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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 = 0.037

wR factor = 0.102

Data-to-parameter ratio = 18.8

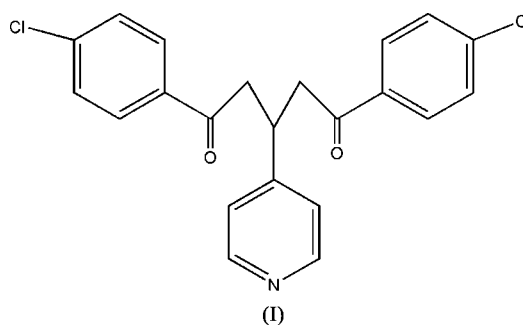
For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1,5-Bis(4-chlorophenyl)-3-(4-pyridyl)-
pentane-1,5-dioneThe title compound, $\text{C}_{22}\text{H}_{17}\text{Cl}_2\text{NO}_2$, has been synthesized and characterized by single-crystal X-ray diffraction. In the crystal structure, the two benzene rings and the pyridyl group lie in a propeller arrangement around the central C atom, thereby minimizing the steric effects between these rings.

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Comment

As part of an investigation of the structures of multi-ring compounds, we report here the structure of the title compound, (I) (Fig. 1).



In the molecule of (I), all bond lengths are within normal ranges (Allen *et al.*, 1987). The two benzene rings and the pyridyl group lie in a propeller arrangement around C9, thereby minimizing the steric effects between these rings. The dihedral angle between the two benzene rings is $15.4(4)^\circ$. The dihedral angles between the pyridine ring and the rings C1–C6 and C12–C17 are $69.3(4)$ and $89.6(4)^\circ$, respectively.

There are no short intermolecular contacts in the crystal structure of (I).

Experimental

The title compound was synthesized by the reaction of equivalent amounts of (*E*)-1-(4-chlorophenyl)-3-(pyridin-4-yl)prop-2-en-1-one (0.1 mmol), isonicotinohydrazide (0.1 mmol) and 1-(4-chlorophenyl)ethanone (0.1 mmol) in ethanol solution (30 ml) for 3 h at 373–383 K. Single crystals suitable for X-ray diffraction analysis were obtained by evaporation of an ethanol solution.

Crystal data

 $\text{C}_{22}\text{H}_{17}\text{Cl}_2\text{NO}_2$ $M_r = 398.27$ Orthorhombic, $P2_12_12_1$ $a = 5.6553(11) \text{ \AA}$ $b = 15.754(3) \text{ \AA}$ $c = 21.191(4) \text{ \AA}$ $V = 1888.0(6) \text{ \AA}^3$

Z = 4

 $D_x = 1.401 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.36 \text{ mm}^{-1}$

T = 298(2) K

Block, yellow

 $0.34 \times 0.13 \times 0.08 \text{ mm}$

Data collection

Bruker SMART APEX area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.941$, $T_{\max} = 0.969$

17030 measured reflections
 4606 independent reflections
 1774 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$
 $\theta_{\text{max}} = 28.2^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.102$
 $S = 0.76$
 4606 reflections
 245 parameters
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0544P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.14 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -0.18 \text{ e } \text{Å}^{-3}$
 Extinction correction: SHELXL97 (Sheldrick, 1997a)
 Extinction coefficient: 0.0066 (10)
 Absolute structure: Flack (1983), with 1918 Friedel pairs
 Flack parameter: 0.01 (7)

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C—H distances in the range 0.93–0.98 Å, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); 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.

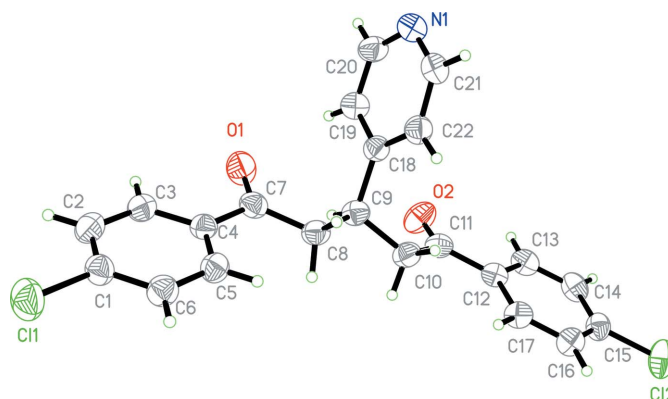


Figure 1
 The structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

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