

(E)-N'-[4-(4-Chlorobenzoyloxy)-3-methoxybenzylidene]thiophene-2-carbohydrazide**Yan-Li Zhao,* Qiao-Zhen Zhang,
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In the title compound, $C_{20}H_{17}ClN_2O_3S$, the vanillin group makes dihedral angles of 73.65 (11) and 2.16 (16) $^\circ$ with the chloro-substituted benzene ring and the thiophene mean plane, respectively. The crystal packing is stabilized by an intermolecular N—H \cdots O hydrogen bond that forms a centrosymmetric dimer.

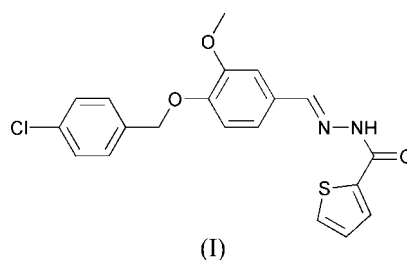
Received 23 October 2006
Accepted 2 November 2006**Key indicators**

Single-crystal X-ray study
 $T = 294$ K
 Mean $\sigma(C-C) = 0.007$ Å
 R factor = 0.056
 wR factor = 0.176
 Data-to-parameter ratio = 13.3

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

Comment

For the background to this study, see the first paper in this series of three (Zhang *et al.*, 2006). We report here the synthesis and structure of the title compound, (I) (Fig. 1).



All bond lengths and angles in (I) are within normal ranges (Allen *et al.*, 1987). The vanillin group (C8–C13/C15/O1/O2) is nearly planar, with an r.m.s. deviation for fitted atoms of 0.0280 Å. This plane makes dihedral angles of 2.16 (16) and 73.65 (11) $^\circ$ with the thiophene ring (C17–C20/S1) and the terminal benzene ring (C1–C6), respectively. The dihedral angle between the thiophene and benzene rings is 72.24 (14) $^\circ$.

The crystal packing is stabilized by an intermolecular N—H \cdots O=C hydrogen bond (Table 1) that forms a centrosymmetric dimer (Fig. 2).

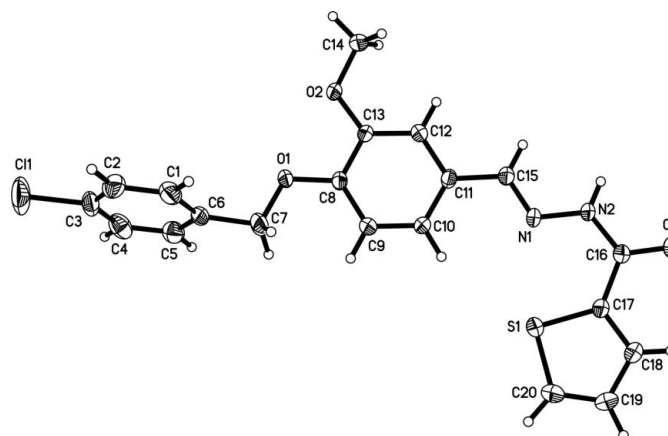


Figure 1
The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level.

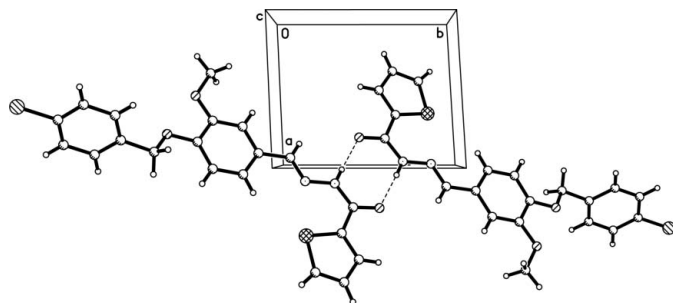


Figure 2
A packing diagram of (I), viewed along the *c* axis, showing hydrogen bonds as dashed lines.

Experimental

An anhydrous ethanol solution (50 ml) of 4-(4-chlorobenzoyloxy)-3-methoxybenzaldehyde (2.77 g, 10 mmol) was added to an anhydrous ethanol solution (50 ml) of thiophene-2-carbohydrazide (1.42 g, 10 mmol) and the mixture was stirred at 350 K for 5 h under nitrogen, giving a white precipitate. The product was isolated, recrystallized from ethanol and dried in a vacuum to give (I) in 82% yield. Colourless single crystals suitable for X-ray analysis were obtained by slow evaporation of an acetonitrile solution.

Crystal data

$C_{20}H_{17}ClN_2O_3S$	$V = 934.6 (5) \text{ \AA}^3$
$M_r = 400.88$	$Z = 2$
Triclinic, $P\bar{1}$	$D_x = 1.424 \text{ Mg m}^{-3}$
$a = 8.275 (3) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 9.884 (3) \text{ \AA}$	$\mu = 0.34 \text{ mm}^{-1}$
$c = 11.584 (3) \text{ \AA}$	$T = 294 (2) \text{ K}$
$\alpha = 89.503 (5)^\circ$	Block, colourless
$\beta = 81.062 (5)^\circ$	$0.14 \times 0.12 \times 0.08 \text{ mm}$
$\gamma = 86.951 (5)^\circ$	

Data collection

Bruker SMART APEX CCD area-detector diffractometer	4778 measured reflections
φ and ω scans	3264 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1803 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.931$, $T_{\max} = 0.973$	$R_{\text{int}} = 0.035$
	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.176$
 $S = 1.06$
 3264 reflections
 246 parameters
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0663P)^2 + 0.5087P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.39 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.39 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2\cdots O3^i$	0.86	1.95	2.811 (5)	175

Symmetry code: (i) $-x + 2, -y + 3, -z$.

H atoms were included in calculated positions, with $C-H = 0.93-0.97 \text{ \AA}$ and $N-H = 0.86 \text{ \AA}$, and refined as riding, with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C,N)$ or $1.5U_{\text{eq}}(\text{methyl } C)$.

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

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