

State of LA Fungi

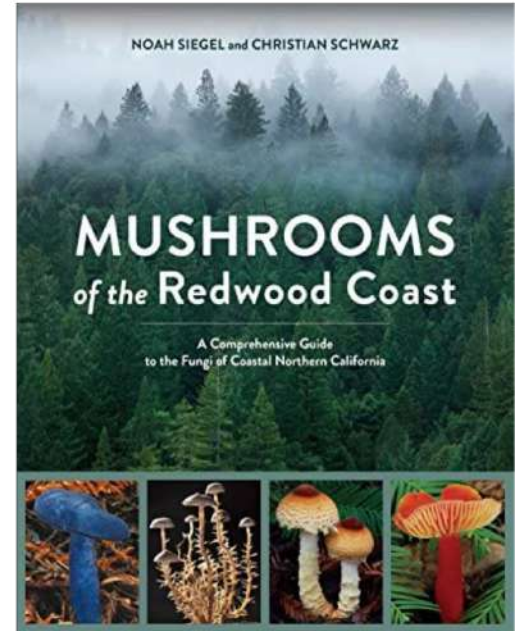
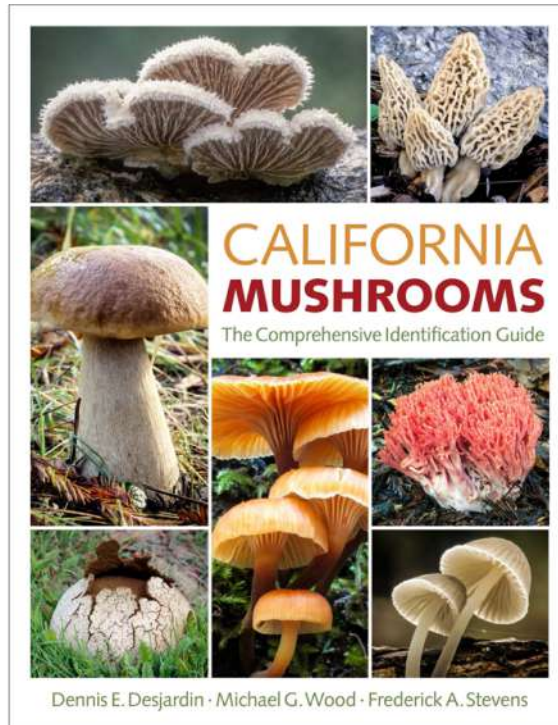
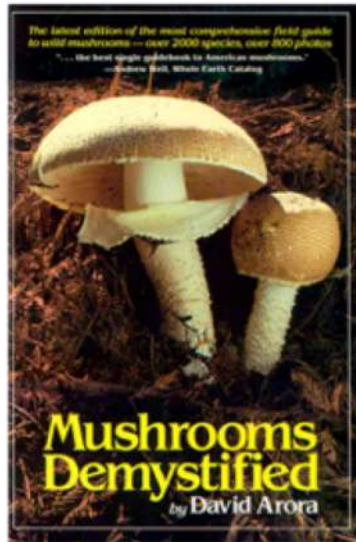
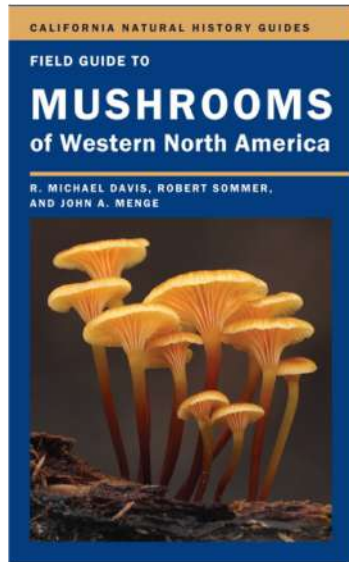
March 28, 2023



Chlorophyllum rhacodes

Los Angeles Mycological Society

- I. Look back on March
- II. Reproduction and mating types in Basidiomycota
- III. Mushrooms in the chaparral
- IV. Evolution and ecology of psilocybin in nature





Southern California

7,819

OBSERVATIONS

578

SPECIES

300

IDENTIFIERS

1,588

OBSERVERS



233 observations

CC

Common Fieldcap

(Agaricus pinivorus)

171 observations

CC

Hairy Curtain Crust

(Stereum hirsutum)

132 observations

CC

Splitgill Mushroom

(Schizophyllum commune)

112 observations

CC

Oak-loving Elf Saddle

(Hebelia dryophila)

106 observations

CC

Yellow Fieldcap

(Bolbitis platensis)

101 observations

CC

Blewit

(Lepista nuda)

92 observations

CC

Mushrooms

(Larariumycia pinnatifida)

86 observations

CC

Stubble Rosegill

(Volvotruncus glaucophyllus)

86 observations

CC

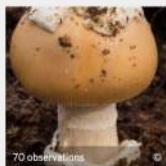
Pale Brittistern

(Cantharellus cantharellus)

82 observations

CC

Golden Ear

(Naematelia aurantia)

70 observations

CC

Springtime Amanita

(Amanita vesia)

60 observations

CC

Turkey-Tail

(Trametes versicolor)

55 observations

CC

Western Destroying A...

(Amanita ocreata)

53 observations

CC

Crystal Brain Fungus

(Myrmecium nucitatum)

51 observations

CC

Deer Mushroom

(Pluteus cervinus)

49 observations

CC

Black Witches' Butter

(Exidia glandulosa)

48 observations

CC

Flocculose Inkcap

(Coprinella flocculosa)

48 observations

CC

Giraffe Spots

(Pezizopora alluobadia)

47 observations

CC

Hare's Foot Inkcap

(Coprinopsis lagopus)

47 observations

CC

Desert Shaggymane

(Phollex pilularis)

https://www.inaturalist.org/observations?d1=2023-03-01&d2=2023-03-28&place_id=51727&taxon_id=47170&view=species

iNaturalist: Most Common Fungal Species in SoCal, March 2023



Southern California

4,181

OBSERVATIONS

492

SPECIES

283

IDENTIFIERS

1,011

OBSERVERS



122 observations

CC

Stubble Rosegill
(*Vivipholus glaucophyllus*)

83 observations

CC

Splitgill Mushroom
(*Schizophyllum commune*)

81 observations

CC

Hairy Curtain Crust
(*Shiitake pinastri*)

61 observations

CC

Oak-loving Elf Saddle
(*Chelvelia dryophila*)

64 observations

CC

Western Jack-O'-Lantern
(*Clathrellus oliveosus*)

51 observations

CC

Turkey-Tail
(*Trametes versicolor*)

51 observations

CC

Rosy Navel
(*Cantharellus roseus*)

48 observations

CC

Golden Ear
(*Naematolus aurantia*)

46 observations

CC

Blewit
(*Lepista nuda*)

44 observations

CC

Springtime Amanita
(*Amanita vespa*)

40 observations

CC

Desert Shaggymane
(*Peziza pilosella*)

37 observations

CC

Western Destroying A...
(*Amanita ocreata*)

34 observations

CC

Yellow Cobblestone Li...
(*Acanthopora socialis*)

33 observations

CC

Rufous Candy Cap
(*Lactarius rufus*)

32 observations

CC

Giraffe Spots
(*Peniophora alboadusta*)

30 observations

CC

Blushing Bride Amanita
(*Amanita rostrata*)

26 observations

CC

Field Bird's Nest Fungus
(*Cyathus olivaceus*)

26 observations

CC

Stork's-bill Chytrid
(*Synchytrium papillatum*)

24 observations

CC

Yellow Fieldcap
(*Boletus edulis*)

23 observations

CC

Browwit
(*Clitocybe brunneocapitata*)

https://www.inaturalist.org/observations?d1=2023-03-01&d2=2023-03-28&place_id=51727&taxon_id=47170&view=species

iNaturalist: Most Common Fungal Species in SoCal, February 2023

Fungal reproduction.

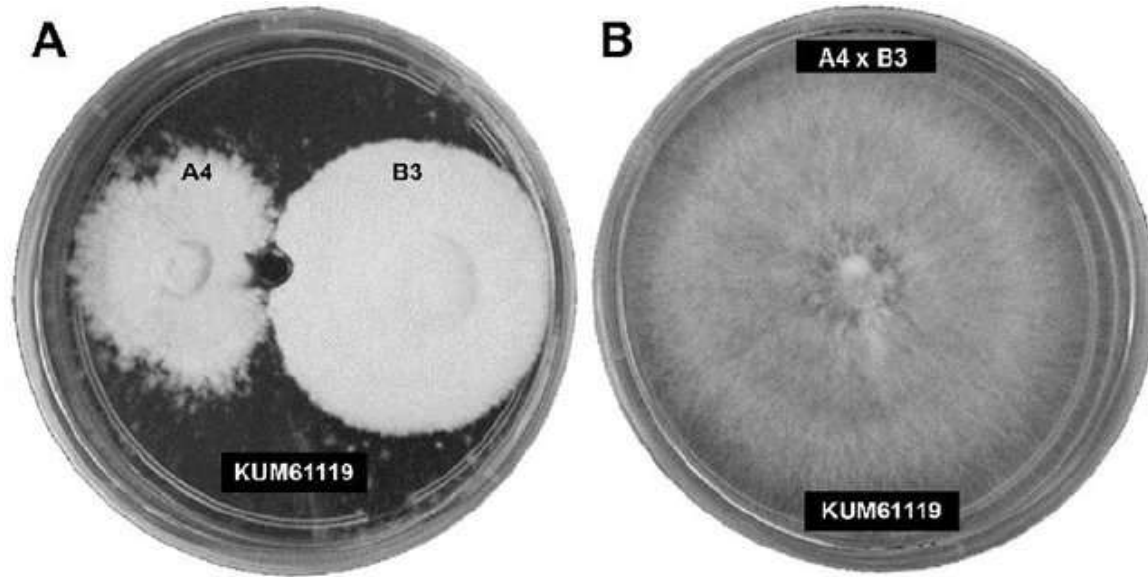


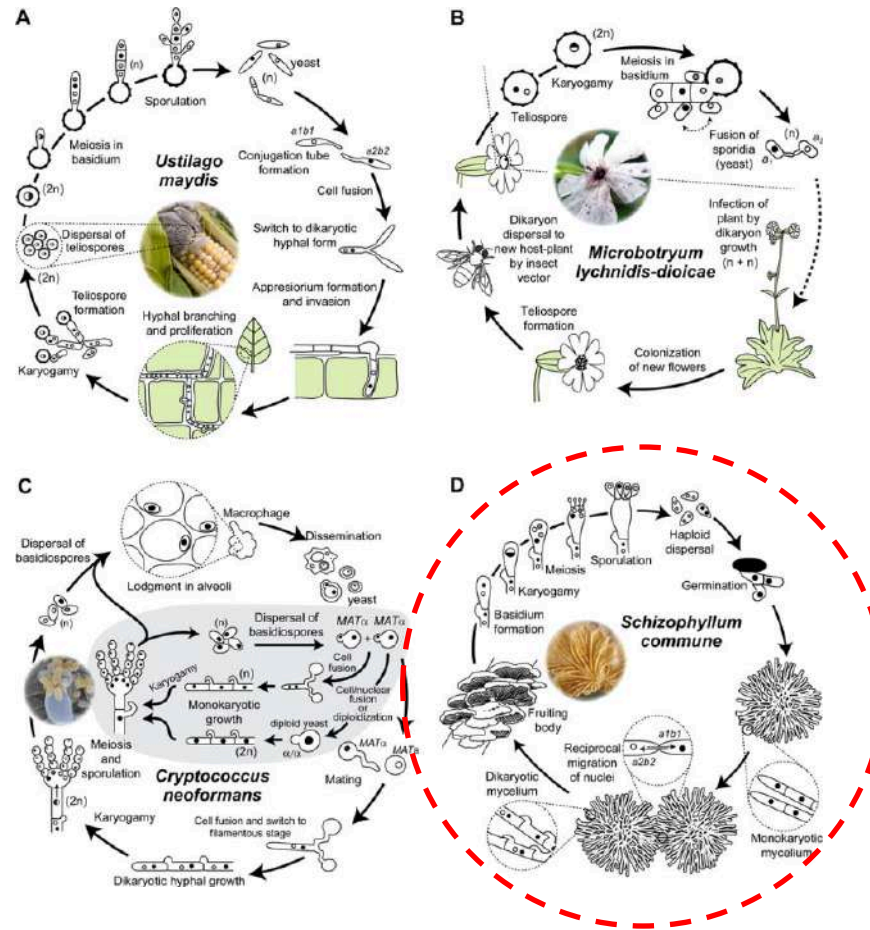
Fig. 1: Cross compatibility test. (A) A compatible mating between two selected monokaryotic isolates of A4 and B3 (obtained from KUM61119 strain). (B) Successfully crossed strain (dikaryotic). Avin et al., 2014.

Diversity in mating system complexity *just* in Basidiomycota

<https://en.wikipedia.org/wiki/Basidiomycota>

Mating in other fungal lineages:

https://en.wikipedia.org/wiki/Mating_in_fungi



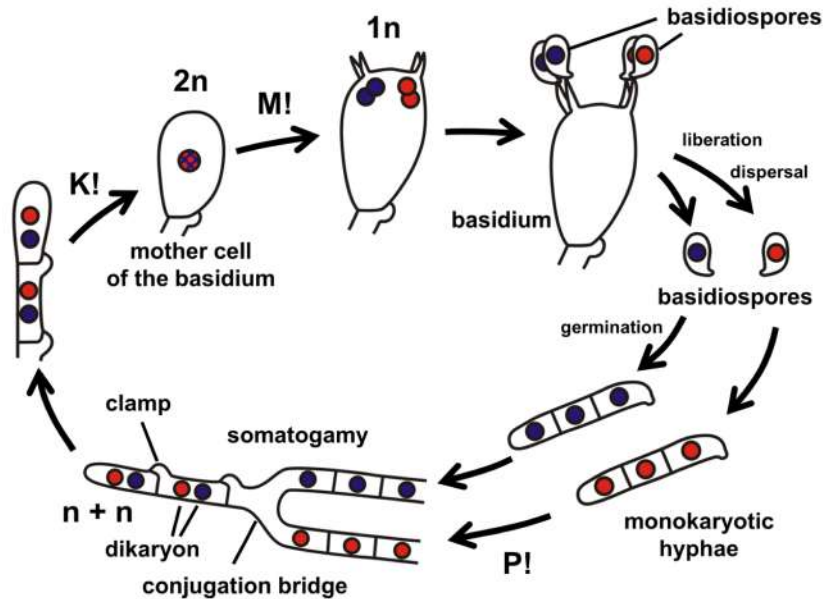
Most common system in mushroom-forming fungi.

Figure: Coelho MA, Bakkeren G, Sun S, Hood ME, Giraud T. Fungal Sex: The Basidiomycota. Microbiol Spectr. 2017 Jun;5(3):10.

<https://journals.asm.org/doi/10.1128/microbiolspec.FUNK-0046-2016>

Basidiomycete Mating Types

David Bermudes



© M. Piepenbring, CC BY-SA



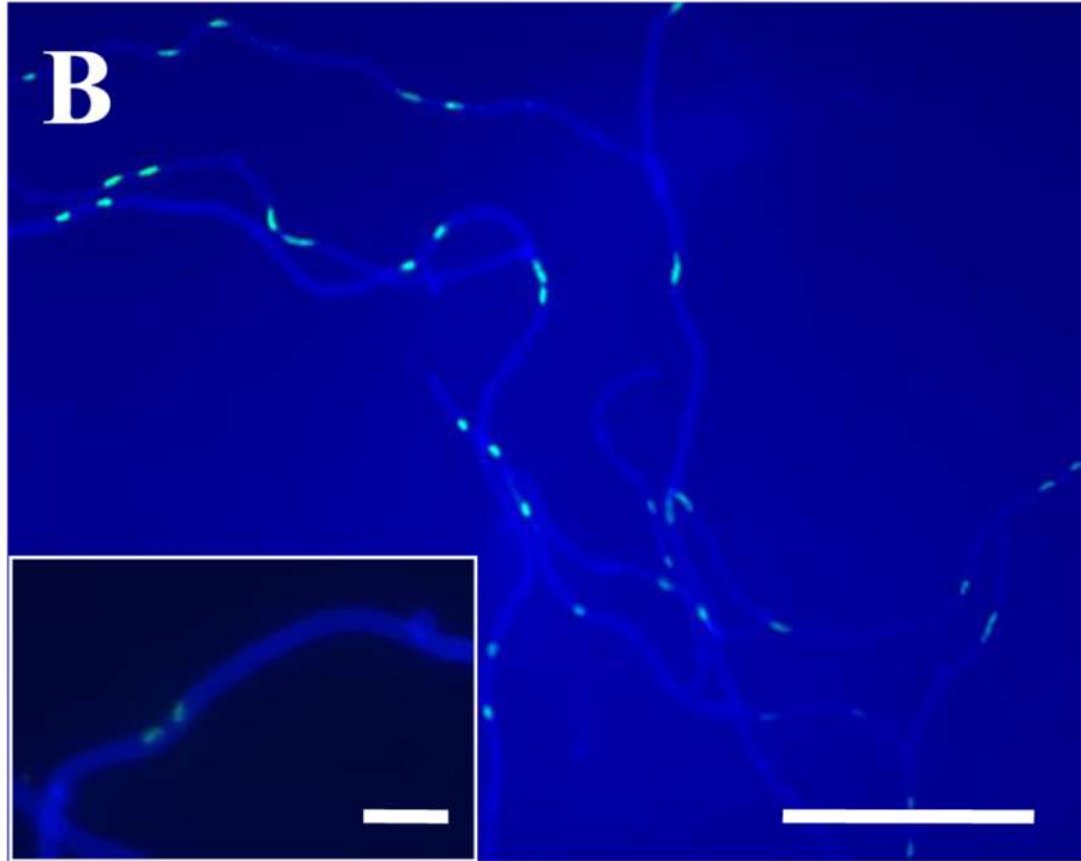
Omphalotus olivascens

https://bio.libretexts.org/Bookshelves/Botany/A_Photographic_Atlas_for_Botany_%28Morrow%29/03%3A_Fungi_and_Lichens/3.06%3A_Basidiomycota_%28Club_Fungi%29/3.6.03%3A_Life_Cycles_of_Basidiomycetes

Kwoniella mangrovensis

Haploid nuclei – alone but together.

The mycelium of a mushroom that has mated spends its life with two haploid nuclei that are paired within the same septal division (genetically referred to as haploid dikaryotic or $n + n$)



Guerreiro et al., 2013
Eukaryotic Cell
 12: 5 746-760

How does a basidiomycete go from haploid to haploid dikaryotic?

Several functions need to be performed to become a dikaryon:

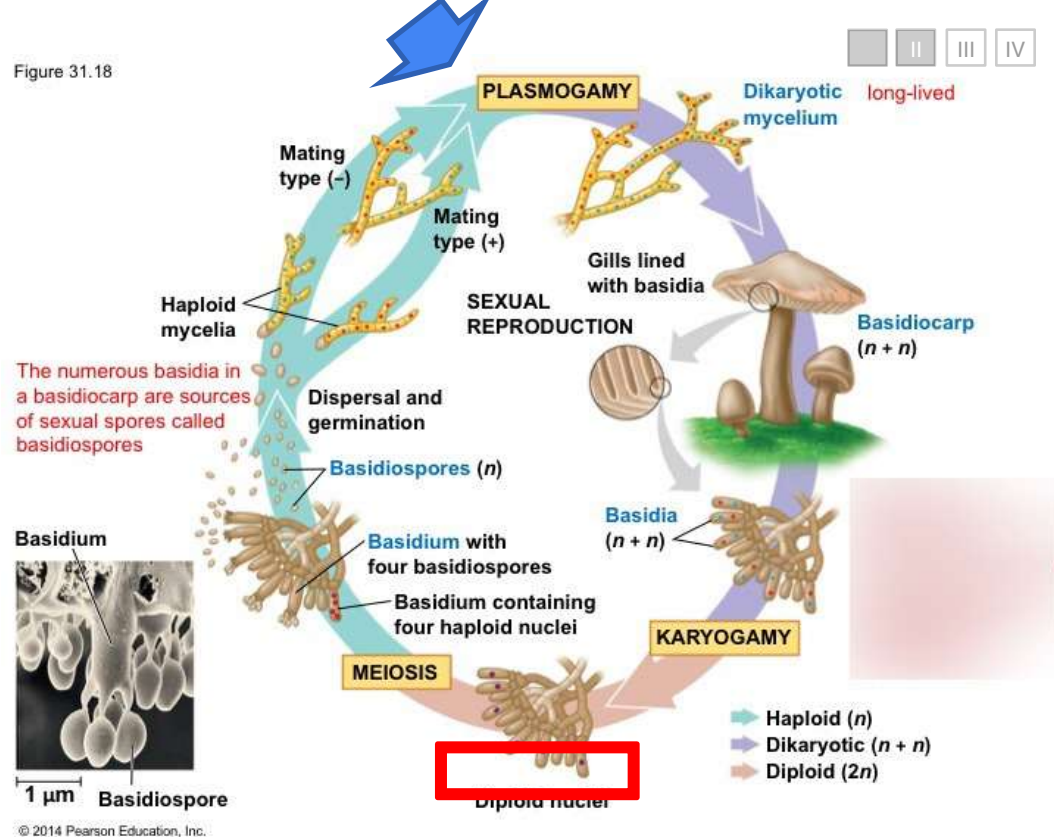
Haploids need to sense each other and grow toward each other (pheromone production and pheromone receptors)

Cell wall dissolution at the tips needs to occur followed by cell fusion (plasmogamy)

Nuclei of each haploid need to migrate to the opposing cell

Clamp connections (many but not all basidiomycetes) need to form that are also involved with septal dissolution and nuclear migration

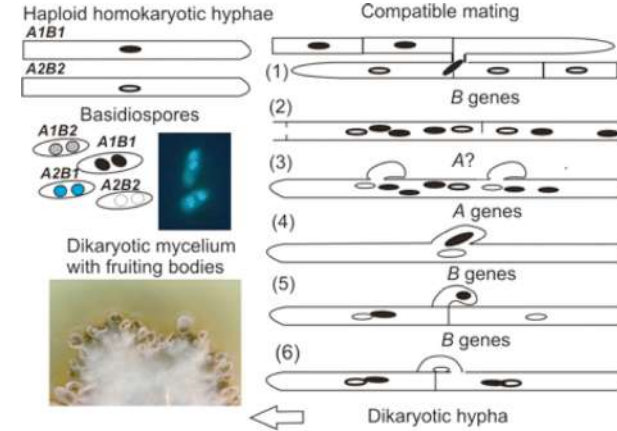
Figure 31.18



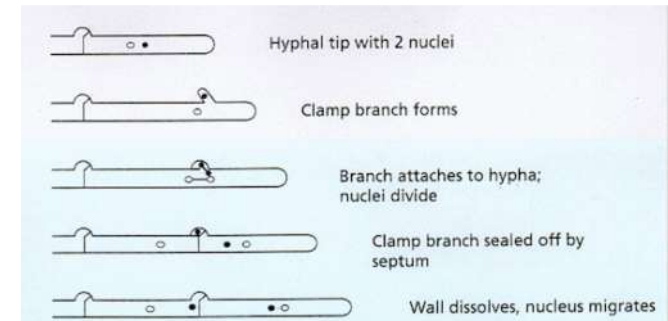
<https://quizlet.com/287338501/basidiomycete-mushroom-forming-life-cycle-diagram/>

Ascomycetes and Basidiomycetes (The Dikarya)

- 90% are heterothallic
- Many are **tetrapolar**, having two unlinked mating loci
 - E.g., *A1 A2, B1 B2*
- Mating compatibility requires different specificities in the functions relating to the process of dikaryonization
- Dikaryonization requires the entire set (*A1 B1 A2 B2*) which the individual haploids don't have.
- This is a process of complementation that results in a complete set of genes required for the process. The loci are often referred to as "opposite", but complementary is more accurate
 - E.g., two haploids that are complementarity are *A1 B1* and *A2 B2*
 - Thought to promote outcrossing
 - "A" genes are involved in clamp initiation and "B" genes are involved in the completion of clamp connections. Some of the "A" genes are signal transduction factors.
 - "B" genes are involved in the initial recognition through pheromones and pheromone receptors. "B" genes regulate the reciprocal nuclear migration, illustrated by movement of a nucleus via a hyphal fusion through broken septa from one haploid hypha into the other.



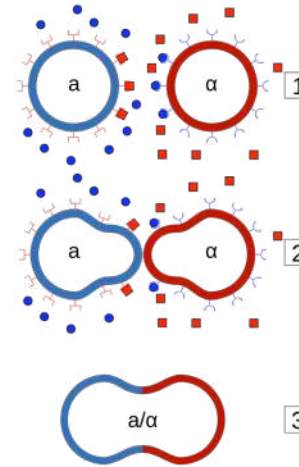
[Eukaryot Cell. 2010 Jun; 9\(6\): 847–859.](http://www.nature.com/naturecell/2010/0601)



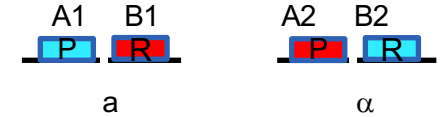
<http://archive.bio.ed.ac.uk/jdeacon/microbes/basidio.htm>
<https://www.youtube.com/watch?v=FTtQtYZFBow>

Hormonal (Pheromone) Control of Mating

- Mating (cell fusion) is dictated by mating types that are controlled by:
 - 1) Cell surface receptors
 - 2) Pheromones
 - 3) Transcription factors
 - Cells must be of compatible (sometimes referred to as opposite) mating types
 - E.g., a and α
 - (arbitrary designations)



What does this look like genetically?

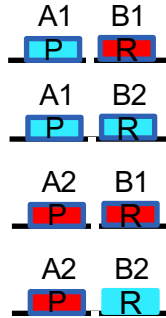


The two different cells have two different sets of mating type genes on two different chromosomes

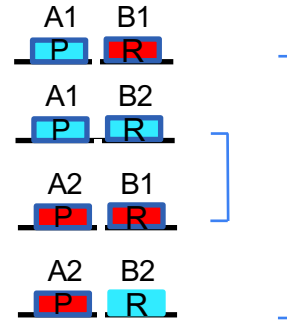
In yeast, type a produces a pheromone ● that is specific for the receptor on α type. Type α produces a pheromone ■ that is specific for the receptor on a type. There are also two different receptor types 𐀀 and 𐀁 that are specific for the pheromones.

<https://www.youtube.com/watch?v=Qy2HPA3W9JM>

What are all the possibilities
from spore production in
basidia?



Which ones are
compatible?



Each one is only
compatible with 1 out
of 4?

Why Are “Opposite” (Complementary) Loci Required?

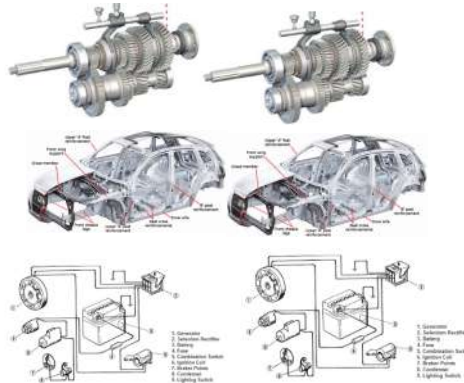
Analogy: Working Together to Build a Car

- Requirements include:
 - List 1 – List 2
 - Wheels • Transmission
 - Engine • Chassis
 - Fuel • Electrical

Two people with
the same list 1



Two people with
the same list 2



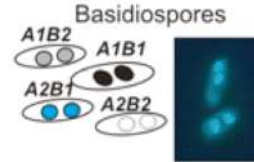
Two people with
List 1 and 2



This one is the
only functional
one

MAT Loci Control of Mating in Fungi: Effects Observable at the Macroscopic and Microscopic Levels

- The roles of combined (complementary) A and B genes
- Hyphae may fuse, but the process of creating a dikaryon is incomplete
- This can be observed on a petri plate and with a microscope



Pairing of strains with:	Events observed
Different A, different B idiomorphs (dikaryon)	1. Septal dissolution
Fertile	2. Nuclear migration
	3. Clamp branches arise and fuse with hypha
Common A, different B idiomorphs	1. Septa dissolve
Flat	2. Nuclei migrate
Common B, different A idiomorphs	1. Septa remain intact
Barrage	2. No nuclear migration
	3. Clamp branches arise but do not fuse
Common A, Common B idiomorphs	1. No septal dissolution
Overlap	2. No nuclear migration
	3. No clamp connections

Deacon 2006, Fungal Biology, Table 5.2.

Observations on *Omphalotus olivascens*



Arbitrarily: A1 B1 x A2 B2

This picture shows a progression of a mating between *Omphalotus olivascens* that produced clamp connections and septa. Mycelial contact zone was robust. Initially only the colony on the left was pigmented, but after enough growth both sides turned out to be highly pigmented. Pictures taken about two weeks apart

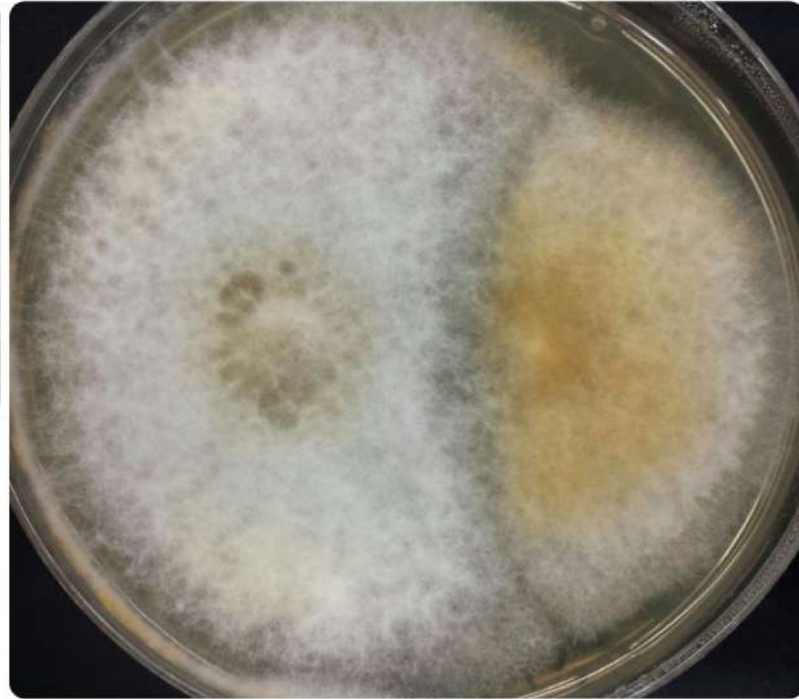
Photo courtesy of Brenda Whitney,
BIOL437/L/492V Spring 2016

Observations on *Omphalotus olivascens*



A1 B1 x A1 B2

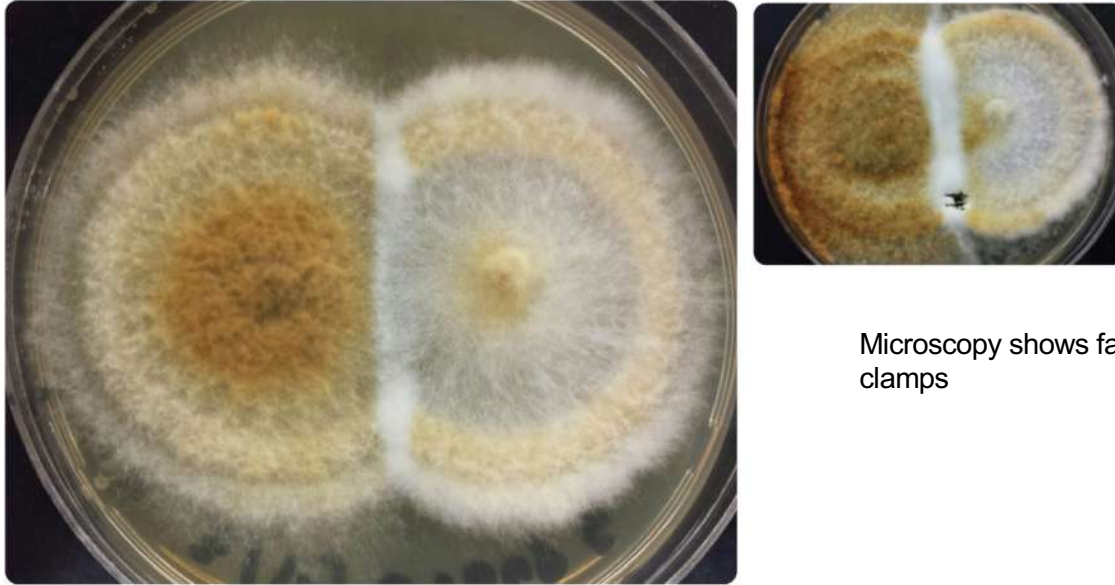
(same A different B)



Two weeks progression of mating type growth that produced a flat reaction.

Photo courtesy of Brenda Whitney,
BIOL437/L/492V Spring 2016

Observations on *Omphalotus olivascens*



Microscopy shows false
clamps

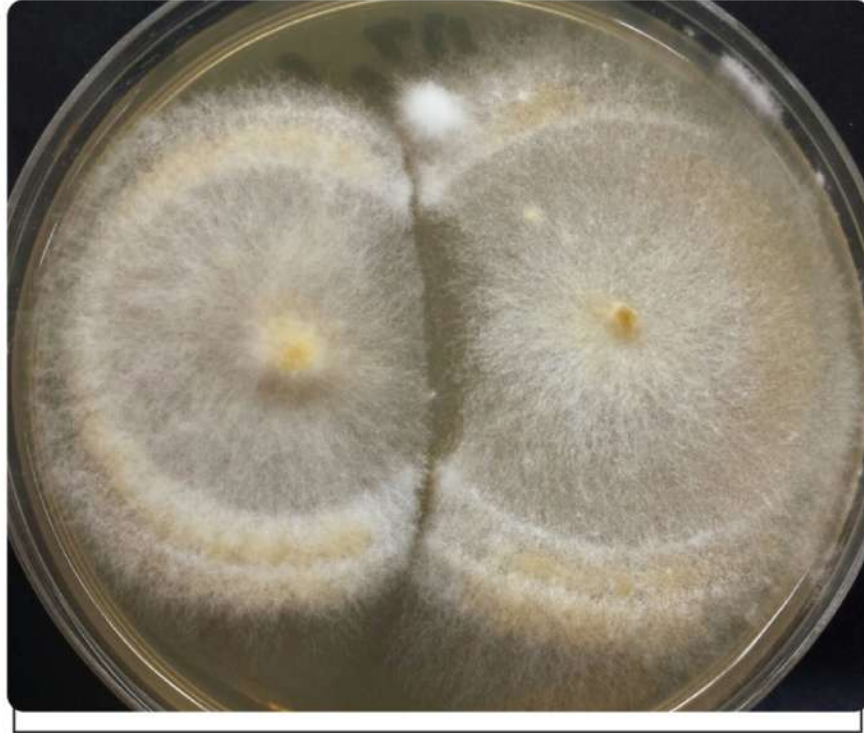
A1 B1 x A2 B1 (same B)

This photo shows a two week progression between mating types which produced false clamps. Mycelial contact zone is barrage.

Observations on *Omphalotus olivascens*



A1 B1 x A1 B1 (same A same B)
"Overlap"
No clamp connections
No septal dilution
No nuclear migration



Mating type growth between same mating types, two week progress

Tetrapolar fungal mating types: Sexes by the thousands

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Received 28 August 1995; accepted 16 January 1996

THE MAGAZINE SHOP

DISCOVER

LOGIN

REGISTER

STAY
CURIOUS

THE SCIENCES | MIND | TECHNOLOGY | HEALTH | ENVIRONMENT | PLANET EARTH

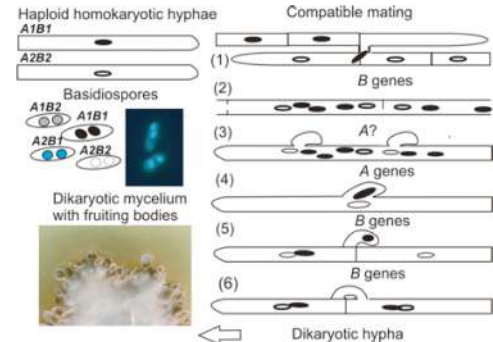
PLANET EARTH

Why This Fungus Has Over 20,000 Sexes

D-brief | By Nathaniel Scharping | Nov 6, 2017 1:23 PM



-
- The diagram illustrates the signaling pathway for hyphal fusion in yeast. It begins with pheromone binding to a GPCR (Receptor), which activates G proteins (α and βγ subunits). This leads to the activation of MAPK, Ras, and Rho/Cdc42. MAPK activates cAMP, which activates Pka. Ras also activates Pka. Rho/Cdc42 promotes cytoskeletal organization. These factors then lead to nuclear migration, which is coupled with septal dissolution and hyphal fusion.



Higher numbers of mating types is due to variations in some of the individual genes. Example, Y and Z signal transduction proteins

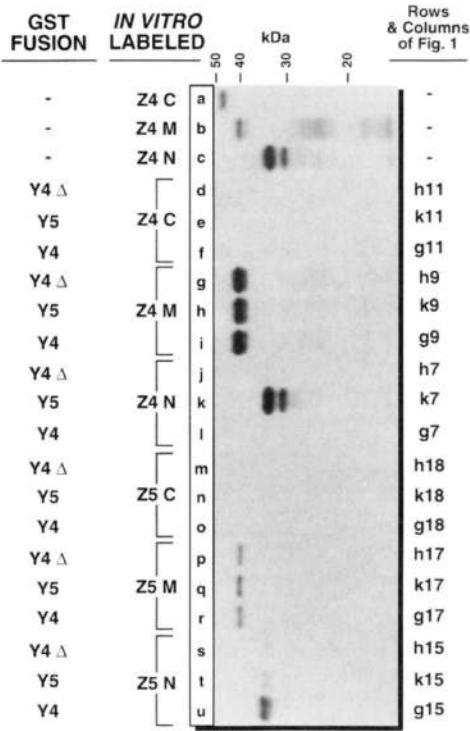
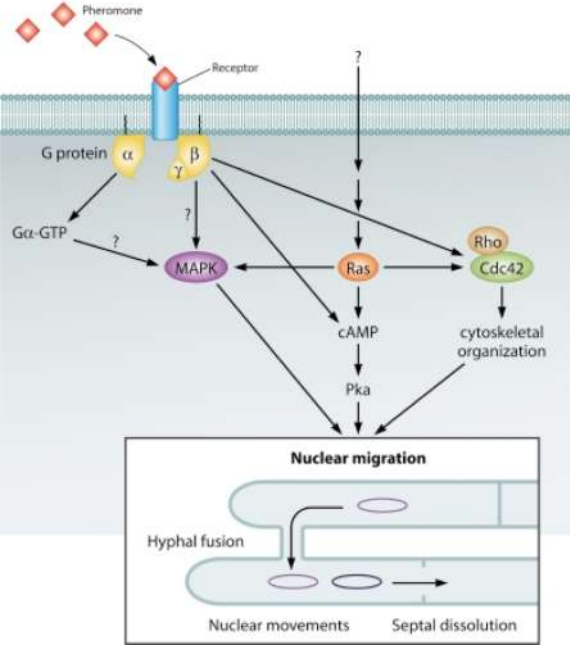



FIGURE 3.—Autoradiogram of electrophoretic gels displaying the results of protein affinity assays between full-length GST-fusion proteins Y4, Y5 and Y4_{ΔFL 1-9}, 81,928 and *in vitro* translated proteins: Z5N, N-terminal amino acids 1–289; Z5M, mid-region amino acids 288–494; Z5C, C-terminal amino acids 493–755; Z4N, N-terminal amino acids 1–271; Z4M, mid-region amino acids 270–544; Z4C, C-terminal amino acids 543–930. Procedures as described for Figure 2.

BIPOLAR MATING TYPES

zygomycetes


 alternative pathways for biosynthesis of trisporic acid leads to physiological complementation

ascomycetes

Saccharomyces cerevisiae

mating type a encodes a1 and a2
 mating type α encodes a1 and a2

diploids form a1/a2 heterodimer which is an active homeodomain transcription factor

mating type switching is seen

Neurospora crassa

mating type A *mtA-1* induces A pheromone and a-factor receptor
 mating type a *mta-1* induces a pheromone and A-receptor

the mating type loci are ideomorphic and show no homology

basidiomycetes

Ustilago hordei

linkage of independent a and b loci lead to a phenotypically bipolar mating behaviour

TETRAPOLAR MATING TYPES

basidiomycetes

Ustilago maydis

bi-allelic a locus
 a1
 a2

each locus encodes a pheromone and a pheromone receptor for the other pheromone

receptor activation leads to cell fusion

multiallelic b locus

b1
 .
 b30
 .
 .

homeodomain transcription factors *b_{East}* and *b_{West}* form active heterodimers if b loci of mates differ; heterodimers induce filamentous growth

Coprinus cinereus

multiallelic A

Aa1-1 ...
 Ab1-1 ...
 Ac1-1 ...
 Ad1-1 ...

archaetype with 4 pairs 2of homeodomain transcription factors; active heterodimers induce pseudoclamp formation

multiallelic B

multiple sets of pheromones and receptors; activation essential for fully compatible mating and spore formation

Schizophyllum commune

multiallelic A

Aa1 ... 9
 Ab1 ... 32

each locus encodes a pair of homeodomain transcription factors; active heterodimers induce clamp cell formation

multiallelic B

Ba1 ... 9
 Bb1 ... 9

pheromone receptors and multiple pheromones; receptor activation leads to nuclear migration and fully compatible mating

Based on the subvariants, there are a large number of theoretical sexes. Some of the variations such as having multiple pheromones and multiple pheromone receptors expand compatibility, while others such as incompatible within signal transduction factors such as Y and Z diminish compatibility.

Fig. 1. Mating systems in higher fungi. Bipolar and tetrapolar mating type systems of some fungi are listed. The function of the mating type genes is indicated.

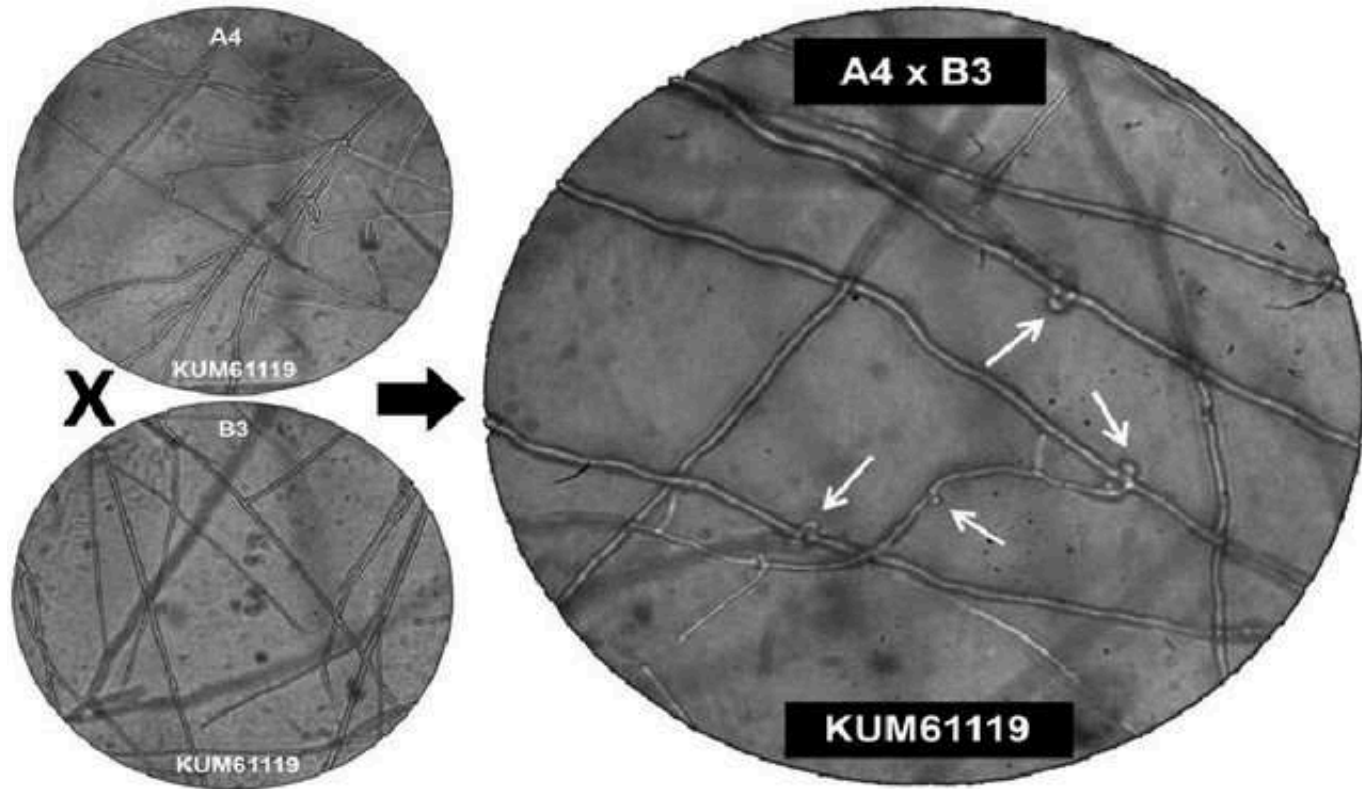


Fig. 2: Detection of compatibility by mating test. The crosses were scored as compatible if clamp connections observed. Avin et al., 2014.

Evolutionary forces in mating type multiplicity:

Gene duplication – Sequence divergence – Chromosome rearrangements



Functional integration

How could it be the case?

- Fungi have an extreme capacity for tolerating internal and external (genetic and environmental) disturbance
- After it has appeared, redundant genetic features could become linked with novel genetic variation that fulfills important life history and ecological functions.

Recent conditions for mushrooms in the chaparral.



Arrhenia chlorocyanea
Hygrophoraceae, Agaricales

Common moss mushroom look-alikes.



Contunyces rosellus
Hygrophoraceae, Agaricales

San Gabriel Mts. March 2023. SP <https://www.inaturalist.org/observations/150748901>



***Omphalina* sp.**
Hygrophoraceae, Agaricales

Murrieta, CA. March 2023. RD <https://www.inaturalist.org/observations/149713015>



Temecula, CA. 17 March 2023 SP



Pleuroflammula
Crepidotaceae, Agaricales
On wood of *Artemisia*.





Calocybe onychia
Lyophyllaceae, Agaricales



Entoloma* subg. *Cyanula
Entolomataceae, Agaricales





***Cystolepiota* sp.**
Verrucosporaceae, Agaricales

Under *Eriogonum fasciculatum*.



Lepiota cf. boudieri
Verrucosporaceae, Agaricales





Cortinarius (Telamonia)
Cortinariaceae, Agaricales

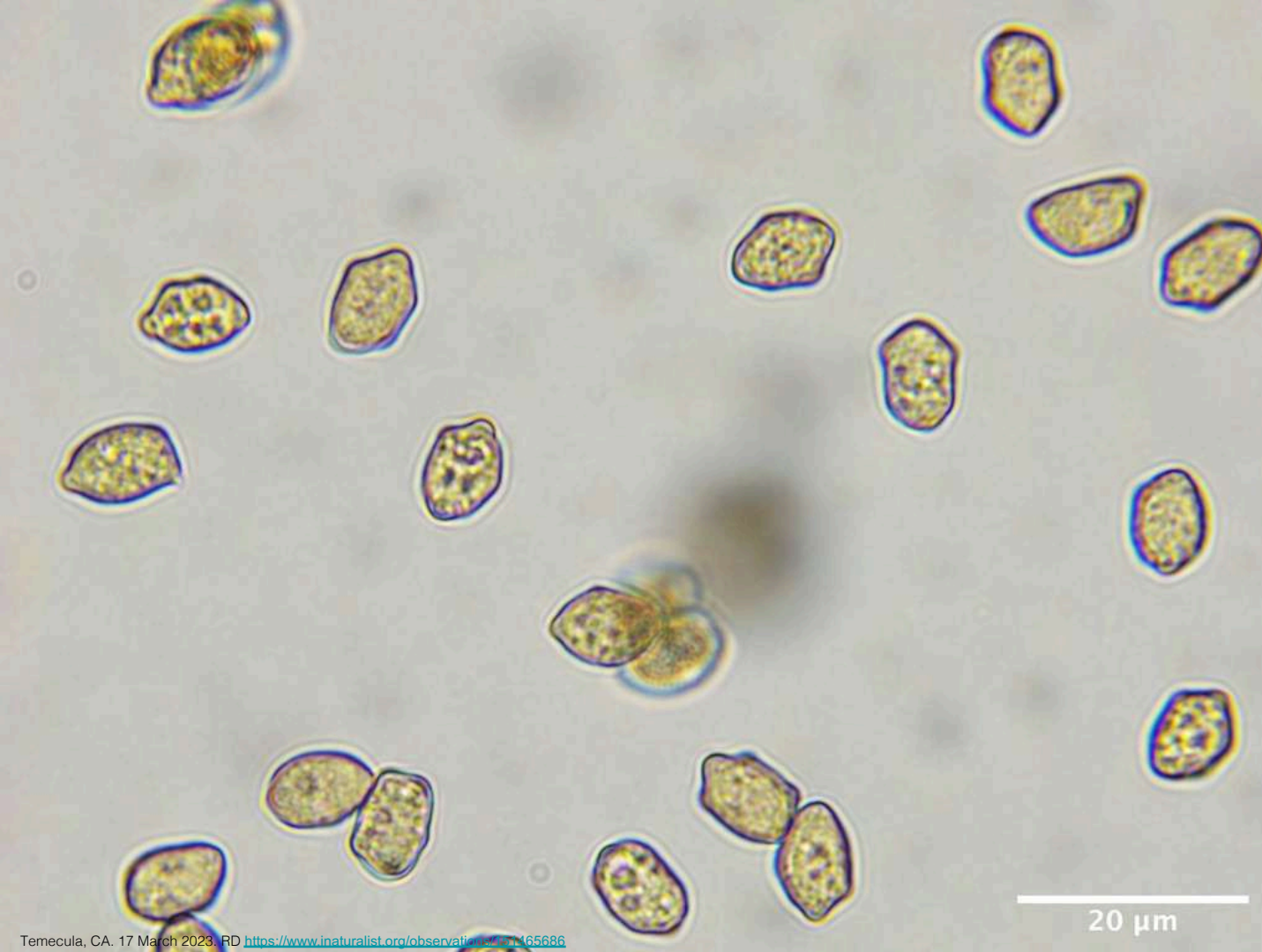
Under *Adenostoma fasciculatum* (chamise).



***Inocybe* sp.**
Inocybaceae, Agaricales

Under scrub oak.





20 μm



***Balsamia* sp.**
 Helvellaceae, Pezizales

Under scrub oak.





***Melanogaster* sp.**
 Melanogastraceae, Boletales

NOT POOP.

III IV

Melanogaster sp.
Melanogastraceae, Boletales
Under scrub oak.





Synchytrium papillatum
Chytridiomycota
Parasite on *Erodium*.

Psilocybin-producing fungi



Panaeolus cinctulus, RD



Gymnopilus spp. https://en.wikipedia.org/wiki/Gymnopilus_luteofolius



Inocybe corydalina

Alan Rockefeller <https://www.inaturalist.org/observations/4675516>



Pluteus americanus https://en.wikipedia.org/wiki/Pluteus_americanus#/media/File:Pluteus_americanus.jpg



Massospora spp.

<https://www.sciencedirect.com/science/article/pii/S1754504819300352>



Psilocybe spp.

Psilocybe species diversity. Basidiomata of (A) *Psilocybe serbica* (jonagraska, CC-BY-SA 3.0), (B) *P. mescaleñoensis* (Alan Rockefeller, CC BY-SA 4.0), (C) *P. cubensis* (Ricardo Arredondo, CC BY-NC), (D) *P. ovoideocystidiata* (Shroomydan, CC-BY-SA 3.0), (E) *P. allenii* (Alan Rockefeller, CC-BY-SA 3.0), (F) *P. azurescens* (Shroom360, CC BY-SA 3.0), (G) *P. cyanescens* (Alan Rockefeller, CC BY-SA 3.0), (H) *P. subaeruginosa* (ericos bob, CC BY-SA 3.0), (I) *P. angulospora* (Inski, CC BY-NC-SA), (J) *P. baecystis* (Caleb Brown, CC-BY-SA 3.0), (K) *P. pelliculosa* (Scottdarbey, CC-BY-SA 3.0), (L) *P. semilanceata* (Alan Rockefeller, CC-BY-SA 3.0), (M) *P. hoogshagenii* (Brayan [Cora](#), Jaramillo, CC BY-SA 3.0), (N) *P. mexicana* (Alan Rockefeller, CC BY-SA 4.0), (O) *P. neoxalapensis* (David Morales, CC-BY-NC 4.0), (P) *P. zapotecorum* (Alan Rockefeller, CC-BY-SA 3.0).

<https://doi.org/10.1016/j.funbio.2022.01.003>

Genera with psilocybin-producing species

Genus	Family	Lifestyle	Morphology	Sclerotia
<i>Panaeolus</i>	Bolbitiaceae	Saprobic	Agaricoid, Secotiid	Variable
<i>Pholiotina</i>	Bolbitiaceae	Saprobic	Agaricoid	No
<i>Gymnopilus</i>	Hymenogastraceae	Saprobic	Agaricoid	No
<i>Psilocybe</i>	Hymenogastraceae	Saprobic	Agaricoid, Secotiid	Variable
<i>Pluteus</i>	Pluteaceae	Saprobic	Agaricoid	Variable
<i>Inocybe</i>	Inocybaceae	Ectomycorrhizal	Agaricoid	Variable
<i>Massospora</i>	Entomophthoraceae	Insect Pathogenic	Abdominal Spore Mass	No

Table 1, Meyer & Slot 2023: https://papers.ssrn.com/sol3/papers.cfm?abstract_id=4384673

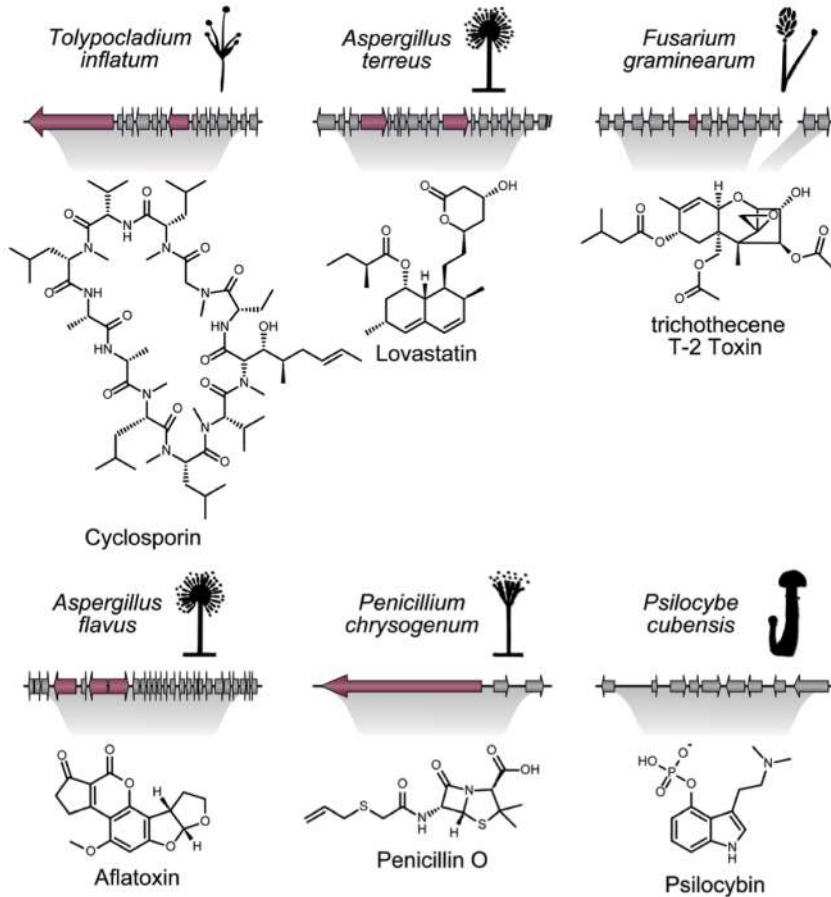
“Secondary metabolites (SMs) are generally defined as small organic molecules produced by an organism that are not essential* for their growth, development and reproduction.”

From: Biotechnology and Biology of Trichoderma, 2014

* i.e., a molecule *in addition* to the standard toolkit shared by most branches of the tree of life.



Psilocybe ovoideocystidiata – West LA, Sept. 2022. RD

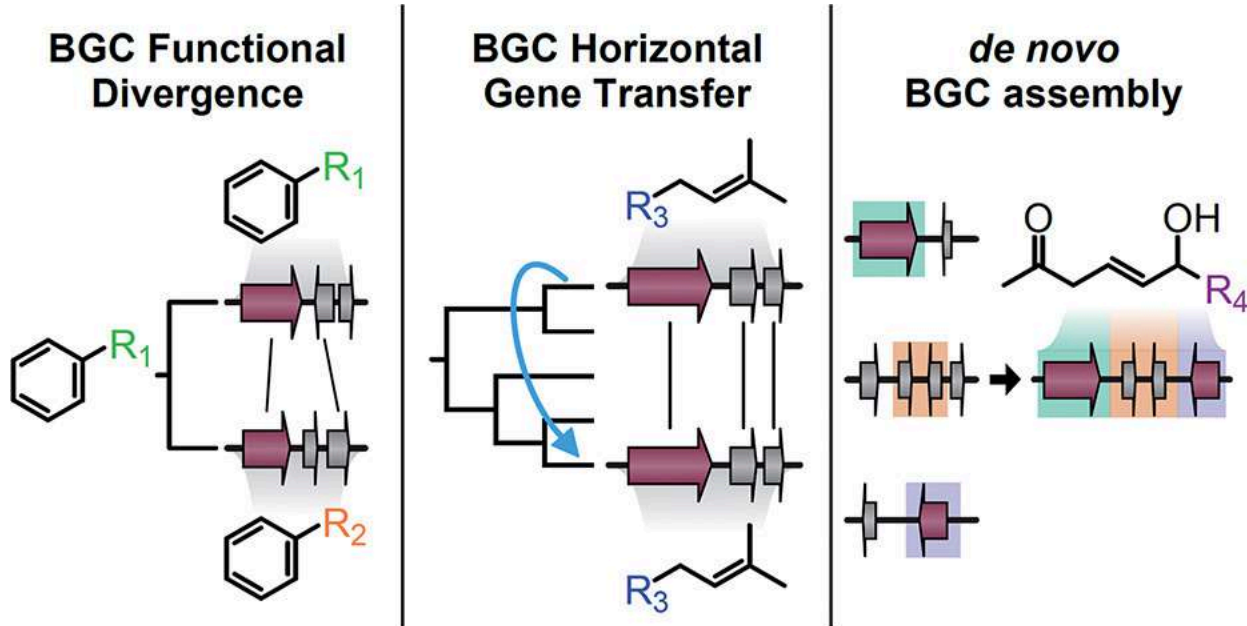


Secondary metabolites are end products in a careful process of construction and modification, encoded genetically in groups of genes called biosynthetic gene clusters (BGCs).

These genes encode different proteins with different functions in the constructive process.

They include enzymes, transporters, and transcription factors, as well as enzymes that confer resistance to the final metabolite.

Mechanisms for the evolution of BGCs and fungal chemodiversity:



Horizontal gene cluster transfer increased hallucinogenic mushroom diversity

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Patrick Brandon Matheny,³ and Jason C. Slot^{1,4} 2018

Abstract

Secondary metabolites are a heterogeneous class of chemicals that often mediate interactions between species. The tryptophan-derived secondary metabolite, psilocin, is a serotonin receptor agonist that induces altered states of consciousness. A phylogenetically disjunct group of mushroom-forming fungi in the Agaricales produce the psilocin prodrug, psilocybin. Spotty phylogenetic distributions of fungal compounds are sometimes explained by horizontal transfer of metabolic gene clusters among unrelated fungi with overlapping niches. We report the discovery of a psilocybin gene cluster in three hallucinogenic mushroom genomes, and evidence for its horizontal transfer between fungal lineages. Patterns of gene distribution and transmission suggest that synthesis of psilocybin may have provided a fitness advantage in the dung and late wood-decay fungal niches, which may serve as reservoirs of fungal indole-based metabolites that alter behavior of mycophagous and wood-eating invertebrates. These hallucinogenic mushroom genomes will serve as models in neurochemical ecology, advancing the (bio)prospecting and synthetic biology of novel neuropharmaceuticals.

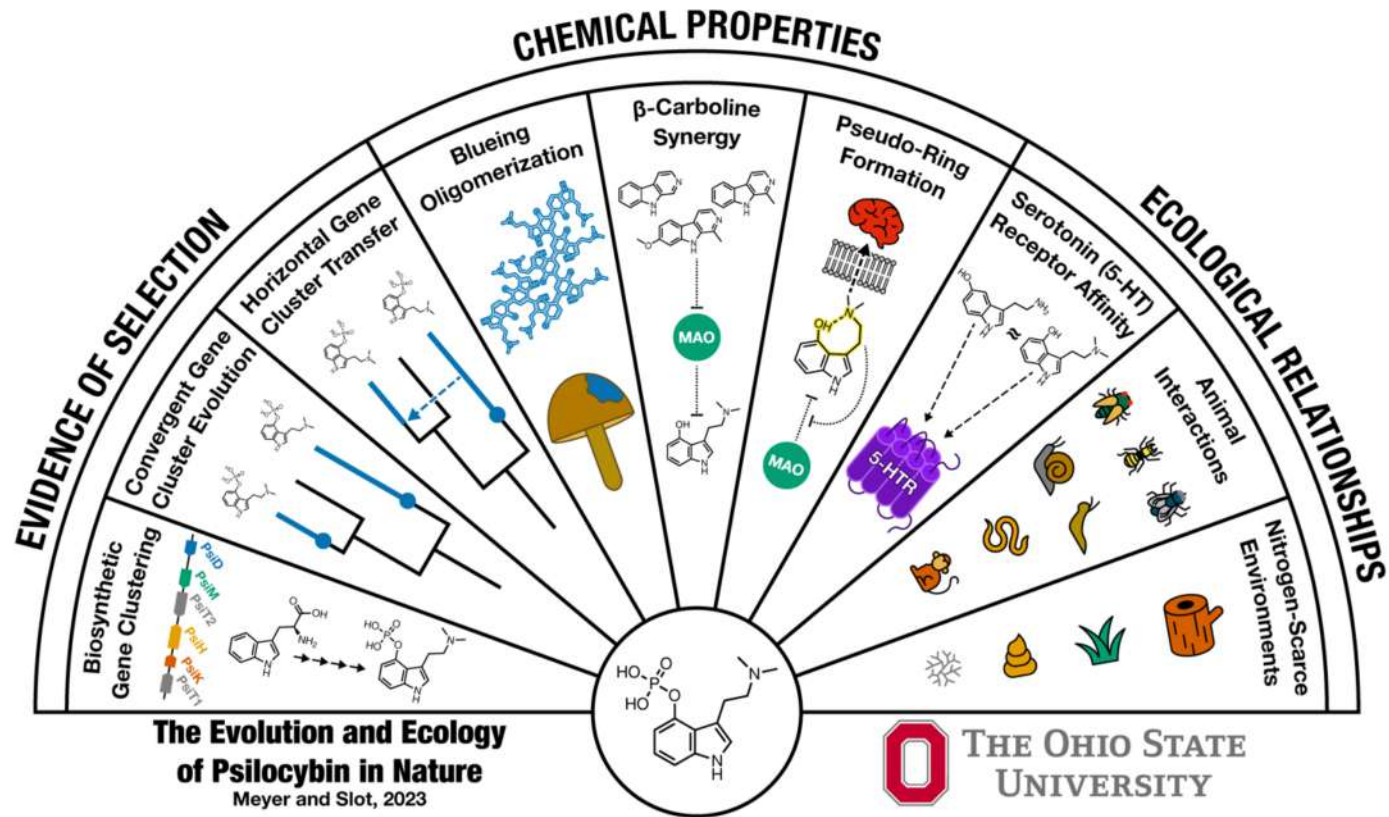
Convergent evolution of psilocybin biosynthesis by psychedelic mushrooms

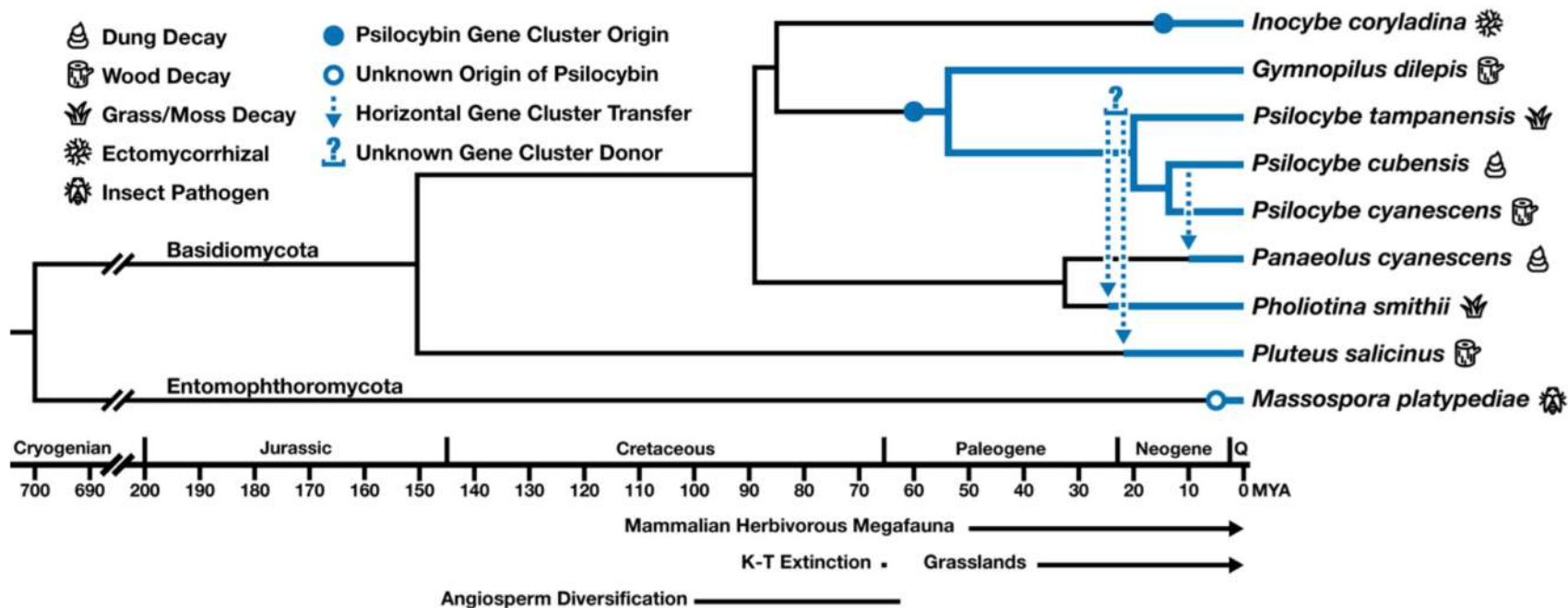
Ali R. Awan, Jaclyn M. Winter, Daniel Turner, William M. Shaw, Laura M. Suz, Alexander J. Bradshaw, Tom Ellis, Bryn T.M. Dentinger 2018

<https://www.biorxiv.org/content/10.1101/374199v2.full>

Abstract

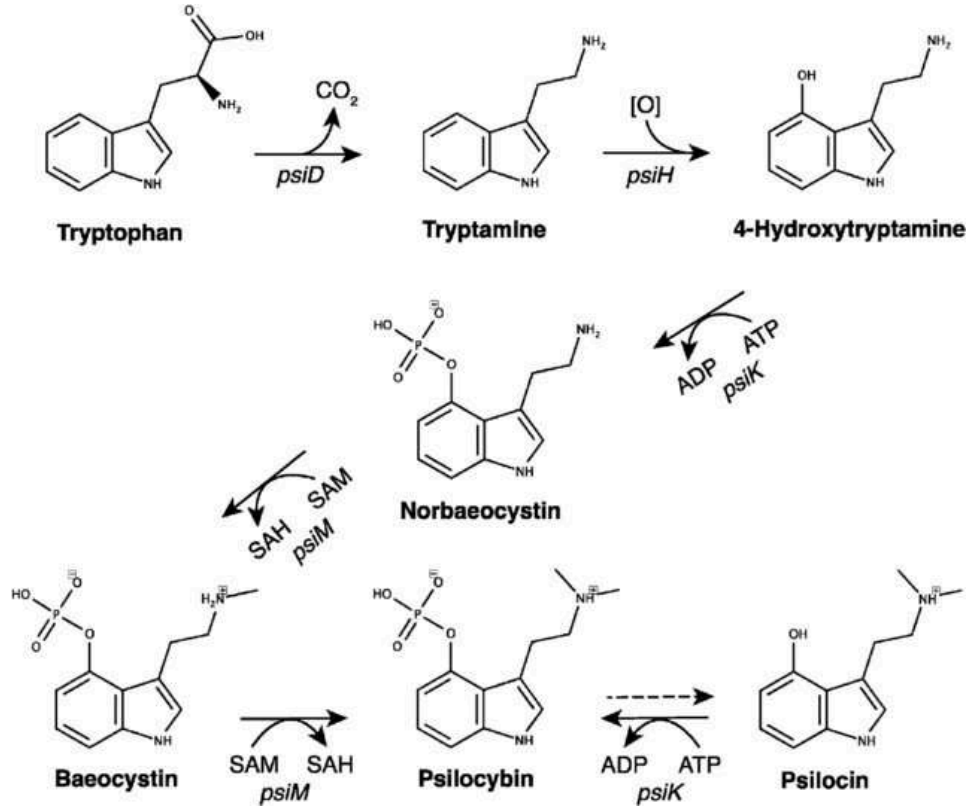
Psilocybin is a psychoactive compound with clinical applications produced by dozens of mushroom species¹. There has been a longstanding interest in psilocybin research with regard to treatment for addiction², depression³, and end-of-life suffering⁴. However, until recently very little was known about psilocybin biosynthesis and its ecological role. Here we confirm and refine recent findings⁵ about the genes underpinning psilocybin biosynthesis, discover that there is more than one psilocybin biosynthesis cluster in mushrooms, and we provide the first data directly addressing psilocybin's ecological role. By analysing independent genome assemblies for the hallucinogenic mushrooms *Psilocybe cyanescens* and *Pluteus salicinus* we recapture the recently discovered psilocybin biosynthesis cluster^{5,6} and show that a transcription factor previously implicated in its regulation is actually not part of the cluster. Further, we show that the mushroom *Inocybe corydalina* produces psilocybin but does not contain the established biosynthetic cluster, and we present an alternative cluster. Finally, a meta-transcriptome analysis of wild-collected mushrooms provides evidence for intra-mushroom insect gene expression of flies whose larvae grow inside *Psilocybe cyanescens*. These larvae were successfully reared into adults. Our results show that psilocybin does not confer complete protection against insect mycophagy, and the hypothesis that it is produced as an adaptive defense compound may need to be reconsidered.





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The most common factor among the various ecological niches of psilocybin-producing fungi appears to be their low-nitrogen substrates. For example, the late stages of wood and dung decay are nitrogen-poor environments (Chen et al., 2013; Cowling and Merrill, 1966; Hao et al., 2004; Hess et al., 2021; Petersen et al., 1998), and ectomycorrhizal species are effective nitrogen scavengers for their host plants (Hernández et al., 2002; Mbarki et al., 2008; Stamets, 1996). Yet, despite environmental limitations, a large portion of nitrogen is allocated to psiloids. For example, psilocybin can make up to 1.6% of a mushroom's total nitrogen content (Borner and Brenneisen, 1987; Braaksma and Schaap, 1996; Gartz, 1994; Kamata et al., 2005). The high nitrogen allocation to psilocybin production suggests that its benefits outweigh any cost to nitrogen-limited growth and reproduction processes.



Psilocybin construction starts with the essential amino acid tryptophan,

which then undergoes a series of modifications through its interactions with specific proteins.

Van Court, R. C. *et al.* Diversity, biology, and history of psilocybin-containing fungi: Suggestions for research and technological development. *Fungal Biol.* **126**, 308–319 (2022)

<https://doi.org/10.1016/j.funbio.2022.01.003>

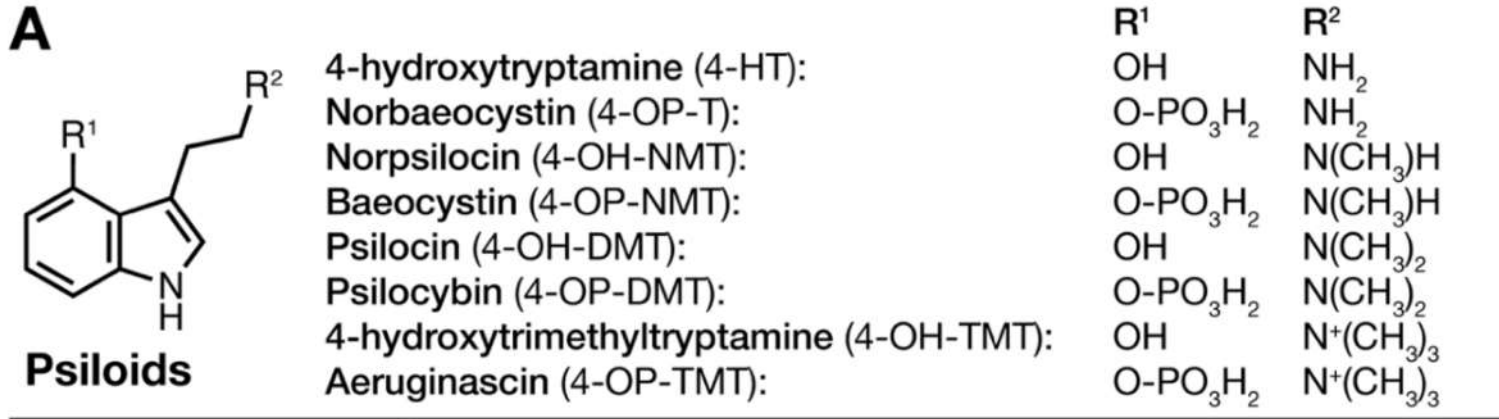


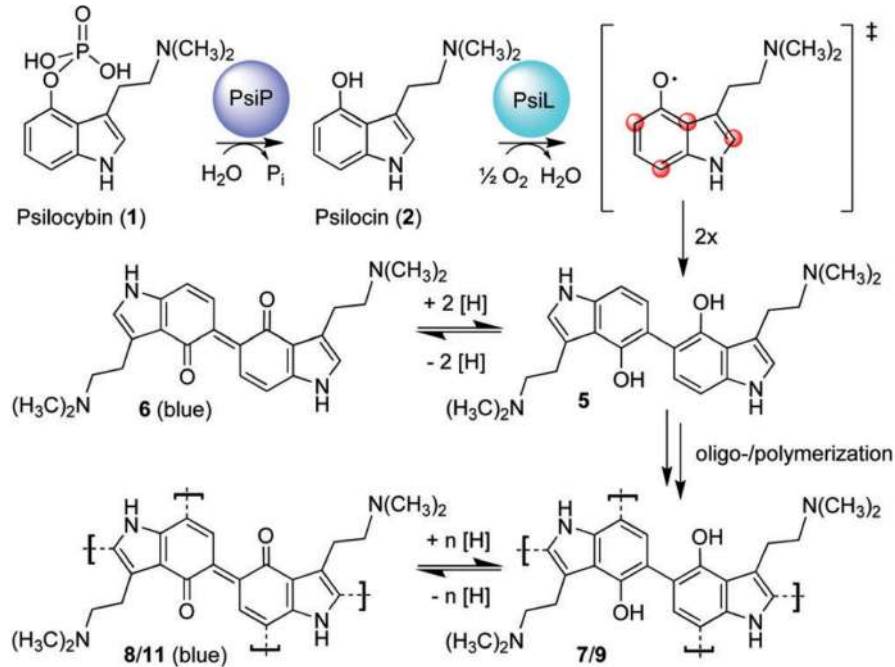
Fig. 1. Tryptophan-derived compounds in psychedelic/magic mushrooms (Gurevich, 1993; Lenz et al., 2021b). (A) Psiloids: the eight known natural 4-substituted tryptamine metabolites of the psilocybin biosynthesis pathway (Fricke et al., 2017; Stijve, 1984) found primarily in mushroom fruiting bodies (Blei et al., 2020). (B) Serotonin: a key signaling molecule found in all domains of life (Erland et al., 2019) and a key neurotransmitter in animals (Andrés et al., 2007).

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Another tryptophan-derived molecule is serotonin, which fulfills diverse cell signaling functions across all lifeforms.

The structural similarity of Psilocin to Serotonin allows Psilocin to interact with animal Serotonin-receptor proteins.

Evolutionary genomics is a powerful tool that can provide insights into the ecological role of psilocybin production. By studying rare molecular evolutionary events surrounding psilocybin's biosynthetic pathway (i.e., gene clustering, convergent evolution, HGT), we are given clues as to how these species have adapted to their environments. Two convergent origins of the psilocybin gene cluster in Agaricales have been established in wood-decay (the inferred ancestral state of *Gymnopilus* and *Psilocybe*) and in the ectomycorrhizal *Inocybe*, with another possible convergence in Entomophthoromycota (Awan et al., 2018; Boyce et al., 2019; Reynolds et al., 2018). Most other known psilocybin producers are saprobes and likely acquired the wood-decay-associated cluster through HGT (Fig. 2A) (Reynolds et al., 2018). This suggests that psilocybin may have different functions in different ecological niches but is most useful in decay niches. The ancestral ecologies of the cluster donor species were likely similar saprobic niches to their recipients, leading to physical contact between them and increasing the chances of HGT (Gluck-Thaler and Slot, 2015; Reynolds et al., 2018). Horizontal acquisition of psilocybin production may also facilitate expansion into new ecological niches (Slot, 2017). For instance, the HGT between the *Ps. cubensis* and *Pa. cyanescens* lineages likely occurred in their shared dung decay niche (Reynolds et al., 2018), suggesting both species faced similar pressures that psilocybin may have at least partially alleviated. Psilocybin production may defend against cohabiting animals, especially insects, that compete for resources or consume the fungus. An ecological association with insects is further supported by the likely convergent origin of psilocybin in *M. levispora/platypedia*, a fungus that is directly antagonistic to an insect.



Ecological function in the blue bruising reaction?

Caused by “rapid conversion of phosphorylated psiloids into their hydroxylated counterparts, followed by oxidative coupling that forms blue oligomers.”

“Chromophoric oligomers may serve a defensive ecological function due to their polyphenolic and aryl-coupling properties. When ingested, these compounds generate reactive oxygen species, causing intestinal lesions in insects (Barbehenn and Peter Constabel, 2011; Salminen and Karonen, 2011).”

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https://papers.ssrn.com/sol3/papers.cfm?abstract_id=4384673

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7004109/pdf/ANIE-59-1450.pdf>

Do we why know why mushrooms produce psilocybin?

No. But it must be important different ways in different fungal lineages,
whatever they're doing...