

Ethyl 2-amino-4-isopropyl-1,3-thiazole-5-carboxylate

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Key indicators

Single-crystal X-ray study
 $T = 295\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
R factor = 0.038
wR factor = 0.109
Data-to-parameter ratio = 19.4

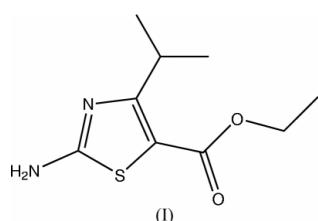
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Both the molecular and the crystal structures of the title compound, $\text{C}_9\text{H}_{14}\text{N}_2\text{O}_2\text{S}$, are similar to those of its 4-phenyl analogue. The supramolecular network is based upon $\text{N}-\text{H}\cdots\text{N}$ hydrogen-bonded centrosymmetric dimers linked by $\text{N}-\text{H}\cdots\text{O}$ contacts.

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Comment

The quest for bioactive compounds led us to the synthesis of a variety of heterocyclic compounds, among them the title compound, (I). There are many compounds in nature incorporating the thiazole moiety in their structure (Ikemoto *et al.*, 2003; Kumar *et al.*, 2002; El-Meligie & El-Awady, 2002), that have useful bioactivities. For example, Leucamide A was first extracted from the Australian marine sponge *Leucetta microraphis* and showed cytotoxicity toward several tumour cell lines (Wang & Nan, 2003). Thiazoles containing an isopropyl group were recently incorporated in the synthesis of minor-groove binders and this has led to a new class of potent antibacterial and antifungal compounds (Khalaf *et al.*, 2004; Antony *et al.*, 2004).

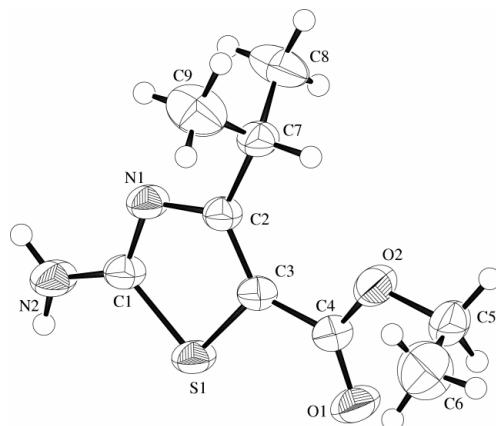


The molecular structure of (I) is unexceptional, with all ring bond lengths and angles (Table 1) close to the mean values obtained from 22 related fragments in the Cambridge Structural Database (Version 5.25, with updates to April 2004; Allen, 2002). Steric repulsion between the adjacent isopropyl and ester groups causes the main deviation from ideal geometry, widening the angles C2–C3–C4 and C3–C2–C7 to 133.90 (14) and 127.20 (14) $^\circ$, respectively. However, all geometric parameters are in excellent agreement with those found for the 4-phenyl analogue of (I) (Lynch & McClenaghan, 2000). Indeed, the similarity of these two structures extends to their hydrogen-bonding motifs. Compound (I) mimics its analogue in forming hydrogen-bonded centrosymmetric dimers *via* near-linear $\text{N}-\text{H}\cdots\text{N}$ contacts (Table 2), the supramolecular network being completed by $\text{N}-\text{H}\cdots\text{O}$ contacts.

Experimental

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Bromine (10.5 g, 65.3 mmol) was added to a stirred suspension of ethyl 4-methyl-3-oxopentanoate (10.0 g, 63.2 mmol) in water (50 ml)

**Figure 1**

The molecular structure of (I), with 50% probability displacement ellipsoids.

at 273 K over a period of 45 min. After a further 30 min at 273 K, the reaction mixture was extracted with diethyl ether (150 ml). The organic layer was then dried over magnesium sulfate and filtered. The solvent was removed under reduced pressure to give ethyl 2-bromo-4-methyl-3-oxopentanoate as an oil (14.6 g, 61.1 mmol). This oil was added to a solution of thiourea (4.7 g, 61.1 mmol) in ethanol (50 ml). The reaction mixture was kept under reflux for 1 h. Ice-water (250 ml) was added and the mixture was basified with 18 M aqueous ammonia with vigorous stirring. The insoluble material was filtered off, washed with water and dried under reduced pressure at 303 K overnight. This gave the desired product as a pale-yellow crystalline material [6.7 g, 49% yield; m.p. 449–451 K, literature m.p. 449–451 K (Barton *et al.*, 1982)]. ^1H NMR (CDCl_3): 1.25 (6H, *d*, $J = 8.0$ Hz), 1.33 (3H, *t*, $J = 7.1$ Hz), 3.88 (1H, *hept*, $J = 6.9$ Hz), 4.27 (2H, *q*, $J = 7.1$ Hz), 5.41 (2H, *s*). IR (KBr): 3393, 3112, 1665, 1531, 1509, 1466, 1307, 1134, 1034 cm^{-1} .

Crystal data

$\text{C}_9\text{H}_{14}\text{N}_2\text{O}_2\text{S}$	$D_x = 1.275 \text{ Mg m}^{-3}$
$M_r = 214.28$	Mo $\text{K}\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 24 reflections
$a = 7.8757 (10) \text{ \AA}$	$\theta = 18.7\text{--}19.8^\circ$
$b = 9.1080 (11) \text{ \AA}$	$\mu = 0.27 \text{ mm}^{-1}$
$c = 15.8434 (12) \text{ \AA}$	$T = 295 (2) \text{ K}$
$\beta = 100.853 (9)^\circ$	Plate, colourless
$V = 1116.1 (2) \text{ \AA}^3$	$0.65 \times 0.40 \times 0.18 \text{ mm}$
$Z = 4$	

Data collection

Rigaku AFC-7S diffractometer	$R_{\text{int}} = 0.023$
$\omega/2\theta$ scans	$\theta_{\text{max}} = 28.0^\circ$
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	$h = 0 \rightarrow 10$
$T_{\text{min}} = 0.760$, $T_{\text{max}} = 0.950$	$k = 0 \rightarrow 11$
2849 measured reflections	$l = -20 \rightarrow 19$
2674 independent reflections	3 standard reflections every 150 reflections
2119 reflections with $I > 2\sigma(I)$	intensity decay: none

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0493P)^2 + 0.2832P]$
$R[F^2 > 2\sigma(F^2)] = 0.038$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.109$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.23 \text{ e \AA}^{-3}$
2674 reflections	$\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$
138 parameters	

H atoms treated by a mixture of independent and constrained refinement

Table 1
Selected geometric parameters (\AA , $^\circ$).

S1—C1	1.7350 (15)	N1—C1	1.3226 (18)
S1—C3	1.7468 (15)	N1—C2	1.371 (2)
O1—C4	1.2149 (19)	N2—C1	1.331 (2)
O2—C4	1.3357 (19)	C2—C3	1.3707 (19)
C1—S1—C3	88.68 (7)	C3—C2—C7	127.20 (14)
C1—N1—C2	111.26 (12)	N1—C2—C7	117.60 (12)
N1—C1—N2	123.58 (14)	C2—C3—C4	133.90 (14)
N1—C1—S1	114.78 (12)	C2—C3—S1	110.09 (12)
N2—C1—S1	121.64 (12)	C4—C3—S1	116.01 (11)
C3—C2—N1	115.18 (13)		
C2—N1—C1—S1	-1.08 (18)	N1—C2—C3—S1	0.45 (18)
C3—S1—C1—N1	1.13 (13)	C1—S1—C3—C2	-0.85 (12)
C1—N1—C2—C3	0.4 (2)		

Table 2
Hydrogen-bonding geometry (\AA , $^\circ$).

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
N2—H1 \cdots O1 ⁱ	0.85 (2)	2.08 (2)	2.902 (2)	163 (2)
N2—H2 \cdots N1 ⁱⁱ	0.86 (2)	2.15 (2)	3.006 (2)	177 (2)

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

The amine H atoms were located in a difference map and refined isotropically; all other H atoms were constrained to idealized geometry with a riding model, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$. C—H distances: $\text{CH}_3 = 0.96 \text{ \AA}$, $\text{CH}_2 = 0.97 \text{ \AA}$ and $\text{CH} = 0.98 \text{ \AA}$.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1988); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1992); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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(I)

Crystal data

C₉H₁₄N₂O₂S
 $M_r = 214.28$
Monoclinic, P2₁/n
 $a = 7.8757$ (10) Å
 $b = 9.1080$ (11) Å
 $c = 15.8434$ (12) Å
 $\beta = 100.853$ (9)°
 $V = 1116.1$ (2) Å³
 $Z = 4$

$F(000) = 456$
 $D_x = 1.275$ Mg m⁻³
Mo K α radiation, $\lambda = 0.71069$ Å
Cell parameters from 24 reflections
 $\theta = 18.7\text{--}19.8^\circ$
 $\mu = 0.27$ mm⁻¹
 $T = 295$ K
Plate, colourless
0.65 × 0.40 × 0.18 mm

Data collection

Rigaku AFC-7S
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 $\omega/2\theta$ scans
Absorption correction: ψ scan
(North et al., 1968)
 $T_{\min} = 0.760$, $T_{\max} = 0.950$
2849 measured reflections

2674 independent reflections
2119 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\max} = 28.0^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = 0 \rightarrow 10$
 $k = 0 \rightarrow 11$
 $l = -20 \rightarrow 19$
3 standard reflections every 150 reflections
intensity decay: none

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.109$
 $S = 1.03$
2674 reflections
138 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 atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[c^2(F_o^2) + (0.0493P)^2 + 0.2832P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻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>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.35390 (5)	0.22977 (5)	0.31256 (2)	0.04375 (14)
O1	0.47481 (17)	-0.03494 (16)	0.23705 (8)	0.0606 (4)
O2	0.70278 (16)	-0.08324 (14)	0.34096 (8)	0.0531 (3)
N1	0.55929 (16)	0.32156 (15)	0.44919 (8)	0.0420 (3)
N2	0.3123 (2)	0.46619 (18)	0.40776 (11)	0.0540 (4)
H1	0.222 (3)	0.483 (2)	0.3702 (13)	0.056 (6)*
H2	0.345 (3)	0.526 (2)	0.4493 (14)	0.061 (6)*
C1	0.41151 (19)	0.35188 (17)	0.39715 (10)	0.0402 (3)
C2	0.63246 (18)	0.19615 (17)	0.42422 (9)	0.0370 (3)
C3	0.54117 (18)	0.12966 (17)	0.35213 (9)	0.0379 (3)
C4	0.5675 (2)	-0.00187 (19)	0.30463 (10)	0.0421 (3)
C5	0.7327 (3)	-0.2197 (2)	0.29860 (13)	0.0588 (5)
H5A	0.8551	-0.2427	0.3109	0.088*
H5B	0.6975	-0.2083	0.2369	0.088*
C6	0.6343 (4)	-0.3418 (3)	0.3283 (2)	0.0880 (8)
H6A	0.6587	-0.3454	0.3900	0.132*
H6B	0.6678	-0.4329	0.3057	0.132*
H6C	0.5128	-0.3261	0.3085	0.132*
C7	0.8035 (2)	0.14796 (19)	0.47675 (10)	0.0446 (4)
H7	0.8388	0.0574	0.4514	0.067*
C8	0.9424 (2)	0.2634 (3)	0.47442 (16)	0.0726 (6)
H8A	0.9522	0.2819	0.4159	0.109*
H8B	1.0511	0.2284	0.5058	0.109*
H8C	0.9115	0.3525	0.5001	0.109*
C9	0.7856 (3)	0.1150 (3)	0.56843 (13)	0.0739 (7)
H9A	0.7498	0.2021	0.5943	0.111*
H9B	0.8949	0.0826	0.6006	0.111*
H9C	0.7008	0.0393	0.5685	0.111*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0385 (2)	0.0470 (2)	0.0396 (2)	0.00162 (16)	-0.00809 (15)	-0.00464 (16)
O1	0.0576 (7)	0.0646 (8)	0.0504 (7)	0.0054 (6)	-0.0135 (6)	-0.0192 (6)
O2	0.0550 (7)	0.0518 (7)	0.0471 (6)	0.0117 (6)	-0.0041 (5)	-0.0141 (5)
N1	0.0384 (6)	0.0419 (7)	0.0405 (7)	0.0034 (5)	-0.0062 (5)	-0.0049 (5)
N2	0.0453 (8)	0.0508 (9)	0.0565 (9)	0.0124 (6)	-0.0141 (7)	-0.0123 (7)
C1	0.0366 (7)	0.0402 (8)	0.0402 (8)	-0.0005 (6)	-0.0021 (6)	-0.0012 (6)
C2	0.0340 (7)	0.0391 (7)	0.0355 (7)	-0.0006 (6)	0.0006 (5)	-0.0009 (6)
C3	0.0348 (7)	0.0413 (8)	0.0350 (7)	-0.0008 (6)	0.0004 (5)	-0.0003 (6)

C4	0.0401 (7)	0.0461 (8)	0.0386 (7)	-0.0031 (6)	0.0031 (6)	-0.0049 (6)
C5	0.0586 (11)	0.0555 (11)	0.0601 (11)	0.0114 (9)	0.0052 (9)	-0.0196 (9)
C6	0.0876 (17)	0.0597 (14)	0.118 (2)	-0.0006 (13)	0.0237 (16)	-0.0137 (14)
C7	0.0382 (8)	0.0476 (9)	0.0429 (8)	0.0063 (6)	-0.0055 (6)	-0.0062 (7)
C8	0.0383 (9)	0.0989 (17)	0.0746 (14)	-0.0089 (10)	-0.0046 (9)	0.0152 (12)
C9	0.0611 (12)	0.0956 (17)	0.0594 (12)	0.0074 (12)	-0.0033 (9)	0.0329 (12)

Geometric parameters (\AA , $^\circ$)

S1—C1	1.7350 (15)	C5—H5A	0.970
S1—C3	1.7468 (15)	C5—H5B	0.970
O1—C4	1.2149 (19)	C6—H6A	0.960
O2—C4	1.3357 (19)	C6—H6B	0.960
O2—C5	1.453 (2)	C6—H6C	0.960
N1—C1	1.3226 (18)	C7—C9	1.516 (3)
N1—C2	1.371 (2)	C7—C8	1.522 (3)
N2—C1	1.331 (2)	C7—H7	0.980
N2—H1	0.85 (2)	C8—H8A	0.960
N2—H2	0.86 (2)	C8—H8B	0.960
C2—C3	1.3707 (19)	C8—H8C	0.960
C2—C7	1.5093 (19)	C9—H9A	0.960
C3—C4	1.450 (2)	C9—H9B	0.960
C5—C6	1.482 (3)	C9—H9C	0.960
C1—S1—C3	88.68 (7)	C5—C6—H6A	109.5
C4—O2—C5	117.46 (13)	C5—C6—H6B	109.5
C1—N1—C2	111.26 (12)	H6A—C6—H6B	109.5
C1—N2—H1	118.9 (14)	C5—C6—H6C	109.5
C1—N2—H2	119.7 (14)	H6A—C6—H6C	109.5
H1—N2—H2	121 (2)	H6B—C6—H6C	109.5
N1—C1—N2	123.58 (14)	C2—C7—C9	110.59 (14)
N1—C1—S1	114.78 (12)	C2—C7—C8	110.90 (15)
N2—C1—S1	121.64 (12)	C9—C7—C8	110.91 (17)
C3—C2—N1	115.18 (13)	C2—C7—H7	108.1
C3—C2—C7	127.20 (14)	C9—C7—H7	108.1
N1—C2—C7	117.60 (12)	C8—C7—H7	108.1
C2—C3—C4	133.90 (14)	C7—C8—H8A	109.5
C2—C3—S1	110.09 (12)	C7—C8—H8B	109.5
C4—C3—S1	116.01 (11)	H8A—C8—H8B	109.5
O1—C4—O2	122.72 (16)	C7—C8—H8C	109.5
O1—C4—C3	122.75 (15)	H8A—C8—H8C	109.5
O2—C4—C3	114.53 (13)	H8B—C8—H8C	109.5
O2—C5—C6	110.80 (18)	C7—C9—H9A	109.5
O2—C5—H5A	109.5	C7—C9—H9B	109.5
C6—C5—H5A	109.5	H9A—C9—H9B	109.5
O2—C5—H5B	109.5	C7—C9—H9C	109.5
C6—C5—H5B	109.5	H9A—C9—H9C	109.5
H5A—C5—H5B	108.1	H9B—C9—H9C	109.5

C2—N1—C1—N2	178.68 (16)	C5—O2—C4—O1	-2.8 (3)
C2—N1—C1—S1	-1.08 (18)	C5—O2—C4—C3	177.58 (15)
C3—S1—C1—N1	1.13 (13)	C2—C3—C4—O1	-174.00 (18)
C3—S1—C1—N2	-178.64 (16)	S1—C3—C4—O1	6.4 (2)
C1—N1—C2—C3	0.4 (2)	C2—C3—C4—O2	5.6 (3)
C1—N1—C2—C7	179.22 (14)	S1—C3—C4—O2	-173.96 (11)
N1—C2—C3—C4	-179.14 (16)	C4—O2—C5—C6	-87.8 (2)
C7—C2—C3—C4	2.2 (3)	C3—C2—C7—C9	-119.71 (19)
N1—C2—C3—S1	0.45 (18)	N1—C2—C7—C9	61.6 (2)
C7—C2—C3—S1	-178.24 (13)	C3—C2—C7—C8	116.8 (2)
C1—S1—C3—C2	-0.85 (12)	N1—C2—C7—C8	-61.8 (2)
C1—S1—C3—C4	178.82 (13)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N2—H1···O1 ⁱ	0.85 (2)	2.08 (2)	2.902 (2)	163 (2)
N2—H2···N1 ⁱⁱ	0.86 (2)	2.15 (2)	3.006 (2)	177 (2)

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