

1 **Phytotoxins produced by *Phoma chenopodiicola*, a**  
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4 **fungal pathogen of *Chenopodium album***  
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## Abstract

1  
2 Three phytotoxins were isolated from the liquid culture of *Phoma chenopodiicola*, a fungal  
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4 pathogen proposed for the biological control of *Chenopodium album*, a common worldwide weed of  
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6 arable crops. The three phytotoxins appeared to be a new tetrasubstituted furopyran, a new  
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8 tetrasubstituted isocoumarin and a new *ent*-pimaradiene. They were characterized by using  
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10 spectroscopic (essentially 1D and 2D NMR and HR ESI MS) and chemical methods as 3-(3-  
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12 methoxy-2,6-dimethyl-7*aH*-furo[2,3-*b*]pyran-4-yl)-but-2-en-1-ol, 4,5,7-trihydroxy-3-methyl-  
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14 isochroman-1-one, (1*S*,2*S*,3*S*,4*S*,5*S*,9*R*,10*S*,12*S*,13*S*)- 1,3,12-triacetoxy-2,hydroxy-6-oxo-*ent*-  
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16 pimara-7(8),15-dien-18-oic acid 2,18-lactone, and consequently named chenopodolan D,  
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18 chenisocoumarin and chenopodolin B. The absolute configuration to chenisocoumarin was assigned  
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20 applying an advanced Mosher's method through the derivatization of its secondary hydroxylated  
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22 carbon C-4, while that of chenopodolan D by application of quantum mechanical calculations of  
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24 chiroptical (ECD and ORD) properties. Assayed by leaf puncture to non-host weeds Chenopodolan  
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26 D and chenopodolin B showed phytotoxicity while chenisocoumarin and the 9-*O*-acetyl derivative  
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28 of chenopodolan D were inactive. These results confirm that the nature of the side chain at C-4 in  
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30 chenopodolans and in particular its hydroxylation are important features to impart activity. The  
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32 activity of chenopodolin B could also be explained by its possible hydrolyzation into chenopodolin.  
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43 *Keywords:* *Chenopodium album*; *Phoma chenopodiicola*; phytotoxins; furopyrans; *ent*-pimaradiene;  
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45 isocoumarines; herbicides  
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## 1. Introduction

Fungal pathogens responsible for foliar and stem diseases of *Chenopodium album* L., a worldwide diffused weed of arable crops (Holm et al., 1977) also known as common lambsquarters or fat hen, have been studied both as potential biocontrol agents, and as sources of bioactive metabolites. *Ascochyta caulina* (P. Karst.) v.d. Aa and v. Kest has been first extensively studied for both purposes and, more recently, *Phoma chenopodiicola*. From this latter organism a new phytotoxic unrearranged *ent*-pimaradiene diterpene, named chenopodolin (Cimmino et al., 2013a), and successively three novel tetrasubstituted furopyrans, named chenopodolans A-C (Cimmino et al., 2013b) were recently isolated from its liquid culture; the relation between the structural features and the phytotoxic activity against some weeds was also determined by assaying the natural toxins and some of their hemisynthetic derivatives (Cimmino et al., 2013a; Cimmino et al., 2013b).

A further analysis of the organic extracts of the fungus, aimed both at re-isolating chenopodolans A-C to assign their absolute configuration, and to understand the phytotoxicity of the extracts, higher than that expectable by the content of chenopodolin and chenopodolans A-C, allowed to ascertain the presence of further bioactive metabolites and to identify them.

In particular this manuscript reports on: (a) the isolation and the chemical characterization of three new metabolites produced by *P. chenopodicola*, named chenopodolan D (**1**), chenisocoumarin (**2**) and chenopodolin B (**3**) (Fig. 1); (b) the preliminary studies to evaluate their potential as natural herbicides; (c) the assignment of the absolute configuration of **1** by quantum mechanical calculations of Optical Rotatory Dispersion (ORD) and Electronic Circular Dichroism (ECD), and of **2** by applying an advanced Mosher's method.

## 2. Results And Discussion

Compounds **1-3** were obtained from the purification of the organic extract of *P. chenopodiicola* as reported in details in the Experimental. Chenopodolan D (**1**, Fig. 1) has the molecular formula

C<sub>14</sub>H<sub>18</sub>O<sub>4</sub> as deduced by its HR ESIMS spectrum and consistent with six hydrogen deficiencies. Compared to chenopodolans A-C (Cimmino et al., 2013b), its <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1) showed proton and carbon signals very similar to the 2,3,4,6-tetrasubstituted furopyran moiety. Also the signals for the methoxy and the two methyl groups at C-3, C-2 and C-6, respectively, were very similar to those of the other chenopodolans A-C, while the signals of the side chain attached to C-4 differed substantially. In particular the <sup>1</sup>H and <sup>13</sup>C NMR spectra showed a broad singlet and a triplet typical of a hydroxymethylene group (HO-CH<sub>2</sub>-11) at δ 4.38 (2H)/59.8 (t) (Pretsch et al., 2000; Breitmaier and Voelter 1987), instead of the signals of the terminal vinylic methyl group (Me-11) of chenopodolan C (Cimmino et al., 2013b). The presence of a hydroxymethylene is also consistent with the hydroxy group band observed in the IR spectra (Nakanishi and Solomons 1977). These findings suggested for **1** the structure of 11-hydroxychenopodolan C, which was confirmed by the assignment of all the protons and the corresponding carbon chemical shifts based on the couplings observed in the COSY and HSQC spectra (Berger and Braun 2004). Furthermore, the quaternary carbons C-3, C-3a, C-4, C-6, C-7a and C-9 as reported in Table 1 were assigned by the long range couplings observed in the HMBC spectrum (Table 1) (Berger and Braun 2004). Thus chenopodolan D was formulated as 3-(3-methoxy-2,6-dimethyl-7a*H*-furo[2,3-*b*]pyran-4-yl)-but-2-en-1-ol.

This structure was confirmed by all the long range couplings observed in the HMBC spectrum (Table 1) and in particular those between H<sub>2</sub>-11 with both C-9 and C-10. Further supports to the structure assigned to **1** were obtained from its HR ESI MS spectrum which showed: the potassium [2M + K]<sup>+</sup>, sodium [2M + Na] and protonated [2M + H]<sup>+</sup> dimer forms at *m/z* 539, 523 and 501, respectively; the potassium [M + K]<sup>+</sup> and sodium [M + Na]<sup>+</sup> clusters at *m/z* 289 and 273, respectively; the pseudomolecular ion [M + H]<sup>+</sup> *m/z* at 251.1217.

The structure assigned to **1** was finally confirmed by preparing its 11-*O*-acetyl derivative (**4**, Fig. 1) by usual acetylation with pyridine and acetic anhydride. The IR spectrum of **4** showed the absence of any hydroxy group band while its <sup>1</sup>H NMR spectrum (Table 1) differed from that of **1**

essentially for the downfield shift ( $\Delta\delta$  0.41 and 0.39) of the H<sub>2</sub>C-11 signals resonating as two broad singlets at  $\delta$  4.73 and 4.71, respectively, and for the presence of the singlet of the acetyl group at  $\delta$  2.08. Its ESIMS spectrum showed potassium [2M + K]<sup>+</sup>, sodium [2M + Na]<sup>+</sup> and protonated [2M + H]<sup>+</sup> dimer forms, potassium cluster [M + K]<sup>+</sup> and the pseudomolecular ion [M + H]<sup>+</sup> at *m/z*: 623, 607, 585, 331, 293, respectively. Thus, **1** was formulated as 3-(3-methoxy-2,6-dimethyl-7*aH*-furo[2,3-*b*]pyran-4-yl)-but-2-en-1-ol.

The absolute configuration of (-)-**1** was determined by *ab initio* computational analyses of its ORD and ECD spectra (Autschbach, 2009, 2012). This approach was demonstrated to be particularly straightforward and reliable for the absolute configuration assignment of natural products (Mazzeo et al., 2013; Santoro et al., 2013; Santoro et al., 2014), allowing analyses in solution and sometimes on a microscale, using very dilute solutions (Berova et al., 2010). To this end, conformational analysis was carried out on the randomly chosen (*R*)-**1** stereoisomer. The conformational search was first carried out at MM level with MMFF94s force field, then refined by optimization at DFT/B3LYP/TZVP level of theory in gas phase, providing 15 appreciably populated conformers (Fig. S28 and Table S1 in SI). The conformers differ in the rotation of the OCH<sub>3</sub> moiety and of the C4-C9 bond which affect the relative position of the exocyclic double bond in respect to the bicyclic system. UV and ECD spectra of (-)-**1** were recorded in THF in the 210-350 nm range (Fig. 2). The UV spectrum shows a maximum at 210 nm ( $\epsilon$  44600), followed by a shoulder at 229 nm ( $\epsilon$  29900), two weak broad bands at 254 nm ( $\epsilon$  11500) and 286 nm ( $\epsilon$  6090) and a very broad band centered at 326 nm ( $\epsilon$  7800). The ECD spectrum shows relatively intense negative Cotton effect (CE) at 220 nm ( $\Delta\epsilon$  -6.0) followed weaker bands in the low energy spectral range: two positive CEs at 232 nm ( $\Delta\epsilon$  +0.4) and 254 ( $\Delta\epsilon$  +1.1), a negative band at 272 nm ( $\Delta\epsilon$  -0.4) and a broad positive one centred at 288 nm ( $\Delta\epsilon$  +0.62).

The UV and ECD spectra of (*R*)-**1**, were then calculated in gas phase at TDDFT/CAM-B3LYP/aug-cc-pVDZ level on previously found conformers and Boltzmann averaged over

1 conformers populations. Computed ECD spectrum for (*R*)-**1** was then compared with experimental  
2 one for (-)-**1** (Fig. 2). Such comparison shows that, although the low energy range of the theoretical  
3 spectrum is not in full agreement with the experimental, computations correctly reproduced the  
4 strongest Cotton effect at 231 nm in sign and position, suggesting a (*R*) absolute configuration for (-  
5 )-**1**. A close look into the ECD spectra of single conformers (Fig. S29) reveals that, although they  
6 show quite different ECD spectra, most of them display a negative Cotton effect between 210 and  
7 230 nm, pointing out that the sign of this band can be considered as a reliable probe for absolute  
8 configuration assignment of this compound. In order to get a better agreement between computed  
9 and experimental ECD spectra, both conformational analysis and ECD calculation were carried out  
10 taking into account solvation effects by IEFPCM implicit solvation model (Tomasi et. al., 2005).  
11 For this purpose, the outcome from MM conformational analysis on (*R*)-**1** was submitted to  
12 optimization at DFT/B3LYP/TZVP theory level by IEFPCM model (THF) leading again to 15  
13 significantly populated conformers (Fig. S28 and Table S1 in SM). As inferred from Table S1  
14 solvent effects give rise to a redistribution of conformers population with decreasing of the most  
15 populated conformer 2 and increasing of population of conformers 8 and 11. The UV and ECD  
16 spectra for (*R*)-**1** were then calculated at TDDFT/CAM-B3LYP/aug-cc-pVDZ/IEFPCM(THF) level  
17 and Boltzmann averaged over conformers populations. Comparison of computed ECD spectrum for  
18 (*R*)-**1** with experimental one for (-)-**1** (Fig. 2) shows a better agreement than that with the spectrum  
19 computed in gas phase. In fact the IEFPCM computed spectrum better reproduce the low energy  
20 spectral range. Thus, when the solvent effects are taken into account the ECD spectral analysis does  
21 allow to clearly establish a (*R*) absolute configuration for natural (-)-**1**. The good agreement in  
22 position and relative intensity of the ECD CEs with experimental spectrum supports also the  
23 accuracy in the theoretical reproduction of conformers population.

24 To further confirm such configurational assignment computational analysis of ORD curve  
25 was then undertaken. The ORD data of (-)-**1** were recorded in CHCl<sub>3</sub> at *c* = 0.2 g/100ml obtaining a  
26 plane negative curve (Fig. 3). ORD data for (*R*)-**1** were then calculated on previously found  
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1 conformers at TDDFT/B3LYP/aug-cc-pVDZ level in gas phase and Boltzmann averaged over  
2 conformers populations. Comparison of computed and experimental data shows (Fig. 3) an  
3 agreement in sign, both giving rise to a negative plane curve, thus confirming the assignment of (*R*)  
4 absolute configuration for (-)-**1**. However, a difference in one order of magnitude of optical rotation  
5 values was observed. To obtain a better numerical agreement between computed and experimental  
6 data ORD was calculated also taking into account chloroform solvent effects. Therefore,  
7 conformational analysis was carried out by DFT/B3LYP/TZVP/IEFPCM(CHCl<sub>3</sub>) computations,  
8 obtaining 14 stable conformers with populations quite similar to those found in THF. For each  
9 conformer ORD data were computed at TDDFT/B3LYP/aug-cc-pVDZ/IEFPCM(CHCl<sub>3</sub>) level of  
10 theory and Boltzmann averaged over populations, obtaining the ORD curve in Figure 3. As inferred  
11 from Figure 3 computations in solution allow to obtain a slightly better numerical agreement with  
12 experimental, although with data still one order of magnitude greater than experimental. In  
13 conclusion, both ECD and ORD analyses indicate the same assignment of (*R*) absolute  
14 configuration for (-)-**1**. However, computed ORD curve does not match (both in gas phase and  
15 solvent) the experimental data in order of magnitude.

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36 Chenisocoumarin (**2**, Fig. 1) has the molecular formula C<sub>10</sub>H<sub>10</sub>O<sub>5</sub>, as deduced by its HR  
37 ESIMMS and consistent with six hydrogen deficiencies. Its IR spectrum showed bands typical for  
38 hydroxy, ester carbonyl and aromatic groups (Nakanishi and Solomons 1977). Compared to that of  
39 6-hydroxymellein, which was previously isolated as the main phytotoxic metabolite from the  
40 culture filtrates of the same fungus (Cimmino et al., 2013b), the <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 2)  
41 of **2** showed very similar signal systems. Indeed, the <sup>1</sup>H NMR and COSY spectra showed the  
42 presence of two *meta*-coupled aromatic protons (H-6 and H-8) resonating as two doublets (*J* = 1.7  
43 Hz) at δ 6.35 and 6.33, respectively. Their corresponding carbons were observed at δ 106.8 and  
44 99.2, respectively, and were also assigned for their couplings observed in the HSQC spectrum.  
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46 Furthermore, the significant signal of the phenolic protons at C-8 hydrogen bonded with the  
47 carbonyl group at C-1, previously observed in the <sup>1</sup>H NMR spectrum of 6-hydroxymellein, was  
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absent in **2**. Thus, the two hydroxy phenolic groups were located at C-5 and C-7 also on the basis of the chemical shifts of their corresponding quaternary carbons resonating as two singlets at  $\delta$  164.1 and 155.3 (C-7 and C-5, respectively) (Breitmaier and Voelter 1987). These latter were assigned also on the basis of the couplings observed in the HMBC spectrum (see Table 2), as well as the other quaternary carbons at  $\delta$  142.5, 135.8 and 169.0, which were assigned to C-4a, C-8a and the carbonyl group C-1, respectively. Furthermore, the signal systems typical for 4-hydroxymellein were observed in the  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra (Table 2) at  $\delta$  4.57 (H-3, *dq*,  $J = 6.6$  and 1.8 Hz)/67.2(C-3, *d*), 4.38 (H-4, *d*,  $J = 1.8$  Hz)/77.5(C-4, *d*) and 1.48 (Me-9, *d*,  $J = 6.6$  Hz)/15.8 (Me-19, *q*).

Based on these findings, **2** was formulated as 4,5,7-trihydroxy-3-methyl-isochroman-1-one. This structure was supported from all the couplings observed in the HMBC spectrum (Table 2) and the data obtained from its HR ESIMS spectrum. This latter showed the sodium cluster  $[\text{M} + \text{Na}]^+$  and the pseudomolecular ion  $[\text{M} + \text{H}]^+$  at  $m/z$  233 and 211.1218, respectively. The *trans* relative configuration was assigned to **2** on the basis of the significant correlation observed in its NOESY spectrum between H-3 and Me-9, (Berger and Braun 2004).

The absolute configuration of **2** was assigned by applying an advanced Mosher's method (Othani et al, 1991). Besides the suitable secondary hydroxyl group at C-3, **2** also contained two phenol hydroxy groups at C-5 and C-7. First, **2** was converted into the corresponding 5,7-*O,O'*-dimethyl derivative **5** by treatment with an ethereal solution of diazomethane. **5** showed the significant presence of the hydroxy group into the IR spectrum (Nakanishi and Solomon, 1987), while its  $^1\text{H}$  NMR spectrum differed from that of **2** for the presence of the two singlets typical of methoxy groups at  $\delta$  3.92 and 3.89, respectively (Pretsch et al., 2000). Its ESI MS spectrum showed: the potassium  $[2\text{M} + \text{K}]^+$  and the sodium  $[2\text{M} + \text{Na}]^+$  dimeric forms at  $m/z$  515, 499, respectively; the potassium  $[\text{M} + \text{K}]^+$  and sodium  $[\text{M} + \text{Na}]^+$  clusters at  $m/z$  277, 261, respectively; and the pseudomolecular ion  $[\text{M} + \text{H}]^+$  at  $m/z$  239. By reaction with *R*-(-)- $\alpha$ -methoxy- $\alpha$ -trifluoromethylphenylacetyl (MTPA) and *S*-(+)-MTPA chlorides, **5** was converted into the

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corresponding diastereomeric *S*-MTPA and *R*-MTPA monoesters at C-4 (**6** and **7**, respectively), whose spectroscopic data were consistent with the structure assigned to **2**. Subtracting the chemical shift of the protons (Table 3) of the 3-*O*-*R*-MTPA (**7**) from that of 3-*O*-*S*-MTPA (**6**) esters, the  $\Delta\delta$  (**6-7**) values for all of the protons were determined as reported in Figure 4. The positive  $\Delta\delta$  values were located on the right-hand side, and those with negative values on the left-hand side of the model A as reported in Othani et al. (1991). This model allowed the assignment of the *R* configuration at C-4. Considering the relative configuration assigned by NMR as above reported, **2** was formulated as (4*R*,3*S*)-4,5,7-trihydroxy-3-methyl-isochroman-1-one.

Chenopodolin B (**3**) had a molecular formula of C<sub>26</sub>H<sub>32</sub>O<sub>9</sub> as deduced by its HR ESI MS spectrum and consistent with eleven hydrogen deficiencies. The preliminary investigation of its <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 4) showed that they are very similar to those of chenopodolin (Cimmino et al., 2013a). Indeed **3** showed the signal systems typical of the tetracyclic *ent*-pimaradiene lactone and also those corresponding to the  $\alpha,\beta$ -unsaturated ketone at C(6)-C(8), the vinylic and the tertiary methyl both bonded to C-13, as well as the acetoxy groups at C-1 and C-12. The only difference observed by the comparison of the above cited spectra was the downfield shift ( $\Delta\delta$  1.07) of the proton of C-3, appearing as broad singlet at  $\delta$  5.21 while its corresponding carbon resonated at  $\delta$  78.1. The singlet of a further acetyl group was observed at  $\delta$  2.10 (MeCO, *s*)/174.8 (MeCO, *s*) and 20.7 (MeCO, *q*). Thus, **3** appeared to be the 3-*O*-acetyl chenopodolin for which the chemical shifts of all the protons and the corresponding carbons were assigned on the basis of the couplings observed in the COSY, HSQC and HMBC spectra (Table 4) and the NOESY spectrum. In fact in this latter spectrum significant correlations were observed between H-15 with both H<sub>2</sub>-16 and Me-17, H-9 with H-3, and H-12 and H<sub>2</sub>-11 and Me-17. The structure assigned to **3** was supported by the data obtained from its HR ESIMS spectrum, which showed the sodium dimer form [2M + Na]<sup>+</sup> and the sodiated cluster [M + Na]<sup>+</sup> at *m/z* 999 and 511.1953, respectively. When compared to the chenopodolin 1-*O*-acetyl derivative prepared by its usual acetylation with pyridine and acetic anhydride (Cimmino et al., 2013a), **3** showed the same chromatographic behaviour, also

1 when eluted in mixture, by TLC analysis solvent systems A and C) and the same  $^1\text{H}$  NMR data as  
2 previously reported (Cimmino et al., 2013a). Thus, **3** was formulated as  
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4 (1*S*,2*S*,3*S*,4*S*,5*S*,9*R*,10*S*,12*S*,13*S*)-1,3,12-triacetoxy-2,-6-oxo-*ent*-pimara-7(8),15-dien-18-oic acid  
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6 2,18-lactone. This is the first isolation of **3** as a naturally occurring compound. It was not an artefact  
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8 generated by the transesterification of chenopodolin with ethyl acetate during the extraction process  
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10 because chenopodolin remained unaltered when dissolved and left in EtOAc (the solvent used to  
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12 extract the fungal culture filtrates) for 2-3 days at room temperature.  
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16 Assayed at  $4 \times 10^{-3}$  M on punctured detached leaves, **3** proved to be active in particular on  
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18 *Sonchus arvensis*, causing the fast appearance of wide necrosis (scored +++) and on *Urtica dioica*  
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20 and *Parietaria officinalis*, with smaller but still evident necrosis (+). This result differed from the  
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22 not toxic activity previously observed when **3** was tested on different plants (*Mercurialis annua*,  
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24 *Cirsium arvense* and *Setaria viride*) using the same bioassay (Cimmino et al., 2013a). The different  
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26 behaviour could be due to different plant sensitivity. Furthermore, it is also important to consider  
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28 that when **3** crosses the plant cell membrane it could be partially or fully hydrolyzed at the  
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30 physiological pH, according to the well know lethal metabolism (Hassal, 1990), into chenopodolin.  
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32 Compound **1** was less toxic compared to **3**, as it caused medium-size necrosis (++) on *Stellaria*  
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34 *media* and *U. dioica*, whereas was inactive against other plants. Finally, both 9-*O*-acetyl  
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36 chenopodolan D (**4**) and chenisocoumarin (**2**) proved to be inactive on all the tested plants. The  
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38 phytotoxicity observed for **1** was in agreement with the results previously obtained testing  
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40 chenopodolans A-C (Cimmino et al., 2013b) confirming that the nature of the side chain at C- 4,  
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42 and in particular its hydroxylation, is an important feature to impart biological activity. Its 9-*O*  
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44 acetyl derivative probably is inactive because it is not completely hydrolyzed into **1**.  
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53 In conclusion, a new furopyran, named chenopodolan D (**1**), a new isocoumarin, named  
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55 chenisocoumarin (**2**) and a new unrearranged *ent*-pimaradiene, named chenopodolin B (**3**), were  
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57 isolated from *P. chenopodiicola*, a fungal pathogen proposed for the biocontrol of *C. album*. The  
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59 absolute configuration of **1** was established by quantum mechanical calculations of ECD and ORD  
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1 spectra, while that of **2** by applying an advanced Mosher's method. Both **1** and **3** showed  
2 phytotoxic activity. So that it is possible also hypothesised a synergism between them and  
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4 chenopodolin and chenopodolans B and C which contribute to the phytotoxicity showed by the  
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6 culture filtrates of *P. chenopodiicola* and its organic extracts. This could be an important finding in  
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8 the view of a practical application as herbicides of these phytotoxic metabolites in the management  
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10 of *C. album* .  
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### 16 3. EXPERIMENTAL

#### 19 3.1. General

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22 Optical rotations were measured in CHCl<sub>3</sub> on Jasco P-1010 and Jasco DIP-370 digital  
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24 polarimeters; absorption and ECD spectra were recorded at room temperature on a JASCO J815  
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26 spectropolarimeter, using 0.5 mm cells and concentrations of about  $1 \times 10^{-3}$  M. IR spectra were  
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28 recorded as glassy film on a Perkin-Elmer Spectrum One FT-IR Spectrometer and UV spectra were  
29  
30 recorded in MeOH solution on a Perkin-Elmer Lambda 25 UV/Vis spectrophotometer. <sup>1</sup>H and <sup>13</sup>C  
31  
32 NMR spectra were recorded at 400/100 MHz in CDCl<sub>3</sub> on Bruker spectrometers. The same solvent  
33  
34 was used as internal standard. Carbon multiplicities were determined by DEPT spectra (Berger and  
35  
36 Braun, 2004) DEPT, COSY-45, HSQC, HMBC and NOESY experiments (Berger and Braun, 2004)  
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38 were performed using Bruker microprograms. HRESI and ESIMS spectra were recorded on Waters  
39  
40 Micromass Q-TOF Micro and Agilent Technologies 6120 Quadrupole LC/MS instruments,  
41  
42 respectively. Analytical and preparative TLC were performed on silica gel plates (Merck, Kieselgel  
43  
44 60 F<sub>254</sub>, 0.25); or reverse phase (Whatman, Stratocrom RP-18, 0.20 mm) the spots were visualized  
45  
46 by exposure to UV light and/or by spraying first with 10% H<sub>2</sub>SO<sub>4</sub> in MeOH and then with 5%  
47  
48 phosphomolybdic acid in EtOH, followed by heating at 110 °C for 10 min. Column  
49  
50 chromatography was performed on silica gel (Merck, Kieselgel 60, 0.063-0.200 mm). Solvent  
51  
52 systems: (A) CHCl<sub>3</sub>-*i*-PrOH (97:3); (B) EtOH-H<sub>2</sub>O (6:4); (C) CHCl<sub>3</sub>-*i*-PrOH (95:5); (D) *n*-hexane-  
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54 Me<sub>2</sub>CO (7:3); (E) *n*-hexane-Me<sub>2</sub>CO (1:1).  
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### 3.2. Fungal strain.

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2 The fungus was isolated from diseased leaves of *C. album* and identified as *P. chenopodiicola*  
3 (Berestetskyi, unpublished). A monoconidial isolate was deposited in the culture collection of both  
4  
5 the All-Russian Research Institute of Plant Protection, Pushkin, Saint-Petersburg, Russia (I-13.2)  
6  
7 and the Institute of Sciences of Food Production, Bari, Italy (ITEM 12534). The isolate was  
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9 routinely grown and maintained in plates and slants containing potato-dextrose agar (PDA, Sigma-  
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11 Aldrich, Chemie GmbH, Buchs, Switzerland).  
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### 3.3. Production, Extraction and Purification of chenopodolan D, cheniscumarine and 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 *chenopodolin B (1-3).*

The fungus was grown in 1 l Erlenmeyer flasks containing 300 mL of a defined mineral  
medium (Pinkerton and Strobel, 1976) as recently described (Cimmino et al., 2013b). The culture  
filtrates (4.6 l) were lyophilized, dissolved in distilled water (1/10 of its original volume) and then  
extracted by EtOAc (4 x 1 l). The organic extract (780 mg), having high phytotoxic activity, was  
fractionated by CC eluted with the solvent system A, as already described (Cimmino et al., 2013b).  
Six homogeneous fraction groups were collected. The residue (43.3 mg) of the fourth fraction was  
further purified by preparative TLC, eluted with the solvent system A, giving five fractions. The  
residue of the second fraction, obtained as a yellow oil, was identified as cheniscumarin (**2**, *R<sub>f</sub>*  
0.24, 7.8 mg, 1.69 mg/l). The residue (11 mg) of the third fraction was further purified by reverse  
phase TLC (eluent system B) yielding a yellow oil identified as chenopodolan D (**1**, *R<sub>f</sub>* 0.66, 4.5  
mg, 0.98 mg/l). The residue (138.7 mg) of the second fraction of the original column was purified  
by CC eluted with the system C, yielding five homogeneous fractions. The residue (67.6 mg) of the  
second fraction was washed with *n*-hexane followed by CHCl<sub>3</sub> to separated 6-hydroxymellein. The  
residue (4.9 mg) was further purified by preparative TLC (eluent system D) obtaining a white  
amorphous solid, which was identified as chenopodolin B (**3**, *R<sub>f</sub>* 0.41, 4 mg, 0.85 mg/l).

### 3.4. *Chenopodolan D (1)*

Compound **1**:  $[\alpha]_{\text{D}}^{25}$ : -6.8 ( $c = 0.19$ ); IR  $\nu_{\text{max}}$  3324, 1733, 1656, 1622, 1545, 1465, 1354, 1243  $\text{cm}^{-1}$ ; UV  $\lambda_{\text{max}}$  nm ( $\log \epsilon$ ) 337 (3.6), 252 (sh);  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra: see Table 1; HRESI MS (+)  $m/z$ : 539  $[\text{2M} + \text{K}]^+$ , 523  $[\text{2M} + \text{Na}]^+$ , 501  $[\text{2M} + \text{H}]^+$ , 289  $[\text{M} + \text{K}]^+$ , 273  $[\text{M} + \text{Na}]^+$ , 251.1217  $[\text{M} + \text{H}]^+$  (calcd for  $\text{C}_{14}\text{H}_{18}\text{O}_4$  251.1208).

### 3.5. *Chenisocoumarin (2)*

Compound **2**:  $[\alpha]_{\text{D}}^{25}$ : -1.6 ( $c = 0.36$ , MeOH); IR  $\nu_{\text{max}}$  3153, 1625, 1500, 1461, 1374  $\text{cm}^{-1}$ ; UV  $\lambda_{\text{max}}$  nm ( $\log \epsilon$ ) 303 (3.6), 266 (4.1), 229 (sh);  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra: see Table 2; HRESI MS (+)  $m/z$ : 233  $[\text{M} + \text{Na}]^+$ , 211.1218  $[\text{M} + \text{H}]^+$  (calcd for  $\text{C}_{10}\text{H}_{10}\text{O}_5$  211.1208).

### 3.6. *Chenopodolin B (3)*

Compound **3**:  $[\alpha]_{\text{D}}^{25}$ : -55.5 ( $c = 0.20$ ); IR  $\nu_{\text{max}}$  1798, 1719, 1460, 1368, 1269, 1233  $\text{cm}^{-1}$ ; UV  $\lambda_{\text{max}}$  nm ( $\log \epsilon$ ) 277 (3.2), 229 (3.8);  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra: see Table 4; HRESI MS (+)  $m/z$ : 999  $[\text{2M} + \text{Na}]^+$ , 511.1953  $[\text{M} + \text{Na}]^+$  (calcd for  $\text{C}_{26}\text{H}_{32}\text{NaO}_9$  511.1944).

### 3.7. *11-O-acetylchenopodolan D (4)*

Chenopodolan D (**1**, 2.5 mg) was acetylated with pyridine (50  $\mu\text{l}$ ) and  $\text{Ac}_2\text{O}$  (50  $\mu\text{l}$ ) at room temperature for 1 h. The reaction was stopped by the addition of MeOH and the azeotrope, obtained by the addition of benzene, was evaporated by an  $\text{N}_2$  stream. The oily residue (5.0 mg) was purified by preparative TLC eluted with the solvent system E, to give the 11-*O*-acetyl derivative **4** of chenopodolan D as a homogeneous compound ( $R_f$  0.62, 2.4 mg). Derivative **4** had: IR  $\nu_{\text{max}}$  1732, 1689, 1622, 1551, 1459, 1370, 1232  $\text{cm}^{-1}$ ; UV  $\lambda_{\text{max}}$  nm ( $\log \epsilon$ ) 336 (3.4), 253 (3.6);  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra: see Table 1. HRESI MS (+)  $m/z$ : 623  $[\text{2M} + \text{K}]^+$ , 607  $[\text{2M} + \text{Na}]^+$ , 585  $[\text{2M} + \text{H}]^+$ , 331  $[\text{M} + \text{K}]^+$ , 293  $[\text{M} + \text{H}]^+$ .

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2 3.8. 5,7-*O,O'*-dimethylchenisocoumarin (**5**)  
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5 Chenisocoumarin (**2**, 4 mg), dissolved CH<sub>3</sub>OH (2 ml) was added with an ethereal solution of  
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7 diazomethane (200  $\mu$ l). The reaction was carried out overnight at room temperature in the dark. The  
8  
9 reaction was stopped by evaporation under N<sub>2</sub> stream. The residue (7.6 mg) was purified by TLC,  
10  
11 eluted with the solvent system A, yielding the 5,7-*O,O'*-demethyl derivative of **2** (**5**) as an  
12  
13 homogeneous compound (2.6 mg, *Rf* 0.46). Derivative **5** had: IR  $\nu_{\max}$  3406, 1698, 1604, 1560 cm<sup>-1</sup>;  
14  
15 UV  $\lambda_{\max}$  nm (log  $\epsilon$ ) 299 (3.5), 262 (4.2), 230 (sh); <sup>1</sup>H and <sup>13</sup>C NMR spectra: see Table 2. ESIMS  
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17 (+) *m/z*: 515 [2M + K]<sup>+</sup>, 499 [2M + Na]<sup>+</sup>, 277 [M + K]<sup>+</sup>, 261 [M + Na]<sup>+</sup>, 239 [M + H]<sup>+</sup>.  
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24 3.9. 11-*O*-(*S*)- $\alpha$ -Methoxy- $\alpha$ -trifluoromethyl- $\alpha$ -phenylacetate (MTPA) Ester of **5** (**6**)  
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27 (R)-(-)-MPTA-Cl (20  $\mu$ l) was added to **5** (1.0 mg) dissolved in dry pyridine (20  $\mu$ l). The  
28  
29 mixture was kept at room temperature for 1 h and then the reaction stopped by adding MeOH.  
30  
31 Pyridine was removed by a N<sub>2</sub> stream. The residue (2.5 mg) was purified by preparative TLC,  
32  
33 eluted with the solvent system D, yielding **6** as a homogeneous oil (*Rf* 0.71, 1.2 mg). It had: IR  $\nu_{\max}$   
34  
35 1722, 1606, 1581 cm<sup>-1</sup>; UV  $\lambda_{\max}$  nm (log  $\epsilon$ ) 301 (3.9), 261 (4.3), 232 (sh); <sup>1</sup>H NMR spectrum see  
36  
37 Table 3; ESIMS (+) *m/z*: 947 [2M + K]<sup>+</sup>, 931 [2M + Na]<sup>+</sup>, 493 [M + K]<sup>+</sup>, 477 [M + Na]<sup>+</sup>, 455 [M +  
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39 H]<sup>+</sup>.  
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47 3.10. 11-*O*-(*R*)- $\alpha$ -Methoxy- $\alpha$ -trifluoromethyl- $\alpha$ -phenylacetate (MTPA) Ester of **5** (**7**)  
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50 (S)-(+)-MPTA-Cl (20  $\mu$ l) was added to **5** (1.0 mg) dissolved in dry pyridine (20  $\mu$ l). The  
51  
52 reaction was carried out under the same conditions used for preparing **6** from **5**. The purification of  
53  
54 the crude residue (2.2 mg) by preparative TLC eluted with solvent system D, allowed to obtain **7** as  
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56 a homogeneous oil (*Rf* 0.71, 1.2 mg). It had: IR  $\nu_{\max}$  1720, 1605, 1581 cm<sup>-1</sup>; UV  $\lambda_{\max}$  nm (log  
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ε) 303 (3.9), 262 (4.2), 229 (sh); <sup>1</sup>H NMR spectrum see Table 3; ESIMS (+) *m/z*: 947 [2M + K]<sup>+</sup>, 931 [2M + Na]<sup>+</sup>, 493 [M + K]<sup>+</sup>, 477 [M + Na]<sup>+</sup>, 455 [M + H]<sup>+</sup>.

### 3.11. Computational methods

Preliminary conformational analysis was performed by Spartan02 package (SPARTAN 02) employing MMFF94s molecular mechanics (MM) force field with Monte Carlo searching and assuming the (*R*) absolute configuration for **1**. All possible conformers were searched, considering the degrees of freedom of the system within a range of 30 kcal/mol and retaining only the structures in an energy range of 10 kcal/mol with respect to the most stable one. The minimum energy conformers found by MM were further fully optimized by Gaussian09 package (Frisch et al., 2009) using the Density Functional Theory (DFT) at the DFT/B3LYP/TZVP level either in gas phase or in THF and CHCl<sub>3</sub> by IEFPCM solvation model (Tomasi et. al. 2005). All conformers are real minima, no imaginary vibrational frequencies have been found and the free energy values have been calculated and used to get the Boltzmann population of conformers at 298.15 K. The DFT geometries were then employed as input for Time Dependent DFT (TDDFT) calculations of UV, ECD, and ORD spectra both in gas phase and by IEFPCM implicit solvation model in THF and CHCl<sub>3</sub>. Optical rotations were computed by using the TDDFT/B3LYP/aug-cc-pVDZ level on the DFT optimized geometries. The latter were also employed as input geometries for calculation of UV and ECD spectra at the TDDFT/CAM-B3LYP/aug-cc-pVDZ level and taking into account the lowest 30 states. TDDFT calculations employing the long-range corrected CAM-B3LYP functional (Yanai et al., 2004) provided good reproduction of Cotton effects observed in the theoretical ECD spectra (Evidente et. al., 2011). The theoretical ORD, UV, and ECD spectra were obtained as average over the conformers Boltzmann populations. The ECD spectra were obtained from calculated excitation energies and rotational strengths, as a sum of Gaussian functions centered at the wavelength of each transition, with a parameter σ (width of the band at ½ height) of 0.2 eV using SpecDis v1.60 program (Bruhn et al., 2013). The ECD spectrum was also calculated both in

1 the length and velocity formalisms to guarantee origin independence and to evaluate the quality of  
2 the molecular wave functions employed (Moscowitz, 1965). These calculated spectra were almost  
3 coincident, indicating a good level of calculation. Therefore, in all figures only the velocity-form  
4 predicted spectra are reported.  
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### 10 11 3.12. Leaf puncture assay

12 Each metabolite was first dissolved in MeOH and then diluted with distilled water to the  
13 desired concentrations (final concentration of methanol: 2%). The compounds were tested by using  
14 a leaf puncture assay on 6 different plant species, namely *Stellaria media*, *Urtica dioica*, *Sonchus*  
15 *arvensis*, *Parietaria officinalis*, *Lactuca serriola*, *Helianthus annuus*, with the weedy plants (the  
16 first four in the list) collected in local open fields, and the crop plants grown in pots. Pure  
17 compounds were tested at  $4 \times 10^{-3}$  M by applying a droplet (20  $\mu$ l) of solution to detached leaves  
18 previously punctured with a needle. Five replications were used for each metabolite and for each  
19 plant species tested. Leaves were kept in a moistened chamber under continuous fluorescent lights.  
20 Symptoms were estimated visually between three to five days after droplet application, by using a  
21 visual scale from 0 (no symptoms) to 4 (necrosis wider than 1 cm). Control treatments were carried  
22 out by applying droplets not containing the metabolites.  
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46 Biomolecolare del CNR, Pozzuoli, Italy”.  
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### 56 **Appendix A. Supplementary material**

57 Supplementary data associated with this article can be found, in the online version at <http://:.....>  
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**Table 1.**<sup>1</sup>H and <sup>13</sup>C NMR data of chenopodolan D (**1**) and <sup>1</sup>H NMR data of its 11-*O*-acetyl derivative (**4**)<sup>a,b</sup>

Position	<b>1</b>			<b>4</b>
	$\delta\text{C}^c$	$\delta\text{H}$ ( <i>J</i> in Hz)	HMBC	$\delta\text{H}$ ( <i>J</i> in Hz)
2	163.5 <i>s</i>		Me-8	
3	165.9 <i>s</i>		H-7a, OMe, Me-8	
3a	101.9 <i>s</i>		H-7a, Me-8	
4	126.2 <i>s</i>		H-7a, Me-13	
5	135.7 <i>d</i>	6.98 <i>s</i>	Me-13	6.96 <i>s</i>
6	160.1 <i>s</i>		H-5, H-7a, Me-13	
7a	93.7 <i>d</i>	6.18 <i>s</i>		6.19 <i>s</i>
8	14.4 <i>q</i>	1.95 (3H) <i>s</i>		1.93 (3H) <i>s</i>
9	135.9 <i>q</i>		H-10, H <sub>2</sub> -11, Me-12	
10	131.7 <i>d</i>	5.68 <i>br t</i> ( <i>J</i> = 6.6) <sup>d</sup>	H <sub>2</sub> -11	
11	59.8 <i>t</i>	4.32 (2H) <i>br s</i>		4.73 <i>br s</i> 4.71 <i>br s</i>
12	16.5 <i>q</i>	1.85 (3H) <i>s</i> <sup>d</sup>	H-10	1.88 (3H) <i>s</i>
13	13.7 <i>q</i>	2.05 (3H) <i>s</i>	H-5	2.04 (3H) <i>s</i>
OMe	56.1 <i>q</i>	3.92 <i>s</i>		3.92 (3H) <i>s</i>
MeCO				2.08 (3H) <i>s</i>

<sup>a</sup>The chemical shifts are in  $\delta$  values (ppm) from TMS. <sup>b</sup>2D <sup>1</sup>H, <sup>1</sup>H (COSY) <sup>13</sup>C, <sup>1</sup>H (HSQC) NMR experiments delineated the correlations of all the protons and the corresponding carbons. <sup>c</sup>Multiplicities were assigned by DEPT spectrum. <sup>d</sup>Allylic coupling (< 1 Hz) was observed in the COSY spectrum between H-10 and Me-12

**Table 2**<sup>1</sup>H and <sup>13</sup>C NMR data of chenisocoumarin (**2**)<sup>a,b</sup> and its .5,7-*O,O'*-dimethyl derivative (**5**)

Position	<b>2</b>	<b>5</b>		
	$\delta C^c$	$\delta H$ ( <i>J</i> in Hz)	HMBC	$\delta H$ ( <i>J</i> in Hz)
1	169.0 <i>s</i>		H-8, H-3, Me-9	
3	67.2 <i>d</i>	4.57 <i>dq</i> (6.6, 1.8)	H-3, Me-9	4.54 <i>dq</i> (6.6, 1.3)
4	77.5 <i>d</i>	4.38 <i>d</i> (1.8)	Me-9	4.47 <i>d</i> (1.3)
4a	142.5 <i>s</i>		H-4, H-3	
5	155.3 <i>s</i>		H-6, H-4	
6	106.8 <i>d</i>	6.35 <i>d</i> (1.7)	H-8, H-4	6.54 <i>br s</i>
7	164.1 <i>s</i>		H-8	
8	99.2 <i>s</i>	6.33 <i>d</i> (1.7)	H-6, H-4	6.50 <i>br s</i>
8a	135.8 <i>s</i>			
9	15.8 <i>q</i>	1.48 (3H) <i>d</i> (6.6)	H-4, H-3,	1.50 (3H) <i>d</i> (6.6)
OMe				3.92 (3H) <i>s</i>
OMe				3.89 (3H) <i>s</i>

<sup>a</sup>The chemical shifts are in  $\delta$  values (ppm) from TMS. <sup>b</sup>2D <sup>1</sup>H, <sup>1</sup>H (COSY) <sup>13</sup>C, <sup>1</sup>H (HSQC) NMR experiments delineated the correlations of all the protons and the corresponding carbons.

<sup>c</sup>Multiplicities were assigned by DEPT spectrum.

**Table 3**

<sup>1</sup>H NMR data of 4-*O*- (*S*)- and 4-*O*- (*R*)-MTPA esters of cheniscumarine 5,7-*O,O'*-dimetilderivative (**6** and **7**, respectively)

	<b>6</b>	<b>7</b>
	$\delta$ (H ( <i>J</i> in Hz))	$\delta$ (H ( <i>J</i> in Hz))
3	4.702 <i>br q</i> (6.5)	4.661 <i>br q</i> (6.7)
4	5.945 <i>br s</i>	5.907 <i>br s</i>
6	6.637 <i>br s</i>	6.694 <i>br s</i>
8	6.544 <i>br s</i>	6.578 <i>br s</i>
9	1.448 (3H) <i>d</i> (6.5)	1.417 (3H) <i>d</i> (6.7)
OMe <sup>a</sup>	3.920 (3H) <i>s</i>	3.945 (3H) <i>s</i>
OMe <sup>a</sup>	3.870 (3H) <i>s</i>	3.874 (3H) <i>s</i>
OMe	3.430 (3H) <i>s</i>	3.371 (3H) <i>s</i>
Ph	7.381-7.312 <i>m</i>	7.414-7.307 <i>m</i>

<sup>a</sup>These assignments can be reversed

**Table 4.**  $^1\text{H}$  and  $^{13}\text{C}$  NMR Data of chenopodolin B (**3**)<sup>a,b</sup>

Position	$\delta\text{C}^c$	$\delta\text{H}$ (J in Hz)	HMBC
1	72.4 <i>d</i>	5.07 (1H) <i>d</i> (4.9)	Me-20
2	76.9 <i>d</i>	4.62 (1H) <i>d</i> (4.9)	H-1, Me-19
3	78.1 <i>d</i>	5.21 (1H) <i>br s</i>	H-5, H-1
4	46.7 <i>s</i>		H-5, Me-19
5	58.7 <i>d</i>	2.96 (1H) <i>s</i>	H-1, H-3, Me-19
6	194.2 <i>s</i>		H-5
7	126.7 <i>d</i>	5.85 (1H) <i>s</i>	H <sub>2</sub> -14
8	155.7 <i>s</i>		H <sub>2</sub> -14, H <sub>2</sub> -11, H-9
9	48. <i>d</i>	3.00 (1H) <i>m</i>	
10	42.1 <i>s</i>		H-5, H-2, Me-20
11	11.9 <i>t</i>	1.43 (2H) <i>m</i>	Me-20
12	76.8 <i>d</i>	4.71 (1H) <i>t</i>	H-14, H <sub>2</sub> -11, Me-17
13	41.5 <i>s</i>		H <sub>2</sub> -16, H <sub>2</sub> -14, Me-17
14	43.3 <i>t</i>	2.73 (1H) <i>d</i> (16.0) 2.31 (1H) <i>d</i> (16.0)	Me-17
15	139.3 <i>d</i>	5.87 (1H) <i>dd</i> (17.8,11.2)	H-16B, H <sub>2</sub> -14, H-12, Me-17
16	116.8 <i>t</i>	5.19 (1H) <i>d</i> (11.2) 5.03 (1H) <i>d</i> (17.8)	Me-17
17	25.5 <i>q</i>	1.08 (3H) <i>s</i>	H-12
18	175.4 <i>s</i>		H-5, H-3, Me-19
19	12.1 <i>q</i>	1.47 (3H) <i>s</i>	H-5
20	17.3 <i>q</i>	1.12 (3H) <i>s</i>	H-1, H-5
MeCO-12	170.6 <i>s</i>		H-12, MeCO-12
MeCO-3	174.8 <i>s</i>		H-5, H-1, H-2, Me-19
MeCO-1	169.4 <i>s</i>		H-1, MeCO-1
MeCO-1	20.1 <i>q</i>	2.27 (3H) <i>s</i>	
MeCO-3	20.7 <i>q</i>	2.10 (3H) <i>s</i>	
MeCO-12	20.7 <i>q</i>	2.07 (3H) <i>s</i>	

<sup>a</sup>The chemical shifts are in  $\delta$  values (ppm) from TMS. <sup>b</sup>2D  $^1\text{H}$ ,  $^1\text{H}$  (COSY)  $^{13}\text{C}$ ,  $^1\text{H}$  (HSQC) NMR experiments delineated the correlations of all the protons and the corresponding carbons.

<sup>c</sup>Multiplicities were assigned by DEPT spectrum.

## Figure Legend

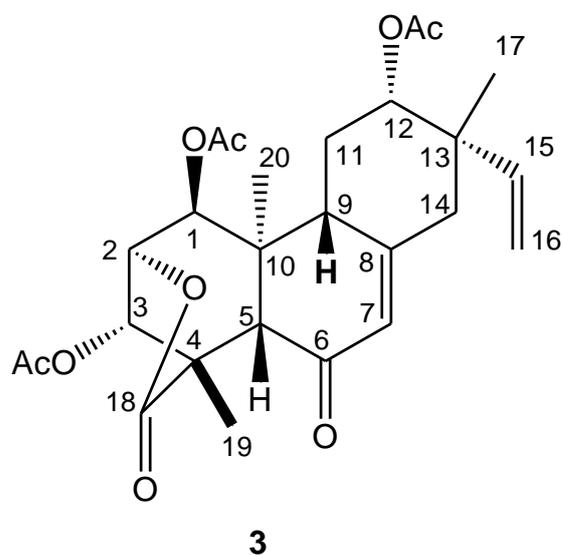
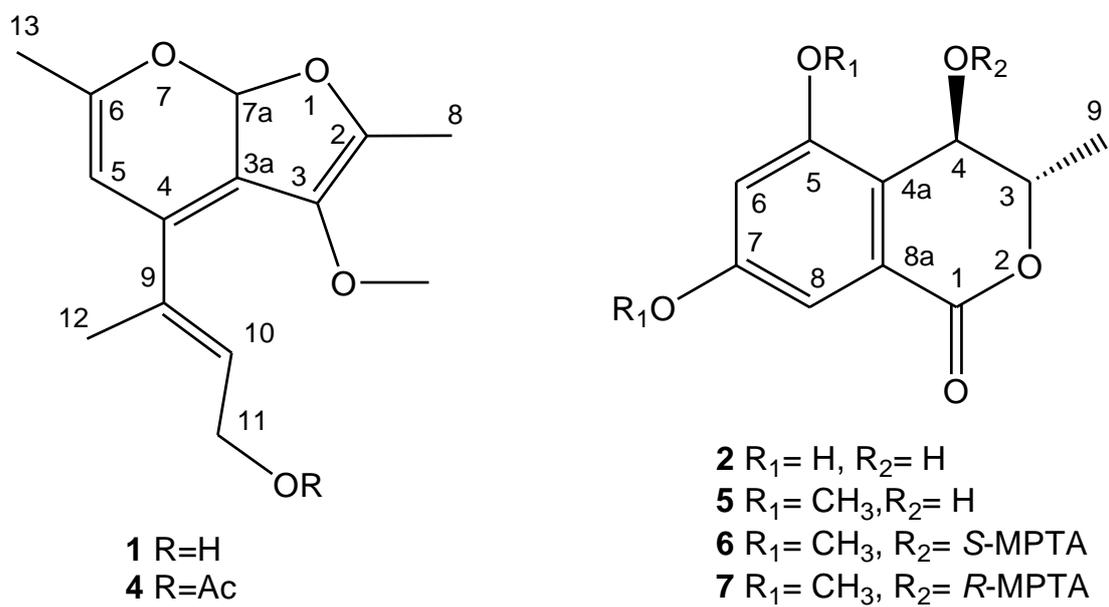
**Fig. 1.** Structures of: chenopodolan D (**1**) chenisocoumarin (**2**) chenopodolin B (**3**), 11-*O*-acetyl-chenopodolan D (**4**), 5,7-*O,O'*-dimetilderivative-, 4-*O-S*-MTPA- and 4-*O-R*-MTPA esters of chenisocoumarin (**5**, **6** and **7**, respectively).

**Fig. 2.** Experimental UV and ECD spectra of (-)-**1** [solid red line, THF] and calculated UV and ECD spectra for (*R*)-**1** [TDDFT/CAM-B3LYP/aug-cc-pVDZ//DFT/B3LYP/TZVP/gas phase dotted green line; TDDFT/CAM-B3LYP/aug-cc-pVDZ//DFT/B3LYP/TZVP/IEFPCM(THF) dashed blue line; 0.2 eV bandwidth]. Calculated UV and ECD spectra are blue shifted of +20.0 nm and reduced 20 times.

**Fig. 3.** Experimental ORD curve of (-)-**1** [solid red line (◆), CHCl<sub>3</sub>] and calculated ORD for (*R*)-**1** [TDDFT/CAM-B3LYP/aug-cc-pVDZ//DFT/B3LYP/TZVP/gas phase, dotted green line (▲); TDDFT/CAM-B3LYP/aug-cc-pVDZ//DFT/B3LYP/TZVP/IEFPCM(CHCl<sub>3</sub>), dashed blue line (●)].

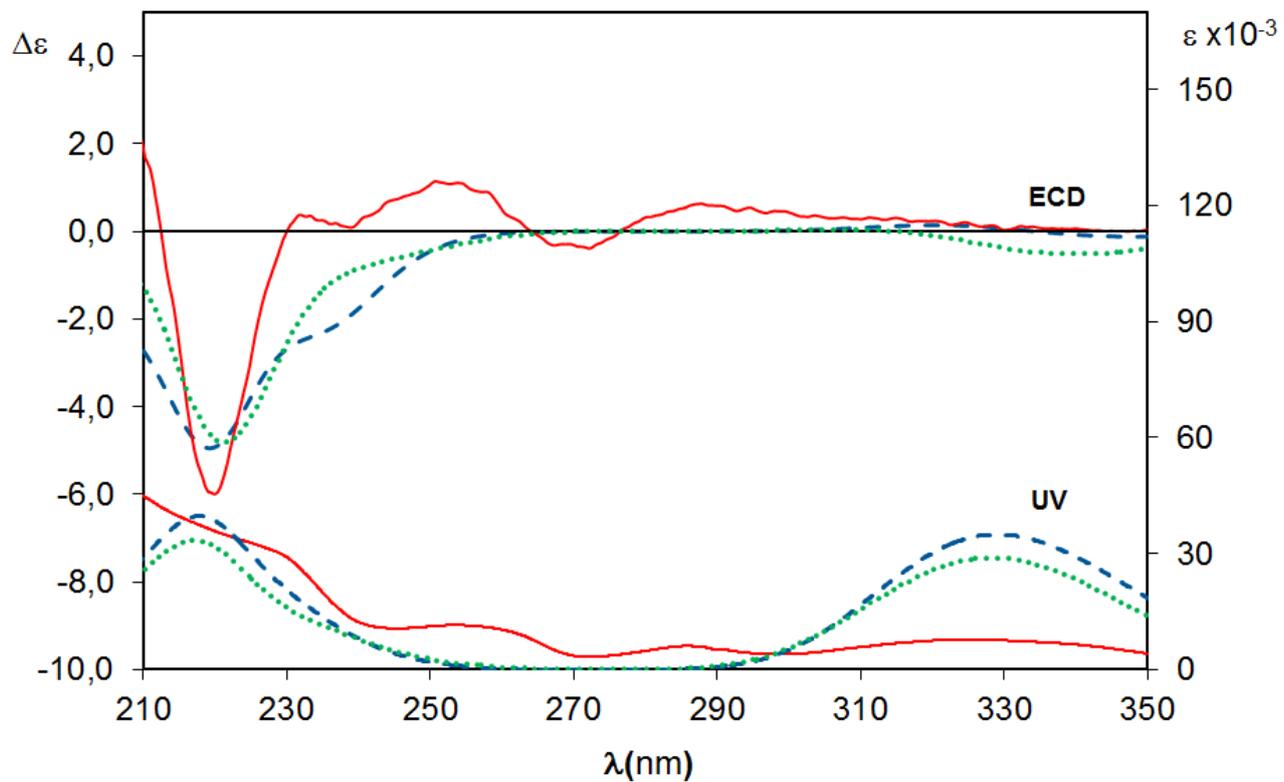
**Fig. 4.** Structures of 4-*O-S*- and 4-*O-R*-MTPA esters of 5,7-*O,O'*-dimethyl chenisocoumarin. (**6** and **7**, respectively), reporting the  $\Delta\delta$  value obtained by comparison (**6-7**) of each proton system.

Figure 1



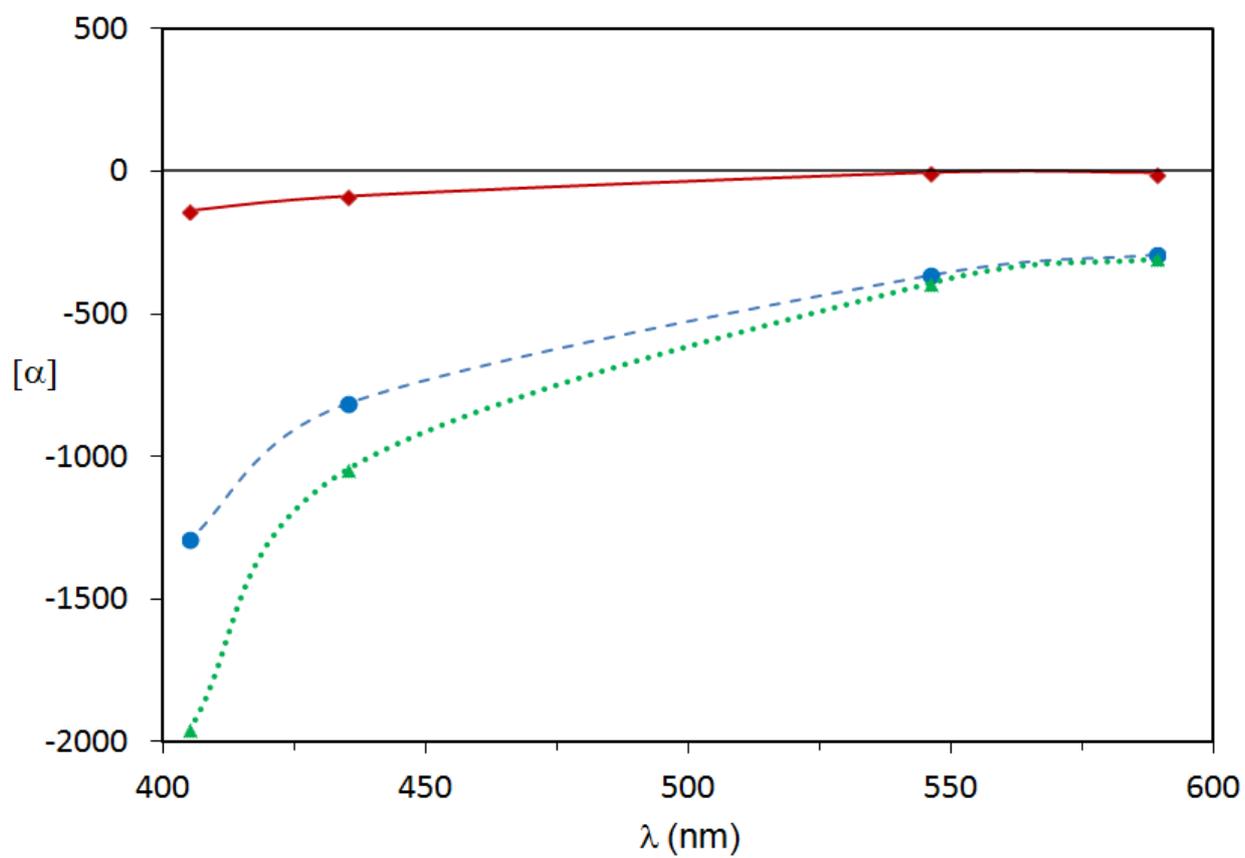
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Figure. 2



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Figure. 3



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Fig. 4

