



3S_rRNA_gene

okonta__1.00__f_Holozoa__1.00__g_Choanozoa__1.00__s_Meta:
i__1.00__f_Perkinsidae__0.98__g_A31__0.98__s_uncultured_euka
__f_Intramacronucleata__0.94__g_Conthreep__0.94__s_Oligohym
__f_Intramacronucleata__1.00__g_Conthreep__1.00__s_Oligohym
__f_Intramacronucleata__0.90__g_Conthreep__0.90__s_Oligohym
__f_Intramacronucleata__0.73__g_Conthreep__0.72__s_Oligohym
es__0.31__o_Teleaulax__0.12__f_uncultured_marine_eukaryote__
__f_Intramacronucleata__0.71__g_Conthreep__0.69__s_Prostc
__f_Intramacronucleata__0.79__g_Conthreep__0.79__s_Oligohym
thokonta__0.65__f_Nuclelmycea__0.65__g_Fungi__0.65__s_Dikar
lales__0.79__o_Rhodomonas__0.59__f_Rhodomonas_marina__0.
lales__1.00__o_Rhodomonas__1.00__f_Rhodomonas_marina__0.
lales__0.73__o_Rhodomonas__0.73__f_Rhodomonas_marina__0.
__0.23__f_Polycystinea__0.23__g_Spumellaria__0.23__s_unculture
yta__0.78__f_Phragmoplastophyta__0.78__g_Streptophyta__0.78
yta__0.95__f_Phragmoplastophyta__0.95__g_Streptophyta__0.95_
__f_Intramacronucleata__0.39__g_Conthreep__0.39__s_Oligohym
thokonta__0.83__f_Nuclelmycea__0.83__g_Fungi__0.83__s_Dikar
__0.78__f_Chrysophyceae__0.78__g_Ochromonadales__0.78__s_C
__0.99__f_Chrysophyceae__0.99__g_Ochromonadales__0.98__s_C
__1.00__f_Chrysophyceae__1.00__g_Ochromonadales__0.97__s_C
__0.85__f_Chrysophyceae__0.85__g_Ochromonadales__0.82__s_C
__0.78__f_Chrysophyceae__0.78__g_Ochromonadales__0.77__s_C
__0.82__f_Chrysophyceae__0.82__g_Ochromonadales__0.82__s_C
__0.90__f_Chrysophyceae__0.89__g_Ochromonadales__0.88__s_C
thokonta__0.93__f_Nuclelmycea__0.91__g_Fungi__0.91__s_Dikar
thokonta__0.86__f_Nuclelmycea__0.86__g_Fungi__0.81__s_Dikar
thokonta__1.00__f_Nuclelmycea__1.00__g_Fungi__1.00__s_Dikar
thokonta__0.97__f_Nuclelmycea__0.96__g_Fungi__0.96__s_Dikar
thokonta__0.66__f_Nuclelmycea__0.61__g_Fungi__0.61__s_Dikar
akobida__0.48__o_uncultured_dinoflagellate__0.48
akobida__1.00__o_uncultured_dinoflagellate__1.00
akobida__0.20__o_uncultured_dinoflagellate__0.20
yta__0.72__f_Phragmoplastophyta__0.72__g_Streptophyta__0.72_
yta__1.00__f_Phragmoplastophyta__1.00__g_Streptophyta__1.00_
yta__0.94__f_Phragmoplastophyta__0.94__g_Streptophyta__0.94
yta__0.56__f_Phragmoplastophyta__0.56__g_Streptophyta__0.56_
thokonta__0.87__f_Nuclelmycea__0.87__g_Fungi__0.87__s_Dikar
s__1.00__o_Rhodomonas__0.89__f_Rhodomonas_sp._CCMP743_
s__0.91__o_Rhodomonas__0.91__f_Rhodomonas_sp._CCMP743_
yta__0.15__f_Phragmoplastophyta__0.15__g_Streptophyta__0.15_
s__0.57__o_Rhodomonas__0.57__f_Rhodomonas_sp._CCMP743_
s__0.97__o_Rhodomonas__0.97__f_Rhodomonas_sp._CCMP743_
s__0.99__o_Rhodomonas__0.99__f_Rhodomonas_sp._CCMP743_
monadales__0.95__o_Rhodomonas_sp._KMMCC_574__0.77
s__0.89__o_Rhodomonas__0.83__f_Rhodomonas_sp._CCMP743_
s__1.00__o_Rhodomonas__1.00__f_Rhodomonas_sp._CCMP743_
konta__1.00__f_Nuclelmycea__1.00__g_Fungi__0.96__s_Cryptom