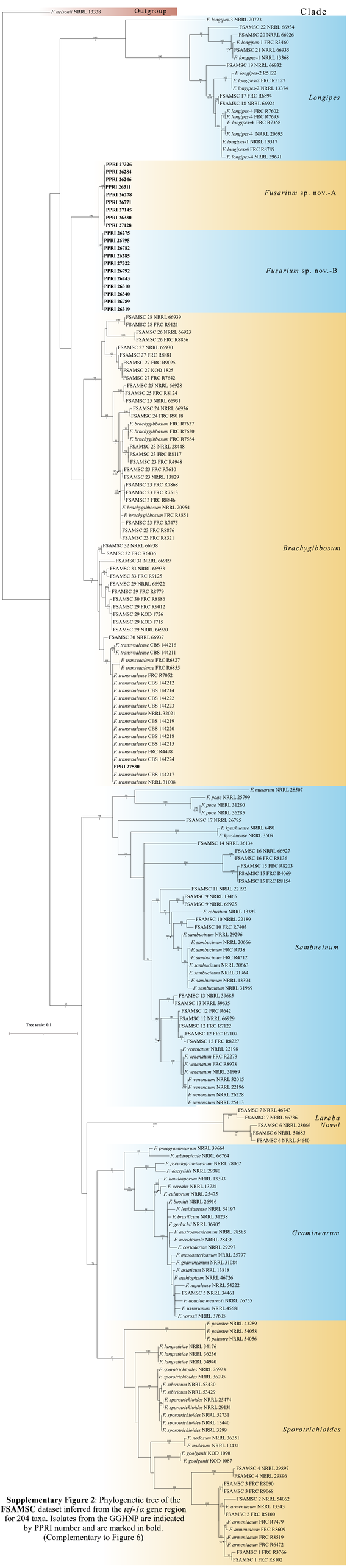

Supplementary Figure 2: Phylogenetic tree of the **FSAMSC** dataset inferred from the *tef-1α* gene region for 204 taxa. Isolates from the GGHNP are indicated by PPRI number and are marked in bold. (Complementary to Figure 6)