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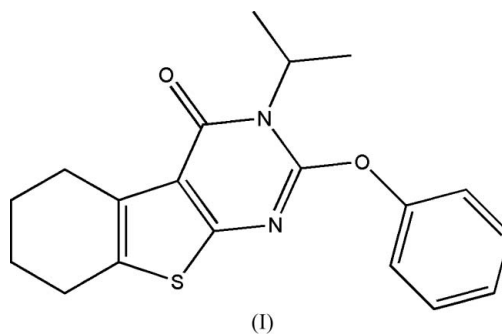
## Key indicators

Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.007$  Å  
Disorder in main residue  
 $R$  factor = 0.060  
 $wR$  factor = 0.169  
Data-to-parameter ratio = 15.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.3-Isopropyl-2-phenoxy-5,6,7,8-tetrahydro-1-benzo-  
thieno[2,3-*d*]pyrimidin-4(3*H*)-one

In the title compound,  $\text{C}_{19}\text{H}_{20}\text{N}_2\text{O}_2\text{S}$ , the central thienopyrimidine ring system is essentially planar. The molecular structure is stabilized by  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds. In the crystal packing, the molecules are linked only by van der Waals forces.

## Comment

Pyrimidine derivatives are attracting the increasing attention of the synthetic community because of the important role played by such systems in many natural products, antibiotics and drugs (Ding *et al.*, 2004). In recent years, we have been engaged in the preparation of derivatives of heterocycles *via* the aza-Wittig reaction. The title compound, (I), was synthesized and structurally characterized in this context.



In the title molecule (Fig. 1), the central thienopyrimidine ring system is essentially planar, with a maximum displacement of 0.077 (3) Å for atom N1. Selected bond lengths and angles are listed in Table 1. The C14–C19 phenyl ring is twisted by an angle of 75.90 (13)° with respect to the mean plane of the thienopyrimidine ring system.

There are three intramolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds which stabilize the molecular structure (Table 2). The crystal packing is determined by van der Waals forces, neither intermolecular hydrogen-bonding interactions nor  $\pi-\pi$  stacking interactions being observed.

## Experimental

To a solution of iminophosphorane (1.45 g, 3 mmol) in anhydrous dichloromethane (15 ml) was added isopropyl isocyanate (3 mmol) under dry nitrogen at room temperature. The reaction mixture was left unstirred for 48 h at room temperature and then the solvent was removed under reduced pressure and diethyl ether/petroleum ether (1:3 *v/v*, 20 ml) was added to precipitate triphenylphosphine oxide. After filtration, the solvent was removed, and the residue was dissolved in  $\text{CH}_3\text{CN}$  (15 ml). PhOH (3.1 mmol) and excess  $\text{K}_2\text{CO}_3$  were then added to the solution of the carbodiimide. After stirring the mixture for 18 h at room temperature, the solution was evapo-

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rated and the residue was recrystallized from EtOH to give the title compound, (I), in a yield of 70% (m.p. 422 K).  $^1\text{H}$  NMR ( $\text{CDCl}_3$ , 400 MHz): 7.46–7.17 (*m*, 5H, Ar-H), 5.63 (*m*, 1H, CH), 2.99–2.66 (*m*, 4H, 2CH<sub>2</sub>), 1.83–1.82 (*m*, 4H, 2CH<sub>2</sub>), 1.59–1.58 (*d*, *J* = 6.8 Hz, 6H, 2CH<sub>3</sub>); MS (EI, 70 eV) *m/z* (%): 340 (*M*<sup>+</sup>, 100), 298 (53), 270 (45), 255 (76), 227 (42). Elemental analysis calculated for C<sub>19</sub>H<sub>20</sub>N<sub>2</sub>O<sub>2</sub>S: C 67.03, H 5.92, N 8.23%; found: C 66.85, H 5.85, N 8.17%. Crystals suitable for single-crystal X-ray diffraction were obtained from a hexane/dichloromethane solution (1:3 v/v) at room temperature.

#### Crystal data

C<sub>19</sub>H<sub>20</sub>N<sub>2</sub>O<sub>2</sub>S

*M<sub>r</sub>* = 340.43

Monoclinic, *P*2<sub>1</sub>

*a* = 12.4411 (15) Å

*b* = 5.8030 (7) Å

*c* = 13.2870 (16) Å

$\beta$  = 113.994 (2)°

*V* = 876.37 (18) Å<sup>3</sup>

*Z* = 2

*D<sub>x</sub>* = 1.290 Mg m<sup>-3</sup>

Mo *K*α radiation

Cell parameters from 1462

reflections

$\theta$  = 2.9–21.1°

$\mu$  = 0.20 mm<sup>-1</sup>

*T* = 292 (2) K

Block, colourless

0.30 × 0.20 × 0.20 mm

#### Data collection

Bruker SMART CCD area-detector diffractometer

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

*T<sub>min</sub>* = 0.943, *T<sub>max</sub>* = 0.962

5473 measured reflections

3473 independent reflections

2596 reflections with *I* > 2σ(*I*)

*R<sub>int</sub>* = 0.025

$\theta_{\text{max}}$  = 27.0°

*h* = −13 → 15

*k* = −7 → 7

*l* = −16 → 16

#### Refinement

Refinement on *F*<sup>2</sup>

*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.060

*wR* (*F*<sup>2</sup>) = 0.169

*S* = 1.05

3473 reflections

231 parameters

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0921P)^2]$

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

( $\Delta/\sigma$ )<sub>max</sub> < 0.001

$\Delta\rho_{\text{max}} = 0.27 \text{ e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$

Absolute structure: Flack (1983),

1386 Friedel pairs

Flack parameter: −0.01 (12)

**Table 1**

Selected geometric parameters (Å, °).

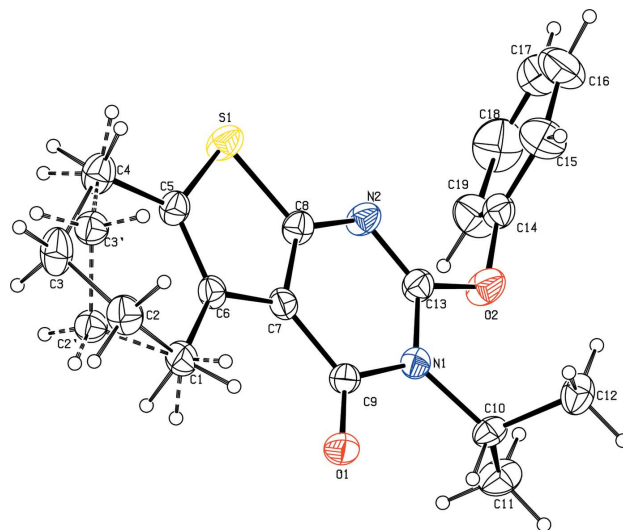
C5–C6	1.367 (5)	C8–N2	1.370 (4)
C5–S1	1.745 (4)	C8–S1	1.707 (4)
C6–C7	1.415 (5)	C9–O1	1.220 (4)
C7–C8	1.391 (4)	C9–N1	1.416 (4)
C7–C9	1.444 (5)		
C8–S1–C5	91.48 (17)		

**Table 2**

Hydrogen-bond geometry (Å, °).

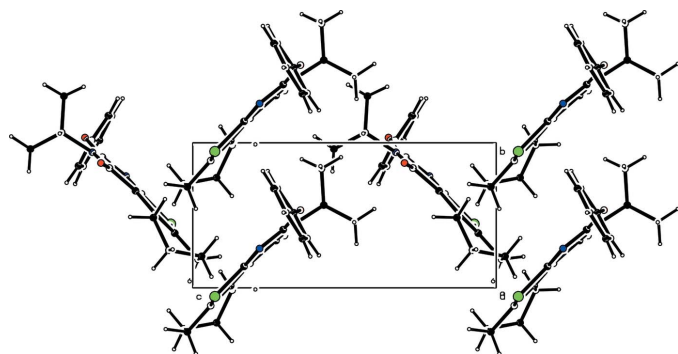
<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
C10–H10...O1	0.98 (4)	2.21 (4)	2.733 (5)	112 (3)
C11–H11C...O2	0.96	2.27	2.846 (6)	118
C12–H12A...O2	0.96	2.43	2.981 (5)	116

The positional parameters of H10, bonded to C10, were refined freely. All other H atoms were placed at calculated positions and treated as riding atoms, with C–H = 0.93–0.97 Å, and *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C) for aromatic and methylene H atoms or 1.5*U*<sub>eq</sub>(C) for methyl H atoms. The terminal cyclohexene ring is disordered. There are two possible conformations, C1–C6 and C1/C2'/C3'/C4–C6, with an occupancy ratio of 0.690 (12):0.310 (12).



**Figure 1**

View of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary size. Both disorder components are shown.



**Figure 2**

Packing diagram of the title compound, viewed along the *a* axis. Only the major conformation of the disordered cyclohexene ring is shown.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXTL (Sheldrick, 2001).

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