Sesquiterpenes and other constituents of *Xylaria* sp. NC1214, a fungal endophyte of the moss *Hypnum* sp.

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ABSTRACT

Oxygenated guaiane-type sesquiterpenes, xylaguaianols A–D (**1–4**), an iso-cadinane-type sesquiterpene

1 isocadinanol A (**5**), and an α -pyrone 9-hydroxyxylarone (**6**), together with five known sesquiterpenes
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3 (**7–11**), and four known cytochalasins (**12–15**) were isolated from a culture broth of *Xylaria* sp. NC1214,
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5 a fungal endophyte of the moss *Hypnum* sp. The structures of all compounds were elucidated by the
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7 analysis of their spectroscopic data and relative configurations of **1–5** were determined with the help of
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9 NMR NOESY experiments. Cytochalasins C (**12**), D (**13**), and Q (**14**) were investigated for their cytotoxic
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11 activity against five tumor cell lines. Cytochalasin D showed significant cytotoxicity against all five cell
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13 lines, with IC₅₀s ranging from 0.22 to 1.44 μ M, whereas cytochalasins C and Q exhibited moderate, but
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15 selective cytotoxicity.
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23 **Keywords:** *Xylaria* sp., fungal endophyte, *Hypnum* sp., xylaguanols, sesquiterpenes, α -pyrone,
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1. Introduction

Endophytic fungi constitute rich sources of secondary metabolites with novel structures and biomedical potential (Schulz et al., 2002; Gunatilaka, 2006; Zhang et al., 2006). *Xylaria* species, often isolated as endophytes from decaying plant tissue, occur worldwide from arctic to tropical regions (Schuëffler et al., 2007). Many *Xylaria* species are known to produce diverse secondary metabolites (Song et al., 2014), which include succinic acid derivatives (Klaiklay et al., 2012), cytochalasins (Espada et al., 1997), α -pyrones (Schuëffler et al., 2007), terpenoids (Li et al., 2010; Wu et al., 2014; Yan et al., 2011), xanthones (Davis and Pierens, 2006; Healy et al., 2004), cyclopeptides (Lin et al., 2001; Wu et al., 2011), and lactones (Jimenez-Romero et al., 2008). In an ongoing search for bioactive natural products from fungal endophytes, a cytotoxic EtOAc extract of *Xylaria* sp. NC1214, a fungal endophyte of the moss *Hypnum* sp., cultured in potato dextrose broth (PDB) containing 0.25 mM CuSO₄ (Paranagama et al., 2007) was investigated. Fractionation of this extract resulted in the isolation of fifteen metabolites including four cytotoxic cytochalasins. Herein reported are the isolation and structure elucidation of four new oxygenated guaiane-type sesquiterpenes (**1–4**), a rare bicyclic sesquiterpene (**5**) and a new α -pyrone (**6**), as well as the known sesquiterpenes, *epi*-guaidiol A (**7**) (Xu et al., 2009), hydroheptelidic acid (**8**) (Calhoun et al., 1992), gliocladic acid (**9**) (Itoh et al., 1982), bullatantriol (**10**) (Sung et al., 1992), and $1\beta,4\beta,7\alpha$ -trihydroxyeudesmane (**11**) (Sung et al., 1992), and four known cytochalasins, cytochalasins C (**12**), D (**13**), Q (**14**), and R (**15**) (Edwards et al., 1989). Also reported herein is the cytotoxic activity of cytochalasins **12–14** against five sentinel cancer cell lines.

2. Results and discussion

Xylaguaianols A (**1**) and B (**2**) (Figure 1) were determined to have the same molecular formula, C₁₅H₂₈O₄, on the basis of their HRMS and ¹³C NMR spectroscopic data, indicating that they are sesquiterpenes. Comparison of their ¹H and ¹³C NMR spectroscopic data (Tables 1 and 2, respectively) with those reported for bicyclic sesquiterpenes (Wu et al., 2014) suggested that **1** and **2** are tetrahydroxy derivatives of guaiane-type sesquiterpenes. The ¹³C and DEPT NMR spectra of xylaguaianol A (**1**) (Table

2) indicated fifteen carbons consisting of three primary, six secondary (of which one was oxygenated; δ_C 68.8), three tertiary, and three oxygenated quaternary carbons (δ_C 75.3, 77.6 and 81.9). The analysis of the 1H NMR and HSQC data (Table 1) established that **1** possessed three tertiary methyl groups (δ_H 1.08 s, δ_C 33.3; δ_H 1.18 s, δ_C 24.5; δ_H 1.25 s, δ_C 26.6), six methylene groups of which one is oxygenated (δ_H 3.28–3.33 m and 3.59–3.65 m, δ_C 68.8), and three methine groups (δ_H 1.95 m, δ_C 33.4; δ_H 2.17 m, δ_C 49.3; δ_H 2.86 m, δ_C 55.0). Detailed analysis of 1H – 1H COSY and HMBC spectra (Figure 2) confirmed that **1** contained a guaiane-type sesquiterpene skeleton bearing four hydroxyl groups. Careful inspection of the HMBC spectrum suggested the presence of long-range correlations of H-15 to C-4 (δ_C 81.9), H-14 to C-10 (δ_C 75.3), H-13 to C-11 (δ_C 77.6), and H-13 to C-12 (δ_C 68.8) confirming that these four hydroxyls were attached to C-4, C-10, C-11 and C-12. These data indicated that **1** had the same overall skeletal structure as that of 4 β -hydroxyxylaranol recently encountered in an endophytic fungus of a mangrove plant (Zeng et al., 2015). However, comparison of their $[\alpha]_D$ and NMR data suggested that they are not identical but may be stereoisomers. The relative configuration of **1** was assigned by the NOESY experiment combined with MM2 energy-minimized three-dimensional molecular modeling. The NOE correlations of H-1 with H-5/H-14, and H-5 with H-14/H-15 established that they are on the same face of the molecule (Figure 3). Additional NOEs were observed between H-6 α and 4-OH/H-12/H-13, indicating that the protons at C-6 α , C-12 and C-13 as well as the hydroxyl at C-4 were on the same side of the ring system. Thus, the structure of xylaguaianol A was established as pseudo-1 β ,5 β ,7 β (H)-guaiane-4 α ,10 α ,11,12-tetraol (**1**). The stereoconfiguration at C-11 of **1** was defined by comparison with the data for xylaguaianol B (**2**) as described below.

Analysis of the 1H (Table 1), ^{13}C (Table 2) and 2D NMR data of xylaguaianol B (**2**) suggested that it shared similar structural features with those of **1**. In addition, the NOESY data of **2** indicated that its bicyclic ring system had the same relative configuration as in **1**. Careful comparison of the ^{13}C NMR chemical shifts of **1** and **2** suggested that the major differences between these existed at C-7 (δ_C 33.4 for **1**, δ_C 32.2 for **2**), C-12 (δ_C 68.8 for **1**, δ_C 70.2 for **2**) and C-13 (δ_C 26.6 for **1**, δ_C 23.8 for **2**), indicating that **1**

1 and **2** were a pair of epimers of *R/S* configurations at C-11. Careful examination of the ^1H and ^{13}C NMR
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3 data recently reported for a series of guaiane-type sesquiterpenes epimeric at C-11, whose absolute
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5 configurations have been determined by X-ray crystallographic analysis (Wu et al., 2014), suggested that
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7 the ^{13}C NMR data may be used to define the absolute stereochemistry at C-11 for this type of
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9 sesquiterpenes. The reported data indicated that the γ -*gauche* interaction (Whitesell et al., 1987) between
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11 C-6/C-8 and 11-OH/C-12/C-13 resulted in the difference of chemical shifts for these carbons, especially
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13 for C-13. The *7R/11R* epimer always exhibited δ_{C} for C-13 at a higher field compared to that of the
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15 corresponding *7R/11S* epimer. It was found that the δ_{C} for C-13 of **2** (23.8) was at a higher field than that
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17 of **1** (26.6) and this allowed us to distinguish the *11R*^{*} and *11S*^{*} configurations for **1** and **2**, respectively, by
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19 assuming *7S*^{*}-configuration of **1** and **2**. Thus, xylaguaianol A and xylaguaianol B were identified as
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21 *1\beta,5\beta,7\beta*(H)-guaiane-*4\alpha,10\alpha,11R*^{*},12-tetraol (**1**) and *1\beta,5\beta,7\beta*(H)-guaiane-*4\alpha,10\alpha,11S*^{*},12-tetraol (**2**),
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23 respectively.
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30 Xylaguaianol C (**3**), obtained as a colorless oil, was determined to have the molecular formula
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32 $\text{C}_{15}\text{H}_{28}\text{O}_3$ by HRESIMS and NMR data, suggesting two degrees of unsaturation. The ^{13}C NMR data (Table
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34 2) established the presence of fifteen sp^3 carbons of which three were oxygenated (δ_{C} 69.2, 74.0, 75.9).
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36 Further analysis of the ^1H (Table 1), ^{13}C (Table 2), and ^1H - ^1H COSY NMR data together with its
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38 molecular formula suggested the presence of a trihydroxylated guaiane-type sesquiterpene skeleton in **3**.
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40 The HMBC correlations of H-13 to C-11 and C-12, and H-14 to C-10 confirmed that three of these
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42 hydroxyls were located at C-10, C-11 and C-12. The NOESY correlations of H-1 with H-14/H-5 and H-5
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44 with H-15 indicated that these protons were co-facial. Additional correlations of H-*8\alpha* with H-12/H-13
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46 were also observed, suggesting that these protons were on the same side of the guaiane-ring. However,
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48 with the data available it was not possible to define the configuration at C-11 of **3**, and because attempted
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50 crystallization failed, X-ray crystallography could not be applied for this purpose. Thus, the structure
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52 xylaguaianol C was determined as *1\beta,4\alpha,5\beta,7\beta*(H)-guaiane-*10\alpha,11,12*-triol (**3**).
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59 Xylaguaianol D (**4**), obtained as a colorless oil, was assigned the molecular formula $\text{C}_{15}\text{H}_{26}\text{O}_2$ by a
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combination of HRESIMS and NMR data, suggesting three degrees of unsaturation. The ^{13}C NMR data of **4** (Table 2) demonstrated the presence of fifteen signals consisting of thirteen sp^3 carbons including two oxygenated (δ_{C} 69.0 and 75.2) and two quaternary sp^2 carbons (δ_{C} 129.7 and 142.0). Comparison of the ^1H and ^{13}C NMR data of **4** (Tables 1 and 2, respectively) with those of **1–3** combined with its ^1H – ^1H COSY data established the presence of a double bond bearing guaiane-type sesquiterpene ring system in **4**. This double bond was located between C-1 and C-10 based on strong HMBC correlations observed for methyl protons at δ 1.63 (H-14) and two olefinic carbons at δ_{C} 129.7 (C-1) and 142.0 (C-10) (Figure 2). The NOE correlations of H-4 with H-5 and H-6 α with H-15/H-12 suggested that these protons are on the same side of the molecule. Thus, xylaguaianol D was identified as 4 β ,5 β ,7 β (H)-1,10-dehydroxyguaiane-11,12-diol (**4**).

Isocadinanol A (**5**), obtained as a colorless oil, was determined to have the molecular formula $\text{C}_{15}\text{H}_{26}\text{O}_3$ by its HRESIMS and NMR data, suggesting three degrees of unsaturation. The ^{13}C NMR spectrum of **5** revealed the presence of fifteen carbons suggesting that it was a sesquiterpene. Analysis of ^1H and HSQC NMR data indicated that **5** possessed a tertiary methyl (δ_{H} 1.31 s, δ_{C} 27.8), three secondary methyls [δ_{H} 0.86 d ($J = 6.4$ Hz), δ_{C} 21.0; δ_{H} 0.97 d ($J = 6.8$ Hz), δ_{C} 21.2; δ_{H} 1.05 d ($J = 6.4$ Hz), δ_{C} 14.9], three methylenes [δ_{H} 1.08 m and 1.54 m, δ_{C} 19.7; δ_{H} 1.13 m and 1.67 m, δ_{C} 37.2; δ_{H} 1.66 m and 2.12 ddd ($J = 13.2, 4.4, 2.4$ Hz), δ_{C} 34.8], and six methines of which two were oxygenated [δ_{H} 2.92 dd ($J = 16.0, 1.6$ Hz), δ_{C} 68.3; δ_{H} 3.43 m, δ_{C} 71.3]. These data also indicated that **5** contained two oxygenated quaternary carbons of an oxirane moiety (δ_{C} 65.9 and 66.4). The presence of the fragments –CH(CH₃)CH(O)CH₂CHCH(CH₃)₂ and –CH(O)CH₂CH₂CH– in **5** was deduced by the analysis of its ^1H – ^1H COSY data and the connectivity between these two fragments and with the two epoxy quaternary carbons and a tertiary methyl was established by the analysis of its HMBC data (Figure 2). These data suggested that **5** contained an isocadinane ring derived from a cadinane-like ring system in which the methyl group at C-4 has undergone rearrangement to C-5. The relative configuration of **5** was determined based on its NOESY spectrum combined with MM2 energy-minimized three-dimensional molecular modeling. The

NOE correlation of H-1 with H-10 confirmed that these had the same relative configuration. Additional NOEs observed between H-9/H-14, H-14/H-4, H-4/H-15, and H-15/H-7 (Figure 3) indicated that H-1, hydroxyls at C-4 and C-9, the C-5(6) epoxy group and the isopropyl group at C-7 were all on the same side of the ring system. Thus, the structure of isocadinanol A was established as 1 α ,7 β ,10 α (H)-5 α (6 α)-epoxy-isocadinane-4 α ,9 α -diol (**5**).

9-Hydroxyxylarone (**6**), obtained as a white amorphous solid, was assigned the molecular formula C₁₃H₁₈O₄ by a combination of HRESIMS and NMR data, indicating five degrees of unsaturation. Inspection of its ¹H, ¹³C and HMBC NMR data suggested the presence of thirteen carbons consisting of four methyl carbons of which one was oxygenated, one methylene carbon, three methine carbons of which two were aromatic and/or olefinic and five sp² quaternary carbons. Analysis of ¹H NMR and HSQC data demonstrated the presence of three tertiary methyls of which one was oxygenated (δ_{H} 1.93 s, δ_{C} 8.7; δ_{H} 1.94 s, δ_{C} 13.0; δ_{H} 3.90 s, δ_{C} 56.2), one secondary methyl [δ_{H} 0.92 t (J = 7.2 Hz), δ_{C} 9.6], one methylene (δ_{H} 1.55–1.75 m, δ_{C} 30.4) and three methines [δ_{H} 4.43 dt (J = 8.6, 6.6 Hz), δ_{C} 70.0; δ_{H} 6.18 s, δ_{C} 92.9; δ_{H} 6.49 dd (J = 8.8, 1.2 Hz), δ_{C} 135.8]. Further analysis of the HMBC spectrum (Figure 2) indicated the existence of an α -pyrone system in **6** and comparison of the ¹H and ¹³C NMR data of **6** with the known α -pyrone, xylarone (Schüffler et al., 2007), established that they were structurally related, except for the presence of an oxymethine moiety (δ_{H} 4.43 m, δ_{C} 70.0) in **6** instead of the methylene moiety in xylarone. These findings together with HRMS data suggested that **6** was a hydroxylated derivative of xylarone. The HMBC correlations of H-10 and H-11 to C-9 and H-9 to C-7 and C-8 suggested that this hydroxyl group was at C-9. However, lack of sufficient material precluded us from performing additional studies to determine its configuration. Thus, the structure of this metabolite was elucidated as 9-hydroxyxylarone (**6**).

Metabolites **7–15** were identified as *epi*-guaidiol A (**7**) (Xu et al., 2009), hydroheptelidic acid (**8**) (Calhoun et al., 1992), gliocladic acid (**9**) (Itoh et al., 1982), bullatantriol (**10**) (Sung et al., 1992), 1 β ,4 β ,7 α -trihydroxyeudesmane (**11**) (Sung et al., 1992), cytochalasin C (**12**) (Edwards et al., 1989), cytochalasin D (**13**) (Edwards et al., 1989), cytochalasin Q (**14**) (Edwards et al., 1989), and cytochalasin R

(15) (Edwards et al., 1989), respectively, by comparison of their MS and NMR spectroscopic data with those reported. Since many cytochalasins have been reported to be cytotoxic (Van Goietsenoven et al., 2011; Xu et al., 2015), cytochalasins C (12), D (13) and Q (14) encountered in this study were tested against a panel of five human tumor cell lines including prostate adenocarcinoma (PC-3M), non-small cell lung cancer (NCI-H460), CNS glioma (SF-268), breast adenocarcinoma (MCF-7), and metastatic breast adenocarcinoma (MDA-MB-231). As depicted in Table 3, cytochalasin D (13) exhibited strong cytotoxicity against all five cell lines with moderate selectivity to NCI-H460. Cytochalasins C (12) and Q (14) showed moderate cytotoxicity against all tumor cell lines except MCF-7.

3. Experimental

3.1. General experimental procedures

1D and 2D NMR spectra were recorded on a Bruker Avance III 400 NMR instrument at 400 MHz for ^1H NMR and 100 MHz for ^{13}C NMR. Chemical shift values (δ) are given in parts per million (ppm), and the coupling constants are in Hz. Low-resolution and high-resolution MS were recorded on Shimadzu LCMS-DQ8000 α and JEOL HX110A spectrometers, respectively. Optical rotations were measured at 25 $^\circ\text{C}$ with a JASCO Dip-370 digital polarimeter using MeOH or acetone as solvent. UV spectra were recorded in MeOH using a Shimadzu UV-1601 UV-Vis spectrometer. Column chromatography (CC) was performed using Baker silica gel 40 μm flash chromatography packing (J. T. Baker) or Sephadex LH-20 (25–100 μm ; GE Healthcare). Analytical and preparative thin-layer chromatography (TLC) were performed on pre-coated 0.20 mm thick plates of silica gel 60 F254 (Merck). HPLC purifications were carried out on a 10 $\times$ 250 mm Phenomenex Luna 5 μm C18 column with a Waters Delta Prep system consisting of a PDA 996 detector. MM2 energy minimizations of possible conformations of compounds were performed using CambridgeSoft ChemBio3D Ultra.

3.2. Fungal material

Endophyte NC1214 was isolated on 2% malt extract agar from surface-sterilized, photosynthetic tissue of a freshly collected sample of *Hypnum* sp. (Hypnaceae; *Hypnum* moss) obtained from a granite

1 bald in mixed deciduous forest in the southern Appalachian Mountains near Highlands, North Carolina
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3 (for isolation details, see U'Ren et al., 2012a). The strain was accessioned as a living mycelial voucher at
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5 the Robert L. Gilbertson Mycological Herbarium (MYCO-ARIZ, NC1214). Total genomic DNA was
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7 isolated from fresh mycelium and the nuclear ribosomal internal transcribed spacers and 5.8s gene (ITS
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9 rDNA; ca. 600 base pairs [bp]) and the adjacent portion of the nuclear ribosomal large subunit (LSU
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11 rDNA) was amplified as single fragment by PCR (U'Ren et al., 2012b). The product (1147 bp) was
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13 visualized on a 1% agarose gel, cleaned, normalized, and sequenced as described previously (U'Ren et al.,
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15 2012b). Sequencing reads were assembled, bases were called, and quality scores were assigned by *phred*
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17 (Ewing and Green, 1998) and *phrap* (Ewing et al., 1998) through Mesquite (Maddison and Maddison,
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19 2011), followed by manual editing in Sequencher (Gene Codes Corp.). The resulting bidirectional
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21 sequence has been deposited in GenBank (accession JQ761854.1). Phylogenetic analyses of >2400
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23 endophytic fungi and related strains from lichens for a related study (Chen et al., 2015), coupled with rich
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25 taxon sampling from GenBank and focusing only on the conserved 5.8S and LSUrDNA sequences, placed
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27 the isolate within Xylariaceae (Xylariales, Sordariomycetes, Pezizomycotina, Ascomycota). Multilocus
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29 analyses with sampling across the Xylariaceae (U'Ren et al., 2012) indicated that NC1214 was part of a
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31 clade consisting of endophytes and endolichenic fungi that is sister to the clade containing *Xylaria*
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33 *coccophora*. Pending description of this novel clade, we refer to this strain conservatively as *Xylaria* sp.
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44 3.3. Fermentation, extraction and isolation

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46 A seed culture of *Xylaria* sp. NC1214 grown on PDA for 2 weeks was used for inoculation. Mycelia
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48 were scraped out and vortexed with sterile PDB (90 mL) and filtered through a 100 µm sterile filter to
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50 separate spores and/or fragments of hyphae from the mycelia. Absorbance of the filtrate was measured (at
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52 600 nm) and adjusted to 0.6. This filtrate was used to inoculate 10 × 2.0 L Erlenmeyer flasks, each holding
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54 1.0 L of the medium (PDB) containing 0.25 mM aq. CuSO₄ and incubated at 160 rpm and 28 °C. After 14
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56 days, mycelia were separated by filtration, and the filtrate was extracted with EtOAc (3 × 5 L).
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Evaporation of the EtOAc extract under reduced pressure afforded the crude extract (3.24 g). A portion (3.04 g) of this extract was partitioned between MeOH:H₂O (80:20) and hexanes and the resulting MeOH:H₂O (80:20) fraction was diluted with H₂O to MeOH:H₂O (50:50) and extracted (x 3) with CHCl₃. The CHCl₃ extracts were combined and evaporated under reduced pressure to obtain the CHCl₃ fraction (2.78 g). The MeOH:H₂O (80:20) fraction after extraction with CHCl₃ was evaporated to afford the aq. MeOH fraction (0.20 g). A portion of the above CHCl₃ fraction (2.73 g) was subjected to Sephadex LH-20 (100 g) gel-permeation chromatography and eluted sequentially with 350 mL each of hexanes:CH₂Cl₂ (4:1), CH₂Cl₂:acetone (4:1), CH₂Cl₂:acetone (2:3), and MeOH, respectively. Evaporation of these yielded fractions A (0.60 g), B (1.97 g), C (0.11 g), and D (0.02 g), respectively. Fraction A (0.59 g) was subjected to chromatography over a column of RP C-18 and eluted with a gradient of MeOH:H₂O (60:40, 70:30, 80:20, 90:10, 100:0; each 120 mL) to afford five fractions A1–A5, respectively. Fraction A1 (135.4 mg) was subjected to silica gel chromatography and eluted with CHCl₃:MeOH (from 98:2 to 90:10) to give five fractions A1–1 to A1–5. The major fraction A1–3 (50.3 mg) was further fractionated by C-18 (20 g) CC, eluted with a gradient initially of MeOH:H₂O (50:50) to MeOH, followed by HPLC purification [C-18; MeOH:H₂O (55:45)] to afford **6** (0.5 mg; *R*_f 0.44, CHCl₃–MeOH, 14:1) and **12** (2.4 mg; *R*_f 0.69, CHCl₃–MeOH, 14:1). Fraction A2 (386.8 mg) was further fractionated by silica gel CC using CHCl₃–MeOH (from 98:2 to 90:10) as the eluent to give 15 sub-fractions A2–1 to A2–15. Evaporation of fractions A2–8, A2–11, and A2–12 under reduced pressure yielded **7** (0.6 mg, *R*_f 0.26, CHCl₃–MeOH, 14:1), **1** (0.8 mg; *R*_f 0.34, CHCl₃–MeOH, 10:1) and **3** (1.8 mg; *R*_f 0.30, CHCl₃–MeOH, 10:1), respectively. Further fractionation of A2–1 (78.2 mg) by silica gel (12 g) CC and elution with CHCl₃–MeOH (from 99:1 to 95:5) followed by prep-HPLC purification [C-18; MeOH:H₂O (55:45)] afforded **14** (25.3 mg, *R*_t 20.0 min) and **15** (0.7 mg, *R*_t 38.1 min). Purification of fraction A2–7 (21.4 mg) by prep-HPLC [C-18; MeOH:H₂O (60:40)] afforded **5** (2.0 mg; *R*_f 0.26, CHCl₃–MeOH, 14:1). Further fractionation of fraction B (1.97 g) by a RP C-18 CC and elution with a gradient of MeOH:H₂O (50:50, 60:40, 70:30, 80:20, 90:10, 100:0; each 350 mL) afforded six fractions B1–B6, respectively. Fraction B1 (279.4 mg) was further

separated using silica gel CC and eluted with CHCl₃–MeOH (from 98:2 to 85:15) followed by purification RP C-18 CC [MeOH:H₂O (50:50 to 70:30)] to give **11** (2.4 mg; *R_f* 0.32, CH₂Cl₂–acetone, 2:1). Fraction B2 (762.2 mg) was separated by silica gel CC and eluted with CHCl₃–MeOH (from 98:2 to 85:15) to give 13 sub-fractions, B2–1 to B2–13. Fraction B2–5 (45.6 mg) was separated by C-18 (10 g) CC and eluted with MeOH:H₂O (50:50 to 70:30) to afford **2** (2.0 mg; *R_f* 0.40, CH₂Cl₂–acetone, 5:2). B2–9 (32.0 mg) was further purified over a column (10 g) of RP C-18 and eluted with MeOH:H₂O (50:50 to 70:30) to afford **10** (2.3 mg; *R_f* 0.30, CH₂Cl₂–acetone, 2:1). Fraction B3 (494.6 mg) was separated by silica gel (80 g) CC and eluted with CHCl₃–MeOH (from 98:2 to 85:15) to give 18 sub-fractions, B3–1 to B3–18. Evaporation of B3–4 under reduced pressure afforded **13** (137.1 mg; *R_f* 0.47, CHCl₃–MeOH, 5:1). Fraction B3–13 (23.5 mg) was further separated by RP C-18 CC and eluted with MeOH:H₂O (70:30 to 80:20) to give **4** (1.6 mg; *R_f* 0.23, CH₂Cl₂–acetone, 1:1). In order to isolate the polar metabolites of this fungal strain, the MeOH:H₂O (50:50) fraction and the fractions C–D obtained from the CHCl₃ fraction resulting from solvent–solvent partitioning of the original EtOAc extract which had similar TLC profiles were combined (0.33 g) and separated by C18 RP prep-HPLC with solvent gradients of 10% to 58% aqueous MeCN (0–30 min), 58% aqueous MeCN to MeCN (30–35 min), and 100% MeCN (35–40 min), yielding **8** (148.0 mg; *R_t* 17.8 min) and **9** (43.2 mg; *R_t* 23.6 min).

3.4. Spectroscopic data of metabolites

3.4.1. Xylaguaianol A (**1**)

Colorless oil; [α]_D²⁵ +30.2 (*c* 0.08, MeOH); for ¹³C NMR and ¹H NMR spectroscopic data, see Tables 1 and 2; APCIMS *m/z* 277 [M–H₂O+Na]⁺ (25), 237 [M–2H₂O+H]⁺ (20), 219 [M–3H₂O+H]⁺ (100); positive HRESIMS *m/z* 255.1956 [M–H₂O+H]⁺ (calcd for C₁₅H₂₇O₃, 255.1960).

3.4.2. Xylaguaianol B (**2**)

Colorless oil; [α]_D²⁵ +1.2 (*c* 0.12, MeOH); for ¹³C NMR and ¹H NMR spectroscopic data, see Tables 1 and 2; positive APCIMS *m/z* 277 [M–H₂O+Na]⁺ (20), 237 [M–2H₂O+H]⁺ (34), 219 [M–3H₂O+H]⁺

(100), 201 $[M-4H_2O+H]^+$ (55); positive HRESIMS m/z 255.1954 $[M-H_2O+H]^+$ (calcd for $C_{15}H_{27}O_3$, 255.1960).

3.4.3. Xylaguaianol C (3)

Colorless oil; $[\alpha]_D^{25} +22.2$ (c 0.16, MeOH); for ^{13}C NMR and 1H NMR spectroscopic data, see Tables 1 and 2; positive APCIMS m/z 257 $[M+H]^+$ (10), 221 $[M-2H_2O+H]^+$ (85), 207 $[M-H_2O-CH_2OH]^+$ (100), 203 $[M-3H_2O+H]^+$ (68); positive HRESIMS m/z 279.1924 $[M+Na]^+$ (calcd for $C_{15}H_{28}O_3Na$, 279.1931).

3.4.4. Xylaguaianol D (4)

Colorless oil; $[\alpha]_D^{25} +14.7$ (c 0.09, MeOH); UV (MeOH) λ_{max} (log ϵ) 212 (3.7) nm; for ^{13}C NMR and 1H NMR spectroscopic data, see Tables 1 and 2; positive APCIMS m/z 233 $[M-C_2H_4+Na]^+$ (100); positive HRESIMS m/z 261.1876 $[M+Na]^+$ (calcd for $C_{15}H_{26}O_2Na$, 261.1831).

3.4.5. Isocadinanol A (5)

Colorless oil; $[\alpha]_D^{25} +10.1$ (c 0.10, MeOH); 1H NMR (400 MHz, $CDCl_3$) δ 3.43 (1H, m, H-9), 2.92 (1H, dd, $J = 16.0, 1.6$ Hz, H-4), 2.12 (1H, ddd, $J = 13.2, 4.4, 2.4$ Hz, H-8 α), 1.74 (1H, m, H-11), 1.67 (1H, m, H-3 α), 1.66 (1H, m, H-8 β), 1.54 (1H, m, H-2 α), 1.44 (1H, m, H-1), 1.34 (1H, m, H-10), 1.31 (3H, s, H-15), 1.13 (1H, m, H-3 β), 1.08 (1H, m, H-2 β), 1.05 (3H, d, $J = 6.4$ Hz, H-14), 0.97 (3H, d, $J = 6.8$ Hz, H-13), 0.91 (1H, m, H-7), 0.86 (3H, d, $J = 6.4$ Hz, H-12); ^{13}C NMR (100 MHz, $CDCl_3$) δ 71.3 (CH, C-9), 68.3 (CH, C-4), 66.4 (C, C-6), 65.9 (C, C-5), 49.5 (CH, C-7), 43.9 (CH, C-10), 37.7 (CH, C-1), 37.2 (CH₂, C-3), 34.8 (CH₂, C-8), 27.8 (CH₃, C-15), 27.5 (CH, C-11), 21.2 (CH₃, C-13), 21.0 (CH₃, C-12), 19.7 (CH₂, C-2), 14.9 (CH₃, C-14); positive APCIMS m/z 201 $[M-3H_2O+H]^+$ (38), 219 $[M-2H_2O+H]^+$ (100), 237 $[M-H_2O+H]^+$ (70); HRESIMS m/z 277.1773 $[M+Na]^+$ (calcd for $C_{15}H_{28}O_3Na$, 277.1774).

3.4.6. 9-Hydroxyxylarone (6)

White amorphous solid; $[\alpha]_D^{25} +8.5$ (c 0.06, MeOH); UV (MeOH) λ_{max} (log ϵ) 309.5 (1.8), 225.5 (3.9) nm; 1H NMR (400 MHz, $CDCl_3$) δ 6.49 (1H, dd, $J = 8.8, 1.2$ Hz, H-8), 6.18 (1H, s, H-5), 4.43 (1H, dt, $J = 8.6, 6.6$ Hz, H-9), 3.90 (3H, s, OCH₃-4), 1.94 (1H, s, H-13), 1.93 (3H, s, H-12), 1.55–1.75 (2H, m,

H-10), 0.92 (3H, t, $J = 7.2$ Hz, H-11); ^{13}C NMR (100 MHz, CDCl_3) δ 164.1 (C, C-4), 163.8 (C, C-2), 159.1 (C, C-6), 135.8 (CH, C-8), 127.7 (C, C-7), 102.8 (C, C-3), 92.9 (CH, C-5), 70.0 (CH, C-9), 56.2 (CH₃, OCH₃-4), 30.4 (CH₂, C-10), 13.0 (CH₃, C-13), 9.6 (CH₃, C-11), 8.7 (CH₃, C-12); positive APCIMS m/z 239 $[\text{M}+\text{H}]^+$ (100); positive HRESIMS m/z 239.1271 $[\text{M}+\text{H}]^+$ (calcd for $\text{C}_{13}\text{H}_{19}\text{O}_4$, 239.1278).

3.5. Cytotoxicity Assay

The resazurin-based colorimetric (AlamarBlue) assay was employed for evaluation of cytotoxicity *in vitro* of samples against human prostate adenocarcinoma (PC-3M), human non-small cell lung (NCI-H460), human CNS glioma (SF-268), human breast (MCF-7), and human metastatic breast adenocarcinoma (MDA-MB-231) as described previously (Wijeratne et al., 2012; Wijeratne et al., 2014). Doxorubicin and DMSO were used as positive and negative controls, respectively.

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FIGURES

Fig. 1. Structures of metabolites 1–15.

Fig. 2. Key HMBC and ^1H - ^1H COSY correlations for compounds **1**, **4**, **5**, and **6**.

Fig. 3. Key NOE correlations for compounds **1** and **5**.

Table 1.¹H NMR (400 MHz) spectroscopic data for compounds **1–4** (acetone-*d*₆, δ in ppm, *J* in Hz).

position	1	2	3	4
1	2.86 m	2.90 m	2.05 m	-
2 α	1.50 m	1.53 m	1.50 m	2.31 m
2 β	1.71 m	1.70 m	1.64 m	2.16 m
3 α	1.63 m	1.67 m	1.66 m	1.35 m
3 β	1.91 m	1.93 m	1.21 m	1.64 m
4	-	-	1.97 m	2.08 m
5	2.17 m	2.24 m	1.88 m	2.38 m
6 α	2.17 m	2.13 m	1.77 m	0.81 m
6 β	1.12 m	1.20 m	0.90 m	1.76 m
7	1.95 m	1.99 m	1.89 m	1.67 m
8 α	2.12 m	2.08 m	1.82 m	1.13 m
8 β	1.45 m	1.44 m	1.14 m	2.05 m
9 α	1.85 m	1.70 m	1.88 m	2.08 m
9 β	1.72 m	1.53 m	1.37 m	2.08 m
12a	3.28–3.33 m	3.17–3.22 m	3.29–3.36 m	3.35–3.40 m
12b	3.59–3.65 m	3.41–3.50 m	3.40–3.50 m	3.47–3.52 m
13	1.25 s	1.23 s	0.98 s	1.00 s
14	1.08 s	1.07 s	1.10 s	1.63 d (1.2)
15	1.18 s	1.20 s	0.93 d (6.8)	0.88 d (7.2)
12-OH	3.50–3.56 m			3.54–3.60 m
13-OH				3.12 s
14-OH			3.03 s	
15-OH	3.18 s	3.19 s

Table 2.¹³C NMR (100 MHz) spectroscopic data for compounds **1–4** (acetone-*d*₆, δ in ppm).

position	1	2	3	4
1	55.0, CH	54.9, CH	56.6, CH	142.0, C
2	26.2, CH ₂	26.3, CH ₂	26.8, CH ₂	30.9, CH ₂
3	38.9, CH ₂	39.0, CH ₂	31.7, CH ₂	33.7, CH ₂
4	81.9, C	81.8, C	40.0, CH	39.9, CH
5	49.3, CH	49.3, CH	47.9, CH	47.3, CH
6	36.0, CH ₂	36.3, CH ₂	22.3, CH ₂	29.6, CH ₂
7	33.4, CH	32.2, CH	44.9, CH	50.3, CH
8	25.9, CH ₂	25.8, CH ₂	25.0, CH ₂	27.6, CH ₂
9	29.7 ^a , CH ₂	28.9 ^a , CH ₂	34.3, CH ₂	35.8, CH ₂
10	75.3, C	75.2, C	74.0, C	129.7, C
11	77.6, C	77.6, C	75.9, C	75.2, C
12	68.8, CH ₂	70.2, CH ₂	69.2, CH ₂	69.0, CH ₂
13	26.6, CH ₃	23.8, CH ₃	19.3, CH ₃	21.0, CH ₃
14	33.3, CH ₃	33.2, CH ₃	31.9, CH ₃	22.5, CH ₃
15	24.5, CH ₃	24.6, CH ₃	16.8, CH ₃	15.6, CH ₃

^aOverlapping with signals due to the solvent.

Table 3.

Cytotoxicity data for cytochalasins C (**12**), D (**13**) and Q (**14**).^a

Compound	Tumor Cell Line ^b				
	PC-3M	NCI-H460	SF-268	MCF-7	MDA-MB-231
12	1.65 ± 0.33	1.06 ± 0.03	0.96 ± 0.08	> 5	1.72 ± 0.27
13	1.03 ± 0.08	0.22 ± 0.02	0.43 ± 0.06	1.44 ± 0.26	1.01 ± 0.08
14	1.53 ± 0.12	1.51 ± 0.16	1.31 ± 0.08	> 5	1.32 ± 0.13
doxorubicin	0.25 ± 0.02	0.05 ± 0.01	0.45 ± 0.07	0.32 ± 0.09	0.67 ± 0.11

^aResults are expressed as IC₅₀ value ± standard deviation in μM. Doxorubicin and DMSO were used as positive and negative controls.

^bCell lines used: PC-3M = metastatic human prostate adenocarcinoma; NCI-H460 = human non-small cell lung cancer; SF-268 = human CNS glioma; MCF-7 = human breast adenocarcinoma; MDA-MB-231 = metastatic human breast adenocarcinoma.

Xylaguaianols A–D, isocadinanol A, and 9-hydroxyxylarone, together with several known sesquiterpenes and cytochalasins were encountered in *Xylaria* sp. NC1214, a fungal endophyte of the moss *Hypnum* sp. Cytochalasin D showed significant cytotoxicity against five cancer cell lines. Cytochalasins C and Q exhibited moderate, but selective cytotoxicity.