

Supplementary data

A Comprehensive Phylogenetic and Bioinformatics Survey of Lectins in the Fungal kingdom

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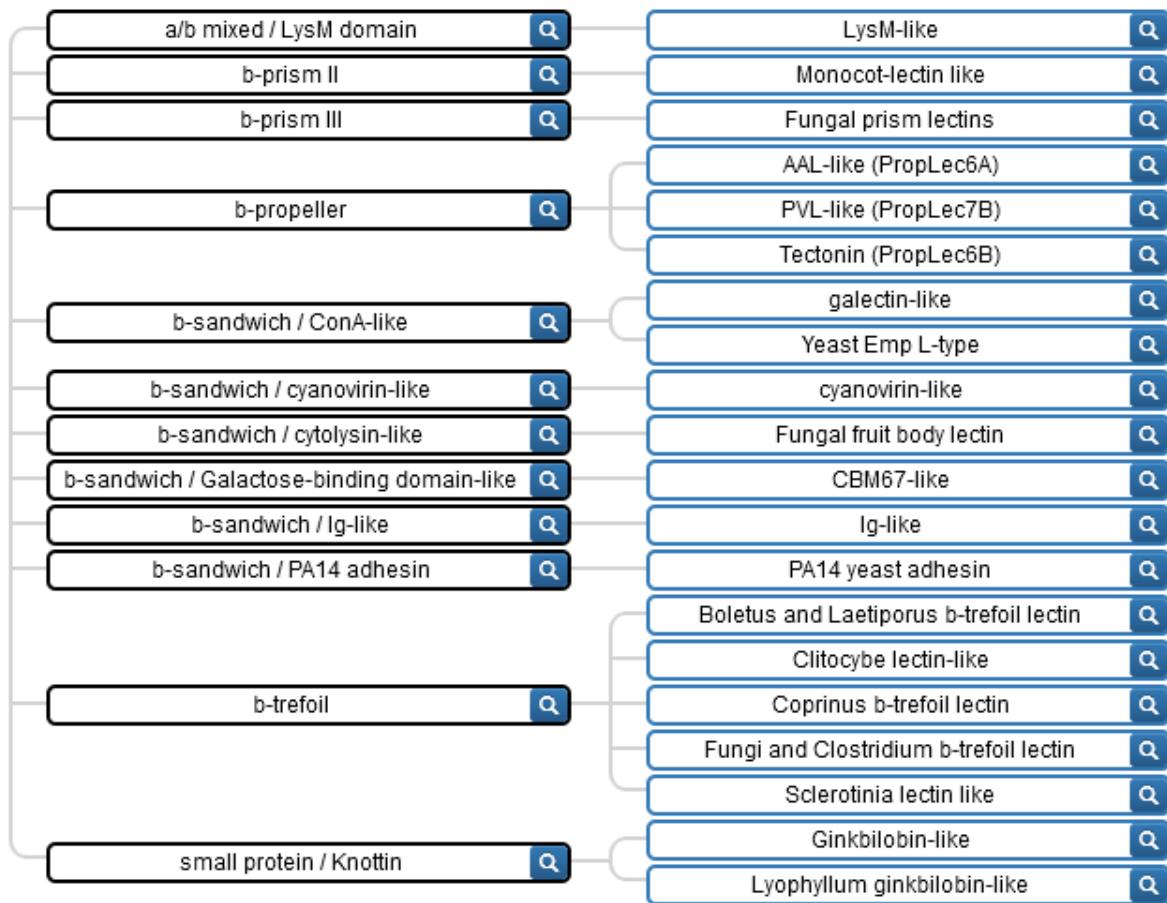


Figure S1: Distribution of lectin folds and classes of fungal lectin with 3D structures in Unilectin3D database.

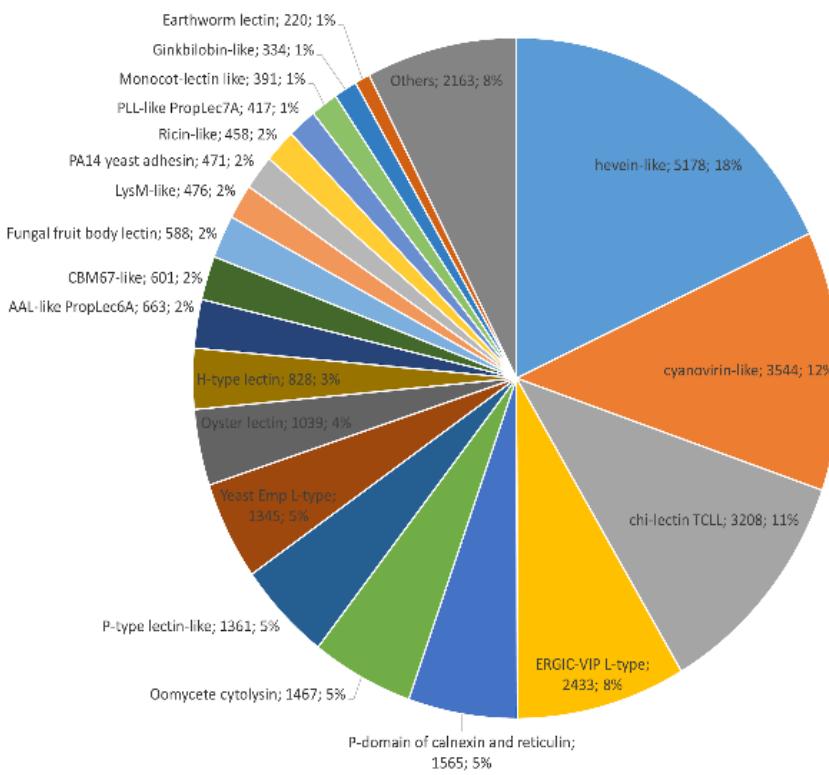
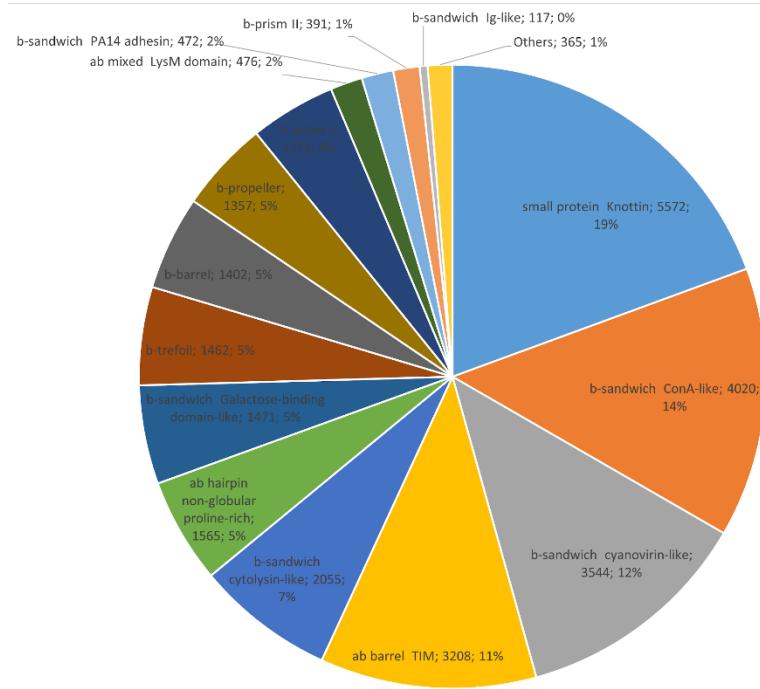


Figure S2: Distribution of folds (left) and classes (right) of predicted lectin sequences in MycoLec. Only lectin sequences with a similarity score > 25% were used.

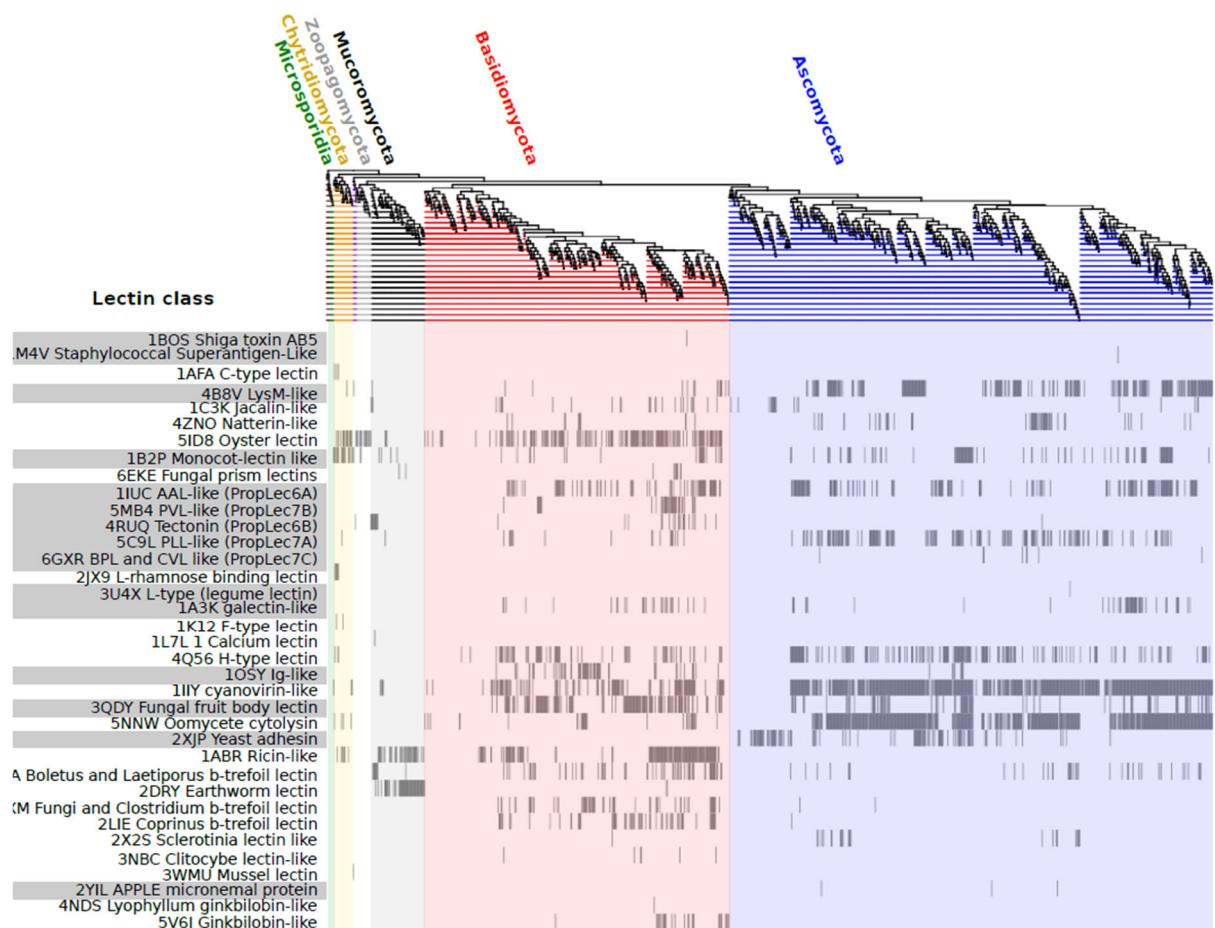


Figure S3: Distribution of predicted lectins by species in MycoLec. Each vertical line corresponds to a fungal strain organized according to their phylogenetic relationship as displayed by the tree. Left, Clustering of lectin classes. Lectins with a similarity score > 25% were used to detect the presence of the different lectin classes found in the MycoCosm genomes.

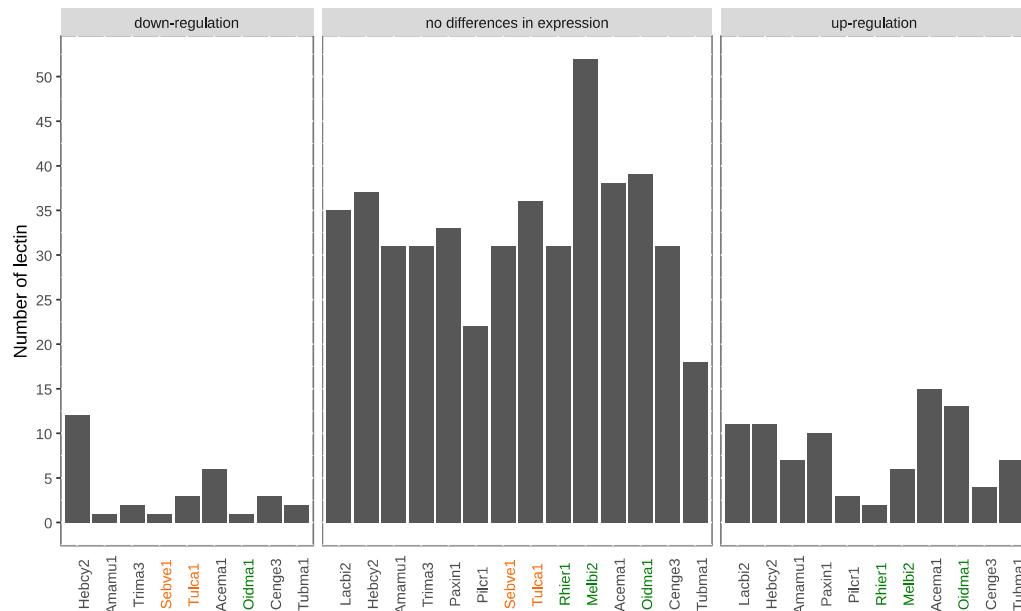


Figure S4: Impact of mycorrhization of 14 fungal strains with their corresponding host plant on lectin expression. Each bar corresponds to a strain annotated by a tag used in the MycoCosm database to refer to the specific strain and genomic assembly. Tags are colored according to the mycorrhizae type: grey ectomycorrhizae, green ericoid mycorrhizae, orange orchid mycorrhizae.

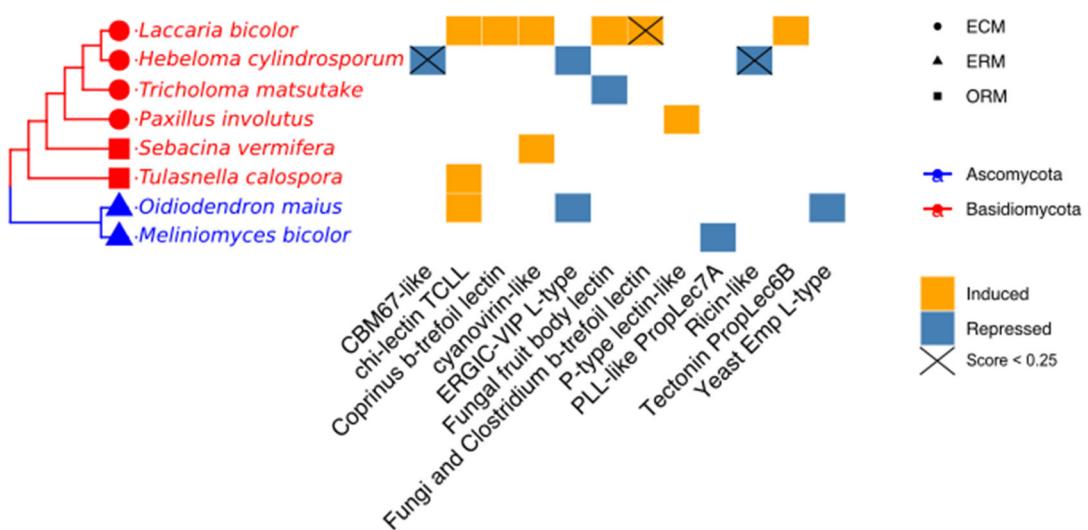


Figure S5: Differential expression of lectins within mycorrhizal fungi upon plants interaction. Lectins with invariable expression are not represented. Three species are ericoid mycorrhizae (ERM), two are orchids mycorrhizae (ORM) and the other are ectomycorrhizae (ECM).

Table S1: Fungal species investigated in the exploration of the lectins induced and repressed during their mycorrhization with a compatible plant host. ECM: ectomycorrhizae, ORM: orchids mycorrhizae, ERM: ericoid mycorrhizae

JGI ID	Fungal species	Plant host	Analysis Method	Mycorrhizae type	Ref
Amamu1	<i>Amanita muscaria</i>	<i>Populus tremula tremolooides</i>	CLC	ECM	[1]
Hebcy2	<i>Hebeloma cylindrosporum</i>	<i>Pinus pinaster</i>	CLC	ECM	[1]
Paxin1	<i>Paxillus involotus</i>	<i>Fagus sylvatica</i>	CLC	ECM	[1]
Pilcr1	<i>Piloderma croeus</i>	<i>Quercus robur</i>	CLC	ECM	[1]
Oidma1	<i>Oidiodendron maius</i>	<i>Vaccinium myrtillus</i>	CLC	ERM	[1]
Sebve1	<i>Sebacina vermifera</i>	<i>Arabidopsis thaliana</i>	CLC	ORM	[1]
Tulca1	<i>Tulasnella calospora</i>	<i>Serapias vomeracea</i>	CLC	ORM	[1]
Melbi2	<i>Melinomyces bicolor</i>	<i>Vaccinium myrtillus</i>	CLC	ERM	[2]
Rhier1	<i>Rhizoscyphus ericaceae</i>	<i>Vaccinium myrtillus</i>	CLC	ERM	[2]
Tubma1	<i>Tuber magnatum</i>	<i>Quercus robur</i>	CLC	ECM	[3]
Cenge3	<i>Cenococcum geophilum</i>	<i>Pinus sylvestris</i>	CLC	ECM	[4]
Acema1	<i>Acephala macrosclerotiorum</i>	<i>Pinus sylvestris</i>	HISAT/ DESeq2	ECM	[5]
Lacbi2	<i>Laccaria bicolor</i>	<i>Populus tremula x alba</i>	HISAT/DESeq2	ECM	[6]
Trima3	<i>Tricholoma matsutake</i>	<i>Pinus sylvestris</i>	CLC	ECM	[5]

Table S2: Lectin content in the predicted proteomes of the Agaricomycetes fungal class sorted by ecological niche.

<i>Coniophora olivacea</i>		1	2	2		1		6	4
<i>Coniophora puteana</i>		1	1			1		3	3
<i>Crucibulum laeve</i>		1	3	3		1		20	7
<i>Cylindrobasidium torrendii</i>	1		1					2	2
<i>Dentipellis sp.</i>			1	3	3	1		8	4
<i>Dichomitus squalens</i>	3		1	1	3	4	1	13	6
<i>Exidia glandulosa</i>			20	2	1	1		31	7
<i>Fibroporia radiculosa</i>			1			1		1	3
<i>Fistulina hepatica</i>			1					1	1
<i>Fomitopsis pinicola</i>			1			1	2	4	3
<i>Galerina marginata</i>	2	2	1	6	1	7	1	1	28
<i>Ganoderma sp.</i>			1		1			3	3
<i>Gloeophyllum trabeum</i>				2				2	1
<i>Heliocybe sulcata</i>			1	1				2	2
<i>Hypholoma sublateritium</i>	1		5	1	3	1	1	14	8
<i>Lentinellus vulpinus</i>	1		1		1	1	3	7	5
<i>Lentinula edodes</i>	1							1	2
<i>Lentinus tigrinus</i>				1		2	1	4	3
<i>Neolentinus lepideus</i>				1				1	1
<i>Obba rivulosa</i>	4	1		1		2		8	4
<i>Omphalotus olearius</i>					1	1		2	2
<i>Peniophora sp.</i>	1		2		2	2	5	12	5
<i>Phanerochaete carnosa</i>		2			6			10	3
<i>Phanerochaete</i>									
<i>chrysosporium</i>		6		1	2	1		10	4
<i>Phlebia brevispora</i>		1		1	1	2	1	19	9
<i>Phlebia centrifuga</i>				3	1	6		10	3
<i>Phlebia radiata</i>		3			1	8	1	32	8
<i>Phlebiopsis gigantea</i>		4		1	1	2	1	9	5
<i>Plicaturopsis crista</i>				1				2	2
<i>Pluteus cervinus</i>	1		38	1	1	1	1	48	8
<i>Polyporus brumalis</i>				1		1	1	1	4
<i>Postia placenta</i>	1			1	3	4	2	23	8
<i>Punctularia strigosozonata</i>				3		1		4	2
<i>Pycnoporus cinnabarinus</i>					1			2	2
<i>Ramaria rubella</i>	1		1	3				2	4
<i>Schizophyllum commune</i>		1	2		1			3	4
<i>Schizopora paradoxa</i>	2			1	6	1	1	1	13
<i>Serpula himantoides</i>			1		3		1	8	4
<i>Stereum hirsutum</i>				1	1	1		3	3
<i>Trametes pubescens</i>						2		2	2
<i>Trametes versicolor</i>						1		2	2
Plant pathogen/Wood decayer								0	
<i>Armillaria cepistipes</i>		2			16	1	4	2	26
<i>Armillaria gallica</i>		3			16	1	3	3	27
<i>Fomitiporia mediterranea</i>				1		2		3	2
<i>Laetiporus sulphureus</i>		1		1	1			5	4
Plant pathogen								0	
<i>Armillaria mellea</i>	1	1			1	3	5	2	14
<i>Armillaria ostoyae</i>		2			16	2	3	2	25
<i>Armillaria solidipes</i>	1	2			17	1	4	1	26
<i>Heterobasidion annosum</i>			1	2				3	2
<i>Moniliophthora perniciosa</i>	1				3			4	2
<i>Rhizoctonia solani</i>	1			8		3	4	3	22
<i>Wolfiporia cocos</i>					1	1		2	2
Overall total	12	8	13	16	3	109	18	34	141
Number of species	4	7	5	8	2	32	5	10	11
% species	4%	7%	5%	7%	2%	30%	5%	9%	10%
	14%	3%	29%	41%	50%	7%	19%	34%	18%
	21%	3%	29%	41%	50%	7%	19%	34%	18%
	23%	3%	29%	41%	50%	7%	19%	34%	18%
	25%	3%	29%	41%	50%	7%	19%	34%	18%
	5%	21%	22%	24%	25%	5%	21%	22%	14%
	13%	5%	14%	15%	16%	5%	13%	14%	15%
	22%	5%	14%	15%	16%	5%	13%	14%	15%
	100%	7%	14%	15%	16%	5%	13%	14%	15%

Table S3 : Details of lectins identified in the genome of *Laccaria bicolor*

Lectin class	#	Mycosm AC (score)	NCBI AC	Protein name
Tectonin PropLec6B	3	Lacbi2:399271 (0.73) Lacbi2:399270 (0.67) Lacbi2:322629 (0.27)	XP_001876432.1	tectonin 2
Coprinus β-trefoil lectin	3	Lacbi2:330799 (0.48) Lacbi2:327918 (0.42) Lacbi2:691792 (0.25)	XP_001885184.1	predicted protein [<i>Laccaria bicolor</i> S238N-H82]
Galectin like	2	Lacbi2:236913 (0.36) Lacbi2:312069 (0.35)	XP_001883510.1	galectin [<i>Laccaria bicolor</i> S238N-H82]
Physarium lectin	2	Lacbi2:381649 (0.39) Lacbi2:322629 (0.39)	XP_001875654.1	ricin-containing lipase tectonin-like
Oyster lectin	2	Lacbi2:585014 (0.28) Lacbi2:448672 (0.27)	XP_001880964.1	predicted protein [<i>Laccaria bicolor</i> S238N-H82]
Fungal fruit body lectin	1	Lacbi2:185716 (0.54)	XP_001885326.1	predicted protein, partial
Boletus and Laetiporus β-trefoil lectin	1	Lacbi2:318163 (0.30)	XP_001879265.1	predicted protein [<i>Laccaria bicolor</i> S238N-H82]
Cyanovirin like	1	Lacbi2:327824 (0.42)	XP_001881773.1	predicted protein [<i>Laccaria bicolor</i> S238N-H82]
P domain of calnexin and reticulin	1	Lacbi2:399410 (0.50)	XP_001874124.1	calnexin [<i>Laccaria bicolor</i> S238N-H82]
Ergic vip L type	1	Lacbi2:399414 (0.48)	XP_001888824.1	ERGIC53, mannose lectin
P-type lectin like	1	Lacbi2:642707 (0.29)	XP_001874815.1	predicted protein [<i>Laccaria bicolor</i> S238N-H82]
PVL like PropLec7B	1	Lacbi2:692684 (0.67)	XP_001891161.1	predicted protein, partial

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