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ITS4-ITS5 D1-D2,

[www.ncbi.com](http://www.ncbi.com)

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Description	Max score	Total score	Query cover	E value	Ident	Accession
Microsporium canis strain CHUS14303 18S ribosomal RNA gene, part	1301	1301	100%	0.0	99%	EF581130.1
Microsporium canis strain CHUS65602 18S ribosomal RNA gene, part	1301	1301	100%	0.0	99%	EF581129.1
Microsporium canis small subunit ribosomal RNA gene, partial sequ	1301	1301	100%	0.0	99%	AF168127.1
Microsporium canis mRNA for 5.8S ribosomal RNA, internal transcri	1301	1301	100%	0.0	99%	AJ252329.1
Microsporium ferrugineum strain CHUS95602 18S ribosomal RNA ge	1290	1290	100%	0.0	99%	EF581133.1
Microsporium ferrugineum mRNA for 5.8S ribosomal RNA, internal	1290	1290	100%	0.0	99%	AJ252338.1
Microsporium canis strain LM 2174 18S ribosomal RNA gene, partial	1288	1288	99%	0.0	99%	FJ181444.1
Microsporium canis strain ATCC 23828 18S ribosomal RNA gene, par	1288	1288	99%	0.0	99%	AY213652.1
Microsporium canis strain ATCC MYA-4605 18S ribosomal RNA gene,	1282	1282	98%	0.0	99%	GU291265.1
Microsporium canis mRNA for 5.8S ribosomal RNA, internal transcri	1282	1282	100%	0.0	99%	AJ252328.1
Microsporium ferrugineum isolate 651 18S ribosomal RNA gene, pa	1277	1277	99%	0.0	99%	JN134136.1
Microsporium ferrugineum isolate 650 18S ribosomal RNA gene, pa	1277	1277	99%	0.0	99%	JN134135.1
Microsporium canis 5.8S rRNA gene, ITS1 and ITS2, complete sequ	1277	1277	100%	0.0	99%	AJ000619.1
Microsporium canis strain BMU02865 18S ribosomal RNA gene, parti	1273	1273	97%	0.0	99%	EF631606.1
Microsporium distortum strain ATCC 26224 18S ribosomal RNA gene	1273	1273	97%	0.0	99%	EF631808.1
Microsporium ferrugineum mRNA for 5.8S ribosomal RNA, internal	1271	1271	100%	0.0	99%	AJ252335.1

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Microsporium audouinii strain CBS 109478 18S ribosomal RNA	1805	1805	98%	0.0	99%	GU213362.1
Arthroderma curreyi partial 18S rRNA gene, strain CBS 138.26	1797	1797	100%	0.0	99%	AJ315165.1
Microsporium canis strain BMU02865 18S ribosomal RNA gene,	1792	1792	97%	0.0	99%	EF631606.1
Microsporium distortum strain ATCC 26224 18S ribosomal RNA	1792	1792	97%	0.0	99%	EF631608.1
Microsporium canis strain ATCC 36299 18S ribosomal RNA gene	1792	1792	97%	0.0	99%	EF631605.1
Microsporium canis 18S ribosomal RNA gene, partial sequence	1790	1790	97%	0.0	99%	AY083227.1
Arthroderma ciferrii isolate AFTOL-ID 428 18S ribosomal RNA	1788	1788	99%	0.0	99%	AB015770.1
Arthroderma ciferrii isolate AFTOL-ID 428 18S ribosomal RNA	1781	1781	100%	0.0	99%	EF413624.1
T.rubrum DNA for 18S ribosomal RNA, partial	1781	1781	100%	0.0	99%	XS8570.1
Ctenomyces serratus 18S ribosomal RNA gene, partial sequen	1775	1775	100%	0.0	99%	U29391.1
Arthroderma racemosum strain ATCC 38556 18S ribosomal RN	1770	1770	97%	0.0	99%	EF631613.1
Microsporium gallinae strain ATCC 12108 18S ribosomal RNA g	1770	1770	97%	0.0	99%	EF631610.1
Arthroderma cajetani strain ATCC 34662 18S ribosomal RNA ge	1764	1764	97%	0.0	99%	EF631607.1
Arthroderma ciferrii 18S rRNA gene, isolate CBS 272.66, partia	1760	1760	99%	0.0	99%	AB015789.2
Trichophyton longifusum strain ATCC 22397 18S ribosomal RN	1759	1759	97%	0.0	99%	EF631616.1
Arthroderma obtusum strain ATCC 42129 18S ribosomal RNA f	1759	1759	97%	0.0	99%	EF631612.1
Trichophyton rubrum strain CBS 118892 18S ribosomal RNA ge	1755	1755	100%	0.0	98%	X431933.1
Arthroderma uncinatum strain ATCC 28454 18S ribosomal RN	1753	1753	97%	0.0	99%	EF631614.1
Arthroderma gypseum strain ATCC 24102 18S ribosomal RNA f	1753	1753	97%	0.0	99%	EF631611.1
Trichophyton tonsurans 18S ribosomal RNA gene, partial sequ	1751	1751	97%	0.0	99%	AY083229.1
Ctenomyces serratus 18S rRNA gene, isolate CBS 187.61, parti	1749	1749	99%	0.0	99%	AB015771.1
Trichophyton megninii strain ATCC 12106 18S ribosomal RNA j	1748	1748	97%	0.0	99%	EF631617.1
Arthroderma benhamiae strain ATCC 28446 18S ribosomal RN	1748	1748	97%	0.0	99%	EF631615.1
Trichophyton mentagrophytes var. erinacei 18S ribosomal RN	1746	1746	97%	0.0	99%	AY083228.1
Epidermophyton floccosum strain ATCC 52066 18S ribosomal f	1744	1744	97%	0.0	99%	EF631604.1
Trichophyton schoenleinii strain ATCC 22775 18S ribosomal R	1742	1742	97%	0.0	99%	EF631620.1
Microsporium persicolor 18S ribosomal RNA gene, partial sequ	1742	1742	96%	0.0	99%	AY083228.1
Trichophyton verrucosum strain BMU03157 18S ribosomal RN	1736	1736	97%	0.0	99%	EF631623.1
Trichophyton rubrum strain BMU01672 18S ribosomal RNA ge	1736	1736	97%	0.0	99%	EF631621.1

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