

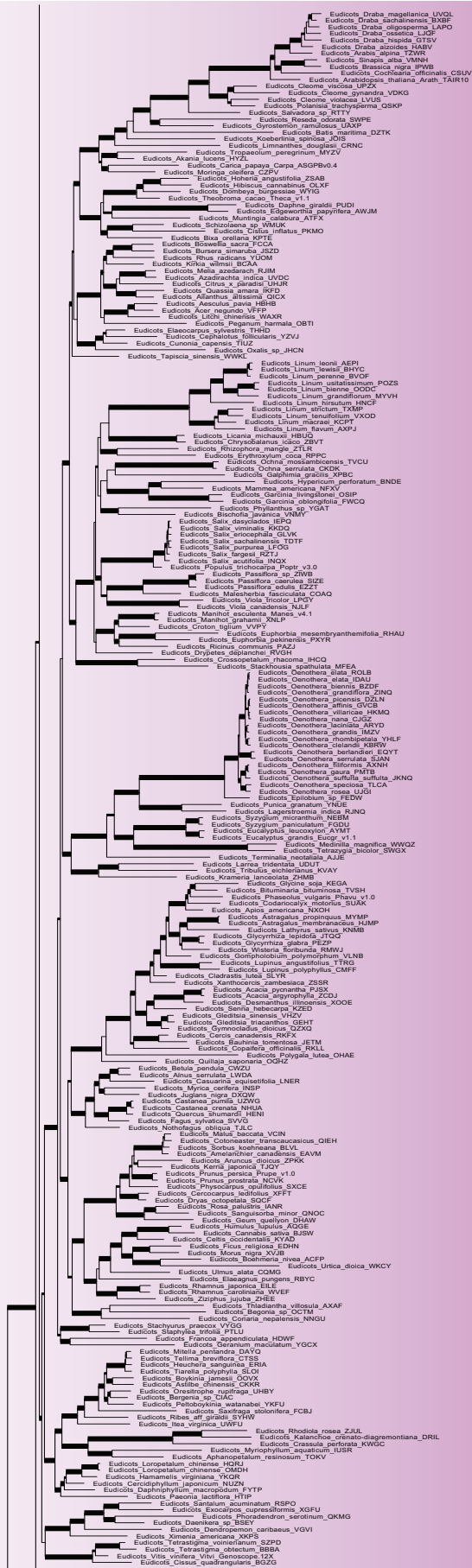
APPENDIX S5. Phylogram from a maximum likelihood analysis of a supermatrix made by concatenating 1KP transcriptome sequences from the gene regions covered by the GoFlag 451 probe set. Due to the extremely long branch leading to the closest outgroups, the tree was rooted between the bryophytes and vascular plants.

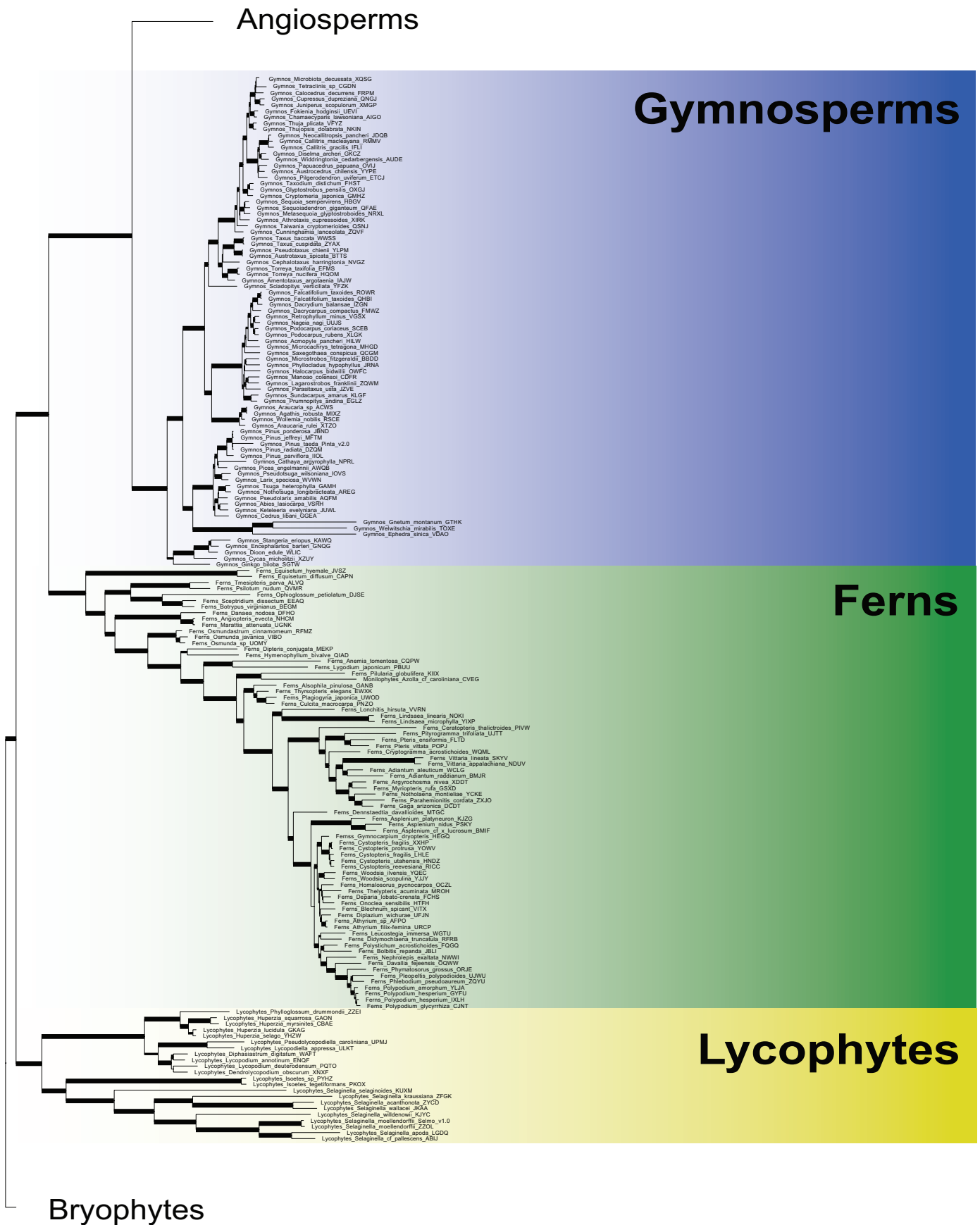
Bootstrap

— x = 100
— 70 < x < 100
— x ≤ 70



Angiosperms





Vascular plants

