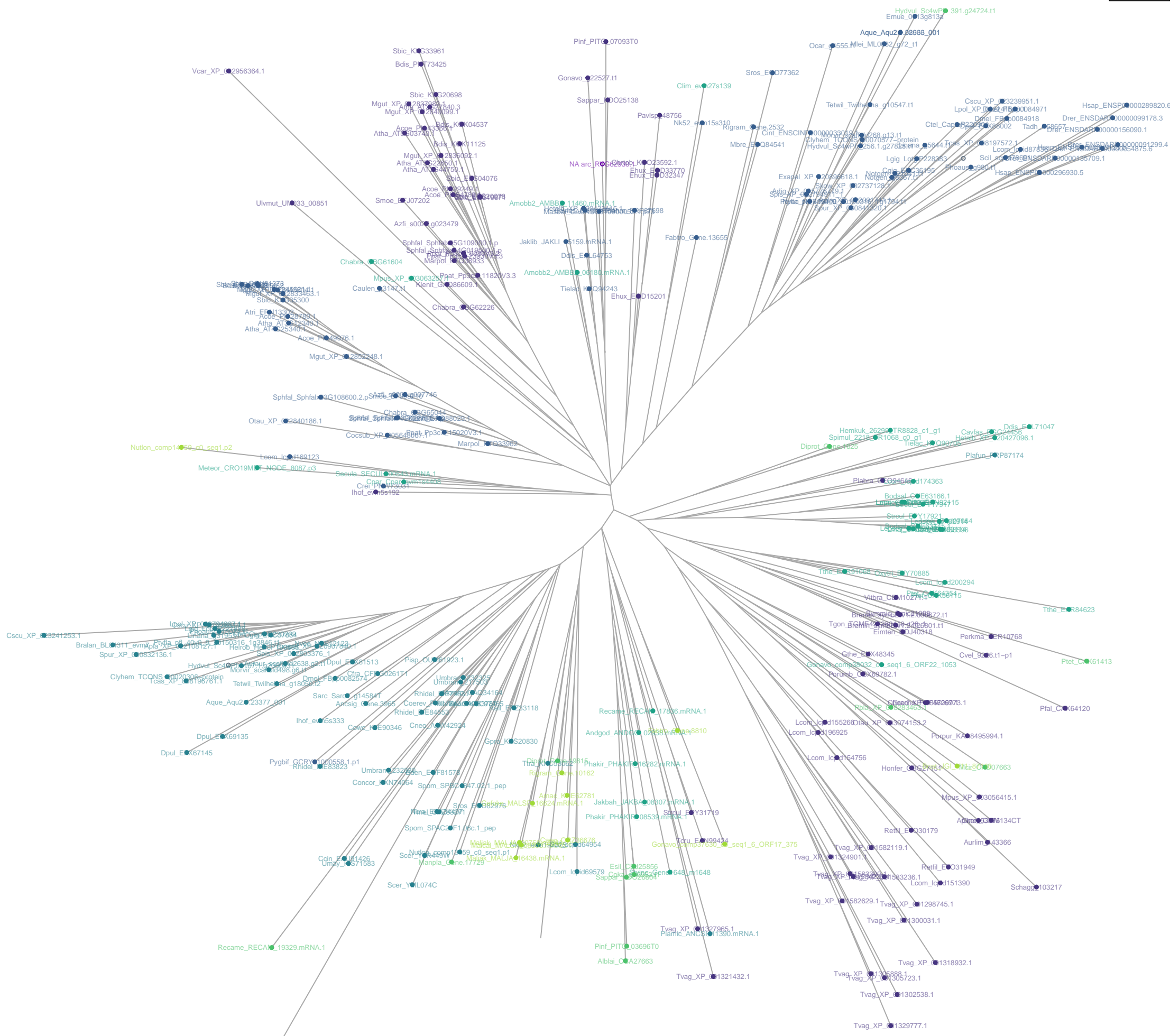


euk.Nucleoplasmin.phy.HG1.seqs.iqtree.treefile
n=302 sequences

eukaryotes
archaea
bacteria



● Nucleoplasmin.HG1.4:like:NPM1/NPM2/NPM3:likeclu:5 n = 84
● Nucleoplasmin.HG1.5:NPM1/NPM2/NPM3 n = 83
● Nucleoplasmin.HG1.3:like:NPM1/NPM2/NPM3:likeclu:5 n = 64
● Nucleoplasmin.HG1.1:like:NPM1/NPM2/NPM3:likeclu:5 n = 41
● Nucleoplasmin.HG1.0:like:NPM1/NPM2/NPM3:likeclu:5 n = 13
● Nucleoplasmin.HG1.2:like:NPM1/NPM2/NPM3:likeclu:5 n = 12
● other n = 0

● NA n = 1

