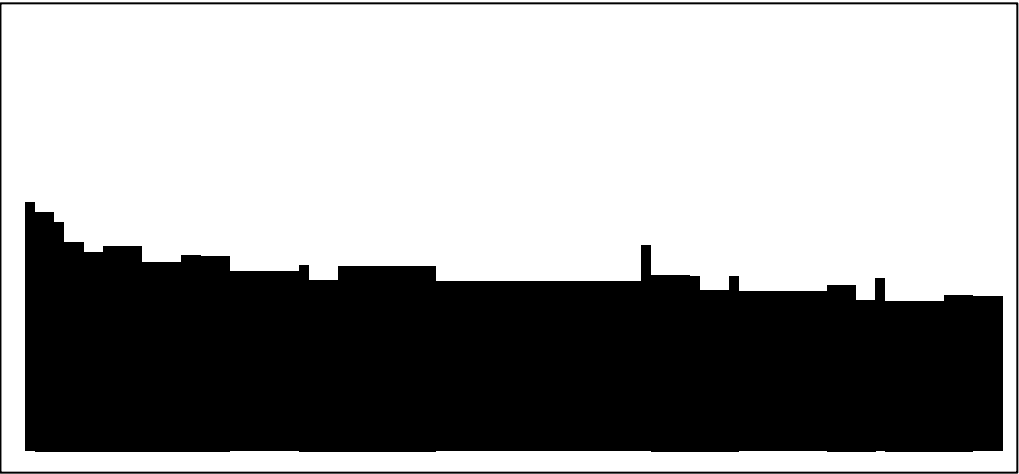


OTU_105;size=23182;.csv

uncultured fungus

91.67



Uncultured fungus mitochondrial partial cox1 gene	FN640494	91.67	99.54	1.32955e-80	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN640537	91.2	99.54	5.65341e-79	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599574	91.2	99.54	5.65341e-79	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600272	90.74	99.54	6.88726e-78	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN640525	89.81	99.54	3.56769e-75	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599598	89.81	99.54	3.56769e-75	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600228	89.35	99.54	1.51702e-73	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600172	89.35	99.54	1.51702e-73	uncultured fungus
Leohumicola sp. DAOM 239516 cytochrome oxidase sub	EU678468	89.62	97.7	5.29492e-73	Leohumicola sp. DAOM 239516
Leohumicola verrucosa strain DAOM 231142 cytochrom	EU678458	89.62	97.7	5.29492e-73	Leohumicola verrucosa
Leohumicola verrucosa strain DAOM 239536 cytochrom	EU678454	89.62	97.7	5.29492e-73	Leohumicola verrucosa
Leohumicola verrucosa strain DAOM 239497 cytochrom	EU678418	89.62	97.7	5.29492e-73	Leohumicola verrucosa
Uncultured fungus mitochondrial partial cox1 gene	FN640520	88.89	99.54	1.84811e-72	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600185	88.89	99.54	1.84811e-72	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599586	88.89	99.54	1.84811e-72	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599552	88.89	99.54	1.84811e-72	uncultured fungus
Pseudogymnoascus bhattii isolate C10 cytochrome ox	FJ590558	89.2	98.16	6.45053e-72	Pseudogymnoascus bhattii
Leohumicola minima strain DAOM 232587 cytochrome o	EU678466	89.2	98.16	6.45053e-72	Leohumicola minima
Leohumicola verrucosa strain DAOM 231144 cytochrom	EU678459	89.15	97.7	2.25146e-71	Leohumicola verrucosa
Leohumicola verrucosa strain DAOM 230085 cytochrom	EU678456	89.15	97.7	2.25146e-71	Leohumicola verrucosa
Leohumicola verrucosa strain HNLHM117 cytochrome o	EU678449	89.15	97.7	2.25146e-71	Leohumicola verrucosa
Uncultured fungus mitochondrial partial mRNA for c	FN600327	88.43	99.54	7.85836e-71	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600319	88.43	99.54	7.85836e-71	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600308	88.43	99.54	7.85836e-71	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599611	88.43	99.54	7.85836e-71	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599583	88.43	99.54	7.85836e-71	uncultured fungus
Tetracladium apiense strain CCM F-23099 cytochrome	EU883398	88.43	99.54	7.85836e-71	Tetracladium apiense
Tetracladium breve strain CCM F-10501 cytochrome c	EU883397	88.43	99.54	7.85836e-71	Tetracladium breve
Leohumicola levissima strain HNLHM2 cytochrome oxi	EU678419	88.73	98.16	7.85836e-71	Leohumicola levissima
Uncultured fungus mitochondrial partial cox1 gene	FN640486	88.02	100	2.74284e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600279	88.02	100	2.74284e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599602	88.02	100	2.74284e-70	uncultured fungus
Gymnostellatospora sp. Cr1 cytochrome oxidase subu	FJ590567	88.68	97.7	2.74284e-70	Gymnostellatospora sp. Cr1
Pseudogymnoascus roseus isolate Cr26 cytochrome ox	FJ590566	88.68	97.7	2.74284e-70	Pseudogymnoascus roseus
Pseudogymnoascus roseus isolate Cr4 cytochrome oxi	FJ590565	88.68	97.7	2.74284e-70	Pseudogymnoascus roseus
Geomyces pannorum isolate Cr3 cytochrome oxidase s	FJ590564	88.68	97.7	2.74284e-70	Pseudogymnoascus pannorum
Pseudogymnoascus roseus isolate C12 cytochrome oxi	FJ590563	88.68	97.7	2.74284e-70	Pseudogymnoascus roseus
Gymnostellatospora alpina isolate C11 cytochrome o	FJ590562	88.68	97.7	2.74284e-70	Gymnostellatospora alpina
Pseudogymnoascus roseus isolate C1 cytochrome oxid	FJ590560	88.68	97.7	2.74284e-70	Pseudogymnoascus roseus
Tetracladium furcatum strain CCM F-11883 cytochrom	EU883411	88.68	97.7	2.74284e-70	Tetracladium furcatum
Leohumicola terminalis strain DAOM 231145 cytochro	EU678464	88.68	97.7	2.74284e-70	Leohumicola terminalis
Leohumicola verrucosa strain DAOM 231141 cytochrom	EU678457	88.68	97.7	2.74284e-70	Leohumicola verrucosa
Uncultured fungus mitochondrial partial cox1 gene	FN640542	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN640489	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600335	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600323	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600307	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600304	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600269	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600250	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600241	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599610	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599608	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599606	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599592	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599591	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599577	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599573	87.96	99.54	9.57344e-70	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599563	87.96	99.54	9.57344e-70	uncultured fungus
Tetracladium sp. CCM F-10008 cytochrome c oxidase	EU883412	87.96	99.54	9.57344e-70	Tetracladium sp. CCM F-10008
Tetracladium setigerum strain CCM F-20987 cytochro	EU883408	87.96	99.54	9.57344e-70	Tetracladium setigerum
Tetracladium setigerum strain CCM F-10186 cytochro	EU883405	87.96	99.54	9.57344e-70	Tetracladium setigerum
Tetracladium apiense strain CCM F-23299 cytochrome	EU883400	87.96	99.54	9.57344e-70	Tetracladium apiense
Myxotrichum deflexum strain UAMH6365 cytochrome ox	EU678467	89.66	93.55	3.34146e-69	Myxotrichum deflexum
Leohumicola sp. DAOM 231148 cytochrome oxidase sub	EU678461	88.26	98.16	3.34146e-69	Leohumicola sp. DAOM 231148
Leohumicola atra strain DAOM 239535 cytochrome oxi	EU678450	88.26	98.16	3.34146e-69	Leohumicola atra
Leohumicola levissima strain HNLHM41 cytochrome ox	EU678429	88.26	98.16	3.34146e-69	Leohumicola levissima
Leohumicola atra strain DAOM 239515 cytochrome oxi	EU678425	88.26	98.16	3.34146e-69	Leohumicola atra
Pseudogymnoascus pannorum strain NN050741 mitochon	KR055655	88.21	97.7	1.16628e-68	Pseudogymnoascus pannorum
Uncultured fungus mitochondrial partial cox1 gene	FN599605	87.56	100	1.16628e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599564	87.56	100	1.16628e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599557	87.56	100	1.16628e-68	uncultured fungus
Pseudogymnoascus sp. BM-2009-4 cytochrome oxidase	FJ590561	88.21	97.7	1.16628e-68	Pseudogymnoascus sp. BM-2009-4
Arachnida sp. BOLD:ACM9770 voucher BIOUG11746-C06	KR099129	87.5	99.54	4.07073e-68	Arachnida sp. BOLD:ACM9770
Phialocephala europaea strain 3_122_3 cytochrome o	JN091450	87.5	99.54	4.07073e-68	Phialocephala europaea
Uncultured fungus mitochondrial partial mRNA for c	FN600318	87.5	99.54	4.07073e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599599	87.5	99.54	4.07073e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599593	87.5	99.54	4.07073e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599590	87.5	99.54	4.07073e-68	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599558	87.5	99.54	4.07073e-68	uncultured fungus
Tetracladium palmatum strain CCM F-10001 cytochrom	EU883403	87.5	99.54	4.07073e-68	Tetracladium palmatum
Tetracladium marchalianum strain CCM F-26299 cytoc	EU883392	87.5	99.54	4.07073e-68	Tetracladium marchalianum
Leohumicola lenta strain DAOM 231149 cytochrome ox	EU678465	87.79	98.16	4.07073e-68	Leohumicola lenta
Leohumicola sp. DAOM 230084 cytochrome oxidase sub	EU678455	87.79	98.16	4.07073e-68	Leohumicola sp. DAOM 230084
Leohumicola levissima strain HNLHM4 cytochrome oxi	EU678422	87.79	98.16	4.07073e-68	Leohumicola levissima
Uncultured fungus mitochondrial partial cox1 gene	FN599616	87.1	100	1.42082e-67	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599601	87.1	100	1.42082e-67	uncultured fungus
Tetracladium furcatum strain CCM F-06983 cytochrom	EU883404	88.1	96.77	1.42082e-67	Tetracladium furcatum
Uncultured fungus mitochondrial partial cox1 gene	FN640524	87.04	99.54	4.95916e-67	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN640498	87.04	99.54	4.95916e-67	uncultured fungus
Uncultured fungus mitochondrial partial mRNA for c	FN600287	87.04	99.54	4.95916e-67	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599565	87.04	99.54	4.95916e-67	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN599562	87.04	99.54	4.95916e-67	uncultured fungus
Tetracladium breve strain CCM F-12505 cytochrome c	EU883410	87.04	99.54	4.95916e-67	Tetracladium breve
Leohumicola incrustata strain DAOM 239500 cytochro	EU678448	87.32	98.16	1.73092e-66	Leohumicola incrustata
Leohumicola incrustata strain DAOM 239504 cytochro	EU678441	87.32	98.16	1.73092e-66	Leohumicola incrustata
Leohumicola incrustata strain HNLHM91 cytochrome o	EU678439	87.32	98.16	1.73092e-66	Leohumicola incrustata
Marssonina brunnea f. sp. 'multigermtubi' mitochon	JN204424	87.26	97.7	6.0415e-66	Marssonina brunnea f. sp. 'multigermtubi'
Uncultured fungus mitochondrial partial cox1 gene	FN640540	87.26	97.7	6.0415e-66	uncultured fungus
Uncultured fungus mitochondrial partial cox1 gene	FN640511	87.26	97.7	6.0415e-66	uncultured fungus