

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

Shou-Xin Liu,^{a*} Jian-Rong Han,^{b,‡}
Xiao-Li Zhen^b and Xia Tian^b^aCollege of Chemical & Pharmaceutical Engineering, Hebei University of Science & Technology, Shijiazhuang 050018, People's Republic of China, and ^bCollege of Sciences, Hebei University of Science & Technology, Shijiazhuang 050018, People's Republic of China‡ Second contact author,
email: han_jianrong@163.com

Correspondence e-mail: liu_shouxin@163.com

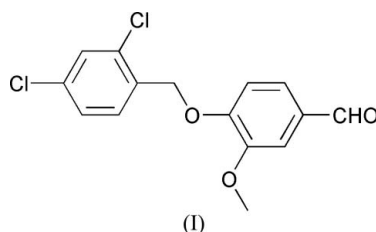
Key indicators

Single-crystal X-ray study
 $T = 294$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.044
 wR factor = 0.104
Data-to-parameter ratio = 16.2For 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 vanillin group makes a dihedral angle of $4.47(8)^\circ$ with the dichlorobenzene ring. Intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds help to consolidate the crystal structure.

Comment

As part of our interest in the coordination properties of Schiff bases functioning as ligands, we have investigated the title compound, (I), which is used as a precursor in the preparation of Schiff bases.



In (I) (Fig. 1), the bond lengths and angles are within their normal ranges (Allen *et al.*, 1987). The vanillin group (atoms C1–C6/C8/O1/O2) is almost planar, with an r.m.s. deviation for the fitted atoms of 0.016 Å. This plane makes a dihedral angle of $4.47(8)^\circ$ with the mean plane of the C10–C15 benzene ring.

The crystal structure is stabilized by weak $\text{C}-\text{H}\cdots\text{O}$ interactions that link adjacent molecules into one-dimensional chains running along the b axis (Table 1 and Fig. 2).

Experimental

An anhydrous acetonitrile solution (50 ml) of 4-hydroxy-3-methoxybenzaldehyde (1.52 g, 10 mmol) was added dropwise to a

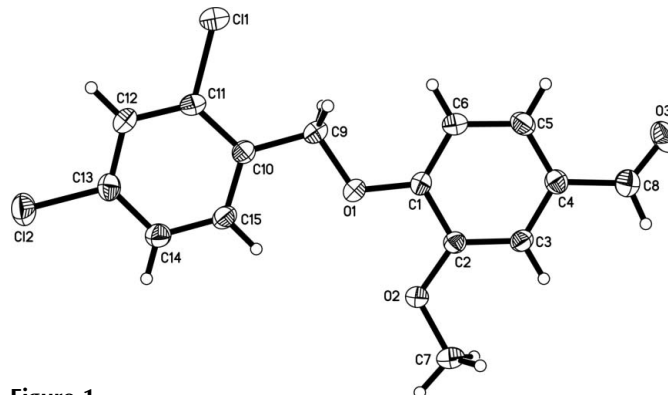


Figure 1
The molecular structure of (I), with displacement ellipsoids for the non-H atoms drawn at the 30% probability level.

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 was refluxed for 24 h under a nitrogen atmosphere. The solvent was removed and the resultant mixture poured into ice–water (100 ml). The resulting white precipitate was isolated and recrystallized from acetonitrile, and then dried in a vacuum to give the pure compound in 52% yield. Colorless single crystals of (I) suitable for X-ray crystallographic analysis were obtained by slow evaporation of an acetonitrile solution.

Crystal data

C₁₅H₁₂Cl₂O₃ Z = 4
M_r = 311.15 D_x = 1.429 Mg m⁻³
Monoclinic, P₂₁/c Mo K α radiation
a = 7.507 (4) Å μ = 0.45 mm⁻¹
b = 24.436 (15) Å T = 294 (2) K
c = 8.386 (5) Å Block, colorless
 β = 109.945 (9)° 0.20 × 0.16 × 0.10 mm
V = 1446.1 (15) Å³

Data collection

Bruker SMART APEX CCD 8038 measured reflections
diffractometer 2952 independent reflections
 φ and ω scans 1663 reflections with I > 2 σ (I)
Absorption correction: multi-scan R_{int} = 0.045
(SADABS; Sheldrick, 1996) θ_{max} = 26.5°
T_{min} = 0.903, T_{max} = 0.956

Refinement

Refinement on F² H-atom parameters constrained
R[F² > 2 σ (F²)] = 0.044 w = 1/[$\sigma^2(F_o^2) + (0.0423P)^2$]
wR(F²) = 0.104 where P = (F_o² + 2F_c²)/3
S = 1.03 (Δ/σ)_{max} = 0.003
2952 reflections $\Delta\rho_{max}$ = 0.22 e Å⁻³
182 parameters $\Delta\rho_{min}$ = -0.21 e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

D–H...A	D–H	H...A	D...A	D–H...A
C12–H12...O3 ⁱ	0.93	2.62	3.512 (3)	161

Symmetry code: (i) -x + 1, y - 1/2, -z + 1/2.

The H atoms were included in calculated positions (C–H = 0.93–0.97 Å) and refined as riding, with U_{iso}(H) = 1.2U_{eq}(C) or 1.5U_{eq}(methyl C).

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve

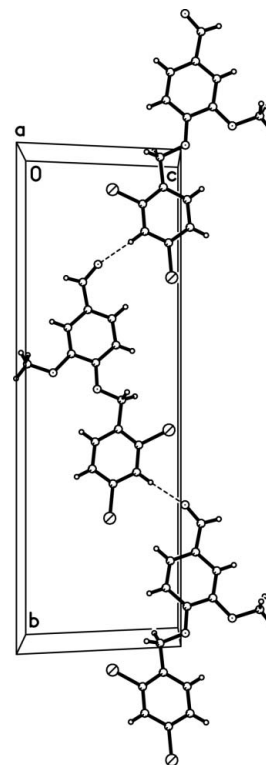


Figure 2

Partial packing diagram for (I), with hydrogen bonds shown as dashed lines.

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.

The project was supported by the Foundation of the Education Department of Hebei Province (grant No. 606022).

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
Bruker (1999). SMART (Version 5.0) and SAINT (Version 4.0) for Windows NT. Bruker AXS Inc., Madison, Wisconsin, USA.
Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
Sheldrick, G. M. (1997a). SHELXS97 and SHELXL97. University of Göttingen, Germany.
Sheldrick, G. M. (1997b). SHELXTL. Version 5.10 for Windows NT. Bruker AXS Inc., Madison, Wisconsin, USA.