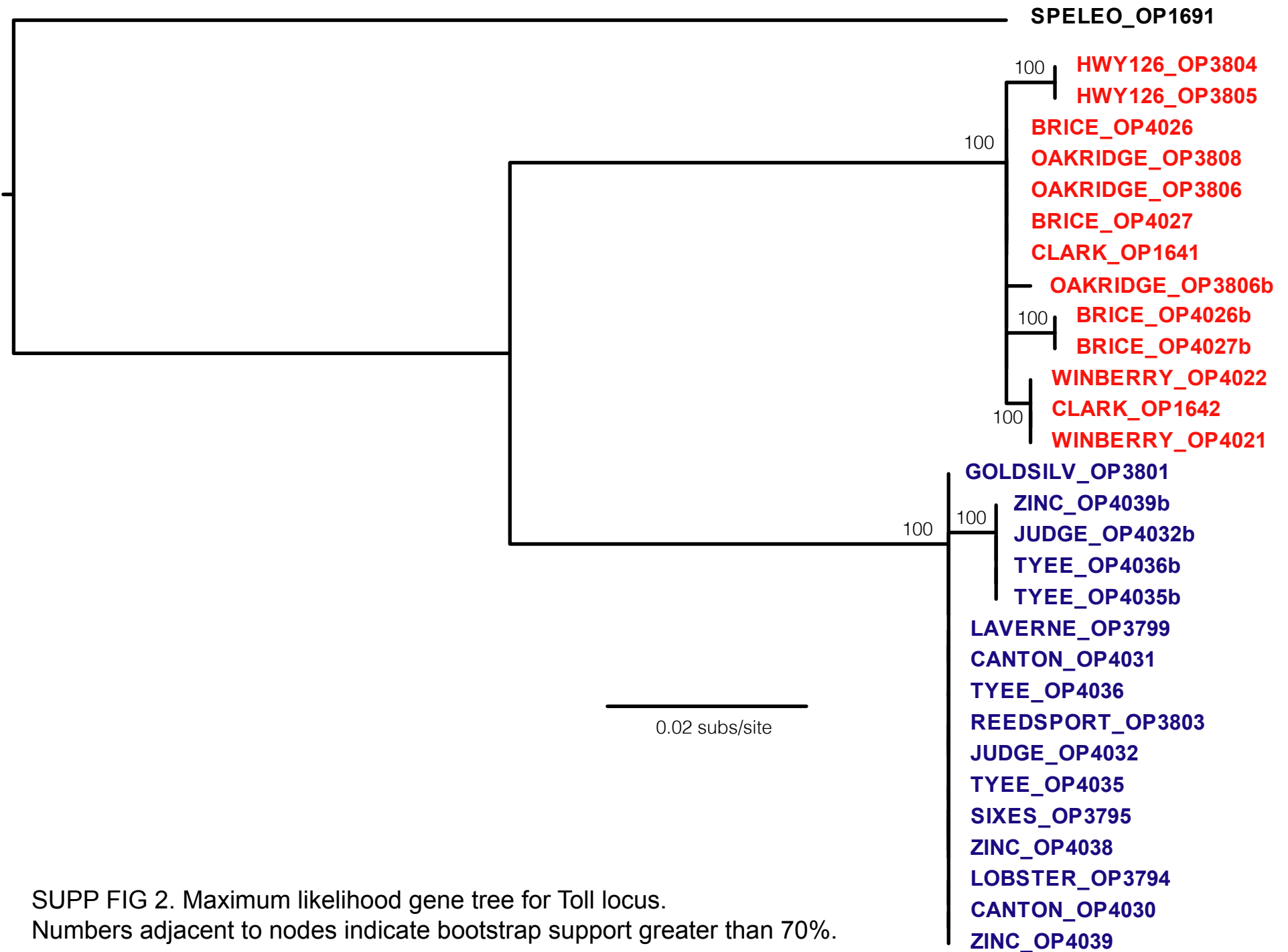
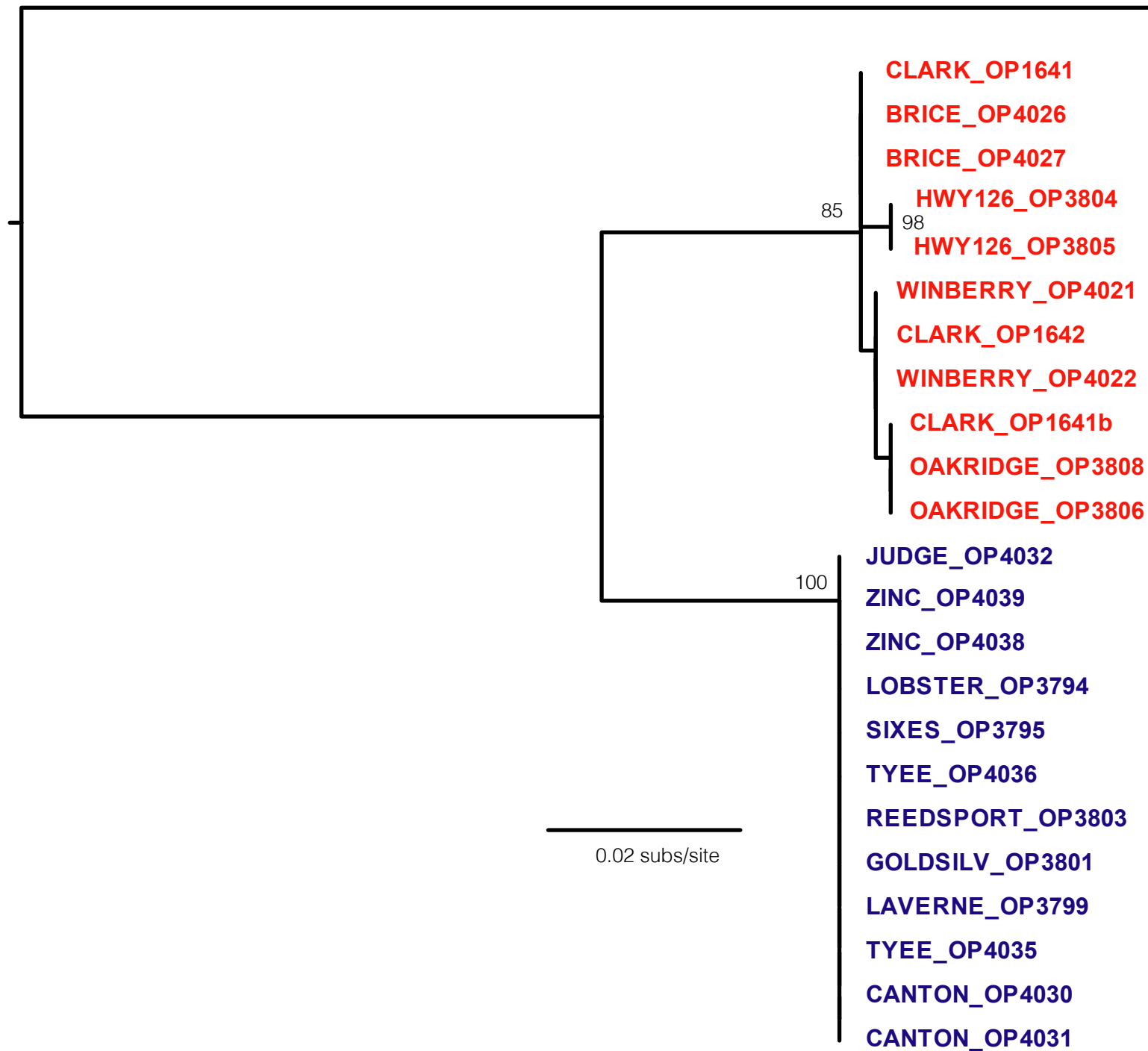


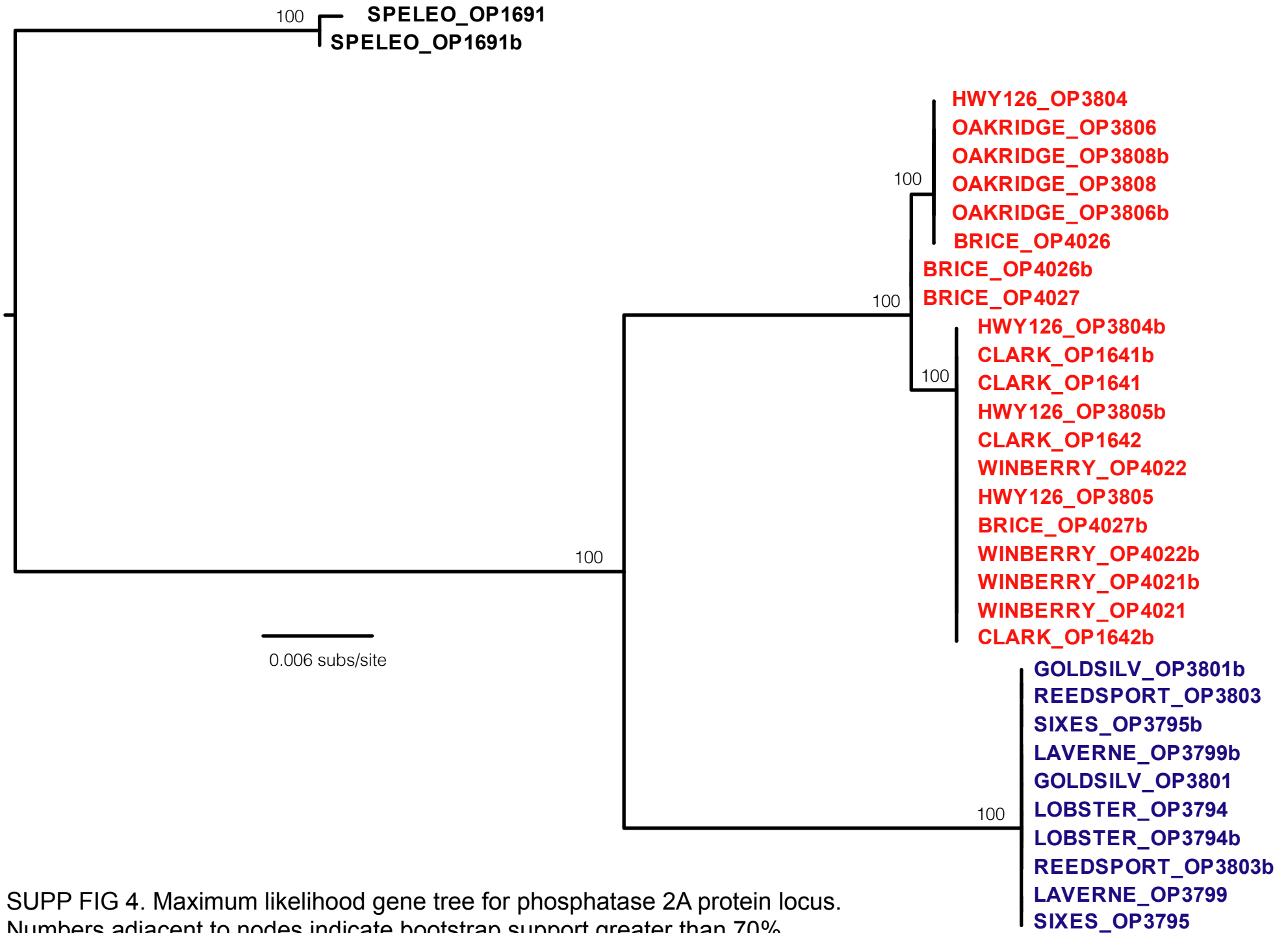
SUPP FIG 1. Maximum likelihood gene tree for cytochrome c oxidase 1 (COI) locus. Numbers adjacent to nodes indicate bootstrap support greater than 70%.



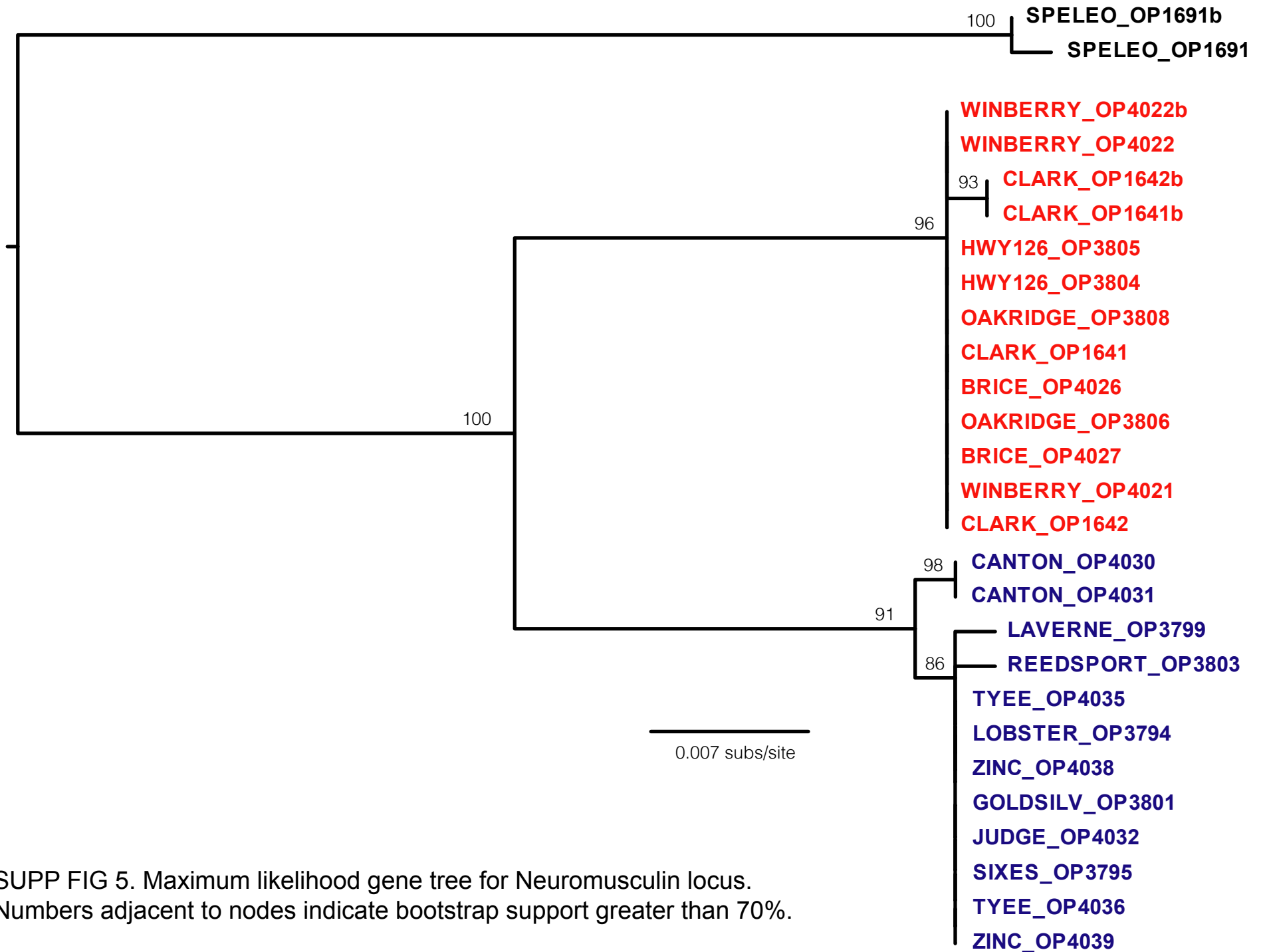
SUPP FIG 2. Maximum likelihood gene tree for Toll locus.
 Numbers adjacent to nodes indicate bootstrap support greater than 70%.



SUPP FIG 3. Maximum likelihood gene tree for F-box/LRR-repeat protein locus. Numbers adjacent to nodes indicate bootstrap support greater than 70%.



SUPP FIG 4. Maximum likelihood gene tree for phosphatase 2A protein locus. Numbers adjacent to nodes indicate bootstrap support greater than 70%.



SUPP FIG 5. Maximum likelihood gene tree for Neuromusculin locus.
 Numbers adjacent to nodes indicate bootstrap support greater than 70%.