

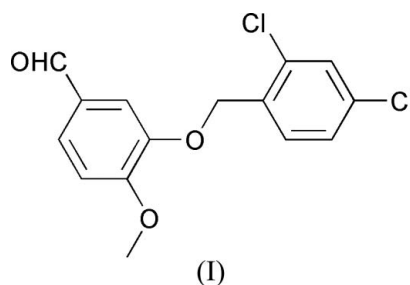
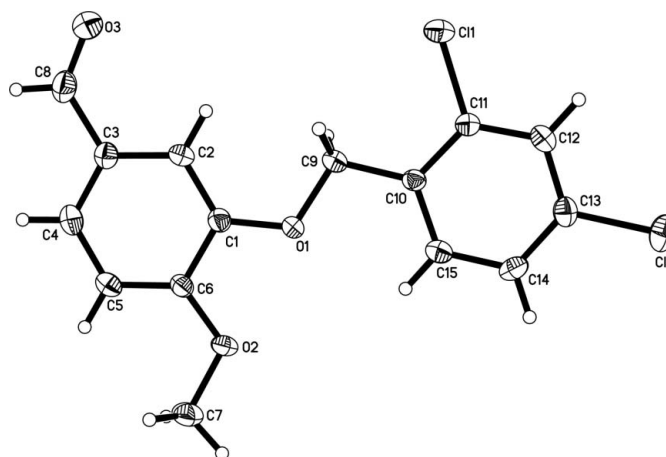
## 3-(2,4-Dichlorobenzoyloxy)-4-methoxybenzaldehyde

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## Key indicators

Single-crystal X-ray study  
 $T = 294$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.006$  Å  
 $R$  factor = 0.062  
 $wR$  factor = 0.132  
Data-to-parameter ratio = 13.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.In the title compound,  $\text{C}_{15}\text{H}_{12}\text{Cl}_2\text{O}_3$ , the isovanillin group  
makes a dihedral angle of  $3.12$  ( $12$ ) $^\circ$  with the dichlorobenzene  
ring. Intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds help to  
consolidate the crystal packing.Received 16 November 2006  
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## Comment

As part of our interest in the coordination properties of Schiff  
bases functioning as ligands, we investigated the title  
compound, (I), used as a precursor in the preparation of Schiff  
bases.In the crystal structure of (I) (Fig. 1), the bond lengths and  
angles are within their normal ranges (Allen *et al.*, 1987). The  
isovanillin group (atoms C1–C6/C8/O1/O2) is essentially  
planar, with an r.m.s. deviation for the fitted atoms of 0.016 Å.  
This plane makes a dihedral angle of  $3.12$  ( $12$ ) $^\circ$  with the mean  
plane of the C10–C15 benzene ring.The crystal packing of (I) is stabilized by weak  $\text{C}-\text{H}\cdots\text{O}$   
interactions that link adjacent molecules into zigzag chains  
running along the  $c$  axis (Table 1 and Fig. 2).**Figure 1**  
The molecular structure of (I), with displacement ellipsoids for non-H  
atoms drawn at the 30% probability level.

Experimental

An anhydrous acetonitrile solution (50 ml) of 3-hydroxy-4-methoxybenzaldehyde (1.52 g, 10 mmol) was added dropwise to a solution (100 ml) of 1-(bromomethyl)-2,4-dichlorobenzene (2.40 g, 10 mmol) and potassium carbonate (1.38 g, 10 mmol) in acetonitrile, over a period of 30 min, and the mixture refluxed for 24 h under a nitrogen atmosphere. The solvent was removed and the resultant mixture poured into ice-water (100 ml). The white precipitate was then isolated and recrystallized from acetonitrile, and then dried in a vacuum to give the pure compound in 50% yield. Colorless single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an acetonitrile solution.

Crystal data

$C_{15}H_{12}Cl_2O_3$   $Z = 4$   
 $M_r = 311.15$   $D_x = 1.466 \text{ Mg m}^{-3}$   
 Monoclinic,  $P2_1/n$  Mo  $K\alpha$  radiation  
 $a = 7.456 (3) \text{ \AA}$   $\mu = 0.46 \text{ mm}^{-1}$   
 $b = 24.850 (9) \text{ \AA}$   $T = 294 (2) \text{ K}$   
 $c = 8.370 (3) \text{ \AA}$  Block, colorless  
 $\beta = 114.616 (6)^\circ$   $0.24 \times 0.20 \times 0.16 \text{ mm}$   
 $V = 1409.9 (9) \text{ \AA}^3$

Data collection

Bruker SMART APEX CCD 6657 measured reflections  
 diffractometer 2436 independent reflections  
 $\varphi$  and  $\omega$  scans 1493 reflections with  $I > 2\sigma(I)$   
 Absorption correction: multi-scan  $R_{int} = 0.048$   
 (SADABS; Sheldrick, 1996)  $\theta_{max} = 25.0^\circ$   
 $T_{min} = 0.878, T_{max} = 0.929$

Refinement

Refinement on  $F^2$   $w = 1/[\sigma^2(F_o^2) + (0.0298P)^2 + 2.615P]$   
 $R[F^2 > 2\sigma(F^2)] = 0.062$  where  $P = (F_o^2 + 2F_c^2)/3$   
 $wR(F^2) = 0.132$   $(\Delta/\sigma)_{max} < 0.001$   
 $S = 1.03$   $\Delta\rho_{max} = 0.23 \text{ e \AA}^{-3}$   
 2436 reflections  $\Delta\rho_{min} = -0.25 \text{ e \AA}^{-3}$   
 182 parameters  
 H-atom parameters constrained

Table 1 Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C4-H4\cdots O3^i$	0.93	2.60	3.526 (5)	175

Symmetry code: (i)  $x + \frac{1}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$ .

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

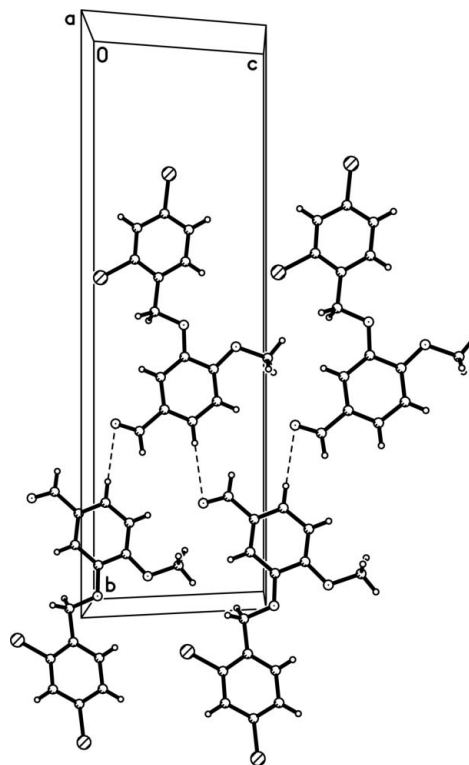


Figure 2 A partial packing diagram for (I), with hydrogen bonds shown as dashed lines.

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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