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N-(Thiazol-2-yl)acetamide

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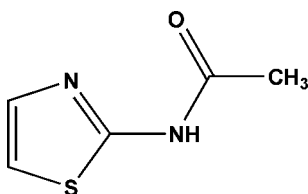
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å;
 R factor = 0.036; wR factor = 0.104; data-to-parameter ratio = 18.6.

The title compound, $\text{C}_5\text{H}_6\text{N}_2\text{OS}$, was synthesized from acetyl chloride and 2-aminothiazole in dry acetone. The asymmetric unit contains two molecules. The crystal structure is stabilized by $\text{N}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related literature, see: Raman *et al.* (2000); Wang *et al.* (2008); Yunus *et al.* (2007, 2008).



Experimental

Crystal data

$\text{C}_5\text{H}_6\text{N}_2\text{OS}$
 $M_r = 142.18$
Monoclinic, $P2_1/c$
 $a = 16.0650$ (12) Å
 $b = 11.3337$ (8) Å
 $c = 7.0670$ (5) Å
 $\beta = 101.908$ (10)°

$V = 1259.04$ (16) Å³
 $Z = 8$
Mo $K\alpha$ radiation
 $\mu = 0.42$ mm⁻¹
 $T = 173$ (2) K
 $0.30 \times 0.26 \times 0.22$ mm

Data collection

Bruker SMART1000 CCD
diffractometer
Absorption correction: multi-scan
(*SADABS*; Bruker, 1999)
 $T_{\min} = 0.830$, $T_{\max} = 1.000$
(expected range = 0.757–0.911)

7429 measured reflections
3024 independent reflections
2602 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.104$
 $S = 1.05$
3024 reflections

163 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.34$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2B}\cdots\text{N3}^{\text{i}}$	0.88	2.04	2.897 (2)	163
$\text{N4}-\text{H4A}\cdots\text{N1}^{\text{ii}}$	0.88	2.07	2.938 (2)	171
$\text{C2}-\text{H2A}\cdots\text{O2}^{\text{iii}}$	0.95	2.41	3.350 (2)	171
$\text{C7}-\text{H7A}\cdots\text{O1}^{\text{iv}}$	0.95	2.46	3.382 (2)	165

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

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WK2088).

References

- Bruker (1999). *SMART*, *SAINTE* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Macrae, C. F., Edgington, P. R., McCabe, P., Pidcock, E., Shields, G. P., Taylor, R., Towler, M. & van de Streek, J. (2006). *J. Appl. Cryst.* **39**, 453–457.
- Raman, R., Razavi, H. & Kelly, J. W. (2000). *Org. Lett.* **2**, 3289–3292.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Wang, X.-J., Yang, Q., Liu, F. & You, Q.-D. (2008). *Synth. Commun.* **38**, 1028–1035.
- Yunus, U., Tahir, M. K., Bhatti, M. H., Ali, S. & Helliwell, M. (2007). *Acta Cryst.* **E63**, o3690.
- Yunus, U., Tahir, M. K., Bhatti, M. H. & Wong, W.-Y. (2008). *Acta Cryst.* **E64**, o722.

supporting information

Acta Cryst. (2008). E64, o1516 [doi:10.1107/S1600536808021442]

N-(Thiazol-2-yl)acetamide

Uzma Yunus, Muhammad Kalim Tahir, Moazzam Hussain Bhatti and Wai-Yeung Wong

S1. Comment

The thiazole ring and its derivatives are of great importance in biological systems due to their vast range of biological activities such as anti-inflammatory, analgesic and antipyretic (Raman *et al.*, 2000). On the other hand amide compounds have extensive applications in the pharmaceutical industry (Wang *et al.*, 2008). As a part of our research the title compound (I) has been synthesized and its crystal structure is reported herein (Yunus *et al.*, 2007; 2008).

The title compound (I) crystallizes in a monoclinic space group with two molecules in asymmetric unit. All the bond lengths and angles are within the normal ranges. The molecules are stabilized by intermolecular hydrogen bonds N—H \cdots N, and C—H \cdots O (Table 1, Fig 2).

S2. Experimental

A mixture of acetyl chloride (26 mmol) and 2-aminothiazole (26 mmol) was refluxed in dry acetone (60 ml) for two hours. After cooling, the mixture was poured into acidified cold water. The resulting yellow solid was filtered and washed with cold acetone. Single crystals of the title compound suitable for single-crystal *x*-ray analysis were obtained by recrystallization of the yellow solid from ethyl acetate.

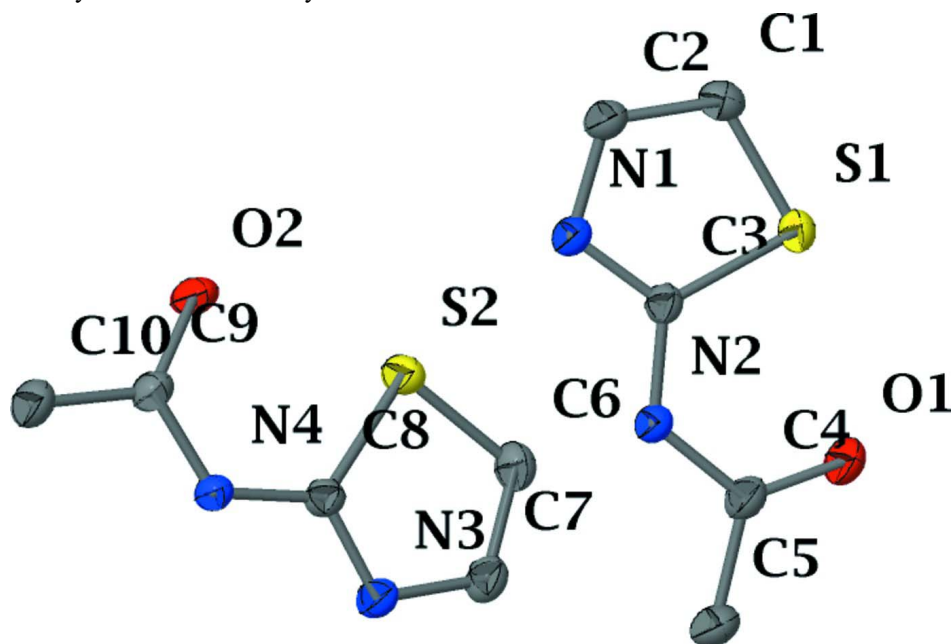
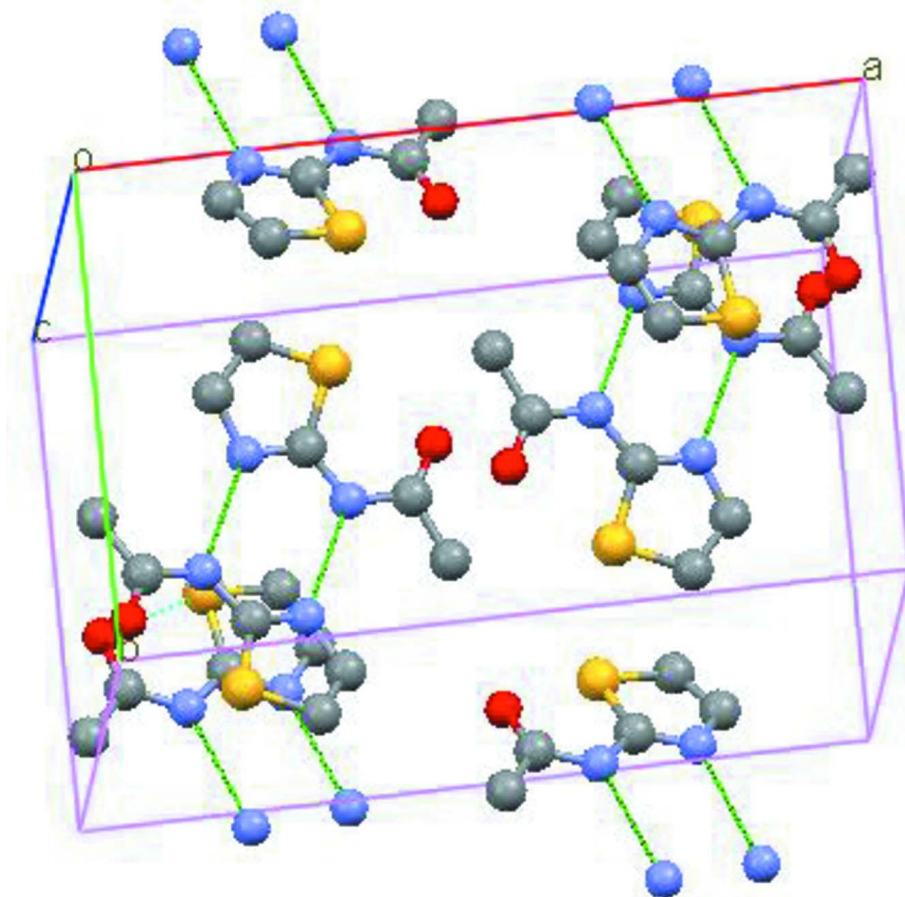


Figure 1

The molecular structure of (I) with ellipsoids drawn at the 50% probability level.

**Figure 2**

A packing diagram for (I) showing N—H...N hydrogen bonding.

***N*-(Thiazol-2-yl)acetamide**

Crystal data

$C_5H_6N_2OS$

$M_r = 142.18$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 16.0650$ (12) Å

$b = 11.3337$ (8) Å

$c = 7.0670$ (5) Å

$\beta = 101.908$ (10)°

$V = 1259.04$ (16) Å³

$Z = 8$

$F(000) = 592$

$D_x = 1.500$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 7429 reflections

$\theta = 2.6$ – 28.3 °

$\mu = 0.42$ mm⁻¹

$T = 173$ K

Block, pale yellow

$0.30 \times 0.26 \times 0.22$ mm

Data collection

Bruker SMART1000 CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and φ scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 1999)

$T_{\min} = 0.830$, $T_{\max} = 1.000$

7429 measured reflections

3024 independent reflections

2602 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.024$
 $\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 2.6^\circ$
 $h = -21 \rightarrow 18$

$k = -15 \rightarrow 12$
 $l = -9 \rightarrow 9$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.104$
 $S = 1.05$
 3024 reflections
 163 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0544P)^2 + 0.5928P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.44 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.34 \text{ e } \text{\AA}^{-3}$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.22544 (11)	0.26865 (16)	0.3481 (3)	0.0287 (4)
H1A	0.2073	0.1965	0.2843	0.034*
C2	0.17369 (11)	0.35988 (15)	0.3648 (3)	0.0276 (4)
H2A	0.1143	0.3575	0.3127	0.033*
C3	0.29472 (10)	0.43764 (14)	0.5196 (2)	0.0223 (3)
C4	0.43344 (10)	0.50546 (16)	0.6861 (3)	0.0281 (4)
C5	0.48077 (11)	0.61150 (18)	0.7800 (3)	0.0362 (4)
H5A	0.5407	0.5910	0.8272	0.054*
H5B	0.4560	0.6372	0.8887	0.054*
H5C	0.4765	0.6755	0.6852	0.054*
C6	0.28449 (12)	0.53644 (16)	1.0659 (3)	0.0313 (4)
H6A	0.3085	0.4598	1.0867	0.038*
C7	0.32581 (11)	0.63693 (16)	1.1264 (3)	0.0290 (4)
H7A	0.3831	0.6374	1.1960	0.035*
C8	0.20282 (10)	0.71411 (14)	0.9848 (2)	0.0227 (3)
C9	0.06151 (10)	0.78011 (16)	0.8300 (3)	0.0277 (4)
C10	0.00676 (12)	0.88668 (17)	0.7755 (3)	0.0381 (4)
H10A	-0.0506	0.8618	0.7116	0.057*
H10B	0.0312	0.9361	0.6872	0.057*
H10C	0.0038	0.9318	0.8922	0.057*
N1	0.21293 (9)	0.45737 (13)	0.4630 (2)	0.0254 (3)
N2	0.34869 (8)	0.52329 (13)	0.6143 (2)	0.0252 (3)

H2B	0.3273	0.5934	0.6293	0.030*
N3	0.27956 (8)	0.73976 (13)	1.0806 (2)	0.0255 (3)
N4	0.14391 (8)	0.80175 (12)	0.9218 (2)	0.0246 (3)
H4A	0.1602	0.8756	0.9418	0.030*
O1	0.46679 (8)	0.40975 (12)	0.6735 (2)	0.0386 (3)
O2	0.03563 (8)	0.67920 (12)	0.7966 (2)	0.0380 (3)
S1	0.32900 (3)	0.30062 (4)	0.45821 (7)	0.02647 (13)
S2	0.18167 (3)	0.56560 (4)	0.94536 (7)	0.02913 (13)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0280 (8)	0.0239 (8)	0.0334 (9)	-0.0035 (6)	0.0047 (7)	-0.0030 (7)
C2	0.0213 (8)	0.0266 (8)	0.0329 (9)	-0.0019 (6)	0.0008 (6)	-0.0002 (7)
C3	0.0200 (7)	0.0211 (7)	0.0250 (8)	0.0019 (6)	0.0030 (6)	0.0021 (6)
C4	0.0200 (8)	0.0303 (9)	0.0322 (9)	0.0014 (6)	0.0016 (7)	0.0026 (7)
C5	0.0232 (8)	0.0368 (10)	0.0448 (11)	-0.0038 (7)	-0.0015 (7)	-0.0044 (8)
C6	0.0300 (9)	0.0250 (8)	0.0376 (10)	0.0057 (7)	0.0038 (7)	0.0037 (7)
C7	0.0237 (8)	0.0296 (9)	0.0321 (9)	0.0033 (7)	0.0024 (7)	0.0052 (7)
C8	0.0217 (7)	0.0215 (7)	0.0247 (8)	-0.0020 (6)	0.0044 (6)	0.0014 (6)
C9	0.0184 (7)	0.0280 (9)	0.0352 (9)	-0.0010 (6)	0.0021 (6)	0.0005 (7)
C10	0.0240 (8)	0.0327 (10)	0.0540 (12)	0.0040 (7)	-0.0004 (8)	0.0011 (9)
N1	0.0185 (6)	0.0249 (7)	0.0314 (8)	0.0000 (5)	0.0016 (5)	-0.0005 (6)
N2	0.0184 (6)	0.0222 (7)	0.0329 (8)	0.0005 (5)	0.0005 (5)	-0.0019 (6)
N3	0.0200 (6)	0.0242 (7)	0.0305 (8)	-0.0007 (5)	0.0009 (5)	0.0032 (6)
N4	0.0182 (6)	0.0195 (7)	0.0342 (8)	-0.0011 (5)	0.0008 (5)	-0.0003 (5)
O1	0.0232 (6)	0.0317 (7)	0.0557 (9)	0.0058 (5)	-0.0039 (6)	-0.0007 (6)
O2	0.0220 (6)	0.0276 (7)	0.0590 (9)	-0.0047 (5)	-0.0042 (6)	-0.0023 (6)
S1	0.0229 (2)	0.0210 (2)	0.0349 (2)	0.00316 (14)	0.00479 (16)	-0.00010 (15)
S2	0.0261 (2)	0.0205 (2)	0.0385 (3)	-0.00185 (15)	0.00132 (17)	-0.00081 (16)

Geometric parameters (Å, °)

C1—C2	1.347 (2)	C6—S2	1.7271 (19)
C1—S1	1.7236 (18)	C6—H6A	0.9500
C1—H1A	0.9500	C7—N3	1.384 (2)
C2—N1	1.385 (2)	C7—H7A	0.9500
C2—H2A	0.9500	C8—N3	1.311 (2)
C3—N1	1.311 (2)	C8—N4	1.381 (2)
C3—N2	1.379 (2)	C8—S2	1.7284 (17)
C3—S1	1.7326 (16)	C9—O2	1.223 (2)
C4—O1	1.221 (2)	C9—N4	1.371 (2)
C4—N2	1.366 (2)	C9—C10	1.497 (2)
C4—C5	1.502 (3)	C10—H10A	0.9800
C5—H5A	0.9800	C10—H10B	0.9800
C5—H5B	0.9800	C10—H10C	0.9800
C5—H5C	0.9800	N2—H2B	0.8800
C6—C7	1.343 (3)	N4—H4A	0.8800

C2—C1—S1	110.78 (13)	N3—C7—H7A	122.2
C2—C1—H1A	124.6	N3—C8—N4	121.06 (15)
S1—C1—H1A	124.6	N3—C8—S2	115.59 (12)
C1—C2—N1	115.51 (15)	N4—C8—S2	123.35 (12)
C1—C2—H2A	122.2	O2—C9—N4	121.01 (16)
N1—C2—H2A	122.2	O2—C9—C10	123.13 (16)
N1—C3—N2	121.20 (15)	N4—C9—C10	115.86 (15)
N1—C3—S1	115.26 (12)	C9—C10—H10A	109.5
N2—C3—S1	123.49 (12)	C9—C10—H10B	109.5
O1—C4—N2	121.52 (16)	H10A—C10—H10B	109.5
O1—C4—C5	123.60 (15)	C9—C10—H10C	109.5
N2—C4—C5	114.88 (15)	H10A—C10—H10C	109.5
C4—C5—H5A	109.5	H10B—C10—H10C	109.5
C4—C5—H5B	109.5	C3—N1—C2	109.91 (14)
H5A—C5—H5B	109.5	C4—N2—C3	123.68 (15)
C4—C5—H5C	109.5	C4—N2—H2B	118.2
H5A—C5—H5C	109.5	C3—N2—H2B	118.2
H5B—C5—H5C	109.5	C8—N3—C7	109.65 (15)
C7—C6—S2	110.75 (13)	C9—N4—C8	123.68 (14)
C7—C6—H6A	124.6	C9—N4—H4A	118.2
S2—C6—H6A	124.6	C8—N4—H4A	118.2
C6—C7—N3	115.69 (15)	C1—S1—C3	88.54 (8)
C6—C7—H7A	122.2	C6—S2—C8	88.31 (8)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N2—H2B...N3 ⁱ	0.88	2.04	2.897 (2)	163
N4—H4A...N1 ⁱⁱ	0.88	2.07	2.938 (2)	171
C2—H2A...O2 ⁱⁱⁱ	0.95	2.41	3.350 (2)	171
C7—H7A...O1 ^{iv}	0.95	2.46	3.382 (2)	165

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