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## The epichloae: alkaloid diversity and roles in symbiosis with grasses

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### Abstract

Epichloae (*Epichloë* and *Neotyphodium* species; Clavicipitaceae) are fungi that live in systemic symbioses with cool-season grasses, and many produce alkaloids that are deterrent or toxic to herbivores. The epichloae colonize much of the aerial plant tissues, and most benignly colonize host seeds to transmit vertically. Of their four chemical classes of alkaloids, the ergot alkaloids and indole-diterpenes are active against mammals and insects, whereas peramine and lolines specifically affect insects. Comparative genomic analysis of Clavicipitaceae reveals a distinctive feature of the epichloae, namely, large repeat blocks in their alkaloid biosynthesis gene loci. Such repeat blocks can facilitate gene losses, mutations, and duplications, thus enhancing diversity of alkaloid structures within each class. We suggest that alkaloid diversification is selected especially in the vertically transmissible epichloae.

### Ecology and agricultural implications of the epichloae

Epichloae (*Epichloë* and *Neotyphodium* species; Clavicipitaceae) are symbiotic fungi (endophytes) that grow within, and have apparent co-evolutionary relationships with, cool season grasses (Poaceae, subfamily Poöideae) [1,2••]. Epichloae are typically host-specific, and grow in intercellular spaces in a highly regulated fashion without eliciting responses from, or damage to, adjacent host cells [3•,4•,5••]. They are best known for their production of various combinations of alkaloids in four different classes, which help protect their host plants by deterring vertebrate and invertebrate herbivores (Figure 1; Table 1).

Grass-epichloë symbiotes consist of individual plants systemically harboring single clones of their *Epichloë* or *Neotyphodium* species. As such, a symbiotum is a genetic entity with phenotypes that are manifested under specific environmental conditions. Documented phenotypes of grass plants with symbiotic epichloae vary among species and genotypes, and include improved survival and growth, enhanced tolerance to stresses such as water deficit, and increased resistance to vertebrate or invertebrate herbivores [1,6-9]. Most epichloae are vertically transmitted in host maternal lineages by virtue of their abilities to grow benignly in seeds [3•,10•]. However, some can produce fruiting bodies (stromata) around inflorescence primordia, and actually suppress maturation of host inflorescences (choke disease). In the culmination of sexual development, mature stromata produce contagious meiotic spores. With one possible exception described recently [11], sexual *Epichloë* species

are haploid. In contrast, most asexual, strictly seed-borne epichloae described to date are polyploid interspecific hybrids [12].

The vast majority of studies on grass-epichloë interactions have focused on ryegrasses (*Lolium* spp.) and fescues (*Festuca*, *Lolium* and *Schedonorus* spp.) that are important pasture and forage grasses in temperate climates, and have been motivated by episodes of toxicosis suffered by livestock grazing some of those grasses. For example, *Neotyphodium coenophialum*, an asexual, strictly seed-transmitted symbiont of tall fescue (*Lolium arundinaceum* = *Schedonorus arundinaceus* = *Festuca arundinacea*), is responsible for fescue toxicosis, generally believed to be an effect of its ergot alkaloids [13,14•] (Figure 1b). Additionally, *N. coenophialum* produces peramine [15 •] (Figure 1c) and lolines [16] (Figure 1a), which are more specifically active against insects. Also, strains of *Neotyphodium lolii* in perennial ryegrass (*Lolium perenne*) produce tremorgenic indole diterpenes implicated in ryegrass staggers (Figure 1d) [17,18•], as well as peramine and ergot alkaloids. Although it has been suggested that alkaloids are an agronomically selected feature of epichloae in forage grasses [19], the widespread capability of epichloae in wild grasses to produce various combinations and often very high levels implies that they have important ecological roles in defensive mutualisms [12,20•,21••].

## Activities against grazing mammals

Structural similarities of ergot alkaloids to neurotransmitters that control motility, body temperature, and appetite enable the alkaloids to interact with the dopamine, serotonin and noradrenaline receptors in central and peripheral nervous systems (reviewed in [22]). Symptoms of fescue toxicosis in cattle can include poor conception rates, agalactia, poor weight gain, increased respiration rate and salivation, and loss of blood flow to the extremities resulting in necrosis of the hooves and tail (dry gangrene). In horses, reproductive problems include placental abnormalities and fetal asphyxia [13,23]. Toxic effects of ergot alkaloids are exacerbated by heat or cold stress to the animal [13,23,24]. Also, because concentrations of the major ergot alkaloid, ergovaline, are highest in the crown area of vegetative tall fescue plants, toxicosis appears to be particularly common when the animals overgraze infested fields [23].

Effects of ergot alkaloids on rabbits have been tested with perennial ryegrass symbiotic with each of three epichloae — the natural *N. lolii* × *Epichloëtyphina* hybrid strain Lp1, a marker-exchange *dmaW* mutant lacking ergot alkaloids, and an *lpsA* disruption mutant that lacks ergovaline but produces simpler ergot alkaloids — as well as endophyte-free perennial ryegrass [25••]. Rabbits preferred plants symbiotic with the *dmaW* knockout to the plants with wild-type Lp1, the *lpsA* knockout, and endophyte-free plants, suggesting that simple ergot alkaloids counteract an endophyte enhancement of herbivore preference.

The various ergot alkaloids have distinct pharmacological properties. Hence, ergotamine is used for migraines, brominated ergocryptine is used for parkinsonism, and ergonovine has been used as an aid for parturition [22]. Therefore, it is reasonable to expect that, at higher doses, they also have different spectra of activity against natural herbivores. Much less information is available about the biological spectra and relative activities of different indole-diterpenes, which have a huge range of structural diversity [18•]. Though their tremorgenic activities are not directly lethal, affected animals lose their balance and can suffer accidental trauma or drowning. Published studies to date are limited, but so far this class of alkaloids has been demonstrated to affect neurotransmitter release and to block smooth muscle high-conductance calcium-activated potassium (maxi-K) channels in vertebrates [26].

## Activities against insects

Among the four classes of alkaloids produced by epichloae, lolines and peramine are best documented to provide protection against insects, and neither alkaloid class exhibits appreciable activity against livestock [27•,28]. Peramine was identified as a feeding deterrent that is crucial for protection of perennial ryegrass pastures from Argentine stem weevil (*Listronotus bonariensis*; Coleoptera) in New Zealand pastures [29]. Loline alkaloids (reviewed in [16]) show a particularly broad spectrum of activity, and are approximately as potent as nicotine against insects. Lolines have been shown to be active against Japanese beetle (*Popillia japonica*; Coleoptera) [30], fall armyworm (*Spodoptera frugiperda*; Lepidoptera), European corn borer (*Ostrinia nubilalis*; Lepidoptera) [31•], Argentine stem weevil [32], birdcherry oat aphid (*Rhopalosiphum padi*; Homoptera), and greenbug aphid (*Schizapus graminis*; Homoptera) [33]. *In vitro* tests have also shown that *N*-acetyllooline (NAL) and *N*-formyllooline (NFL) deter feeding of Japanese beetle larvae (*P. japonica*, Coleoptera) at concentrations comparable to those measured in roots of tall fescue-*N. coenophialum* symbiota [34].

Different lolines may vary in specificity to different insect species and on certain development stages, and the decorations of the 1-amine are important in the spectrum of anti-insect activity of lolines. Significant effects of NFL and NAL have been observed in tests with *S. frugiperda* [31•]. Both affected mean larva weight in a no-choice test, and NFL also had a deterrent effect in a feeding choice experiment. In contrast, the two *N*-methyl forms, loline and *N*-methyllooline (NML), had no significant activity in similar tests. When sprayed onto greenbug aphids, NAL and NML exhibited greater insecticidal activity than NFL and loline [31•]. Meadow fescue (*Lolium pratense*) with *N. uncinatum*, a symbiotum with high levels of NFL and NAL, deterred oviposition by Argentine stem weevil [35]. Interestingly, *N*-acetylnoorloline (NANL) and NFL had different effects in deterrence and mortality of weevil larvae [32,35]. At high concentrations NFL reduced their growth, development and survival, whereas NANL caused high mortality but little effect on larval growth and development.

The pharmacological properties of lolines and peramine on insects remain to be investigated. Furthermore, several grass-epichloë symbiota accumulate 1-acetamidopyrrolizidine (AcAP; Figure 1a) instead of fully cyclized lolines, suggesting that AcAP may have a biological activity yet to be determined [21••]. Also relatively under-studied are the mechanisms of anti-insect activities of ergot alkaloids and indole-diterpenes (reviewed in [36]). Tests with *dmaW* and *lpsA* knockout endophytes indicate an effect of ergot alkaloids, probably ergovaline, against the black cutworm *Agrotis ipsilon* (Lepidoptera: Noctuidae) [37]. Also, the indole-diterpene, nodulisporic acid, is a potent and broad-spectrum insecticide, and has been shown to modulate glutamate-gated chloride channels in insects [38].

## Genetic basis for alkaloid diversity

Much of the basis for diversity within each class of alkaloids (Table 1) is due to presence-absence polymorphism of middle-pathway or late-pathway genes. In addition, variations in certain genes for steps in ergot alkaloid and indole-diterpene biosynthesis affect the activity or specificity of certain biosynthetic enzymes, thereby contributing additional chemotypic diversity [21••].

So far, variation in loline alkaloids is strictly associated with presence or absence of late-pathway genes. Epichloae with all 11 *LOL* genes, such as *Epichloë festucae* E2368 (Figure 2a), make loline, NML, and NFL (and when symbiotic with certain plants, NAL is also

produced). Strains lacking functional *lolO* accumulate 1-acetamidopyrrolizidine (AcAP), and strains lacking *lolN*, *lolM*, and *lolP* accumulate NANL [21••].

Ergot alkaloids include clavines, lysergic acid and simple lysergic acid amides, as well as the complex ergopeptines derived from lysergic acid and three common amino acids by a series of condensations and cyclizations (Figure 1b) [22]. Considerable diversity in ergot alkaloid profiles is associated with variation in organization, gene content and gene sequences in the *EAS* clusters. Figure 2b compares *EAS* loci between *E. festucae* F11, which makes the same ergopeptine (ergovaline) found in toxic tall fescue, *Neotyphodium gansuense* var. *inebrians*, a seed-borne symbiont of drunken horse grass (*Achnatherum inebrians*) that makes ergonovine but no ergopeptine, and *Periglandula ipomoeae*, a related seed-transmitted symbiont of morning glories that makes another ergopeptine (ergobalansine) as well as ergonovine and lysergic acid  $\alpha$ -hydroxyethylamide [21••]. The *EAS* cluster of *N. gansuense* var. *inebrians* resembles that of *Periglandula ipomoeae*, except that it lacks *lpsA* and *easH*, and the cluster is divided into two with a telomere immediately downstream of *cloA*. All other epichloae characterized to date have *EAS* clusters that more closely resemble that of *E. festucae*, though they vary considerably in content. Some lack *EAS* genes completely, some have all genes for ergopeptine biosynthesis, and some lack all but the four genes required for chanoclavine biosynthesis [21••]. The structural differences between ergopeptines, such as ergobalansine and ergovaline, are determined by sequence variations in *lpsA*, which determine which amino acids are specified by each of the three modules of the LpsA subunit in the nonribosomal peptide synthetase (NRPS) [22,39•] (Figures 1b and 2b). So far, ergopeptines have been identified that are derived from 20 different permutations of amino acids [22,40]. However, ergovaline is the only ergopeptine definitively identified as a product of symbiotic epichloae.

Peramine biosynthesis also involves a multifunctional NRPS, the product of the *perA* gene [15•]. Intriguingly, two forms of *perA* have been found, one that specifies seven enzymatic domains, and a variant (*perA- $\Delta R^*$* ) that lacks the sequence for the C-terminal reductase domain (Figure 2c) [21••]. Symbiotic *E. festucae* E2368 has, and expresses, the *perA- $\Delta R^*$*  variant [21••], raising the possibility that it may play a role in synthesis of a heretofore unknown metabolite.

The extreme diversity of indole-diterpenes, similarly to ergot-alkaloid diversity, is determined by presence-absence polymorphism as well as sequence variations in *idtP* and *idtQ* that affect specificity of enzymes encoded by those genes (Figure 2c) [18•]. Furthermore, *ltmE* and *ltmJ*, which specify late steps in lolitrem B biosynthesis, are derived by duplication and neofunctionalization of other *IDT* genes [21••].

## Is alkaloid diversity selectively favored?

Clearly the Clavicipitaceae, and particularly the epichloae, have the capability to produce a wide variety of alkaloids, both among and within chemical classes. The facts that various forms of ergot alkaloids, indole-diterpenes and lolines are produced by different strains and exhibit different biological specificities suggest that diversity of chemical profiles with many structural variants may be inherently beneficial. Several lines of evidence suggest that chemotypic variation is selectively favored. First, there is considerable variation between and even within *Epichloë/Neotyphodium* species for alkaloids in the various classes [12,41]. For example, of the two *E. festucae* strains with sequenced genomes, F11 produces ergot alkaloids (end-product, ergovaline), peramine, and indole-diterpenes (end-product, lolitrem B), whereas E2368 produces only lolines (end-product, NFL)[21••]. Even different *E. festucae* isolates from *Festuca rubra* subsp. *commutata* differ in production of ergot alkaloids [12] or indole-diterpenes [42•]. In a survey of *E. festucae* from different host

grasses [42•], there was considerable variation in the end-product indole-diterpenes (terpendole C, lolitriol, and various lolitremes) based on presence or absence of functional *ltmE*, *ltmF*, and *ltmJ* genes.

Second, the *EAS*, *IDT* and *LOL* gene clusters in the epichloae contain large blocks of transposon-derived repeat sequences, which would be expected to promote rearrangements and partial or complete deletions of the genes. Comparisons of the alkaloid loci among various species and strains [21••] indicates that the lack of some or all of the genes in a cluster is most commonly due to deletions, because gene remnants are often present. An exception is in the *E. festucae* *IDT* cluster, where *ltmE* and *ltmJ* are uniquely found, and no remnants of those genes have been found in the other sequenced genomes. The abundance of repeats within each cluster is not indicative of other secondary metabolism gene clusters [21••], suggesting that unstable alkaloid clusters may be a selected trait.

Third, as described above, different forms of alkaloids within each class exhibit differences in biological activity. Therefore, it may be expected that different alkaloid profiles are selected according to the context of prevailing herbivores on the different host grasses and in different environments. It has been suggested that ecological and life-cycle variation among the epichloae has selected for alkaloid loci imbued with blocks of transposon-derived repeats that facilitate both gene losses and, as in the case of *ltmE* and *ltmJ*, gene duplication and neofunctionalization [21••].

Fourth, those symbionts that produce extremely high levels of certain alkaloids are strictly vertically transmitted, asexual species. Loline alkaloids in plants with strictly seed-borne epichloae range from 500 to nearly 20 000  $\mu\text{g g}^{-1}$  of plant dry mass [43•], whereas levels in grasses with horizontally transmissible, sexual *Epichloë* species accumulate less than 500  $\mu\text{g g}^{-1}$  plant dry mass [12]. Levels of the other alkaloids tend to be comparable between plants with sexual and asexual species, although alkaloids are less frequently found in plants with sexual epichloae [44,45]. However, levels of ergot alkaloids in drunken horse grass (*A. inebrians*) with strictly seed-borne *N. gansuense* var. *inebrians* are 2–3 orders of magnitude higher than in most other grasses with ergot-alkaloid-producing epichloae [46]. The high levels of alkaloids may simply relate to the importance of host protection for vertically transmitted symbionts, and might also compensate for the relatively low potential for asexual species to evolve diverse alkaloid profiles.

Fifth, an unexpected pattern emerged when characterizing alkaloid gene clusters in hybrid epichloae [47]; those that had received redundant copies of alkaloid gene clusters consistently lacked a subset of genes from one of the gene clusters. *Neotyphodium chisosum* and *N. uncinatum* had independently obtained two *LOL* clusters, and in both, one cluster was complete and the other lacked *lolN*, *lolM*, and a functional *lolP*. One of the two *IDT* clusters in the hybrid designated *Neotyphodium* sp. FaTG-2 lacked *ltmE* and *ltmJ*. Finally, two strains of *N. coenophialum*, probably both originating from an ancestor with complete, redundant *EAS* clusters, had independent gene losses. Strain e19 had an inactivating frame-shift in *lpsB* of the *EAS2* cluster (*lpsB2*), whereas strain e4163 had an apparently functional *lpsB2*, but had lost *lpsB1* and the adjacent *easE1* gene. We speculate that these gene losses and mutations may reduce efficiency of the biosynthetic pathways such that intermediates as well as end-products accumulate to substantial levels. Panaccione [48••] has suggested that such a situation may be selectively favored based on the diversity of biological activities of different structures in each alkaloid class.

## Conclusions

Two features of the epichloae are of paramount importance in establishment of mutualistic symbioses with grasses: their capability of, and sometimes dependence upon, vertical transmission *via* seeds, and their production of alkaloids that protect host plants from vertebrate and invertebrate herbivores. Both characteristics are variable in the epichloid clade. *Epichloë/Neotyphodium* species in some hosts are incapable of vertical transmission, and rely on spores formed in their fruiting structures for contagious (horizontal) transmission, others can use both transmission methods, and others, including a large number of interspecific hybrids, are asexual and mainly or solely seed-transmitted. On the basis of reports to date, alkaloids are most frequently associated with strictly seed-transmitted epichloae, less often with mixed-transmission strains, and rarest among those that only transmit horizontally. Also, strictly seed-borne epichloae can produce much higher levels of alkaloids — particularly lolines — than horizontally transmissible epichloae. Chemotypes of epichloae vary widely both with respect to presence or absence of each alkaloid class, and structural variation within each class, with implications for their spectra of biological activity. Also, selection may favor alkaloid profiles that include substantial levels of intermediates over those dominated by pathway endproducts [48••]. Evidence for selection on alkaloid levels and chemotypic diversification is particularly strong for symbiots with vertically transmissible epichloae.

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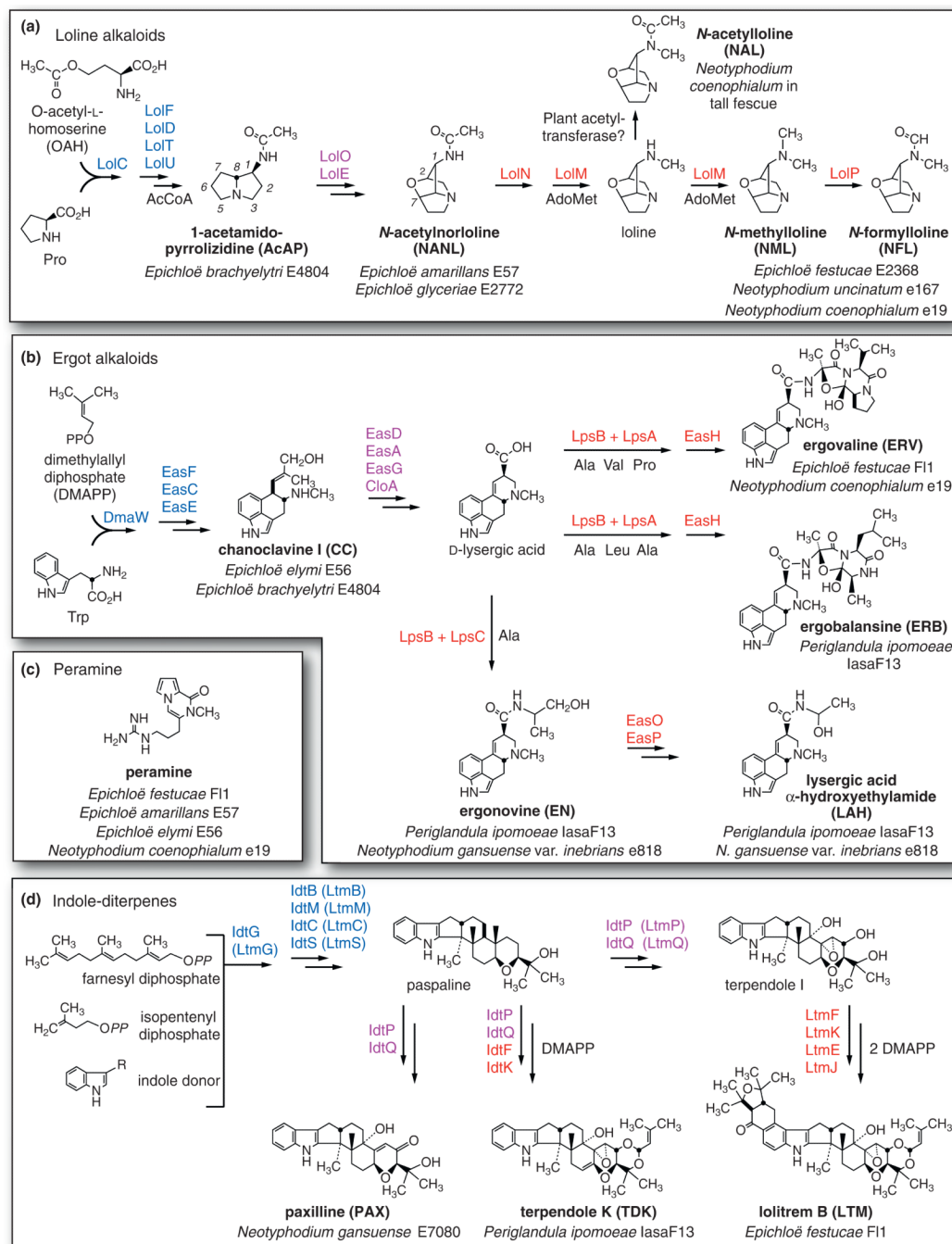
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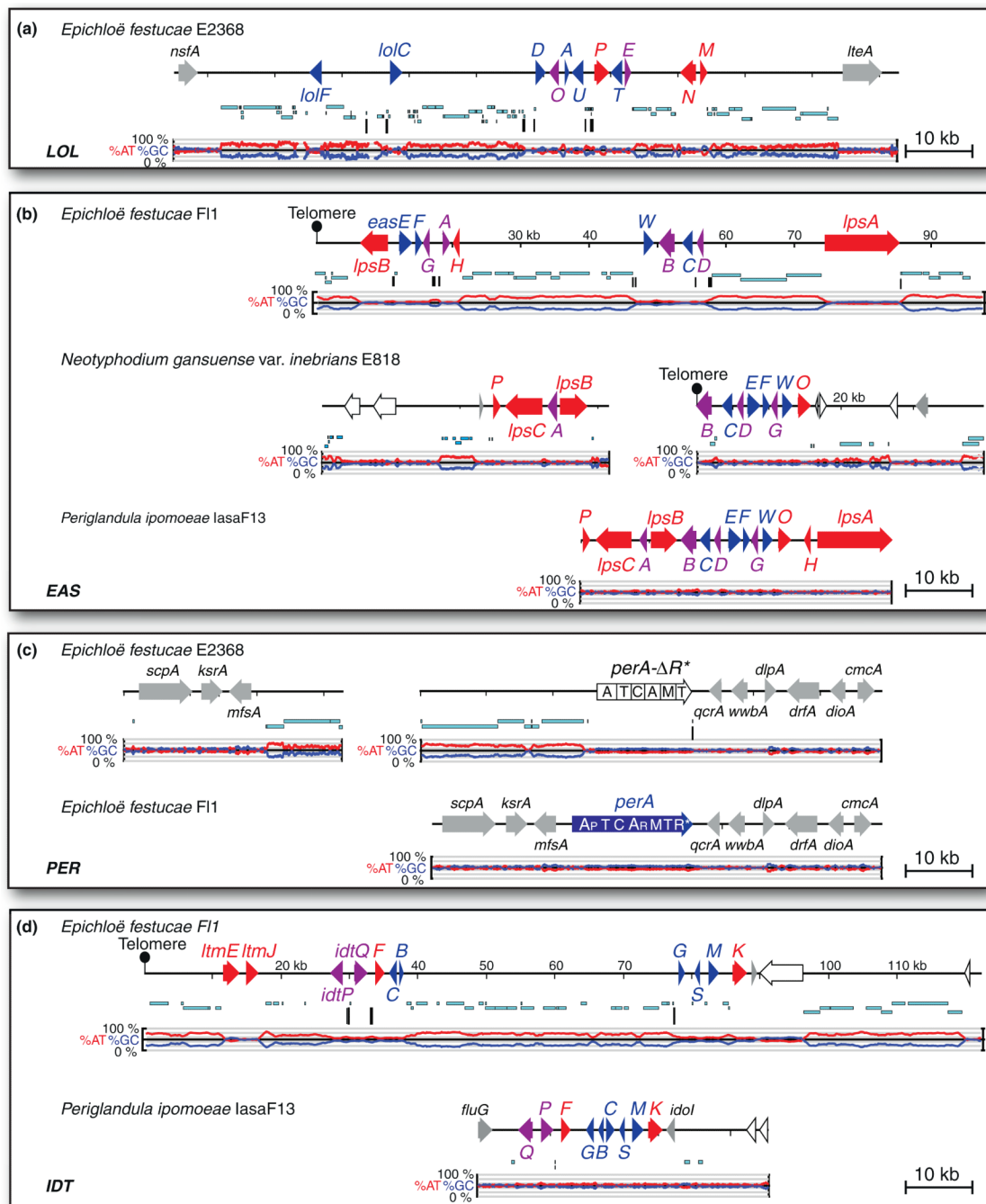
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alkaloid biosynthetic pathways may be selectively favored such that pathway intermediates as well as end-products accumulate to levels at which they can exert various biological effects.



**Figure 1.** Structures of alkaloids and proposed pathways for their biosynthesis in *Epichoë*, *Neotyphodium* and *Periglandula* species. Panel A: Proposed pathway for lolines. Panel B: Proposed pathway for ergot alkaloids. Panel C: Structure of peramine. Panel D: Proposed pathway for indole-diterpenes. Enzymes are designated according to names of genes that encode them. Enzyme designations are in blue type for early steps, purple type for intermediate steps, and red type for late steps. Substrates and cofactors indicated beneath or to the right of arrows are abbreviated as follows: AcCoA = acetyl-CoA, AdoMet = S-adenosylmethionine, Ala = L-alanine, DMAPP = dimethylallyl diphosphate, Leu = L-leucine,

Pro = *L*-proline, Trp = *L*-tryptophan, Val = *L*-valine. Major pathway end products are designated in bold, and species and strains that produce them *in symbio* are given beneath each.

**Figure 2.**

Structures of biosynthesis gene loci for four classes of alkaloids in several sequenced genomes of *Epichoë*, *Neotyphodium* and *Periglandula* species. Panel A: Loline alkaloid biosynthesis gene loci (LOL). Tracks from top to bottom of each map represent genes (arrows and letters), repeats (cyan bars), miniature inverted-repeat transposable elements (MITEs) (vertical bars), and graphs of AT (red) and GC (blue) contents. Genes are color coded according to Figure 1. Additionally, gray arrows indicate housekeeping and other genes not known to be involved in alkaloid biosynthesis, and open arrows indicate pseudogenes. Gene names are abbreviated as follows: A = *lolA*, C = *lolC*, D = *lolD*, E = *lolE*, F = *lolF*, M = *lolM*, N = *lolN*, O = *lolO*, P = *lolP*, T = *lolT*, and U = *lolU*. Panel B:

Ergot alkaloid biosynthesis gene loci (*EAS*). Gene names are abbreviated as follows: *A* = *easA*, *B* = *cloA*, *C* = *easC*, *D* = *easD*, *E* = *easE* (= *ccsA*), *F* = *easF*, *G* = *easG*, *H* = *easH*, *O* = *easO*, *P* = *easP*, and *W* = *dmaW*. Tracks and color coding are as in panel A. Panel C: The peramine synthetase gene (*perA*) and flanking genes. Domains of the *perA* gene product are indicated, and are discussed in references [15•,21••]. The *perA* gene of *E. festucae* E2368 lacks the 3' region encoding the reductase domain (R\*), thought to be required for peramine biosynthesis, but the possibility that it may encode an enzyme for biosynthesis of a related alkaloid has not been explored. Tracks and color coding are as in panel A. Panel D: Indole-diterpene biosynthesis gene loci (*IDT*). Gene names are abbreviated as follows: *B* = *idtB/ltmB*, *C* = *idtC/ltmC*, *F* = *idtF/ltmF*, *G* = *idtG/ltmG*, *K* = *idtK/ltmK*, *M* = *idtM/ltmM*, *P* = *idtP/ltmP*, *Q* = *idtQ/ltmQ*, *S* = *idtS/ltmS*. Tracks and color coding are as in panel A.

**Table 1**  
***Epichloë* and *Neotyphodium* species with sequenced genomes, and their inferred alkaloid pathway end-products**

Species	Isolate <sup>a</sup>	Host	Transmission	EA <sup>b</sup>	IDT <sup>b</sup>	LOL <sup>b</sup>	PER <sup>b</sup>
<i>Epichloë amarillans</i>	E57 = ATCC 200744	<i>Agrostis hyemalis</i>	Mixed	–	–	NANL	PER
<i>E. amarillans</i>	E4668	<i>Agrostis hyemalis</i>	Mixed	ERV	–	–	PER
<i>E. baconii</i>	E1031, = ATCC 200745	<i>Calamagrostis villosa</i>	Horizontal	–	–	–	PER
<i>E. brachyelytri</i>	E4804	<i>Brachyelytrum erectum</i>	Mixed	CC	–	AcAP	PER
<i>E. bromicola</i>	E502 = ATCC 200750	<i>Bromus erectus</i>	Horizontal	–	–	–	–
<i>E. elymi</i>	E56 = ATCC 201551	<i>Elymus virginicus</i>	Mixed	CC	–	–	PER
<i>E. festucae</i>	E2368	<i>Festuca rubra</i> and <i>Lolium giganteum</i> (ascospore isolate)	Mixed	ERV	–	NFL	(–)
<i>E. festucae</i>	F11	<i>Festuca trachyphylla</i>	Mixed	ERV	LTB	–	PER
<i>E. glyceriae</i>	E277	<i>Glyceria striata</i>	Horizontal	ERV	–	AcAP	(–)
<i>E. “mollis”</i>	E3601 = AL9924	<i>Holcus mollis</i>	Mixed	ERV	–	–	PER
<i>E. poae</i>	E4646	<i>Poa nemoralis</i>	Mixed	–	–	–	(–)
<i>E. poae</i>	E5819	<i>Poa nemoralis</i>	Mixed	ERV	–	–	(–)
<i>E. typhina</i>	E8 = ATCC 200736	<i>Lolium perenne</i>	Horizontal	–	–	–	PER
<i>Neotyphodium aotearoae</i>	e899 = MYA-1229	<i>Echinopogon ovatus</i>	Vertical	–	TD	NFL	–
<i>N. chisosum</i>	e3609 = ATCC 64037	<i>Achnatherum eminens</i>	Vertical	–	–	NFL	PER
<i>N. coenophialum</i>	e19 = ATCC 90664	<i>Lolium arundinaceum</i>	Vertical	ERV	–	NFL	PER
<i>N. coenophialum</i>	e4163	<i>Lolium</i> sp. (4×)	Vertical	ERV	TD	NFL	PER
<i>N. funkii</i>	e4096	<i>Achnatherum robustum</i>	Vertical	CC	TD	–	PER
<i>N. gansuense</i> var. <i>inebrians</i>	e818 = MYA-1228	<i>Achnatherum inebrians</i>	Vertical	LAH	–	–	–
<i>N. gansuense</i>	e7080	<i>Achnatherum inebrians</i>	Vertical	–	PAX	–	–
<i>N. uncinatum</i>	e167 = CBS 102646	<i>Lolium pratense</i>	Vertical	–	–	NFL	–

<sup>a</sup>Strains listed with ATCC and MYA prefixes are in the American Type Culture Collection. CBS = Centraalbureau voor Schimmelcultures, Utrecht, The Netherlands.

<sup>b</sup>Abbreviations: AcAP = 1-acetamidopyrrolizidine, CC = chanoclavine, EN = ergonovine, ERV = ergovaline, LAA = lysergic acid amide, LAH = lysergic acid  $\alpha$ -hydroxyethylamide, LTB = lolitrem B, NANL = *N*-acetyl norlooline, NFL = *N*-formyllooline, nt=not tested, PAX = paxilline, PER = per-amine, TD = terpendoles. – = none detected. (–) indicates *perA*- $\Delta$ R\* locus (see text).