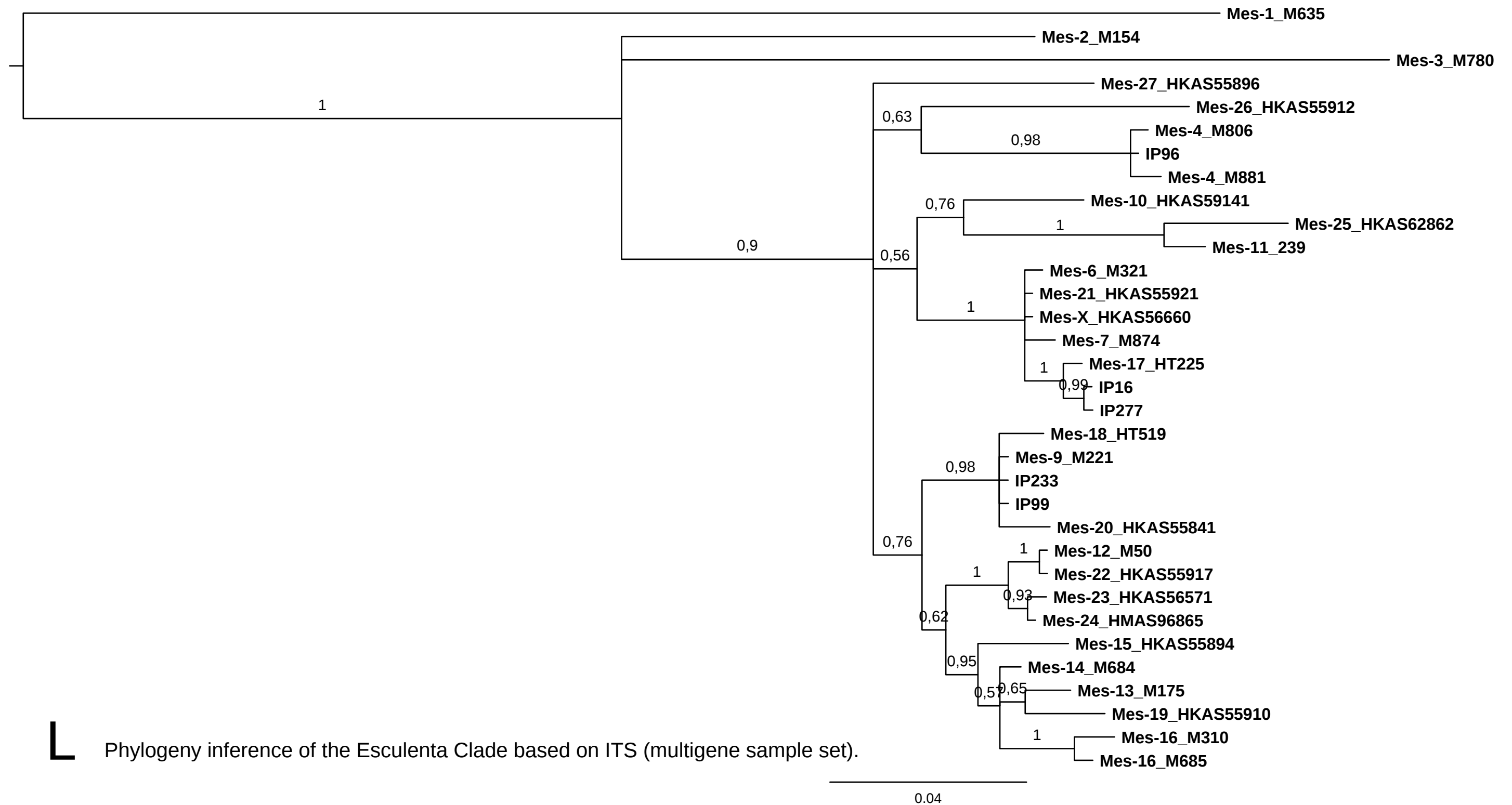
Phylogeny inference of the Esculenta Clade based on RPB1.





K

Phylogeny inference of the Esculenta Clade based on LSU.



Phylogeny inference of the Esculenta Clade based on ITS (multigene sample set).