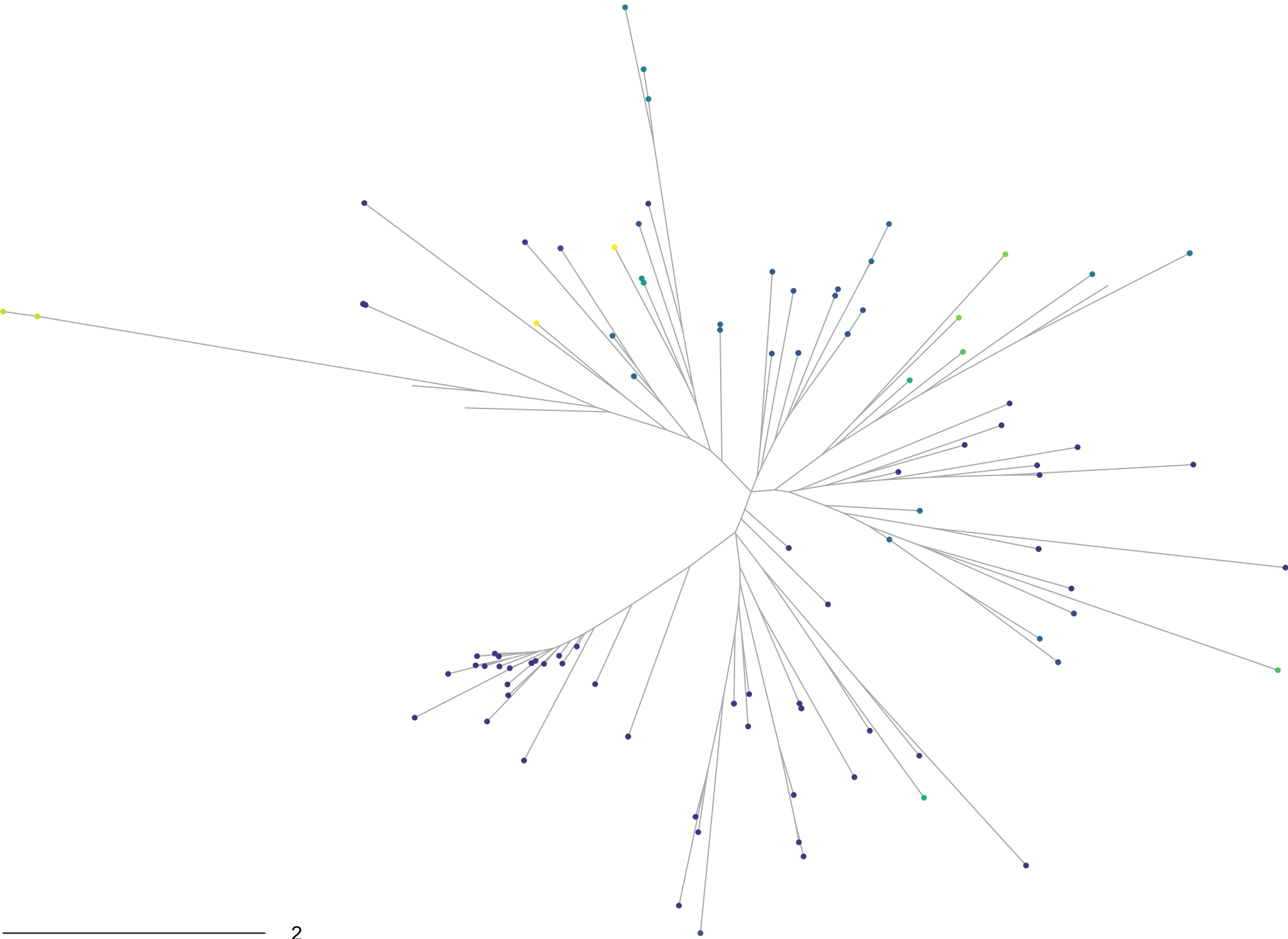


vir.zf-CXXC.phy.HG1.seqs.iqtree.treefile
n=113 sequences

eukaryotes
archaea
bacteria
viruses



2

zf-CXXC.HG1.0:CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/MBD1/TET1/TET3 n = 64
zf-CXXC.HG1.5:AP001267.5/KMT2A/KMT2B n = 18
zf-CXXC.HG1.4:like:AP001267.5/KMT2A/KMT2B:likeclu:5 n = 10
zf-CXXC.HG1.8:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 6
zf-CXXC.HG1.1:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 2
zf-CXXC.HG1.2:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 2
zf-CXXC.HG1.3:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 2
zf-CXXC.HG1.7:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 2
zf-CXXC.HG1.9:like:AP001267.5/CXXC1/CXXC4/CXXC5/DNMT1/FBXL19/KDM2A/KDM2B/KMT2A/KMT2B/MBD1/TET1/TET3:likeclu:0/5 n = 2
other n = 2

n =