

1-(4-Hydroxyphenyl)-3-(2,4,6-trimethoxyphenyl)propan-1-one

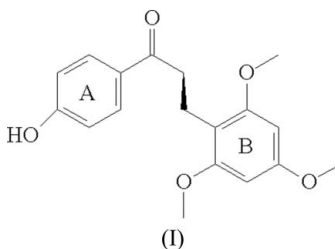
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Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.076
 wR factor = 0.184
Data-to-parameter ratio = 8.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.The two benzene rings of the title compound, $\text{C}_{18}\text{H}_{20}\text{O}_5$, are almost parallel, with a dihedral angle of $1.3(2)^\circ$. The hydroxy group forms intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds which link the molecules into infinite chains along the c axis.

Comment

The title compound, (I), also known as loureirin B, was extracted from the herb *Sanguis draxonis* with ethanol (Zhou *et al.*, 2001). The compound was crystallized from methanol:toluene (3:1) and its structure is reported here. The dihedral angle between the planes defined by atoms C1–C6 and C1'–C6' is $1.3(2)^\circ$. The hydroxyphenyl and trimethoxyphenyl ring planes are separated by 1.414 Å, and the C7=O2 carbonyl group is essentially coplanar with the hydroxyphenyl ring. The hydroxy group forms a strong $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond to the carbonyl O atom, linking the molecules into infinite chains along the c axis.

Experimental

The title compound was extracted according to the procedure of Zhou *et al.* (2001) from the herb *Sanguis draxonis*. The compound was crystallized from methanol/toluene (3:1), yielding colorless block-like crystals after a week at room temperature.

Crystal data

 $\text{C}_{18}\text{H}_{20}\text{O}_5$
 $M_r = 316.34$
Monoclinic, Pc
 $a = 5.6200(11)$ Å
 $b = 14.447(3)$ Å
 $c = 10.149(2)$ Å
 $\beta = 96.04(3)^\circ$
 $V = 819.4(3)$ Å³
 $Z = 2$ $D_x = 1.282$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 1803
reflections
 $\theta = 4.2\text{--}27.2^\circ$
 $\mu = 0.09$ mm⁻¹
 $T = 298(2)$ K
Block, colorless
 $0.30 \times 0.20 \times 0.20$ mm

Data collection

MAC DIP 2030K diffractometer
 ω scans
Absorption correction: none
3054 measured reflections
1803 independent reflections
1778 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.029$
 $\theta_{\text{max}} = 27.2^\circ$
 $h = 0 \rightarrow 7$
 $k = -18 \rightarrow 18$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.076$
 $wR(F^2) = 0.184$
 $S = 1.19$
 1803 reflections
 209 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0669P)^2 + 0.3852P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.20 \text{ e } \text{\AA}^{-3}$
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.84 (7)

Table 1
 Selected geometric parameters (\AA , $^\circ$).

| | | | |
|--------------|------------|----------------|-----------|
| O1—C4 | 1.347 (6) | C1—C2 | 1.392 (7) |
| O2—C7 | 1.236 (6) | C7—C8 | 1.498 (7) |
| O3—C6' | 1.366 (6) | C8—C9 | 1.537 (7) |
| O3—C7' | 1.428 (6) | C9—C1' | 1.494 (7) |
| C6'—O3—C7' | 119.2 (4) | C1'—C9—C8 | 113.2 (4) |
| C1—C7—C8 | 120.3 (4) | | |
| O1—C4—C5—C6 | −179.1 (5) | C8—C9—C1'—C6' | 83.9 (6) |
| C2—C1—C7—C8 | −177.1 (5) | C9'—O5—C2'—C3' | −2.2 (8) |
| C7—C8—C9—C1' | 177.5 (5) | | |

Table 2
 Hydrogen-bond geometry (\AA , $^\circ$).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------------|-------|-------------|-------------|---------------|
| O1—H1A \cdots O2 ⁱ | 0.82 | 1.89 | 2.708 (5) | 179 |

Symmetry code: (i) $x + 1, -y + 1, z - \frac{1}{2}$.

All H atoms were refined using a riding model with C—H = 0.93 \AA , $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic CH; 0.97 \AA , $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for CH₂; 0.96 \AA , $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for CH₃; and 0.82 \AA , $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ for OH atoms. In the absence of significant anomalous dispersion effects, Friedel pairs were merged.

Data collection: *DENZO* (Otwinowski & Minor, 1997); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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References

Otwinowski, Z. & Minor, W (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.

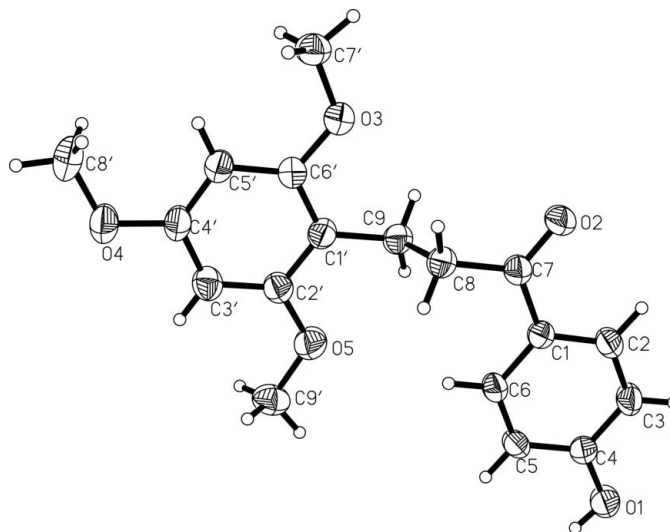


Figure 1
 A view of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented by circles of arbitrary size.

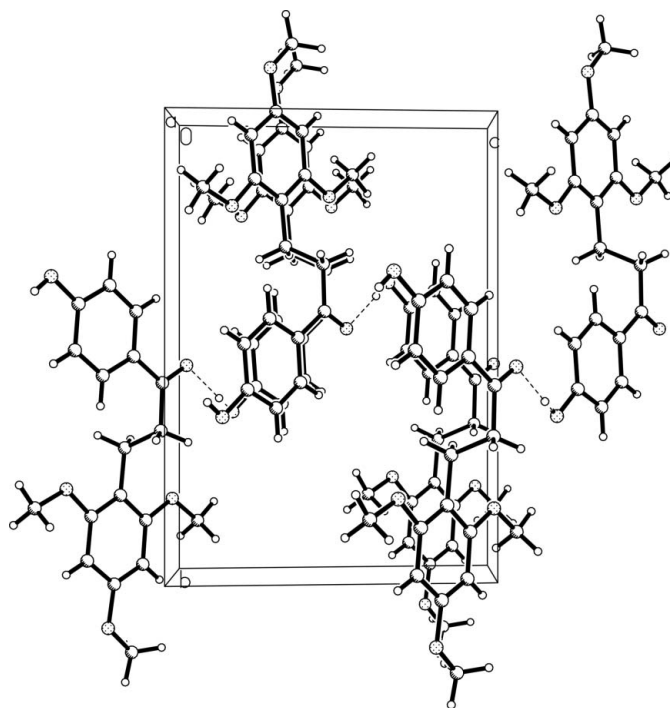


Figure 2
 The molecular packing of (I), viewed along the a axis. Dashed lines indicate hydrogen-bonding interactions.

Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.
 Zhou, Z. H., Wang, J. L. & Yang, C. R. (2001). *Chin. Tradit. Herb. Drugs*, **32**, 484–486.