

**Table S2.** Ten Ascomycota fungal genomes with high-quality genome annotations were chosen to annotate the *Ceratobasidium* sp. C02 assembly

Genome	Accession	Reference	Source	Evidence
<i>Laccaria bicolor</i>	Lacbi2 (16097)	<a href="https://doi.org/10.1038/nature06556">https://doi.org/10.1038/nature06556</a>	JGI	<i>ab initio</i> 38,913 ESTs protein homology - UniProtKB fungi
<i>Mycena indigotica</i>	GCF_014461135.1	<a href="https://doi.org/10.1073/pnas.2010761117">https://doi.org/10.1073/pnas.2010761117</a>	NCBI	<i>ab initio</i> RNA sequencing protein homology - UniProtKB fungi
<i>Coprinopsis cinerea</i>	Copci1	<a href="https://doi.org/10.1073/pnas.1003391107">https://doi.org/10.1073/pnas.1003391107</a>	JGI	<i>ab initio</i> 5,612 ESTs
<i>Schizophyllum commune</i>	Schco3	<a href="https://doi.org/10.1038/nbt.1643">https://doi.org/10.1038/nbt.1643</a>	JGI	<i>ab initio</i> 31,000 ESTs
<i>Sparassis crispa</i>	GCF_003851025.1	<a href="https://doi.org/10.1038/s41598-018-34415-6">https://doi.org/10.1038/s41598-018-34415-6</a>	NCBI	<i>ab initio</i> RNA sequencing
<i>Serpula lacrymans</i>	SerlaS7_9_2	<a href="https://doi.org/10.1126/science.1205411">https://doi.org/10.1126/science.1205411</a>	JGI	<i>ab initio</i> undisclosed ESTs protein homology - GenBank nr proteins
<i>Postia placenta</i>	Pospl1	<a href="https://doi.org/10.1073/pnas.0809575106">https://doi.org/10.1073/pnas.0809575106</a>	JGI	<i>ab initio</i> 44,520 ESTs protein homology - 'reliable homology-based gene models'

<i>Dichomitus squalens</i>	Dicsqu463_1	<a href="https://doi.org/10.1128/MRA.00264-19">https://doi.org/10.1128/MRA.00264-19</a>	JGI	<i>ab initio</i> RNA sequencing protein homology - 'homology-based'
<i>Agaricus bisporus</i> var. <i>bisporus</i>	99190	<a href="https://doi.org/10.1073/pnas.1206847109">https://doi.org/10.1073/pnas.1206847109</a>	JGI	<i>ab initio</i> 79,271 ESTs protein homology - GenBank nr proteins

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