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Hydrogen bonding and π - π stacking in dimethylgenistein

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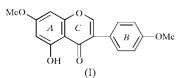
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The title compound, 5-hydroxy-4',7-dimethoxyisoflavone, $C_{17}H_{14}O_5$, is composed of a benzopyranone moiety, a phenyl moiety and two methoxy groups. The benzopyranone ring is not coplanar with the phenyl ring, the dihedral angle between them being 56.28 (3)°. The two methoxy groups are nearly coplanar with their corresponding rings, having C-C-O-C torsion angles of 2.9 (2) and 5.9 (2)°. The molecules are linked by C-H···O hydrogen bonds into sheets containing classical centrosymmetric $R_2^2(8)$ rings. The sheets are further linked by aromatic π - π stacking interactions and C-H···O hydrogen bonds into a supramolecular structure.

Comment

Hydrogen bonds and $\pi-\pi$ stacking interactions are an important research area in supramolecular chemistry and crystal engineering (MacDonald & Whitesides, 1994). These interactions play an important role in self-assembly and recognition of aromatic compounds (Janiak, 2000; Hunter & Sanders, 1990) as auxiliary stabilizing short contacts (William *et al.*, 1999; Luque *et al.*, 2001; Kaafarani *et al.*, 2001). In biomacromolecular systems, stacking interactions and hydrogen bonds are important for the double-helical DNA structure (Hunter, 1993); they can direct the intercalation of drugs into DNA (Wang *et al.*, 1984) and they contribute to the stability of the tertiary structure of proteins (Burley & Petsko, 1985).



Genistein, a natural soy isoflavone, has potential phytoestrogen (Hua *et al.*, 2003; Warren, 2002) and antioxidant activities (Ian *et al.*, 1995). Studies have also found genistein effective in inhibiting cardiovascular disease (Hwang *et al.*, 2001), tyrosine kinases (Nevala *et al.*, 2002) and cancer cell growth (Yuan *et al.*, 2003; W. F. Chen *et al.*, 2003), and in accelerating the formation of bone cells (X. W. Chen *et al.*, 2003). The title compound, namely 5-hydroxy-4',7-dimethoxyisoflavone, (I), is a derivative of genistein and has potential medical applications. We report here the crystal structure of (I).

The title compound is composed of a benzopyranone moiety, a phenyl moiety and two methoxy groups (Fig. 1). The geometry of the isoflavone skeleton of (I) is similar to that of its analogue dalspinin (Lakshmi, *et al.*, 1996) with respect to most of the bond distances and angles. The atoms of the benzopyranone moiety, composed of rings A (C1–C6) and C (O1/C1/C6–C9), are almost coplanar, the dihedral angle between the rings being 1.37 (8)°. To avoid steric conflicts, the two rigid ring systems, *viz.* benzene ring B (C10–C15) and the benzopyranone moiety, are rotated by 56.28 (3)° with respect to one another. The methoxy group at atom C3 is nearly coplanar with ring A, as indicated by the C16–O4–C3–C2 torsion angle [2.9 (2)°]; the methoxy group at atom C13 is also coplanar with the attached ring, the C17–O5–C13–C12 torsion angle being 5.9 (2)°.

Fig. 2 shows how a cyclic dimer is formed through a supramolecular synthon, $R_2^2(8)$. Methoxy atom O4 acts as a hydrogen-bond acceptor, via atom H4, to atom C4 of ring A. In this manner, a centrosymmetric $R_2^2(8)$ ring is formed. Hydroxy atoms O2 from the two molecules linked by the $R_2^2(8)$ ring act as hydrogen-bond acceptors, via atoms H14, to atoms C14 of rings B in adjacent molecules. The combination of the C14-H14···O2 interaction and the $R_2^2(8)$ supramolecular synthon generates a $(10\overline{1})$ sheet, which includes two A, two B and two C rings from four molecules, and these six rings are almost coplanar; furthermore, these dimers are also linked into (100) chains by $C11 - H11 \cdots O3$ interactions (Fig. 3). The combination of the $(10\overline{1})$ sheets and the (100) chains generates a three-dimensional framework. An independent O2-H2O···O3 intramolecular hydrogen bonds generates a characteristic intramolecular S(6) motif. Details of the hydrogen bonding are given in Table 1.

Intermolecular stacking *via* aromatic π - π interactions is also present (Fig. 3), the two molecules being offset by partial overlap of rings *B* (π rich) and *C* (π deficient). Ring *B* of one

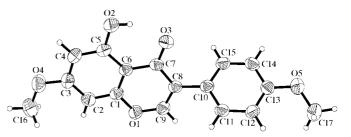


Figure 1

A view of the molecule of (I), showing the atom-numbering scheme and 50% probability displacement ellipsoids. H atoms are shown as spheres of arbitrary radii.

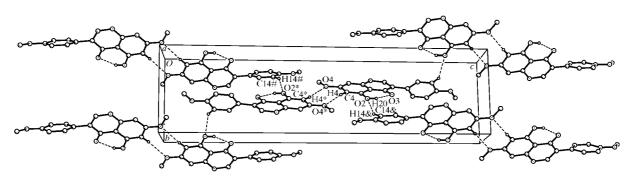


Figure 2

Part of the crystal structure of (I), showing the formation of the $(10\overline{1})$ sheets *via* hydrogen bonds. Atoms marked with an asterisk (*), hash (#) or ampersand (&) are at the symmetry positions (1 - x, 1 - y, 1 - z), $(x - \frac{1}{2}, \frac{1}{2} - y, z - \frac{1}{2})$ and $(\frac{3}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z)$, respectively. For clarity, some H atoms have been omitted.

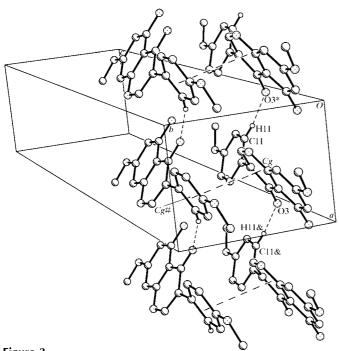


Figure 3

Part of the crystal structure of (I), showing the formation of the (100) chains *via* hydrogen bonds and π - π stacking interactions. Labels *Cg* represent the centroids of rings *A* and *C*. Atoms marked with an asterisk (*), hash (#) or ampersand (&) are at the symmetry positions (1 - x, y, z), $(\frac{1}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z)$ and (1 + x, y, z), respectively. For clarity, some H atoms have been omitted.

molecule and ring *C* of a neighbouring molecule are almost parallel, with a dihedral angle between them of 8.02 (7)°. The perpendicular plane-to-plane distance between the rings is 3.311 Å, and the corresponding $Cg \cdots Cg^{\#}$ distance is 3.693 Å [*Cg* represents the centroids of rings *A* and *C*; symmetry code: (#) $\frac{1}{2} - x$, $\frac{1}{2} + y$, $\frac{3}{2} - z$], indicating that a strong $\pi - \pi$ stacking interaction exists in the title compound. Hydrogen bonds and aromatic $\pi - \pi$ stacking interactions play a key role in assembling the supramolecular structure.

Experimental

Genistein (1.0 g) was dissolved in Na₂CO₃ (20 ml, 5%) and dimethyl sulfate (0.5 ml) was added dropwise to the solution with stirring. The

mixture was stirred for 4 h at room temperature and a colourless precipitate began to appear. The precipitate was filtered off and washed with water until the pH of the filtrate was 8. After recrys-tallization from ethyl acetate, the product had a melting point of 428 K. Crystals of (I) suitable for X-ray analysis were obtained by slow evaporation from ethyl acetate after 7 d at room temperature.

 $D_x = 1.441 \text{ Mg m}^{-3}$

Cell parameters from 34

Mo $K\alpha$ radiation

reflections

 $\theta = 2.7 - 15.9^{\circ}$ $\mu = 0.11 \text{ mm}^{-1}$

T = 296 (2) K

 $h = -6 \rightarrow 6$

 $l = -36 \rightarrow 36$

 $k = 0 \rightarrow 9$

Prism, colourless

 $0.58\,\times\,0.54\,\times\,0.50\;\mathrm{mm}$

3 standard reflections

every 97 reflections

intensity decay: 0.4%

Crystal data

 $C_{17}H_{14}O_5$ $M_r = 298.28$ Monoclinic, P_{21}/n a = 5.7754 (8) Å b = 7.9446 (12) Å c = 30.044 (5) Å $\beta = 93.807$ (12)° V = 1375.4 (4) Å³ Z = 4

Data collection

Siemens *P*4 diffractometer ω scans 3089 measured reflections 2485 independent reflections 1673 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.010$ $\theta_{\text{max}} = 25.3^{\circ}$

Refinement

 Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0522P)^2]$
 $R[F^2 > 2\sigma(F^2)] = 0.035$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.087$ $(\Delta/\sigma)_{max} = 0.001$

 S = 0.90 $\Delta\rho_{max} = 0.19 \text{ e Å}^{-3}$

 2485 reflections
 $\Delta\rho_{min} = -0.15 \text{ e Å}^{-3}$

 203 parameters
 Extinction correction: SHELXL97

 H-atom parameters constrained
 Extinction coefficient: 0.0233 (18)

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O2−H2O···O3	0.82	1.86	2.590 (2)	148
$C4-H4\cdots O4^{i}$	0.93	2.54	3.459 (2)	171
$C11 - H11 \cdot \cdot \cdot O3^{ii}$	0.93	2.56	3.464 (2)	164
C14-H14···O2 ⁱⁱⁱ	0.93	2.58	3.399 (2)	147
			, ,	

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

H atoms were placed at calculated positions and treated as riding, with C–H distances in the range 0.93–0.96 Å and $U_{iso}(H)$ values of $1.2U_{eq}$ of the attached C atom $[1.5U_{eq}(C)$ for methyl H atoms].

Data collection: *XSCANS* (Siemens, 1996); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Siemens, 1997); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1473). Services for accessing these data are described at the back of the journal.

References

- Burley, S. K. & Petsko, G. A. (1985). Science, 229, 23-28.
- Chen, W. F., Huang, M. H., Tzang, C. H., Yang, M. S. & Wong, M. S. (2003). Biochim. Biophys. Acta, 1638, 187–196.
- Chen, X. W., Sanford, C. G. L., Quarles, D. & Anderson, J. B. (2003). J. Nutr. Biochem. 14, 342–349.
- Hua, P., Tsai, W. J. & Kuo, S. M. (2003). *Biochim. Biophys. Acta*, **1627**, 63–70. Hunter, C. A. (1993). *J. Mol. Biol.* **230**, 1025–1054.
- Hunter, C. A. & Sanders, J. K. (1990). J. Am. Chem. Soc. 112, 5525-5534.

- Hwang, J. L., Hodis, H. N. & Sevanian, A. (2001). J. Agric. Food Chem. 49, 308–314.
- Ian, R. R., Ivor, D. E. & Jennifer, K. M. (1995). J. Nutr. Biochem. 6, 481–485. Janiak, C. (2000). J. Chem. Soc. Dalton Trans. pp. 3885–3896.
- Kaafarani, B. R., Pinkerton, A. A. & Neckers, D. C. (2001). *Tetrahedron Lett.* 42, 8137–8139.
- Lakshmi, S., Kumar, S., Senthilselvan, J. & Subramanian, K. (1996). Acta Cryst. C52, 2873–2875.
- Luque, A., Sertucha, J., Castillo, O. & Roman, P. (2001). New J. Chem. pp. 1208–1214.
- MacDonald, J. C. & Whitesides, G. M. (1994). Chem. Rev. 94, 2383-2420.
- Nevala, R., Lassila, M., Finckenberg, P., Paukku, K., Korpela, R. & Vapaatalo, H. (2002). Eur. J. Pharmacol. 452, 87–96.
- Sheldrick, G. M. (1990). Acta Cryst. A46, 467-473.
- Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany.
- Siemens (1996). XSCANS. Version 2.21. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
- Siemens (1997). SHELXTL. Version 5.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Wang, A. H., Ughetto, G., Quigley, G. J., Hakoshima, T., van der Marel, G. A., van Boom, J. H. & Rich, A. (1984). *Science*, 225, 1115–1121.
- Warren, N. R. (2002). Life Sci. 71, 865-877.
- William, O. G., Barbour, L. J. & Atwood, J. L. (1999). Science, 285, 1049–1051.
- Yuan, L., Wagatsuma, C., Yoshida, M., Miura, T., Mukoda, T., Fujii, H., Sun, B. X., Kim, J. H. & Surh, Y. J. (2003). *Mutat. Res.* 523–524, 55–62.