



# Three new koninginins from *Trichoderma neokongii* 8722



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## ABSTRACT

Three new fungal metabolites, named koninginins I (**1**), J (**2**) and K (**3**) together with four known koninginins A (**4**), B (**5**), D (**6**) and E (**7**), were isolated from solid fermentation products of *Trichoderma neokongii* 8722. Three new structures were elucidated by extensive spectroscopic methods, including 1D NMR and 2D NMR, and HR-ESI-MS experiments.

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## 1. Introduction

*Trichoderma* can produce a lot of secondary metabolites using for biological control (Reino et al., 2008; El-Hasan et al., 2009; Stoppacher et al., 2010; Mukherjee et al., 2012). A new polyketide type compound, koniginin, was obtained from *Trichoderma* sp. (Reino et al., 2008). Koninginins A, B, C, E and G showed a growth inhibition of etiolating wheat coleoptiles (Cutler et al., 1989, 1991, 1999; Parker et al., 1995a, 1995b). Koninginins A, B, D and G also could affect the growth of some plant pathogen fungi (Dunlop et al., 1989; Ghisalberti and Rowland, 1993; Cutler et al., 1999). In order to search for new structure koniginin type compounds against plant diseases, the extract of *Trichoderma neokongii* 8722 was investigated and three new koniginin type compounds, together with four known koninginins were obtained. This report describes three new compounds structures.

## 2. Results and discussion

From the extracts of solid fermentation products of *T. neokongii* 8722, seven compounds including three new koninginins I–K (**1–3**) (Fig. 1) were identified. The structures of the known compounds were determined to be koninginins A (**4**) (Cutler et al., 1989), B (**5**) (Cutler et al., 1991), D (**6**) (Dunlop et al., 1989; Song et al., 2010), and E (**7**) (Parker et al., 1995a).

Compound **1** was obtained as colorless amorphism. The HR-ESI-MS data indicated a molecular formula of C<sub>16</sub>H<sub>24</sub>O<sub>5</sub> based on the [M+H]<sup>+</sup> ion signal at *m/z* 297.1699 (calc. 297.1697). The NMR data (Table 1) revealed one quaternary carbon at  $\delta_C$  199.5, 175.1 and 112.9, five methines at  $\delta_C$  67.1 ( $\delta_H$  4.39), 66.9 ( $\delta_H$  4.27), 78.1 ( $\delta_H$  4.11), 73.4 ( $\delta_H$  3.70) and 68.7 ( $\delta_H$  3.73), and seven methylenes and one methyl, which suggested compound **1** was koniginin type compound (Dunlop et al., 1989; Reino et al., 2008). According to the NMR and MS spectra, compound **1** had one more hydroxyl than koniginin D (Cutler et al., 1991; Song et al., 2010). A preliminary linear skeleton bearing two branches was deduced to be C-2–C-3–C-4 (–branch)–C-7–C-8–C-9–C-10–C-11 (–branch) from complete interpretation of key cross-peaks in the COSY spectrum (H-2/H-3/H-4; H-7/H-8/H-9/H-10/H-11) and key correlations in the HMBC spectrum: H-2 ( $\delta_H$  2.32 and 2.62) correlated with the carbons at  $\delta_C$  199.5 (C-1), 30.6 (C-3), 67.1 (C-4), H-3 ( $\delta_H$  1.97 and 2.18) with the carbons at  $\delta_C$  199.5 (C-1), 34.3 (C-2), 67.1 (C-4) and 175.1 (C-5), H-4 ( $\delta_H$  4.39) with the carbons at  $\delta_C$  175.1 (C-5), 112.9 (C-2), 34.3 (C-2) and 30.6 (C-3), and H-7 ( $\delta_H$  4.27) with the carbons at  $\delta_C$  199.5 (C-1), 175.1 (C-5), 112.9 (C-6), 78.1 (C-9), and 68.7 (C-15) (w), H-9 ( $\delta_H$  4.11) with the carbons at  $\delta_C$  66.9 (C-7) and 28.6 (C-8), H-16 ( $\delta_H$  1.16) with the carbons at  $\delta_C$  40.3 (C-14), 68.7 (C-15), and other correlations (Fig. 2). In the <sup>1</sup>H NMR spectrum, two intermediate coupling constants (*J* = 4.8, 6.6 Hz) were observed for H-4, which confirmed a pseudoequatorial position for H-4. Except for the signals of H-15 and C-15 for **1**, the NMR data were very similar to those of koniginin D (Dunlop et al., 1989; Song et al., 2010). According to a biogenetic perspective, and comparing the specific rotation, chemical shift and coupling constant of **1** with that of the

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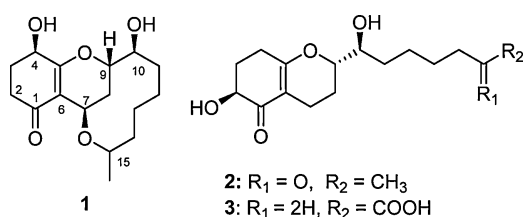


Fig. 1. The structures of compounds 1–3.

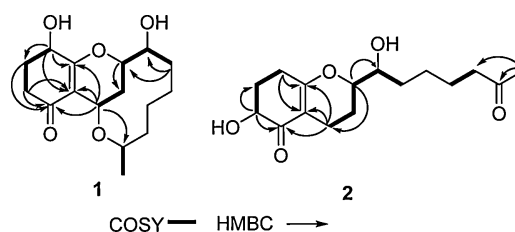


Fig. 2. HMBC and key correlations of compounds 1–2.

literature data (Song et al., 2010), the relative configuration of **1** was same with koniginin D (Dunlop et al., 1989; Song et al., 2010). Based on above data, compound **1** was elucidated to be koniginin I.

Compound **2** was obtained as colorless powder. The HR-ESI-MS data indicated a molecular formula of C<sub>16</sub>H<sub>24</sub>O<sub>5</sub> based on the [M+Na]<sup>+</sup> ion signal at *m/z* 319.1519 (calc. 319.1516). The MS and NMR spectroscopic data of compound **2** were very similar to those of koniginin B except that one methylene (C-15) in koniginin B was oxidized to the ketone in compound **2** (Cutler et al., 1991; Liu and Wang, 2001). The 2D-NMR data (Table 2) showed the detail: H-2 ( $\delta_{\text{H}}$  2.14) correlated with the carbons at  $\delta_{\text{C}}$  209.0 (C-15) and 43.5 (C-14), H-14 ( $\delta_{\text{H}}$  2.46) with the carbons at  $\delta_{\text{C}}$  209.0 (C-15), 30.0 (C-16) and 23.4 (C-13), and other correlations (Fig. 2). In the <sup>1</sup>H NMR spectrum, two intermediate coupling constants (*J* = 6.8, 16.4 Hz) were observed for H-2, which confirmed a axial position for H-2. Except that one methylene (C-15) in koniginin B was oxidized to the ketone in compound **2**, the NMR data were very similar to those of koniginin B (Cutler et al., 1991; Liu and Wang, 2001). From a biogenetic perspective, the configuration of **2** should be identical to that of the co-occurring koniginin B. So, compound **2** was identified to be koniginin J.

Compound **3** was obtained as colorless oil. The HR-ESI-MS data indicated a molecular formula of C<sub>16</sub>H<sub>24</sub>O<sub>6</sub> based on the [M+Na]<sup>+</sup> ion signal at *m/z* 335.1466 (calc. 335.1465). The MS and NMR spectroscopic data of compound **3** were very similar to those of koniginin B except that terminal methyl (CH<sub>3</sub>-16) in koniginin B was oxidized to the carboxyl group in compound **3** (Cutler et al., 1991; Liu and Wang, 2001). The 2D-NMR data (Table 2) showed the correlations between H-14 ( $\delta_{\text{H}}$  1.65) and carbons at  $\delta_{\text{C}}$  178.3 (C-16), 33.8 (C-15) and 29.1 (C-13), H-15 ( $\delta_{\text{H}}$  2.35) and carbons at  $\delta_{\text{C}}$  178.3 (C-16), 33.8 (C-15) and 24.7 (C-14). In the <sup>1</sup>H NMR spectrum, two intermediate coupling constants (*J* = 6.0, 18.0 Hz) were observed for H-2, which confirmed a axial position for H-2. Except that

terminal methyl (CH<sub>3</sub>-16) in koniginin B was oxidized to the carboxyl group in compound **3**, the NMR data were very similar to those of koniginin B (Cutler et al., 1991; Liu and Wang, 2001). From a biogenetic perspective, the configuration of **3** should be identical to that of the co-occurring koniginin B. So, compound **3** was identified to be koniginin K.

Compounds **1**–**7** were assayed for antifungal activity (*Gaeumannomyces graminis*, *Fusarium moniliforme*, *Verticillium cinnabarium* and *Phyricularia oryzae*), but all compounds did not show the inhibition activity to the tested phytopathogenic fungi at 100  $\mu\text{g}/\text{disk}$ . Nematicidal activity result indicated that only koniginin A (**4**) had weak activity against *Panagrellus redivivus* and *Caenorhabditis elegans*.

### 3. Experimental

#### 3.1. General

UV spectra were measured on a Shimadzu UV-2401PC spectrophotometer,  $\lambda_{\text{max}}$  (log  $\epsilon$ ) in nm. NMR experiments were carried out on Bruker AM-400 and Bruker DRX-500 NMR spectrometers with TMS as internal standard. ESI-MS and HR-ESI-MS were recorded on a Finnigan LCQ-Advantage mass spectrometer and a VG Auto-Spec-3000 mass spectrometer. Optical rotations were measured on a Jasco DIP-370 digital polarimeter. Column chromatography was carried out on silica gel (G, 200–300 mesh and H, Qingdao Marine Chemical Factory, Qingdao, China), and Sephadex LH-20 (Pharmacia). Precoated silica gel GF254 plates (Qingdao Marine Chemical Factory, Qingdao, China) were used for thin layer chromatography (TLC).

#### 3.2. Fungal material

The strain of *T. neokongii* 8722 was deposited in Southwest Forestry University. Four pathogenic fungi (*Gaeumannomyces graminis*, *Fusarium moniliforme*, *Verticillium cinnabarium* and *Phyricularia oryzae*) were provided by Dr. Fan L. M. at Yunnan Agricultural University. The culture medium consisted of potato (peeled, 200 g), agar (15 g) and glucose (20 g), per L of deionized H<sub>2</sub>O. *T. neokongii* 8872 (20 L) was cultured on potato-dextrose agar dish at temperature of 26 °C for 21 days.

#### 3.3. Extraction and isolation

Solid fermentation products of *T. neokongii* 8722 (20 L) was cut into small pieces and extracted with mixture solution (EtOAc:MeOH:HAc = 80:15:5, v/v/v) by three times to afford of rude extracts. The extracts were dissolved in water, and extracted with EtOAc and then *n*-butanol three times, respectively.

The EtOAc (31.0 g) residue was subjected to a column of silica gel G (200–300 mesh) using petroleum ether–EtOAc and CHCl<sub>3</sub>–MeOH gradient solvent system to produce 11 fractions (Fr.1–Fr.11). The fraction Fr.5 (320 mg) was subjected to Sephadex LH-20 CHCl<sub>3</sub>–MeOH (1:1) and subsequent purified by preparative TLC to give compound **5** (11 mg). The fraction Fr.8 (1.70 g) was subjected

**Table 1**  
NMR data of compounds **1** (in CD<sub>3</sub>OD, *J* in Hz).

Position	<sup>1</sup> H	<sup>13</sup> C	HMBC
1	–	199.5	–
2	2.32 (1H, ddd, 4.8, 8.4, 16.8) 2.62 (1H, ddd, 4.8, 7.8, 16.8)	34.3	1, 3, 4 1, 3, 4
3	1.97 (1H, m) 2.18 (1H, m)	30.6	1, 2, 4 1, 2, 4
4	4.39 (1H, dd, 4.8, 6.6)	67.1	2, 3, 5, 6
5	–	175.1	–
6	–	112.9	–
7	4.27 (brs)	66.9	1, 5, 6, 9, 15 (w)
8	2.06 (1H, m) 1.57 (1H, m)	28.6	6, 7 9
9	4.11 (1H, m)	78.1	7, 8, 5 (w)
10	3.70 (1H, m)	73.4	11, 12
11	1.66 (2H, m)	33.8	9, 10, 12
12	1.54 (2H, m)	26.9	9
13	1.44 (2H, m)	27.0	12, 14, 15
14	1.44 (2H, m)	40.3	12, 15
15	3.73 (1H, m)	68.7	13
16	1.16 (3H, d, 6.18)	23.7	14, 15

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