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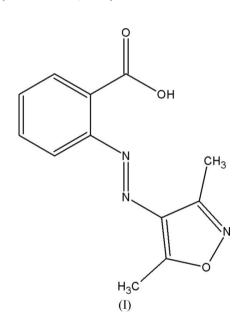
#### **Key indicators**

Single-crystal X-ray study T = 298 K Mean  $\sigma(C-C) = 0.004 \text{ Å}$ R factor = 0.043 wR factor = 0.074 Data-to-parameter ratio = 9.5

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

## Comment

In the past few years, synthesis of isoxazole-related compounds has been the subject of many investigations due to the versatility of their properties. Some uses or properties are anti-inflammatory, analgesic and ulcerogenic (Daidone et al., 1999), antimicrobial and antifungal (Bhatt et al., 1998), inhibition of cyclooxygenase-2 (Talley, 1999; Talley et al., 2000), anticancer activity (Li et al., 2003), selective agonist of dopamine D4 receptors (Rowley et al., 1996) and antagonist of GABA (Frolund et al., 2002).



A perspective view of the title compound is shown in Fig. 1. The dihedral angle between the mean planes formed by the two rings is  $25.20 (15)^\circ$ . The bond lengths of the isoxazole ring (Table 1) are in very good agreement with the usual values for isoxazoles (Allen et al., 1987).

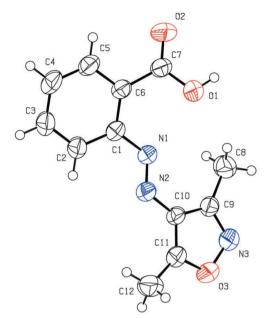
Zigzag chains, which run in the [010] direction (Fig. 2), are formed via an intermolecular  $O-H \cdots N$  hydrogen bond. The structure is further stabilized by  $\pi - \pi$  stacking interactions. The isoxazole ring at (x, y, z) shows two stacking interactions: with the benzene ring at  $(\frac{1}{2} + x, \frac{1}{2} - y, 1 - z)$ , with a distance of 3.9343 (18) Å between the ring centroids, and with that at  $\left(-\frac{1}{2}+x, \frac{1}{2}-y, 1-z\right)$ , with a distance between the ring centroids of 3.6700 (18) Å. Columns are formed along the aaxis via these  $\pi$ - $\pi$  stacking interactions (Fig. 3).

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# Double $\pi$ - $\pi$ stacking in 2-[(*E*)-(3,5-dimethylisoxazol-4-yl)diazenyl]benzoic acid

Molecules of the title compound, C<sub>12</sub>H<sub>11</sub>N<sub>3</sub>O<sub>3</sub>, are linked into zigzag chains by O-H···N hydrogen bonds. The crystal structure is further stabilized by  $\pi - \pi$  stacking interactions.

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#### Figure 1

A view of the molecule of (I), with displacement ellipsoids drawn at the 50% probability level and H atoms drawn as circles of arbitrary size.

# Experimental

In a 100 ml round-bottomed flask were mixed 2-[(2Z)-2-(1-methyl-3oxobutylidene)hydrazino]benzoic acid (0.005 mol), ethanol (25 ml), glacial acetic acid (2 ml) and hydroxylamine hydrochloride (0.005 mol, 0.35 g). The mixture was then stirred and heated under reflux for 18 h. After cooling to room temperature, H<sub>2</sub>O (30 ml) was added, allowing the precipitation of an abundant quantity of orange solid. The product was collected by suction, washed twice with H<sub>2</sub>O and dried under vacuum at 313 K. The crude compound was recrystallized by diffusion of *n*-hexane into a concentrated solution in *n*-hexane/chloroform (1:3 v/v).

#### Crystal data

 $\begin{array}{l} C_{12}H_{11}N_{3}O_{3}\\ M_{r} = 245.24\\ \text{Orthorhombic, } P_{2_{1}2_{1}2_{1}}\\ a = 7.3550 \ (10) \ \text{\AA}\\ b = 11.6182 \ (15) \ \text{\AA}\\ c = 13.6335 \ (17) \ \text{\AA}\\ V = 1165.0 \ (3) \ \text{\AA}^{3}\\ Z = 4\\ D_{x} = 1.398 \ \text{Mg m}^{-3} \end{array}$ 

#### Data collection

Bruker CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: none 8600 measured reflections 1569 independent reflections

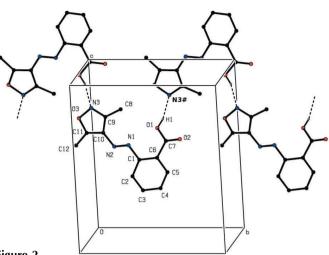
#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.043$   $wR(F^2) = 0.074$  S = 0.911569 reflections 165 parameters Cell parameters from 999 reflections  $\theta = 2.3-27.9^{\circ}$  $\mu = 0.10 \text{ mm}^{-1}$ T = 298 KPolyhedron, orange  $0.26 \times 0.20 \times 0.18 \text{ mm}$ 

Mo  $K\alpha$  radiation

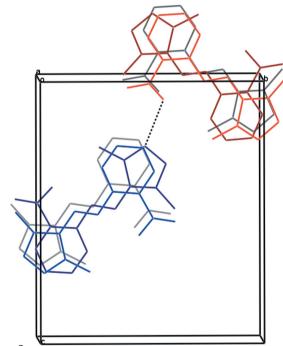
945 reflections with $I > 2c$ $R_{int} = 0.073$	$\sigma(I)$
$\theta_{\rm max} = 27.9^{\circ}$	
$ \begin{array}{l} h = -9 \rightarrow 9 \\ k = -14 \rightarrow 15 \end{array} $	
$l = -17 \rightarrow 17$	

H-atom parameters constrained  $w = 1/[\sigma^2(F_o^2) + (0.0217P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$   $(\Delta/\sigma)_{max} = 0.019$   $\Delta\rho_{max} = 0.14 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{min} = -0.10 \text{ e } \text{\AA}^{-3}$ 





Part of the crystal structure, showing the formation of a zigzag chain along [010]. The dashed lines represent hydrogen bonds. H atoms not involved in hydrogen bonds have been omitted. [Symmetry code: (#) -x,  $\frac{1}{2} + y$ ,  $\frac{3}{2} - z$ .]





View approximately along the [100] direction, showing the  $\pi$ - $\pi$  stacking. H atoms have been omitted.

## Table 1

Selected geometric parameters (Å, °).

O3-N3	1.424 (3)	C9-C10	1.418 (4)
O3-C11	1.352 (3)	C10-C11	1.347 (3)
N3-C9	1.301 (3)	N1-N2	1.254 (3)
N2-N1-C1	113.0 (2)	N1-N2-C10	113.2 (2)
112 111 01	115.0 (2)	111 112 010	115.2 (2)
G4 114 119 G40			
C1-N1-N2-C10	176.6 (2)		

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\overline{O1\!-\!H1\!\cdot\cdot\cdot\!N3^i}$	0.82	1.94	2.735 (3)	163
Commentary and as (i)				

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

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with aromatic C-H = 0.93 Å, methyl C-H = 0.96 Å and O-H = 0.82 Å. The methyl groups were allowed to rotate but not to tip. In the absence of significant anomalous dispersion effects, Friedel pairs were merged prior to final refinement.

Data collection: *SMART-NT* (Bruker, 2001); cell refinement: *SAINT-NT* (Bruker, 2000); data reduction: *SAINT-NT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL-PC* (Sheldrick, 1994); software used to prepare material for publication: *PLATON* (Spek, 2003).

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