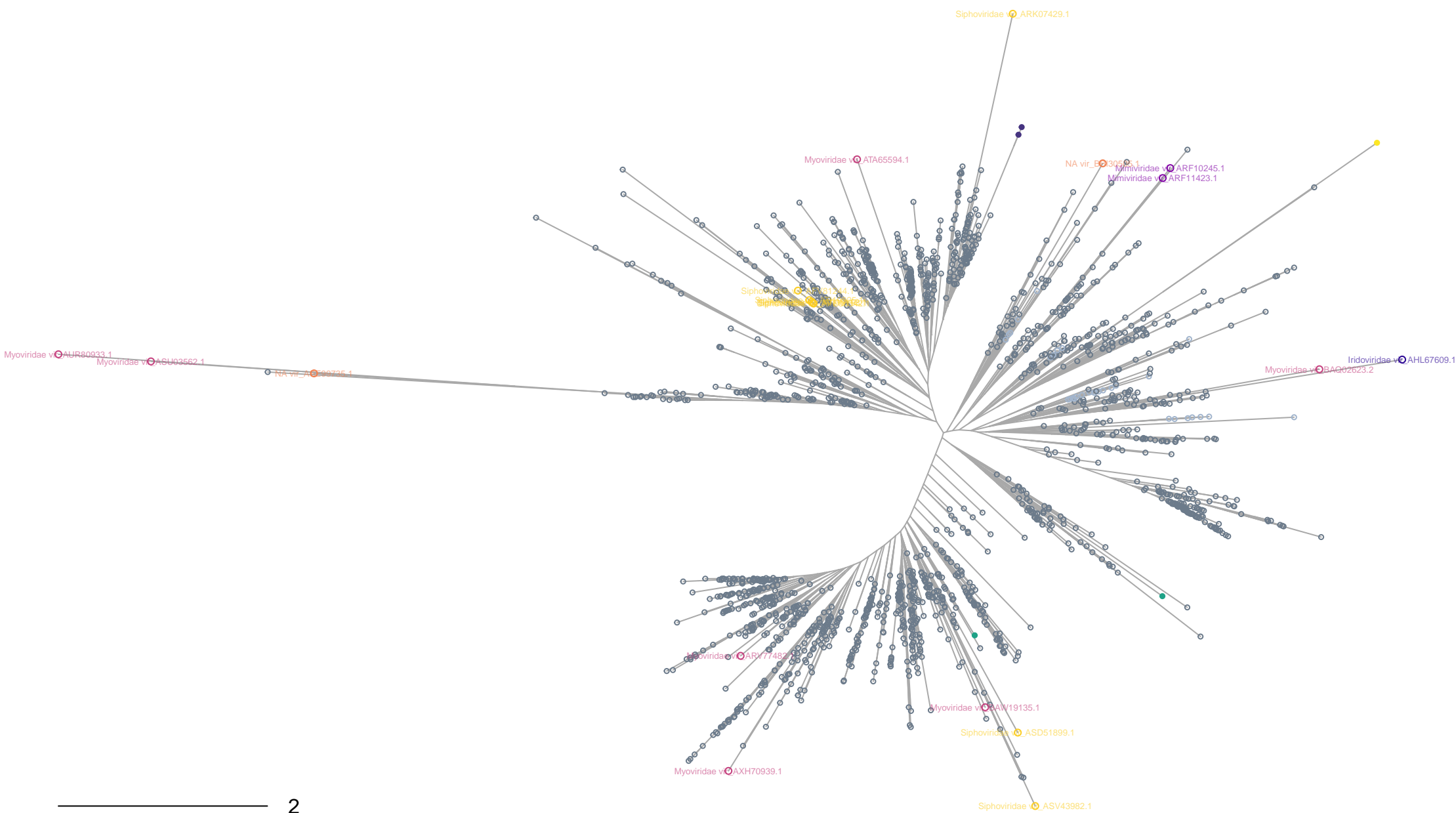


PTIP
vir.PTIP.phy.HG1.seqs.iqtree.treefile
n=1740 sequences

- eukaryotes
- archaea
- bacteria
- viruses



2

- PTIP.HG3.0:like:RFC1:iikeclu:4 n = 2
- PTIP.HG3.4:RFC1 n = 2
- other n = 1

- Iridoviridae n = 1
- Mimiviridae n = 2
- Myoviridae n = 7
- NA n = 2
- Siphoviridae n = 8

PTIP

☐

7

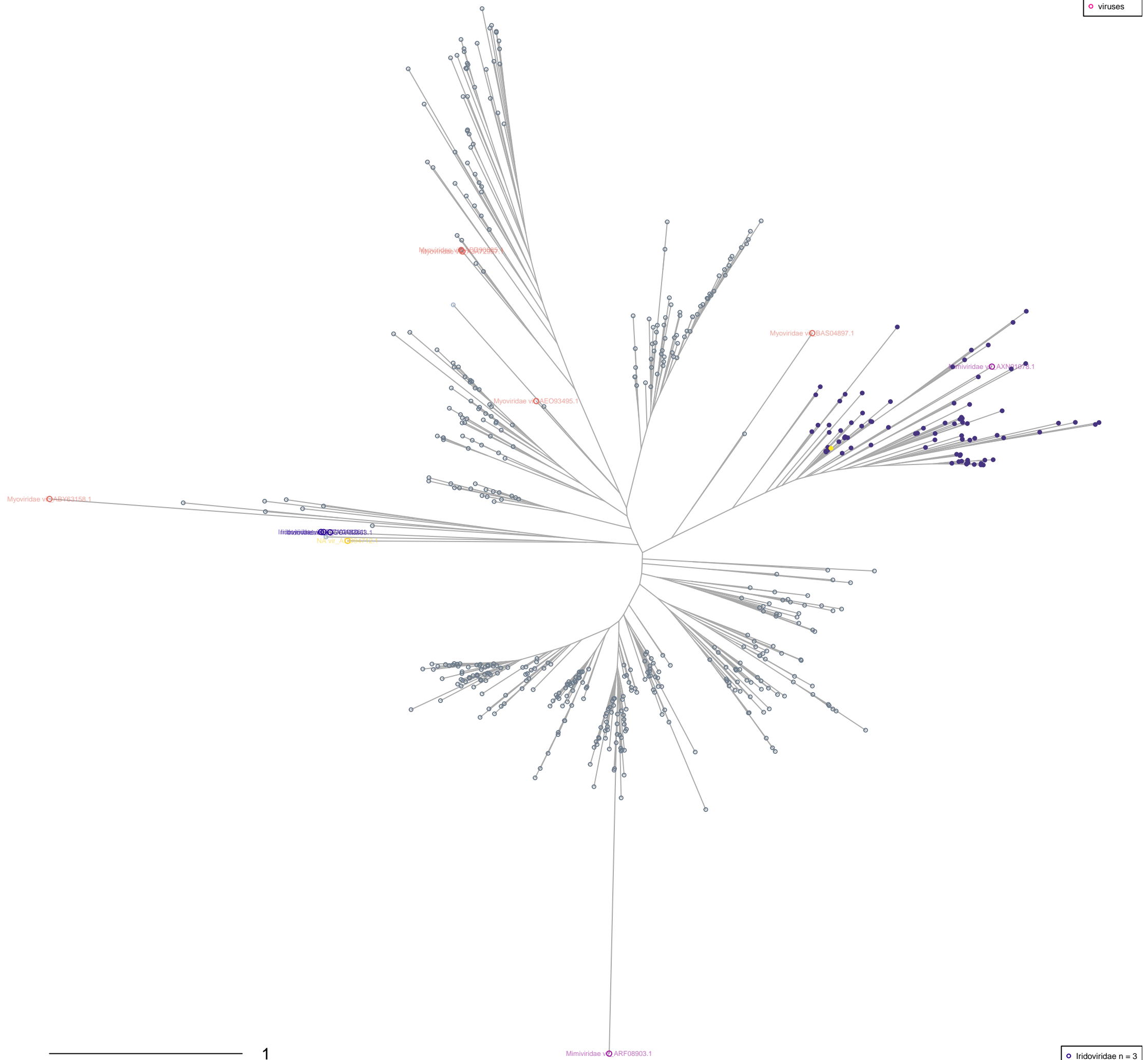
1

1

1

PTIP
vir.PTIP.phy.HG3.seqs.iqtree.treefile
n=554 sequences

- eukaryotes
- archaea
- bacteria
- viruses

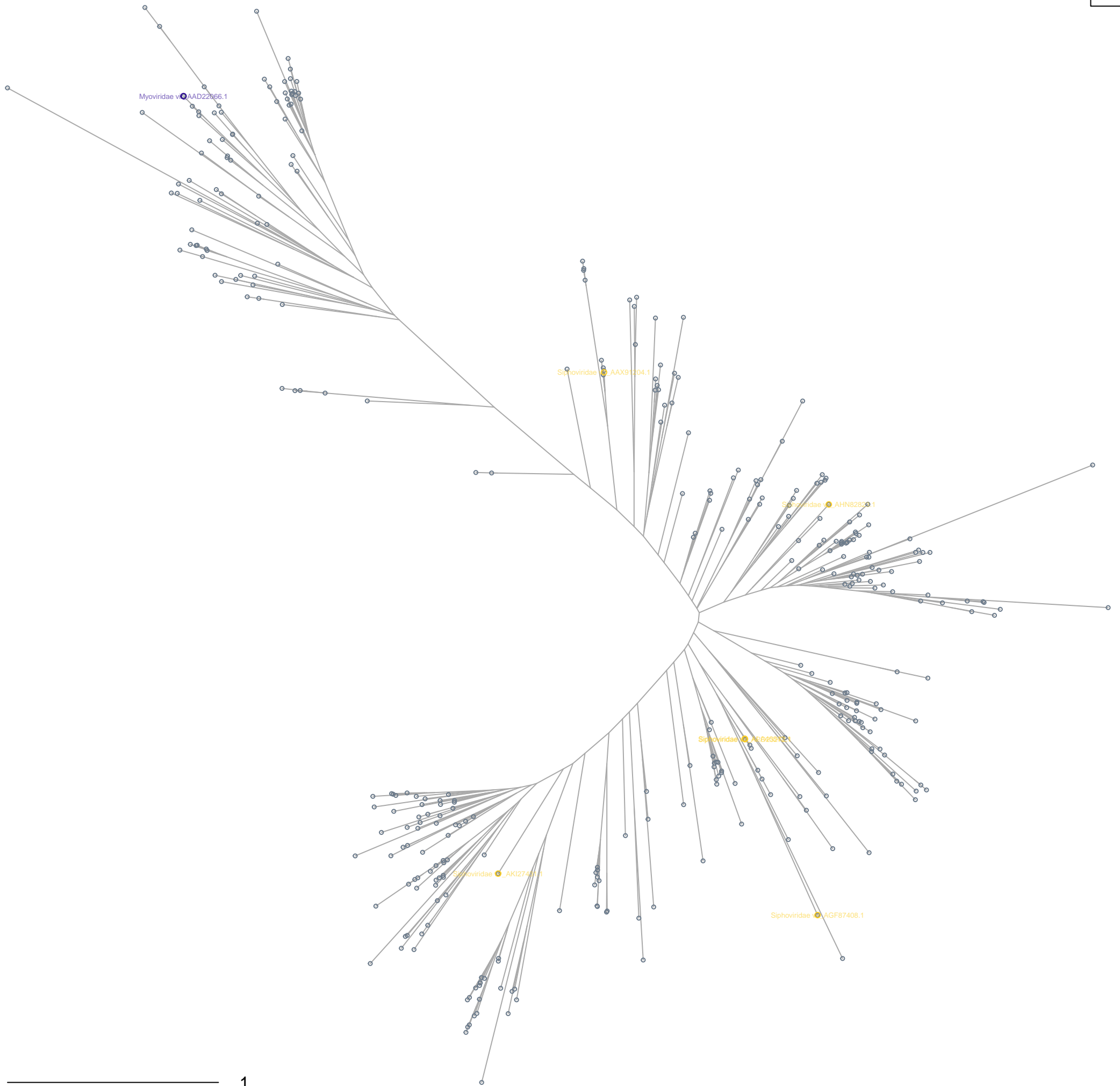


PTIP.HG3.4:RFC1 n = 85
other n = 1

Iridoviridae n = 3
Mimiviridae n = 2
Myoviridae n = 5
NA n = 1

PTIP
vir.PTIP.phy.HG4.seqs.iqtree.treefile
n=372 sequences

- eukaryotes
- archaea
- bacteria
- viruses



1

- Myoviridae n = 1
- Siphoviridae n = 6