BPTP SPECIES DELIMITATION

bPTP web server (/ptp/)

Look up jobs (/findjob/)

Help (/help/)

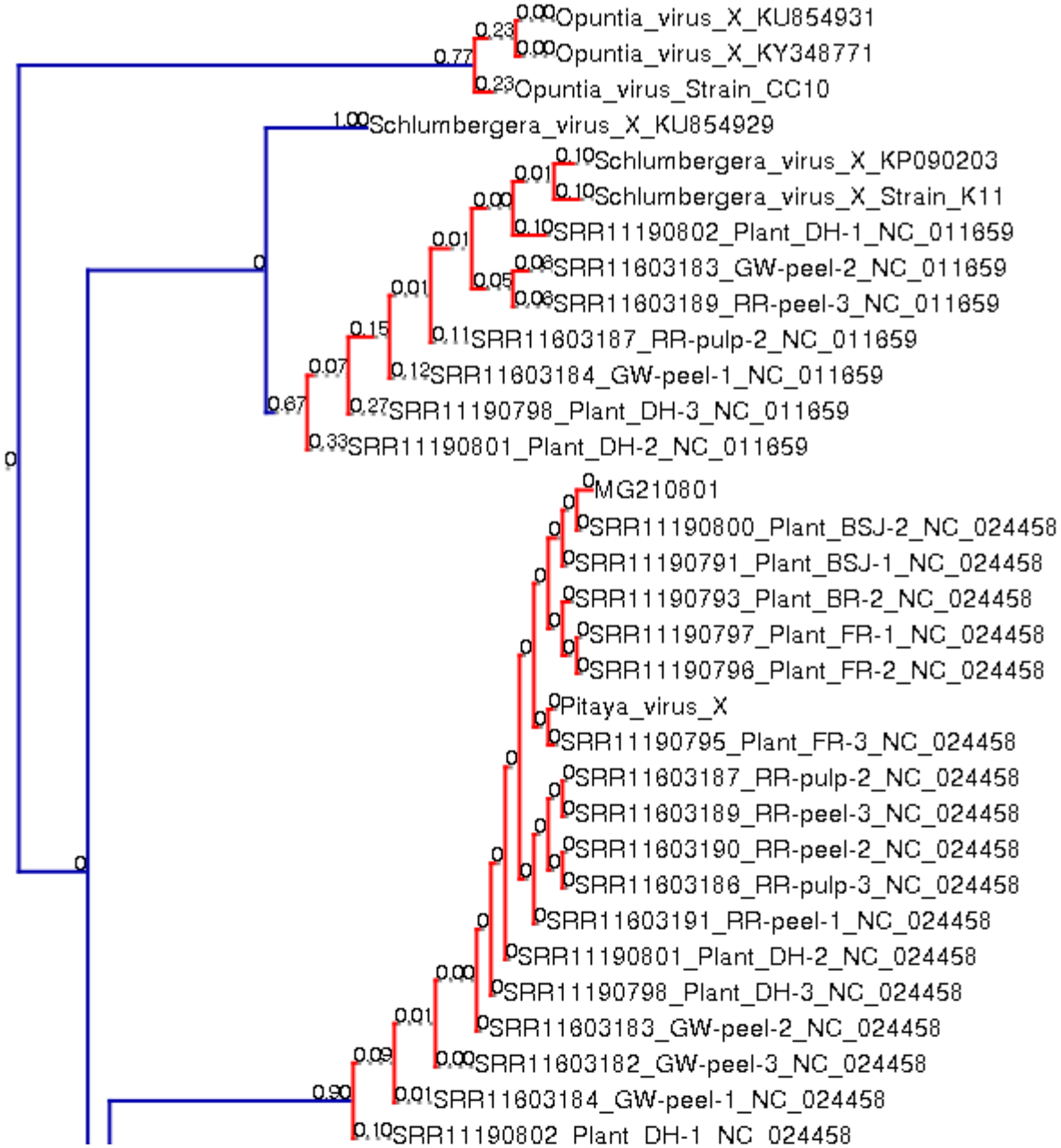
About PTP and bPTP (http://cme.h-its.org/exelixis/web/software/PTP/index.html)

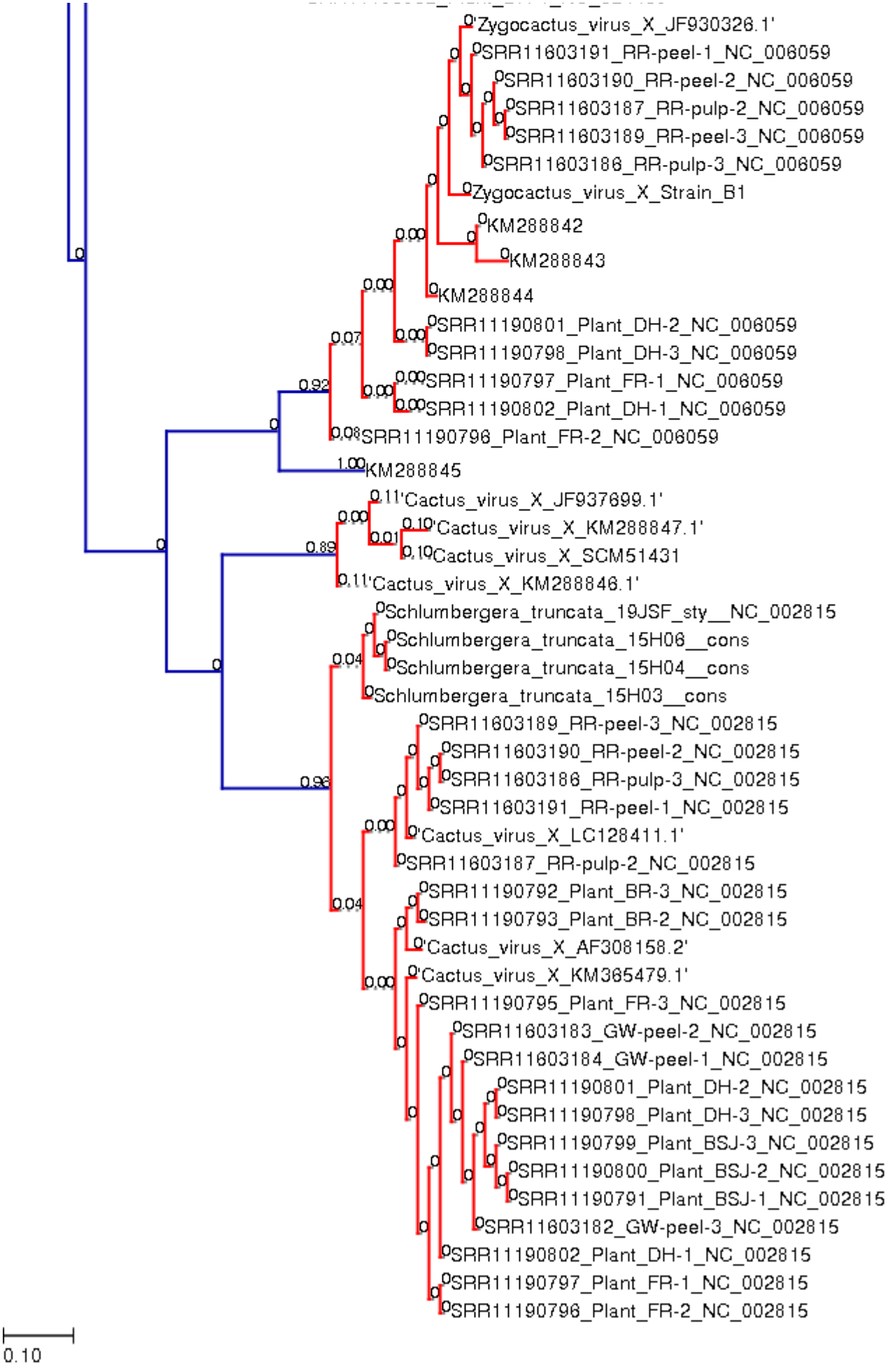
PTP paper (http://bioinformatics.oxfordjournals.org/content/29/22/2869)

Download PTP and bPTP (https://github.com/zhangjiajie/PTP)

PTP species delimitation results

Maximum likelihood solution:



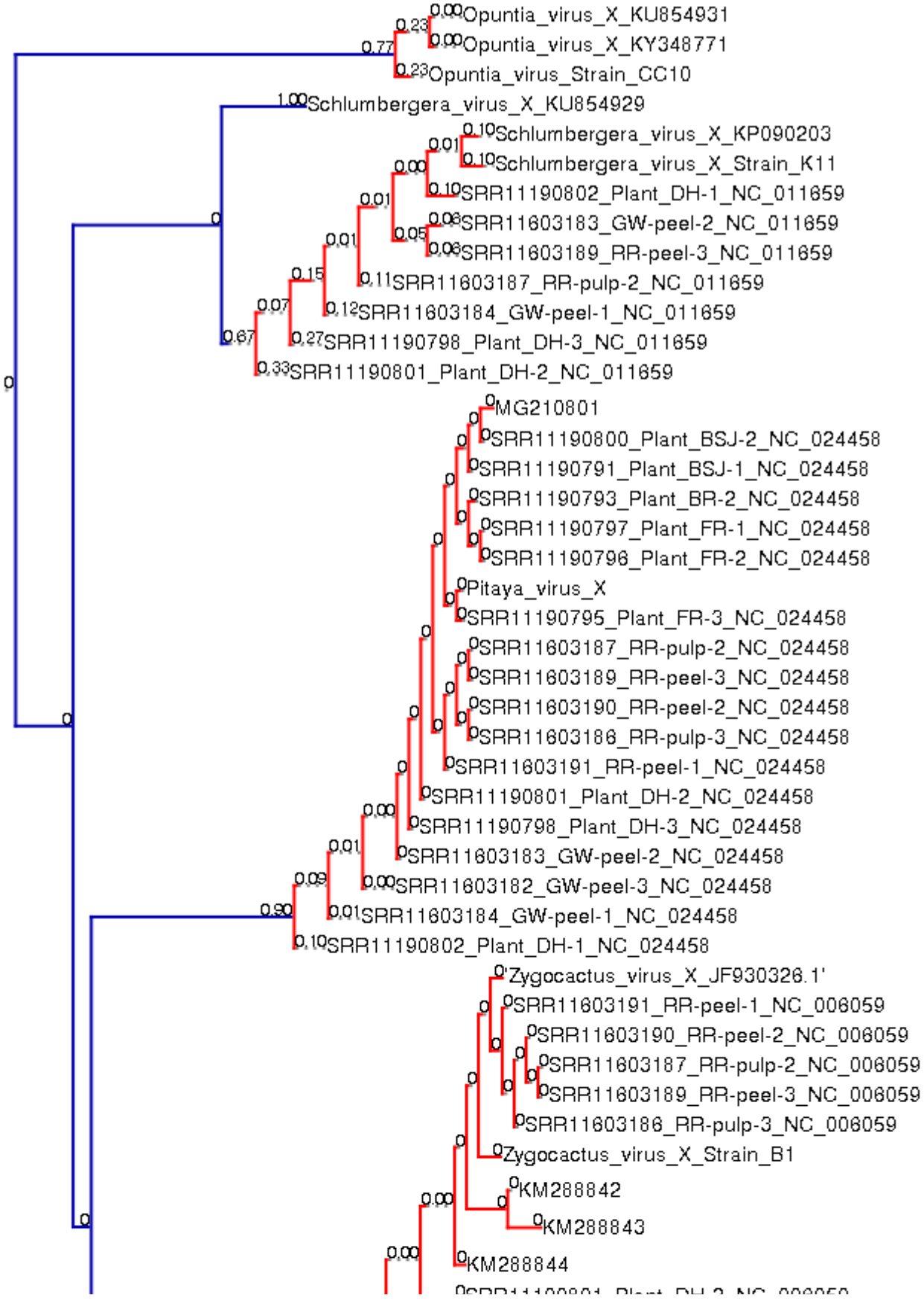

'Zygocactus_virus_X_JF930326.1'
SRR11603191_RR-peel-1_NC_006059
SRR11603190_RR-peel-2_NC_006059
SRR11603187_RR-pulp-2_NC_006059
SRR11603189_RR-peel-3_NC_006059
SRR11603186_RR-pulp-3_NC_006059
Zygocactus_virus_X_Strain_B1
KM288842
KM288843
KM288844
SRR11190801_Plant_DH-2_NC_006059
SRR11190798_Plant_DH-3_NC_006059
SRR11190797_Plant_FR-1_NC_006059
SRR11190802_Plant_DH-1_NC_006059
SRR11190796_Plant_FR-2_NC_006059
KM288845
'Cactus_virus_X_JF937699.1'
'Cactus_virus_X_KM288847.1'
Cactus_virus_X_SCM51431
'Cactus_virus_X_KM288846.1'
Schlumbergera_truncata_19JSF_sty__NC_002815
Schlumbergera_truncata_15H06__cons
Schlumbergera_truncata_15H04__cons
Schlumbergera_truncata_15H03__cons
SRR11603189_RR-peel-3_NC_002815
SRR11603190_RR-peel-2_NC_002815
SRR11603186_RR-pulp-3_NC_002815
SRR11603191_RR-peel-1_NC_002815
'Cactus_virus_X_LC128411.1'
SRR11603187_RR-pulp-2_NC_002815
SRR11190792_Plant_BR-3_NC_002815
SRR11190793_Plant_BR-2_NC_002815
'Cactus_virus_X_AF308158.2'
'Cactus_virus_X_KM365479.1'
SRR11190795_Plant_FR-3_NC_002815
SRR11603183_GW-peel-2_NC_002815
SRR11603184_GW-peel-1_NC_002815
SRR11190801_Plant_DH-2_NC_002815
SRR11190798_Plant_DH-3_NC_002815
SRR11190799_Plant_BSJ-3_NC_002815
SRR11190800_Plant_BSJ-2_NC_002815
SRR11190791_Plant_BSJ-1_NC_002815
SRR11603182_GW-peel-3_NC_002815
SRR11190802_Plant_DH-1_NC_002815
SRR11190797_Plant_FR-1_NC_002815
SRR11190796_Plant_FR-2_NC_002815
0.10

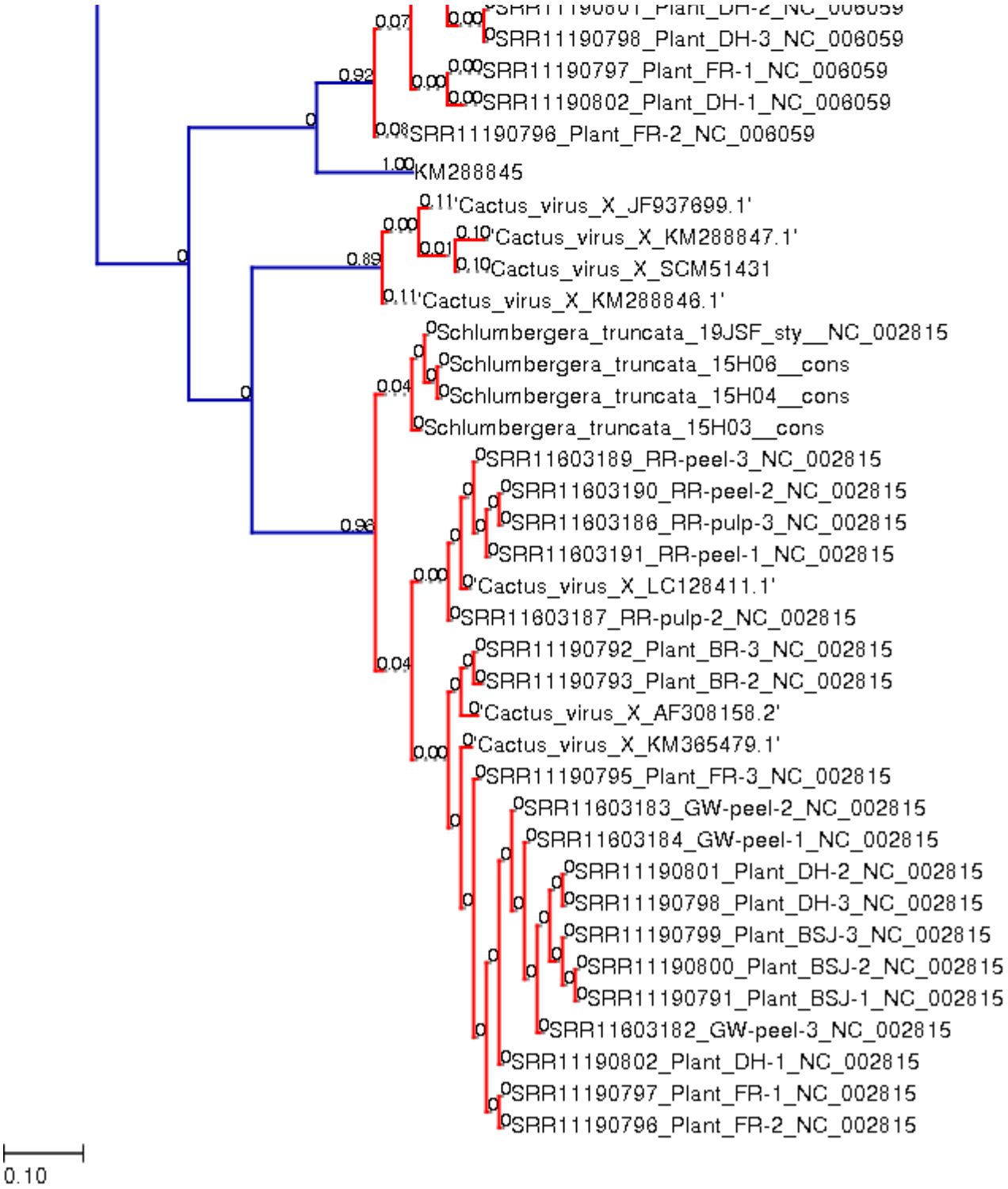
Download output in SVG (/download/99378/output.PTPMLPartition.txt.svg)

Download delimitation results (/download/99378/output.PTPMLPartition.txt)

Download annotated tree (/download/99378/output.PTPMLPartition.txt.ml.tre)

Highest Bayesian supported solution:





Download output in SVG (/download/99378/output.PTPhSupportPartition.txt.svg)

Download delimitation results (/download/99378/output.PTPhSupportPartition.txt)

Download annotated tree (/download/99378/output.PTPhSupportPartition.txt.sh.tre)

Download Bayesian results (/download/99378/output.PTPPartitions.txt)

Download likelihood trace file (/download/99378/output.PTPllh.txt)

Download likelihood trace plot in pdf (/download/99378/output.llh.pdf)

Download delimitation summary (/download/99378/output.PTPPartitonSummary.txt)

Accptance rate: 0.17002999999999999
Merge: 49995
Split: 50005
Estimated number of species is between 8 and 24
Mean: 10.00

PhyloMap-PTP

Show PhyloMap visualization of the maximum likelihood PTP species delimitation (/ptp/phylomap/?job_id=99378&email=boris-mptp-server@opayq.com)

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Contact through google-group (https://groups.google.com/forum/?hl=en#!forum/ptp-species-delimitation)