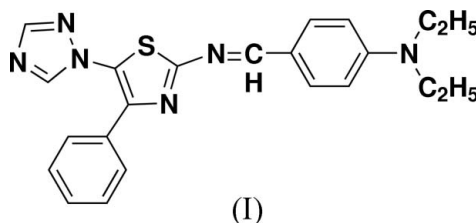


**Xin Zhou, Ling Shao, Zhong Jin,  
Jian-Bing Liu and Jian-Xin Fang\***State Key Laboratory and Institute of Elemento-  
Organic Chemistry, Nankai University, Tianjin  
300071, People's Republic of ChinaCorrespondence e-mail:  
xzhouxi\_cn@yahoo.com.cn**Key indicators**Single-crystal X-ray study  
 $T = 294$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.045  
 $wR$  factor = 0.115  
Data-to-parameter ratio = 16.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.***N*-[4-(*N,N*-Diethylamino)benzylidene]-4-phenyl-  
5-(1*H*-1,2,4-triazol-1-yl)thiazol-2-amine**The title compound,  $\text{C}_{22}\text{H}_{22}\text{N}_6\text{S}$ , has been synthesized as a  
potent fungicidal agent and its crystal structure investigated.  
The dihedral angles between the plane of the thiazole ring and  
those of the triazole, phenyl and substituted phenyl rings are  
53.0 (2), 37.5 (3) and 14.6 (2)°, respectively.

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**Comment**Thiazoles exhibit a wide variety of biological activities as  
antitumour, antifungal, antibiotic and antiviral agents  
(Hodgetts & Kershaw, 2002). In our previous work, we have  
synthesized some novel 2-aminothiazole derivatives by  
incorporating a triazole ring into 2-aminothiazole derivatives  
in order to improve the biological activity of the parent  
compounds (Shao *et al.*, 2004). We have recently designed and  
synthesized the title compound, (I), as part of our continuing  
investigation into potentially potent fungicides and report its  
structure here (Fig. 1).The title compound contains four planar rings, namely a  
thiazole ( $P1$ ), a triazole ( $P2$ ), a phenyl ( $P3$ ) and a substituted  
phenyl ( $P4$ ). The dihedral angles between  $P1$  and  $P2$ ,  $P3$  and  
 $P4$  are 53.0 (2), 37.5 (3) and 14.6 (2)°, respectively. The C11—  
N5 distance [1.389 (3) Å] is slightly longer than the corre-  
sponding distance in 2-amino-4-(2,5-dichlorophenyl)-5-(1*H*-  
1,2,4-triazol-1-yl)-1,3-thiazole [1.339 (3) Å; Shao *et al.*, 2004].  
The C11—N5—C12—C13 torsion angle is 177.70 (18)°, indi-  
cating that the substituted phenyl ring and the thiazole ring  
have a *trans* configuration.**Experimental**Three drops of piperidine were added to a mixture of 4-phenyl-5-  
(1*H*-1,2,4-triazol-1-yl)-thiazol-2-yl-amine (0.49 g, 2 mmol) and 4-  
*N,N*-diethylaminobenzaldehyde (0.36 g, 2 mmol) in anhydrous  
benzene. The solution was then refluxed for 8 h while being  
monitored using thin-layer chromatography. The solvent was concentrated  
under reduced pressure at low temperature and the precipitate was  
filtered and recrystallized from ethanol. Single crystals of (I), suitable  
for X-ray analysis, were obtained by slow diffusion of petroleum  
ether and ethyl acetate (3:1).

Crystal data

C<sub>22</sub>H<sub>22</sub>N<sub>6</sub>S  
*M<sub>r</sub>* = 402.53  
 Monoclinic, *P*2<sub>1</sub>/*c*  
*a* = 9.0753 (16) Å  
*b* = 20.924 (4) Å  
*c* = 11.518 (2) Å  
 $\beta$  = 108.639 (3)°  
*V* = 2072.5 (6) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.290 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 2309 reflections  
 $\theta$  = 2.4–23.4°  
 $\mu$  = 0.18 mm<sup>-1</sup>  
*T* = 294 (2) K  
 Parallelepiped, yellow  
 0.22 × 0.20 × 0.18 mm

Data collection

Bruker SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  
*T<sub>min</sub>* = 0.969, *T<sub>max</sub>* = 0.969  
 11567 measured reflections

4220 independent reflections  
 2444 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.052  
 $\theta_{\max}$  = 26.4°  
*h* = -11 → 11  
*k* = -26 → 21  
*l* = -14 → 8

Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.045  
*wR* (*F*<sup>2</sup>) = 0.115  
*S* = 1.00  
 4220 reflections  
 264 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0511P)^2 + 0.005P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.002$   
 $\Delta\rho_{\max} = 0.20 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.27 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

S1–C3	1.719 (2)	N4–C4	1.382 (2)
S1–C11	1.738 (2)	N5–C12	1.287 (3)
N1–C2	1.312 (3)	N5–C11	1.389 (3)
N1–C1	1.355 (3)	N6–C16	1.365 (3)
N2–N3	1.368 (2)	N6–C19	1.466 (3)
N3–C2	1.341 (3)	C3–C4	1.363 (3)
N3–C3	1.415 (2)	C4–C5	1.478 (3)
N4–C11	1.304 (3)	C12–C13	1.437 (3)
C3–S1–C11	88.17 (10)	C4–C3–S1	111.68 (15)
C2–N1–C1	101.58 (19)	C3–C4–N4	113.82 (18)
C1–N2–N3	101.45 (18)	C3–C4–C5	127.39 (18)
C2–N3–N2	109.08 (17)	N4–C11–N5	128.6 (2)
C2–N3–C3	129.98 (18)	N4–C11–S1	115.10 (16)
C11–N4–C4	111.22 (18)	N5–C12–C13	123.3 (2)
C12–N5–C11	117.26 (18)	N5–C12–H12	118.4
N2–N3–C3–C4	-126.5 (2)	C12–N5–C11–S1	-169.54 (16)
N4–C4–C5–C6	-144.2 (2)	C11–N5–C12–C13	177.70 (18)
C4–N4–C11–N5	-175.45 (19)	N5–C12–C13–C18	-174.0 (2)

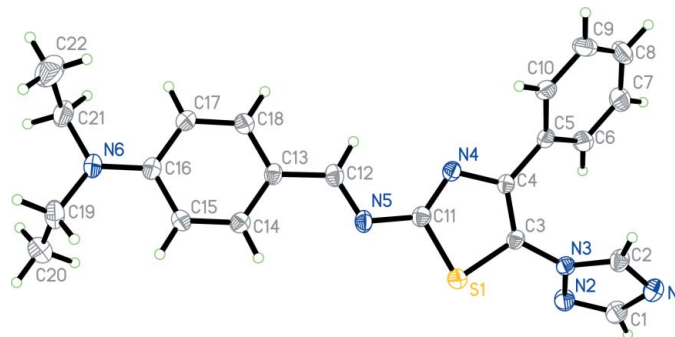


Figure 1

A view of the title compound, with displacement ellipsoids drawn at the 30% probability level.

All H atoms were placed in calculated positions, with C–H = 0.93 Å, and included in the final cycles of refinement using a riding model, with *U<sub>iso</sub>*(H) = 1.2*U<sub>eq</sub>*(C).

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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