



## Article

# Allelopathic Effects of Foliar *Epichloë* Endophytes on Belowground Arbuscular Mycorrhizal Fungi: A Meta-Analysis

Rui Zhong<sup>1,2,\*</sup>, Lin Zhang<sup>1,\*</sup>  and Xingxu Zhang<sup>2</sup> 

<sup>1</sup> College of Resources and Environmental Sciences, National Academy of Agriculture Green Development, Key Laboratory of Plant-Soil Interactions, Ministry of Education, China Agricultural University, Beijing 100193, China

<sup>2</sup> State Key Laboratory of Grassland Agro-Ecosystems, College of Pastoral Agriculture Science and Technology, Lanzhou University, Lanzhou 730020, China

\* Correspondence: ruizhong@cau.edu.cn (R.Z.); linzhang@cau.edu.cn (L.Z.)

**Abstract:** Many grasses are simultaneously symbiotic with *Epichloë* fungal endophytes and arbuscular mycorrhizal fungi (AMF). *Epichloë* endophytes are a group of filamentous fungi that colonize and grow within aerial plant tissues, such as leaves and stems. Infection and hyphal growth of *Epichloë* endophytes confer fitness advantages to the host plants. In addition to producing fungal alkaloids and altering host metabolic/genetic profiles, it is proven that symbiosis of plants with root/foliar endophytes affects the plant–soil relationship. We propose that the *Epichloë* presence/infection results in variations of soil and root AMF through allelopathic effects. We performed a meta-analysis that integrated the allelopathic effects of *Epichloë* endophytes on grass–AMF development. In the pre-symbiotic phase of grass–AMF symbiosis, root exudation from *Epichloë*-infected plants positively affected AMF growth, whereas the shoot exudates of *Epichloë*-infected plants inhibited AMF growth. In the symbiotic phase of grass–AMF symbiosis, the *Epichloë* infection was found to reduce root mycorrhizal colonization in plants. No pattern in the response of soil AMF to *Epichloë* presence was found. This study should improve our understanding of the impact of *Epichloë* endophytes on belowground microbial symbionts within the same host plant. Grass–*Epichloë*–AMF symbiosis may become an important model for studying above–belowground interactions.

**Keywords:** *Epichloë* endophytes; allelopathy; arbuscular mycorrhizal fungus; host grass



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## 1. Introduction

Above–belowground (ABG) interaction is one of the frontiers in studying and understanding the key biochemical and ecological processes in agricultural and natural ecosystems [1,2]. Plants interact with a wide range of microbes—both above- and belowground—that influence plant fitness through nutrient cycling and a wide array of signaling compounds [3–6]. Grass species are the main components of grassland and agricultural ecosystems, and their fungal symbionts may exert an important effect on plant fitness and biological processes [7–10]. Many types of grass in Poaceae are symbiotic with *Epichloë* (Ascomycota, Clavicipitaceae) fungal endophytes [7,9,10]. *Epichloë* fungal endophytes grow asymptotically within the intercellular spaces of the aerial tissues of the grass hosts and vertically transmit their progeny through seeds. *Epichloë* endophytes are a source of secondary metabolites [4,6,11], such as protective fungal alkaloids that are toxic to herbivores and inhibit pathogen colonization, including lolines, indole diterpenes, ergot alkaloids, and peramines. The infection and growth of *Epichloë* endophytes modify plant immune and metabolic profiles to establish a hyphal network within aerial tissues to promote plant persistence and fitness [10,12].

*Epichloë*-infected grass plants may offer a competitive advantage to non-infected grass by influencing the root-associated (bulk soil, rhizosphere soil, and root endosphere) microbes [13–15]. A previous study suggested that aboveground *Epichloë* endophytes also

generate shifts and variations in belowground biochemical processes and components [8]. Arbuscular mycorrhizal fungi (AMF), belonging to the subphylum Glomeromycotina, can form complex associations with host plants and play important roles in promoting plant fitness and productivity [5]. The symbiosis of AMF with roots begins with spore germination, and the development processes of pre-symbiotic and symbiotic phases are also determined by the host plants. Previous studies on *Epichloë*-infected grass plants reported shifts in the soil AMF community structure and root AMF colonization [15,16]. Some studies have assessed the potential pathways of *Epichloë* infection/presence on the developmental status of grass-AMF symbiosis, including root/*Epichloë* exudation and shoot litter [17–19]. *Epichloë* infection also changed the root exudation profiles [20,21] and shoot decomposition. However, the *Epichloë*-facilitated soil microbial processes and the subsequent soil microbial responses that contribute to greater tolerance or resistance to grass hosts are unknown. Determining this may carry significant economic and ecological importance in the development of sustainable agricultural practices [13,22].

Understanding the influences of grass-*Epichloë* interactions on the plant-soil-microbe interactions is of ecological and agricultural importance in grassland and agricultural ecosystems. Based on the transmission of *Epichloë* endophytes and AMF, we propose a model that integrates the effects of grass-*Epichloë*-AMF association. Here, we took complementary approaches to validate the multiple tripartite interactions and the predictions of our model. We reviewed the literature related to the interactions between *Epichloë* endophytes and AMF. A meta-analysis critically analyzed the potential pathways of *Epichloë* infection on pre-symbiotic and symbiotic phases of grass-AMF development, including (1) exploring the allelopathic effects of *Epichloë* endophytes on the pre-symbiotic phase of grass-AMF symbiosis, and (2) accessing the effects of *Epichloë* infection on the symbiotic phase of belowground grass-AMF symbiosis.

## 2. Material and Methods

We performed a quantitative analysis of published results about the effects of *Epichloë* endophyte infection on the pre-symbiotic and symbiotic phases of grass-AMF symbiosis. We obtained and reviewed the related literature reporting the multiple interactions between *Epichloë* and AMF, and performed a standard meta-analysis. Papers in journals were collected from the Web of Science and the China National Knowledge Infrastructure (CNKI) databases using the keywords “*Epichloë*/*Neotyphodium* and mycorrhiza” in May 2022.

These papers were screened based on the following criteria: (1) these fungal endophytes belong to the *Epichloë* (formerly *Neotyphodium*) genus, and the host plants of *Epichloë* endophytes are the cool-season grass species; (2) the experiments had to examine the effects of *Epichloë* endophyte infection/presence on growth and/or colonization of AMF; (3) the references of these selected papers were also checked to search for further appropriate papers; (4) the selected papers had to include the means of AMF growth and colonization for both the endophyte-infected and endophyte-free grasses. Here a total of 32 papers (21 papers in English and 11 papers in Chinese) were obtained from the databases and references of the selected papers for the subsequent meta-analysis.

The extracted values were obtained from the graphs using GetData Graph Digitizer 2.22 and tables in the selected papers. The mean and standard deviations (= standard errors  $\times \sqrt{n}$  or the unreported standard deviations were estimated at 10% of the mean), and the number of repetitions ( $n$ ) of *Epichloë*-infected (experimental group) and non-infected (control group) grass plants were extracted from the selected papers. The data were grouped into the following categories: (1) exudation resources (shoot, root, and *Epichloë* strains) or AMF growth (spore germination and hyphal growth) indices in the pre-symbiotic phase of grass-AMF development, and (2) grass classification (grass genus level) or root/soil AMF (root AMF colonization, root AMF concentration, and length of the extraradical hyphae of soil AMF) in the symbiotic phase of grass-AMF symbiosis. For each group as a categorical variable, this group had to be reported in at least five cases. The AMF growth indices in the pre-symbiotic phase of grass-AMF symbiosis were determined by laboratory experiments,

and the root/soil AMF indices in the symbiotic phase of grass–AMF symbiosis were determined by controlled pot and field experiments. The meta-analysis was performed using MetaWin software (version 2.1, an intellectual property of Michael S. Rosenberg, Dean C. Adams and Jessica Gurevitch), and these effects were calculated using the log of the response ratio ( $\ln R$ ) using the following Equation according to a study [23].

$$\ln R = \ln\left(\frac{\bar{X}_e}{\bar{X}_c}\right) = \ln(\bar{X}_e) - \ln(\bar{X}_c)$$

where  $\bar{X}_e$  and  $\bar{X}_c$  represent the mean AMF growth and colonization of the experimental and control groups, respectively.

The variance of the natural logarithm of the response ratio ( $\ln R(v)$ ) was approximated using the following Equation according to a study [24].

$$\ln R(v) = \frac{S_e^2}{n_e \bar{X}_e^2} + \frac{S_c^2}{n_c \bar{X}_c^2}$$

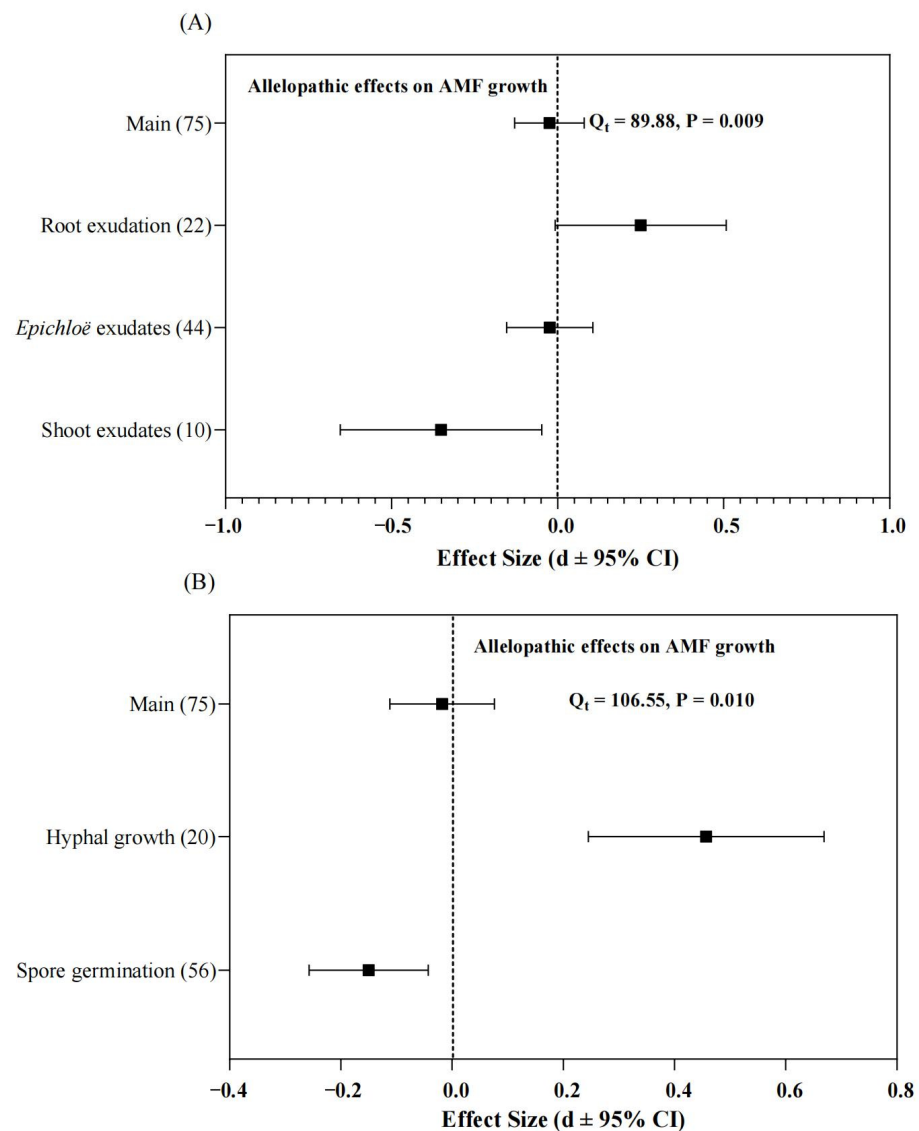
where  $n_e$  and  $n_c$  indicate the repetitions of the experimental and control groups, respectively, and  $S_e$  and  $S_c$  indicate the standard deviations of the experimental and control groups, respectively [24]. The  $\ln R$  means of the effects of *Epichloë* infection/presence on AMF growth and root/soil AMF were calculated using the combination of  $\ln R$  and  $\ln R(v)$ , and  $\ln R(v)$  was weighted by the inverse variance of  $\ln R$  for each observation. The 95% bootstrap confidence intervals (95% CI) for the mean  $\ln R$  were calculated using 9999 iterations of bootstrapping, according to a previous study [24].

### 3. Results

From 142 articles obtained from the Web of Science (124 articles) and CNKI (18 articles) databases, only 32 articles were selected and included in the meta-analysis. A total of 198 cases were obtained from these selected references, including: (1) allelopathic effects of *Epichloë* fungal endophytes on AMF growth (78 cases in 5 studies), and (2) *Epichloë* infection on soil and root AMF colonization in plants (120 cases of 31 studies). The present meta-analysis aimed to assess whether foliar *Epichloë* presence/infection could result in some changes to the belowground AMF components (spore germination, hyphal growth, root AMF colonization/concentration, and soil AMF length in pre-symbiotic and symbiotic phases of grass–AMF development) of the grass hosts.

#### 3.1. Does *Epichloë* Presence Affect the Pre-Symbiotic Phase of Grass-Arbuscular Mycorrhizal Fungi Development?

The results of this present meta-analysis showed there were significant ( $Q_t = 89.88$ ,  $p = 0.009$ ,  $df = 75$ ) and negative (effect size =  $-0.025$ , 95% CI =  $-0.129$  to  $0.008$ ) effects of *Epichloë* fungal endophytes on the pre-symbiotic phase of grass–AMF development within a grass plant (Figure 1A,B). Exudates of *Epichloë* fungal strains had a weakly negative (effect size =  $-0.024$ , 95% CI =  $-0.154$  to  $0.106$ ) effect on AMF growth (spore germination, hyphal length, and branch numbers) compared to the control (equal volume of medium) group (Figure 1A). Meanwhile, the shoot exudation from *Epichloë*-infected grass plants negatively (effect size =  $-0.351$ , 95% CI =  $-0.1655$  to  $-0.048$ ) affected the AMF growth in the pre-symbiotic phase of grass–AMF development than that of non-infected grass plants (Figure 1A). However, root exudation of *Epichloë*-infected grass plants stimulated (effect size =  $0.250$ , 95% CI =  $-0.007$  to  $0.508$ ) AMF growth compared to that of non-infected plants (Figure 1A).



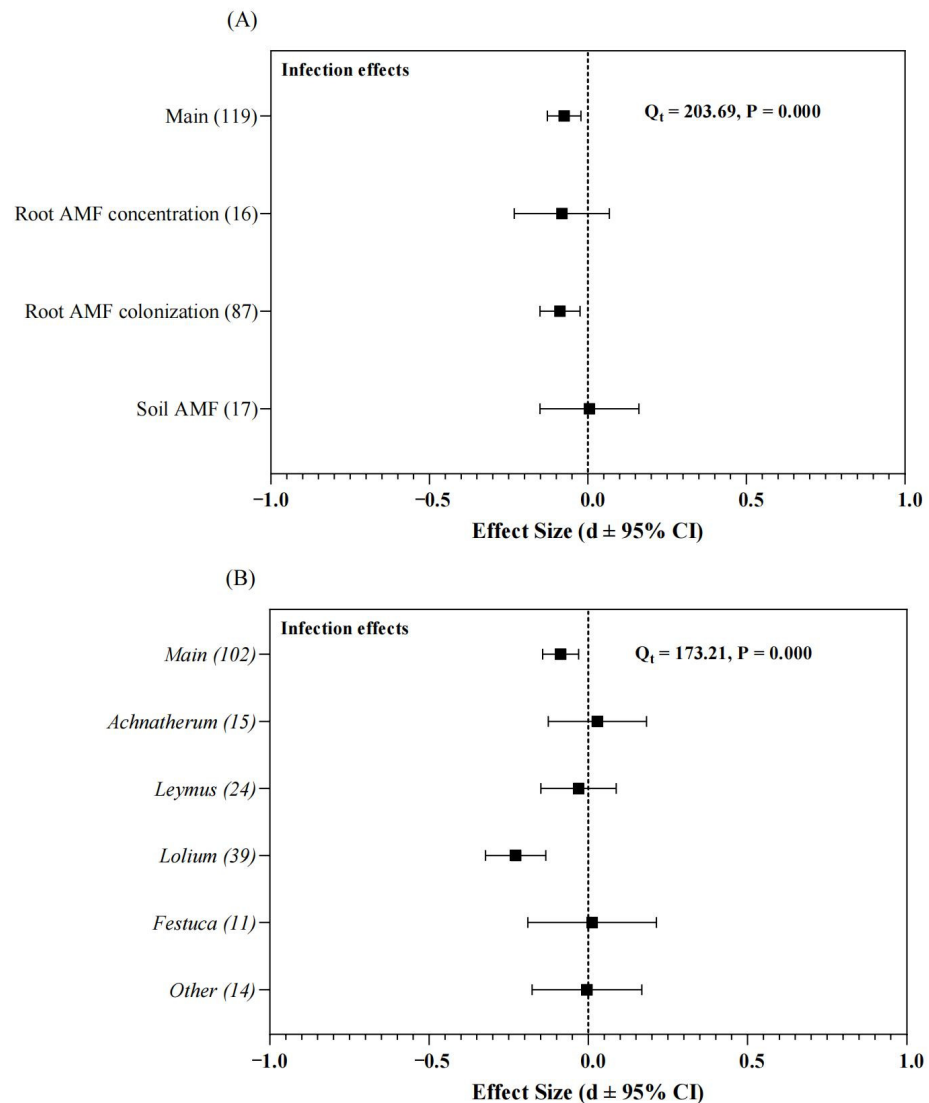
**Figure 1.** The cumulative effect size of allelopathic effects of *Epichloë* endophytes on the pre-symbiotic phase of the grass and arbuscular mycorrhizal fungi (AMF) development among (A) different allelopathic (exudates from root, shoot, and *Epichloë* strains) effects, or (B) AMF germination and hyphal growth. Squares indicate the mean values and bars denote the 95% confidence interval (CI).

The present meta-analysis indicated there was an overall negative effect ( $Q_t = 106.55$ ,  $p = 0.010$ ,  $df = 75$ , effect size = 0.018, 95% CI = −0.112 to 0.076) of *Epichloë*-associated (shoot, root and *Epichloë* strains) exudates on the spore germination and hyphal growth of AMF (Figure 1B). The *Epichloë*-associated exudates reduced (effect size = 0.150, 95% CI = −0.256 to −0.043) the spore germination of AMF (Figure 1B). However, the *Epichloë*-associated exudates increased (effect size = 0.457, 95% CI = 0.245 to 0.669) the hyphal growth of AMF (Figure 1B).

### 3.2. Does *Epichloë* Infection Affect the Symbiotic Phase of Grass–Arbuscular Mycorrhizal Fungi Development?

The present meta-analysis showed an overall negative effect of *Epichloë* infection on belowground AMF ( $Q_t = 203.69$ ,  $p = 0.000$ ,  $df = 119$ , effect size = −0.075, 95% CI = −0.128 to −0.022) and root AMF colonization ( $Q_t = 173.21$ ,  $p = 0.000$ ,  $df = 102$ , effect size = −0.087, 95% CI = −0.143 to −0.030) of grass plants (Figure 2A,B). The *Epichloë* infection reduced root AMF colonization (effect size = −0.088, 95% CI = −0.151 to −0.025) and root AMF concentration (effect size = −0.082, 95% CI = −0.232 to 0.067) of grass plants (Figure 2A). In line with the results of allelopathic effects on AMF growth via root exudations and shoot

litter, there was no change (effect size = 0.005, 95% CI =  $-0.151$  to  $0.161$ ) in the length of extraradical AMF hyphae in the growing soil of *Epichloë*-infected versus non-infected plants (Figure 2A).



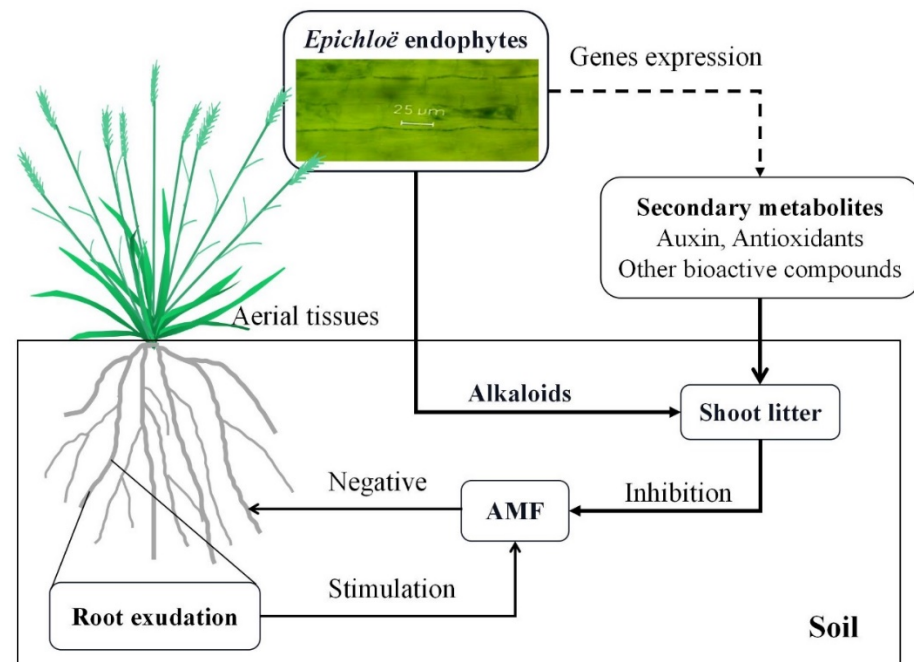
**Figure 2.** The cumulative effect size of *Epichloë* endophytes infection on the symbiotic phase of the grass and arbuscular mycorrhizal fungi (AMF) development among (A) soil and root AMF effects or (B) root AMF colonization of different plant genera. The squares indicate the mean values and the bars denote the 95% confidence interval (CI).

The present meta-analysis also indicated that the root AMF colonization was significantly ( $Q_t = 203.69, p = 0.000, df = 119$ ) decreased (effect size =  $-0.087$ , 95% CI =  $-0.143$  to  $-0.030$ ) in the roots of *Epichloë*-infected versus non-infected plants (Figure 2B). In line with the main effects, the same effects were observed in the *Lolium* (effect size =  $-0.228$ , 95% CI =  $-0.322$  to  $-0.133$ ) and *Leymus* (effect size =  $-0.031$ , 95% CI =  $-0.149$  to  $0.088$ ) grass plants (Figure 2B). However, the opposite results were found in *Festuca* (effect size =  $0.012$ , 95% CI =  $-0.190$  to  $0.214$ ) and *Achnatherum* (effect size =  $0.029$ , 95% CI =  $-0.126$  to  $0.183$ ) grass plants (Figure 2B).

### 3.3. Synthesis of *Epichloë* Effects on the Development of Grass and Arbuscular Mycorrhizal Fungi

We addressed two perspectives to understand or explain the effects of *Epichloë*-infection on AMF from changes of root exudation and shoot litter mediated by *Epichloë*

endophytes, and these variations of AMF in root and soil could contribute to the improvement of plant growth and resistance (Figure 3). The root exudation and shoot litter from the *Epichloë* infected grass plants positively and negatively affected the growth of AMF in the pre-symbiotic phase of grass AMF development compared with the non-infected grass plants (Figure 3). Therefore, there were no obvious changes in the growing soil of *Epichloë* infected versus non-infected grass plants (Figure 3). Collectively, the presence of foliar *Epichloë* reduced root colonization by AMF (Figure 3).



**Figure 3.** The synthesis of potential mechanisms for the effects of *Epichloë* endophytes infection/presence on the developmental processes of grass roots and soil arbuscular mycorrhizal fungi (AMF).

#### 4. Discussion

The ABG interaction considers the spatiotemporal scales involved in the interactions between aboveground *Epichloë* fungal endophytes and belowground AMF within a plant [25]. The spatiotemporal scales could help to understand the coexistence interactions between foliar *Epichloë* endophyte and root AMF in a plant [25]. *Epichloë* systemic endophytes are found in the aerial tissues of 20% ~ 30% of all grasses [7], and AMF colonizes the roots of 80% of angiosperms [26], suggesting that these three-way plant–fungal interactions are ecologically common. Based on the transmission modes of *Epichloë* (vertically transmitted through seeds) and AMF (horizontally colonized grass roots), *Epichloë* endophytes had higher spatiotemporal priority to modify host metabolic and genetic profiles than co-infected AMF [25]. Here, we addressed two perspectives to investigate how foliar *Epichloë* endophyte presence/infection affects AMF growth and the development of grass–AMF symbiosis by altering (1) root exudates and (2) shoot litter decomposition (Figure 3).

##### 4.1. The Effects of *Epichloë* Presence on the Pre-Symbiotic Phase of Grass–AMF Development

The development of AMF symbiosis with plants begins with spore germination, and the related developmental processes depend on the host plants, such as root exudation and plant litter. Root exudates provide chemical signaling to regulate rhizosphere microbial biomass and diversity [3]. Previous meta-analysis and experiment cases showed that the presence of *Epichloë* endophytes stimulated root exudation [8] and altered the composition of root exudates of the host grass [20,21]. For example, *Epichloë*-infected grass roots release more carbon (in the form of phenolic compounds and amines) into the rhizosphere through root exudation and volatiles than non-infected grasses [20,27]. Some secondary metabolites (such

as lipids, flavonoids, and phenolic compounds) could regulate the pre-symbiotic phase of plant-AMF symbiosis [28,29] at the start of the connection of AMF initial mycelium with plant roots [30]. Our meta-analysis showed that root exudation of *Epichloë*-infected plants stimulated AMF growth (Figure 1). Additionally, higher AMF diversity in the rhizosphere has been reported in *Achnatherum inebrians* plants hosting the endophyte *Epichloë gansuensis* in field experiments under drought stress [14]. However, the shoot litter of *Epichloë*-infected plants also affected soil microbes compared to non-infected plants [25,31]. Shoot exudates of *Epichloë*-infected grasses inhibited AMF growth in the present meta-analysis (Figure 1). There was a reasonable explanation that *Epichloë* fungal alkaloids were present in the litter of *Epichloë*-infected plants [32,33] and the soil for more than 50 days [34]. Additionally, flavonoids (recognized as inhibitors of AMF [35]) were higher in the roots of *Epichloë*-infected than in non-infected *Lolium multiflorum* plants [36]. This study showed that shoot and root exudation had opposite effects on AMF growth; therefore, there was no obvious change in soil AMF in *Epichloë*-infected versus non-infected grasses (Figures 1–3). This is in line with a study showing that aboveground *Epichloë*-grass associations do not affect belowground AMF symbionts [37]. The shifts of root-associated soil fungal and bacterial community structures in the *Epichloë*-infected versus non-infected grass plants had been reported in many field and pot cases [8,38–41]. There were different *Epichloë*-associated effects on the AMF growth—*Epichloë* presence reduced the spore germination of AMF strains while promoting the hyphal growth of AMF (Figure 1). This could help explain the variations of soil AMF associated with the presence of an *Epichloë* endophyte. However, the response mechanism of the root-associated microbiome of grass plants to the infection of *Epichloë* endophyte is still unclear. Further research should consider the potential mechanisms underlying these selections of soil microbiome by the infection of *Epichloë* endophytes.

#### 4.2. The Effects of *Epichloë* Infection on the Symbiotic Phase of Grass–AMF Development

The symbiotic phase in which the AMF mycelium contacts and starts to grow along the root surface and enters the root endosphere is determined by the plant's nutritional status and defenses [30]. Root mycorrhizal colonization was lower in *Epichloë*-infected plants than in non-infected plants (Figure 2). In line with two meta-studies, AMF negatively colonized the grass roots of *Epichloë*-infected plants [8,13]. *Epichloë* infection promotes auxin content [42], nutrients [42,43], and biomass accumulation [4,44] in the roots of grass hosts. Therefore, we propose that the presence of *Epichloë* may reduce the beneficial effects of AMF in grass hosts. A previous study supports our suggestion that *Epichloë* endophytes facilitate the invasion of grass hosts by reducing the benefits derived from other mutualisms [13]. Additionally, the hyphal growth of *Epichloë* endophytes in aerial tissues also modified the metabolic profiles (promoting the accumulation of flavonoids, benzoic acids, and cinnamic acids) of the roots of grass hosts [36,45], which may inhibit root colonization by AMF. We proposed that the decrease of AMF colonization in the roots of *Epichloë*-infected versus non-infected grass plants was partly explained by the higher bioactive compounds in the roots of plants infected with *Epichloë* fungal endophytes. Some indirect evidence has been reported that *Epichloë* infection altered the root exudation [20,21], root metabolic profiles [36,42], and root/rhizosphere AMF community [14,37], while whether the relationship among root exudation, root metabolic profiles and root/rhizosphere AMF are shifted in the root–soil relations of *Epichloë*-infected versus non-infected grass plants were still not clear. Generally, the aboveground *Epichloë* endophytes could result in changes to belowground AMF, especially AMF growth and root colonization. This meta-analysis also provides insight into the connection between chemical changes in root exudates and shoot litter for plant–soil interactions in ecosystem processes.

## 5. Conclusions and Further Perspectives

In conclusion, we propose that the presence/infection and hyphal growth of *Epichloë* endophytes in aboveground tissues negatively affected belowground AMF. Root exudation

of *Epichloë*-infected plants stimulated AMF growth in the pre-symbiotic phase, whereas shoot exudates of *Epichloë*-infected grass plants inhibited AMF growth. Additionally, AMF negatively colonized grass hosts hosting an *Epichloë* endophyte. There were no obvious changes in soil AMF. This study should help improve the understanding of the multiple plant–microbe interactions in ABG interactions. This study provides insights and evidence of the capability of the belowground plant and soil processes to be altered by aboveground *Epichloë* endophytes. There was a positive relationship between the AMF colonization rate and the *Epichloë* infection rate of the *Poa bonariensis* population [46]. The *Epichloë*-infected *Hordeum comosum* population had greater AMF colonization than non-infected plants [47]. The concentration of aboveground *Epichloë* endophyte is positively related to AMF concentration in the individual plant [48]. This means that the concentration of both *Epichloë* fungal endophytes and AMF within a grass plant was decreased compared with grass plants infected with either *Epichloë* endophytes or AMF. Some studies also reported that AMF colonization also reduced the concentration of *Epichloë* endophytes and their fungal alkaloid production [49]. Future research should consider (1) how variations in belowground processes from the *Epichloë* endophytes present feedback to the *Epichloë*–grass associations; (2) genetic and metabolic profiles of grass hosts to respond to simultaneous infections and environmental stresses; and (3) simultaneous infections on plant growth and defenses and the applications of multiple plant–microbe interactions in sustainable agriculture.

**Author Contributions:** R.Z. and L.Z. developed the concept; R.Z. collected and analyzed the data, R.Z., L.Z. and X.Z. wrote this manuscript, and all authors revised this manuscript. All authors have read and agreed to the published version of the manuscript.

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