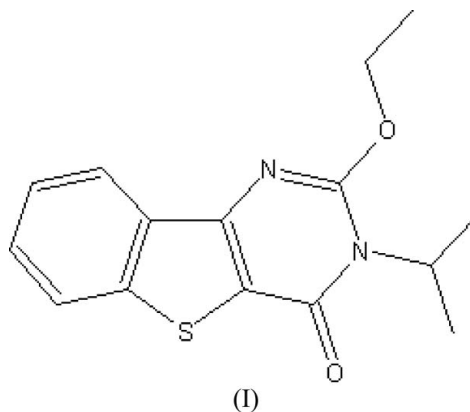
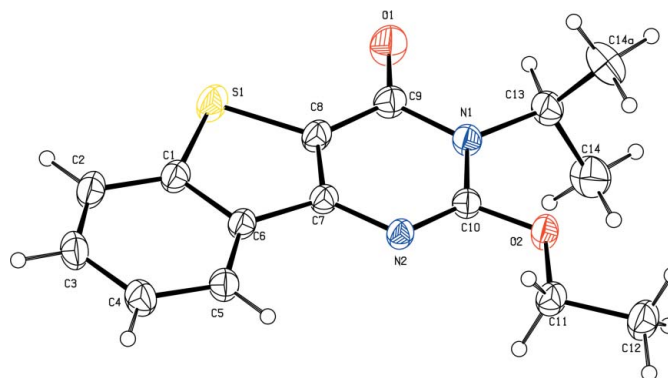


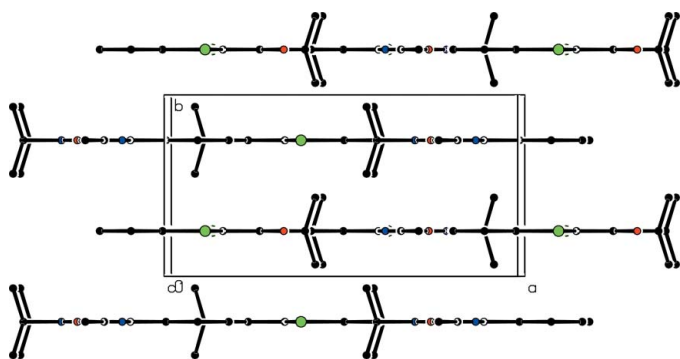
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of ChinaCorrespondence e-mail:  
mwding@mail.ccnu.edu.cn,  
mwding@mail.ccnu.edu.cn**Key indicators**Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.044  
 $wR$  factor = 0.122  
Data-to-parameter ratio = 14.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**2-Ethoxy-3-isopropylbenzo[4,5]thieno[3,2-*d*]-  
pyrimidin-4(3*H*)-one**The molecule of the title compound,  $\text{C}_{15}\text{H}_{16}\text{N}_2\text{O}_2\text{S}$ , has  
crystallographic mirror symmetry. The crystal structure is  
stabilized by inter- and intramolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen-  
bonding interactions.

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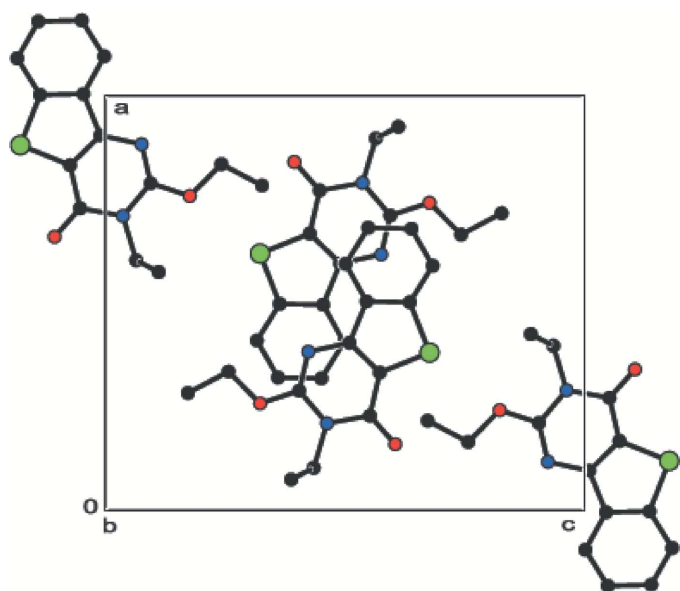
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**Comment**Thienopyrimidine derivatives are of interest as possible anti-  
viral agents, and because of their other biological properties,  
including antibacterial, antifungal, antiallergic and anti-  
inflammatory. We have recently focused on the synthesis of  
fused heterocyclic systems containing thienopyrimidine *via*  
the aza-Wittig reaction at room temperature (Ding *et al.*,  
2004). We present the X-ray crystallographic analysis of the  
title compound, (I), in this paper.As shown in Fig. 1, compound (I) contains three fused rings.  
All the ring atoms in the benzothieno[3,2-*d*]pyrimidine system  
are essentially in the same plane. Selected bond lengths and**Figure 1**  
View of the molecule of (I), showing the atom-labeling scheme.  
Displacement ellipsoids are drawn at the 50% probability level. H atoms  
are represented by circles of arbitrary size [symmetry code: (a)  $x, \frac{1}{2} - y, z$ ].



**Figure 2**  
The crystal packing, viewed along the *c* axis, showing the planar sheets of molecules parallel to (010).



**Figure 3**  
View of the packing of (I) along the *b* axis. H atoms bonded to C atoms have been omitted for clarity.

torsion angles are presented in Table 1. Some C—H...O inter- and intramolecular hydrogen bonds are present in the crystal structure (Table 1). The molecules are packed as layers parallel to the *ac* plane (Figs. 2 and 3).

## Experimental

To a solution of ethyl 3-triphenylphosphoranylidenaminobenzo[*b*]thiophene-2-carboxylate (3 mmol) in dry dichloromethane (5 ml) was added isopropyl isocyanate (3 mmol) under nitrogen at room temperature. After the reaction mixture was left to stand for 8–12 h at 273–278 K, the solvent was removed under reduced pressure and diethyl ether/petroleum ether (1:2 *v/v*, 12 ml) was added to precipitate triphenylphosphine oxide. After filtration, anhydrous ethanol (10 ml) was added with several drops of EtONa in EtOH. The mixture was stirred for 1–6 h at room temperature. The solution was then concentrated under reduced pressure and the residue was recrystallized from ethanol to afford compound (I) (yield: 65%, m.p.

446 K). <sup>1</sup>H NMR (CDCl<sub>3</sub>, 400 MHz): δ 8.19–7.45 (4H, Ar—H), 5.58 (*m*, 1H, N—CH), 4.68–4.62 (*m*, 2H, O—CH<sub>2</sub>) 1.54–1.50 (*m*, 9H, CH<sub>3</sub>). Crystals suitable for single-crystal X-ray diffraction were grown from dichloromethane at 300 K.

### Crystal data

C<sub>15</sub>H<sub>16</sub>N<sub>2</sub>O<sub>2</sub>S  
*M<sub>r</sub>* = 288.36  
 Orthorhombic, *Pnma*  
*a* = 13.3930 (16) Å  
*b* = 6.8922 (8) Å  
*c* = 15.4970 (18) Å  
*V* = 1430.5 (3) Å<sup>3</sup>  
*Z* = 4  
*D<sub>x</sub>* = 1.339 Mg m<sup>-3</sup>

Mo Kα radiation  
 Cell parameters from 1811 reflections  
 θ = 2.6–24.4°  
 μ = 0.23 mm<sup>-1</sup>  
*T* = 292 (2) K  
 Block, colorless  
 0.30 × 0.20 × 0.20 mm

### Data collection

Bruker SMART 4K CCD area-detector diffractometer  
 φ and ω scans  
 Absorption correction: none  
 8299 measured reflections  
 1768 independent reflections

1339 reflections with *I* > 2σ(*I*)  
*R*<sub>int</sub> = 0.039  
 θ<sub>max</sub> = 27.5°  
*h* = -17 → 16  
*k* = -8 → 8  
*l* = -20 → 15

### Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.044  
*wR*(*F*<sup>2</sup>) = 0.122  
*S* = 1.06  
 1768 reflections  
 120 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0653P)^2 + 0.1581P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 (Δ/σ)<sub>max</sub> < 0.001  
 Δρ<sub>max</sub> = 0.29 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.24 e Å<sup>-3</sup>

**Table 1**  
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>
C14—H14A...O2	0.96	2.40	2.941 (2)	115
C13—H13...O1	0.98	2.22	2.744 (3)	112
C2—H2...O1 <sup>i</sup>	0.93	2.53	3.407 (3)	157

Symmetry code: (i)  $x - \frac{1}{2}, -y + \frac{1}{2}, -z + \frac{1}{2}$ .

H atoms were positioned geometrically and treated as riding, with C—H = 0.93–0.98 Å and *U*<sub>iso</sub>(H) values set equal to *xU*<sub>eq</sub>(carrier atom), with *x* = 1.2 for *Csp*<sup>2</sup> and *x* = 1.5 for methyl C parent atoms.

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

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