

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (\AA^2)

	x	y	z	B_{eq}
N1	-0.2128 (7)	0.7957 (2)	0.6965 (2)	4.11 (8)
C1A	-0.3028 (8)	0.8593 (2)	0.7484 (2)	3.70 (8)
C1B	-0.2556 (9)	0.8249 (2)	0.8234 (2)	4.7 (1)
C1G	-0.3154 (13)	0.8826 (4)	0.8856 (3)	6.7 (2)
C1D1	-0.5833 (19)	0.9098 (8)	0.8868 (4)	11.9 (3)
C1D2	-0.237 (2)	0.8377 (5)	0.9536 (3)	9.9 (3)
C1'	-0.1541 (7)	0.9386 (2)	0.7356 (2)	3.61 (8)
O1'	0.0765 (5)	0.9372 (2)	0.7248 (2)	4.51 (7)
N2	-0.2915 (6)	1.0080 (2)	0.7374 (2)	4.05 (8)
C2A	-0.1791 (8)	1.0911 (2)	0.7326 (2)	4.2 (1)
C2B	-0.1516 (18)	1.1148 (3)	0.6537 (3)	7.9 (2)
C2'	-0.3464 (9)	1.1533 (2)	0.7713 (2)	4.6 (1)
O21'	-0.2578 (7)	1.2251 (1)	0.7785 (2)	6.1 (1)
O22'	-0.5644 (7)	1.1329 (2)	0.7906 (3)	7.1 (1)
Solvent molecule				
S1†	0.0659 (6)	0.1503 (2)	0.0329 (1)	7.88 (7)
S2†	-0.0570 (9)	0.0861 (2)	0.0137 (2)	6.9 (8)
O3	-0.2197 (12)	0.1414 (4)	0.0590 (2)	9.9 (2)
C1	0.020 (2)	0.1416 (5)	-0.0631 (4)	11.7 (4)
C2	0.207 (2)	0.0541 (7)	0.0562 (5)	11.6 (3)

† S1 and S2 were refined with site-occupancy factors of 0.62 and 0.38, respectively.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: VJ1038). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

References

- Benedetti, E., Morelli, G., Nemethy, G. & Scheraga, H. A. (1983). *Int. J. Peptide Protein Res.* **22**, 1–15.
 Enraf–Nonius (1979). *Structure Determination Package*. Enraf–Nonius, Delft, The Netherlands.
 Motherwell, W. D. S. & Clegg, W. (1978). *PLUTO. Program for Plotting Molecular and Crystal Structures*. University of Cambridge, England.
 Nardelli, M. (1983). *Comput. Chem.* **7**, 95–98.
 North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). *Acta Cryst.* **A24**, 351–359.
 Ramachandran, G. N. & Sasisekharan, V. (1968). *Advances in Protein Chemistry*, Vol. 23, p. 238. New York: Academic Press.
 Sheldrick, G. M. (1976). *SHELX76. Program for Crystal Structure Determination*. University of Cambridge, England.
 Sheldrick, G. M. (1985). *SHELXS86. Program for the Solution of Crystal Structures*. University of Göttingen, Germany.
 Vickovic, I. (1994). *J. Appl. Cryst.* **27**, 437.
 Winkler, F. K. & Dunitz, J. D. (1971). *J. Mol. Biol.* **59**, 169–182.

Table 2. Selected torsion angles ($^\circ$) involving non-H atoms

N1—C1A—C1'—N2	(ψ_1)	137.3 (3)
C1A—C1'—N2—C2A	(ω_1)	174.4 (3)
N1—C1A—C1B—C1G	($^1\chi_1$)	175.6 (4)
C1A—C1B—C1G—C1D1	($^1\chi_{21}$)	59.1 (7)
C1A—C1B—C1G—C1D2	($^1\chi_{22}$)	-176.1 (5)
C1'—N2—C2A—C2'	(φ_2)	-151.8 (4)
N2—C2A—C2'—O21'	(ψ_{21})	171.9 (3)
N2—C2A—C2'—O22'	(ψ_{22})	-11.8 (5)

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Dalspinosin

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Abstract

The title compound, 3-(3,4-dimethoxyphenyl)-5,7-dihydroxy-6-methoxy-4H-1-benzopyran-4-one, $\text{C}_{18}\text{H}_{16}\text{O}_7$, consists of two phenyl rings (*A* and *B*) and a heterocyclic ring *C*. Rings *A* and *B* are planar and ring *C* is slightly puckered. The packing of the molecules in the unit cell is governed by van der Waals interactions and hydrogen bonds.

Comment

Dalspinosin (**I**) is an isoflavone derivative having a unique 3',4' arrangement of the methoxy groups in ring *B*. Isoflavonoids have oestrogenic, insecticidal, pesticidal and antifungal properties (Harborne, Mabry & Mabry, 1975). Fig. 1 is a perspective view of the molecular geometry showing numbering scheme adopted.

Table 3. Hydrogen-bonding geometry (\AA , $^\circ$)

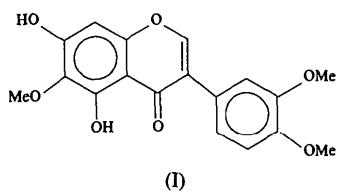
$D—H \cdots A$	$D—H$	$H \cdots A$	$D \cdots A$	$D—H \cdots A$
N1—H1—O21''	1.02 (7)	1.74 (7)	2.737 (5)	163 (5)
N1—H2—O3''	0.79 (7)	2.02 (7)	2.788 (6)	164 (7)
N1—H3—O22'''	0.69 (7)	2.21 (7)	2.875 (5)	161 (7)
N2—H6—O22'	0.79 (7)	2.33 (6)	2.654 (5)	106 (6)

Symmetry codes: (i) $-x, y - \frac{1}{2}, \frac{3}{2} - z$; (ii) $-\frac{1}{2} - x, 1 - y, \frac{1}{2} + z$; (iii) $-1 - x, y - \frac{1}{2}, \frac{3}{2} - z$.

H atoms of terminal groups in side chains showed abnormal B_j 's and hence were only included in the structure-factor calculations. The other H atoms in the peptide molecule were refined isotropically.

Data collection: *SDP* (Enraf–Nonius, 1979). Cell refinement: *SDP*. Data reduction: *SDP*. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELX76* (Sheldrick, 1978). Molecular graphics: *PLUTO* (Motherwell & Clegg, 1976); *ORTEP* (Vickovic, 1994). Software used to prepare material for publication: *PARST* (Nardelli, 1983).

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The two phenyl rings (*A* and *B*) are planar. The dihedral angle between rings *A* and *B* is 31.55 (6) $^{\circ}$. Ring *C* is slightly puckered with $q_2 = 0.070$ (2) \AA , $\theta_2 = 94$ (1) $^{\circ}$ and $\varphi_2 = -109$ (2) $^{\circ}$ (Cremer & Pople, 1975) and ring *A* is twisted from the mean plane of ring *C* by 4.62 (5) $^{\circ}$. The methoxy groups at C6, C3' and C4' are oriented slightly out of the plane of the rings to which they are attached (Shoja, 1992).

The mean bond lengths averaged over each type of bond agree well with the values observed in similar compounds (Kaneda, Itaka & Shibata 1973; Breton, Precigoux, Courseille & Hospital, 1975). The structure is stabilized by van der Waals interactions and O—H \cdots O and C—H \cdots O hydrogen bonds (Desiraju, 1991).

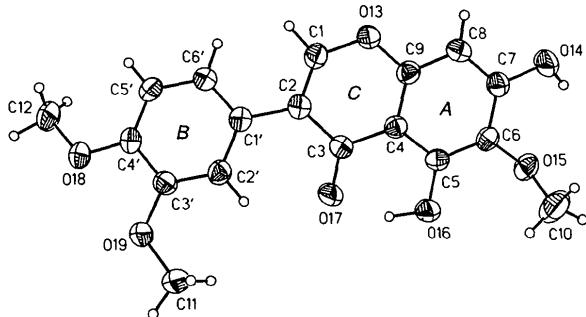


Fig. 1. Molecular structure showing 50% probability displacement ellipsoids.

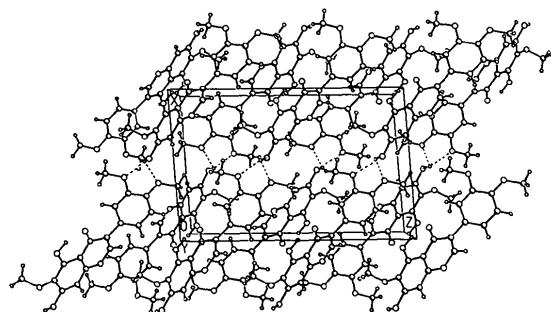


Fig. 2. Packing in the crystal viewed down the *b* axis.

Experimental

The compound was isolated from extracts of the roots of the plant *Dalbergia Spinoso*. Single crystals were obtained by slow evaporation of a methanol solution.

Crystal data

C ₁₈ H ₁₆ O ₇	Mo K α radiation
$M_r = 344.31$	$\lambda = 0.71073 \text{ \AA}$
Monoclinic	Cell parameters from 23 reflections
$P2_1/c$	$\theta = 20\text{--}34^{\circ}$
$a = 10.752$ (1) \AA	$\mu = 0.114 \text{ mm}^{-1}$
$b = 8.711$ (2) \AA	$T = 293 \text{ K}$
$c = 16.646$ (1) \AA	Rectangular
$\beta = 94.4$ (1) $^{\circ}$	$0.1 \times 0.1 \times 0.05 \text{ mm}$
$V = 1554.5$ (4) \AA^3	Pale yellow
$Z = 4$	
$D_x = 1.471 \text{ Mg m}^{-3}$	
$D_m = 1.472 \text{ Mg m}^{-3}$	

Data collection

Enraf-Nonius CAD-4 diffractometer	$R_{\text{int}} = 0.0307$
$\omega/2\theta$ scans	$\theta_{\text{max}} = 25^{\circ}$
Absorption correction:	$h = 0 \rightarrow 12$
none	$k = 0 \rightarrow 10$
2888 measured reflections	$l = -19 \rightarrow 19$
2732 independent reflections	2 standard reflections
2116 observed reflections	frequency: 60 min
	intensity decay: <2%
	[$I > 2\sigma(I)$]

Refinement

Refinement on F^2	$(\Delta/\sigma)_{\text{max}} = 0.075$
$R(F) = 0.0399$	$\Delta\rho_{\text{max}} = 0.198 \text{ e \AA}^{-3}$
$wR(F^2) = 0.0998$	$\Delta\rho_{\text{min}} = -0.225 \text{ e \AA}^{-3}$
$S = 1.048$	Extinction correction: none
2732 reflections	Atomic scattering factors
290 parameters	from International Tables for Crystallography (1992, Vol. C) Tables 4.2.6.8 and 6.1.1.4
H atoms refined isotropically	where $P = (F_o^2 + 2F_c^2)/3$
$w = 1/[\sigma^2(F_o^2) + (0.0612P)^2 + 0.5485P]$	

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (\AA^2)

	x	y	z	B_{eq}
C1	0.9235 (2)	0.0711 (2)	0.1484 (1)	3.17 (5)
C2	1.0056 (1)	-0.0454 (2)	0.1484 (1)	2.71 (5)
C3	0.9891 (1)	-0.1579 (2)	0.0831 (1)	2.63 (4)
C4	0.8777 (1)	-0.1415 (2)	0.0294 (1)	2.54 (4)
C5	0.8458 (1)	-0.2495 (2)	-0.0323 (1)	2.64 (5)
C6	0.7348 (1)	-0.2362 (2)	-0.0792 (1)	2.71 (5)
C7	0.6527 (1)	-0.1154 (2)	-0.0652 (1)	2.87 (5)
C8	0.6823 (2)	-0.0068 (2)	-0.0062 (1)	3.16 (5)
C9	0.7936 (1)	-0.0215 (2)	0.0396 (1)	2.80 (5)
C1'	1.1054 (1)	-0.0600 (2)	0.2149 (1)	2.65 (5)
C2'	1.2204 (1)	-0.1303 (2)	0.2036 (1)	2.77 (5)
C3'	1.3097 (1)	-0.1477 (2)	0.2671 (1)	2.82 (5)
C4'	1.2878 (1)	-0.0917 (2)	0.3439 (1)	2.78 (5)
C5'	1.1763 (1)	-0.0208 (2)	0.3549 (1)	2.99 (5)
C6'	1.0857 (1)	-0.0059 (2)	0.2912 (1)	2.96 (5)
C10	0.7568 (3)	-0.3376 (4)	-0.2088 (1)	5.49 (9)
C11	1.4473 (2)	-0.2868 (3)	0.1868 (1)	4.34 (7)
C12	1.3536 (2)	-0.0844 (3)	0.4832 (1)	4.04 (7)
O13	0.8206 (1)	0.0883 (1)	0.0974 (1)	3.45 (4)
O14	0.5432 (1)	-0.1013 (2)	-0.1096 (1)	3.83 (4)
O15	0.6951 (1)	-0.3435 (1)	-0.1363 (1)	3.49 (3)
O16	0.9225 (1)	-0.3685 (1)	-0.0444 (1)	3.64 (4)
O17	1.0638 (1)	-0.2649 (1)	0.0742 (1)	3.81 (4)
O18	1.3805 (1)	-0.1183 (1)	0.4031 (1)	3.49 (4)
O19	1.4220 (1)	-0.2199 (2)	0.2623 (1)	3.82 (4)

Table 2. Selected geometric parameters (\AA)

C1—C2	1.345 (3)	C8—C9	1.374 (3)
C1—O13	1.350 (3)	C9—O13	1.372 (2)
C2—C3	1.464 (3)	C1'—C2'	1.405 (3)
C2—C1'	1.486 (3)	C1'—C6'	1.387 (3)
C3—C4	1.446 (3)	C2'—C3'	1.380 (3)
C3—O17	1.247 (3)	C3'—C4'	1.405 (3)
C4—C5	1.415 (3)	C3'—O19	1.370 (3)
C4—C9	1.401 (3)	C4'—C5'	1.373 (3)
C5—C6	1.381 (3)	C4'—O18	1.366 (2)
C5—O16	1.349 (3)	C5'—C6'	1.390 (3)
C6—C7	1.404 (3)	C10—O15	1.422 (4)
C6—O15	1.377 (2)	C11—O19	1.429 (3)
C7—C8	1.384 (3)	C12—O18	1.417 (3)
C7—O14	1.347 (2)		

Table 3. Hydrogen-bonding geometry (\AA , $^\circ$)

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
C2'—H2'...O17	0.94 (2)	2.32 (2)	2.880 (2)	118 (2)
C10—H10A...O16	0.95 (4)	2.64 (4)	3.159 (3)	115 (3)
O14—H14...O15	0.86 (4)	2.28 (4)	2.725 (2)	113 (3)
O16—H16...O17	1.00 (3)	1.64 (3)	2.562 (2)	150 (3)
C8—H8...O14'	0.88 (3)	2.68 (3)	3.346 (3)	133 (2)
C5'—H5'...O16"	0.94 (2)	2.73 (2)	3.655 (3)	170 (2)
C12—H12B...O15 ⁱⁱ	0.97 (2)	2.77 (3)	3.373 (3)	121 (2)
C11—H11B...O15 ⁱⁱⁱ	1.03 (3)	2.75 (3)	3.636 (3)	145 (2)
O16—H16...O16 ⁱⁱⁱⁱ	1.00 (3)	2.64 (3)	3.135 (2)	110 (2)
C11—H11C...O14 ^v	1.00 (3)	2.72 (3)	3.621 (4)	150 (2)
O14—H14...O18 ^v	0.86 (4)	2.46 (4)	3.022 (2)	123 (3)
O14—H14...O19 ^v	0.86 (4)	2.13 (3)	2.871 (2)	145 (2)

Symmetry codes: (i) $1 - x, -y, -z$; (ii) $2 - x, \frac{1}{2} + y, \frac{1}{2} - z$; (iii) $2 - x, 1 - y, -z$; (iv) $2 - x, -y, -z$; (v) $x - 1, -\frac{1}{2} - y, z - \frac{1}{2}$.

The structure was solved by direct methods using *SHELXS86* (Sheldrick, 1985). The initial *R* factor for the model proposed was 0.19. After a few cycles of full-matrix least-squares refinement, the *R* factor reduced to 0.11. All H atoms were located from the difference Fourier map and were refined isotropically.

Data collection: *Enraf–Nonius CAD-4 Software* (Enraf–Nonius, 1989). Cell refinement: *SDP* (Frenz, 1978). Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *ORTEP* (Johnson, 1965). Software used to prepare material for publication: *PARST* (Nardelli, 1983).

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: DE1027). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

References

- Breton, M., Precigoux, G., Courseille, Ch. & Hospital, M. (1975). *Acta Cryst.* **B31**, 921–923.
 Cremer, D. & Pople, J. A. (1975). *J. Am. Chem. Soc.* **97**, 1354–1359.
 Desiraju, G. R. (1991). *Acc. Chem. Res.* **24**, 290–296.
 Enraf–Nonius (1989). *CAD-4 Software*. Version 5.0. Enraf–Nonius, Delft, The Netherlands.
 Frenz, B. A. (1978). *The Enraf–Nonius CAD-4 SDP – a Real-Time System for Concurrent X-ray Data Collection and Crystal Structure Solution. Computing in Crystallography*, edited by H. Schenk, R. Olthof-Hazekamp, H. van Koningsveld & G. C. Bassi, pp. 64–71. Delft University Press.
 Harborne, J. B., Mabry, T. J. & Mabry, H. (1975). In *The Flavonoids*. London: Chapman & Hall.

- Johnson, C. K. (1965). *ORTEP*. Report ORNL-3794. Oak Ridge National Laboratory, Tennessee, USA.
 Kaneda, M., Itaka, Y. & Shibata, S. (1973). *Acta Cryst.* **B29**, 2827–2832.
 Nardelli, M. (1983). *Comput. Chem.* **7**, 95–96.
 Sheldrick, G. M. (1985). *SHELXS86. Program for the Solution of Crystal Structures*. University of Göttingen, Germany.
 Sheldrick, G. M. (1993). *SHELXL93. Program for the Refinement of Crystal Structures*. University of Göttingen, Germany.
 Shoja, M. (1992). *Acta Cryst.* **C48**, 2033–2035.

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Acetoxymethyl 4-Chloro-N-furfuryl-5-sulfamoylanthranilate, an Absorption Furosemide Prodrug

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Abstract

The title compound, $C_{15}H_{15}ClN_2O_7S$, which was synthesized and characterized as the acetoxymethyl ester of 4-chloro-N-furfuryl-5-sulfamoylanthranilic acid (furosemide) is an absorption furosemide prodrug. The molecule crystallized in a triclinic unit cell, space group $P\bar{1}$. The crystal structure is stabilized by one intramolecular and two intermolecular hydrogen bonds.

Comment

Furosemide is a strong diuretic agent widely used in hypertension crisis. The use of some acyloxymethyl esters of furosemide as prodrugs to improve the therapeutic success of this drug has been studied by Prandi, Fagiolino, Manta & Llera (1992).

