

# Highly oxygenated isoprenylated cyclohexanoids from the fungus Parastagonospora nodorum SN15

Amr El-Demerdash, Grégory Genta-Jouve, Margot Bärenstrauch, Caroline Kunz, Emmanuel Baudouin, Soizic Prado

## ▶ To cite this version:

Amr El-Demerdash, Grégory Genta-Jouve, Margot Bärenstrauch, Caroline Kunz, Emmanuel Baudouin, et al.. Highly oxygenated isoprenylated cyclohexanoids from the fungus Parastagonospora nodorum SN15. Phytochemistry, 2019, 166, pp.112056 -. 10.1016/j.phytochem.2019.112056 . hal-03487373

## HAL Id: hal-03487373 https://hal.science/hal-03487373

Submitted on 20 Dec 2021

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial 4.0 International License

## Highly oxygenated isoprenylated cyclohexanoids from the fungus

## Parastagonospora nodorum SN15

Amr El-Demerdash,<sup>1,2</sup> Grégory Genta-Jouve,<sup>1,3</sup> Margot Bärenstrauch,<sup>1</sup> Caroline Kunz,<sup>1,4</sup> Emmanuel

Baudouin,<sup>5</sup> Soizic Prado<sup>1\*</sup>

<sup>1</sup>Muséum National d'Histoire Naturelle, Unité Molécules de Communication et Adaptation des Microorganismes, UMR 7245, CP 54, 57 rue Cuvier, 75005 Paris, France

<sup>2</sup>Organic Chemistry Division, Chemistry Department, Faculty of Science, Mansoura University, Mansoura 35516, Egypt

<sup>3</sup>Université Paris Descartes, Laboratoire de Chimie-Toxicologie Analytique et Cellulaire (C-TAC), UMR CNRS 8638, COMETE, 4 Avenue de l'Observatoire 75006 Paris, France

<sup>4</sup>Sorbonne Université, Faculté des Sciences et Ingénierie, UFR 927, F-75005 Paris, France

<sup>5</sup>Sorbonne Université, CNRS, Laboratoire de Biologie du Développement, Institut de Biologie Paris Seine (LBD-IBPS), Paris F-75005, France **Abstract**: The chemical investigation of the wheat plant pathogen *Parastagonospora nodorum* SN15 led to the purification of seven highly oxygenated acetylenic cyclohexanoids named stagonosporynes A-H (1-7). Their structures were determined on the basis of extensive NMR and the relative and absolute configurations by an array of computational methods including simulation of NOESY spectrum and electronic circular dichroism (ECD). All compounds were evaluated for their herbicidal activity and stagonosporyne G (7) displayed the most significant herbicidal activity.

#### 1 1. Introduction

2 The Dothideomycetes represent a large and diverse class of fungi including many plant pathogens. Dothideomycetes are also prolific producers of specialised metabolites 3 (Chomcheon et al., 2006; Chomcheon et al., 2009; Hewage et al., 2014; Senadeera et al., 4 5 2012), which may be required for pathogenicity (Muria-Gonzalez et al., 2015; Stergiopoulos et al., 2013). Among members of Dothideomycete plant pathogens, Parastagonospora 6 nodorum (Teleomorph: Phaeosphaeria, syn. Septoria) is known as the causative agent of 7 Stagonospora (or Septoria) nodorum blotch leading to major losses in wheat crops in many 8 9 parts of the word (Figueroa et al., 2018; Solomon et al., 2006). Accordingly, P. nodorum has emerged as a Dothideomycete necrotrophic pathogen model and has been sequenced in 2007 10 11 (Hane et al., 2007). Its genome revealed the presence of 38 specialised metabolite biosynthetic gene clusters (BGC) including genes coding for polyketide synthases (PKS); 12 13 non-ribosomal peptide synthases (NRPS), terpene synthases and prenyl transferases 14 suggesting a strong potential of specialised metabolite production (Chooi et al., 2014; Ipcho et al., 2012). 15

16 Interestingly, the authors showed that the expression of several BGCs was upregulated 17 during *in planta* infection (Ipcho et al., 2012) such as *SNOG* 00477 and *SNOG-5791-5797* 

gene clusters. Deletion of SN477 gene combined with its heterologous expression in 18 19 Saccharomyces cerevisiae revealed that it encodes for a PR-PKS responsible of the biosynthesis of (R)-(-)-mellein, a well know dihydroisocoumarine compound previously 20 21 described as phytotoxic and widely widespread in fungi (Arakawa, 1968; Blair and Newbold, 1955; Chooi et al., 2015; Nishikawa, 1933). Recently heterologous expression of SNOG-22 23 5791-5797 in Aspergillus nidulans led to the isolation of new phytotoxic  $\alpha$ -pyrones (Li et al., 2018). Previous chemical investigations of *P. nodorum* led also to the identification of several 24 other metabolites (Devys et al., 1994; Devys et al., 1992; Tan et al., 2009; Yang et al., 2013) 25 but their number remains quite low compared to the huge biosynthetic potential of the strain. 26

Accordingly, and as a part of our ongoing research program on fungal natural products, we investigated the chemistry of the fungus *Parastagonospora nodorum* SN15 cultivated on rice medium. This led to the identification of seven previously undescribed highly oxygenated cyclohexanoids named stagonosporynes A-G (1-7) along with the known compound siccayne (8) (Figure 1). Herein we report their isolation and structural elucidation along with their herbicidal activity.

33

#### 35 2. Results and discussion

Molecular formula of compound 1 was established as  $C_{11}H_{16}O_3$  from its positive 36 HRESIMS *m/z* 219.0984 [M+Na]<sup>+</sup> (calc. for C<sub>11</sub>H<sub>16</sub>O<sub>3</sub>Na 219.0997), indicating four 37 unsaturation degrees. The IR spectrum exhibited characteristic absorption bands for hydroxyl 38 functionalities (3443 cm<sup>-1</sup>). The <sup>13</sup>C-J modulated NMR spectrum recorded in CDCl<sub>3</sub> revealed 39 the resonance of 11 carbons including one methyl at ( $\delta_C$  23.4, CH<sub>3</sub>-11), two oxymethines at 40  $(\delta_{\rm C}$  74.3, C-1;  $\delta_{\rm C}$  65.7, C-4), three methylenes ( $\delta_{\rm C}$  25.9, C-2;  $\delta_{\rm C}$  31.9, C-3;  $\delta_{\rm C}$  44.1, C-5) and 41 one olefinic carbon at ( $\delta_{\rm C}$  122.9, C-10). Three quaternary resonances including two acetylenic 42 carbons ( $\delta_c$  89.7, C-7 and  $\delta_c$  85.6, C-8) and one sp<sup>2</sup> carbon ( $\delta_c$  125.7, C-9) along with one 43 oxygenated sp<sup>3</sup> carbon at ( $\delta_C$  70.6, C-6) were also depicted (**Table 3**). Taking into account the 44 four degrees of unsaturation elucidated from the HRMS, compound 1 must be cyclic. The <sup>1</sup>H 45 NMR spectrum recorded in CDCl<sub>3</sub> (Table 1) revealed the presence of two oxymethine 46 protons ( $\delta_{\rm H}$  3.72, H-1;  $\delta_{\rm H}$  3.97, H-4) and three methylenes ( $\delta_{\rm H}$  1.83/1.68, CH<sub>2</sub>-2;  $\delta_{\rm H}$  1.95/1.35, 47 CH<sub>2</sub>-3;  $\delta_{\rm H}$  2.38/1.66, CH<sub>2</sub>-5) suggesting an oxygenated cyclohexanoid scaffold. Additional 48 signals for two olefinic protons CH<sub>2</sub>-10 observed at  $\delta_{\rm H}$  5.24/5.29 and a terminal methyl CH<sub>3</sub>-49 11 at  $\delta_{\rm H}$  1.86 suggested a methyl-terminal double bond function. <sup>1</sup>H-<sup>1</sup>H COSY spectrum 50 recorded in DMSO- $d_6$  of 1 revealed the presence of one main spin-coupling system showed in 51 bold blue lines (Figure 2), including the oxygenated methine CH-1 ( $\delta_{\rm H}$  3.72), the two 52 methylenes groups CH<sub>2</sub>-2 and CH<sub>2</sub>-3 ( $\delta_{\rm H}$  1.83/1.68,  $\delta_{\rm H}$  1.35/1.95), the oxymethine CH-4 ( $\delta_{\rm H}$ 53 3.97) and the methylene CH<sub>2</sub>-5 ( $\delta_{\rm H}$  1.66/2.38). The <sup>1</sup>H-<sup>1</sup>H COSY spectrum allowed us to 54 identify a second substructure including a terminal methyl ( $\delta_{\rm H}$  1.86, CH<sub>3</sub>-11) with the olefinic 55 protons ( $\delta_{\rm H}$  5.24 and 5.29, CH<sub>2</sub>-10). These two substructures were assembled together based 56 on the cross-peaks correlations observed in HMBC. Indeed, the oxymethine H-1 ( $\delta_{\rm H}$  3.72) and 57 the two diastereotopic protons CH<sub>2</sub>-5 ( $\delta_{\rm H}$  1.66/2.38) displayed key HMBC correlations 58 (Figure 2) with C-6 ( $\delta_C$  76.5) confirming that 1 is a cyclohexane-containing compound. 59 Additional key HMBC correlations (Figure 1) were observed between the olefinic protons 60 CH<sub>2</sub>-10, the terminal methyl CH<sub>3</sub>-11 and the acetylenic carbone C-8 ( $\delta_{\rm C}$  85.6). Furthermore, 61 both H-1 and CH<sub>2</sub>-5 displayed HMBC correlations with the acetylenic quaternary carbon C-7 62  $(\delta_{\rm C} 89.7)$  indicating the connection of the envne unit at C-6. This was further confirmed by 63 key HMBC correlations observed in DMSO- $d_6$  between OH-6 ( $\delta_H$  4.89) and the quaternary 64 carbon C-7 ( $\delta_{C}$  89.3). The relative configuration of 1 was deduced from the values of the 65

proton coupling constants and the NOESY correlations. NOEs observed in DMSO-d<sub>6</sub> and 66 CDCl<sub>3</sub> between H-4 and H-5eq/H-3eq and H-2ax suggest that they are in the same plan. 67 Additionally, NOEs between H-1 and H-3ax, H-5ax and H-2-eq suggest that there are on the 68 other side of the cyclohexane ring. Relative configuration at C-6 was deduced from NOEs 69 observed in DMSO-d6 between OH-6 and H-4, H-5eq (Figure 2). This was confirmed by 70 *trans*-diaxial coupling constants observed on <sup>1</sup>H-spectrum (CDCl<sub>3</sub>)  $J_{a/a}$  10.6 H-1/H-2ax and 71 72 10.1 H-4/H-3ax, which established that H-1 and H-4 were in axial position of a chair conformation for the cyclohexane ring (Figure 3). Therefore, compound 1 was assigned as 73 74 shown (Figure 1) and named stagonosporyne A. To determine the absolute configuration of 1, prediction of the ECD spectrum was undertaken and the experimental and simulated 75 spectra generated by time-dependent density functional theory (TDDFT) were compared. All 76 computational works were performed using Gaussian 16 (Frisch et al., 2016). The absolute 77 78 configuration of 1 was determined as 1*S*, 4*S*, 6*S* as the only Cotton effect (due to the  $\pi \rightarrow \pi^*$ ) 79 transition of the envne moiety) depicted on the spectra shared the same sign (Figure 4).

80 Molecular formula of compound 2 was established as  $C_{11}H_{16}O_4$  from its positive HR-ESI-MS *m/z* 235.09319 [M+Na]<sup>+</sup> (calcd. for C<sub>11</sub>H<sub>16</sub>O<sub>4</sub>Na 235.0946), indicating four 81 unsaturation degrees. The IR spectrum exhibited distinct absorption bands for hydroxyl 82 functionalities (broad band at 3369  $\text{cm}^{-1}$ ). The NMR spectral data of 2 recorded in both 83 DMSO-d<sub>6</sub> and CD<sub>3</sub>OD showed similar features of a highly oxygenated cyclohexanoid 84 compound. The obvious difference was the presence of an oxymethine in C-5 at  $\delta_{\rm H}$  2.95/ $\delta_{\rm C}$ 85 79.5 (DMSO-d<sub>6</sub>) instead of the methylene CH<sub>2</sub>-5 at  $\delta_{\rm H}$  2.06/1.38 and  $\delta_{\rm C}$  46.7 previously 86 described for stagonosporyne A (1). Compound 2 was thus named stagonosporyne B. The 87 relative configuration of 2 was deduced based on the proton coupling constant values and 88 NOESY correlations. The *trans*-diaxial coupling constants observed on <sup>1</sup>H-NMR spectrum 89 (CD<sub>3</sub>OD) (J<sub>a/a</sub> 9.4, 11.4, 11.8) of H-5/H-4, H-4/H-3ax and H-2ax/H-1 allowed us to assign H-90 1, H-5 and H-4 in axial form of a chair conformation for the cyclohexane ring. Moreover, 91 NOEs observed in DMSO-d<sub>6</sub> between H-1 and H-5/OH-6/H-3ax suggest that they are in the 92 same plan while NOEs between H-4 and H-2ax/OH-5 suggest that there are on the other side 93 94 of the six-membered ring (Figure 3). Accordingly, structure 2 was determined as shown in (Figure 1). The absolute configuration of 2 was determined as 1*R*, 4*R*, 5*S*, 6*S* by comparison 95 of the ECD spectrum with the one obtained for compound 1 (only one CE observed for this 96 family of compounds). 97

98 The molecular formula  $C_{13}H_{18}O_5$  was established for compound **3** according its 99 HRESIMS  $[M+K]^+$  *m/z* 293.0796 (Calc. for  $C_{13}H_{18}O_5K$  293.0791) and indicates five degrees

of unsaturation. Comparison of the <sup>1</sup>H NMR spectrum (Table 1) of 3 with this one of 2 100 indicated that the both compounds are closely related structures. Indeed, the only differences 101 102 were the presence of an extra methyl signal ( $\delta_{\rm H}$  2.05,  $\delta_{\rm C}$  21.1, 3H, CH<sub>3</sub>-13) along with the carbonyl C-12 ( $\delta_{\rm C}$  172.7), which was attributed to an acetyl functionality CH<sub>3</sub>CO and satisfied 103 104 the fifth unsaturation degree suggested by the HRMS. This was further confirmed through key HMBC correlations from both the oxymethine H-4 ( $\delta_H$  4.81, 1H) and CH<sub>3</sub>-13 ( $\delta_H$  2.05, 3H), 105 106 with the quaternary carbonyl ester at ( $\delta_C$  172.7, C-12) allowing thus to define the attachment 107 of the acetyl group at C-4 (Figure 2). The relative stereochemistry was determined from the NOESY spectroscopy. The NOEs correlations between H-1/H-5/H-3ax and H-2eq suggested 108 that there are on the same side, whereas further NOEs correlations between H-4/H-2ax/H-3eq 109 indicated that there are in the opposite side. In addition, the large coupling constant  $(J_{4/5} = 9.9)$ 110 111 Hz) between the two oxymethines H-4 and H-5 indicated that these protons are in a transdiaxial orientation. Furthermore, the large coupling constant ( $J_{1/2} = 11.8$  Hz) between H-1 and 112 H-2ax suggested also that H-1 is in axial position. In order to confirm the relative 113 configuration of **3**, we took advantage of the recently introduced software package Spinach 114 115 (Edwards et al., 2014; Hogben et al., 2011). Following a classical procedure used for DP4 calculations, namely, geometry optimization followed by frequency calculations, NMR 116 117 properties were calculated using the Gaussian 16W software using density functional theory at the B3LYP/STO-3G level for all eight diastereoisomers (Figure S34 supporting 118 119 information, compound 3 a-h). In order to stick to the experimental NOESY spectrum, all the 120 parameters (spectral window, magnet strength, number of points and mixing time) were set as for the data acquisition. To facilitate the comparison between experimental and theoretical 121 spectra, the Zeeman interactions were defined as scalars from the experimental data. The 122 identification of the right relative configuration was then deduced using a "spot the 123 difference" approach between experimental NOESY and simulated one. As shown Figure 124 S34, the  $1R^*, 4R^*, 5R^*, 6R^*$ -3a was ruled out based because of the two differences observed, 125 the absence of crosspeak between H-1/H-3ax and the additional cross-peak between H-10b/H-126 13. The diasteroisomers 3-b, 3-d, 3-f and 3-h were dismissed because of the absence of 127 crosspeaks H1/H-3ax and H-5/H-3ax together with the presence H-5/H-3eq. The 128 diastereoisomer 3-c was discarded based on the absence of crosspeak corresponding to the H-129 1/H-3ax. Eventually, 3-e was excluded due to the absence of the intense crosspeak H10a/H-13 130 on the experimental spectrum. The relative configuration was then assigned as 131  $1S^{*}, 4S^{*}, 5R^{*}, 6R^{*}$ -3. This approach was further confirmed by the proton couplings similar to 132

compound 2. The absolute configuration of 3 was also determined as 1*R*, 4*R*, 5*S*, 6*S* as both
experimental and calculated ECD spectra nicely looked alike (Figure S82, *supporting information*). Therefore, the structure 3 was established and named stagonosporyn C (3).

The molecular formula  $C_{12}H_{14}O_5$  was determined for compound 4 according to its HR-136 ESI-MS m/z 237.0755 [M-H]<sup>-</sup> (calc. 237.0763), suggesting six degrees of unsaturation. The IR 137 spectrum exhibited characteristic absorption bands at 3378 and 1790 cm<sup>-1</sup> for hydroxyl and 138 carbonate carbonyl functionality. The <sup>13</sup>C NMR spectrum (**Table 3**) of **4** exhibited 12 carbon 139 resonances consisting of one sp<sup>3</sup> methyl CH<sub>3</sub>-11 ( $\delta_{C}$  22.9), two sp<sup>3</sup> methylenes CH<sub>2</sub>-2 ( $\delta_{C}$ 140 24.9) and CH<sub>2</sub>-3 ( $\delta_C$  20.9), three sp<sup>3</sup> oxymethines CH-1 ( $\delta_C$  73.6), CH-4 ( $\delta_C$  78.9) and CH-5 141 ( $\delta_{\rm C}$  82.9), one oxygenated sp<sup>3</sup> carbon C-6 ( $\delta_{\rm C}$  73.2), one sp<sup>2</sup> methine CH<sub>2</sub>-10 at ( $\delta_{\rm C}$  123.4), 142 two sp<sup>2</sup> carbons C-9 and C-12 corresponding to an olefinic one and a carbonate carbonyl at 143 ( $\delta_C$  127.6) and ( $\delta_C$  157.6), respectively. Two sp acetylenic C-7 and C-8 carbons at ( $\delta_C$  86.7) 144 and ( $\delta_{\rm C}$  90.1), respectively were also depicted on <sup>13</sup>C NMR. The <sup>1</sup>H NMR and <sup>1</sup>H -<sup>1</sup>H COSY 145 data (Table 1 & Figure 2) of 4 suggested that this compound is also an isoprenylated 146 cyclohexanoid structurally related to the aforementioned stagonosporyne A (1) except for the 147 presence of an additional carbonate carbonyl carbon at  $\delta_C$  157.6. This cyclic carbonate 148 functionality was further confirmed by HMBC correlations observed between both 149 150 oxymethine protons H-4 ( $\delta_{\rm H}$  4.95) and H-5 ( $\delta_{\rm H}$  4.72) with the carbon C-12 ( $\delta_{\rm C}$  157.6) establishing the carbonate moiety at C-4 and C-5. Therefore compound 4 was assigned as a 151 carbonate derivative of compound 1. NOEs correlation between H-4, H-5, H2-ax and H-3eq 152 suggested that these protons are orientated on the same side and confirmed the *cis* relationship 153 between H-4 and H-5. This conclusion was confirmed by the coupling constant ( $J_{4/5} = 6.5$ ) 154 between H-4 and H-5 suggesting a torsion angle of ca 40° between these two protons. In 155 addition, coupling constant of H-1 ( $J_{e/a}$  4.9,  $J_{e/e}$  1.5) suggested that H-1 is in equatorial 156 157 position. The relative configuration was assigned using the NOESY simulation (Figure S42, supporting information) approach described above and the  $1S^*$ ,  $4R^*$ ,  $5R^*$ ,  $6S^*$  configuration 158 could be assigned for **4**. The absolute configuration was determined by comparison of the sign 159 of the unique CE on the ECD spectrum with the other compounds (Figure S82, supporting 160 information). Structure 4 was thus elucidated as the stagonosporyne D. 161

Molecular formula  $C_{12}H_{12}O_5$  was established to Stagonosporyne E (**5**) according to its negative HR-ESI-MS at m/z 235.0593 [M-H]<sup>-</sup> (235.0606 calcd. for  $C_{12}H_{11}O_5$ ), indicating seven degrees of unsaturation. The <sup>13</sup>C and <sup>1</sup>H NMR data as well 2D NMR spectral data interpretation including <sup>1</sup>H-<sup>1</sup>H COSY, <sup>1</sup>H-<sup>13</sup>C HSQC and HMBC (**Figure 2**) indicated that

compound 5 displays similar framework as deduced before for stagonosporyne D (4). The key 166 difference between the two compounds was the replacement of the two methylenes CH<sub>2</sub>-2 and 167 CH<sub>2-3</sub> in 4 by an *endo*-cyclic double bond suggesting a 1,4,5,6-tetraoxygenated cyclohexene 168 ring system. This observation was consistent with the molecular formula determined by 169 HRMS. The relative configuration was deduced from NOESY and <sup>1</sup>H NMR coupling 170 constants analyses. Indeed, the coupling constant  $J_{1/2} = 5.0$  Hz suggested that H-1 is in 171 *pseudo*-equatorial position. The coupling constants  $J_{4/5} = 7.6$  Hz as well as the NOE 172 correlations between H-4 and H-5 characterized their cis relationship. The NOE correlation 173 174 observed between H1/H5-ax on the NOESY simulation spectra (Figure S51, supporting *information*) confirmed the  $1S^*$ ,  $4R^*$ ,  $5R^*$ ,  $6S^*$ -5. In agreement with the relative configuration 175 of 4. The ECD spectrum was similar to the one obtained for 5 thus the absolute configuration 176 was assigned accordingly. The structure for compound 5 was proposed and named 177 178 stagonosporyne E.

The molecular formula of compound 6 was determined as  $C_{11}H_{14}O_3$  according to its 179 negative HR-ESI-MS at m/z 193.0854 [M-H]<sup>-</sup>, which requires five double bond equivalents. 180 Examination of its <sup>1</sup>H and <sup>13</sup>C NMR data (Table 2 & 3) indicated the presence of a 1,4,6-181 trihydroxy cyclohexene core bearing an *endo*-cyclic double bond at C2-C3 at ( $\delta_{\rm H}$  5.52/5.75, 182  $\delta_{\rm C}$  129.8/133.4 respectively), satisfying the fifth unsaturation degree. The COSY in 183 combination with HSQC, HMBC (Figure 2) supported the identification of structure 6 as the 184 stagonosporyn F (Figure 1). The analysis of its <sup>1</sup>H NMR spectrum (Table 2) showed that the 185 large coupling constant ( $J_{4/5} = 9.4$  Hz) between H-4 and H-5ax suggests that there are in 186 trans-diaxial position while the coupling constant  $(J_{1/2} = 2.1 \text{ Hz})$  between H-1 and H-2 187 suggests that H-1 is in *pseudo*-axial position. This was further confirmed by key NOEs 188 correlations between H-4 and H-5eq and H-1 and H-5ax and NOESY simulation spectra 189 190 (Figure S60, supporting information). The absolute configuration was thus assigned as 1S\*, 4*R*\*, 6*S*\*-**6**. 191

192 Compound 7 was purified in minute quantity (0.5 mg). It displayed a molecular 193 formula  $C_{11}H_{14}O_3$  at m/z 193.0854, [M-H]<sup>-</sup>, established by HRESIMS, indicating five 194 unsaturation degrees. The <sup>1</sup>H NMR and <sup>13</sup>C (**Table 2 & 3**) of 7 displayed similarities with 195 those recorded for stagonosporyne F (6). Indeed, the key difference was the presence of an 196 additional oxymethine H-5 ( $\delta_H$  3.41,  $\delta_C$  75.6) and as well as a sp<sup>3</sup> methine H-6 ( $\delta_H$  2.62,  $\delta_C$ 197 45.3) instead of the hydroxylated carbon C-6, previously observed for compound 6. The 198 COSY spectrum (**Figure 1**) revealed the presence of a main spin system from H-1 to H-6

corresponding to a 1,4,5-tri-oxygenated cyclohexene system. The HMBC spectrum (Figure 2) 199 displayed key correlations that confirmed this motif. Indeed, the two oxymethines H-1 ( $\delta_{\rm H}$ 200 4.20) and H-5 ( $\delta_H$  3.41) exhibited HMBC correlations with the methine C-6 at ( $\delta_C$  45.3). 201 Further key HMBC correlations were observed from both H-5 ( $\delta_{\rm H}$  3.41) and H-6 ( $\delta_{\rm H}$  2.62) 202 with the quaternary carbon C-7 at  $\delta_{\rm C}$  88.8, confirming the attachment of an isoprenvl side 203 chain at C-6. The <sup>1</sup>H NMR spectrum (**Table 2**) showed characteristic large coupling constants 204  $(J_{5/6} 11.4, J_{5/4} 8.0 \text{ and } J_{6/1} 9.3 \text{ Hz})$  suggesting that H-1 and H-6 as well as H-4 and H-5 are in 205 trans diaxial positions. This was further confirmed by strong NOE correlations between H-1 206 and H-5 as well as between H-4 and H-6. Based on all previously mentioned spectral data, 207 structure 7 was therefore assigned as stagonosporyne G (7). The ECD spectrum was similar to 208 209 the one obtained for 5 thus the absolute configuration was assigned 1S, 4S, 5S, 6R 210 accordingly.

Compound 8 was purified as pale yellow oil material. It had a molecular formula 211  $C_{11}H_{11}O_2$ , established by HRESIMS at m/z 175,0747 [M+H]<sup>+</sup> (calcd for 175,0754), indicating 212 seven degrees on unsaturation. A preliminary investigation of its <sup>1</sup>H NMR spectrum showed 213 214 characteristic signals of a 1,2,4-trisubstituted benzene ring, consisting of a meta-coupled proton H-1 ( $\delta_{\rm H}$  6.70, d, 2.7) and two ortho-coupled aromatic protons H-3 ( $\delta_{\rm H}$  6.64, dd, 8.6, 215 2.7) and H-4 ( $\delta_{\rm H}$  6.67, d, 8.6). The full extensive 2D NMR spectral data analysis involving 216 COSY, HSQC and HMBC support the identified structure 8 as siccayne, a previously isolated 217 compound from the marine fungi Halocyphina villosa (Kupka et al., 1981). 218

The acetylenic containing metabolites display a myriad of important biologicalactivities

(Kuklev et al., 2013) and among them herbicidal and antimicrobial ones that have already 221 been reported. Especially phomentrioloxin B, an acetylenic polyoxyenated cyclohexene 222 structurally related to our isolated compounds and previously isolated from the fungus 223 Diaporthe gulave caused necrotic spots on Papaver rhoes and Urtica dioica at 5mM (Andolfi 224 et al., 2014). Furthermore, closely related compounds were isolated from the Australian native 225 grass, Neurachne alopecuroidea and showed in vitro growth promotion of barley seedlings 226 227 and potent activity against several phytopathogens such as Phytophtora cinnamomi, Rhizoctonia salona, Pythium irregular and Alternaria alternata. 228

Accordingly, the pure isolated natural products stagonosporyne A-G (1-7) along with siccayne (8) were tested for their herbicidal and antifungal activities. Stagonosporynes A-G (1-7) did not display antifungal activity against the two phytopathogens *Fusarium oxysporum* and *Botrytis cinerea* (data not shown). Stagonosporynes A-F (1-6) did not exhibit herbicidal activity while stagonosporyne G (7) displayed a significant herbicidal activity at the highest tested concentration (100  $\mu$ g/mL) against the laboratory model plant *Arabidopsis thaliana* as shown in **Figure 5**.

#### 236 **3.** Conclusion

The chemical investigation of the wheat plant pathogen *Parastagonospora nodorum* SN15 led to the purification of seven highly oxygenated acetylenic cyclohexanoids named stagonosporynes A-H (1-7). All compounds were evaluated for their herbicidal activity and stagonosporyne G (7) displayed the most significant herbicidal activity. Thus, taken together, these results suggest that stagonosporyne G may belong to the chemical arsenal required for the virulence of *P. nodorum* SN15. Further experiments are therefore needed to determine the precise role of this compound in *P. nodorum* wheat colonization.

#### 244 **4. Experimental.**

#### 245 *4.1.General Experimental Procedures.*

Optical rotations  $[\alpha]^{20}$  <sub>D</sub> were determined using a Perkin Elmer 341 polarimeter. IR 246 spectra were performed on a PerkinElmer BX FR-IR spectrometer. The CD spectra were 247 finished on Jasco J-810 spectropolarimeter system. High-resolution mass spectra were 248 249 recorded on an API Q-STAR PULSAR i of Applied Biosystem by direct injection of the purified compounds. 1D and 2D NMR spectra were recorded on Bruker Avance III HD 400 250 251 MHz and 600 MHz spectrometers (Wissembourg, France) equipped with a BBFO Plus Smart probe and a triple resonance TCI cryoprobe, respectively (CNRS-MNHN). The chemical 252 253 shifts are relative to the residual signal solvent (CD<sub>3</sub>OD:  $\delta_{\rm H}$  3.31;  $\delta_{\rm C}$  49.15; CD<sub>3</sub>CN:  $\delta_{\rm H}$  1.94; δ<sub>C</sub> 118.26, CDCl<sub>3</sub>: δ<sub>H</sub> 7.26; δ<sub>C</sub> 77.16; DMSO-d6: δ<sub>H</sub> 2.50; δ<sub>C</sub> 39.52). Preparative HPLC 254 255 experiments were performed on an Agilent system and an Agilent PrepHT XDB-C18 column  $(21.2 \times 150 \text{ mm i.d.}; 5 \mu\text{m}; \text{USA})$ . Column chromatography (CC) was performed using silica 256 257 gel (Geduran Si 60, 40–63 µm, Merck, Germany and Lichroprep RP-18, 40–63µm, Merck KGaA, Germany); TLC silica gel-precoated plates (F254, 20 × 20 cm, Merck KGaA, 258 Germany). 259

#### 260 *4.2. Microorganism.*

The fungus *Parastagonospora nodorum* SN15 was grown in 5L Erlenmeyer flasks (5 x 600 mL inoculated with 5mL fungal suspension in glycerol 10%) at 25 °C for 21 days under a 12 h photoperiod culturing parameters.

#### 264 *4.3. Extraction and isolation.*

The rice culture of the fungus *Parastagonospora nodorum* SN15 was extracted by 2 L 265 of ethyl acetate. Organic phases were combined and dried over anhydrous MgSO<sub>4</sub>, filtrated 266 and concentrated under reduced pressure to afford 19 g of brown gummy material. The 267 268 methanolic soluble part (4.75 g) was subjected to a normal phase silica-gel flash chromatography (35-70  $\mu$ m), using a gradient of cyclohexane/ CH<sub>2</sub>Cl<sub>2</sub> [100/0 $\rightarrow$ 0/100]. The 269 column was further eluted using a gradient of CH<sub>2</sub>Cl<sub>2</sub>/MeOH [100/0 $\rightarrow$ 0/100] and terminated 270 with ethyl acetate/acetone/water/formic acid (5/3/0.5/0.5) to afford 25 fractions. All fractions 271 were monitored by TLC using different eluent systems and combined together according to 272 their polarities in 13 major fractions F1-F13. Promising fractions F8, F9 and F10 were 273 introduced for further reparative chromatographic purifications by reversed phase 274 (preparative/semi-preparative and analytical C<sub>18</sub>-HPLC) using the eluent system 275 water/acetonitrile containing 0.1% formic acid. The detailed isolation scheme of the isolated 276 compounds and their quantities is given in the supporting information. 277

#### 278 *4.4. Computational details.*

All Density functional theory calculations were performed using Gaussian 16W. After a conformational analysis with the GMMX module and the MMFF94 force field ( $\Delta E < 1.5$ kcal/mol), a geometry optimization was realized at the B3LYP/STO-3G level. Rotational strength was calculated using TD DFT at the B3LYP/6.31g level for 20 excited states. Total nuclear spin-spin coupling *J* (Hz) were predicted at the B3LYP/STO-3G level. Zeeman interactions were defined as scalars from the experimental data.

#### 285 *4.5. Biological activities.*

Surface sterilized Arabidopsis thaliana Columbia-0 seeds were sawn in 96-well plates 286 (4-5 seeds per well) and incubated at 22°C under continuous light, in the presence of 200 µL 287 half-strength Murashige and Skoog (MS) medium (Duchefa, France). After 5 days, MS 288 medium was removed and replaced by 200 µL of fresh medium supplemented with 289 stagonosporyne G dissolved in DMSO. For all conditions, the final concentration of DMSO 290 represented 1% of the total volume of medium. For glyphosate controls, a ready-to-use 291 commercial solution of Roundup<sup>®</sup> containing 7.2 g.L<sup>-1</sup> of acid glyphosate was diluted in MS 292 medium to a final concentration of 72µg.mL<sup>-1</sup>. Plantlets were photographed after 7 days and 293 294 survival rate was established as the percentage of fully-green plantlets over the total number 295 of developed seedlings.

297 Stagonosporyne A (1): colorless oil (1.2 mg);  $[\alpha]^{20}$  <sub>D</sub> + 6 (c = 0.80, MeOH), ECD (0.8

298 mg/mL, MeOH); IR (neat)  $v_{max}$  3443, 2955, 2925, 2870, 1725, 1463, 1384, 1270, 1122, 1073,

299 741 cm<sup>-1</sup>; <sup>1</sup>H NMR and <sup>13</sup>C NMR, **Table 1 and 3**; HRESIMS m/z 219.0682 [M+Na]<sup>+</sup> (calcd

 $\ \ \, \text{ for } C_{11}H_{16}O_3Na,\,219.0678).$ 

- 301 Stagonosporyne B (2): colorless oil (1.6 mg);  $[\alpha]^{20} D + 10$  (c = 1.06, MeOH), ECD (0.8
- 302 mg/mL, MeOH); IR (neat)  $v_{max}$  3369, 2922, 2853, 1716, 1601, 1455, 1395, 1384 cm<sup>-1</sup>; <sup>1</sup>H
- 303 NMR and <sup>13</sup>C NMR, **Table 1 and 3**; HRESIMS m/z 235.09319 [M+Na]<sup>+</sup> (calcd for 304 C<sub>11</sub>H<sub>16</sub>O<sub>4</sub>Na, 235.0940).
- 305 Stagonosporyne C (3): colorless oil (1.3 mg);  $[\alpha]^{20} + 14$  (c = 0.86, MeOH); ECD (0.65

306 mg/mL, MeOH); IR (neat)  $v_{max}$  3383, 2923, 2854, 1716, 1601, 1455, 1395, 1384, 1373 cm<sup>-1</sup>;

307 <sup>1</sup>H NMR and <sup>13</sup>C NMR, **Table 1 and 3**; HRESIMS m/z 293.0796 [M+K]<sup>+</sup> (calcd for

- $308 \quad C_{13}H_{18}O_5K, 293.0785).$
- 309 Stagonosporyne D (4): colorless oil (2 mg);  $[\alpha]^{20} _{D}$  + 26 (c = 1.33, MeOH); ECD (0.66
- 310 mg/mL, MeOH); IR (neat) v<sub>max</sub> 3378, 2923, 2855, 1790, 1733, 1602, 1445, 1395, 1384, 1373,
- 311 1265, 1196 cm<sup>-1</sup>; <sup>1</sup>H NMR and <sup>13</sup>C NMR, **Table 1 and 3**; HRESIMS *m/z* 237.0755 [M-H]<sup>-</sup>

312 (calcd for  $C_{12}H_{13}O_5$ , 237.0768).

- Stagonosporyne E (**5**): colorless oil (1.2 mg);  $[\alpha]^{20} _{D}$  + 24 (*c* = 0.8, MeOH); ECD (0.8 mg/mL, MeOH); IR (neat)  $v_{max}$  3372, 2922, 2853, 1733, 1601, 1455, 1395, 1373, 1255, 1099 cm<sup>-1</sup>; <sup>1</sup>H NMR and <sup>13</sup>C NMR, **Table 1 and 3**; HRESIMS *m/z* 235.0593 [M-H]<sup>-</sup> (calcd for C<sub>12</sub>H<sub>11</sub>O<sub>5</sub>, 235.0611).
- 317 Stagonosporyne F (6): pale yellow oil (2.4 mg);  $[\alpha]^{20}$  D + 67 (c = 1.60, MeOH); ECD (0.24
- 318 mg/mL, MeOH); IR (neat)  $v_{max}$  3347, 2926, 1613, 1374, 1266, 1054, 838 cm<sup>-1</sup>; <sup>1</sup>H NMR and
- <sup>13</sup>C NMR, **Table 2 and 3**; HRESIMS m/z 193.0854 [M-H]<sup>-</sup> (calcd for C<sub>11</sub>H<sub>14</sub>O<sub>3</sub>, 193.0870).
- 320 Stagonosporyne G (7): colorless oil (0.5 mg);  $[\alpha]^{20}$  <sub>D</sub> + 18 (c = 0.16, MeOH); ECD (0.5
- 321 mg/mL, MeOH); IR (neat)  $v_{max}$  3343, 2924, 1602, 1448, 1395, 1384, 1373, 1196, 1098, 1035,
- 322 700 cm<sup>-1</sup>; <sup>1</sup>H NMR and <sup>13</sup>C NMR, **Table 2 and 3**; HRESIMS *m/z* 193.0854 [M-H]<sup>-</sup> (calcd for
- 323  $C_{11}H_{14}O_3$ , 193.0870).

#### 324 Acknowledgements

325 This work was supported by ATM "Microorganisms" grant from the Natural History Museum

- of Paris. The 400 MHz and 600 MHz NMR spectrometers used in this study were funded
- 327 jointly by the Région Ile-de-France, the MNHN (Paris, France) and the CNRS (France).
- 328

#### 329 **Réferences**

- 330 Andolfi, A., Maddau, L., Basso, S., Linaldeddu, B. T., Cimmino, A., Scanu, B., Deidda, A.,
- Tuzi, A., Evidente, A., 2014. Diplopimarane, a 20-nor-ent-pimarane produced by the oak
  pathogen *Diplodia quercivora*. J. Nat. Prod. 77, 2352-2360.
- Arakawa, H., 1968. Absolute configuration of mellein. Bull. Chem. Soc. Jap. 41, 2541.
- Blair, J., Newbold, G. T., 1955. The structure of mellein. Chem. Ind. (London, U. K.), 93-94.
- 335 Chomcheon, P., Sriubolmas, N., Wiyakrutta, S., Ngamrojanavanich, N., Chaichit, N.,
- Mahidol, C., Ruchirawat, S., Kittakoop, P., 2006. Cyclopentenones, scaffolds for organic
- syntheses produced by the Eendophytic fungus mitosporic Dothideomycete sp. LRUB20. J
  . Nat. Prod 69, 1351-1353.
- Chomcheon, P., Wiyakrutta, S., Sriubolmas, N., Ngamrojanavanich, N., Mahidol, C.,
  Ruchirawat, S., Kittakoop, P., 2009. Metabolites from the endophytic mitosporic *Dothideomycete* sp. LRUB20. Phytochemistry 70, 121-127.
- 342 Chooi, Y. H., Krill, C., Barrow, R. A., Chen, S., Trengove, R., Oliver, R. P., Solomon, P. S.,
- 2015. An in planta-expressed polyketide synthase produces (*R*)-mellein in the wheat
  pathogen *Parastagonospora nodorum*. Appl. Environ. Microbiol. 81, 177-186.
- Chooi, Y. H., Muria-Gonzalez, M. J., Solomon, P. S., 2014. A genome-wide survey of the
  secondary metabolite biosynthesis genes in the wheat pathogen *Parastagonospora nodorum*. Mycology 5, 192-206.
- Devys, M., Barbier, M., Bousquet, J.-F., Kollmann, A., 1994. Isolation of the (-)-(*3R*)-5hydroxymellein from the fungus *Septoria nodorum*. Phytochemistry 35, 825-826.
- Devys, M., Barbier, M., Kollmann, A., Bousquet, J.-F., 1992. N-Methoxy septorinol, a
  substituted pyrazine from the fungus *Septoria nodorum*. Phytochemistry 31, 4393-4394.
- Edwards, L. J., Savostyanov, D. V., Welderufael, Z. T., Lee, D., Kuprov, I., 2014. Quantum
- mechanical NMR simulation algorithm for protein-size spin systems. J. Magn. Reson. 243,
  107-113. .
- Figueroa, M., Hammond-Kosack, K. E., Solomon, P. S., 2018. A review of wheat diseases—a
  field perspective. Mol. Plant Pathol. 19, 1523-1536.

- Frisch, M. J., Trucks, G. W., Schlegel, H. B., Scuseria, G. E., Robb, M. A., Cheeseman, J. R., 357 Scalmani, G., Barone, V., Petersson, G. A., Nakatsuji, H., Li, X., Caricato, M., Marenich, 358 A. V., Bloino, J., Janesko, B. G., Gomperts, R., Mennucci, B., Hratchian, H. P., Ortiz, J. 359 V., Izmaylov, A. F., Sonnenberg, J. L., Williams, Ding, F., Lipparini, F., Egidi, F., Goings, 360 J., Peng, B., Petrone, A., Henderson, T., Ranasinghe, D., Zakrzewski, V. G., Gao, J., Rega, 361 N., Zheng, G., Liang, W., Hada, M., Ehara, M., Toyota, K., Fukuda, R., Hasegawa, J., 362 Ishida, M., Nakajima, T., Honda, Y., Kitao, O., Nakai, H., Vreven, T., Throssell, K., 363 Montgomery Jr., J. A., Peralta, J. E., Ogliaro, F., Bearpark, M. J., Heyd, J. J., Brothers, E. 364 N., Kudin, K. N., Staroverov, V. N., Keith, T. A., Kobayashi, R., Normand, J., 365 Raghavachari, K., Rendell, A. P., Burant, J. C., Iyengar, S. S., Tomasi, J., Cossi, M., 366 Millam, J. M., Klene, M., Adamo, C., Cammi, R., Ochterski, J. W., Martin, R. L., 367 Morokuma, K., Farkas, O., Foresman, J. B., Fox, D. J., 2016. Gaussian 16 Rev. B.01. 368
- Hane, J. K., Lowe, R. G., Solomon, P. S., Tan, K. C., Schoch, C. L., Spatafora, J. W., Crous,
  P. W., Kodira, C., Birren, B. W., Galagan, J. E., Torriani, S. F., McDonald, B. A., Oliver,
  R. P., 2007. Dothideomycete plant interactions illuminated by genome sequencing and
  EST analysis of the wheat pathogen *Stagonospora nodorum*. Plant Cell 19, 3347-3368.
- Hewage, R. T., Aree, T., Mahidol, C., Ruchirawat, S., Kittakoop, P., 2014. One strain-many
  compounds (OSMAC) method for production of polyketides, azaphilones, and an
  isochromanone using the endophytic fungus *Dothideomycete* sp. Phytochemistry 108, 8794. .
- Hogben, H. J., Krzystyniak, M., Charnock, G. T., Hore, P. J., Kuprov, I., 2011. Spinach--a
  software library for simulation of spin dynamics in large spin systems. J. Magn. Reson.
  208, 179-194.
- Ipcho, S. V., Hane, J. K., Antoni, E. A., Ahren, D., Henrissat, B., Friesen, T. L., Solomon, P.
  S., Oliver, R. P., 2012. Transcriptome analysis of *Stagonospora nodorum*: gene models, effectors, metabolism and pantothenate dispensability. Mol. Plant Pathol. 13, 531-545.
- Kuklev, D. V., Domb, A. J., Dembitsky, V. M., 2013. Bioactive acetylenic metabolites.
  Phytomedicine 20, 1145-1159.
- 386 Kupka, J., Anke, T., Steglich, W., Zechlin, L., 1981. Antibiotics from basidiomycetes. XI.
- 387 The biological activity of siccayne, isolated from the marine fungus *Halocyphina villosa* J.
- 388 & E. Kohlmeyer. J. Antibiot. (Tokyo) 34, 298-304.

369

Wallingford, CT.

- Li, H., Hu, J., Wei, H., Solomon, P. S., Vuong, D., Lacey, E., Stubbs, K. A., Piggott, A. M.,
  Chooi, Y. H., 2018. Chemical ecogenomics-guided discovery of phytotoxic alpha-Pyrones
  from the fungal wheat pathogen *Parastagonospora nodorum*. Org. Lett. 20, 6148-6152.
- Muria-Gonzalez, M. J., Chooi, Y. H., Breen, S., Solomon, P. S., 2015. The past, present and future of secondary metabolite research in the Dothideomycetes. Mol. Plant. Pathol. 16, 92-
- 394 107.
- Nishikawa, E., 1933. Biochemisty of molds. II. A metabolic product of *Aspergillus melleus*Yukawa. Nippon Nogei Kagaku Kaishi 9, 772-774.
- Senadeera, S. P. D., Wiyakrutta, S., Mahidol, C., Ruchirawat, S., Kittakoop, P., 2012. A novel
  tricyclic polyketide and its biosynthetic precursor azaphilone derivatives from the
  endophytic fungus *Dothideomycete* sp. Org. Biomol. Chem. 10, 7220-7226.
- Solomon, P. S., Lowe, R. G., Tan, K. C., Waters, O. D., Oliver, R. P., 2006. *Stagonospora nodorum*: cause of stagonospora nodorum blotch of wheat. Mol. Plant. Pathol. 7, 147-156.
- 402 Stergiopoulos, I., Collemare, J., Mehrabi, R., De Wit, P. J., 2013. Phytotoxic secondary
- 403 metabolites and peptides produced by plant pathogenic *Dothideomycete* fungi. FEMS
  404 Microbiol. Rev. 37, 67-93.
- Tan, K.-C., Trengove, R. D., Maker, G. L., Oliver, R. P., Solomon, P. S., 2009. Metabolite
  profiling identifies the mycotoxin alternariol in the pathogen *Stagonospora nodorum*.
  Metabolomics 5, 330-335.
- Yang, X.-L., Awakawa, T., Wakimoto, T., Abe, I., 2013. Induced biosyntheses of a novel
  butyrophenone and two aromatic polyketides in the plant pathogen *Stagonospora nodorum*. Nat. Prod. Bioprospect. 3, 141-144.
- 411
- 412
- 413
- 414
- 415
- 416
- 417

418

419

- 420
- 421
- 422
- 423
- 424



Stagonosporyne A (1)

OH

Ó

OH





Stagonosporyne B(2)



Stagonosporyne  $E(\mathbf{5})$ 

Stagonosporyne C (3)



Stagonosporyne F (6)



**`O** Stagonosporyne D (**4**)

Stagonosporyne G(7)

Figure 1.



Siccayne (8)















COSY — HMBC 7

Figure 2.



Figure 3.



Figure 4.



Figure 5.

	Table	1.
--	-------	----

pos.	Stagonosporyne A (1)		Stagonosporyne B (2) <sup>a</sup>		Stagonosporyne C (3) <sup>c</sup>	Stagonosporyne D (4) <sup>a</sup>	
1	3.72, dd (10.6, 4.5) <sup>c</sup>	3.37, m <sup>b</sup>	3.46, dd (11.8,4.4) <sup>a</sup>	3.27, m <sup>b</sup>	3.50, dd (11.8, 4.5)	3.96, dd (4.9, 1.5)	
2ax	1.68, m	1.51, m	1.66, dddd (13.7, 11.8, 11.8, 4.4)	1.43, m	1.71, m	2.15, m	
2eq	1.83, m		1.81, m	1.57, m	1.84, m	1.81, m	
3ax	1.35, dddd (2.7, 12.7, 10.1, 3.9)	1.16, m	1.33, m	1.15, m	1.37, m	2.19, m	
3eq	1.95, ddt (12.7, 6.9, 4.2)	1.71, m	1.90, m	1.67, m	1.99, m	2.02, m	
4	3.97, dddd (10.1, 10.1, 4.2, 4.2)	3.60, m	3.59, ddd (11.4, 9.4, 5.0)	3.35, m	4.83, ddd (11.4, 9.9, 5.1)	4.95, m	
5ax	1.66, m	1.38, m	3.17, d (9.4)	2.95, d (9.2,4.4)	3.42, d (9.9)	4.72, d (6.5)	
5eq	2.38, ddd (13.5, 4.2, 2.5)	2.06, m					
6	-	-	-	-	-	-	
7	-	-	-	-	-	-	
8	-	-	-	-	-	-	
9	-	-	-	-	-	-	
10	5.29, m	5.24, m	5.25 dq (3.2,1.5)	5.19, br s	5.36, br s	5.31, m	
	5.24, dq (3.2, 1.5)	5.19, m	5.32 dq (3.2, 1.1)	5.23, br s	5.27, br s	5.29, m	
11	1.86, dd (1.5, 1.1)	1.82, dd (1.5, 1.1)	1.91, dd (1.1,1.5)	1.82, br s	1.93, dd (1.5, 1.1)	1.86, dd (1.1, 1.5)	
12	-	-	-	-	-	-	
13	-	-	-	-	2.05, s	-	
1-OH	2.19, br s	4.59, d (6.6)	-	4.66, d (4.3)	-	-	
4-OH	2.57, br s	4.42, d (5.0)	-	4.56, d (4.4)	-	-	
5-OH	-	-	-	4.85, d (4.4)	-	-	
6-OH		4.89, d (1.3)		5.29, s			

a: Recorded in CD<sub>3</sub>OD, b: recorded in DMSO-d6, c: recorded in CDCl<sub>3</sub>, d: recorded in CD<sub>3</sub>CN,

### Table 2.

pos.	Stagonosporyne E (5) <sup>a</sup>	Stagonosporyne F (6) <sup>a</sup>	Stagonosporyne G (7) <sup>a</sup>
1	4.24, d (5.0)	4.23, ddd (2.1, 2.1, 2.1)	4.20, m
2	6.24, ddd (10.1, 5.0, 1.1)	5.52, dt (10.4, 2.1)	5.64, dt (10.2, 2.0)

3	5.99, ddd (10.1, 3.5, 0.6)	5.75, ddd (10.4, 4.0, 2.1)	5.60, dt (10.2, 1.9)		
4	5.30, m	4.34, m	4.03, m		
5ax	4.99, d (7.4)	1.76, dd (13.3, 9.4)	3.41, dd (11.4, 8.0)		
5eq	-	2.46, ddd (13.3, 5.6, 2.1)	-		
6	-	-	2.62, dd (11.4, 9.3)		
7	-	-	-		
8	-	-	-		
9	-	-	-		
10	5.20-5.30, m	5.26, m	5.26-5.20, m		
		5.24, m			
11	1.84, dd (1.5, 1.1)	1.88, dd (1.5, 1.1)	1.90, dd (1.5, 1.1)		

a: Recorded in CDCl<sub>3</sub>

### Table 3.

pos.		1	2	2	3	<b>4</b> <sup>a</sup>	<b>5</b> <sup>a</sup>	<b>6</b> <sup>a</sup>	<b>7</b> <sup>a</sup>
1	74.3, CH <sup>c</sup>	73.7, CH <sup>b</sup>	76.1, CH <sup>a</sup>	74.3, CH <sup>b</sup>	76.3, CH	73.6, CH	70.5, CH	72.8, CH	72.6, CH
2	25.9, CH <sub>2</sub>	27.9, CH <sub>2</sub>	29.4, CH <sub>2</sub>	28.4, CH <sub>2</sub>	29.1, CH <sub>2</sub>	24.9, CH <sub>2</sub>	134.7, CH	129.6, CH	132.1, CH
3	31.9, CH <sub>2</sub>	33.1, CH <sub>2</sub>	29.9, CH <sub>2</sub>	28.9, CH <sub>2</sub>	26.7, CH <sub>2</sub>	20.9, CH <sub>2</sub>	124.6, CH	133.3, CH	130.8, CH
4	65.7, CH	64.0, CH	72.6, CH	70.4, CH	75.5, CH	78.9, CH	75.5, CH	65.1, CH	73.3, CH
5	44.1, CH <sub>2</sub>	46.7, CH <sub>2</sub>	80.9, CH	79.5, CH	78.1, CH	82.9, CH	80.2, CH	44.5, CH <sub>2</sub>	75.6, CH
6	70.6, C	69.8, C	78.1, C	76.5, C	78.3, C	73.2, C	71.2, C	70.5, C	45.3, CH
7	89.7, C	93.8, C	87.1, C	89.3, C	86.7, C	86.7, C	86.8, C	91.9, C	88.8, C
8	85.6, C	82.5, C	89.6, C	86.5, C	90.1, C	90.1, C	89.6, C	84.7, C	84.9, C
9	125.7, C	126.5, C	128.2, C	126.7, C	128.1, C	127.6, C	127.4, C	127.9, C	128.7, C
10	122.9, CH <sub>2</sub>	121.4, CH <sub>2</sub>	122.2, CH <sub>2</sub>	121.1, CH <sub>2</sub>	122.7, CH <sub>2</sub>	123.4, CH <sub>2</sub>	123.4, CH <sub>2</sub>	122.15, CH <sub>2</sub>	121.3, CH <sub>2</sub>
11	23.4, CH <sub>3</sub>	23.3, CH <sub>3</sub>	23.6, CH <sub>3</sub>	23.4, CH <sub>3</sub>	23.5, CH <sub>3</sub>	22.9, CH <sub>3</sub>	23.1, CH <sub>3</sub>	23.1, CH <sub>3</sub>	23.8, CH <sub>3</sub>
12	-	-		-	172.7, C	157.6, C	156.5, C	72.8, CH	72.6, CH
13	-	-		-	21.1, CH <sub>3</sub>	-	-	-	-

a: Recorded in CD<sub>3</sub>OD, b: recorded in DMSO-d6, c: recorded in CDCl<sub>3</sub>, d: recorded in CD<sub>3</sub>CN,

- Figure 1: Structures of stagonosporynes A-G (1-7) along with siccayne (8)
- Figure 2: Key COSY and HMBC correlations for the isolated compounds 1-7.
- Figure 3. Key NOEs for compounds 1 and 2
- Figure 4: Comparison between both experimental and TDDFT predicted ECD spectra of 1
- **Table 1**. <sup>1</sup>H NMR data  $\delta_{\rm H}$  (*J* in Hz) of stagonosporynes A-D (1-4), 600 MHz
- **Table 2**. <sup>1</sup>H NMR data  $\delta_{\rm H}$  (*J* in Hz) of stagonosporynes E-G (**5-7**), 600 MHz
- Table 3. <sup>13</sup>C NMR data of stagonosporynes A-G (1-7), 150 MHz
- Figure 5. Herbicidal activities of stagonosporyne G (7) at 100 µg/mL compared to glyphosate
- tested at 72  $\mu$ g/mL. A) *In vitro* toxic activity and B) survival rate.



## Parastagonospora nodorum SN15

### Herbicidal stagonosporyne G