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Key indicators

Single-crystal X-ray study

T = 273 K

Mean $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$

R factor = 0.053

wR factor = 0.151

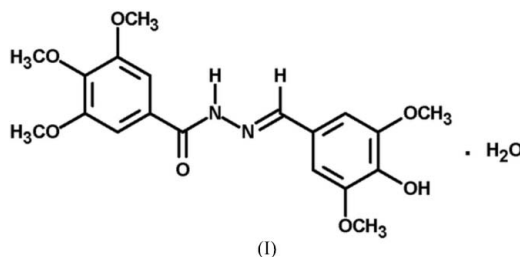
Data-to-parameter ratio = 17.6

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.4-Hydroxy-3,5-dimethoxybenzaldehyde 3,4,5-tri-
methoxybenzoylhydrazone monohydrateThe title compound, $\text{C}_{19}\text{H}_{22}\text{N}_2\text{O}_7 \cdot \text{H}_2\text{O}$, was synthesized as part
of a continuing project involving the structures of hydrazone
derivatives. The molecules pack in a three-dimensional
framework structure by a combination of $\text{O}-\text{H} \cdots \text{O}$, $\text{O}-$
 $\text{H} \cdots \text{N}$ and $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds.

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Comment

Molecules containing hydrazone units are of interest to us
(Sun *et al.*, 2006) owing to their broad spectrum of activities
and applications (Gup & Kirkan, 2005; Ganjali *et al.*, 2006;
Getautis *et al.*, 2006). Moreover, many hydrazones have also
been used as ligands for complexation of metal ions (Kuria-
kose *et al.*, 2007).Compound (I) possesses normal geometric parameters
(Fig. 1) (Allen *et al.*, 1987). The central hydrazone component
is effectively planar with an all-*trans* extended conformation
(Table 1). On the other hand, the two benzene rings form an
angle of $19.1 (1)^\circ$ to one another and the C11 benzene ring is
slightly twisted out of the plane of the central hydrazone unit.
The existence of (I) in the expected keto form is evident from
the C10—O4 bond length of $1.231 (2) \text{ \AA}$. Of the five methoxy
groups only O2/C7 is coplanar with one of the benzene rings,
with the others showing out-of-plane twists ranging from ~ 15
to 85° (Table 1). Similar geometry has been observed in
related hydrazone analogues (Peralta *et al.*, 2007; Raj &
Kurup, 2007).The molecules are linked into a three-dimensional frame-
work structure by a combination of six independent hydrogen
bonds (Table 2, Fig. 2). The water molecule plays a significant
role in the crystal packing as it is the acceptor in one hydrogen
bond and a donor in three others, including one bifurcated
bond (Table 2).

Experimental

A mixture of 4-hydroxy-3,5-dimethoxybenzaldehyde (1 mmol) and
3,4,5-trimethoxybenzoylhydrazide (1 mmol) in anhydrous ethanol
(30 ml) was refluxed for 5 h and then cooled to room temperature.

The precipitate was filtered off and dried. The crude product was recrystallized from ethanol. Colorless crystals were obtained in 76% yield. Analysis calculated for $C_{19}H_{24}N_2O_7 \cdot H_2O$: C 55.88, H 5.92, N 6.86%; found: C 55.83, H 5.95, N 6.92%. A single crystal suitable for an X-ray structural analysis was obtained by slowly evaporating an ethanol solution at room temperature.

Crystal data

| | |
|---------------------------------|---|
| $C_{19}H_{24}N_2O_7 \cdot H_2O$ | $V = 1960.59 (14) \text{ \AA}^3$ |
| $M_r = 408.40$ | $Z = 4$ |
| Monoclinic, $P2_1/c$ | Mo $K\alpha$ radiation |
| $a = 10.1299 (4) \text{ \AA}$ | $\mu = 0.11 \text{ mm}^{-1}$ |
| $b = 13.8108 (6) \text{ \AA}$ | $T = 273 (2) \text{ K}$ |
| $c = 14.8758 (6) \text{ \AA}$ | $0.35 \times 0.24 \times 0.12 \text{ mm}$ |
| $\beta = 109.598 (2)^\circ$ | |

Data collection

| | |
|--|--|
| Bruker SMART CCD area-detector diffractometer | 26378 measured reflections |
| Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996) | 4755 independent reflections |
| $T_{\min} = 0.963$, $T_{\max} = 0.987$ | 2463 reflections with $I > 2\sigma(I)$ |
| | $R_{\text{int}} = 0.071$ |

Refinement

| | |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.053$ | 270 parameters |
| $wR(F^2) = 0.151$ | H-atom parameters constrained |
| $S = 1.03$ | $\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$ |
| 4755 reflections | $\Delta\rho_{\text{min}} = -0.22 \text{ e \AA}^{-3}$ |

Table 1

Selected torsion angles ($^\circ$).

| | | | |
|---------------|--------------|----------------|------------|
| C9–N1–N2–C10 | 170.63 (18) | C1–C2–O2–C7 | –175.5 (2) |
| N1–N2–C10–C11 | –173.93 (16) | N2–C10–C11–C16 | –15.4 (3) |
| C8–O3–C6–C5 | 33.3 (3) | C16–C15–O7–C19 | 15.0 (4) |
| N2–N1–C9–C4 | –177.10 (18) | C14–C13–O5–C17 | 95.6 (3) |
| C5–C4–C9–N1 | –176.0 (2) | C13–C14–O6–C18 | 84.6 (3) |

Table 2

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

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|-----------------------------------|-------|--|--------------|---|
| O1–H1 \cdots O8 | 0.85 | 1.94 | 2.727 (2) | 154 |
| O1–H1 \cdots O3 | 0.85 | 2.26 | 2.704 (2) | 113 |
| N2–H2 \cdots O1 ⁱ | 0.86 | 2.49 | 3.313 (2) | 160 |
| O8–H17 \cdots O4 ⁱⁱ | 0.85 | 2.19 | 3.033 (2) | 171 |
| O8–H18 \cdots O4 ⁱⁱⁱ | 0.85 | 2.17 | 2.956 (2) | 154 |
| O8–H18 \cdots N1 ⁱⁱⁱ | 0.85 | 2.50 | 3.143 (2) | 133 |
| Symmetry codes: | (i) | $x, -y + \frac{3}{2}, z + \frac{1}{2}$ | (ii) | $x - 1, -y + \frac{3}{2}, z - \frac{1}{2}$ |
| | | | (iii) | $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$ |

All H atoms were initially located in a difference Fourier map and were treated as riding atoms, with C–H = 0.93 (C_{sp^2}) and 0.96 \AA (methyl), N–H = 0.86 \AA , and O–H = 0.85 \AA , and with $U_{\text{iso}}(\text{H}) = kU_{\text{eq}}(\text{C,N,O})$, where $k = 1.5$ for the methyl and the hydroxyl groups and $k = 1.2$ for all other H atoms.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT* (Siemens, 1996); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s)

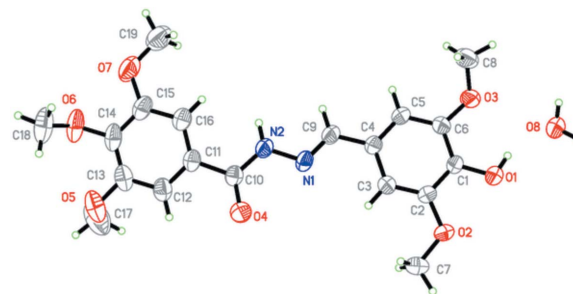


Figure 1

The molecular structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as small spheres of arbitrary radius.

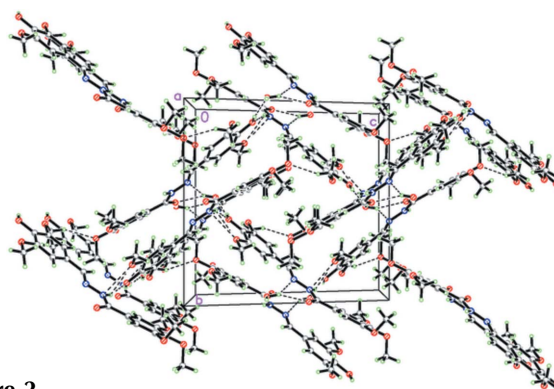


Figure 2

A packing diagram for (I), viewed down the a axis. Dashed lines indicate hydrogen bonds.

used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXTL*.

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