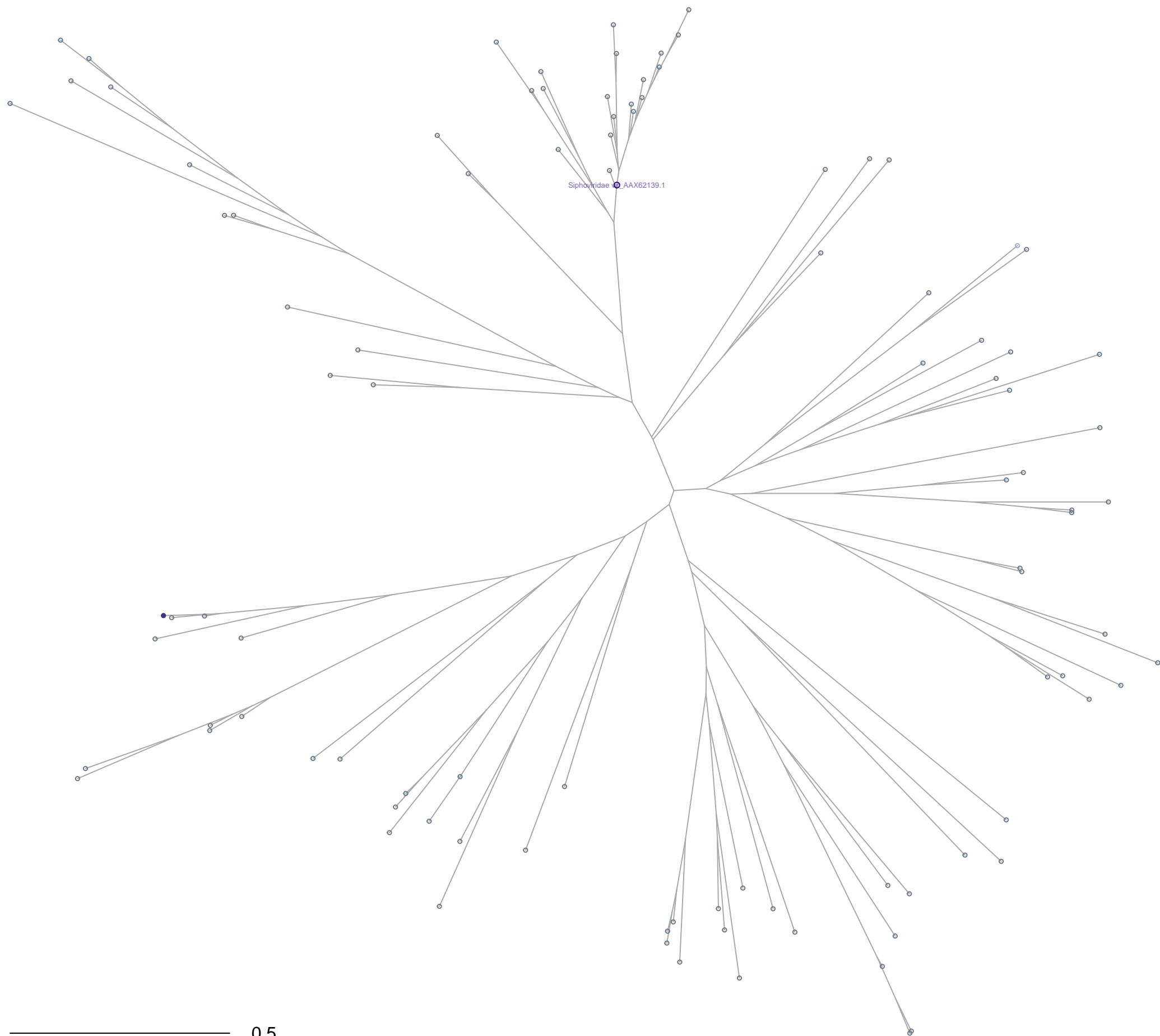


GNAT_acetyltr_2
vir.GNAT_acetyltr_2.phy.HG1.seqs.iqtree.treefile
n=101 sequences

eukaryotes
archaea
bacteria
viruses



0.5

other n = 1

Siphoviridae n = 1