

IS000165

CSF

1310bp

AGAAGTCATACGACCAGCCAAAGCCGGCCTGACTATTTAGCAGGTTAAGGTCTCGTTCGTTATCGGAATTAACC
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 GCCCTGTATTGTTATTTATTGTCACCTCCCCGTGTCGGGATTGGGTAATTTGCGCGCTGCTGCCTTCCTTG
 ATGTGGTAGCCGTTTCTCAGGCTCCCTCTCCGGAATCGAACCCTAATTCCCCGTTACCCGTTGATACCATGGTAG
 GCCTCTATCCTACCATCGAAAGTTGATAGGGCAGATATTTGAATGAAGCATCGCCGGCACAAGGCCATGCGATT
 GAGAAGTTATTATGATTACCATAGAGGGCCGAAACCCATTGGTTTTTATCTAATAAATACACCCCTCCAGAA
 GTCGGGGCTTTTCAGCATGTATTAGCTCTAGAATTACCACAGTTATCCATGTAGCAAGATACCATCAAATAAACTA
 TAACTGATTTAATGAGCCATTCGCAGTTTCACAGTATGAAT

<u>Description</u>	<u>Max Score</u>	<u>Total Score</u>	<u>Query Cover</u>	<u>E value</u>	<u>Per. Ident</u>	<u>Accession</u>
Cryptococcus neoformans var. grubii H99 chromosome 02	2420	7260	100%	0.0	100.00%	CP047903.1
Cryptococcus neoformans var. grubii strain Ze90-1 chromosome 2	2420	32441	100%	0.0	100.00%	CP048073.1
Cryptococcus neoformans strain EN28 chromosome 2, complete sequence	2420	2420	100%	0.0	100.00%	CP025718.1
Cryptococcus neoformans var. grubii strain KN99 chromosome 2, complete sequence	2420	2420	100%	0.0	100.00%	CP022322.1
Cryptococcus neoformans var. grubii H99 18S ribosomal RNA (CNAG 10502), rRNA	2420	2420	100%	0.0	100.00%	XR_001045463.1
Cryptococcus neoformans var. grubii H99 chromosome 2, complete sequence	2420	2420	100%	0.0	100.00%	CP003821.1
Cryptococcus neoformans var. neoformans strain CBS 132 18S ribosomal RNA gene, partial sequence	2420	2420	100%	0.0	100.00%	HQ596559.1
Filobasidiella neoformans var. neoformans 5S ribosomal RNA, 17S ribosomal RNA genes, internal transcribed spacer 1, 5.8S ribosomal RNA gene, internal transcribed spacer 2, and 25S ribosomal RNA gene, complete sequence	2420	2420	100%	0.0	100.00%	AF356652.1
Cryptococcus neoformans var. grubii partial 18S rRNA gene, molecular type VNI, serotype A,	2420	2420	100%	0.0	100.00%	AJ560301.1

<u>Description</u>	<u>Max Score</u>	<u>Total Score</u>	<u>Query Cover</u>	<u>E value</u>	<u>Per. Ident</u>	<u>Accession</u>
specimen voucher WM148						
Cryptococcus gattii VGII R265 chromosome 4, complete sequence	2414	7218	100%	0.0	99.92%	CP025762.1