

**4-Allyl-3-(pyrazin-2-yl)- $\Delta^2$ -1H-1,2,4-triazole-5(4H)-thione****Irena Wawrzycka-Gorczyca,<sup>a\*</sup>**  
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iwawrz@hermes.umcs.lublin.pl**Key indicators**

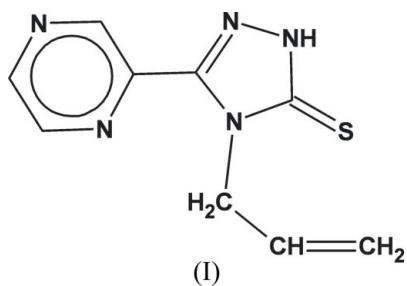
Single-crystal X-ray study  
 $T = 293\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$   
 $R$  factor = 0.043  
 $wR$  factor = 0.131  
Data-to-parameter ratio = 13.0

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

The molecule of the title compound,  $C_9H_9N_5S$ , has a non-planar conformation. In the crystal structure, molecules are connected by  $N-\text{H}\cdots N$  hydrogen bonds along the  $b$  axis. Weak  $C-\text{H}\cdots S$  interactions complete the crystal packing.

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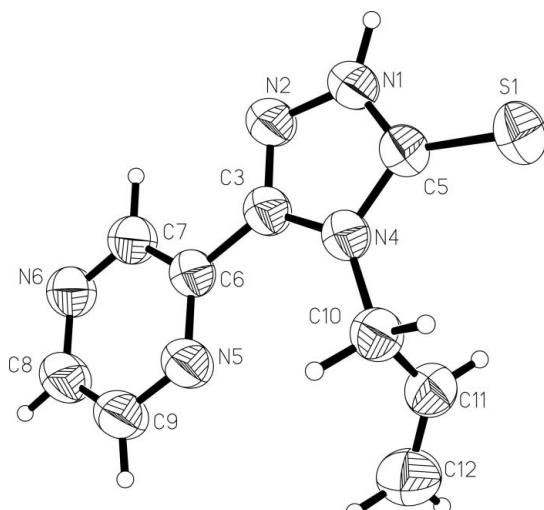
Organic compounds containing aromatic heterocyclic rings have received considerable attention among medicinal chemists because many of them play a role in various biochemical processes. 1,2,4-Triazole and pyrazine derivatives belong to an aromatic heterocyclic group exhibiting a wide range of biological activities, such as antifungal (Demirayak *et al.*, 2000; Doležal *et al.*, 2000; Doležal *et al.*, 2003), antibacterial (Pandeya *et al.*, 2000), anticancer (Invidiata *et al.*, 1991), antiviral (Todoulou *et al.*, 1994), anti-inflammatory (Sahin *et al.*, 2001), antitubercular (Doležal *et al.*, 1996). Chemical modifications leading to the combination of two or more heterocyclic and non-heterocyclic systems produce compounds of significantly enhanced biological profile compared to the parent nuclei. We have therefore combined the 1,2,4-triazole group with the pyrazin-2-yl nucleus, since both of these systems possess well documented biological activities. We present here the crystal structure of 4-allyl-3-(pyrazin-2-yl)- $\Delta^2$ -1,2,4-triazoline-5-thione, (I) (Fig. 1).



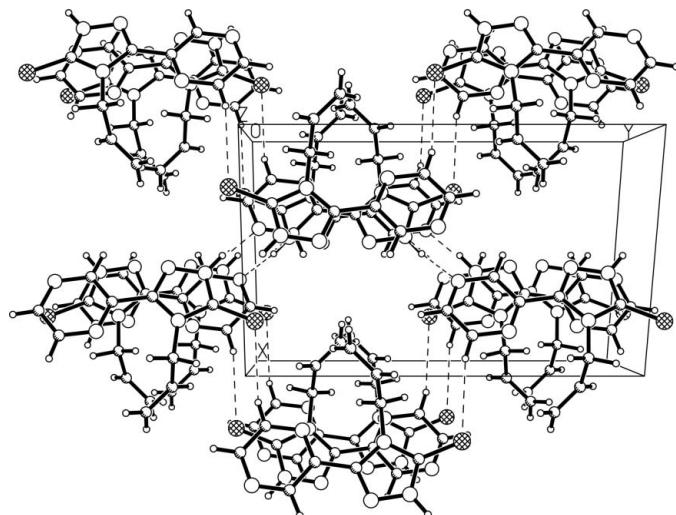
The triazole plane forms dihedral angles of  $76.8(2)^\circ$  and  $6.1(1)^\circ$  with the propene and pyrazine planes, respectively. In the crystal structure, molecules are linked by intermolecular  $N-\text{H}\cdots N$  (triazole $\cdots$ pyrazine) hydrogen bonds and  $C-\text{H}\cdots S$  (pyrazine $\cdots$ triazole) weak interactions (Fig. 2 and Table 2) along the [010] direction.

**Experimental**

Yellow crystals of (I) (Dobosz *et al.*, 2006) were obtained by slow evaporation of an ethanol solution at room temperature.

**Figure 1**

The molecular structure of (I), with the atom numbering. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

The crystal packing of (I). Dashed lines indicate hydrogen bonds.

#### Crystal data

$C_9H_9N_5S$

$M_r = 219.27$

Monoclinic,  $P2_1/c$

$a = 9.736 (2) \text{ \AA}$

$b = 14.750 (2) \text{ \AA}$

$c = 7.913 (2) \text{ \AA}$

$\beta = 112.57 (3)^\circ$

$V = 1049.3 (4) \text{ \AA}^3$

$Z = 4$

#### Data collection

Kuma KM-4 four-circle diffractometer

$\omega$ - $2\theta$  scans

Absorption correction: spherical (Dwiggins, 1975) with modifications.

$T_{\min} = 0.129$ ,  $T_{\max} = 0.236$

2367 measured reflections

2242 independent reflections

$D_x = 1.388 \text{ Mg m}^{-3}$

$\text{Cu K}\alpha$  radiation

Cell parameters from 97 reflections

$\theta = 6\text{--}20.5^\circ$

$\mu = 2.54 \text{ mm}^{-1}$

$T = 293 (2) \text{ K}$

Prism, yellow

$0.6 \times 0.25 \times 0.1 \text{ mm}$

1213 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.055$

$\theta_{\max} = 80.3^\circ$

$h = 0 \rightarrow 11$

$k = -18 \rightarrow 0$

$l = -9 \rightarrow 9$

3 standard reflections

every 100 reflections

intensity decay: 1.4%

#### Refinement

Refinement on  $F^2$

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.131$

$S = 1.03$

2242 reflections

173 parameters

All H-atom parameters refined

$$w = 1/[\sigma^2(F_o^2) + (0.0796P)^2]$$

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

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

$$\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\text{min}} = -0.19 \text{ e \AA}^{-3}$$

Extinction correction: *SHELXL97*

Extinction coefficient: 0.0130 (12)

**Table 1**  
Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

S1—C5	1.671 (2)	N4—C3	1.380 (3)
N1—C5	1.335 (3)	N4—C10	1.475 (3)
N1—N2	1.367 (3)	C10—C11	1.479 (4)
N2—C3	1.303 (3)	C11—C12	1.304 (4)
N4—C5	1.378 (3)		
C10—N4—C3—C6	2.3 (4)	C3—N4—C10—C11	83.9 (3)
N4—C3—C6—N5	5.8 (4)	N4—C10—C11—C12	-128.6 (3)

**Table 2**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1N $\cdots$ N6 <sup>i</sup>	0.95 (3)	1.91 (3)	2.844 (3)	168 (3)
C9—H9 $\cdots$ S1 <sup>ii</sup>	0.92 (3)	2.92 (3)	3.780 (3)	156 (3)

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

All H atoms were found in a difference Fourier map and were refined isotropically.

Data collection: *KM-4 Software* (Kuma, 1991); cell refinement: *KM-4 Software*; data reduction: *KM-4 Software*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL/PC* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL97*.

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