



# MINOR DAMMARANE SAPONINS FROM PANAX NOTOGINSENG

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**Key Word Index**—*Panax notoginseng*; Araliaceae; roots; South China Ginseng; dammarane-saponins; notoginsenosides R<sub>8</sub> and R<sub>9</sub>.

**Abstract**—Further investigation of the roots of *Panax notoginseng* yielded two new minor dammarane saponins, notoginsenosides  $R_8$  and  $R_9$ , whose structures were established by means of spectral evidence as (20S)-dammar-22-ene-3 $\beta$ , 6 $\alpha$ , 12 $\beta$ , 20, 25-pentol 6-O- $\beta$ -D-glucopyranoside and its (20R)-epimer analogue, respectively.

#### INTRODUCTION

Although Ginseng (roots of Panax ginseng C. A. Meyer) is well known for its tonic value worldwide, South China Ginseng (roots of Panax notoginseng (Burk.) F. H. Chen) which has similar medical effects in traditional Chinese medicine appears to be unrecognized in the West. P. notoginseng originally grew in Yunnan province of China, and is called San-Qi or Tian-Qi locally. Because it is used for many medicinal preparations in China, this plant has been widely cultivated. Chemical investigations on South China Ginseng have been reported by our research group [1–8]. In a continuation of investigations on the saponin composition of this important medicinal plant, we report here the isolation and structural elucidation of two new minor saponins, notoginsenosides  $R_8$  (1) and  $R_9$  (2).

### RESULTS AND DISCUSSION

Notoginsenosides  $R_8$  (1) and  $R_9$  (2) were isolated by repeated column chromatography of the saponin fractions in yields of 0.00011 and 0.00003%, respectively. Comparison of their  $^{13}$ C and  $^{1}$ H NMR signals with those of saponins and related derivatives previously obtained from *P. notoginseng* [2,4] indicated that 1 and 2 were panaxatriol-type dammarane saponins with only one glucose in each saponin. The presence of glucose was also supported by the acid hydrolysis results.

Saponin 1 displayed a quasimolecular ion peak at m/z 653 [M - H]<sup>-</sup> and fragment ions at m/z 635 [653 - H<sub>2</sub>O]<sup>-</sup> and 473 [653 - glucose]<sup>-</sup> in the negative ion FAB-mass spectrum. This information, together with the data from the <sup>13</sup>C NMR spectrum allowed its molecular formula to be assigned as  $C_{36}H_{62}O_{10}$ . Saponin 1 showed

a close resemblance with ginsenoside Rh<sub>1</sub>(3) in their <sup>13</sup>C NMR spectra [2]. The only difference between the two saponins was observed in the side-chain. Saponin 3, which has a side-chain common in ginsenosides [1-7], showed resonances at  $\delta$  126.4 (d) and 130.8 (s) for C-24 and C-25, respectively, while saponin 1 had double-bond carbons resonating at  $\delta$  127.4 (d) and 137.7 (d). There are two possibilities for the location of the double bond—between C-23 and C-24, and between C-22 and C-23. However, the chemical shifts of the two olefinic carbons were quite different from those seen in compounds with a double bond between C-23 and C-24, as commonly occurs in ginsenosides, such as chikusetsusaponin L9a [9]. The coupling system of the side-chain was established as follows. In the <sup>1</sup>H-<sup>13</sup>C COSY spectrum of 1, the carbon signals at  $\delta$  127.4 and  $\delta$  137.7 correlated with the proton signals at  $\delta$  6.25 (ddd, J = 15.9, 8.4, 5.6 Hz) and 6.06 (d, J = 15.9 Hz), respectively. In the  ${}^{1}H-{}^{1}H$  COSY spectrum, the proton signal at  $\delta$  6.25 correlated not only with  $\delta$  6.06 but also with two geminal proton signals at  $\delta$  2.78 (dd, J = 14.0, 5.3 Hz) and 2.39 (dd, J = 14.0, 9.6 Hz). The two geminal protons showed cross-peaks with a carbon signal at  $\delta$  40.2 in the  ${}^{1}H^{-13}C$  COSY spectrum. Up to this point it could be concluded that the double bond might be located between C-22 and C-23, and that the carbon signal at  $\delta$  40.2 arises from C-24 in the side chain. This was confirmed by the COLOC and NOESY experiments. The important correlations are shown in Fig. 1. For example, C-24 showed long-range correlations with two tertiary methyl groups assignable to H-26 and H-27. NOE cross-peaks were observed between H-23 and two methyl groups of H-26 and H-27, and between glycosidic linkage positions, H-6 of the aglycone and the anomeric proton. By comprehensive analyses of all two-dimensional NMR spectra, the carbon and proton signals of 1 were unequivocally assigned as shown in Tables 1 and 2. All spectral data are self-consistent. The structure of notoginsenoside  $R_8$  (1) was thus formulated as (20S)-

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glc: \(\beta\)-p-glucopyranosyl

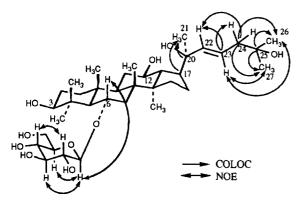


Fig. 1. Important COLOC and NOE correlations of compound

dammar-22-ene-3 $\beta$ ,6 $\alpha$ ,12 $\beta$ ,20,25-pentol 6-O- $\beta$ -D-gluco-pyranoside.

Saponin 2 had the same molecular formula C<sub>36</sub>H<sub>62</sub>O<sub>10</sub> as saponin 1, which was determined by a combination of the information from the FAB-mass spectrum and <sup>13</sup>C NMR spectrum. The <sup>13</sup>C NMR spectra of the two saponins were very similar except for the typical differences at C-17, C-21 and C-24. It is known that the configuration of C-20 may result in a significant difference of the vicinal carbon signal of C-20, such as seen in (20S)- and (20R)-protopanaxadiol (4 and 5) [10]. Table 1 lists the <sup>13</sup>C NMR chemical shifts of compounds 4 and 5. For saponins 1 and 2, the chemical shift variation at the vicinal carbons of C-20 are in accordance with those of compounds 4 and 5 (Table 1), suggesting that saponin 2 should be the (20R)-epimer of 1. In addition, two-dimensional NMR spectra of <sup>1</sup>H-<sup>1</sup>H COSY, <sup>1</sup>H-<sup>13</sup>C COSY, NOESY and COLOC were employed to confirm the proposed structure. The important COLOC and NOE correlations are shown in Fig. 2. Therefore, the structure of notoginsenoside R<sub>9</sub> (2) was established as (20R)-dammar-22-ene- $3\beta$ ,  $6\alpha$ ,  $12\beta$ , 20, 25-pentol 6-O- $\beta$ -D-glucopyranoside.

Table 1. <sup>13</sup>C NMR spectral data of compounds 1-5 in pyridine- $d_{\kappa}(\delta)^*$ 

pyridine- $a_5$ (o)*						
С	1	2	3 [2]	<b>4</b> [10]	5 [10]	
Agly	cone					
1	39.6	39.5	39.5	39.5	39.5	
2	28.0	28.0	28.0	28.2	28.2	
3	78.8	78.7	78.7	77.9	78.0	
4	40.4	40.4	40.4	39.5	39.5	
5	61.6	61.5	61.5	56.3	56.4	
6	80.1	80.1	80.1	18.8	18.7	
7	45.4	45.3	45.3	35.2	35.3	
8	41.3	41.2	41.2	40.0	40.1	
9	50.4	50.3	50.3	50.4	50.6	
10	39.8	39.8	39.8	37.3	37.4	
11	32.3	32.3	32.2	32.0	32.2	
12	71.2	71.0	71.1	70.9	70.8	
13	48.7	49.1	48.4	48.5	49.2	
14	51.6	51.8	51.7	51.6	51.7	
15	31.4	31.4	31.3	31.3	31.5	
16	26.7	26.5	26.9	26.8	26.6	
17	54.2	51.2	54.8	54.7	50.6	
18	17.6	17.5	17.5	16.2	16.3	
19	17.8	17.7	17.7	15.8	15.9	
20	73.4	73.9	73.1	72.9	72.9	
21	27.8	22.5	27.1	26.9	22.7	
22	137.7	138.1	35.9	35.8	43.2	
23	127.4	126.9	23.1	22.9	22.7	
24	40.2	46.4	126.4	126.2	126.0	
25	81.4	81.4	130.8	130.6	130.6	
26	25.4	25.4	25.9	25.8	25.9	
27	25.2	25.3	17.7	17.6	17.7	
28	31.8	31.8	31.8	28.6	28.7	
29	16.5	16.4	16.4	16.4	16.5	
30	17.0	17.1	16.9	17.0	17.3	
Gluc	osyl					
1	106.0	106.0	106.0			
2	75.6	75.5	75.6	_	_	
3	79.6	79.6	79.6		_	
4	72.1	72.1	72.1	_	_	
5	78.1	78.1	78.1			
6	63.3	63.3	63.3		_	

<sup>\*</sup>Assignments of 1 and 2 are based on <sup>1</sup>H-<sup>13</sup>C COSY, <sup>1</sup>H-<sup>1</sup>H COSY, COLOC and NOESY spectra.

Table 2. <sup>1</sup>H NMR spectral data of saponins 1 and 2 in pyridine $d_5$  ( $\delta$ )\*

43 (0)				
¹H	1	2		
Agl	ycone			
1	1.71 (1H, m)	1.70 (1H, m)		
	$1.03\ (1H, m)$	0.98 (1H, m)		
2	1.86 (1H, m)	1.86 (1H, m)		
_	1.42 (1H, br s)	1.42 (1H, br s)		
3	3.53 (1H, dd, 10.8, 5.0)	3.52 (1H, dd, 10.8, 5.2)		
5	$1.44 \ (1H, br \ s)$	$1.43 \ (1H, br \ s)$		
6	4.45 (1H, ddd, 10.2, 10.2, 3.0)	4.43 (1H, ddd, 10.2, 10.2, 3.0)		
7	2.54 (1H, dd, 12.9, 2.7)	2.53 (1H, dd, 13.0, 2.9)		
	1.94 (1H, t, 10.8)	1.96 (1H, t, 10.6)		
9	1.59 (1H, m)	1.61 (1H, m)		
11	2.14 (1H, dd, 9.2, 3.6)	2.13 (1H, dd, 8.8, 3.4)		
	1.51 (1H, d, 4.8)	$1.48 \; (1H, br \; s)$		
12	3.90 (1H, m)	3.91 (1H, m)		
13	2.03 (1H, d, 10.6)	2.01 (1H, m)		
15	1.66 (1H, m)	1.66 (1H, m)		
	1.11 (1H, t, 10.4)	1.15 (1H, t, 10.2)		
16	1.77 (1H, ddd, 12.2, 9.9, 3.0)	1.86†		
	1.35 (1H, d, 8.0)	1.26 (1H, m)		
17	2.30 (1H, dd, 10.8, 7.2)	2.36 (1H, dd, 10.8, 7.0)		
18	1.26 (3H, s)	1.21 (3H, s)		
19	1.06 (3H, s)	1.06 (3H, s)		
21	1.39 (3H, s)	1.40 (3H, s)		
22	6.06 (1H, d, 15.9)	6.05 (1H, d, 15.8)		
23	6.25 (1H, ddd, 15.9, 8.4, 5.6)	6.27 (1H, ddd, 15.8, 8.5, 5.6)		
24	2.78 (1H, dd, 13.9, 5.3)	2.51†		
	2.39 (1H, dd, 14.0, 9.6)	2.43†		
26	1.56 (3H, s)	1.54 (3H, s)		
27	1.56 (3H, s)	1.54 (3H, s)		
28	2.07 (3H, s)	2.06 (3H, s)		
29	1.62 (3H, s)	1.64 (3H, s)		
30	0.80 (3H, s)	0.81 (3H, s)		
Glu	cosyl			
1	5.04 (1H, d, 7.6)	5.03 (1H, d, 7.8)		
2	4.10 (1H, t, 8.8)	4.10 (1H, t, 8.8)		
3	4.27 (1H, t, 9.0)	4.30 (1H, t, 9.4)		
4	4.21 (1H, t, 9.2)	4.23 (1H, t, 9.4)		
5	3.95 (1H, m)	3.96 (1H, m)		
6	4.57 (1H, dd, 11.6, 3.2)	4.54 (1H, dd, 11.6, 3.2)		
	4.37 (1H, dd, 11.6, 6.0)	4.38 (1H, dd, 11.6, 6.0)		

<sup>\*</sup>Assignments are based on <sup>1</sup>H-<sup>13</sup>C COSY, <sup>1</sup>H-<sup>1</sup>H COSY, COLOC and NOESY spectra. Coupling constants (*J* values in Hz) are shown in parentheses.

## **EXPERIMENTAL**

General procedures. NMR spectra were taken in pyridine-d<sub>5</sub> using TMS as int. standard on a Bruker AM-400 spectrometer; <sup>13</sup>C NMR (DEPT) at 100 MHz and <sup>1</sup>H NMR at 400 MHz. 2D-NMR experiments were carried out with standard pulse sequences. FAB-MS: negative ion mode, glycerol matrix, VG Autospec 3000 system, atom source Cs, 35 kV, emitter current 2 μA. Optical rotations were measured on Horiba SEPA-300 polarimeter. For HPLC (Beckman gold system), YMC-

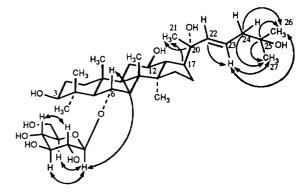


Fig. 2. Important COLOC and NOE correlations of compound **2**.

Pack A 312 ODS column (250 × 16 mm i.d.) was used; solvent: 55% MeOH, flow rate: 4 ml min<sup>-1</sup>; detection: UV at 205 nm. For CC, silica gel H (10–40  $\mu$ , Qingdao) was used. Hydrolysis of saponins with mineral acid and identification of the resulting sugar with TLC were performed as described in ref. [8].

Extraction and separation of saponins. Dried roots of P. notoginseng were extracted and sepd as described in the ref. [8]. The crude saponin fr. (213 g) was chromatographed on silica gel using CHCl<sub>3</sub>-MeOH-H<sub>2</sub>O (50:10:1  $\rightarrow$  20:10:1) to yield frs 1-3 in increasing order of polarity. Fr. 1 was subjected to repeated chromatography on silica gel with CHCl<sub>3</sub>-MeOH (15:1  $\rightarrow$  9:1) to yield a mixture of saponins, which was further purified by prep. HPLC with 55% MeOH to give 1 (55 mg) and 2 (15 mg).

Notoginsenoside  $R_8$  (1). White powder,  $[\alpha]_D^{18} + 29^\circ (\text{MeOH}; c 0.45). FAB-MS (neg.) <math>m/z$ : 653  $[M(C_{36}H_{62}O_{10}) - H]^-$ , 635  $[(M-H) - H_2O]^-$ , 473  $[(M-H-H_2O) - Glc]^-$ . <sup>13</sup>C and <sup>1</sup>H NMR data: Tables 1 and 2.

Notoginsenoside  $R_9$  (2). White powder,  $[\alpha]_D^{18} + 27^\circ$  (MeOH; c=0.32). FAB-MS (neg.) m/z: 653  $[M(C_{36}H_{62}O_{10}) - H]^-$ , 635  $[(M-H)-H_2O]^-$ , 473  $[(M-H-H_2O)-Glc]^-$ . <sup>13</sup>C and <sup>1</sup>H NMR data: Tables 1 and 2.

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<sup>†</sup>Coupling patterns overlap.

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