# organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 295 KMean  $\sigma$ (C–C) = 0.003 Å 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.

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

Both the molecular and the crystal structures of the title compound,  $C_9H_{14}N_2O_2S$ , are similar to those of its 4-phenyl analogue. The supramolecular network is based upon N-H···N hydrogen-bonded centrosymmetric dimers linked by N-H···O contacts.

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## Comment

The quest for bioactive compounds led us to the synthesis of a variety of heterocylic 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)°, 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 N-H···N contacts (Table 2), the supramolecular network being completed by  $N-H \cdots O$ contacts.

### **Experimental**

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved 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)]. <sup>1</sup>H NMR (CDCl<sub>3</sub>): 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,  $1034 \text{ cm}^{-1}$ .

#### Crystal data

$C_9H_{14}N_2O_2S$	$D_x = 1.275 \text{ Mg m}^{-2}$
$M_r = 214.28$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters fro
a = 7.8757 (10)  Å	reflections
b = 9.1080(11) Å	$\theta = 18.7  19.8^{\circ}$
c = 15.8434 (12) Å	$\mu = 0.27 \text{ mm}^{-1}$
$\beta = 100.853 (9)^{\circ}$	T = 295 (2)  K
V = 1116.1 (2) Å <sup>3</sup>	Plate, colourless
Z = 4	$0.65 \times 0.40 \times 0.18$
Data collection	
Rigaku AFC-7S diffractometer	$R_{\rm int} = 0.023$
$\omega/2\theta$ scans	$\theta_{\rm max} = 28.0^{\circ}$
Absorption correction: $\psi$ scan	$h = 0 \rightarrow 10$
(North <i>et al.</i> 1968)	$k = 0 \rightarrow 11$

orth *et al.*, 1968  $T_{\rm min}=0.760,\ T_{\rm max}=0.950$ 2849 measured reflections 2674 independent reflections 2119 reflections with  $I > 2\sigma(I)$ 

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.038$  $wR(F^2) = 0.109$ S = 1.032674 reflections 138 parameters H atoms treated by a mixture of independent and constrained refinement

om 24 mm

 $l = -20 \rightarrow 19$ 3 standard reflections every 150 reflections intensity decay: none

 $w = 1/[\sigma^2(F_o^2) + (0.0493P)^2]$ + 0.2832P] where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.23 \text{ e} \text{ Å}^{-3}$  $\Delta \rho_{\rm min} = -0.26 \text{ e} \text{ Å}^{-3}$ 

Table 1			
Selected	geometric parameters	(Å,	°).

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)	$C_{3}-C_{2}-C_{7}$	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	(Å,	°).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$\frac{1}{N2-H1\cdots O1^{i}}$	0.85 (2) 0.86 (2)	2.08 (2) 2.15 (2)	2.902 (2) 3.006 (2)	163 (2) 177 (2)
Symmetry codes: (i)	$\frac{1}{2} - x, \frac{1}{2} + y, \frac{1}{2} -$	z; (ii) $1 - x, 1 - z$	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  $U_{iso}(H) = 1.5U_{eq}(C)$ . C-H distances:  $CH_3 = 0.96$  Å,  $CH_2 = 0.97$  Å and CH = 0.98 Å.

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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