THE CONFIGURATION AND CONFORMATION OF THE ARABINOSE MOIETY IN PLATYCODINS, SAPONINS ISOLATED FROM PLATYCODON GRANDIFLORUM, AND MI-SAPONINS FROM MADHUCA LONGIFOLIA BASED ON CARBON-13

AND HYDROGEN-1 NMR SPECTROSCOPIC EVIDENCE: THE TOTAL STRUCTURES OF THE SAPONINS

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Abstract. ^{13}C and ^{1}H FT NMR spectroscopy provided confirmatory evidence for the anomeric α configuration and the predominance of the $^{1}\text{C}_{4}$ conformation of the L-arabinopyranose moiety in all saponins hitherto isolated from the title plants.

Shibata and coworkers first isolated platycodin-D ($\underline{1}$) from the root of <u>Platycodon grandiflorum</u> A. De C<u>andolle</u> (Japanese name, Kikyo) and determined its structure except for the anomeric configuration of the L-arabinopyranose (Ara) moiety. Recently, Shoji and coworkers determined the configuration to be α , by measuring the ${}^{1}J(C-1,H-1)(Ara)$ value of $\underline{1}$ to be 168 Hz in its ${}^{1}H$ non-decoupled 25-MHz ${}^{13}C$ FT NMR spectrum in pyridine-d₅ (C_5D_5N). However, the Shionogi group suggested the β -configuration, tentatively assigning all 15-MHz ${}^{13}C$ NMR signals of $\underline{1}$, platycodin-D₂ ($\underline{2}$) and -D₃ ($\underline{3}$), deapioplatycodin-D ($\underline{4}$) and -D₃ ($\underline{5}$), polygalacin-D ($\underline{6}$) and -D₂ ($\underline{7}$), platyconic acid-A ($\underline{8}$), and several mono-O-acetyl derivatives in their L-rhamnopyranose (Rha) moiety, because the Ara carbon signals indicated the β -L nature as the C-5 signals appear at relatively higher fields (see the TABLE) when the Ara-ring assumes the ${}^{4}C_1$ conformation.

On the other hand, the Osaka University group⁵ elucidated the structures of Mi-saponin-A (9) and -B (10) isolated from the seed kernels of Madhuca longifolia (L.) Macbride; the anomeric α configuration of the L-Ara moiety was determined by the 3 J(H-1,H-2)(Ara) value of 5.5 Hz obtained from the anomeric 1 H doublet signal (100 MHz) of an enzymatic hydrolysis product (16) and its hexa-O-acetyl derivative (16a).

In order to solve the controversial problem of the anomeric configuration, we investigated the 13 C and 1 H FT NMR spectra of platycodins (1-4), Mi-saponins (9 and 10), crude-hesperidinase hydrolysis products from platycodins (11 and 13), those from Mi-saponins (14 and 16), takadiastase hydrolysis products from 11 and 14 (12 and 15, respectively), and their peracetates (1a, 9a, and 11a-16a). We report here confirmatory evidence for the configuration and solution conformations of the Ara-rings in these saponins.

Previous tentative ^{13}C signal assignments $^{3\,14}$ for the C-28 side-chain sugar carbons were first re-examined using the complete ^{1}H -decoupled 25-MHz ^{13}C FT and PRFT spectra of $\underline{1}$ - $\underline{4}$ in $\text{C}_5\text{D}_5\text{N}$ at 100°C . The inversion recovery rates of the sugar CH signals were found to be in the order of Ara < Rha < D-xylopyranose (Xyl) [< D-apiofuranose (Api)] in $\underline{1}$ - $\underline{3}$ [$\underline{4}$], and the same orders were also found for their ring-CH₂ signals. These observations are consistent with the

orders of magnitudes of T_1 values of the carbon signals expected from the sugar sequence.⁷ Thus, the tentative ¹³C signal assignments for 1-4 previously reported³⁺⁴ were shown to be almost valid. In a similar manner, the ¹³C signals of the sugar side-chain in 9 and 10 were assigned in comparison with those of 1-4 and with the aid of the order of the inversion recovery rates of the CH and ring-CH₂ signals; <u>i.e.</u>, Ara < Rha < Xyl < (Api) < Rha. The chemical shift $\delta_{\rm C}$ data for the Ara moieties are listed in the TABLE. Thus, ¹³C signal assignments of sugar moieties of 11, 12, 14, and 15 were straightforward, and the $\delta_{\rm C}$ values for Ara are also listed in the TABLE. In addition, ³J(H-1,H-2)(Ara) values observed (100 MHz) were about 2-3 Hz for the compounds mentioned here (see the TABLE). Therefore, in all the cases examined, the Ara seemed to be β -L rather than α -L, when the Ara-rings adopt the ⁴C₁ conformation.

Next, we examined ^1H non-decoupled ^{13}C spectra of the above-mentioned compounds in C_5D_5N at $100\,^{\circ}\text{C}$, from which $^1\text{J}(\text{C-1,H-1})(\text{Ara})$ values were obtained (see the TABLE). These values were about 171-173 Hz, which differ from that (168 Hz) reported by Shoji and coworkers, 2 and may be taken as evidence for either the equatorial or axial conformation of the anomeric Ara proton as compared with the values for tetra-0-acetyl- α - (17a, 168 Hz) and tetra-0-acetyl- β -L-arabino-pyranoses (18a, 176 Hz). Consequently, these $^1\text{J}(\text{C-1,H-1})(\text{Ara})$ values should not be taken as clear evidence for the anomeric α configuration. Note here that all ^{13}C spectral data concerning the Ara moieties are almost identical among the compounds examined.

¹³C and ¹H NMR Spectroscopic Data for L-Arabinose Moieties of Platycodins, Mi-Saponins, and Their Derivatives^a TABLE.

			Platy	tycodigenin series	in ser	ies					Prot	Protobassate series	te ser	ies				
	$\frac{1-3}{2}$	1-3 4,11 12	12	113	<u>1a</u>	11a	12a	<u>13a</u>	$\frac{9}{10}$	14	15	16	9a	14a	<u>15a</u>	<u>16a</u>	<u>17a</u>	18a
		CS	CSDSN			CDC13	13			CSDSN	SN S			ĕ	CDC13		CD	CDC13
	93.7 75.6	93.7 93.7 93.7 75.6 75.8 75.8	93.7 75.8	95.7	93.2	92.7 73.8 _f	93.0 73.8 _f		93.5 75.4	93.5 75.5	93.5	95.8	93.2 73.4 _f			91.9	92.5	90.6
6-10 6-10 6-10	70.4° 65.9°	70.3	70.4	73.5	68.3 68.3	68.1 [±] 68.3 [±]	67.8 [±] 68.8 [±]	68.7 66.5	70.5 66.1	70.9	70.8	73.6 67.5	67.4 ⁷ 67.8 ⁴	67.7 [±] 67.9 [±]	67.1 ² 67.6 ⁴	69.6 67.1	70.0 67.5 63.7	68.9 67.2 69.0
-1) ^d	172	172	172	165	173	173	173		173	171	171	167	169			169	168 (168	177 176) ⁸
δ _H H-1 H-2	6.29 b	6.29 6.27 6.36 b b b	6.36 b	6.18 b	5.85	5.84	5.85	5.63	6.29 b	6.30 b	6.31 b	6.19 b	5.63	5.61	5.54	5.59	5.67	6.36 5.37
3J(H-1,H-2) ^d 3J(H-2,H-3) ^d 3J(H-4,H-5) ^d	2.8 b b	3.0 b b	3.0 b b	4.5 b	2.9 3.5 7.5	2.9 4.0 7.3	3.3 4.9 7.0	5.0 6.5 5.5	3.0 b	3.0 b b	3.3 b b	4.9 b b	5.5 6.8 b	5.8 7.6 b	5.2 9.0 2.5	6.2 8.5 4.5	6.5 8.5 3.5	2.3 b 1.8
Conformation ^e C	ပ	ပ	ပ	ф	ပ	၁	ပ	щ	ပ	ပ	၁	В	В	æ	В	A	A	¥

tubes with TMS as an internal reference (δC 0) at $100^{\circ}C$ in C_5D_5N and at $60^{\circ}C$ in $CDCl_3$. Complete ^{1}H -decoupled and ^{1}H nona 13C FT and PRFT NMR spectra were recorded on a JEOL FX-100 FT NMR spectrometer at 25.05 MHz in 10-mm spinning spherical (flipping angle 90°); pulse repetition time 1 s (2 s for the gated mode); number of data points, 8K. ¹H FT NMR spectra were recorded on a Varian XL-100-12A and/or a Bruker WH-270 FT NMR spectrometer in 5-mm spinning tubes with TMS as an decoupled (in the gated mode with NOE) FT NMR measurement conditions were: spectral width, 5 KHz; pulse width, 14 µs internal reference (6H 0) at 100°C in C₅D₅N at 100.058 MHz for saponins and at 23°C in CDCl₃ at 100 and 270 MHz for acetates. ¹H FT NMR measurement conditions at 100 and 270 MHz

(in parentheses) were: spectral width, 2 KHz $(3.012~{\rm KHz})$; pulse width, 3 µs [flipping angle 10°] $(10~{\rm µs}~[75^\circ])$; acquisition time, 5.0 s $(2.7~{\rm s})$, numbers of data points, $20{\rm K}~(16{\rm K})$.

b Not observed because of signal overlappings.

^c These assignments were reversed in refs. 3 and 4. ^d Predicted values for ¹J(C-1,H-1), ³J(H-1,H-2), ³J(H-2,H-3), ³J(H-4,H-5) are: ~165, ⁷⁻⁸, ⁷⁻¹⁰, ²⁻³ Hz for α -L-Ara in ⁴C₁; ~177, 2-3, 2-3, ⁷⁻¹⁰ Hz for α -L in ¹C₄; ~177, 3-5, ⁷⁻¹⁰, 2-3 Hz for β -L-Ara in ⁴C₁; ~165, 3-5, 2-3, ⁷⁻¹⁰ Hz for β -L in ¹C₄,

e Conformations assigned for $\alpha\text{-}L\text{-}Ara:$ A, 4C_1 >> $^1C_4;~B,~^4C_1$ > $^1C_4;~C,~^4C_1$ < $^1C_4.$

respectively.

f Assignments may be reversed.

 R_{3} R_{4} R_{5} R_{5

Next, we examined 13 C spectra of the acetates ($\underline{1a}$, $\underline{9a}$, $\underline{11a}$, $\underline{12a}$, $\underline{14a}$, and $\underline{15a}$) in CDCl₃. The data obtained for Ara are shown in the TABLE. Unexpectedly, both δ_{C} and $^{1}J(C-1,H-1)(Ara)$ values for the protobassate (Pro) series (9a, 14a, and 15a) differed from those for the platycodigenin (Pla) series (1a, 11a, and 12a). Also, the ³J(H-1,H-2)(Ara) values at 100 MHz were about 3 and 5.5 Hz for the Pla and Pro series, respectively. The data observed for the Pro series indicate the α -L feature for Ara in contrast with the Pla series.

Thus, we further investigated the 13 C spectra of 13, 16, 5 and their peracetates 6 (13a and 16a⁵). In these spectra, the Ara signals appear to arise from those of α -L- rather than β -L-Ara as compared with those of 17a (see the TABLE). Moreover, their 3J(H-1,H-2)(Ara) values were found to be about 5-6 Hz as reported earlier. 5 These findings suggest that the anomeric configuration is α, as already pointed out.⁵ Rather smaller ³J(H-1,H-2)(Ara) values for ³J(ax, ax) may be due to the considerable contribution of the ${}^{1}C_{4}$ conformation.

The discrepancy of the above results seemed to originate from the conformational difference in their Ara rings in solution. We measured the 270- and/or 100-MHz ¹H FT NMR spectra of the compounds. The Ara ¹H signals were assigned by ¹H-decoupling experiments and by comparisons of the spectra of Pla, Pro, and their peracetates.⁶ The confirmatory evidence for the anomeric configuration together with the Ara-ring conformation was thus provided by ³J(H-1,H-2), ³J(H-2, H-3), ${}^{3}J(H-4,H-5)$, and ${}^{1}J(C-1,H-1)$ values obtained for Ara (see the TABLE for the observed and predicted values). In view of these values, we concluded that the anomeric configuration of Ara is α in all the saponins examined here, and that the Ara-ring predominantly adopts the ${}^{1}\text{C}_{4}$ conformation in saponins 1-4, 9-12, 14, and 15, and peracetates 1a, 11a, and 12a, whereas in saponins 13 and 16, and acetates 9a and 13a-16a, the Ara-ring predominantly adopts the 4C_1 conformation. Thus, the total structures of all saponins from the title plants have been elucidated.

These results suggest that the O-Rha substitution at C-2 in Ara increases the population of the ${}^{1}C_{4}$ conformation. O-Acetylation slightly stabilizes the ${}^{4}C_{1}$ conformation in the Pro series, whereas acetylation in the Pla series may lead to the 16α -OAc group hindering this conformational change by a non-bonded interaction with the Ara moiety. These trends can also be seen in the saponins in each Pro and Pla series; in the latter, the 1C4 population seems to increase a little by the interaction of the 16α -OH group.

Although the present platycodin case might be the most sophisticated one, much caution should be exercised in determining the configuration of sugars such as L-Ara. Variable-temperature NMR studies as well as more detailed data will be reported in a full paper.

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