

Absolute Configuration of *epi*-Rhododendrin and (–)-Rhododendrol [= (–)-Betuligenol] and X-Ray Crystal and Molecular Structure of Rhododendrin [= Betuloside], a Hepatoprotective Constituent of *Taxus baccata*¹

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Rhododendrin (= betuloside) has been isolated from the leaves of *Taxus baccata* L. From its X-ray diffraction studies, the absolute configuration at the chiral centre in the aglucone portion has been found to be *R*. These results establish the stereochemical assignments to different samples of this glucoside isolated from different plants. Our sample of rhododendrin exhibited hepatoprotective activity against two hepatotoxins in rats.

Different *Taxus* species have evoked a good deal of interest among phytochemists in recent years because they have been found to possess a range of biological activities, *i.e.* cytotoxic and antileukaemic;² sedative and antiseptic;³ tranquilising,⁴ and antimitotic.⁵ Khan *et al.*⁶ have isolated betuloside as the major component of the aqueous extract of the leaves of *Taxus baccata*; in view of the fact that the aqueous extract possesses good tranquilising activity,⁴ we planned to carry out a detailed study on the structure and biological activities of betuloside. On close scrutiny of the literature, we found an interesting series of papers relating to the occurrence and structure of this compound. This glycoside, with the name rhododendrin, along with its aglucone rhododendrol, was isolated for the first time from the leaves of *Rhododendron chrysanthum* Linn. in 1901.⁷ Later, a similar glucoside, named betuloside, was isolated from the bark of the white birch, *Betula alba* L.^{8,9} Rhododendrin isolated from *Rhododendron fauriae* var. *rufescens*,¹⁰ and its aglucone rhododendrol, were subsequently shown to be identical with betuloside and betuligenol, respectively.¹¹ More recently the distribution of rhododendrin in the genera *Rhododendron*^{12–14} and *Betula*¹⁵ has been surveyed and it has also been found in *Bergenia* species,¹² *Abies webbiana* Lindl.,¹⁶ *Rhodiola* species,¹⁷ *Alnus glutinosa* Gaertn.,¹⁸ *Taxus baccata* L.,⁶ and *Cotyledon wallichii*.¹⁹

The biosynthesis of rhododendrin has been studied¹⁸ and it was established that it is formed through the phenylpropane pathway *via p*-coumaryl alcohol, dihydro-*p*-coumaryl alcohol, and *C*-methylation of the γ -C-atom of the C₆–C₃ unit with methionine supplying the methyl group. It was demonstrated that the *pro*-(*S*) hydrogen atom of dihydro-*p*-coumaryl alcohol is replaced stereospecifically by the methyl group. There is ambiguity about the configuration at C-2; however, in the previously reported structural formula of rhododendrin the configuration at this chiral centre was depicted as *S*.¹⁸

The aglucone, (–)-rhododendrol, is found to co-occur in many species, but it is interesting that (+)-rhododendrol has been isolated from *Rhododendron maximum*.²⁰ Furthermore (+)-rhododendrol and its glucoside *epi*-rhododendrin were isolated in 1978 from *Acer nikoense* MAXIM. and were proposed to have the (2*S*)-configuration from both the empirical rule on optical rotation–chirality relations²¹ and ¹³C NMR studies.²²

In the light of these different findings concerning configuration, it was of interest to carry out an X-ray study of

Table 1 Bond angles (°) for rhododendrin

Atom 1, 2 and 3	Angle	Atom 1, 2 and 3	Angle
C(1)–O(1)–C(5)	111.0(1)	C(1)–O(6)–C(7)	112.3(1)
O(1)–C(1)–C(2)	109.7(1)	O(6)–C(7)–C(8)	109.0(2)
O(1)–C(1)–O(6)	108.6(1)	O(6)–C(7)–C(16)	109.3(1)
C(2)–C(1)–O(6)	108.8(1)	C(8)–C(7)–C(16)	112.2(2)
C(1)–C(2)–C(3)	108.7(1)	C(7)–C(8)–C(9)	113.4(2)
C(1)–C(2)–O(2)	111.1(1)	C(8)–C(9)–C(10)	114.4(2)
C(3)–C(2)–O(2)	108.8(1)	C(9)–C(10)–C(11)	120.1(2)
C(2)–C(3)–C(4)	110.5(1)	C(9)–C(10)–C(15)	122.3(2)
C(2)–C(3)–O(3)	111.4(1)	C(11)–C(10)–C(15)	117.6(2)
C(4)–C(3)–O(3)	112.2(1)	C(10)–C(11)–C(12)	122.1(2)
C(3)–C(4)–C(5)	110.3(1)	C(11)–C(12)–C(13)	119.8(2)
C(3)–C(4)–O(4)	112.1(1)	C(12)–C(13)–C(14)	119.6(2)
C(5)–C(4)–O(4)	111.4(1)	C(12)–C(13)–O(7)	118.8(2)
O(1)–C(5)–C(4)	108.2(1)	C(14)–C(13)–O(7)	121.7(2)
O(1)–C(5)–C(6)	108.0(1)	C(13)–C(14)–C(15)	120.2(2)
C(4)–C(5)–C(6)	113.5(1)	C(10)–C(15)–C(14)	120.8(2)
C(5)–C(6)–O(5)	112.4(1)		

Numbers in parentheses are estimated standard deviations in the least significant digits.

Table 2 Bond distances (Å) in rhododendrin

Atom 1 and 2	Distance	Atom 1 and 2	Distance
O(1)–C(1)	1.421(2)	O(6)–C(7)	1.455(2)
O(1)–C(5)	1.435(2)	C(7)–C(8)	1.519(2)
C(1)–C(2)	1.523(2)	C(7)–C(16)	1.517(3)
C(1)–O(6)	1.384(2)	C(8)–C(9)	1.528(3)
C(2)–C(3)	1.533(2)	C(9)–C(10)	1.520(2)
C(2)–O(2)	1.411(2)	C(10)–C(11)	1.388(3)
C(3)–C(4)	1.522(2)	C(10)–C(15)	1.403(3)
C(3)–O(3)	1.423(2)	C(11)–C(12)	1.388(3)
C(4)–C(5)	1.544(2)	C(12)–C(13)	1.379(3)
C(4)–O(4)	1.422(2)	C(13)–C(14)	1.397(3)
C(5)–C(6)	1.522(2)	C(13)–O(7)	1.379(2)
C(6)–O(5)	1.439(2)	C(14)–C(15)	1.391(2)

Numbers in parentheses are estimated standard deviations in the least significant digits.

rhododendrin (betuloside) isolated by us from *Taxus baccata* L. This work, and the conclusions concerning the configuration, are reported in the present paper.

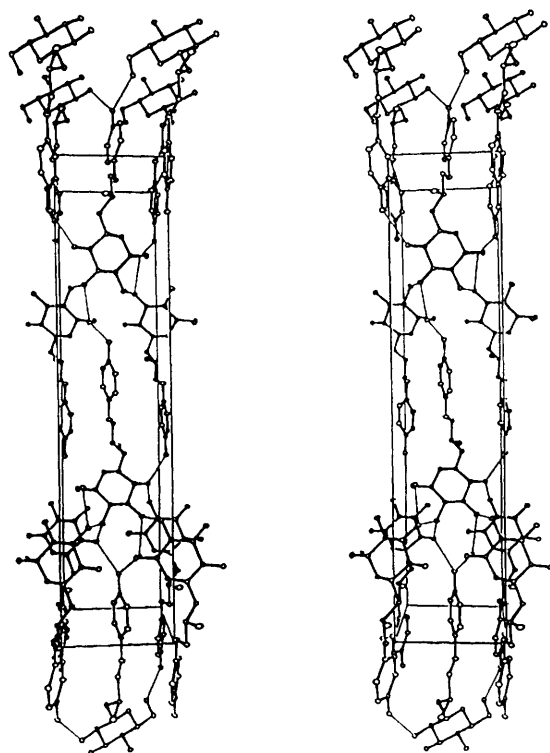
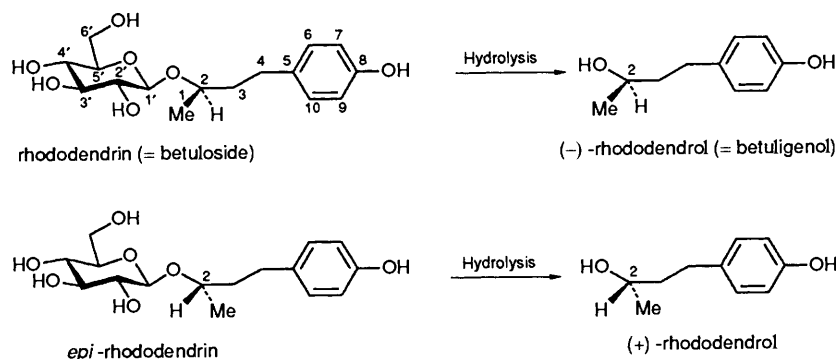


Fig. 2 Stereo-pair packing of rhododendrin

Single-crystal X-Ray Structure Determination of Rhododendrin.—A single crystal, of dimensions $0.12 \times 0.22 \times 0.30$ mm, was selected for the X-ray analysis. The crystal was cooled to 110 K during the experiment with an Enraf-Nonius gas-flow low-temperature device. A CAD4 diffractometer was used for the data collection using Cu-K α radiation obtained from a graphite monochromator. Rhododendrin crystallises from ethyl acetate in the orthorhombic space group $P2_12_12_1$. The setting angles for 25 reflections ($37^\circ < \theta < 41^\circ$) were used in a least-squares refinement to determine the unit-cell parameters: $a = 6.3488(9)$; $b = 7.6033(12)$; $c = 33.074(5)$ Å; $V = 1596.5(7)$ Å³. The empirical formula for betuloside $C_{16}H_{24}O_7$, formula weight 328.35, gives $Z = 4$ with $D_{\text{calc}} = 1.366$ g cm⁻³. The ω scan mode, $\Delta\omega = 1.6^\circ$, was employed for the data collection, where the maximum scan time was 120 s. The intensities of three standard reflections recorded every 10 000 s remained constant throughout the experiment. Diffraction data were measured in the range $1^\circ < \theta \leq 75^\circ$. Data reduction included corrections for Lorentz and polarisation effects and symmetry-related reflections were averaged according to the symmetry of the Laue group. This gave 1943 independent reflections, 1809 of which had $|F|^2 > 2[\sigma(|F|^2) + (0.05 \times |F|^2)^2]$

and were used in the structure solution and refinement. The structure was solved by application of the SHELXS-86²⁵ program, which gave starting co-ordinates for all the non-hydrogen atoms.

The least-squares method was used to refine the structure by minimising $\sum w(|F_o| - |F_c|)^2$. The positions of all the hydrogen atoms were shown clearly in a difference electron density. Only the positions of the hydrogen atoms bonded to oxygen were taken from the difference Fourier; the remaining hydrogen atoms were introduced in idealised positions. In the final refinement cycles their parameters were fixed and they were given a common isotropic displacement parameter $B = 2.5$ Å². The final residuals were $R = 0.041$ and $R_w = 0.067$. The maximum shift was 0.02 σ and the residual electron density was featureless.

Rhododendrin pentaacetate* was obtained by acetylation of rhododendrin with dry pyridine and acetic anhydride at 25 °C for 24 h and was crystallised from methanol as needles, m.p. 82–85 °C; δ_H ([²H₆]acetone) 1.15 (3 H, d, J 7, Me), 1.65–1.90 (2 H, m, 3-H₂), 1.95–2.10 (12 H, 4 s, 4 \times OAc of sugar moiety), 2.25 (3-H, s, 8-OAc), 2.65–2.90 (2 H, m, 4-H₂), 3.80–5.35 (8 H, 7 m, 2-, 1'-, 2'-, 3'-, 4'- and 5'-H and 6'-H₂), 7.0 (2 H, d, J 9, 6- and 10-H) and 7.28 (2 H, d, J 9, 7- and 9-H); δ_C ([²H₆]acetone) 20.12, 20.40, 20.45, 20.50 and 20.81 (5 s, C-1 and 5 \times OCOMe), 31.17 (s, C-3), 39.33 (s, C(4)), 62.75 (s, C-6'), 69.44 (s, C-4'), 72.17 (d, C-2), 73.46, (s, C-2'), 75.26 (s, C-3' and -5'), 99.62 (s, C-1'), 122.16 (s, C-7 and -9), 130.04 (s, C-6 and -10), 140.46 (s, C-5) and 149.85 (s, C-8) and 169.45, 169.57, 169.86, 170.16 and 170.60 (5 s, 5 \times OCOMe); m/z (rel. intensity) 538 (M^+ , 1), 497 (3), 496 (12), 332 (3), 331 (14), 303 (2), 229 (3), 219 (4), 191 (12), 190 (21), 169 (34), 149 (53), 148 (100), 133 (8), 115 (7), 109 (17), 107 (67), 98 (6), 81 (4) and 43 (64).

(-)-Rhododendrol was obtained by hydrolysis of rhododendrin with 5% aq. HCl by the method of Khan *et al.*⁶ and was crystallised from benzene as needles, m.p. 79–80 °C; $[\alpha]_D^{21} -17^\circ$ (c 0.5, MeOH); δ_H (CDCl₃) 1.32 (3 H, d, J 7, Me),

* The pentaacetyl derivative of this glucoside had previously been reported by Khan *et al.*,⁶ the m.p. of our pentaacetate (82–85 °C) is different to that reported by them (185–186 °C).⁶ The pentaacetyl derivative of rhododendrin has also been prepared by Kawaguchi *et al.*,¹⁰ but only the m.p. (96–97 °C) was reported, which is more in line with that of our acetyl derivative. Khan *et al.*⁶ have reported only ¹H NMR data for their derivative, while we have fully characterised our betuloside pentaacetate from its ¹H NMR, ¹³C NMR, and mass spectra. It is noteworthy that in its MS the pentaacetate exhibits the molecular ion peak at m/z 538 (1%).

† The $[\alpha]_D$ -values of different samples of the aglucone reported by different groups of workers are: -20° (c 1.25, EtOH) for betuligenol from *T. baccata*;⁶ -18.5° (c 3.88, EtOH) for betuligenol from *B. alba*;⁸ $+20^\circ$ (c 1.0, EtOH) for (+)-rhododendrol from *A. nikoense* MAXIM;²¹ and $+17.1^\circ$ (c 2.0, EtOH) for *d*-betuligenol from *R. maximum*,²⁰ while the respective m.p.s (°C) are: 80, 6, 81.5, 9, 81–82^{14,21} and 81–83.²⁰

Table 4 Protection (%) provided by rhododendrin in the levels of different enzymes in rats

Enzyme	Galactosamine model rhododendrin (10 mg kg ⁻¹)	Thioacetamide model rhododendrin (25 mg kg ⁻¹)
1 Glutamate oxaloacetate transaminase (GOT)	0	21
2 Glutamate pyruvate transaminase (GPT)	0	26
3 Acid phosphatase (ACP)	11	20
4 Alkaline phosphatase (ALP)	44	36
5 Glutamate dehydrogenase (GDH)	1	100

1.57 (br s, 2-OH), 1.67–1.85 (2 H, m, 3-H₂), 2.52–2.77 (2 H, m, 4-H₂), 3.75–3.92 (1 H, m, 2-H), 5.40 (br s, 8-OH), 6.75 (2 H, d, J 9, 7- and 9-H) and 7.05 (2 H, d, J 9, 6- and 10-H); $\delta_c(\text{CDCl}_3)$ 23.43 (s, C-1), 31.09 (s, C-3), 40.87 (s, C-4), 67.63 (s, C-2), 115.20 (s, C-7 and -9), 129.32 (s, C-6 and -10), 133.82 (s, C-5) and 153.74 (s, C-8); m/z (rel. intensity) 167 [(M + 1)⁺, 4], 166 (M⁺, 36), 149 (4), 148 (26), 134 (8), 133 (76), 121 (6), 120 (6), 108 (19), 107 (100), 105 (4), 103 (4), 94 (14), 91 (10), 78 (6), 77 (22), 65 (6), 55 (5), 45 (19), 43 (8) and 39 (10).

Hepatobiliary Effect of Rhododendrin.—For both the experiments, male Sprague-Dawley rats (100–125 g) were caged separately in groups of five animals each. Group I consisted of normal animals. Group II animals were administered the hepatotoxin galactosamine hydrochloride (800 mg kg⁻¹, i.p. \times 1) or thioacetamide (200 mg kg⁻¹, s.c. \times 1). Group III animals were fed the respective doses of rhododendrin and were injected with the hepatotoxin on day 7.

Animals of all three groups were sacrificed 24 h after administration of the hepatotoxin and their blood was collected and subjected to centrifugation (Remi C-23 cooling centrifuge) at 2000 rpm for 0.2 h. In the serum, different enzymatic parameters as described in Table 4 were analysed according to standard laboratory methods. The hepatoprotective activity of the drug was assessed by the percentage reversal effected by the drug on various enzyme levels elevated by the hepatotoxin.

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