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#### Key indicators

Single-crystal X-ray study  
T = 293 K  
Mean  $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$   
R factor = 0.042  
wR factor = 0.100  
Data-to-parameter ratio = 8.4

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

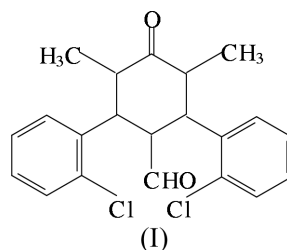
## 3,5-Bis(3-chlorophenyl)-4-formyl-2,6-dimethyl- cyclohexan-1-one

The structure of the title molecule,  $\text{C}_{21}\text{H}_{20}\text{Cl}_2\text{O}_2$ , reveals a chair conformation for the cyclohexane ring in which all the substituents occupy equatorial positions.

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#### Comment

During attempts to prepare a pyran derivative by the Michael condensation of diethyl ketone and *o*-chlorobenzaldehyde, an unexpected product, the title compound, (I) (Fig. 1), was obtained as a minor product. The present investigation was undertaken to ascertain the structure and conformation of (I), since they cannot be assigned conclusively from  $^1\text{H}$  NMR data.



The cyclohexane ring is in a chair conformation, as shown by the torsion angles (Table 1) around the C–C bonds involving the six C atoms (C1–C6). The torsion angles deviate from the angle of  $56^\circ$  expected for a perfect chair conformation (Kalsi, 1997) as a result of the  $sp^2$ -hybridized atom C1. The bond lengths and angles within the cyclohexane ring are in the ranges 1.503 (6)–1.551 (6)  $\text{\AA}$  and  $106.6$  (3)– $116.9$  (4) $^\circ$ , respectively. The equatorial orientations of all the substituents are confirmed by the torsion angles of about  $180^\circ$  formed by the external atom and the other three ring atoms. The dihedral angle between the planes of the phenyl rings is  $61.9$  (2) $^\circ$ . The chair conformation observed in (I) is comparable to that observed in the thioketal of *N*-methyl-2,6-diphenyl-3-iso-

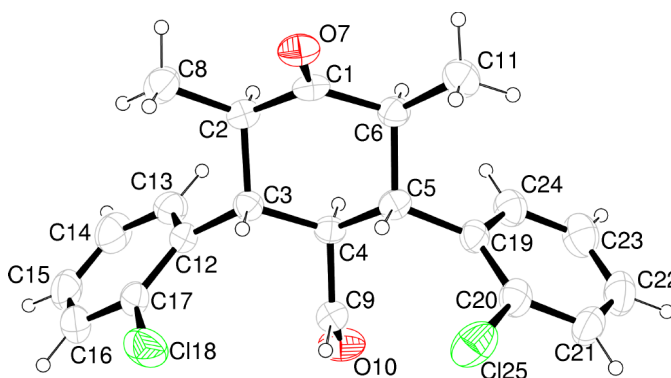


Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

propylpiperidin-4-one (Sujatha, 1995), but contrasts with the twist-boat conformation of the *N*-nitroso derivative of 2,6-bis(2-chlorophenyl)-3,5-dimethylpiperidin-4-one (Sukumar *et al.*, 1993). Atoms O10, C9, C4, O7 and C1 are coplanar, the maximum deviation being 0.014 (2) Å for O10; the molecule is nearly symmetrical with respect to this plane.

The crystal structure is stabilized by C—H···O interactions, the most significant of which are (C9)H9···O7(−1 + *x*, *y*, *z*) and (C16)H16···O7(1 − *x*,  $\frac{1}{2} + y$ ,  $\frac{1}{2} - z$ ) with H···O distances of 2.45 and 2.42 Å, and C—H···O angles of 131 and 147°, respectively. The former interactions contribute to chain formation along the *a* axis; the chains are interlinked by the latter interactions. Another intermolecular interaction exists between atoms C4 and O10( $\frac{1}{2} + x$ ,  $\frac{1}{2} - y$ , −*z*) through atom H4, with an H···O distance of 2.56 Å and an angle of 161° at H4. The formation of (I), although unexpected, can be explained by the condensation of the dibenzylideneacetone derivative formed *in situ* with the acetaldehyde present in ethanol.

## Experimental

The title compound was obtained by the reaction of diethyl ketone with *o*-chlorobenzaldehyde in 80% ethanol. Diffraction quality crystals were obtained by recrystallization from ethanol. <sup>1</sup>H NMR (CDCl<sub>3</sub>, p.p.m.): δ 0.92 (*d*, 6H, *J* = 6.3), 2.87 (*m*, 2H), 3.93 (*t*, 2H, *J* = 12.0), 3.38 (*dt*, 1H, <sup>3</sup>*J* = 4.8, <sup>2</sup>*J* = 12.2), 7.1–7.8 (*m*, 8H), 9.21 (*d*, 1H, *J* = 4.5).

### Crystal data

C<sub>21</sub>H<sub>20</sub>Cl<sub>2</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 375.27  
 Orthorhombic, *P*2<sub>1</sub>2<sub>1</sub>2<sub>1</sub>  
*a* = 7.635 (2) Å  
*b* = 12.1814 (15) Å  
*c* = 20.175 (4) Å  
*V* = 1876.4 (7) Å<sup>3</sup>  
*Z* = 4  
*D<sub>x</sub>* = 1.328 Mg m<sup>−3</sup>

Mo *K*α radiation  
 Cell parameters from 25 reflections  
 $\theta = 2-12^\circ$   
 $\mu = 0.36 \text{ mm}^{-1}$   
*T* = 293 (2) K  
 Translucent block, colourless  
 0.2 × 0.2 × 0.1 mm

### Data collection

Enraf–Nonius CAD-4 diffractometer  
 $\omega-2\theta$  scans  
 Absorption correction:  $\psi$  scan (North *et al.*, 1968)  
 $T_{\min} = 0.982$ ,  $T_{\max} = 0.999$   
 1904 measured reflections  
 1904 independent reflections

1399 reflections with  $I > 2\sigma(I)$   
 $\theta_{\max} = 25.0^\circ$   
 $h = 0 \rightarrow 9$   
 $k = 0 \rightarrow 14$   
 $l = 0 \rightarrow 23$   
 3 standard reflections every 100 reflections  
 intensity decay: negligible

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.100$   
 $S = 1.15$   
 1904 reflections  
 226 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0344P)^2 + 0.6059P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.18 \text{ e } \text{Å}^{-3}$$

$$\Delta\rho_{\min} = -0.18 \text{ e } \text{Å}^{-3}$$

**Table 1**

Selected torsion angles (°).

|             |           |             |           |
|-------------|-----------|-------------|-----------|
| C6—C1—C2—C3 | 57.4 (5)  | C3—C4—C5—C6 | −61.8 (4) |
| C1—C2—C3—C4 | −56.5 (4) | C2—C1—C6—C5 | −58.4 (5) |
| C2—C3—C4—C5 | 61.5 (4)  | C4—C5—C6—C1 | 57.9 (4)  |

Since the Friedel pairs were not measured during the data collection, we were unable to determine the absolute structure; in any case, the molecule is achiral. All the H atoms were fixed geometrically and allowed to ride on the parent C atoms, with aromatic C—H = 0.93 Å, tertiary C—H = 0.97 Å and methyl C—H = 0.96 Å. The displacement parameters  $U_{\text{iso}}(\text{H})$  were set at  $1.5U_{\text{eq}}(\text{C})$  for the methyl H atoms and at  $1.2U_{\text{eq}}(\text{C})$  for other H atoms.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *MolEN* (Fair, 1990); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ZORTEP* (Zsolnai, 1997); software used to prepare material for publication: *SHELXL97*.

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