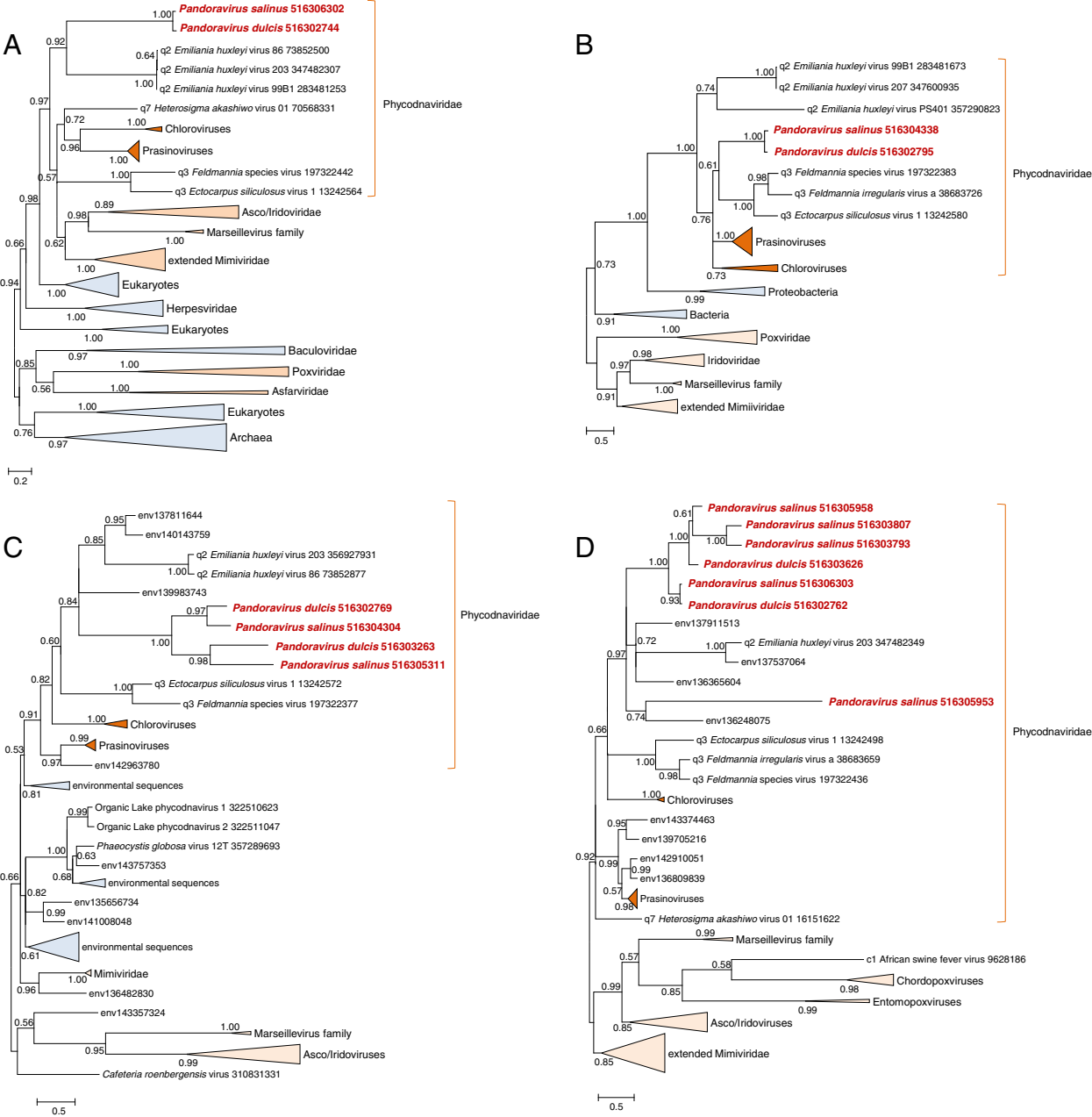
A
Pandoravirus salinus 516306302
Pandoravirus dulcis 516302744
q2 Emiliania huxleyi virus 86 73852500
q2 Emiliania huxleyi virus 203 347482307
q2 Emiliania huxleyi virus 99B1 283481253
q7 Heterosigma akashiwo virus 01 70568331
Chloroviruses
Prasinoviruses
q3 Feldmannia species virus 197322442
q3 Ectocarpus siliculosus virus 1 13242564
Phycodnaviridae
Asco/Iridoviridae
Marseillevirus family
extended Mimiviridae
Eukaryotes
Herpesviridae
Eukaryotes
Baculoviridae
Poxviridae
Asfarviridae
Eukaryotes
Archaea
0.2
B
q2 Emiliania huxleyi virus 99B1 283481673
q2 Emiliania huxleyi virus 207 347600935
q2 Emiliania huxleyi virus PS401 357290823
Pandoravirus salinus 516304338
Pandoravirus dulcis 516302795
q3 Feldmannia species virus 197322383
q3 Feldmannia irregularis virus a 38683726
q3 Ectocarpus siliculosus virus 1 13242580
Prasinoviruses
Chloroviruses
Phycodnaviridae
Proteobacteria
Bacteria
Poxviridae
Iridoviridae
Marseillevirus family
extended Mimiiviridae
0.5
C
env137811644
env140143759
q2 Emiliania huxleyi virus 203 356927931
q2 Emiliania huxleyi virus 86 73852877
env139983743
Pandoravirus dulcis 516302769
Pandoravirus salinus 516304304
Pandoravirus dulcis 516303263
Pandoravirus salinus 516305311
q3 Ectocarpus siliculosus virus 1 13242572
q3 Feldmannia species virus 197322377
Chloroviruses
Prasinoviruses
Phycodnaviridae
env142963780
environmental sequences
Organic Lake phycodnavirus 1 322510623
Organic Lake phycodnavirus 2 322511047
Phaeocystis globosa virus 12T 357289693
env143757353
environmental sequences
env135656734
env141008048
environmental sequences
Mimiviridae
env136482830
env143357324
Marseillevirus family
Asco/Iridoviruses
Cafeteria roenbergensis virus 310831331
0.5
D
Pandoravirus salinus 516305958
Pandoravirus salinus 516303807
Pandoravirus salinus 516303793
Pandoravirus dulcis 516303626
Pandoravirus salinus 516306303
Pandoravirus dulcis 516302762
env137911513
q2 Emiliania huxleyi virus 203 347482349
env137537064
env136365604
Pandoravirus salinus 516305953
env136248075
q3 Ectocarpus siliculosus virus 1 13242498
q3 Feldmannia irregularis virus a 38683659
q3 Feldmannia species virus 197322436
Chloroviruses
env143374463
env139705216
env142910051
env136809839
Prasinoviruses
q7 Heterosigma akashiwo virus 01 16151622
Phycodnaviridae
Marseillevirus family
c1 African swine fever virus 9628186
Chordopoxviruses
Entomopoxviruses
Asco/Iridoviruses
extended Mimiviridae
0.5