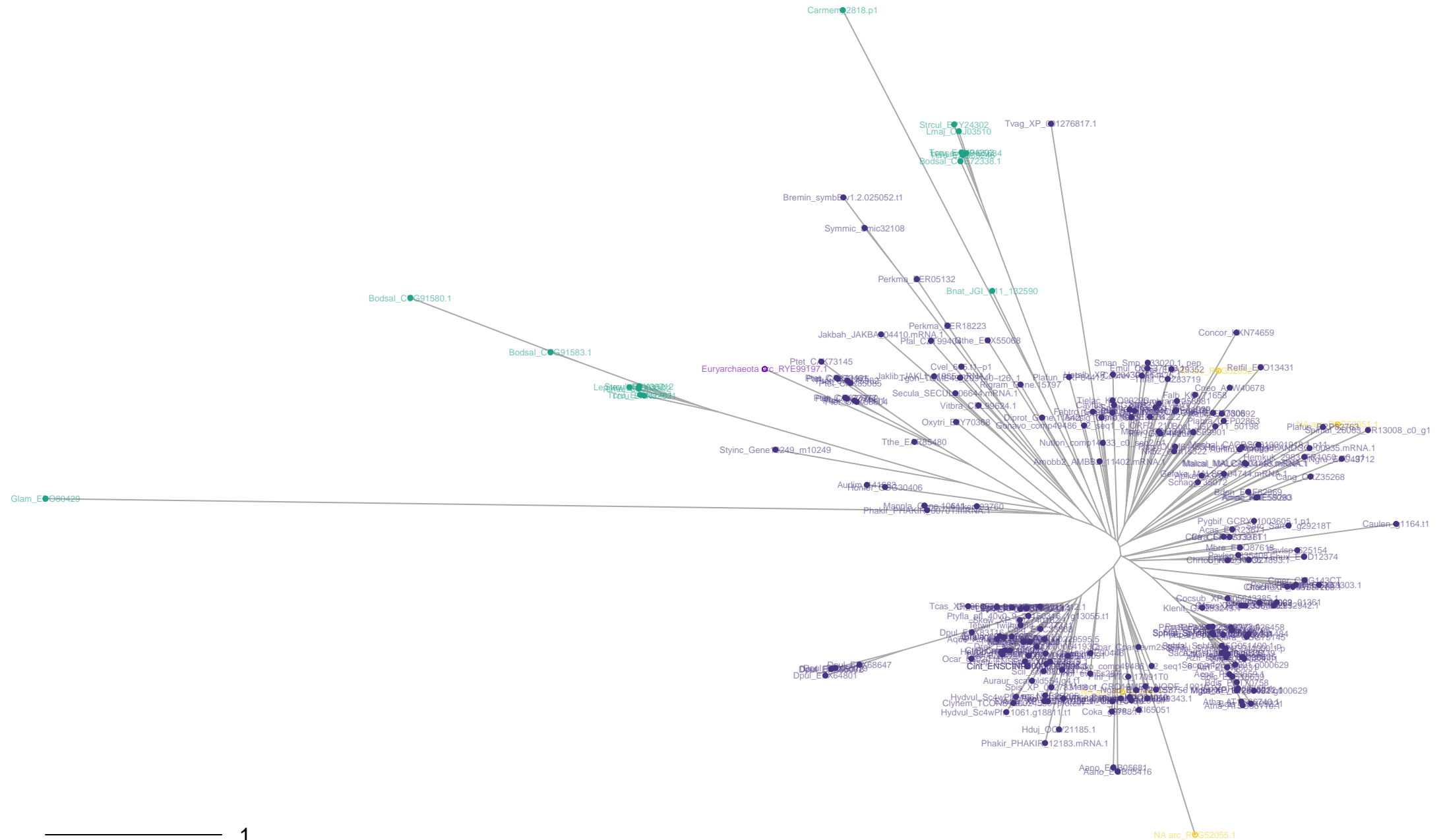


euk.ASF1_hist_chap.phy.HG1.seqs.iqtree.treefile
n=233 sequences

- eukaryotes
- archaea
- bacteria



- ASF1_hist_chap.HG1.1:ASF1A/ASF1B n = 210
- ASF1_hist_chap.HG1.0:like:ASF1A/ASF1B:likeclu:1 n = 16
- other n = 0

- Euryarchaeota n = 1
- NA arc n = 4

