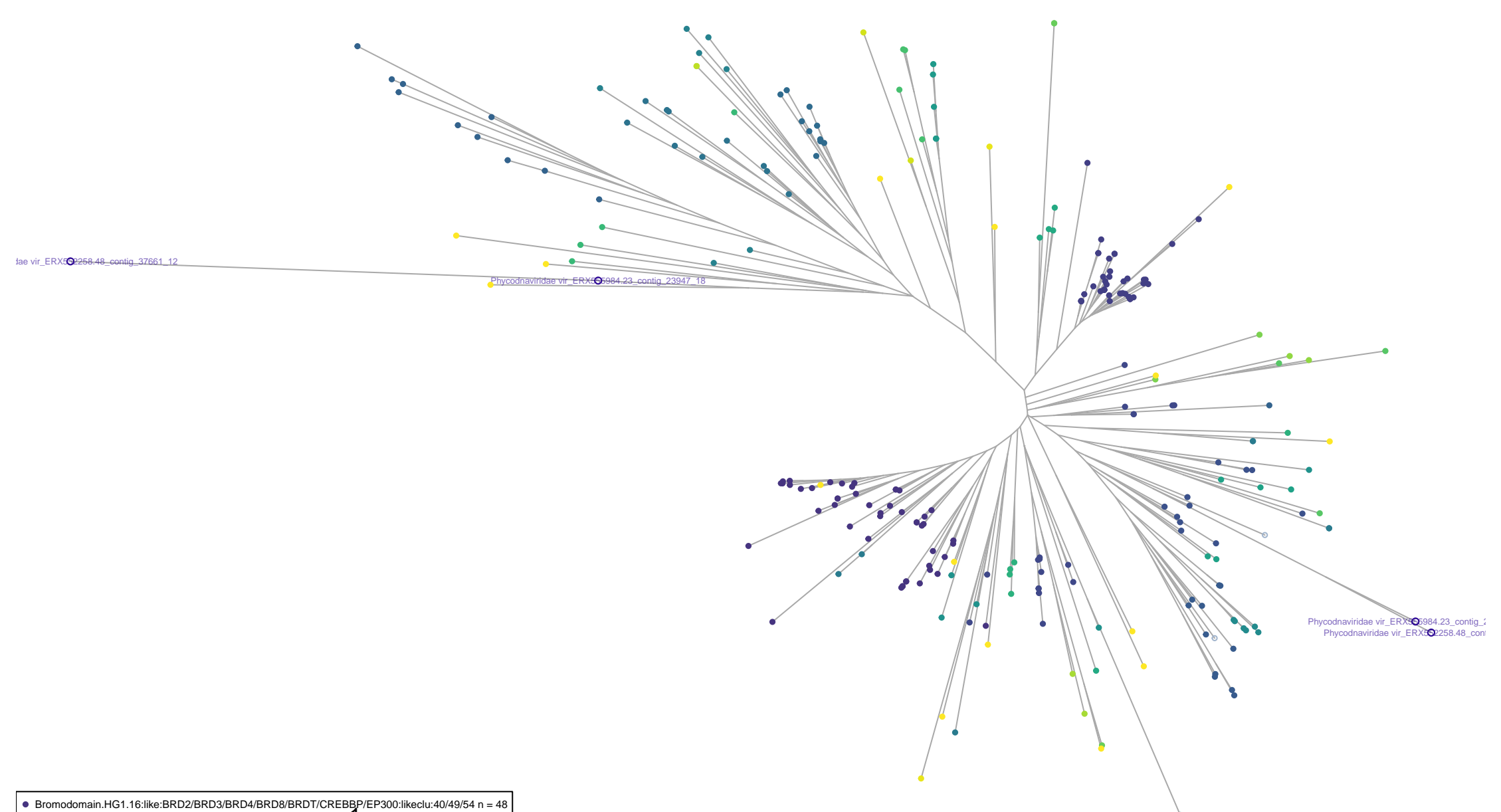


Bromodomain
vir.Bromodomain.phy.HG1.seqs.iqtree.treefile
n=259 sequences

- eukaryotes
- archaea
- bacteria
- viruses



- Bromodomain.HG1.16:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 48
- Bromodomain.HG1.49:BRD2/BRD3/BRD4/BRDT n = 44
- Bromodomain.HG1.15:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 18
- Bromodomain.HG1.32:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 11
- Bromodomain.HG1.38:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 11
- Bromodomain.HG7.1:BRWD1/BRWD3/PHIP n = 11
- Bromodomain.HG2.1:like:BRD1/BRD7/BRD9/BRPF1/BRPF3:likeclu:3/5 n = 10
- Bromodomain.HG2.5:BRD1/BRPF1/BRPF3 n = 10
- Bromodomain.HG1.19:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 7
- Bromodomain.HG2.0:like:BRD1/BRD7/BRD9/BRPF1/BRPF3:likeclu:3/5 n = 7
- Bromodomain.HG1.46:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 6
- Bromodomain.HG1.54:CREBBP/EP300 n = 6
- Bromodomain.HG3.6:BPTF n = 6
- Bromodomain.HG1.27:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 5
- Bromodomain.HG1.37:like:BRD2/BRD3/BRD4/BRDT:likeclu:49 n = 5
- Bromodomain.HG1.5:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 5
- Bromodomain.HG2.3:BRD7/BRD9 n = 4
- Bromodomain.HG3.10:KAT2A/KAT2B n = 4
- Bromodomain.HG1.11:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 3
- Bromodomain.HG3.3:like:BPTF/CECR2/KAT2A/KAT2B:likeclu:6/7/10 n = 3
- Bromodomain.HG1.25:like:BRD2/BRD3/BRD4/BRDT:likeclu:49 n = 2
- Bromodomain.HG1.28:like:BRD8:likeclu:40 n = 2
- Bromodomain.HG1.9:like:BRD2/BRD3/BRD4/BRD8/BRDT/CREBBP/EP300:likeclu:40/49/54 n = 2
- Bromodomain.HG2.2:like:BRD1/BRD7/BRD9/BRPF1/BRPF3:likeclu:3/5 n = 2
- Bromodomain.HG3.7:CECR2 n = 2
- Bromodomain.HG5.2:BAZ1A/BAZ1B/BAZ2A/BAZ2B n = 2
- other n = 15

Phycodnaviridae vir_ERX56984.23_contig_1
Phycodnaviridae vir_ERX562258.48_con

Phycodnaviridae n = 4