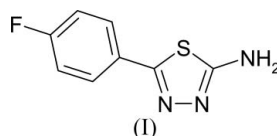
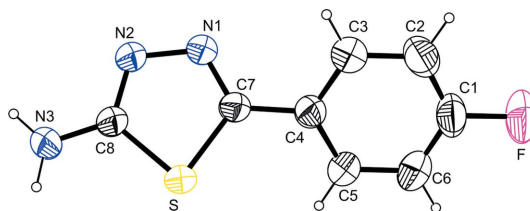


**Rong Wan,* Feng Han, Feng Wu,
Jin-Jun Zhang and Jin-Tang Wang**Department of Applied Chemistry, College of
Science, Nanjing University of Technology, No.
5 Xinmofan Road, Nanjing, Nanjing 210009,
People's Republic of China

Correspondence e-mail: rwan01@jlonline.com

Key indicatorsSingle-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
 R factor = 0.048
 wR factor = 0.133
Data-to-parameter ratio = 14.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**5-(4-Fluorophenyl)-1,3,4-thiadiazol-2-ylamine**The title compound, $\text{C}_8\text{H}_6\text{FN}_3\text{S}$, was synthesized by the
reaction of 4-fluorobenzoic acid and thiosemicarbazide. The
two H atoms of the amine are involved in $\text{N}-\text{H}\cdots\text{N}$ hydrogen
bonding, resulting in the formation of layers parallel to the
(100) plane.Received 19 September 2006
Accepted 27 September 2006**Comment**1,3,4-Thiadiazole derivatives represent an interesting class of
compounds possessing broad-spectrum biological activities
(Nakagawa *et al.*, 1996). These compounds are known to
exhibit diverse biological effects, such as insecticidal and
fungicidal activities (Wang *et al.*, 1999) and are widely used in
the field of medicine (Sato *et al.*, 1991), for example as anti-
cancer drugs (Omar *et al.*, 1986). We are focusing our synthetic
and structural studies on thiadiazole derivatives and have
recently published the structure of 3-[5-(4-fluorophenyl)-1,3,4-
thiadiazol-2-yl]-2-phenylthiazolidin-4-one, (II) (Wan *et al.*,
2006).The structure of the title compound, (I), is shown in Fig. 1.
The thiadiazole and the fluorophenyl rings make a dihedral
angle of $30.1(2)^\circ$, much larger than that observed in the
structure of (II) [$12.7(2)^\circ$]. $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds form a pseudo-dimer using one
H atom of the NH_2 group whereas the other H atom links the
dimers, forming an infinite layer parallel to the (100) plane
(Table 1 and Fig. 2).**Experimental**4-Fluorobenzoic acid (5 mmol) and thiosemicarbazide (5 mmol) were
added to toluene (50 ml), which was heated under reflux for 4 h. The**Figure 1**A view of the molecular structure of (I), showing displacement ellipsoids
drawn at the 50% probability level. H atoms are represented as spheres of
arbitrary radii.

reaction mixture was left to cool to room temperature, poured into ice–water and filtered. The filter cake was crystallized from acetone to give pure compound (I) (m.p. 513–514 K). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of an acetone solution.

Crystal data

$C_8H_6FN_3S$
 $M_r = 195.22$
 Monoclinic, $P2_1/c$
 $a = 8.8020$ (18) Å
 $b = 8.9610$ (18) Å
 $c = 11.035$ (2) Å
 $\beta = 97.19$ (3)°
 $V = 863.5$ (3) Å³

$Z = 4$
 $D_x = 1.502$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 0.34$ mm⁻¹
 $T = 293$ (2) K
 Plate, yellow
 $0.40 \times 0.30 \times 0.10$ mm

Data collection

Enraf–Nonius CAD-4
 diffractometer
 $\omega/2\theta$ scans
 Absorption correction: ψ scan
 (North *et al.*, 1968)
 $T_{\min} = 0.875$, $T_{\max} = 0.967$
 1810 measured reflections

1699 independent reflections
 1378 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$
 $\theta_{\text{max}} = 26.0^\circ$
 3 standard reflections
 every 200 reflections
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.133$
 $S = 1.03$
 1699 reflections
 118 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0831P)^2 + 0.1422P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.38$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.33$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|------------------------|-------|-------------|-------------|---------------|
| $N3-H3A\cdots N2^i$ | 0.86 | 2.14 | 2.987 (3) | 170 |
| $N3-H3B\cdots N1^{ii}$ | 0.86 | 2.27 | 3.065 (3) | 155 |

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

All H atoms were positioned geometrically and treated as riding on their parent atoms, with $C-H = 0.93$ Å, $N-H = 0.86$ Å and $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C,N)$.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-III* (Burnett & Johnson, 1996), *ORTEP-3 for Windows* (Farrugia, 1997) and

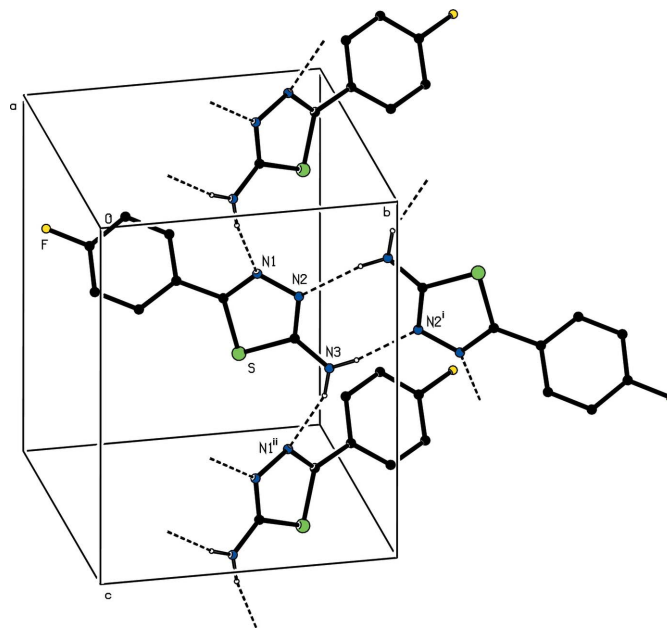


Figure 2

Partial packing view showing the hydrogen-bonded (dashed lines) network. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry codes: (i) $1 - x, 2 - y, 1 - z$; (ii) $x, \frac{3}{2} - y, \frac{1}{2} + z$.]

PLATON (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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