

STRUCTURAL ELUCIDATION IN ANTHRAQUINONES USING ^1H NMR GLYCOSYLATION AND ALKYLATION SHIFTS

SURAJ B. KALIDHAR

Department of Chemistry and Biochemistry, Haryana Agricultural University, Hisar-125 004, India

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Key Word Index—Emodin; citreoresin; quetin; fallacinol; nalgiovessin; frangilin; islandicin; catenarin; glucoside; primveroside; ^1H NMR; alkylation/glycosylation shifts; peracetates.

Abstract—The trend shown by the alkylation/glycosylation ^1H NMR shifts in the A ring in anthraquinones is not affected by B ring substituents and this observation helps in structural elucidation.

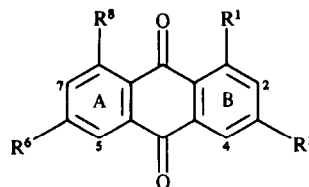
INTRODUCTION

The position of *O*-glycosylation/alkylation in anthraquinones may be determined with the help of ^1H NMR acylation shifts [1]. In an earlier report [2], an alternative procedure was proposed. It involves ^1H NMR comparison of the peracetates of the aglycone and glycoside. The proton, *para* to the site of *O*-glycosylation, undergoes a larger shift. In the absence of a *para* proton, the *ortho* protons undergo larger shifts. The position of *O*-alkylation can be determined in a similar way. In the new method [2], the two anthraquinones being compared differ at one site only, and the shifts pinpoint the site of *O*-glycosylation/alkylation. In the former [1]*, the two anthraquinones (pertrimethylsilyl ether and peracetate) differ at many sites and the shifts cannot be attributed to *O*-glycosylation/alkylation alone. It is now shown that the trend of the shifts in the A ring is not affected by the B ring substitution pattern and this observations helps in structural elucidation in anthraquinones.

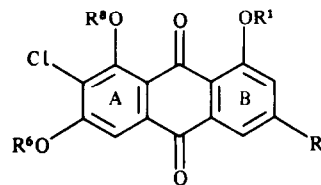
DISCUSSION

A comparison of 8 and 9, having same A ring, shows $\Delta\text{H}-2 = +0.08$ and $\Delta\text{H}-4 = +0.32$ for 1-*O*-alkylation (Tables 1 and 2). Shifts from 2 and 7, with different A rings are $\Delta\text{H}-2 = +0.17$ and $\Delta\text{H}-4 = +0.31$. A trend $\Delta\text{H}-4 > \Delta\text{H}-2$ is true in all of the 39 comparisons made (Table 2). A similar trend, $\Delta\text{H}-4 > \Delta\text{H}-2$, is observed for 1-*O*-glycosylation in 28 comparisons (Table 2). The shifts from 31 and 17 are $\Delta\text{H}-2 = -0.18$ and $\Delta\text{H}-4 = -0.02$ (Table 2) and considering the signs of the shifts, $\Delta\text{H}-4 > \Delta\text{H}-2$, in this extreme case with lowest $\Delta\text{H}-4$ (Table 2).

The 6-*O*-alkylation shifts from 30 comparisons are shown in Table 3. The shifts, $\Delta\text{H}-5$ and $\Delta\text{H}-7$, are $> +0.20$ in 29 comparisons. There is deviation in one case only, that is, comparison of 34 and 6 shows $\Delta\text{H}-5 = +0.18$ ($\leq +0.2$) and $\Delta\text{H}-7 = +0.25$. Despite slight deviation ($\Delta\text{H}-5 \leq +0.20$) in this particular case, it cannot be confused with 8-*O*-alkylation (Table 4) which requires $\Delta\text{H}-5 > \Delta\text{H}-7$ (here $\Delta\text{H}-7 > \Delta\text{H}-5$). Of the four 6-*O*-glycosylation shifts shown in Table 3, three are as predic-



- 1 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{OH}, \text{R}^3 = \text{CH}_2\text{OH}$
- 2 $\text{R}^1, \text{R}^6 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^8 = \text{OMe}$
- 3 $\text{R}^1, \text{R}^6 = \text{OH}, \text{R}^3 = \text{CH}_2\text{OH}, \text{R}^8 = \text{OMe}$
- 4 $\text{R}^1, \text{R}^8 = \text{OH}, \text{R}^3 = \text{CH}_2\text{OH}, \text{R}^6 = \text{OMe}$
- 5 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{OH}, \text{R}^3 = \text{Me}$
- 6 $\text{R}^1, \text{R}^8 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{OMe}$
- 7 $\text{R}^1, \text{R}^6 = \text{OMe}, \text{R}^3 = \text{Me}, \text{R}^8 = \text{OH}$
- 8 $\text{R}^1 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6, \text{R}^8 = \text{OMe}$
- 9 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{OMe}, \text{R}^3 = \text{Me}$
- 10 $\text{R}^1 = \text{Me}, \text{R}^3, \text{R}^6, \text{R}^8 = \text{OMe}$
- 11 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{OMe}, \text{R}^3 = \text{CH}_2 \cdot \text{CH}(\text{OH}) \cdot \text{Me}$
- 12 $\text{R}^1, \text{R}^8 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{H}$
- 13 $\text{R}^1, \text{R}^6 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^8 = \text{O-glc}$
- 14 $\text{R}^1, \text{R}^6 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^8 = \text{-O-gent}$
- 15 $\text{R}^1, \text{R}^6 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^8 = \text{O-prim}$
- 16 $\text{R}^1, \text{R}^8 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{O-glc}$
- 17 $\text{R}^1 = \text{-O-glc}, \text{R}^3 = \text{Me}, \text{R}^6, \text{R}^8 = \text{OH}$
- 18 $\text{R}^1 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{OMe}, \text{R}^8 = \text{-O-glc}$
- 19 $\text{R}^1 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{OMe}, \text{R}^8 = \text{O-gent}$
- 20 $\text{R}^1 = \text{OH}, \text{R}^3 = \text{Me}, \text{R}^6 = \text{OMe}, \text{R}^8 = \text{-O-prim}$

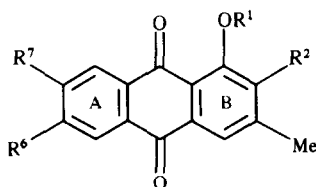


- 21 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{Me}, \text{R}^3 = \text{CH}_2 \cdot \text{CHOH} \cdot \text{Me}$
- 22 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{H}, \text{R}^3 = \text{Me}$
- 23 $\text{R}^1, \text{R}^3, \text{R}^6, \text{R}^8 = \text{Me}$
- 24 $\text{R}^1, \text{R}^6, \text{R}^8 = \text{H}, \text{R}^3 = \text{CH}_2\text{OH}$

*This method [1] is less sensitive than the new one [25].

Table 1. Chemical shifts of aromatic protons in the ^1H NMR of the peracetates of anthraquinones

Peracetates of anthraquinones	Chemical shifts† (δ , CDCl_3)			
	H-2	H-4	H-5	H-7
Citreoresin (1) [3]	7.42	8.19	7.99	7.29
Questin (2) [3]	7.22	8.01	7.65	7.11
Questinol (3) [3]	7.42	8.18	7.67	7.14
Fallacinol (4) [4]	7.35	8.15	7.65	6.88
Emodin (5) [1]	7.24	8.03	7.98	7.26
Physcion (6) [1]	7.24	8.03	7.69	6.92
1- <i>O</i> -Methylphyscion (7) [5]	7.05	7.70	7.60	6.95
8- <i>O</i> -Methylphyscion (8) [1]	7.18	7.96	7.36	6.77
*1,8-Di- <i>O</i> -methylphyscion (9) [6]	7.10	7.64	7.33	6.77
*3,6,8-Tri- <i>O</i> -methyl-1-methylantraquinone (10)	6.92	7.53	7.32	6.74
*Di- <i>O</i> -methylalgiovensin (11) [8]	7.17	7.68	7.28	6.75
Chrysophanol (12) [9]	7.21	8.01	8.21	7.38‡
Emodin-8- <i>O</i> -glucoside (13) [1]	7.23	7.97	7.76	7.31
Emodin-8- <i>O</i> -gentiobioside (14) [10]	7.19	7.97	7.78	7.29
Emodin-8- <i>O</i> -primveroside (15) [11]	7.17	7.96	7.78	7.21
Emodin-6- <i>O</i> -glucoside (16) [1]	7.19	7.91	7.54	6.98
Emodin-1- <i>O</i> -glucoside (17) [1]	7.35	7.88	7.97	7.29
Physcion-8- <i>O</i> -glucoside (18) [1]	7.15	7.90	7.42	6.94
Physcion-8- <i>O</i> -gentiobioside (19) [12]	7.18	7.92	7.50	6.88
Physcion-8- <i>O</i> -primveroside (20) [13]	7.13	7.88	7.45	6.97
*Di- <i>O</i> -methylalgiolaxin (21) [8]	7.18	7.68	7.53	—
Norfrangilin (22) [14]	7.23	7.98	8.0	—
*Di- <i>O</i> -methylfrangilin (23) [15]	7.07	7.64	7.55	—
1,6,8-Trihydroxy-3-hydroxymethyl-7-chloroanthraquinone (24) [14]	7.40	8.16	8.03	—
1,3,8-Trihydroxy-6- <i>O</i> -methyl-2-methylantraquinone (25) [16]	—	7.89	7.65	6.83
*2-Carbomethoxy-3,6,8-tri- <i>O</i> -methyl-1-methylantraquinone (26) [7]	—	7.63	7.31	6.77
Alaternin (27) [17]	—	8.11	7.95	7.24
6- <i>O</i> -Methylalaternin (28) [18]	—	7.95	7.50	6.80
Xanthorin (29) [19]	7.17	7.90	—	6.95
*Tri- <i>O</i> -methylaxanthorin (30) [19]	7.05	7.55	—	6.80
Dermocybin (31) [1]	7.17	7.86	—	—
Dermocybin-1- <i>O</i> -glucoside (32) [1]	7.24	7.70	—	—
Islandicin (33) [20]	7.24	—	8.05	7.32‡
Catenarin (34) [21]	7.23	—	7.87	7.17

* ^1H NMR of the unacetylated compound.† Multiplicities and J values have been excluded in this Table.‡ H-6 = δ 7.74 and 7.68 for **12** and **33** respectively.**35** $\text{R}^1, \text{R}^2, \text{R}^7 = \text{H}, \text{R}^6 = \text{OH}$ **36** $\text{R}^1, \text{R}^2, \text{R}^6 = \text{H}, \text{R}^7 = \text{OH}$ **37** $\text{R}^1 = \text{Me}, \text{R}^2 = \text{OH}, \text{R}^6 = \text{OMe}, \text{R}^7 = \text{H}$ **38** $\text{R}^1 = \text{Me}, \text{R}^2 = \text{OH}, \text{R}^6 = \text{H}, \text{R}^7 = \text{OMe}$

ted ($> +0.20$). The fourth has a slight deviation, i.e. shifts from **34** and **16** are $\Delta\text{H-5} = +0.33$ and $\Delta\text{H-7} = +0.19$ ($\leq +0.2$). Although $\Delta\text{H-5}$ is larger than $\Delta\text{H-7}$ it cannot be confused with 8-*O*-glycosylation which has $\Delta\text{H-7} <$

$+0.10$ (Table 4) (here $\Delta\text{H-7} > +0.10$). When shifts for H-5 and H-7 are both $> +0.20$, there are indications for 6-*O*-alkylation/glycosylation and in such cases no attention is required whether $\Delta\text{H-5} > \Delta\text{H-7}$ or $\Delta\text{H-7} > \Delta\text{H-5}$ etc.

The 8-*O*-alkylation shifts from 28 comparisons are shown in Table 4. The trend $\Delta\text{H-5} > \Delta\text{H-7}$ is always true. There is a similar trend for 8-*O*-glycosylation ($\Delta\text{H-5} > \Delta\text{H-7}$) from a comparison of 24 pairs. The glycosylation shifts from **28** and **19** are $\Delta\text{H-5} = 0.00$ and $\Delta\text{H-7} = -0.08$ and $\Delta\text{H-5} > \Delta\text{H-7}$ in this peculiar case with no shift for H-5. The 6,8-di-*O*-alkylation shifts from 24 comparisons are shown in Table 5. The trend is $\Delta\text{H-5} > +0.40$ and $\Delta\text{H-7} > +0.40$ and $\Delta\text{H-5}$ is always $> \Delta\text{H-7}$.

There are reports of partial structures 1,6-dihydroxy-3-methylantraquinone **35**/1,7-dihydroxy-(**36**) [21] and 2-hydroxy-1,6-di-*O*-methyl-3-methylantraquinone (**37**)/2-hydroxy-1,7-di-*O*-methyl-(**38**) [22]. Of these, those preferred on biogenetic considerations are **35** [21] and **37** [23].

Table 2. 1-*O*-Alkylation/glycosylation shifts

1- <i>O</i> -Alkylation (Δ^*H)											
Anthra- quinones	Δ^*H-2	Δ^*H-4	Anthra- quinones	Δ^*H-2	Δ^*H-4	Anthra- quinones	Δ^*H-2	Δ^*H-4	Anthra- quinones	Δ^*H-2	Δ^*H-4
2, 7	+0.17	+0.31	8, 9	+0.08	+0.32	15, 9	+0.07	+0.32	19, 30	+0.13	+0.37
2, 9	+0.12	+0.37	8, 30	+0.13	+0.41	15, 30	+0.12	+0.41	20, 7	+0.08	+0.18
2, 30	+0.17	+0.46	12, 30	+0.16	+0.46	16, 7	+0.14	+0.21	20, 9	+0.03	+0.24
5, 7	+0.19	+0.33	13, 7	+0.16	+0.27	16, 9	+0.09	+0.27	20, 30	+0.08	+0.33
5, 9	+0.14	+0.39	13, 9	+0.13	+0.33	16, 30	+0.14	+0.36	22, 30	+0.18	+0.43
5, 30	+0.19	+0.48	13, 30	+0.18	+0.42	18, 7	+0.10	+0.20	29, 30	+0.12	+0.35
6, 7	+0.19	+0.33	14, 7	+0.14	+0.27	18, 9	+0.05	+0.26	31, 7	+0.12	+0.16
6, 9	+0.14	+0.39	14, 9	+0.09	+0.33	18, 30	+0.10	+0.35	31, 9	+0.07	+0.22
6, 30	+0.19	+0.48	14, 30	+0.14	+0.42	19, 7	+0.13	+0.22	31, 30	+0.12	+0.31
8, 7	+0.13	+0.26	15, 7	+0.12	+0.26	19, 9	+0.08	+0.28			
1- <i>O</i> -glycosylation ($\Delta H^\dagger$)											
	$\Delta H-2$	$\Delta H-4$		$\Delta H-2$	$\Delta H-4$		$\Delta H-2$	$\Delta H-4$		$\Delta H-2$	$\Delta H-4$
2, 17	-0.13	+0.13	12, 32	-0.03	+0.31	16, 17	-0.16	+0.03	20, 32	-0.11	+0.18
2, 32	-0.02	+0.31	13, 17	-0.12	+0.09	16, 32	-0.05	+0.21	22, 17	-0.12	+0.10
5, 17	-0.11	+0.15	13, 32	-0.01	+0.27	18, 17	-0.20	+0.02	22, 32	-0.01	+0.28
5, 32	0.0	+0.33	14, 17	-0.16	+0.09	18, 32	-0.09	+0.20	29, 17	-0.18	+0.02
6, 17	-0.11	+0.15	14, 32	-0.05	+0.27	19, 17	-0.17	+0.04	29, 32	-0.07	+0.20
6, 32	0.0	+0.33	15, 17	-0.18	+0.08	19, 32	-0.06	+0.22	31, 17	-0.18	-0.02
12, 17	-0.14	+0.13	15, 32	-0.07	+0.26	20, 17	-0.22	0.0	31, 32	-0.07	+0.16

* $\Delta^*H = \delta$ value of the aromatic proton in anthraquinone peracetate minus that in *O*-alkyl peracetate.

$\dagger\Delta H = \delta$ value of the aromatic proton in aglycone peracetate minus that in *O*-glycosyl peracetate.

Table 3. 6-*O*-Alkylation/glycosylation shifts.

6- <i>O</i> -alkylation (Δ^*H)								
Anthra- quinones	Δ^*H-5	Δ^*H-7	Anthra- quinones	Δ^*H-5	Δ^*H-7	Anthra- quinones	Δ^*H-5	Δ^*H-7
1, 4	+0.34	+0.41	3, 8	+0.31	+0.37	13, 18	+0.32	+0.37
1, 6	+0.30	+0.37	3, 9	+0.34	+0.37	14, 19	+0.28	+0.41
1, 7	+0.39	+0.34	3, 10	+0.35	+0.40	15, 20	+0.33	+0.24
1, 25	+0.34	+0.46	3, 11	+0.39	+0.39	27, 6	+0.26	+0.32
1, 28	+0.49	+0.49	3, 26	+0.36	+0.37	27, 7	+0.35	+0.29
2, 8	+0.29	+0.38	5, 4	+0.33	+0.38	34, 4	+0.22	+0.29
2, 9	+0.32	+0.34	5, 6	+0.29	+0.34	34, 6	+0.18	+0.25
2, 10	+0.33	+0.37	5, 7	+0.38	+0.31	34, 7	+0.27	+0.22
2, 11	+0.37	+0.36	5, 25	+0.33	+0.43	34, 25	+0.22	+0.34
2, 26	+0.34	+0.34	5, 28	+0.48	+0.46	34, 28	+0.37	+0.37
6- <i>O</i> -glycosylation (ΔH)								
	$\Delta H-5$	$\Delta H-7$		$\Delta H-5$	$\Delta H-7$		$\Delta H-5$	$\Delta H-7$
1, 16	+0.45	+0.31	5, 16	+0.44	+0.28	27, 16	+0.41	+0.26
						34, 16	+0.33	+0.19

Table 4. 8-*O*-Alkylation/glycosylation shifts

8- <i>O</i> -alkylation ($\Delta'H$)								
Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7	Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7	Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7
1, 2	+0.34	+0.18	6, 9	+0.36	+0.15	27, 3	+0.28	+0.10
1, 3	+0.32	+0.15	6, 10	+0.37	+0.18	28, 8	+0.14	+0.03
4, 8	+0.29	+0.11	6, 11	+0.41	+0.17	28, 9	+0.17	+0.03
4, 9	+0.32	+0.11	6, 26	+0.38	+0.15	28, 10	+0.18	+0.06
4, 10	+0.33	+0.14	25, 8	+0.29	+0.06	28, 11	+0.22	+0.05
4, 11	+0.37	+0.13	25, 9	+0.32	+0.06	28, 26	+0.19	+0.03
4, 26	+0.34	+0.11	25, 10	+0.33	+0.09			
5, 2	+0.33	+0.15	25, 11	+0.37	+0.08	34, 2	+0.22	+0.06
5, 3	+0.31	+0.12	25, 26	+0.34	+0.06	34, 3	+0.20	+0.03
6, 8	+0.36	+0.15	27, 2	+0.30	+0.13			
8- <i>O</i> -glycosylation (ΔH)								
	ΔH -5	ΔH -7		ΔH -5	ΔH -7		ΔH -5	ΔH -7
1, 13	+0.23	-0.02	5, 15	+0.20	+0.05	27, 14	+0.17	-0.05
1, 14	+0.21	0.0	6, 18	+0.27	-0.02	27, 15	+0.17	+0.03
1, 15	+0.21	+0.08	6, 19	+0.19	+0.04	28, 18	+0.08	-0.14
4, 18	+0.23	+0.06	6, 20	-0.24	-0.05	28, 19	0.0	-0.08
4, 19	+0.15	0.0	25, 18	+0.23	-0.11	28, 20	+0.05	-0.17
4, 20	+0.20	+0.09	25, 19	+0.15	-0.05	34, 13	+0.11	-0.14
5, 13	+0.22	-0.05	25, 20	+0.20	-0.14	34, 14	+0.09	-0.12
5, 14	+0.20	-0.03	27, 13	+0.19	-0.07	34, 15	+0.09	-0.04

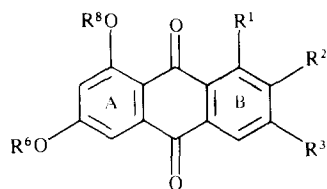
Table 5. 6,8-Di-*O*-alkylation shifts

Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7	Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7	Anthra-quinones	$\Delta'H$ -5	$\Delta'H$ -7
1, 8	+0.63	+0.52	5, 11	+0.70	+0.51	27, 10	+0.63	+0.50
1, 9	+0.66	+0.52	5, 26	+0.67	+0.49	27, 11	+0.67	+0.49
1, 10	+0.67	+0.55	22, 21	+0.47	—	27, 26	+0.64	+0.47
1, 11	+0.71	+0.54	22, 23	+0.45	—	34, 8	+0.51	+0.40
1, 26	+0.68	+0.52	24, 23	+0.48	—	34, 9	+0.54	+0.40
5, 8	+0.62	+0.49	24, 21	+0.50	—	34, 10	+0.55	+0.43
5, 9	+0.65	+0.49	27, 8	+0.59	+0.47	34, 11	+0.59	+0.42
5, 10	+0.66	+0.52	27, 9	+0.62	+0.47	34, 26	+0.56	+0.40

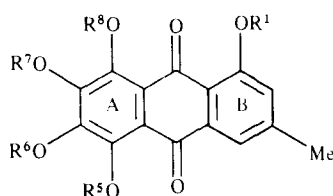
Table 6. Chemical shifts of A ring protons in the ¹H NMR of anthraquinones and alkylation shifts

Anthraquinones	Chemical shifts (δ)			
	H-5	H-6	H-7	H-8
† 1,6-Dihydroxy-3-methylanthraquinone (35) [20]	7.95	—	7.98	8.25
‡ 2-Hydroxy-1,6- <i>O</i> -methyl-3-methylanthraquinone (37) [21]	7.68	—	7.22	8.19
Alkylation shifts ($\Delta'H$)				
35, 37	$\Delta'H$ -5 +0.27	$\Delta'H$ -6 —	$\Delta'H$ -7 +0.26	$\Delta'H$ -8 +0.06

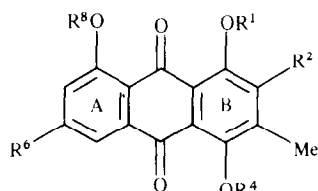
† As peracetate in CDCl₃.
‡ Unacetylated in CDCl₃.



- 25** $R^1, R^3 = \text{OH}, R^2, R^6 = \text{Me}, R^8 = \text{H}$
26 $R^1, R^6 = \text{Me}, R^2 = \text{CO}_2\text{Me}, R^3 = \text{OMe}, R^8 = \text{Me}$
27 $R^1, R^2 = \text{OH}, R^3 = \text{Me}, R^6, R^8 = \text{H}$
28 $R^1, R^2 = \text{OH}, R^3, R^6 = \text{Me}, R^8 = \text{H}$



- 29** $R^1, R^5, R^7, R^8 = \text{H}, R^6 = \text{Me}$
30 $R^1, R^5, R^6, R^8 = \text{Me}, R^7 = \text{H}$
31 $R^1, R^5, R^7, R^8 = \text{H}, R^6 = \text{Me}$
32 $R^1, R^5, R^7 = \text{H}, R^6 = \text{Me}, R^8 = \text{glc}$



- 33** $R^1, R^2, R^4, R^6, R^8 = \text{H}$
34 $R^1, R^4, R^8 = \text{H}, R^2 = \text{H}, R^6 = \text{OH}$

The identity of **35** has been confirmed by comparison with a synthetic sample [24]. The $^1\text{H NMR}$ data are shown in Table 6 and the 6-*O*-alkylation shifts $\Delta\text{H-5} = +0.27$, $\Delta\text{H-7} = +0.26$ are consistent with the preferred structure (**37**).

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