organic compounds

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2-(3-Methyl-5-thioxo-4,5-dihydro-1H-1,2,4-triazol-4-yl)acetic acid

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Key indicators: single-crystal X-ray study; T = 291 K; mean σ (C–C) = 0.003 Å; R factor = 0.034; wR factor = 0.087; data-to-parameter ratio = 13.3.

All interatomic distances in the title compound, C₅H₇N₃O₂S, are normal. The 1,2,4-triazoline ring is planar and it is inclined at 78.61 $(7)^{\circ}$ to the planar acetic acid group. The molecules of the title compound are connected via O-H···N hydrogen bonds into zigzag chains along the [101] direction and by N- $H \cdots S$ hydrogen bonds into a three-dimensional net. In addition, a weak $C-H \cdot \cdot \cdot O$ hydrogen bond can also be found in the structure.

Related literature

For potential applications, see: Bohn & Karow (1981); Potts (1961); Santus (1980). For general synthetic procedures, see: Bany & Dobosz (1972); Veverka & Marchalin (1987). For related literature, see: Bernstein et al. (1995); Desiraju & Steiner (1999).

Experimental

Crystal data

C ₅ H ₇ N ₃ O ₂ S	c = 15.6763 (9) Å
$M_r = 173.20$	$\beta = 99.250 \ (5)^{\circ}$
Monoclinic, $P2_1/n$	V = 758.56 (8) Å ³
a = 5.1706 (3) Å	Z = 4
b = 9.4818 (6) Å	Mo $K\alpha$ radiation

 $\mu = 0.38 \text{ mm}^{-1}$ T = 291 (2) K

Data collection

Kuma KM-4 diffractometer	1354 independent reflections
Absorption correction: numerical	1108 reflections with $I > 2\sigma(I)$
(X-RED; Stoe & Cie, 1999)	3 standard reflections
$T_{\min} = 0.918, \ T_{\max} = 0.983$	every 100 reflections
1354 measured reflections	intensity decay: 2.1%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$	102 parameters
$wR(F^2) = 0.087$	H-atom parameters constrained
S = 1.09	$\Delta \rho_{\rm max} = 0.16 \text{ e } \text{\AA}^{-3}$
1354 reflections	$\Delta \rho_{\rm min} = -0.24 \text{ e } \text{\AA}^{-3}$

Table 1 Hydrogen-bond geometry (Å, °).

 $D - \mathbf{H} \cdot \cdot \cdot A$ D - H $D = H \cdots A$ $H \cdot \cdot \cdot A$ $D \cdots A$ $O1\!-\!H1O\!\cdots\!N2^i$ 0.82 1.96 2.746 (2) 162 $N1 - H1N \cdots S1^{ii}$ 0.86 2.39 3.2499 (18) 176 $C3-H3A\cdots O2^{iii}$ 0.97 2.60 3.567 (2) 176

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

Data collection: KM-4 Software (Kuma, 1993); cell refinement: KM-4 Software; data reduction: DATAPROC (Gałdecki et al., 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP in SHELXTL/PC (Sheldrick, 1990b) and ORTEP-3 for Windows (Version 1.062; Farrugia, 1997); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT2545).

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 $0.22 \times 0.07 \times 0.07 \; \text{mm}$

supplementary materials

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2-(3-Methyl-5-thioxo-4,5-dihydro-1H-1,2,4-triazol-4-yl)acetic acid

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Comment

The title compound, (I), is a member of 1,2,4-triazoline-3-thione derivatives family, known to posses antibacterial, anitimycotical and antivirostatical activity (Veverka and Marchalin, 1987; Bohn and Karow, 1981; Potts, 1961; Santus, 1980).

All interatomic distances in (I) are normal. The 1,2,4-triazoline ring of (I) can be considered as planar in the range of experimental error. The most deviating N1 atom derives 0.0046 (13) Å from weighted least squares plane of the ring. The C3, C5, S1 atoms deviate respectively 0.082 (3), -0.004 (4), -0.049 (3) Å from this plane. The acetic acid moiety is planar (Table 1) and the most deviating atom is C4 [0.0005 (17) Å] from the weighted O1/O2/C3/C4 least-squares plane. The N3 atom deviates 0.106 (4) Å from this plane. The above mentioned weighted least-squares planes are inclined at 78.61 (7)°.

The molecules of (I) are connected *via* O1—H1O···N2 hydrogen bonds (Table 1, $C_1^{1}(7)$ motif (Bernstein *et al.*, 1995)) to zigzag chains extended along the [101] axis. The N1—H1N···S1 hydrogen bonds (Table 1, $R_2^{2}(8)$ motif) expands the chains to a folded three dimensional sheet in the (-101) plane. In (I) can be found also one C—H···O short contact (Table 1), which, according to Desiraju and Steiner (1999), can be classified as weak hydrogen bond ($C_1^{1}(4)$ motif).

Experimental

The title compound was synthesized according to method of Veverka and Marchalin (1987). Crystals were obtained by crystallization from mixture of water, methanol, ethanol and 2-butanone (3:1:5:1).

Refinement

The hydrogen atoms were placed in calculated positions after four cycles of anisotrophic refinement and were refined as riding on the parent atom with $U_{iso}(H) = 1.2U_{eq}(C$ -non-methyl or N) and $U_{iso}(H) = 1.5U_{eq}(C$ -methyl or O). The methyl and hydroxyl group was allowed to rotate about its local threefold axis (AFIX 137 and 147 respectively).

Figures



Fig. 1. Molecular structure of the title compound (I). Displacement ellipsoids are drawn at the 50% probability level.



Fig. 2. A part of the molecular packing of the title compound. Hydrogen bonds are indicated by dashed lines.

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2-(3-Methyl-5-thioxo-4,5-dihydro-1H-1,2,4-triazol-4-yl)acetic acid

Crystal data	
$C_5H_7N_3O_2S$	$F_{000} = 360$
$M_r = 173.20$	$D_{\rm x} = 1.517 {\rm ~Mg~m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation $\lambda = 0.71073$ Å
Hall symbol: -P 2yn	Cell parameters from 99 reflections
a = 5.1706 (3) Å	$\theta = 2 - 20^{\circ}$
<i>b</i> = 9.4818 (6) Å	$\mu = 0.38 \text{ mm}^{-1}$
c = 15.6763 (9) Å	T = 291 (2) K
$\beta = 99.250 \ (5)^{\circ}$	Needle, colourless
$V = 758.56 (8) \text{ Å}^3$	$0.22\times0.07\times0.07~mm$
Z = 4	

Data collection

Kuma KM-4 diffractometer	$R_{\rm int} = 0.0000$
Radiation source: fine-focus sealed tube	$\theta_{\text{max}} = 25.1^{\circ}$
Monochromator: graphite	$\theta_{\min} = 3.4^{\circ}$
T = 291(2) K	$h = -6 \rightarrow 6$
ω -2 θ scans	$k = 0 \rightarrow 11$
Absorption correction: numerical (X-RED; Stoe & Cie, 1999)	$l = 0 \rightarrow 18$
$T_{\min} = 0.918, \ T_{\max} = 0.983$	3 standard reflections
1354 measured reflections	every 100 reflections
1354 independent reflections	intensity decay: 2.1%
1108 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	Secondary atom site location: structure-invariant dir ect methods
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.034$	H-atom parameters constrained
$wR(F^2) = 0.087$	$w = 1/[\sigma^2(F_0^2) + (0.0468P)^2]$ where $P = (F_0^2 + 2F_c^2)/3$
S = 1.09	$(\Delta/\sigma)_{\rm max} < 0.001$

1354 reflections

102 parameters

Primary atom site location: structure-invariant direct Extinction correction: none

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.

 $\Delta \rho_{\text{max}} = 0.16 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.24 \text{ e} \text{ Å}^{-3}$

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 $F^2 > 2 \operatorname{sigma}(F^2)$ 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.

	x	У	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
S1	0.99527 (11)	1.04539 (6)	0.63708 (3)	0.0393 (2)
01	0.7717 (3)	0.88549 (17)	0.88096 (9)	0.0437 (4)
H1O	0.8832	0.8517	0.9185	0.066*
C4	0.8051 (4)	0.8362 (2)	0.80424 (12)	0.0301 (5)
O2	0.9822 (3)	0.76167 (17)	0.79093 (9)	0.0420 (4)
N1	0.7338 (3)	0.8532 (2)	0.52587 (10)	0.0401 (5)
H1N	0.8106	0.8759	0.4831	0.048*
N3	0.6155 (3)	0.84909 (19)	0.64966 (10)	0.0313 (4)
C3	0.5851 (4)	0.8886 (2)	0.73708 (12)	0.0336 (5)
H3A	0.4208	0.8508	0.7495	0.040*
H3B	0.5763	0.9905	0.7407	0.040*
C2	0.7812 (4)	0.9166 (2)	0.60293 (12)	0.0335 (5)
N2	0.5497 (4)	0.7480 (2)	0.52213 (11)	0.0413 (5)
C1	0.4797 (4)	0.7476 (2)	0.59811 (12)	0.0347 (5)
C5	0.2828 (4)	0.6513 (3)	0.62525 (15)	0.0438 (6)
H5A	0.2063	0.5947	0.5769	0.066*
H5B	0.1484	0.7056	0.6456	0.066*
H5C	0.3661	0.5911	0.6708	0.066*
	. 0	2		
Atomic displace	ement parameters (A ²	-)		

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

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U ²³
S 1	0.0452 (3)	0.0464 (4)	0.0274 (3)	-0.0021 (3)	0.0090 (2)	-0.0016 (2)
01	0.0578 (10)	0.0515 (10)	0.0206 (8)	0.0089 (8)	0.0029 (7)	-0.0014 (7)
C4	0.0347 (11)	0.0345 (11)	0.0222 (10)	-0.0032 (9)	0.0082 (8)	0.0018 (9)
O2	0.0366 (8)	0.0546 (10)	0.0356 (8)	0.0104 (8)	0.0087 (7)	0.0068 (7)
N1	0.0495 (11)	0.0521 (12)	0.0202 (9)	-0.0065 (10)	0.0095 (8)	0.0007 (8)
N3	0.0322 (9)	0.0425 (10)	0.0198 (8)	0.0043 (8)	0.0065 (7)	0.0005 (8)

supplementary materials

C3	0.0347 (11)	0.0459 (13)	0.0212 (10)	0.0065 (10)	0.0073 (8)	-0.0003 (9)
C2	0.0339 (11)	0.0451 (13)	0.0216 (10)	0.0079 (10)	0.0049 (8)	0.0022 (9)
N2	0.0481 (11)	0.0523 (12)	0.0231 (9)	-0.0059 (10)	0.0042 (8)	-0.0024 (8)
C1	0.0366 (11)	0.0445 (13)	0.0220 (10)	0.0046 (10)	0.0015 (8)	0.0012 (9)
C5	0.0442 (13)	0.0501 (14)	0.0375 (13)	-0.0011 (11)	0.0077 (10)	0.0021 (11)
Geometric param	neters (Å, °)					
S1—C2		1.677 (2)	N3—0	C1	1.3	374 (3)
O1—C4		1.327 (2)	N3—0	23	1.4	453 (2)
01—H10		0.8200	С3—І	H3A	0.9	9700
C4—O2		1.201 (2)	C3—I	I3B	0.9	9700
C4—C3		1.505 (3)	N2—0	C1	1.2	299 (3)
N1—C2		1.336 (3)	C1—0	25	1.480 (3)	
N1—N2		1.374 (3)	C5—I	15A	0.9600	
N1—H1N		0.8600	C5—H5B		0.9600	
N3—C2		1.372 (3)	C5—H5C		0.9600	
C4—O1—H1O		109.5	НЗА—СЗ—НЗВ		10	7.8
O2—C4—O1		125.34 (19)	N1—0	C2—N3	10	3.51 (18)
O2—C4—C3		125.76 (18)	N1—0	C2—S1	12	9.18 (17)
O1—C4—C3		108.89 (17)	N3—C2—S1		12	7.30 (15)
C2—N1—N2		112.98 (17)	C1—N2—N1		104.81 (17)	
C2—N1—H1N		123.5	N2—C1—N3		11	0.26 (19)
N2—N1—H1N		123.5	N2—C1—C5		124.8 (2)	
C2—N3—C1		108.44 (16)	N3—0	C1—C5	12	4.95 (18)
C2—N3—C3		123.71 (18)	C1—0	С5—Н5А	109.5	
C1—N3—C3		127.75 (17)	C1—C5—H5B		109.5	
N3—C3—C4		112.98 (16)	H5A-	-C5—H5B	109.5	
N3—C3—H3A		109.0	C1—0	С5—Н5С	10	9.5
С4—С3—НЗА		109.0	H5A-	-C5—H5C	10	9.5
N3—C3—H3B		109.0	H5B-	-С5—Н5С	109.5	
C4—C3—H3B		109.0				
N3—C3—C4—O	1	175.46 (17)	N3—0	C3—C4—O2	-4	.6 (3)

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	H…A	$D \cdots A$	$D\!\!-\!\!\mathrm{H}^{\dots}\!A$		
O1—H1O····N2 ⁱ	0.82	1.96	2.746 (2)	162		
N1—H1N····S1 ⁱⁱ	0.86	2.39	3.2499 (18)	176		
C3—H3A···O2 ⁱⁱⁱ	0.97	2.60	3.567 (2)	176		
Symmetry adds: (i) $y + 1/2$, $y + 2/2$, $z + 1/2$; (ii) $y + 2$, $y + 2$, $z + 1$; (iii) $y + 1$, $y = z$						

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





