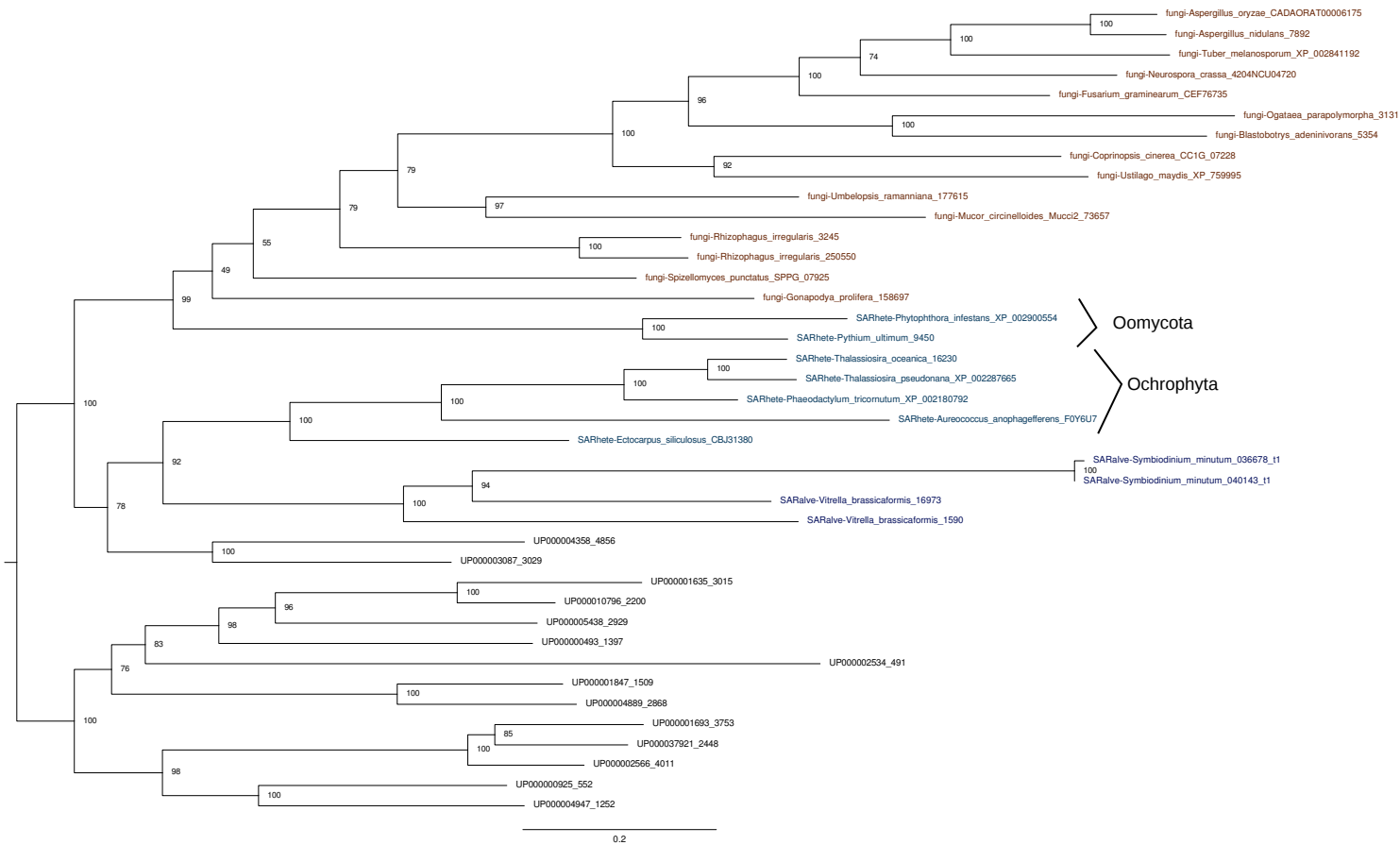


Supplementary figure S24

NAD(P)H-NIR (euks, excluding Labyrinthulea and Ichthyosporea)



Alignment statistics

Number of taxa: 40
Alignment length: 807
Parsimony info. sites: 78.70%
Missing data: 1.43%

Phylogenetic inference

Maximum likelihood
1000 UFBoot replicates
LG+R5

Taxonomy
(sequence names)

- Haptophyta**
- Rhodophyta**
- Chloroplastida**
- Rhizaria**
- Alveolata**
- Stramenopiles**
- Amoebozoa**
- Holozoa**
- Metazoa**
- Holomycota**
- Others**