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3-[3-(Pyridin-3-yl)-1,2,4-oxadiazol-5-yl]-propanoic acid

 Fang Zhang,^a Fei Liu,^{a*} Qifan Chen^b and Mingdong Dong^b

^aCollege of Chemical Engineering & Materials, Eastern Liaoning University, No. 325 Wenhua Road, Yuanbao District, Dandong City, Liaoning Province 118003, People's Republic of China, and ^bExperiment Center, Eastern Liaoning University, No. 325 Wenhua Road, Yuanbao District, Dandong City, Liaoning Province 118003, People's Republic of China

Correspondence e-mail: berylliu8090@sina.com

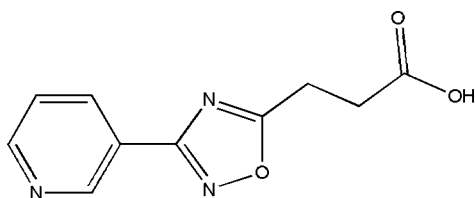
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.034; wR factor = 0.083; data-to-parameter ratio = 7.8.

In the title compound, $\text{C}_{10}\text{H}_9\text{N}_3\text{O}_3$, the benzene ring is almost coplanar with the heterocyclic ring, making a dihedral angle of 11.3 (1)°. The plane of the carboxyl group is rotated by 8.4 (2)° with respect to the 1,2,4-oxadiazole ring plane. The aliphatic chain exhibits an extended conformation. In the crystal, molecules are linked through intermolecular $\text{O}-\text{H}\cdots\text{N}$ bonds, forming a chain structure along the c axis.

Related literature

For the biological activity of 1,2,4-oxadiazoles, see: Jakopin & Dolenc, (2008). For the use of this heterocycle as a core for luminescent liquid crystals, see: Gallardo *et al.* (2008). For related structures, see: Santos *et al.* (2009); Wang *et al.* (2006, 2007)



Experimental

Crystal data

 $\text{C}_{10}\text{H}_9\text{N}_3\text{O}_3$
 $M_r = 219.20$

Orthorhombic, $Pna2_1$
 $a = 6.1298$ (12) Å
 $b = 6.8194$ (14) Å
 $c = 23.426$ (5) Å
 $V = 979.2$ (3) Å³

$Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.11$ mm⁻¹
 $T = 293$ K
 $0.33 \times 0.22 \times 0.21$ mm

Data collection

Siemens P4 diffractometer
 8147 measured reflections
 1138 independent reflections

884 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$
 $wR(F^2) = 0.083$
 $S = 1.00$
 1138 reflections
 146 parameters

1 restraint
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.14$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O1}-\text{H1}\cdots\text{N3}^i$	0.82	1.89	2.704 (3)	172

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

Data collection: *XSCANS* (Bruker, 2003); cell refinement: *XSCANS*; data reduction: *XSCANS* and *SHELXTL* (Sheldrick, 2008); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FK2032).

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supporting information

Acta Cryst. (2011). E67, o193 [https://doi.org/10.1107/S1600536810051639]

3-[3-(Pyridin-3-yl)-1,2,4-oxadiazol-5-yl]propanoic acid**Fang Zhang, Fei Liu, Qifan Chen and Mingdong Dong****S1. Comment**

1,2,4-Oxadiazoles are well known compounds, which exhibit a large number of biological activities (Jakopin & Dolenc, 2008). Recently, the use of this heterocycle as core for luminescent liquid crystals has also been described (Gallardo *et al.*, 2008). Here we report the structure of title compound (Fig. 1), the benzene ring is almost coplanar with the heterocyclic ring, making a dihedral angle of 11.3 (1) °. The torsion angle N2—C5—C6—C10 between the pyridine ring attached to C-5 of the 1,2,4-oxadiazole system is -8.0 (3) °, both rings are almost coplanar. The C-4 side-chain containing a carboxylic acid group shows a zigzag arrangement, having the torsion angle C1—C2—C3—C4 of -178.0 (2) °. In addition, the plane of the carboxylic group is also rotated by 8.4 (2) ° with respect to the mean plane of the 1,2,4-oxadiazole five-membered ring. This makes the molecular structure to be slightly twisted. In the crystal structure, molecules are linked through intermolecular O—H···N, forming a one-dimensional chain structure along the crystallographic *c* axis.

S2. Experimental

To a solution of nitrile (0.2 mol) in ethanol (20 mL) was added hydroxylamine hydrochloride (0.4 mol) in water (40 mL). Then anhydrous sodium carbonate (0.4 mol) in water (120 mL) was slowly added to the resulting solution and the mixture was stirred at 358k for 5 h. The mixture was then concentrated under vacuum to evaporate some water. The resulting suspension was filtered, the amidoxime solid formed was washed with cold water, dried under vacuum. A thoroughly triturated mixture of amidoxime (0.04 mol) and succinic anhydride (0.08 mol) was heated in an oily bath to 403k and kept at this temperature for 4 h. The reaction mixture was cooled to room temperature, and the product was washed with cold water, filtered, and recrystallized from ethanol. Block-shaped crystals suitable for X-ray diffraction were obtained from methanol.

S3. Refinement

H atoms bound to C atoms were placed in calculated positions and treated as riding on their parent atoms, with C—H = 0.93 Å (aromatic C), C—H = 0.97 Å (methylene C), and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The H atom bound to O1 was located from Fourier difference map and refined with O—H = 0.82 Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

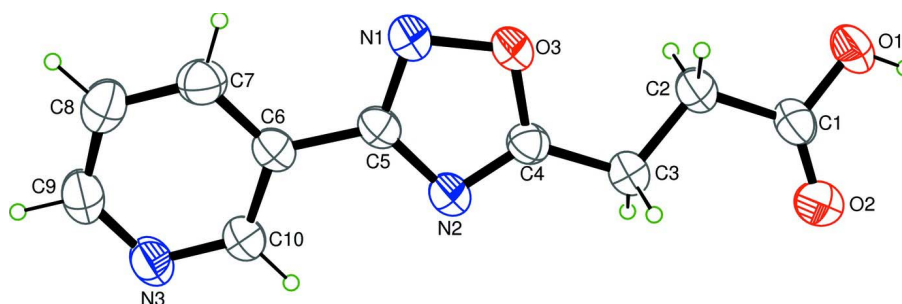


Figure 1

Structure of the title compound showing 50% probability displacement ellipsoids.

3-[3-(Pyridin-3-yl)-1,2,4-oxadiazol-5-yl]propanoic acid

Crystal data

$C_{10}H_9N_3O_3$

$M_r = 219.20$

Orthorhombic, $Pna2_1$

Hall symbol: $P\ 2c\ -2n$

$a = 6.1298\ (12)\ \text{\AA}$

$b = 6.8194\ (14)\ \text{\AA}$

$c = 23.426\ (5)\ \text{\AA}$

$V = 979.2\ (3)\ \text{\AA}^3$

$Z = 4$

$F(000) = 456$

$D_x = 1.487\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 6383 reflections

$\theta = 3.1\text{--}27.5^\circ$

$\mu = 0.11\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Block, colourless

$0.33 \times 0.22 \times 0.21\ \text{mm}$

Data collection

Bruker P4

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

8147 measured reflections

1138 independent reflections

884 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.049$

$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.1^\circ$

$h = -6 \rightarrow 7$

$k = -8 \rightarrow 8$

$l = -30 \rightarrow 30$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.034$

$wR(F^2) = 0.083$

$S = 1.00$

1138 reflections

146 parameters

1 restraint

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: geom and difmap

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0487P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\text{max}} = 0.14\ \text{e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.16\ \text{e \AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O3	0.8809 (3)	0.4377 (3)	0.16313 (7)	0.0450 (5)
O1	0.4239 (4)	0.4072 (4)	0.33554 (8)	0.0628 (6)
H1	0.3237	0.4277	0.3579	0.094*
O2	0.2140 (4)	0.5819 (3)	0.27757 (9)	0.0681 (7)
N3	0.8817 (4)	0.5436 (3)	-0.08242 (9)	0.0468 (6)
C10	0.8297 (5)	0.5499 (4)	-0.02742 (11)	0.0412 (6)
H10	0.6895	0.5885	-0.0173	0.049*
N2	0.6960 (4)	0.5622 (3)	0.09157 (9)	0.0394 (5)
N1	1.0181 (4)	0.4195 (3)	0.11489 (9)	0.0461 (5)
C6	0.9757 (4)	0.5012 (4)	0.01552 (11)	0.0358 (5)
C5	0.8992 (4)	0.4978 (3)	0.07499 (10)	0.0346 (5)
C1	0.3807 (5)	0.4921 (4)	0.28656 (12)	0.0430 (6)
C2	0.5595 (4)	0.4565 (4)	0.24436 (11)	0.0426 (6)
H2A	0.5711	0.3167	0.2373	0.051*
H2B	0.6967	0.5002	0.2606	0.051*
C3	0.5225 (5)	0.5608 (4)	0.18853 (10)	0.0427 (6)
H3A	0.3826	0.5204	0.1731	0.051*
H3B	0.5153	0.7008	0.1956	0.051*
C8	1.2425 (5)	0.4437 (4)	-0.05639 (12)	0.0491 (7)
H8	1.3826	0.4088	-0.0677	0.059*
C7	1.1881 (4)	0.4503 (4)	0.00047 (12)	0.0449 (6)
H7	1.2912	0.4214	0.0284	0.054*
C4	0.6937 (4)	0.5226 (3)	0.14541 (10)	0.0358 (5)
C9	1.0850 (5)	0.4898 (4)	-0.09637 (13)	0.0497 (7)
H9	1.1220	0.4832	-0.1348	0.060*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O3	0.0460 (11)	0.0605 (10)	0.0285 (9)	0.0084 (8)	-0.0026 (8)	0.0037 (8)
O1	0.0615 (14)	0.0953 (16)	0.0315 (10)	0.0191 (11)	0.0058 (9)	0.0162 (10)
O2	0.0625 (15)	0.0933 (16)	0.0484 (13)	0.0305 (12)	0.0078 (11)	0.0187 (11)
N3	0.0530 (14)	0.0543 (12)	0.0331 (12)	-0.0001 (10)	0.0009 (10)	0.0043 (10)
C10	0.0439 (18)	0.0467 (13)	0.0329 (13)	0.0034 (12)	-0.0010 (11)	0.0005 (11)
N2	0.0409 (11)	0.0462 (11)	0.0311 (11)	0.0030 (9)	-0.0008 (10)	0.0040 (9)
N1	0.0466 (12)	0.0580 (14)	0.0336 (11)	0.0079 (10)	0.0017 (10)	0.0048 (9)
C6	0.0411 (14)	0.0344 (10)	0.0319 (12)	-0.0020 (11)	-0.0028 (11)	0.0010 (9)
C5	0.0388 (13)	0.0340 (11)	0.0310 (12)	-0.0018 (11)	-0.0043 (11)	0.0008 (9)
C1	0.0473 (17)	0.0506 (14)	0.0311 (13)	-0.0015 (13)	-0.0036 (12)	0.0005 (10)
C2	0.0437 (15)	0.0525 (14)	0.0316 (13)	0.0027 (10)	-0.0015 (12)	0.0018 (11)

C3	0.0460 (15)	0.0503 (13)	0.0317 (13)	0.0066 (12)	0.0007 (12)	0.0021 (11)
C8	0.0450 (17)	0.0503 (16)	0.0521 (18)	-0.0005 (12)	0.0116 (14)	-0.0025 (12)
C7	0.0399 (15)	0.0470 (14)	0.0479 (17)	0.0019 (12)	-0.0022 (13)	0.0039 (12)
C4	0.0382 (14)	0.0361 (11)	0.0332 (14)	0.0011 (10)	-0.0050 (10)	-0.0013 (10)
C9	0.0575 (19)	0.0544 (15)	0.0371 (15)	-0.0021 (14)	0.0090 (13)	0.0014 (12)

Geometric parameters (Å, °)

O3—C4	1.351 (3)	C6—C5	1.470 (4)
O3—N1	1.414 (3)	C1—C2	1.496 (4)
O1—C1	1.312 (3)	C2—C3	1.506 (3)
O1—H1	0.8200	C2—H2A	0.9700
O2—C1	1.210 (3)	C2—H2B	0.9700
N3—C10	1.328 (4)	C3—C4	1.480 (3)
N3—C9	1.340 (4)	C3—H3A	0.9700
C10—C6	1.387 (4)	C3—H3B	0.9700
C10—H10	0.9300	C8—C7	1.374 (4)
N2—C4	1.290 (3)	C8—C9	1.381 (4)
N2—C5	1.377 (3)	C8—H8	0.9300
N1—C5	1.300 (3)	C7—H7	0.9300
C6—C7	1.393 (4)	C9—H9	0.9300
C4—O3—N1	107.29 (18)	C3—C2—H2B	109.0
C1—O1—H1	109.5	H2A—C2—H2B	107.8
C10—N3—C9	117.9 (2)	C4—C3—C2	113.7 (2)
N3—C10—C6	122.8 (3)	C4—C3—H3A	108.8
N3—C10—H10	118.6	C2—C3—H3A	108.8
C6—C10—H10	118.6	C4—C3—H3B	108.8
C4—N2—C5	102.6 (2)	C2—C3—H3B	108.8
C5—N1—O3	101.84 (19)	H3A—C3—H3B	107.7
C10—C6—C7	118.6 (2)	C7—C8—C9	118.7 (3)
C10—C6—C5	119.1 (2)	C7—C8—H8	120.6
C7—C6—C5	122.3 (2)	C9—C8—H8	120.6
N1—C5—N2	115.8 (2)	C8—C7—C6	118.7 (2)
N1—C5—C6	120.6 (2)	C8—C7—H7	120.6
N2—C5—C6	123.4 (2)	C6—C7—H7	120.6
O2—C1—O1	123.1 (3)	N2—C4—O3	112.4 (2)
O2—C1—C2	125.9 (3)	N2—C4—C3	129.6 (3)
O1—C1—C2	111.0 (2)	O3—C4—C3	117.9 (2)
C1—C2—C3	112.8 (2)	N3—C9—C8	123.2 (3)
C1—C2—H2A	109.0	N3—C9—H9	118.4
C3—C2—H2A	109.0	C8—C9—H9	118.4
C1—C2—H2B	109.0		
C9—N3—C10—C6	-0.8 (4)	O1—C1—C2—C3	-176.8 (2)
C4—O3—N1—C5	1.1 (2)	C1—C2—C3—C4	-178.1 (2)
N3—C10—C6—C7	2.4 (4)	C9—C8—C7—C6	0.7 (4)
N3—C10—C6—C5	-175.4 (2)	C10—C6—C7—C8	-2.3 (4)

O3—N1—C5—N2	-1.3 (3)	C5—C6—C7—C8	175.4 (2)
O3—N1—C5—C6	-176.9 (2)	C5—N2—C4—O3	-0.2 (3)
C4—N2—C5—N1	1.0 (3)	C5—N2—C4—C3	178.2 (2)
C4—N2—C5—C6	176.4 (2)	N1—O3—C4—N2	-0.6 (3)
C10—C6—C5—N1	167.5 (2)	N1—O3—C4—C3	-179.1 (2)
C7—C6—C5—N1	-10.2 (3)	C2—C3—C4—N2	168.1 (2)
C10—C6—C5—N2	-7.7 (3)	C2—C3—C4—O3	-13.6 (3)
C7—C6—C5—N2	174.6 (2)	C10—N3—C9—C8	-0.9 (4)
O2—C1—C2—C3	5.1 (4)	C7—C8—C9—N3	0.9 (4)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O1—H1 \cdots N3 ⁱ	0.82	1.89	2.704 (3)	172

Symmetry code: (i) $-x+1, -y+1, z+1/2$.