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Key indicators

Single-crystal X-ray study

T = 298 K

Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$

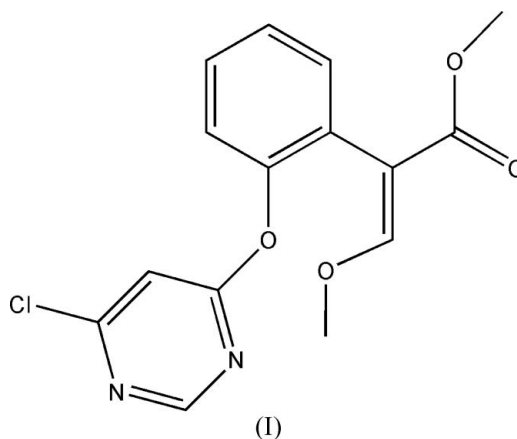
R factor = 0.050

wR factor = 0.136

Data-to-parameter ratio = 14.4

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**(E)-Methyl 2-[2-(6-chloropyrimidin-4-yloxy)-
phenyl]-3-methoxyacrylate**In the molecule of the title compound, $\text{C}_{15}\text{H}_{13}\text{ClN}_2\text{O}_4$, the pyrimidine and benzene rings form a dihedral angle of $69.3(3)^\circ$. No π - π stacking interactions are observed in the crystal structure.Received 16 December 2006
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Comment

The title compound, (I), can be used as an intermediate in the preparation of fungicides (Jones *et al.*, 1998). We report here the crystal structure of (I)

The molecular structure of (I) is shown in Fig. 1. Bond lengths and angles are within normal ranges (Allen *et al.*, 1987). The dihedral angle between the pyrimidine and benzene rings is $69.3(3)^\circ$. The O2–O4/C13–C17 and C7–C12 planes are inclined at an angle of $71.0(1)^\circ$. An intramolecular C–H...O hydrogen bond (Table 1) is observed in the molecular structure.

Experimental

Compound (I) was prepared according to the literature method of Jones *et al.* (1998). Single crystals were obtained by dissolving compound (I) (0.5 g) in ethanol (50 ml) and evaporating the solvent slowly at room temperature for about 10 d.

Crystal data

 $\text{C}_{15}\text{H}_{13}\text{ClN}_2\text{O}_4$ $M_r = 320.72$ Monoclinic, $P2_1/n$ $a = 13.2642(11) \text{ \AA}$ $b = 7.7971(7) \text{ \AA}$ $c = 14.2757(12) \text{ \AA}$ $\beta = 96.88(3)^\circ$ $V = 1465.8(2) \text{ \AA}^3$ $Z = 4$ $D_x = 1.453 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.28 \text{ mm}^{-1}$ $T = 298(2) \text{ K}$

Block, colourless

 $0.40 \times 0.30 \times 0.20 \text{ mm}$

Data collection

Enraf–Nonius CAD-4
diffractometer
 $\omega/2\theta$ scans
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.896$, $T_{\max} = 0.946$
2861 measured reflections

2861 independent reflections
2003 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.000$
 $\theta_{\max} = 26.0^\circ$
3 standard reflections
every 200 reflections
intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.136$
 $S = 1.00$
2861 reflections
199 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.067P)^2 + 0.5P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.26 \text{ e } \text{\AA}^{-3}$

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C17–H17A \cdots O3	0.93	2.45	2.776 (3)	101

H atoms were positioned geometrically [$C-H = 0.93$ (Csp^2) or 0.96 \AA (methyl)] and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C})$, where $x = 1.2$ for Csp^2 and $x = 1.5$ for other H atoms.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1985); 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: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

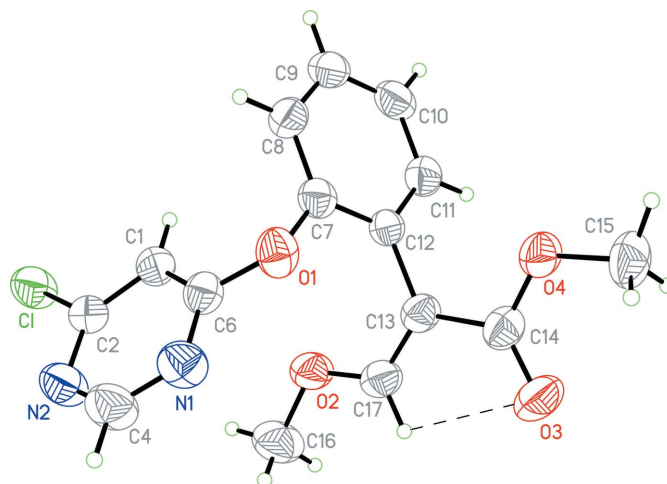


Figure 1

The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. The dashed line indicates a hydrogen bond.

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