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Two new triterpenoid saponins from Ardisia crenata

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Two new triterpenoid saponins, ardisicrenoside K (1) and ardisicrenoside L (2), have been isolated from the roots of *Ardisia crenata* Sims. Their structures have been determined as 3β -O-{ α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)]- α -L-arabinopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)]- α -L-arabinopyranosyl-(1 \rightarrow 4)-[β -D-glucopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 2)-[β -D-glucopyranosyl-(1 \rightarrow 2)-[β -D-glucopyranosyl-(1 \rightarrow 2)-[β -D-glucopyranosyl-(1 \rightarrow 2)-[β -D-glucopyranosyl-(2 \rightarrow 4)-[β -D-glucopyra

Keywords: Ardisia crenata Sims; Triterpenoid saponin; Ardisicrenoside K; Ardisicrenoside L; Anti-fungal

1. Introduction

Previous chemical studies have revealed triterpenoid saponins as the main components of plants of the genus *Ardisia* (Myrsinaceae) [1,2], which have been traditionally used for the treatment of activating blood to eliminate stasis. Recently, more and more structurally novel triterpenoid saponins have been isolated from this genus *Ardisia* [3–10]. They exhibit a wide range of bioactivities: utero-contracting, inhibitory activity on cAMP phosphodiesterase, cytotoxicity, anti-HIV and anti-cancer *etc.* [2].

The roots of *Ardisia crenata* Sims are used as the traditional Chinese medicine "Zhu Sha Gen" for the treatment of respiratory tract infections and menstrual disorders in China [11]. It is a shrub widely distributed in the south of China. The ethanol extract of *Ardisia crenata* Sims also showed good activity against the growth of *Pyricularia oryzae in vitro*, a bioassay system for detecting anti-cancer agents in our preliminary screening. Triterpenoid saponins were detected as the main components in the ethanol extract of *A. crenata* by TLC and color reactions.

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We report here the isolation and structural elucidation of two new triterpenoid saponins, ardisicrenoside K and ardisicrenoside L, obtained from the roots of *A. crenata* Sims, as well as their anti-fungal activity against *Pyricularia oryzae*.

2. Results and discussion

The roots of *A. crenata* Sims were extracted with 60% ethanol. The ethanol extract was then dissolved in water and subsequently partitioned with ethyl acetate and n-butanol. The n-butanol-soluble portion was subjected to column chromatography on Diaion HP-20, MPLC RP-18. The saponin-containing fractions were further chromatographed, by repeated HPLC on an ODS column, to yield two new triterpenoid saponins, ardisicrenoside K (1) and ardisicrenoside L (2).

Ardisicrenoside K (1), white needles, mp 248-250°C gave a positive Liebermann-Burchard reaction and Molish test. The positive HR-ESIMS spectrum revealed a quasimolecular ion peak at m/z 1141.5802 [M + Na]⁺ corresponding to a molecular formula of $C_{55}H_{90}O_{23}$. The IR spectrum of compound 1 exhibits a broad absorption of hydroxyl group at 3417 cm^{-1} , and carbonyl absorption at 1701 cm^{-1} . The ¹H NMR spectrum of compound **1** in pyridine-d₅ exhibits signals of six tertiary methyl groups at δ 1.17, 1.30, 1.09, 1.07, 1.04 and 0.83, a methyl group of rhamnose at δ 1.80 (d, J = 6.0 Hz), two methoxyl groups at δ 3.49 and 3.48, and four anomeric proton signals from sugars at δ 6.40 (brs), 5.38 (d, J = 7.8 Hz). 5.26 (d, J = 7.6 Hz) and 4.95 (brs). The above data suggest that ardisicrenoside K contains four sugars moieties and a triterpenoid aglycone. Complete analysis of the ¹³C NMR and DEPT spectra of compound 1 revealed 55 carbon signals, 32 of which were assigned to the aglycone part, while 23 were assigned to the sugar moiety. ¹³C NMR spectral data of compound 1 are similar to those of the known saponin ardisimamilloside B [4]. As shown in table 1, there is a signal at δ 107.9 (CH, by DEPT) instead of a signal at δ 206.2 due to the 30-CHO group of ardisimamilloside B. Furthermore, in the HMBC spectrum of compound 1, the carbon signal at δ 107.9 not only correlates with Me-29, but also with two methoxyl signals at δ 3.49 (3H, s) and 3.48 (3H, s), confirming that δ 107.9 is the signal of C-30 and the two methoxyl groups are at C-30 (figure 1). The hydroxyl group at C-3 for the aglycone was deduced from the signal at δ 89.1 and its configuration was determined by the NOESY spectrum. The NOE correlations of Hax-3 with H-23 (Me) and H-5 indicate a β configuration for the 3-OH in the aglycone. From the above evidences, the structure of the new sapogenin of compound 1 is established as 3β -hydroxy-16-oxo-13 β , 28-epoxy-30, 30dimethoxyoleanane.

On acid hydrolysis of **1**, arabinose, glucose and rhamnose were identified by co-TLC with authentic samples. The positive-ion ESIMS spectrum exhibits a single predominant peak at m/z 1141, assigned to $[M + Na]^+$. It afforded peaks at m/z 995 $[M + Na - 146]^+$ and 833 $[M + Na - 146 - 162]^+$ in its ESIMS² spectrum. The peak at m/z 833 gave a prominent ion at m/z 671 $[M + Na - 146 - 162 \times 2]^+$ in the ESIMS³ spectrum. In combination with the ESIMS results, it can be concluded that the ratio of arabinose, glucose and rhamnose in **1** is 1:2:1. ¹H $-^1$ H COSY, TOCSY, HMBC and NOESY spectral analysis were used to determine the sequence of the oligosaccharide chain in **1**. From the relatively large H-1 coupling constants (7.8, 7.6 Hz), the anomeric hydroxyl of both glucose moieties should be β -configuration [5]. The small H-1 coupling constants of arabinose and rhamnose, which exhibit a broad anomeric proton singlet in their ¹H NMR spectrum, indicate that they should

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No.	¹³ C	^{I}H	Ardisimamilloside B	No.	¹³ C	^{I}H
1	39.0 t	1.77 (o), 0.89 (o) ^b	39.0	3-O-sugar		
2	26.5 t	1.87 (o), 2.04 (o)	26.5	Ara-1	104.4 d	4.95 (br s)
3	88.9 d	$3.15 (\mathrm{dd}, J = 11.6, 4.0)^{\mathrm{c}}$	89.0	2	80.8 d	4.57 (o)
4	39.6 s		39.6	3	72.4 d	4.47 (o)
5	55.4 d	0.63 (d, $J = 8.3$)	55.5	4	74.6 d	4.58 (o)
5	17.7 t	1.48 (o), 1.36 (o)	17.8	5	63.4 t	4.37 (o), 3.76 (o)
7	33.7 t	1.34 (o), 0.99 (o)	33.8	Glc (terminal)-1	105.4 d	5.38 (d, $J = 7.8$)
8	42.9 s		43.0	2	76.3 d	4.08 (o)
)	50.0 d	1.12 (m)	50.1	3	78.0 d	4.21 (o)
10	36.7 s		36.7	4	71.7 d	4.22 (o)
11	18.8 t	1.48 (o), 1.29 (o)	18.8	5	78.0 d	4.06 (o)
12	31.9 t	2.01 (o), 1.66 (o)	31.6	6	62.9 t	4.39 (o), 4.50 (o)
13	86.3 s		86.2	Glc (inner)-1	103.0 d	5.26 (d, $J = 7.6$)
14	49.8 s		47.9	2	77.2 d	4.28 (o)
15	45.7 t	2.85 (d, $J = 16.1$), 2.02 (m)	45.7	3	79.5 d	4.17 (o)
16	212.6 s		212.6	4	71.8 d	4.13 (o)
17	55.9 s		55.3	5	78.3 d	3.79 (o)
18	53.9 d	2.19 (d, $J = 14.6$)	55.9	6	62.6 t	4.44 (o), 4.28 (o)
19	35.8 t	2.13 (d, $J = 12.4$), 1.38 (o)		Rham-1	101.5 d	6.40 (br s)
20	40.6 s		50.1	2	72.3 d	4.72 (o)
21	31.5 t	1.94 (o), 1.80 (o)	29.6	3	72.7 d	4.67 (dd, J = 9.2, 3.4)
22	24.9 t	2.35 (d, $J = 13.2$), 1.40 (o)	33.8	4	74.8 d	4.27 (o)
23	27.9 q	1.17 (s)	28.0	5	69.3 t	5.06 (o)
24	16.4 q	1.04 (s)	16.4	6	18.9 q	1.80 (d, $J = 6.0$)
25	16.0 q	0.83 (s)	16.1		-	
26	18.7 q	1.30 (s)	18.8			
27	21.7 q	1.09 (s)	21.9			
28	74.9 t	3.94 (d, J = 8.2), 3.59 (m)	74.9			
29	23.9 q	1.07 (s)	23.9			
30	107.9 đ	4.50 (s)	206.2			
30 OCH ₃	58.4 q	3.49 (s)				
30 OCH ₃	58.3 q	3.48 (s)				

Table 1. ¹H and ¹³C NMR data for **1** in C_5D_5N (δ values)^a.

^a Recorded on a Bruker AV 400 (400 MHz for ¹H, 100 MHz for ¹³C) NMR spectrometer.
^b Overlapped signals indicated by (o).
^c Jvalues (in parentheses) are in Hz. The carbon and proton signals were unambiguously assigned through HMQC, HMBC and TOCSY.





Figure 1. Main HMBC correlations for 1.

each possess an α -configuration. Based on these results, and by comparison of the signals from the sugars moieties of **1** in the ¹³C NMR spectrum with the literature [4–6], the four sugars and their anomeric configurations in **1** were determined to be an α -L-arabinopyranose, two β -D-glucopyranoses and an α -L-rhamnopyranose.

The sequence of the oligosaccharide chain was deduced by comparing the chemical shifts of the individual sugar residues with published compounds [12], and confirmed by HMBC and NOESY experiments. The C-1 of arabinose is attached to C-3 of the aglycone, as indicated by the correlation between H-1 (δ 4.95) of arabinose with C-3 (δ 88.9) of the aglycone in the HMBC spectrum, and between H-1 of arabinose and H-3 (δ 3.15) of the aglycone in the NOESY spectrum. An HMBC experiment of compound **1** revealed the following correlations: H-1 (δ 5.38) of the terminal glucose with C-2 (δ 80.8) of arabinose; H-1 (δ 5.26) of the inner glucose with C-4 (δ 74.6) of arabinose; and H-1 (δ 6.42) of rhamnose with C-2 (δ 77.2) of the inner glucose. On the basis of the above analysis, compound **1** was identified as 3β -O-{ α -L-rhamnopyranosyl-($1 \rightarrow 2$)- β -D-glucopyranosyl-($1 \rightarrow 4$)-[β -D-glucopyranosyl-($1 \rightarrow 2$)]- α -L-arabinopyranosyl}-16-oxo-13 β ,28-epoxy-30,30-dimethoxyoleanane, named ardisicrenoside K (figure 2).

Ardisicrenoside L (2), obtained as a white powder, was positive to both the Liebermann– Burchard reaction and the Molish test. The molecular formula $C_{51}H_{84}O_{22}$ was established by analysis of positive HR-ESIMS, which gave a quasi-molecular ion peak at m/z 1071.5396 $[M + Na]^+$. The IR spectrum shows a strong absorption at 3421 cm⁻¹ due to hydroxyl groups and a C—O stretching band at 1041 cm⁻¹.

The ¹H NMR spectrum of compound **2** in pyridine-d₅ displays signals for six tertiary methyl groups at δ 1.58, 1.54, 1.36, 1.21, 1.08 and 0.85, a pair of geminal proton signals at δ 3.43 (*s*), 3.64 (*d*, *J* = 7.41) and four anomeric proton signals at δ 5.47 (*d*, *J* = 7.6 Hz), and 4.79 (*d*, *J* = 5.9 Hz). The above data suggest that **2** contains four sugar moieties and a triterpenoid aglycone. The ¹³C NMR spectrum exhibits 51 carbon signals, 29 of them assigned to the aglycone part and 22 to the sugar moiety. The ¹³C NMR spectral data of **2** are similar to those of the known saponin ardisicrenoside A [6]. As shown in table 2,



Figure 2. Structures of compounds 1 and 2.

a signal at δ 69.7 arising from an oxygenated carbon was observed in the ¹³C NMR of compound **2**, instead of the signals due to the formyl group at C-30 and the quaternary carbon at C-20 of ardisicrenoside A. The existence of the hydroxyl group at C-20 was deduced by the downfield shift at C-19, C-21 and C-29. The oxygenated groups at C-3 and C-16 in the aglycone were deduced from the signals at δ 89.1 and 77.3, respectively. The configuration of oxygenated bond at C-3 and hydroxyl group at C-16 was determined using a NOESY experiment. NOE correlations of Hax-3 (δ 3.17) with H-23 (δ 1.21, Me) and H-5 (δ 0.70), and H-16 (δ 4.28) with H-28 (δ 3.94), indicate a β -configuration for the oxygenated bond at C-3 and α -configuration for 16-OH, respectively. The orientation of 16 α -OH was determined by comparing the chemical shift of C-16 (δ 77.3) with that in the

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No.	¹³ C	^{1}H	Ardisiacripin A	No.	¹³ C	^{l}H	Sugars of ardisiacripin A
1	39.2 t	1.65 (o), 0.75 (o) ^b	38.9 t	3-O-sugar			
2	26.5 t	2.00 (o), 1.86 (o)	26.2 t	Ara-1	104.6 d	4.79 (d, $J = 5.9$)	104.3 d
3	89.1 d	3.17 (0)	88.8 d	2	79.7 d	4.57 (o)	79.3 d
4	39.7 s		39.4 s	3	73.2 d	4.25 (o)	72.7 d
5	55.7 d	0.70 (d, $J = 11.0$) ^c	55.4 d	4	78.5 d	4.23 (o)	78.2 d
6	17.9 t	1.42 (0)	17.6 t	5	64.1 t	4.46 (o), 3.68 (o)	63.8 t
7	37.0 t	2.28 (o), 1.51 (o)	33.9 t	Glc (terminal)-1	104.9 d	5.47 (d, $J = 7.6$)	104.4 d
8	44.7 s		42.2 s	2	76.1 d	4.02 (o)	75.7 d
9	50.5 d	1.33 (o)	50.1 d	3	77.8 d	4.01 (o)	77.6 d
10	36.9 s		36.5 s	4	71.8 d	4.25 (o)	71.3 d
11	19.3 t	1.80 (o), 1.64 (o)	18.8 t	5	77.6 d	4.21 (o)	77.5 d
12	32.8 t	2.15 (o), 1.64 (o)	32.3 t	6	62.9 t	4.55 (o), 4.40 (o)	62.4 t
13	86.4 s		86.1 s	Glc (inner)-1	104.1 d	4.99 (d, $J = 7.9$)	103.8 d
14	42.4 s		44.2 s	2	85.4 d	3.92 (o)	84.9 d
15	34.5 t	1.57 (o), 1.30 (o)	36.3 t	3	77.3 d	4.28 (o)	77.0 d
16	77.3 d	4.28 (o)	76.5 d	4	71.1 d	4.22 (o)	70.5 d
17	44.3 s		43.6 s	5	78.2 d	4.23 (o)	77.8 d
18	50.0 d	2.54 (o)	52.9 d	6	62.3 t	4.43 (o), 4.40 (o)	61.8 t
19	39.5 t	3.00(d, J = 13.9), 1.98 (o)	33.0 t	Xyl-1	107.6 d	4.97 (d, $J = 7.3$)	107.2 d
20	69.7 s		47.9 s	2	76.2 d	4.05 (o)	75.6 d
21	37.4 t	2.81 (d, $J = 13.9$), 2.54 (o)	30.1 t	3	77.3 d	4.28 (o)	77.2 d
22	31.4 t	2.12 (m), 1.99 (o)	31.9 t	4	70.7 d	4.12 (o)	70.2 d
23	28.0 q	1.21 (s)	27.7 q	5	67.4 t	4.56 (o), 3.72 (o)	67.0 d
24	16.6 q	1.08 (s)	16.2 g				
25	16.4 g	0.85 (s)	15.9 g				
26	18.6 g	1.36 (s)	18.1 g				
27	19.6 g	1.58 (s)	19.4 g				
28	77.9 t	3.64(s), 3.43 (d, $J = 7.4$)	77.3 t				
29	32.8 q	1.54 (s)	23.7 q				
30	1		207.3 d				

Table 2. 1 H and 13 C NMR data for compound 2 in C₅D₅N (δ values)^a.

^a Recorded on a Bruker AV 400 (400 MHz for ¹H, 100 MHz for ¹³C) NMR spectrometer.
^b (o) indicates overlapped signals.
^c J values (in parentheses) are in Hz. The carbon and proton signals were unambiguously assigned through HMQC, HMBC and TOCSY.

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literature (16 α -OH, δ 77.0; 16 β -OH, δ 64.0) [5]. From the above evidences, the structure of the new sapogenin was established as 3β ,16 α ,20-trihydroxy-13 β ,28-epoxyoleanane.

The ESIMS spectrum of 2 exhibits a single predominant peak at m/z 1071, assigned to $[M + Na]^+$, which gave ions at m/z 939 $[M + Na - 132]^+$ and 909 $[M + Na - 162]^+$ in its positive-ion ESIMS² spectrum. The ion peaks at m/z 939 and m/z 909 gave the same ion at m/z 777 [M + Na - 132 - 162]⁺ in the MS³ spectrum. The ion peaks at m/z 1047 $[M - H]^{-}$ gave ions at m/z 915 $[M - H - 132]^{-}$ and 735 $[M - H - 132 - 162]^{-}$ in the negative-ion ESIMS² spectrum, and the MS³ spectrum of m/z 735 gave a prominent ion at m/z 591[M – H – 132 – 162 × 2]⁻. On acid hydrolysis, **2** afforded arabinose, glucose and xylose in a ratio of 1:2:1 (analyzed by the same method as 1). By comparing the vicinal coupling constant of anomeric protons with those of published model compounds [5-7], the three sugars were determined to be α -L-arabinopyranose, β -D-glucopyranose and β -Dxylopyranose. Using the same methods as $\mathbf{1}$, the C-1 of arabinose was attached to the C-3 of the aglycone. Long-range coupling correlations were observed between H-1 of xylose and C-2 of the inner glucose, H-1 of the inner glucose and C-4 of arabinose, and H-1 of the terminal glucose and C-2 of the arabinose in the HMBC spectrum (figure 3). Based on the above evidences, the structure of 2 was elucidated as 3β -O-{ β -D-xylopyranosyl- $(1 \rightarrow 2)$ - β -D-glucopyranosyl- $(1 \rightarrow 4)$ - $[\beta$ -D-glucopyranosyl- $(1 \rightarrow 2)$]- α -L-arabinopyranosyl}-13 β ,28-epoxy-16 α ,20-dihydroxyoleanane, named ardisicrenoside L (figure 2).

Ardisicrenosides K and L exhibited weak activity against the plant pathogenic fungus *Pyricularia oryzae* with MMDCs (minimal morphological deformation concentration) of 295 and 320 µM, respectively.



Figure 3. Main HMBC correlations for 2.

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3. Experimental

3.1 General experimental procedures

Melting points were determined on a Yanaco MP-S₃ micromelting point apparatus and are uncorrected. Optical rotations were measured using a P-1020 digital polarimeter (Jasco corporation). ESIMS spectra were obtained using a Bruker esquire 2000 mass spectrometer. IR spectra were recorded on a Shimadzu FT/IR-8400 spectrometer. ¹H and ¹³C NMR, along with 2D NMR spectra, were determined on a Bruker AV-400 (400 MHz for ¹H, 100 MHz for ¹³C) NMR spectrometer, using TMS as an internal standard. Chemical shifts are in δ (ppm) and coupling constants (*J*) are in Hz. TLC was carried out on silica gel 60F₂₅₄ and the spots were visualized by spraying with 10% H₂SO₄ and heating. Diaion HP-20 (Mitsubishi Kasei) and ODS (40–63 µm, Merck) were used for column chromatography. Preparative HPLC was performed using an ODS column (C-18, 250 × 20 mm, Shimadzu Pak; Detector: RID).

3.2 Plant material

The roots of *Ardisia crenata* Sims were collected from Guangxi province of China in 2000, and identified by Professor Qishi Sun. A voucher specimen (YL-2001-113) has been deposited in the Shenyang Pharmaceutical University, Liaoning province of China.

3.3 Extraction and isolation

The air-dried roots of Ardisia crenata Sims (7 kg) were extracted with 60% EtOH (101 \times 2) under reflux. The EtOH extract (500 g) was then suspended in water and subsequently partitioned with ethyl acetate and n-butanol. The n-butanol extract (180 g) was subjected to Diaion HP-20 chromatography using a gradient from H₂O to 95% EtOH. The fraction eluted with 50% EtOH (49 g) was separated by an ODS open column with H₂O-MeOH gradient elution to yield six fractions (1–6). Fraction 4 (2.6 g) was further separated by PHPLC with MeOH–H₂O (3:2) to obtain eight subfractions (41–48). Subfraction 48 (98 mg) was finally purified by PHPLC with MeOH–H₂O (7:3) to afford ardisicrenoside K (15 mg). Ardisicrenoside L (10 mg) was obtained from subfraction 42 (90 mg) by repeated PHPLC with MeOH–H₂O (1:1).

3.3.1 Ardisicrenoside K (1). White needles, mp 248–250°C, $[\alpha]_D^{20} - 17.5$ (c = 0.1, MeOH). IR $\nu_{\text{max}}^{\text{KBr}}$ (cm⁻¹): 3417, 2927, 2869, 1701, 1072, 1045, 891. Positive HR-ESIMS (m/z): $[M + Na]^+$ 1141.5802 (calcd. for C₅₅H₉₀O₂₃Na, 1141.5771). Positive ESIMS (m/z): 1141, 995, 979, 833, 777, 643, 625, 497. Negative ESIMS (m/z): 1153, 1117, 1077, 971, 809, 647, 616. ¹H and ¹³C NMR: see table 1.

3.4 Ardisicrenoside L (2)

White powder, $[\alpha]_D^{20} - 11.4$ (c = 0.1, MeOH). IR ν_{max}^{KBr} (cm⁻¹): 3421, 2933, 2873, 1079, 1041, 887. Positive HR-ESIMS (m/z): $[M + Na]^+$ 1071.5396 (calcd. for C₅₁H₈₄O₂₂Na, 1071.5352). Positive ESIMS (m/z): 1071, 939, 909, 777, 615. Negative ESIMS (m/z): 1047, 915, 753, 591. ¹H and ¹³C NMR: see table 2.

3.5 Acid Hydrolysis of Saponins

Ardisicrenoside K (2 mg) was heated with 6% HCl at 80°C for 10 h. The reaction mixture was then neutralized with 1 M NaOH and filtered. The filtrate was partitioned with CHCl₃. The water layer was concentrated and rhamnose, glucose and arabinose were identified by TLC in comparison with authentic samples.

Ardisicrenoside L (2 mg) was treated with the same way, and xylose, glucose and arabinose were identified by TLC analysis.

3.6 Bioassay

Antifungal assays against the plant pathogenic fungus *Pyricularia oryzae* were carried out as previously reported [13].

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