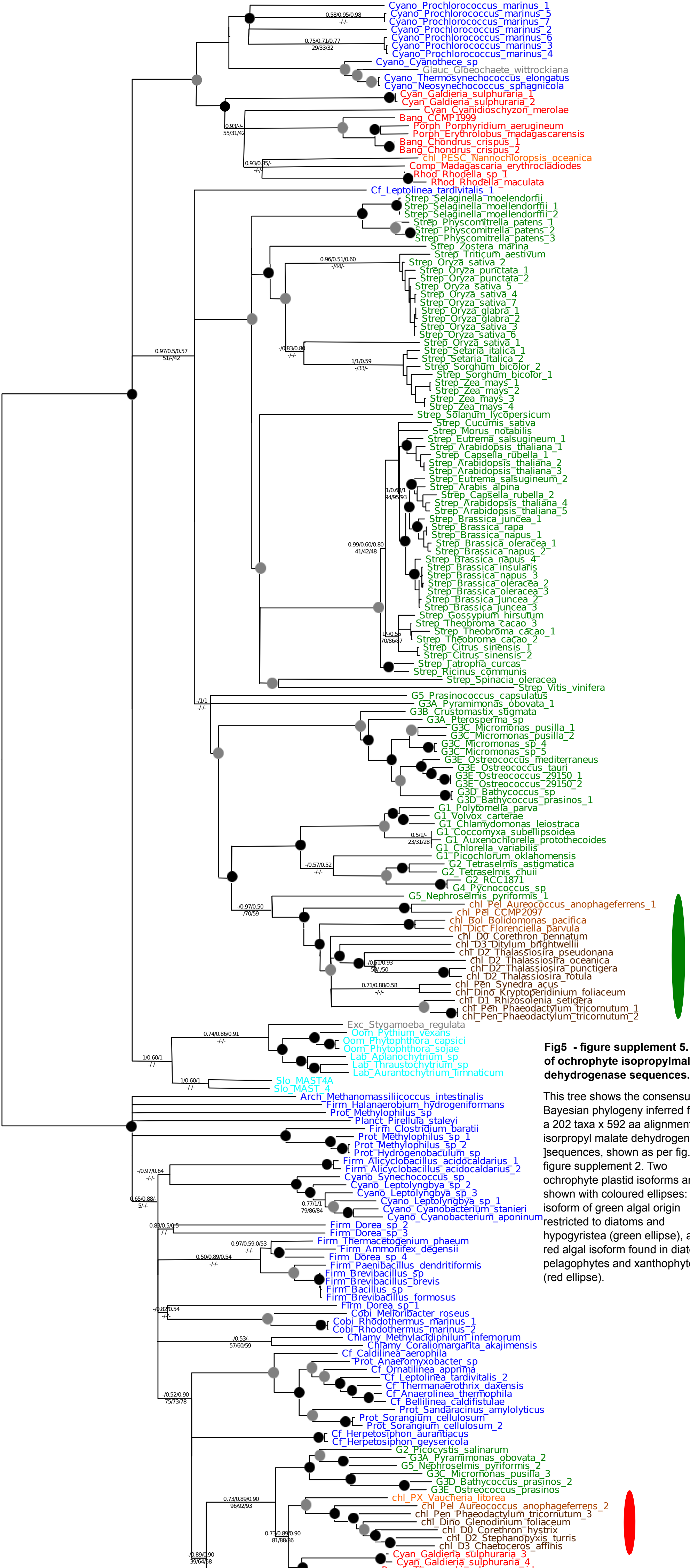

Fig5 - figure supplement 5. Tree of ochrophyte isopropylmalate dehydrogenase sequences.

This tree shows the consensus Bayesian phylogeny inferred for a 202 taxa x 592 aa alignment of isorpropyl malate dehydrogenase]sequences, shown as per fig. 2-figure supplement 2. Two ochrophyte plastid isoforms are shown with coloured ellipses: an isoform of green algal origin restricted to diatoms and hypogyristea (green ellipse), and a red algal isoform found in diatoms, pelagophytes and xanthophytes (red ellipse).