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Article

Biopesticide Compounds of Endolichenic Fungus *Xylaria* sp. Isolated from *Hypogymnia tubulosa*

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Abstract: Endolichenic fungi represent an important ecological group of microorganisms that form associations with photobionts in the lichen thallus. These endofungi that live in and coevolve with lichens are known for synthesizing secondary metabolites with novel structures and diverse chemical skeletons making them an unexplored microbial community of great interest. As part of our search for new phytoprotectants, in this work we studied the endolichenic fungus *Xylaria* sp. isolated from the lichen *Hypogymnia tubulosa*, which grow as epiphyte on the bark of the endemic tree *Pinus canariensis*. From the extract of liquid fermentation, we isolated two new piliformic derivatives, (+)-9-hydroxypiliformic acid (**1**) and (+)-8-hydroxypiliformic acid (**2**), along four known compounds, (+)-piliformic acid (**3**), hexylaconitic acid A anhydride (**4**) and two hydroxyphenylacetic derivatives (**5-6**). Their structures were elucidated based on NMR and HRESIMS data. The extract and compounds were tested for their antifeedant (*Myzus persicae*, *Rhopalosiphum padi* and *Spodoptera littoralis*), antifungal (*Alternaria alternata*, *Botrytis cinerea* and *Fusarium oxysporum*), nematocidal (*Meloidogyne javanica*) and phytotoxic effects on mono and dicotyledonous plant models (*Lolium perenne* and *Lactuca sativa*). Among these compounds **4**, **5** and **6** were effective antifeedants against *M. persicae* and **4** was also active against *R. padi*. Moreover, compound **3** and **4** showed antifungal activity against *B. cinerea* and **4** was the only nematocidal. Phytotoxic test showed that the extract has strong effect against *L. sativa* and *L. perenne* growth. The phytotoxic compounds were **3**, **4** and **5**. (+)-Piliformic acid (**3**) and hexylaconitic A anhydride (**4**) displayed root growth stimulation on low doses and inhibitory effect at higher doses of dicotyledonous specie *L. sativa*.

Keywords: *Xylaria* sp.; endolichenic fungi; (+)-9-hydroxypiliformic acid; (+)-8-hydroxypiliformic acid; biopesticide activity

1. Introduction

Endolichenic fungi represent an important ecological group of microorganisms that form associations with photobionts in the lichen thalli [1]. These fungi live symbiotically inside the lichen, similar to how endophytic fungi inhabit plant tissues [2]. In this symbiosis the lichen provides the living environment and nutrients for the survival of the endolichenic fungi and in turn, they produce a variety of secondary metabolites that help the lichen to accelerate its growth and to protect their host during biotic and abiotic stress. The potential of endolichenic fungi to produce novel structures, diverse skeletons, and extensive bioactivities based on their mutualistic symbiosis with their hosts and unique living environment, has gained significant research attention due to their possible applications in medicine and agriculture [3].

Fungi from the Xylariaceae family are frequent saprotrophs, and have been described as endophytes of phylogenetically diverse plant and lichens species from a variety of ecosystems [4–8]. Furthermore, xylarialean endophytes can be symbiotic or saprotrophic [9]. Endolichenic isolates from

the *Xylaria* genus have demonstrated the ability to produce a great diversity of secondary metabolites with varied bioactivities. Several novel antimicrobial eremophilane sesquiterpenes, named eremoxylarins D-J, have been isolated from *X. hypoxylon* from the lichen *Rhizocarpon geographicum* [10]. Additionally, the nematocidal polyketide grammicin was isolated from *X. grammica* associated with the lichen *Menegazzia sp.* [11]. A variety of compounds, including aplysinopsin, ophiocerin B, and the polyketides piliformic acid and methyl xylariate C, were isolated from *X. venustula*, associated to the lichen *Usnea baileyi* [6]. Furthermore, endolichenic *Xylaria* strains have also been reported to produce novel cyclic peptides, such as the cyclic depsipeptide xylaroamide A from *Xylaria sp.* associated with *Usnea sp.* [12] and a new nitrogen containing phenolic compound with cytotoxic activity has been discovered from *X. psidii* [13].

Secondary metabolites produced by *Xylaria species* have also shown phytoprotectant activities, including herbicidal, fungicidal and insecticidal [14]. Various endophytic isolates of *Xylaria sp.* have been reported to produce griseofulvin [15–17], a metabolite that inhibits the growth of several species of fungal plant pathogens [17]. *Xylaria sp.*, isolated from *Vitis labrusca*, produce diplosporin [18], a compound toxic to the polyphagous insect *Spodoptera frugiperda* when added to artificial diet [19]. Additionally, cytochalasin E isolated from the endophytic fungus *Xylaria sp.* showed phytotoxic effects on *Lactuca sativa* and *Raphanus sativus* seedlings [20].

The excessive use of synthetic pesticides needed to increase crop production and productivity by controlling pathogens that harm the crops [21] has led to pathogenic resistance, environmental problems affecting soil biodiversity, beneficial insects and aquatic life, and has been linked to detrimental effects on human health [22]. Biopesticides, of natural origin, are an alternative to control plant pathogens with less toxic effects on the ecosystem [23]. In this work, the endolichenic fungal strain HYP6 isolated from the lichen *Hypogymnia tubulosa* was fermented in malt liquid medium and extracted with ethyl acetate (EtOAc). The fractionation of the extract gave two new compounds and four known metabolites that have been identified based on their spectroscopic data. The extract and pure compounds were tested for their antifeedant (*Myzus persicae*, *Rhopalosiphum padi* and *Spodoptera littoralis*), nematocidal (*Meloidogyne javanica*), fungicidal (*Alternaria alternata*, *Botrytis cinerea* and *Fusarium oxysporum*) and phytotoxic effects on mono and dicotyledonous plant models (*Lolium perenne* and *Lactuca sativa*).

2. Results and Discussion

2.1. Fungal Identification

The endolichenic fungal strain HYP6 was isolated from the lichen *Hypogymnia tubulosa* from the bark of the endemic tree, *Pinus canariensis*, collected in Tenerife Island (Spain). The rDNA ITS was sequenced and compared with those deposited in the NCBI-Genbank. To refine the taxonomic assignment a total of 55 related sequences were used to conduct maximum likelihood phylogenetic analysis using Tamure-3 parameter (T92+G) model and a bootstrap test with 5000 runs. The results indicated that strain HYP6 is closely related to *Xylaria arbuscula* (Figure 1). Based on multigene phylogenetic analyses [24], *X. arbuscula* is actually considered a species complex, which predominantly including species from the tropical *Xylaria* group [25]. According to these studies and the results described above, the fungal strain HYP6 was identified as *Xylaria sp.* within the *X. arbuscula* complex.

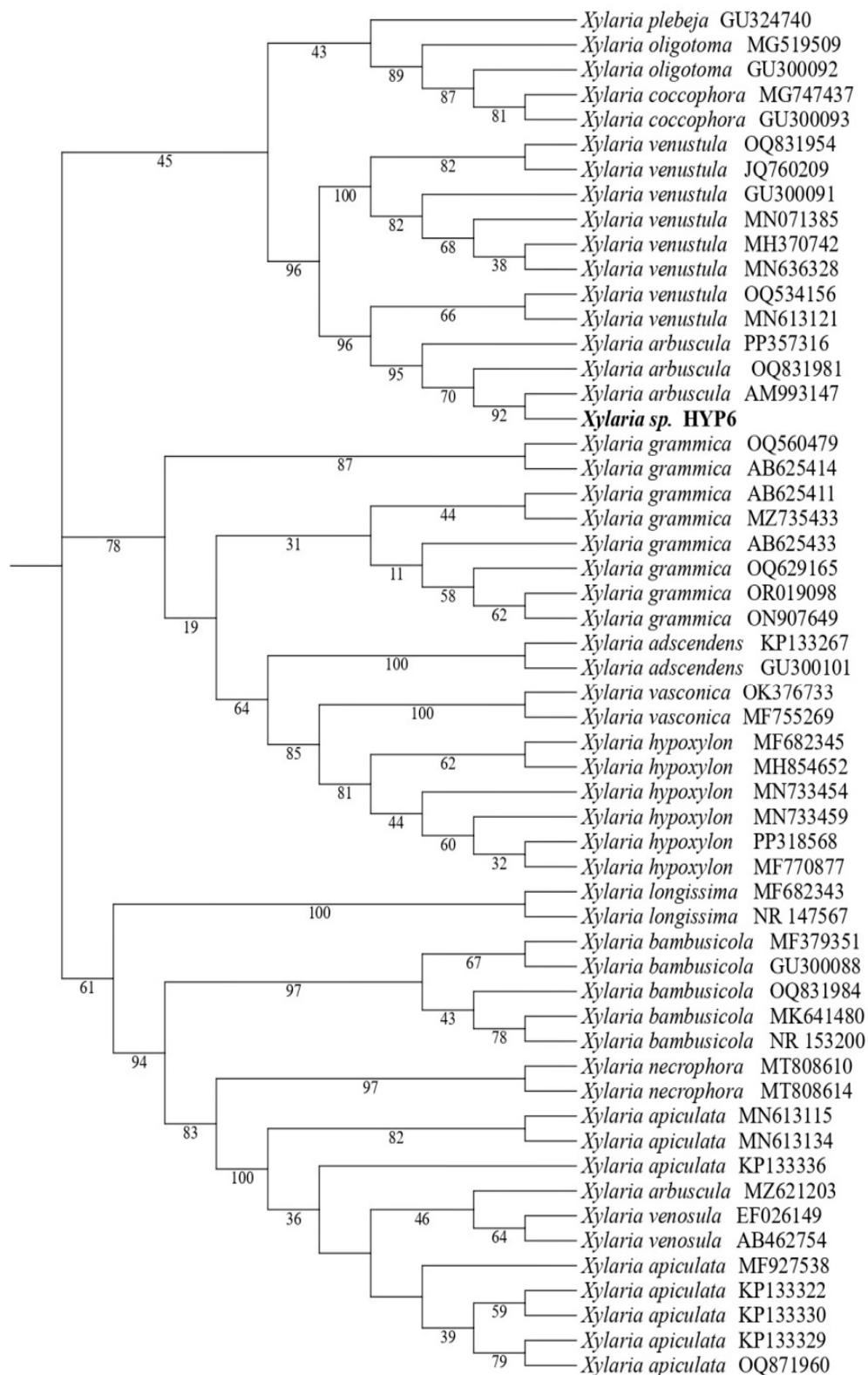


Figure 1. Phylogenetic maximum likelihood tree (T92+G) of *Xylaria* sp. (HYP6) based on selected ITS rDNA sequences obtained from GenBank. Confidence values from a 5000-replicate bootstrap analysis are shown at each branch nodes.

2.2. Structure Elucidation

The fractionation of the extract from the culture broth of endolichenic fungi *Xylaria sp* fermented in malt liquid medium led to the isolation and structural identification of two new piliformic derivatives, (+)-9-hydroxypiliformic acid (**1**) and (+)-8-hydroxypiliformic acid (**2**) together with the known compounds, (+)-piliformic acid (**3**) as the main metabolite of extract, hexylaconitic acid A anhydride (**4**), 2-hydroxyphenylacetic acid (**5**) and 4-hydroxyphenylacetic acid (**6**) (Figure 2).

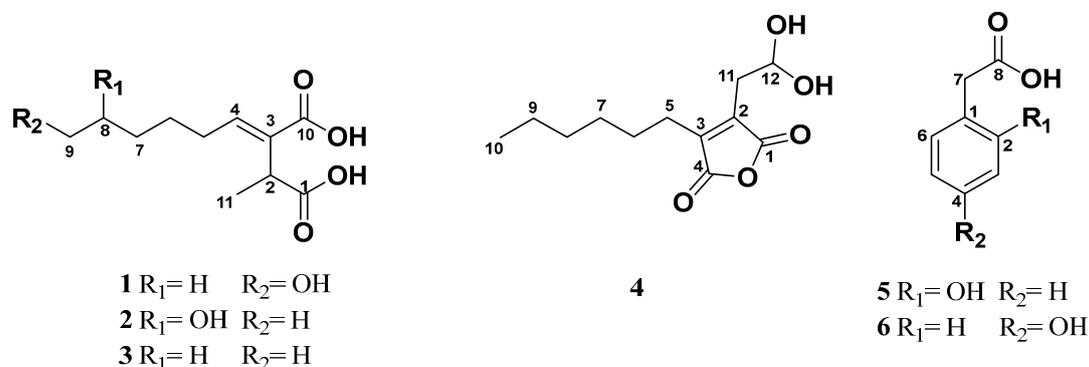


Figure 2. Compounds of endolichenic *Xylaria sp*. (HYP6).

Compound **1** was isolated as colourless oil. Its molecular formula was determined as $C_{11}H_{18}O_5$ by the positive ion mode HRESIMS data at m/z 253.1059 $[M+Na]^+$ (calcd. for $C_{11}H_{18}O_5Na$, 253.1052), indicating three degrees of unsaturation. The IR spectrum showed absorption bands at ν_{max} 3355, 1689 and 1637 cm^{-1} , which indicated the presence of a hydroxyl group, carboxylic acid and double bond, respectively. In the 1H NMR spectrum (Table 1 and Figure S1) were observed resonances of one methyl group at δ 1.31 (d, $J=7.1$ Hz, H-11) coupled with a methine at δ 3.64 (dq, $J=7.1$ Hz, H-2) and one olefinic proton at δ 6.85 (t, $J=7.6$ Hz). Other signals observed were multiplets attributed to the protons of three overlapped methylenes of an aliphatic chain at δ 1.42-1.66 and the signal at δ 3.55 (t, $J=6.6$ Hz, H-9) of a hydroxymethylene group. Analysis the ^{13}C NMR (Table 1 and Figure S2) data in combination with gHSQC spectrum (Figure S4) showed eleven carbon signals, two carbonyl groups at δ 178.0 (C-1) and 170.3 (C-10), two olefinic carbons at δ 134.4 (C-3) and 144.7 (C-4) of a trisubstituted double bond, one oxygenated methylene at δ 62.8 (C-9), one methine at δ 39.0 (C-2), one methyl group at δ 16.4 (C-11), as well as four methylene carbons at δ 29.4 (C-5), 29.6 (C-6), 26.7 (C-7) and 33.4 (C-8) in aliphatic region. The 1H - 1H COSY NMR spectrum of **1** (Figures 3 and S3) showed coupling between the H-2 and H-11 and also established a partial structure from the olefinic proton at δ 6.85 (H-4) along the aliphatic chain with a spin system of methylenic protons H-5/ H-6/ H-7/ H-8/ H-9.

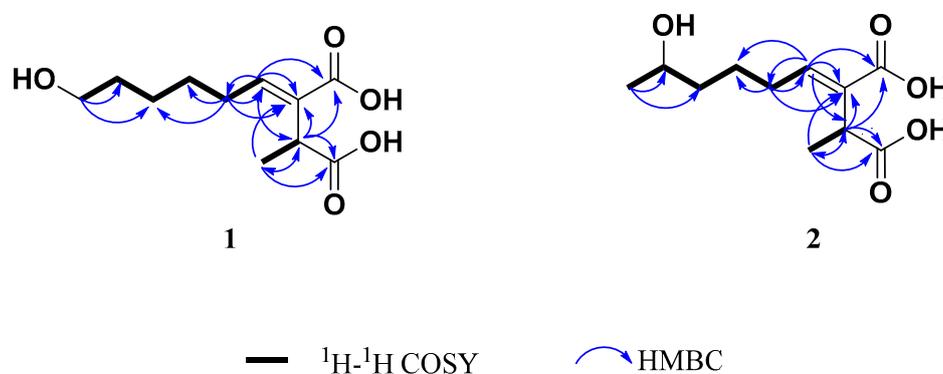


Figure 3. Key 1H - 1H COSY and selected HMBC correlations of compounds **1** and **2**.

Table 1. ¹H NMR and ¹³C NMR spectra of compounds **1** and **2** in CD₃OD.

Position	1		2	
	δ_H , mult (J in Hz)	δ_C , type	δ_H , mult (J in Hz)	δ_C , type
1	-	178.0, C	-	177.8, C
2	3.63, q (7.1)	39.0, CH	3.64, qd (7.1, 1.9)	38.7, CH
3	-	134.5, C		134.3, C
4	6.82, t (7.6)	144.7, CH	6.85, t (7.6)	145.0, CH
5	2.26, m	29.4, CH ₂	2.26, m	29.4, (29.4) ^a , CH ₂
6	1.51, m	29.6, CH ₂	1.61, m 1.52, m	26.0, (26.0) ^a , CH ₂
7	1.43, m	26.7, CH ₂	1.50, m	39.6, (39.7) ^a , CH ₂
8	1.56, m	33.4, CH ₂	3.74, sext (6.1)	68.2, (68.3) ^a , CH
9	3.55, t (6.6)	62.8, CH ₂	1.16, d (6.1)	23.5, CH ₃
10	-	170.3, C	-	169.9, C
11	1.31, d (7.2)	16.4, CH ₃	1.31, d (7.1)	16.3, CH ₃

^a data in () for stereoisomer of hydroxyl group in n-alkyl chain.

The low field resonance of a methine proton at δ 3.64 (H-2) in the CH-CH₃ grouping indicated deshielding effects due to carboxyl groups in the molecule. The carboxyl groups are located on C-1 and C-10, as supported by the HMBC correlations between H-2 and C-1/ C-3/ C-4/C-10/C-11 and between H-11 methyl with C-1/C-2/C-3 (Figure 3). The position of the double bond was confirmed by the correlations previously indicated for H-2 and those observed between H-4 and C-2/ C-3/ C-5/ C-10. The HMBC experiment (Figure S5) showed also connectivities of H-5 with C-2, C-3, C-4, C-6 and C-7, and H-9 with C-7 and C-8, which confirmed the structure of compound **1** closely related to the known fungal metabolite (+) piliformic acid (**3**) [26,27], also isolated from this fungal strain and whose structure has been obtained by chemical synthesis [28]. The configuration of double bond was assigned *E* based on the NOESY between H-3 and H-5 (Figure S6), however with the available spectroscopic data the configuration at C-2 in compound **1** could not be defined. Thus, this new compound was identified as (+)-(*E*)-9-hydroxypiliformic acid (**1**).

Compound **2** exhibited an ion peak in its HRMS (ESI+) at m/z 253.1058 [M+Na]⁺, corresponding to the molecular formula C₁₁H₁₈O₅Na, which was identical to that compound **1**. The ¹H-NMR (Table 1 and Figure S8) spectrum displayed signals similar to those observed in **1**, but now the hydroxymethylene group at C-9 has been substituted in **2** by a new methyl group. Thus, the H-9 protons of the methyl group resonate as doublet at δ 1.16 (3H, d, J=6.2 Hz) coupled with the proton of a new hydroxyl group at δ 3.74 (1H, sext, J= 6.1 Hz). Furthermore, the ¹³C NMR spectrum showed an additional methyl carbon at δ 23.5.8 (C-9) and a methine group at 68.2 (C-8) (Figure S9). The new hydroxyl group was assigned to C-8 considering the ¹H-¹H COSY crosspeak of H-9 (δ 1.16)/ H-8 (δ 3.74) together the HMBC correlations of H-9 methyl group with C-7 and C-8 (Figures 3 and S12). A detailed analysis of the ¹³C NMR (Table 1 and Figure S9) showed double signals for C-5, C-6, C-7 and C-8, which indicated that compound **2** was the mixture of two epimers at C-8 with a ratio 2:1 according to the height of double peaks. These observations, together with a careful analysis of the COSY, HMBC and NOESY experiments, led to the identification compound **2** as a mixture (ratio 2:1) of two epimers of (+)-(*E*)-8-hydroxypiliformic acid.

The known compound **4** was identified as 2-carboxymethyl-3-n-hexyl-maleic acid anhydride and named hexylaconitic acid A anhydride. Its molecular formula $C_{12}H_{16}O_5$ was deduced by HREIMS. The 1H and ^{13}C NMR spectral data are consistent with those reported for 2-carboxymethyl-3-n-hexyl-maleic acid anhydride, which was first isolated as a natural product from *Aspergillus niger* [29] and later obtained by chemical synthesis [30,31]. Now in this work, we have completed the assignments of the proton and carbon resonances in the 1H and ^{13}C spectra using 2D-NMR experiments (COSY, HSQC and HMBC), which had not been previously assigned. The configuration *Z* of double bond was also confirmed by the crosspeak of H₂-5/H₂-11 in the NOESY spectrum (Figure S20).

Another known compounds isolated from this fungus were identified as (+)-piliformic acid (**3**) [26–28], a fungal metabolite isolated previously from different fungi belonging to the *Xylaria* genus [32], and the phenolic compounds, 2-hydroxyphenylacetic acid (**5**) [33] and 4-hydroxyphenylacetic acid (**6**) [34]. In the experimental part we provide complementary NMR and MS data of these compounds.

2.3. Biopesticide Activity

The EtOAc extract and compounds obtained from the fermentation of HYP6 were tested for their biopesticide properties against insect pest (*Myzus persicae*, *Rhopalosiphum padi* and *Spodoptera littoralis*), the plant parasitic nematode *Meloidogyne javanica* and fungal phytopathogens (*Alternaria alternata*, *Botrytis cinerea* and *Fusarium oxysporum*). The extract showed strong antifeedant effects against *M. persicae* (EC₅₀ value of 10.9 $\mu\text{g}/\text{cm}^2$) followed by a moderate activity against *R. padi* and it was not active on *S. littoralis*. This extract was also antifungal to *B. cinerea* and moderately active on *F. oxysporum* and *A. alternata* but it did not have any nematocidal effect (Table 2).

When tested the compounds on insect pests (Table 2), **4** showed a strong antifeedant activity against *M. persicae* (EC₅₀ value of 1.6 $\mu\text{g}/\text{cm}^2$) and *R. padi* (EC₅₀ value of 8.9 $\mu\text{g}/\text{cm}^2$), being more effective than the extract. Additionally, compounds **5** and **6** showed a selective antifeedant activity against *M. persicae* with EC₅₀ value of 4.5 $\mu\text{g}/\text{cm}^2$ and EC₅₀ value of 15.5 $\mu\text{g}/\text{cm}^2$, respectively. None of the tested compounds showed any antifeedant effects against *S. littoralis*. Compound **4** also exhibited a strong inhibition of mycelial growth of *B. cinerea* (EC₅₀ value of 0.12 mg/ml) and moderate against *A. alternata* (EC₅₀ value of 0.24 mg/ml), whereas compound **3** was moderately active on *B. cinerea* (EC₅₀ value of 0.36 mg/ml). When tested against *M. javanica*, compound **4** was very active with an LD₅₀ value of 0.10 mg/ml.

The extract and compounds were also tested for phytotoxic effects on seeds of mono- and dicotyledonous plant species (*L. perenne* and *L. sativa*) (Figure 4). The extract of *Xylaria sp.* strongly inhibited the germination of *L. perenne* with 88.2% inhibition after 7 days compared to the control. The extract also exhibited phytotoxic effects on *L. perenne*, with 89.3% and 100% inhibition of leaf and root growth, respectively and affected the root growth of *L. sativa* (66.8% inhibition). Among the compounds tested, compound **3** affected the germination of *L. sativa* (86.7% inhibition), decreased moderately root growth of *L. perenne* (56.8% inhibition) and affected the root growth of *L. sativa* (72.6% inhibition). Compound **4** reduced significantly the germination of *L. perenne* and *L. sativa* (72.2% and 94.7% inhibition, respectively) with a strong inhibition of root growth (100% and 97.9% inhibition, respectively). Additionally, compound **4** also decreased the leaf growth of *L. perenne* (73.1% inhibition). Compound **5**, with reported phytotoxic effects [35,36], showed significant effect on root growth on *L. perenne* and *L. sativa* root growth (85.4% and 91.3% inhibition, respectively) (Figure 4). The most phytotoxic compounds (**3**, **4**) were further tested in dose-response experiments against the two plant species. At lower concentrations, both compounds stimulated the root growth of *L. sativa*. Compound **3** increased the root length up to 170% at a concentration of 0.1 mg/ml and **4** up to 200% at 0.05 mg/ml.

Table 1. Biocidal effects of extract (EtOAc) and compounds (2–6) against insect pests (*Spodoptera littoralis*, *Myzus persicae*, *Rhopalosiphum padi*), the nematode *Meloidogyne javanica* and phytopathogenic fungi (*Alternaria alternata*, *Botrytis cinerea* and *Fusarium oxysporum*).

Target	Effect	Treatment	Activity (%)	EC ₅₀ ^c /LD ₅₀ ^d
<i>S. littoralis</i>	%FI ^a	Extract	42.9±11.4	
		2	33.0±14.9	
		3	18.0±8.4	
		4	37.8±18.7	
		5	9.1±6.3	
		6	17.7±11.7	
<i>M. persicae</i>	%SI ^a	Extract	81.5±6.8	10.9 (7.2-16.4)
		2	25.5±6.3	
		3	50.4±8.9	
		4	100.0±0.0	1.6 (0.5-4.9)
		5	98.9±0.6	4.5 (2.0-10.0)
		6	91.9±2.3	15.5 (12.0-20.0)
<i>R. padi</i>	%SI ^a	Extract	61.5±6.0	
		2	20.6±6.4	
		3	30.3±8.8	
		4	91.0±3.8	8.9 (4.6-7.5)
		5	61.2±9.2	
		6	43.0±7.9	
<i>M. javanica</i>	%Mortality ^b	Extract	2.0±0.7	
		2	2.2±0.7	
		3	28.4±6.6	
		4	100.0±0.0	0.10 (0.10-0.11)
		5	26.4±5.8	
		6	29.5±5.6	
<i>A. alternata</i>	% MGI ^b	Extract	58.5±2.4	
		2	27.3±3.7	
		3	47.3±5.1	
		4	63.7±1.4	0.24 (0.20-0.28)
		5	32.2±5.6	
		6	42.5±6.0	
<i>B. cinerea</i>	% MGI ^b	Extract	75.9±9.7	
		2	19.7±6.2	
		3	62.1±7.6	0.36 (0.25-0.51)
		4	84.3±4.5	0.12 (0.09-0.15)
		5	0.5±8.8	
		6	5.8±6.2	
<i>F. oxysporum</i>	% MGI ^b	Extract	61.9±6.5	
		2	18.7±3.1	
		3	42.3±3.6	
		4	52.3±2.6	
		5	32.6±9.1	
		6	35.7±3.3	

^a Feeding/Settling Inhibition (%FI / %SI), doses tested at 100µg/cm² or 50.0 µg/cm² for extract or compounds. ^b % Mortality J2 and % MGI, doses tested at 1 mg/ml or 0.5 mg/ml for extract or compounds. ^c Effective dose EC₅₀ (µg/cm²) feeding/settling or EC₅₀ (mg/ml) MGI and 95% confidence limits. ^d Lethal Dose (mg/ml) to give 50% mortality of *M. javanica* and 95% confidence limits.

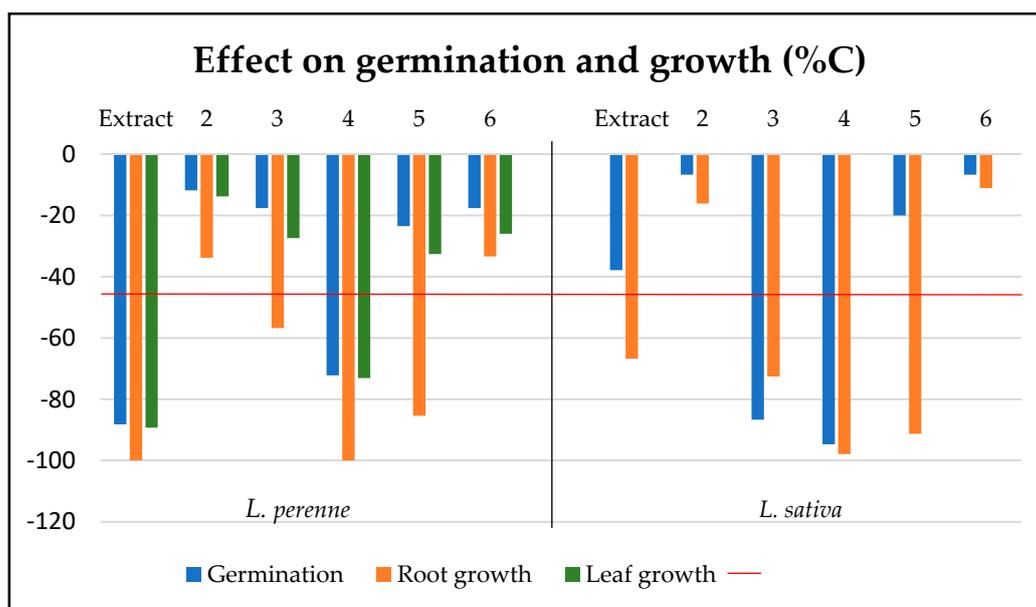


Figure 4. Phytotoxic effects of extract (EtOAc) and compounds (2-6) on *Lolium perenne* and *Lactuca sativa* (doses tested at 0.4 and 0.2 mg/ml for the extract and 2-6 respectively).

In this work, (+)-piliformic acid (**3**) showed moderate antifungal effects against *B. cinerea*. Similarly, previous reports showed moderate activity of **3** against *Colletotrichum gloeosporioides*, one of the phytopathogenic fungus responsible for the anthracnose disease [37]. Compound **4**, with aphid antifeedant, antifungal and nematicidal effects was identified as hexylaconitic A anhydride. This compound (**4**) isolated from *Aspergillus niger* [29,38] and *Aspergillus tubigensis* [39] has been previously reported for its fungicidal activity against *Neurospora crassa* [39], but it was found inactive against other fungal plant pathogens (*Gaeumannomyces graminis* var. *tritici*, *Rhizoctonia solani*, and *Phytophthora cinnamomic*) [38]. This may indicate a species-dependent antifungal effects that could be used to target specific fungal pathogens. However, this is the first report on the aphid antifeedant and nematicidal effects of compound **4**. Phenolic acids related to **5** and **6**, are known for their insecticidal activities [40] and play an important role in the plant resistance against insect pest [41]. For example, previous reports showed that phenylacetic compounds isolated from *Streptomyces gramineus* have insecticidal activity against *Thrips palmi*, also known as melon thrips, a sap-sucking phytophagous insect [42].

The phytotoxic effects of compounds **3** and **4** are consistent with those of the analogous compounds of alkylitaconic acid derivatives, which promoted radicle growth at low doses, while inhibited the seedlings growth at higher doses particularly in dicotyledonous species [35,43]. Additionally, Mondal et al. [44] reported that compound **4** stimulates germination and seedling growth in cauliflower at low ppm concentrations. This behaviour could be explained by hormetic effects of toxic agent showing a biphasic response, promoting early seedling development (and potentially later plant growth) at low concentrations, while inhibiting growth at high concentrations [45,46]. This work has shown for the first time the potent phytotoxic effects of (+)-piliformic acid (**3**) and hexylaconitic A anhydride (**4**) on mono and dicotyledonous plant (*L. perenne* and *L. sativa*) and stimulating effect the growth of *L. sativa* root at lower dose.

3. Materials and Methods

3.1. General Experimental Procedures

Optical rotations were measured on MCP 150-Anton Paar polarimeter (Anton Paar, Seelze-Letter, Germany) at a temperature of 25°C. IR spectra were recorded with a Cary 630 FTIR spectrophotometer (Agilent, Santa Clara, CA, USA). ^1H , ^{13}C and 2D (COSY, gHSQC, HMBC, and NOESY) NMR spectra were recorded in CDCl_3 or CD_3OD solution (Aldrich, St. Louis, MO, USA) on

a Bruker Avance II-500 spectrometer equipped with a 5 mm TCI inverse detection cryo-probe (Bruker Biospin, Fallanden, Switzerland). Chemical shifts are given in ppm (δ), referenced to solvent signal (CDCl_3 , δ_{H} 7.26 and δ_{C} 77.0; CD_3OD_3 , δ_{H} 3.31 and δ_{C} 49.0. The HRESIMS mass spectra were obtained using a Waters LCT Premier XE mass spectrometer (Manchester, UK). Sephadex LH-20 (Sigma-Aldrich, St. Louis, MO, USA) and silica gel 60 (40 \pm 63 μm , Merck, Darmstadt, Germany) was used for column chromatography. TLC was performed with precoated silica gel 60 F254 plates (Merck, Darmstadt, Germany). Flash chromatography separations were carried out on a Biotage Isolera Prime (BIOTAGE, Uppsala, Sweden) apparatus equipped with a UV detector (200-400 nm) using a prepacked RediSep flash column of 35g of silica gel (90 ml, 13 cm \times 3 cm; Teledyne Isco, Lincoln, NEB, USA). Semipreparative HPLC was performed on a Beckman System Gold 125P equipped with a diode-array detector Beckman Coulter 168 (Brea, CA, USA) using a semi-preparative Beckman Ultrasphere ODS (10.0 \times 250 mm, 5 μm) column.

3.2. Lichen Material and Isolation of the Endolichenic Fungus HYP6

The fungal strain HYP6 was isolated from the lichen *Hypogymnia tubulosa* collected in Tenerife Island (Las Lagunetas, 28°24'47.2"N, 16°24'11.2"W) from the bark of endemism *Pinus canariensis*, dominant tree of the Canarian pine forest ecosystem. The samples were placed into sterile polybags and transported under refrigeration in a box container until isolation processing, within 24 hours of collection. Collected lichen thalli were initially washed with distilled water to remove excess dirt. For surface sterilization, thalli were cut into manageable portions followed by successively dipping in a 70% ethanol solution for 1 minute, 1% sodium hypochlorite for 10 minutes and finally washing with 70% ethanol for 1 minute. The surface sterilized thalli were dried on sterilized paper and cut into 1cm pieces. Then, lichen explants were inoculated in Potato Dextrose Agar (PDA) and Malt Extract Agar (MEA) media in petri dishes, containing 50 mg/L of the antibiotic chloramphenicol to avoid bacterial contamination. The plates were incubated at 24°C for 2-3 weeks in darkness with daily observation of the emerging fungal colonies. The new fungal mycelia that grew from the lichen explants were transferred to fresh plates containing the respective medium and various subcultures were performed until pure cultures were obtained.

3.3. Molecular Characterization of HYP6 Strain

For the extraction of DNA of the fungal strain the DNeasy Plant mini kit (Qiagen GmbH, Hilden, Germany, Cat. No 69104) was used. The ITS rDNA region was PCR-amplified using the oligonucleotide primers ITS1 (5'-TCCGTAGGTGAACCTGCGG-3') and ITS4 (5'-TCCTCCGCTTA TTGATATGC-3') with the following steps: Genomic DNA (100–200 ng) was amplified on a PTC-200 Thermal Cycler (MJ Research, San Diego, CA, USA) in a 25 μL final volume with the AmpONE Taq DNA polymerase PCR kit (GeneAll, Seoul, Korea) for 35 cycles (95 °C, 1 min; 50 °C, 20 s; 72 °C, 1.5 min) after an initial denaturation (95 °C, 2 min) and followed by a final extension (72 °C, 7 min). Amplicons were checked by agarose gel (1%) electrophoresis, purified using the EXO-SAP-IT kit (Affimetrix-USB; Thermo Fisher Scientific, Waltham, MS, USA), and sequenced on an AB 3500 Genetic Analyzer (Thermo Fisher Scientific, Waltham, J. *Fungi* **2021**, *7*, 1093 of 11 MS, USA) at the University of La Laguna (La Laguna, Spain) genomic service. The consensus sequence of the rDNA ITS region was generated using aligner software Bioedit (version 5.09; Tom Hall, Department of Microbiology, North Carolina State University, Raleigh, NC, USA) and compared with those published in NCBI (<https://www.ncbi.nlm.nih.gov/>) database by using the online BLAST program (<https://www.ncbi.nlm.nih.gov/BLAST>, accessed on 29 May 2024). The related sequences with high percent identity were aligned using the MEGA11.0 program (Mega Limited, Auckland, New Zealand, Tamura K. 2021) [47] and a phylogenetic tree was constructed using maximum likelihood method with a Tamure-3 parameter (T92+G) model and a bootstrap test with 5000 runs. The display and annotation of the phylogenetic tree was performed using the online tool: The Interactive Tree Of Life (<https://itol.embl.de>, accessed on 31 May, 2024).

3.4. Cultivation of HYP6

The endophytic strain HYP6 was cultivated on PDA solid medium for 10 days at 25 °C. Sterile water (10ml) was added to each petri dishes and the surface of mycelium was gently scraped with a spatula to obtain a suspension of mycelium. This suspension was cultivated in 250 ml Erlenmeyer flasks containing 50 ml of MEB medium (17 g/l of malt extract, 3 g/l of peptone, pH adjusted at 5.4±0.2) for 5 days at 25 °C on a rotary shaker (120 rpm) to prepare the seed culture. Ten Erlenmeyer flasks (500 ml) with 200 ml of fresh MEB medium each were inoculated with 10 ml of seed culture and cultured at 25 °C under continuous agitation (120 rpm) for 14 days.

3.5. Extraction and Isolation of Compounds

The fermentation broth was filtered under reduced pressure through a Buchner funnel using cheesecloth (25 µm pore diameter) to separate the mycelium. The resulting filtrate was subjected to a three-cycle liquid/liquid extraction with ethyl acetate (EtOAc), dried over Na₂SO₄, and the organic phase was concentrated under reduced pressure to yield the crude HYP6 extract (817.5 mg).

The EtOAc extract was fractionated by column chromatography on silica gel eluting with a gradient of increasing polarity of n-hexane/EtOAc and then Cl₂CH₂/MeOH. The fractions obtained were analysed by TLC and those with high similarity were combined to afford five main fractions. Fraction 2 was separated by media pressure chromatography using a Biotage Isolera Prime equipment through on 35g Si pre-packed flash cartridge column, eluted with n-hexane/EtOAc mixtures of increasing polarity (70:30–25:75) at 10 ml/min to afford compound 3 (33.7 mg) and 4 (13.5 mg). Fraction 3 was further purified by Sephadex LH-20 with a mixture of CH₂Cl₂:MeOH (1:1) to give compounds 5 (11.3 mg) and 6 (8.4mg). Fraction 4 was separated by Sephadex-LH-20 using an isocratic mixture of n-hexane:CH₂Cl₂:MeOH (1:1:1) to yield four fractions (Fr. 4.1-Fr.4.4). Fr.4.3 was further separated by reverse semi-preparative HPLC (Ultrasphere ODS, 10.0 × 250 mm, 5 µm) using a gradient solvent system of H₂O/MeOH (80:20-30:70) at a flow rate of 3 ml/min to afford compound 2 (18.4 mg, t_R=42.6 min). Similarly, the reverse semi-preparative HPLC of subfraction FR.4.4 eluted with H₂O:MeOH (60:40) gave compound 1 (4.6 mg, t_R=32.0 min).

3.5.1. (+)-9-Hydroxypiliformic Acid (1)

Colourless oil; $[\alpha]_D^{25} +16$ (c 0.5, CHCl₃); IR ν_{\max} : 3355, 2940, 2834, 1689, 1637, 1215, 1020 cm⁻¹; ¹H NMR data (CD₃OD, 500 MHz,) see Table 1 and Figure S1; ¹³C NMR data (CD₃OD, 125 MHz) see Table 1 and Figure S2; HRMS (ESI⁺) *m/z* 253.1059 [M+Na]⁺ (calcd. C₁₁H₁₈O₅Na, 253.1052).

3.5.2. (+)-8-Hydroxypiliformic Acid (2)

Colourless oil; $[\alpha]_D^{25} +30$ (c 0.5, CHCl₃); IR ν_{\max} : 3324, 2945, 2849, 1684, 1636, 1226, 1015 cm⁻¹; ¹H NMR data (CD₃OD, 500 MHz) see Table 1 and Figure S8; ¹³C NMR data (CD₃OD, 125 MHz) see Table 1 and Figure S9; HRMS (ESI⁺) *m/z* 253.1058 [M+Na]⁺ (calcd. C₁₁H₁₈O₅Na, 253.1052).

3.5.3. (+)-Piliformic Acid (2-Hexylidene-3-methylsuccinic Acid) (3)

White solid; $[\alpha]_D^{25} +24$ (c 0.5, CHCl₃); IR ν_{\max} : 2927, 2859, 1685, 1637, 1418, 1241 cm⁻¹; ¹H NMR (CDCl₃, 500 MHz): δ 0.89 (3H, t, J = 6.9 Hz, H-9), 1.31 (4H, m, H-7, H-8), 1.35 (3H, d, J = 7.1 Hz, H-11), 1.48 (2H, br t, J = 6.8 Hz, H-6), 2.23 (2H, m, H-5), 3.63 (1H, q, J = 7.1 Hz, H-2), 7.01 (1H, t, J = 7.5 Hz, H-4); ¹³C NMR (CDCl₃, 125 MHz): δ 180.4 (C-1), 37.7 (C-2), 131.4 (C-3), 147.4 (C-4), 28.9 (C-5), 28.2 (C-6), 31.6 (C-7), 22.6 (C-8), 14.1 (C-9), 172.3 (C-10), 15.1 (C-11); HRMS (ESI⁻) *m/z* 213.1129 [M-H]⁻ (calcd. C₁₁H₁₇O₄, 213.1127).

3.5.4. Hexylaconitic Acid A Anhydride (2-Carboxymethyl-3-n-hexyl-maleic Acid Anhydride) (4)

Yellowish oil; IR ν_{\max} : 2929, 2859, 1855, 1825, 1763, 1718, 1273 cm⁻¹; ¹H NMR (CDCl₃, 500 MHz): δ 0.88 (3H, t, J=6.7 Hz, H-11), 1.30 (6H, m, H-8, H-9, H-10), 1.59 (2H, quint, J=7.5 Hz, H-7), 2.49 (2H, t, J=7.7 Hz, H-6), 3.56 (2H, s, H-11), 8.45 (1H, br s, -OH); ¹³C NMR (CDCl₃, 125 MHz): δ 165.3 (C-1), 148.3

(C-2), 135.6 (C-3), 165.2 (C-4), 25.1 (C-5), 27.6 (C-6), 29.3(C-7, C-11), 31.4 (C-8), 22.5 (C-9), 14.1 (C-10), 173.2 (C-12); HRMS (ESI) m/z 239.0920 [M-H]⁻ (calcd. C₁₂H₁₅O₅, 239.0919).

3.5.5. 2-Hydroxyphenylacetic Acid (5)

White solid; ¹H NMR (CD₃OD, 500 MHz): δ 3.58 (2H, s, H-7), 6.77 (2H, m, H-3, H-5), 7.08 (1H, dd, J= 7.4, 1.2 Hz, H-4), 7.11 (1H, td, J=7.4, 1.6 Hz, H-6); ¹³C NMR (CD₃OD, 125 MHz): δ 122.8 (C-1), 156.7 (C-2), 115.9 (C-3), 129.3 (C-4), 120.5 (C-5), 132.1 (C-6), 36.6 (C-7), 176.3 (C-8); HRMS (ESI) m/z 151.0400 [M-H]⁻ (calcd. C₈H₇O₃, 151.0395).

3.5.6. 4-Hydroxyphenylacetic Acid (6)

White solid; ¹H NMR (CD₃OD, 500 MHz): δ 3.48 (2H, s, H-7), 6.73 (2H, d, J=8.5 Hz, H-3 and H-5), 7.08 (2H, d, J=8.5 Hz, H-2 and H-6); ¹³C NMR (CD₃OD, 125 MHz): δ 126.8 (C-1), 131.3 (C-2, C-6), 116.2 (C-3, C-5), 157.3 (C-4), 41.1 (C-7), 176.4 (C-8). HRMS (ESI) m/z 151.0395 [M-H]⁻ (calcd. C₈H₇O₃, 151.0395).

3.6. Antifungal Bioassay

To evaluate the antifungal activity three different phytopathogenic fungi were used: *Fusarium oxysporum*, *Alternaria alternata* and *Botrytis cinerea*. These strains come from the fungal collection at Instituto de Productos Naturales y Agrobiología-CSIC (Tenerife, Spain) and the colonies were maintained on PDA medium plates in darkness at 25 °C.

Based on the protocol described before [48], the in vitro mycelial growth inhibition assay was conducted in 12-well plates (Falcon) using a modified PDA-dilution method supplemented with the addition of 0.05 mg/ml of methyltetrazolium salts (MTT). Extract and pure compounds dissolved in ethanol (EtOH) were tested at different concentrations (extract at 1 mg/ml and compounds at 0.5, 0.25, 0.1 and 0.05 mg/ml) and were incorporated into the culture medium before plates were poured. EtOH was used as a negative control and all treatments were replicated four times. Fungal colonies were digitalized and measured using ImageJ (<http://imagej.nih.gov/ij/>, accessed on March 18, 2024) [49] after the incubation of the plates at 25 °C in darkness for 48 h. The mycelial growth inhibition (%MGI) was calculated as: %MGI = (C - T/C) × 100, where C is the diameter of the control colonies and T is the diameter of the test colonies. Data were analysed with STATGRAPHICS statistical analysis software (Centurion XVI, version 16.1.03) and EC₅₀ values (effective dose to obtain 50% of inhibition) were determined by regression analysis (%MGI on Log-dose).

3.7. Antifeedant Bioassay

Spodoptera littoralis colonies were reared on an artificial diet [50], while *Myzus persicae* and *Rhopalosiphum padi* colonies were maintained on bell pepper (*Capsicum annuum*) and barley (*Hordeum vulgare*) plants, respectively. The plants were grown from seeds in pots with commercial substrate and regularly infected for aphid feeding (bell pepper plants were infected at a 4 leaves stage and barley plants when they reached approximately a length of 10 cm). Both the insect colonies and their host plants were maintained in a growth chamber at 22 ± 1 °C, > 70% relative humidity with a 16:8 h light photoperiod.

Bioassays were conducted with 1.0 cm² leaf disks/fragments of *C. annuum* (*M. persicae*, *S. littoralis*) or *H. vulgare* (*R. padi*) as described previously [51]. The tests (10 µL of the solution) were applied at initial doses of 10 or 5 mg/ml (extract or compound) to the upper surface of the leaf fragments. Two sixth-instar larvae (>24 h after moulting) of *S. littoralis* were placed in 6 Petri dishes (9 cm in diameter) with 2 leaf disks (treatment and control) and allowed to feed at room temperature until 75% larval consumption of the paired control or treatment disks. The leaf disk surface consumption was measured using ImageJ (<http://imagej.nih.gov/ij/>, accessed on March 18, 2024) [49]. In the case of aphids, twenty (2x2 cm) ventilated plastic boxes containing 10 apterous aphid adults (24-48 h old) were used. Aphids were allowed to feed in a growth chamber under the described environmental

conditions for 24 hours. Settling was quantified by counting the number of aphids settled on each leaf fragment. All experiments were repeated twice (SE<10%).

The feeding or settling inhibition (%FI or %SI) was calculated as $[1 - (T/C) \times 100]$, where T and C represent treated and control leaf fragments, respectively. The effects (%SI / %FI) were analyzed by the non-parametric Wilcoxon Signed-Rank Test. Extracts and compounds with an effect $\geq 70\%$ were tested in dose-response experiments (3-5 serial dilutions) to calculate their EC₅₀ (the effective dose causing a 50% settling / feeding reduction) with linear regression models (%FI/SI on Log-dose).

3.8. Nematicidal Bioassay

The *Meloidogyne javanica* population was maintained on tomato plants (*Solanum lycopersicum* var. Marmande) cultivated in pot cultures and kept in environmentally controlled growth chambers (at 25 ± 1 °C, >70% relative humidity). Egg masses of *M. javanica* were handpicked from infected tomato roots two months after seedling inoculation. Second-stage juveniles (J2) were obtained by incubating egg masses in a water suspension at 25 °C for 24 h. The tests were carried out in 96-well plates (BD Falcon, San Jose, CA, USA) and the extract and compounds were dissolved in distilled water containing 5% of a DMSO-Tween solution (0.2% Tween 20 in DMSO) according to Andres et al. [52]. The initial concentrations tested were of 1 and 0.5 mg mL for extract and pure compound, respectively, and four replicates were used for each test. Water containing 5% of a DMSO-Tween solution (0.2% Tween 20 in DMSO) was used for the negative control. The mortality rates after 72 h of incubation are presented as percentage of dead J2 corrected according to Scheider-Orelli's formula. Serial dilutions were used to calculate the effective lethal doses (LD₅₀) of the active compound by Probit Analysis (STATGRAPHICS Centurion XVI, version 16.1.03)

3.9. Phytotoxic Bioassay

The phytotoxic test was conducted with *Lolium perenne* and *Lactuca sativa* seeds placed in 12-well microplates (40 seeds for test), as described [53]. The extract or compounds dissolved in EtOH (negative control) were tested at concentrations of 0.4 or 0.2 mg/ml (final concentration in the well) and diluted serially if needed. Juglone (Sigma) was used as positive control (0.1 mg/ml), resulting in 100% germination inhibition. Briefly, the test solution (20 µl) and 300 µl of H₂O were added to a 2.5 cm diameter filter paper into each well plate. The seeds (10/5 of *L. sativa*/*L. perenne* soaked in distilled water for 8 h) were placed in every well and the parafilm-sealed plates were incubated in a plant growth chamber (25°C, 70% RH, 16:8 L:D). Germination was monitored for 7 days and leaf length (for *L. perenne*) and root length (for both species) were measured at the end of the experiment on 25 randomly selected digitalized seedlings with the ImageJ application (<http://rsb.info.nih.gov/ij/>, accessed on May 20, 2024) [49].

4. Conclusions

In this study the endophytic fungus HYP6, isolated from the epiphytic lichen *Hypogymnia tubulosa*, was identified as *Xylaria* sp. belonging to the *X. arbuscula* complex. The EtOAc extract from the liquid culture was chromatographed to afford (+)-piliformic acid (**3**) as the main metabolite and two new hydroxylated derivatives, (+)-9-hydroxypiliformic acid (**1**) and (+)-8-hydroxypiliformic acid (**2**), along the known compounds, hexylaconitic acid A anhydride (**4**) and the hydroxyphenylacetic derivatives (**5-6**). The extract showed significant aphid antifeedant, antifungal and phytotoxic effects. The aphid antifeedant activity of the extract against *M. persicae* can be explained by compounds **4-6**, whereas its moderate activity against *R. padi* is probably due to its content of antifeedant compound **4**. Compound **4** also showed potent fungicide effect against *B. cinerea* and the nematode *M. javanica*. The phytotoxicity observed can be attributed to the compounds **3-5** affecting negatively the growth of *L. perenne* and *L. sativa*. In addition, compound **3** and **4** showed stimulating effect on the growth of *L. sativa* root at low doses. Endolichenic fungus *Xylaria* sp. represents an important source for the biotechnological production of biopesticide compounds for plant disease control. Further

optimization of the fermentation process is needed to increase the production of these active metabolites.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org. Data sequence of *Xylaria* sp.; Figures S1-S7: ¹H-NMR, ¹³C-NMR, COSY, gHSQC, HMBC, NOESY and HREIMS spectra for compound **1**; Figures S8-S14: ¹H-NMR, ¹³C-NMR, COSY, gHSQC, HMBC, NOESY and HREIMS spectra for compound **2**; Figures S15-S20: ¹H-NMR, ¹³C-NMR, COSY, gHSQC, HMBC and NOESY spectra for compound **4**.

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