

Otu	Taxonomy Name	E-Value	Sim	Accession
Otu00023	<i>Helicopedinella sp. RCC2284</i>	8.44E-115	100.00%	LN735338.3
Otu00036	<i>Bacillariophyta sp. 867-32</i>	2.84E-118	99.10%	KP792487.1
Otu00054	<i>Emiliana huxleyi</i>	2.84E-118	99.10%	JN022705.1
Otu00062	Uncultured Marine Eukaryote	5.22E-120	99.60%	KX938088.1
Otu00064	<i>Phaeocystis globosa</i>	5.90E-116	99.10%	KC900889.1
Otu00067	Uncultured Picoeukaryote	4.60E-113	97.80%	JX291751.1
Otu00079	Uncultured Dictyochophyte	1.55E-116	98.70%	EF052136.1
Otu00086	<i>Virgulinema fragilis</i>	1.55E-116	98.70%	JN207213.1
Otu00097	Uncultured Marine Eukaryote	2.84E-118	99.10%	KX935068.1
Otu00100	<i>Planoglabratella opercularis</i>	1.37E-109	97.00%	KP792467.1
Otu00113	<i>Florenciella parvula</i>	1.55E-116	99.10%	LN735274.2
Otu00116	Uncultured Chrysophyte	2.84E-118	99.10%	EF052163.1
Otu00120	Uncultured Marine Eukaryote	8.44E-115	98.30%	KX937565.1
Otu00158	Uncultured Haptophyte	8.44E-115	98.30%	EF052065.1
Otu00162	<i>Braarudosphaera bigelowii</i>	2.84E-118	99.10%	AB847986.2
Otu00218	Uncultured Chrysophyte	2.84E-118	99.10%	EF052251.1
Otu00234	<i>Asterionella formosa</i>	4.06E-106	96.10%	KC509519.1
Otu00238	<i>Prasinophyceae sp. RCC999</i>	1.75E-112	98.30%	LN735517.3
Otu00242	<i>Proboscia inermis</i>	2.79E-96	94.40%	FJ002201.1
Otu00255	Uncultured Marine Eukaryote	3.55E-110	97.80%	KX937647.1
Otu00260	Uncultured Marine Eukaryote	5.22E-120	99.60%	KX938009.1
Otu00268	Uncultured Marine Eukaryote	5.82E-116	100.00%	KX938202.1
Otu00270	Eukaryote SCGC AAA074-P04	2.51E-111	97.40%	JF488733.1
Otu00276	<i>Virgulinema fragilis</i>	4.60E-113	97.80%	KP792491.1
Otu00284	<i>Ralstonia solanacearum</i>	2.51E-111	97.40%	CP011998.1
Otu00319	Uncultured Marine Eukaryote	2.21E-104	96.10%	KX938047.1
Otu00335	<i>Rhizosolenia imbricata</i>	4.60E-113	97.80%	KJ958482.1
Otu00343	<i>Proboscia indica</i>	8.44E-115	98.30%	FJ002241.1
Otu00344	Uncultured Marine Eukaryote	1.55E-116	98.70%	KX937668.1
Otu00364	<i>Syracosphaera pulchra</i>	1.20E-102	96.00%	LN735258.2
Otu00366	<i>Calcidiscus quadriperforatus</i>	2.51E-111	97.40%	LN735221.2
Otu00370	Uncultured Chrysophyte	3.16E-103	96.10%	EF052251.1
Otu00379	<i>Actaea racemosa</i>	1.07E-117	99.60%	KY085920.1
Otu00395	<i>Teleaulax amphioxeia</i>	4.62E-113	99.10%	KX816862.1
Otu00423	Uncultured Diatom	1.21E-102	95.30%	KX938043.1
Otu00494	<i>Virgulinema fragilis</i>	2.51E-111	97.40%	JN207220.1
Otu00502	Uncultured Chrysophyte	5.22E-120	99.60%	KX938184.1
Otu00530	<i>Chaetoceros calcitrans</i>	8.44E-115	98.30%	LN735292.2
Otu00531	<i>Thalassionema sp. CCMP1100</i>	4.06E-106	96.10%	FJ002197.1
Otu00534	Uncultured marine eukaryote	2.84E-118	99.10%	KX934876.1
Otu00548	Eukaryote SCGC AAA074-K22	8.44E-115	98.30%	JF488711.1
Otu00549	<i>Pelagomonas sp. RCC986</i>	1.21E-113	100.00%	LN735511.3
Otu00565	<i>Thalassionema sp. CCMP1100</i>	1.37E-109	97.00%	FJ002197.1
Otu00573	<i>Planoglabratella opercularis</i>	4.60E-113	97.80%	KP792480.1
Otu00581	Uncultured Phototrophic Eukaryote	1.55E-116	98.70%	KP792480.1

Otu00585	Uncultured Marine Eukaryote	4.60E-113	97.80%	KX938161.1
Otu00602	<i>Rhizosolenia imbricata</i>	2.21E-104	95.70%	KJ958482.1
Otu00624	Uncultured Marine Eukaryote	1.55E-116	98.70%	KX937932.1
Otu00636	<i>Helicosphaera carteri</i>	8.44E-115	98.30%	LN735234.2