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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.043 wR factor = 0.140 Data-to-parameter ratio = 12.3

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

3-[*N*'-(4-Hydroxy-3-methoxybenzylidene)hydrazinocarbonyl]-1*H*-1,2,4-triazole

The title compound, $C_{11}H_{11}N_5O_3$, was synthesized by the reaction of 3-(1*H*)-1,2,4-triazole hydrazine with 4-hydroxy-3-methoxybenzaldehyde in ethanol. In the crystal structure, intermolecular N-H···O, O-H···N and N-H···N hydrogen bonds are observed. An intramolecular O-H···O hydrogen bond is also found.

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Comment

Azole derivatives, such as derivatives of pyrazole, imidazole, triazole (including benzotriazole), tetrazole, indole *etc.* exhibit extensive biological activities. They have become a central focus in the study of agricultural chemicals, medicine, adjustment reagents for plant growth and so on (Ernest & William, 1982). A Schiff base is a good type of biologically active substructure and a study of a type of triazole Schiff base has been reported (Sauter *et al.*, 1991). The hydrazonecarbonyl grouping has also been shown to be bioactive (Zhi *et al.*, 2003). In a search for more effective antibacterial medicines, we have synthesized the title compound, (I). We have already reported two structures of triazole compounds containing the hydrazonecarbonyl group (Yang & Pan, 2004; Pan & Yang, 2005).



In (I), there are two independent molecules in the asymmetric unit (Fig. 1). The methoxy groups at C9 and C20 are rotated slightly around the C9–O2 and C20–O5 bonds; the torsion angles C11–O2–C9–C10 and C22–O5–C20–C21 are -11.8 (4) and $-6.8(0.4)^{\circ}$, respectively. The bond lengths and angles are unexceptional (Allen *et al.*, 1987). Intermolecular N–H···O, N–H···N and O–H···N hydrogen bonds are observed (Table 2), which result in seven-membered and five-membered hydrogen-bonded rings (Fig. 2). In addition, an intramolecular O–H···O hydrogen bond forming a five-membered ring is observed.

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Experimental

1H-1,2,4-Triazol-3-ylhydrazine (0.02 mol, 2.54 g) was dissolved in 50 ml of anhydrous ethanol at room temperature. 4-Hydroxy-3methoxybenzaldehyde (0.02 mol, 3.04 g) was added and the mixture was refluxed for 2 h; the precipitate was collected by filtration and washed with ethanol. The product was recrystallized from ethanol and dried under reduced pressure to give the title compound. The compound (2.0 mmol, 0.52 g) was dissolved in 30 ml DMF and was kept at room temperature for 30 days. Brown spike-shaped single crystals formed; they were washed with distilled water.

Z = 4

 $D_x = 1.466 \text{ Mg m}^-$ Mo $K\alpha$ radiation

reflections

 $\mu = 0.11 \text{ mm}^{-1}$ T = 293 (2) KSpike, brown $0.40 \times 0.20 \times 0.10 \text{ mm}$

 $\theta = 9.8 - 14.8^{\circ}$

 $R_{\rm int}=0.032$ $\theta_{\rm max} = 25.2^\circ$ $h = -5 \rightarrow 5$ $k = -17 \rightarrow 1$ $l = -21 \rightarrow 21$

+ 0.0981P]

Cell parameters from 25

4271 independent reflections 2357 reflections with $I > 2\sigma(I)$

Crystal data

$C_{11}H_{11}N_5O_3$
$M_r = 261.25$
Triclinic, $P\overline{1}$
a = 4.473 (1) Å
b = 14.772 (5) Å
c = 18.263 (2) Å
$\alpha = 79.38 \ (2)^{\circ}$
$\beta = 88780 \ (1)^{\circ}$
$\gamma = 86.33 \ (2)^{\circ}$
$V = 1183.6 (5) \text{ Å}^3$

Data collection

Bruker SMART four-circle
diffractometer
φ and ω scans
Absorption correction: multi-scan
SADABS (Bruker, 2002)
$T_{\min} = 0.957, T_{\max} = 0.989$
4819 measured reflections

Refinement

 $w = 1/[\sigma^2(F_o^2) + (0.0831P)^2]$ Refinement on F^2
$$\begin{split} R[F^2 > 2\sigma(F^2)] &= 0.043 \\ wR(F^2) &= 0.140 \end{split}$$
where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ S = 1.01 $\Delta \rho_{\rm max} = 0.21 \text{ e} \text{ } \text{\AA}^{-3}$ 4271 reflections $\Delta \rho_{\rm min} = -0.33 \ \rm e \ \AA^{-3}$ 347 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

O1-C3	1.217 (3)	O5-C20	1.363 (3)
O2-C9	1.362 (3)	O5-C22	1.427 (3)
O2-C11	1.432 (3)	O6-C19	1.354 (3)
O3-C8	1.361 (3)	N4-C3	1.343 (3)
O4-C14	1.213 (3)	N10-C15	1.263 (3)
C11-O2-C9-C10	-11.8 (4)	C22-O5-C20-C21	-6.7 (4)

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O3−H3···O2	0.82	2.18	2.642 (2)	116
O3−H3···N8	0.82	2.25	2.868 (3)	133
$O6-H6A\cdots N3^{i}$	0.82	2.00	2.742 (3)	149
$O6-H6A\cdots O5$	0.82	2.22	2.665 (3)	115
$N2-H2\cdot\cdot\cdot N6^{ii}$	0.86	1.97	2.823 (3)	172
$N2-H2\cdots O4^{ii}$	0.86	2.56	3.016 (3)	115
$N4-H4\cdots O6^{i}$	0.86	2.25	3.089 (3)	166
$N7 - H7 \cdot \cdot \cdot O1^{iii}$	0.86	1.96	2.815 (3)	174
N9−H9···O3	0.86	2.08	2.932 (3)	170

Symmetry codes: (i) -x, 2 - y, 1 - z; (ii) x - 3, 1 + y, z; (iii) -1 - x, 2 - y, -z.



Figure 1

The structure of the asymmetric unit of (I), showing the atomic numbering. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

Packing of (I), showing selected inter- and intramolecular hydrogen bonds as dashed lines. Some of the H atoms have been omitted for clarity. (Symmetry codes are as in Table 2.)

All H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of 0.93-0.96 Å (C-H), 0.86 Å (N-H) and 0.82 Å (O-H), with $U_{iso}(H) = 1.2$ or 1.5 times $U_{eq}(parent$ atom).

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT and SHELXTL (Bruker, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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