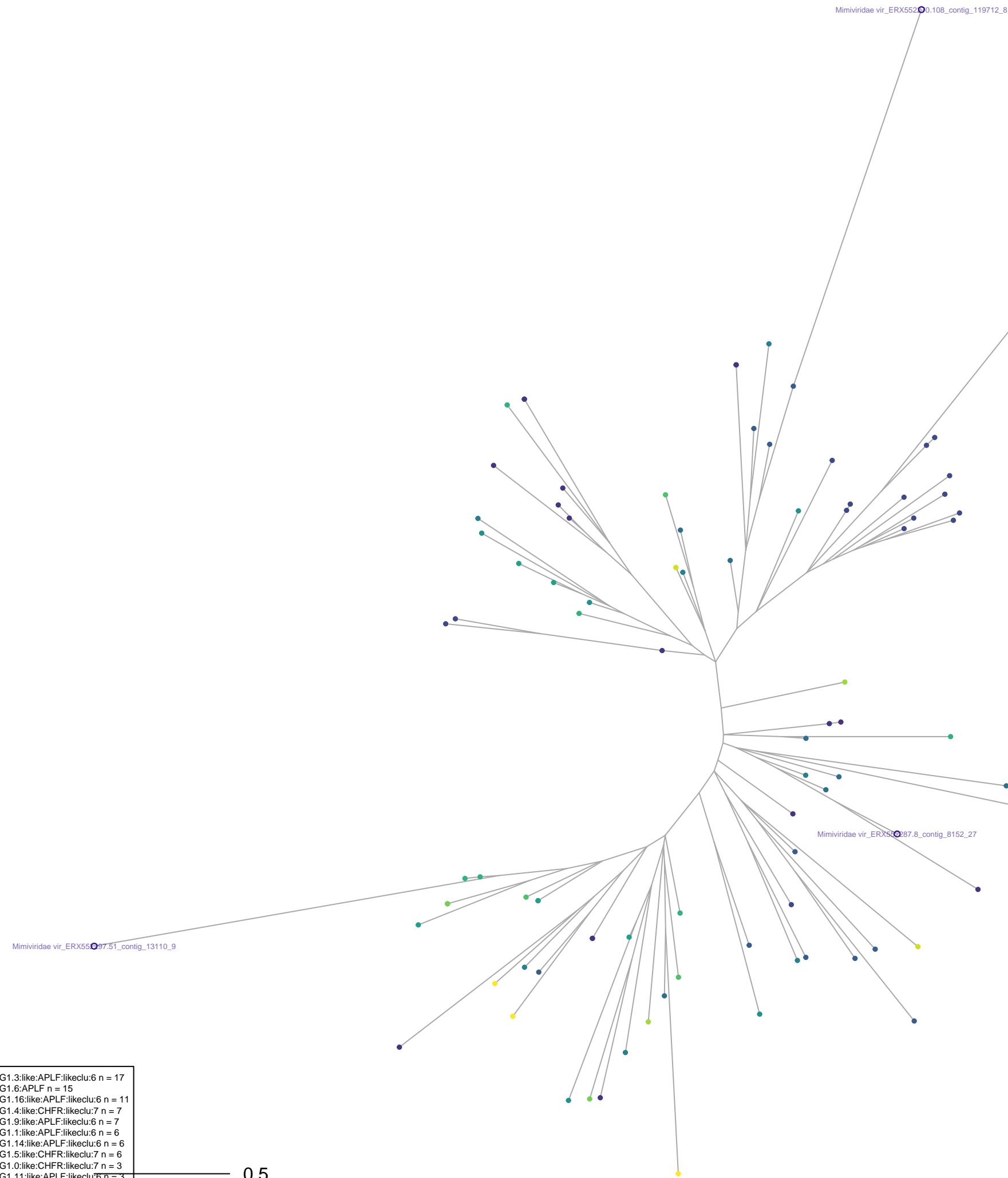


zf-CCHH
vir.zf-CCHH.phy.HG1.seqs.iqtree.treefile
n=91 sequences

- eukaryotes
- archaea
- bacteria
- viruses



- zf-CCHH.HG1.3:like:APLF:likeclu:6 n = 17
- zf-CCHH.HG1.6:APLF n = 15
- zf-CCHH.HG1.16:like:APLF:likeclu:6 n = 11
- zf-CCHH.HG1.4:like:CHFR:likeclu:7 n = 7
- zf-CCHH.HG1.9:like:APLF:likeclu:6 n = 7
- zf-CCHH.HG1.1:like:APLF:likeclu:6 n = 6
- zf-CCHH.HG1.14:like:APLF:likeclu:6 n = 6
- zf-CCHH.HG1.5:like:CHFR:likeclu:7 n = 6
- zf-CCHH.HG1.0:like:CHFR:likeclu:7 n = 3
- zf-CCHH.HG1.11:like:APLF:likeclu:6 n = 3
- zf-CCHH.HG1.10:like:APLF:likeclu:6 n = 2
- zf-CCHH.HG1.2:like:CHFR:likeclu:7 n = 2
- other n = 3

0.5

- Mimiviridae n = 3