The Transformation of Ginsenosides by Acid Catalysis in Gastric pH

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Abstract ☐ 1. The ginsenosides of Korean ginseng decomposed profoundly to produce artifact products of prosapogenin A₁, A₂ and A₃ from ginsenoside Rg_1 , prosapogenin C_1 , C_2 and C_3 from ginsenoside Re, and prosapogenin E₁, E₂ and E₃ from ginsenoside Rb₁ by the acid treatment under physiological condition such as 37°C incubation in 0.1 N HCl. 2. The chemical structures of the artifact substances were determined by the analysis of CMR and mass spectra of TMSderivatives as following;

physiological responces, such as the stimulation on protein synthesis¹⁻³⁾, the growth of nerve cells4), the production of red blood cells^{1-3,5)}, and the activities of anti-stress⁶⁾, anti-fatigue⁷⁾, anti-inflammation^{6,8)} central nervous stimulation or depression⁹⁻¹¹⁾, etc.

These activities were mostly observed by the experiments in which ginsenosides were directly applied to the response systems, or in which ginsenosides were parenterally

R=H: ginsenoside Rg₁

R=H: Pro-A2 R=Rham: Pro-Co R=Rham: cinsenoside Re

 $R=H:Pro-A_3$ R=Rham: Pro-C3

Keywords Ginsenosides, Panax gineseng, 13C-NMR spectroscopy, prosapogenins acid-catalyzed transformation

The ginsenosides of Korean ginseng have been shown to elicit a wide variety of the administered to animals. These experimental designs seem to obviate a due consideration on the possible chemical modification of ginsenosides by the interaction with gastric acid in the gastro-intestinal tract.

Present paper deals with the estimation

on what extent of acid catalyzed modification of ginsenosides may take place at the physiological condition of gastric pH, and also with the structure elucidation of the transformed products of ginsenosides.

EXPERIMENTAL

All melting points were taken on a heat block apparatus and given uncorrected values. CMR data were obtained in pyridine- d_5 or DMSO- d_6 solution using TMS as internal standard on JEOL-PET-100 NMR spectrometer (25.15 MHz) and recorded by ppm. unit.

Mass spectra were taken at 75eV on JEOL 01-SG-2 spectrometer as the TMS-derivatives. Densitogram data were obtained by Dual Wave Length TLC-Scanner.

Purification of Ginsenoside Rg₁, Re and Rb₁ Ginsenoside Rg₁, Re and Rb₁ were isolated from Korean ginseng by the combined methods of solvent fractionation and silica gel column chromatography as followings. All chromatographic effluents were monitored by TLC using the authentic ginsenoside standards as references. The butanol soluble fraction of ginseng methanol extracts was chromatographed over silica gel by using the solvent of chloroform/methanol/water

(75 : 25 : 2.5). Ginsenoside Rg₁ was first eluted and then ginsenoside Re as a single spot. Ginsenoside Rg₁ was isolated as deca-acetate (mp 250°C) from methanol after acetylation by usual way, and then it was deacetylated by ordinary saponification. Ginsenoside Re was purified by repeated

crystallization from water (mp 190-2°C). Ginsenoside Rb₁ was isolated by silica gel column chromatography using the solvent of chloroform/methanol/water (70 : 30 : 4) as eluent. The isolated ginsenosided Rb₁ was crystallized from water (mp 205-9°C).

Time course acid treatment on ginsenosides at gastric pH

Ginsenosides Rg₁, Re and Rb₁ were dissolved in water and then mixed with N-HCl to make the final concentration of 0.5-2.5 mg ginsenosides/ml and 0.1N HCl. respectively. The resulting solutions were incubated at 37°C for the definite time length given in Table I. An aliquot of the incubation mixtures was neutralized by adding 5% Na₂CO₃ solution, and then chromatographed over a silica gel plate by using the solvent of chlorofolm/methanol/water (75: 25: 2.5). The plate was sprayed with d-H₂SO₄ solution and heated in 105°C oven to maximum color development. The color densitogram data were obtained by Shimadzu dual wave length TLC-scanner(Emax 522nm, reference 730nm).

Isolation of Acid-catalized Decomposition Products(prosapogenins)

Ginsenosides Rg₁ (0.9 g) and Re (3 g) were dissolved in water, respectively, and added with 1N-HCl to make the final concentration of 2.5 mg ginsenosides/ml and 0.1N-HCl. The reaction mixtures were incubated at 37°C for 2 hours and then neutralized by the addition of 10% Na₂CO₃ solution. The reaction products were extracted with butanol and concentrated to dryness. The residues were chromatographed

over silica gel column using the eluent of chlorofolm/methanol/water(75: 25: 2.5), and prosapogenins were obtained in a chromatographically pure state by ordinary treatment.

RESULTS AND DISCUSSION

Time Course Rate of Degradation of Ginsenosides.

Ginsenoside Rg₁, Re and Rb₁ were chosen as the representative components of ginsenosides present in Korean ginseng and treated with 0.1N HCL at 37°C for various time intervals. On this treatment, every ginsenosides produced three additional TLC-spots as shown in the thin layer chromatograms of Fig. 1.

They were tentatively designated as prosapogenin A_1 , A_2 , and A_3 (from Rg_1), C_1 , C_2 , and C_3 (from Re), E_1 , E_2 , and E_3

solvent A	solvent B			
A ₁ 0 A ₂ 6 A ₃ 0 Rg1 •	C ₁ ° ° E ₁ C ₂ ° E ₂ C ₃ ° E ₃			
	0 Rb1			
——X	 			

Fig. 1: Thin layer chromatograms of the acid treatment products of dammarene glycosides; ginsenoside Rg₁, Re and Rb₁.

TLC solvent A: CHCl₃/MeOH/H₂O=70: 20:2

B: $CHCl_3/MeOH/H_2O = 70:25:2.5$

Table I: Densitogram of time course acid-hydrolysates of ginsenoside Rg1, Re and Rb1.

Incub. Time (Min) 0 spots		30	45	60	90	120
100%	65.5	37.9	28.5	18.7	14.6	10.6
0	0.7	1.51	3.57	9.3	15.8	28.1
O	24.6	45.4	50.0	53.1	48.8	51.3
0	9.9	15.1	17.9	18.7	20.7	10.0
100%	85.9	75.8	65.6	47.2	38.5	44.5
0	0	1.7	4.5	7.5	12.4	14.8
o	12.5	19.0	23.9	36.7	40.0	33.6
0	1.6	3.5	6.0	8.6	9.1	7.1
100%	65.0	48.3	36.4	23.5	17.1	13.1
0	0	3.5	7.2	7.8	11.4	19.0
0	24.1	35.1	40.0	51.0	54.4	52.4
0	10.9	13.1	16.4	17.9	17.1	15.5
	100% 0 0 100% 0 0 100% 0 0 100% 0 0 0 0	100% 65.5 0 0.7 O 24.6 0 9.9 100% 85.9 0 0 12.5 0 1.6 100% 65.0 0 0 24.1	ne in) 0 15 30 100% 65.5 37.9 0 0.7 1.51 O 24.6 45.4 0 9.9 15.1 100% 85.9 75.8 0 0 1.7 o 12.5 19.0 0 1.6 3.5 100% 65.0 48.3 0 0 3.5 0 24.1 35.1	ne in) 0 15 30 45 100% 65.5 37.9 28.5 0 0.7 1.51 3.57 O 24.6 45.4 50.0 0 9.9 15.1 17.9 100% 85.9 75.8 65.6 0 0 1.7 4.5 0 12.5 19.0 23.9 0 1.6 3.5 6.0 100% 65.0 48.3 36.4 0 0 3.5 7.2 0 24.1 35.1 40.0	ne in) 0 15 30 45 60 100% 65.5 37.9 28.5 18.7 0 0.7 1.51 3.57 9.3 O 24.6 45.4 50.0 53.1 0 9.9 15.1 17.9 18.7 100% 85.9 75.8 65.6 47.2 0 0 1.7 4.5 7.5 0 12.5 19.0 23.9 36.7 0 1.6 3.5 6.0 8.6 100% 65.0 48.3 36.4 23.5 0 0 3.5 7.2 7.8 0 24.1 35.1 40.0 51.0	ne ne 100% 65.5 37.9 28.5 18.7 14.6 0 0.7 1.51 3.57 9.3 15.8 O 24.6 45.4 50.0 53.1 48.8 0 9.9 15.1 17.9 18.7 20.7 100% 85.9 75.8 65.6 47.2 38.5 0 0 1.7 4.5 7.5 12.4 0 12.5 19.0 23.9 36.7 40.0 0 1.6 3.5 6.0 8.6 9.1 100% 65.0 48.3 36.4 23.5 17.1 0 0 3.5 7.2 7.8 11.4 0 24.1 35.1 40.0 51.0 54.4

(from Rb₁) in the order of increasing polarity. The time course rates of the prosapogenin production were monitored by TLC-densitometry using a dual wave length TLC-scanner and the results were illustrated in Table I.

As shown in Table I, the decomposition of ginsenoside Rg₁ took place more than 34% at 15min., 62% at 30 min, and 89.4% at 120 min. Similar tendencies were observed in the data of ginsenoside Re and Rb₁. These results suggest that a significant decomposition of ginsenosides should take place in the gastro-intestinal tract before any appreciable absorption of genuine ginsenosides proceeds.

Isolation of Prosapogenins

We attempted to isolate the prosapogenins produced by the acid catalysis under physiological condition and to determine their chemical structures.

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31.2c)

Ginsenoside Rg₁ (0.9g) and Re (3g) were treated with 0.1N HCl at 37°C for 120 min. and applied to silica gel column to obtain prosapogenin A₁ (Pro-A₁) 100mg, Pro-A₂, 300mg, and Pro-A₃ 190mg (from Rg₁) and $Pro-C_1$ 455mg, $Pro-C_2$ 851mg and $Pro-C_3$ 540mg (from Re) in a chromatographically pure state. Upon concentration of the eluate, Pro-A₁ and Pro-C₁ were further decomposed to yield very complicated mixture of spots on TLC, therefore they could not be subjected to the structure determination.

The Structure of Prosapogenin A2 The CMR spectra of Pro-A2 were assigned by referring to the already assigned CMR spectra of ginsenoside Rh₁⁽²⁾ and tabulated in Table II. As shown in Table II, all the CMR peaks of Pro-A₂ are superimposable with those of ginsenoside Rh₁, except that the CMR spectra of Pro-A2 show additional peaks in the neighbour of the peaks of C_{13-23} . S. Yahara et. al^{12} isolated ginsenoside Rh₁ from the root of Panax ginseng, fully assigned the peaks in CMR spectra of the substance, and demonstrated its chemical structure as 20S-protopanaxatriol-6-0-glucoside. On the other hand, J. Asakawa et al¹³⁾ studied C₂₀ chirality inducd chemical shift changes of ginseng sapogenins. Referring to these data, Pro-A2 was shown to be the mixture of ginsenoside Rh₁ and its C_{20} -**R**-epimer.

The Structure of Prosapogenin A₃

The structure of Pro-A₃ could envisaged by the comparison of CMR spectra of Pro-A3 with those of Pro-A2. Every peaks in the spectra of Pro-A₃ is superimposable with those of Pro-A2 except the peaks of C_{24-27} . The olefinic carbon peaks at 126.2 ppm and 130.6 ppm observed in the CMR spectra of Pro-A2 were replaced in the spectra of Pro-A₃ by the tertiary alcoholic carbon peak at 69.8 ppm. This suggests the presence of additional tertiary alcohol group in carbon-25 of Pro-A₃ produced by the hydration at the olefinic bond of Pro-A2. Identification of the C-25 hydroxyl group in Pro-A₃ is also possible by comparison of

Ginseno Ginseno Pro-A3 Pro-C3 Pro-C2 Pro-A2 Pro-A2 Pro-A3 Comp. side Rh1 side Re (DMSO- d_6) (C_5D_5N) (C_5D_5N) (DMSO- d_6) (DMSO- d_6) (DMSO- d_6) Carbon No. (C_5D_5N) (DMSO-d₆) 38.8 39.4 38.5 38.8 1 38.9 39.5 39.5 38.8 26.8a) 28.8 26.6a) 27.9 26.7a) 2 27.6 27.7 26.9a) 77.6^{d)} 73.6 77.7c) 79.9d) 78.5⁽³⁾ 3 76.6d) 79.9°) 77.7_{p} 40.3 38.5 40.1 40.2 39.0 38.8b 38.8 4 39.1 60.0 61.4 59.7 5 60.4 61.3 61.3 61.3 60.3 73.2 77.5^d78.5€ 78.5^{d} 77.3d) 73.2 73.2 78.0 6 45.5 44.3 44.7 44.8 45.2 44.5 7 44.2 45.0 40.3 41.1 40.2 41.0 41.0 40.2 40.3 8 40.3 50.1 49.2 48.9 48.9 50.2 48.5 9 49.3 50.1 39.6 38.5 39.5 38.6b) 38.3 10 38.9 39.5 38.8

31.1b)

31.8c)

31.5a)

Table II: 13C-NMR Chemical shifts.

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31.1

31.1

3.20

30.1b)

		Han	Sitimation	or Ginsen	osides by Aci	u Catalysis		29
12	69.9	70.9	70.9	69.8	69.8	69.8	71.0	69.0
<u>및</u> 12	47.4	48.0	48.1	47.3	47.3	47.2		
gn	(48.0)	(48.6)	(48.7)	(47.9)	(47.9)	(47.8)	48.2	48.1
رم ن 14	51.5	51.5	51.5	58.9	51.0	51.0	51.6	58.7
ទី 15	30.9°)	31.2	31.2°	30.7ы	31.1	31.1	31.1	31.0b)
Aglycone Sugar 12 12 12 12 12 12 12 12 12 12 12 12 12	25.9*)	26.8	27.0	26.7ª)	26.0	26.0 ^{a)}	27.2	25.7a)
17	53.6	54.5	54.5	53.5	53.5	53.5	54.7	50.7
	(49.7)	(50.4)	(50.5	(40.6)	(49.5)	(49.5)		
18	17.6b)	17.6b)	17.6a)	17.0°	17.4a)	16.9 ^{b)}	17.6a)	16.7
19	17.0ы	17.3ы	17.3a)	17.0°)	16.9ª)	16.5 ^{b)}	17.6a)	16.7
20	72.1	73.0	73.3	72.3	72.0	72.3	73.0	82.1
21	26.6	26.8	26.8	26.0	26.4	26.6	26.8	23.6
	(21.6)	(22.5)	(22.5)	(22.1)	(21.6)	(22.1)		
22	35.0	35.6	36.3	35.6	34.9	35.6	35.8	35.2
	(42.2)	(42.2)	(42.9)	(43.5)	(43.0)	(42.2)	(42.9)	
23	22.0	22.5	19.0	18.1	21.9	18.1	23.0	22.1
	(21.6)	()	(18.6)	(17.6)	()	(17.3)		
24	125.6	126.2.	45.3	44.1	125.5	44.3	126.3	125.2
	(125.4)	(125.9)	(45.5)	(44'6)	129.9	63.9	130.6	130.0
	(125.4)	(125.9)	(45.5'	(44.6)	(125.2)	()		
25	130.0	130.6	69.8	68.9	129.9	68.9	130.6	130.0
26	25.5	17.0 ^{b)}	29.7ы	29.3	25.4	29.3	25.3	25.3
27	17.0 ^{b)}	25.8	30.0ы	29.3	16.5a)	29.3	17.4a)	16.7
28	30.9	31.5	31.5	30.7	31.1	31.1	31.7	30.1
29	16.7 ^{b)}	16.7b)	16.7a)	16.7°	16.5a)	16.5 ^{b)}	16.4a)	16.7
30	15.5b)	16.2b)	10.2a)	15.4c)	16.5 ^{a)}	16.5b)	16.8a)	16.7
Glu-1	104.4	105.7	105.7	104.3	100.0c)	99.9	105.9	99.8
2	74.3	75.1	75.1	74.1	77.0 ^{d)}	77.0 ^{d)}	75.4	77.7°)
3	76.6 ^{d)}	77.8c)	77.8 ^{d)}	76.5^{d}	77.3 ^{d)}	77.4 ^{d)}	79.5b)	77.7°)
4	70.6	71.6	71.6	70.5	71.0	71.1	71.3	70.9
	77.9 ^{d)}	79.2c)	$79.3^{d)}$	77.84)	77.5 ^{d)}	77.6 ^{d)}	80.0b)	77.2°)
6	61.6	62.8	62.9	61.4	61.7	61.7	63.1	61.6
Rhom-1					99.8c)	99.9		99.8
2					70.5	70.6		70.3
3					70.5	70.6		70.3
4					72.2	72.3		72.1
5					86.0	78.0		67.9
6					17.7	17.7		17.7
C-20-qlu	icose							
C-1								96.7
2								73.2
3								76.9°)
4								70.3
5								76.4°)
6								61.3

Fig. 2 Chemical shifts of C₂₃-C₂₇ of betulafolianetriol.¹³⁾

the chemical shifts of the neighbouring carbons in Pro-A₃ with thoe of betulafolianetriol, whose CMR spectra were fully assigned by J. Asagawa *et al.*¹³⁾

All CMR assignments of Pro-A₃ are also supported by the mass spectra of the TMS-derivative of the compound. As shown in Fig. 3, mass spectra of TMS-Pro-A₃ give

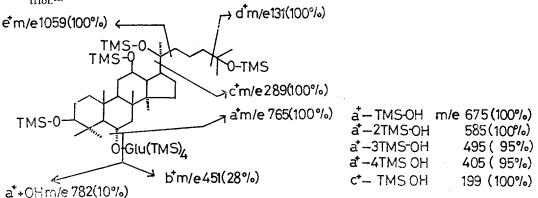
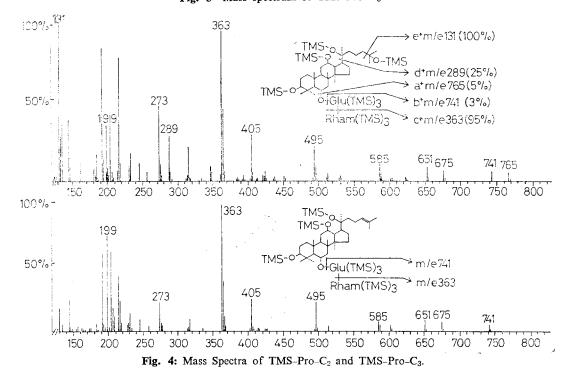


Fig. 3 Mass spectrum of TMS-Pro-A₃.



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the fragment ions of d+ m/e 131 (TMS-isopropanol) and C+ m/e 289 which suggest two TMS-groups in the side chain of TMS-Pro-A₃. Considering also the additional CMR peaks in the neighbour of C_{13-23} , Pro-A₃ was identified as the mixture of C-25-hydroxy derivative and its C-20R epimer produced by the hydration of Pro-A₂ on C_{24-25} olefinic bond.

The Structure of Prosapogenin C_2 and C_3 The structure of Pro- C_2 was determined by comparison of CMR specta of Pro- C_2 and Pro- C_2 . The chemical shifts of side chain carbons are same in both Pro- C_2 and Pro- C_2 . However, those of sugar carbon peaks of Pro- C_2 are almost same with those of ginsenoside Re. The mass spectra of TMS-Pro- C_2 give the fragment ions of TMS-glucosyl rhamnose C_2 m/e 363 (Fig. 4)

Considering the above results, the structure of Pro-C₂ corresponds to ginsenoside Rg₂ and its C-20 R epimer which were produced by the decomposition of C-20 glycoside bond of ginsenoside Re.

The structure of Pro-C₃ was formulated as the C-25-hydroxy derivative of Pro-C₂ by the comparison of the CMR spectra and the mass spectra of Pro-C₃ with those of Pro-C₃. Similar discussions are possible in case of the relation between Pro-A₂ and Pro-A₃. The mass spectra of TMS-Pro-C₃ give fragment ions e⁺ m/e 131 and d⁺ m/e 289 as base peak, suggesting the presence of two TMS-groups in the side chain of Pro-C₃, and fragment ions b⁺ m/e 741 and C⁺ m/e 363 as prominent peaks, suggesting also the

presence of the TMS-rhamno-glucosyl group in Pro-C₃. The fragement ions of e⁺ and d⁺ seem to have a diagnostic value for the structure identification of Pro-C₃ and Pro-A₃ since Pro-C₂ doesn't give those paeaks.

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