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Key indicators

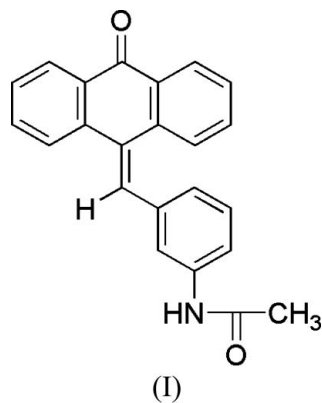
Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.051
 wR factor = 0.150
Data-to-parameter ratio = 9.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

10-(3-Acetamidobenzylidene)anthrone

The title compound, $\text{C}_{23}\text{H}_{17}\text{NO}_2$, was prepared by reacting 10-(3-aminobenzylidene)anthrone and acetic anhydride in the presence of pyridine. X-ray crystal structure analysis shows that the three rings of the anthraquinone system are not coplanar because of steric interactions with the benzylidene group.

Comment

It was reported recently that some 10-substituted benzylideneanthrones possess high antitumour activity (Paull *et al.*, 1992). In our laboratory, some 10-substituted benzylideneanthrones have been prepared and evaluated for antitumour activity. Our study of the structure–activity relationship (SAR) showed that substitution in the phenyl ring of the molecule affects its antitumour activity (Hu & Zhou, 2004). As a continuation of our research work on SARs, we prepared crystals of the title molecule, (I), and investigated its structure.



The molecular structure of (I) is illustrated in Fig. 1. Selected bond lengths and angles are listed in Table 1. As illustrated in Fig. 1, the three rings of the anthraquinone system are not coplanar, because of steric interactions with the benzylidene group; the two outer benzene rings form a dihedral angle of $23.0(1)^\circ$. In the central six-membered ring, atoms C11, C12, C13 and C14 are coplanar to within $0.009(2)$ Å, with atoms C5 and C10 deviating from the plane by $0.125(5)$ and $0.246(5)$ Å, respectively, and the ring adopts an asymmetric boat conformation.

Experimental

10-(3-Aminobenzylidene)anthrone was prepared according to the literature method of Ingram (1950). A mixture of 10-(3-aminobenzylidene)anthrone (0.6 g), acetic anhydride (20 ml) and pyridine

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(0.5 ml) was added to a 100 ml flask and then stirred. The reaction was continued for 1.5 h at room temperature, followed by filtration and washing with ethanol. The crude product was collected and recrystallized from ethanol to afford yellow crystals of (I) (0.4 g, yield 59.0%, m.p. 462–465 K). Spectroscopic analysis: IR (KBr, ν , cm^{-1}): 3319, 1687, 1645, 1602, 1544, 1483, 1417, 1372, 1319, 1277, 932, 776, 683; $^1\text{H NMR}$ (CDCl_3 , δ , p.p.m.): 7.52 (s, 1H, C=CH), 7.00–8.28 (m, 12H), 2.18 (s, 3H, $-\text{CH}_3$). MS (%): 340 ($M+1$, 10), 339 (M , 22), 297 (11), 296 (32), 281 (17), 280 (47), 239 (10), 43 (100).

Crystal data

$\text{C}_{23}\text{H}_{17}\text{NO}_2$	$D_x = 1.264 \text{ Mg m}^{-3}$
$M_r = 339.38$	Mo $K\alpha$ radiation
Monoclinic, $P2_1$	Cell parameters from 25 reflections
$a = 12.3847(3) \text{ \AA}$	$\theta = 3.3\text{--}27.5^\circ$
$b = 7.4853(4) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$c = 12.3351(3) \text{ \AA}$	$T = 296(2) \text{ K}$
$\beta = 128.764(4)^\circ$	Block, yellow
$V = 891.63(8) \text{ \AA}^3$	$0.34 \times 0.23 \times 0.14 \text{ mm}$
$Z = 2$	

Data collection

Enraf–Nonius CAD-4 diffractometer	$R_{\text{int}} = 0.029$
ω - 2θ scans	$\theta_{\text{max}} = 27.5^\circ$
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	$h = -16 \rightarrow 0$
$T_{\text{min}} = 0.963$, $T_{\text{max}} = 0.989$	$k = -9 \rightarrow 9$
3717 measured reflections	$l = -11 \rightarrow 16$
2171 independent reflections	3 standard reflections
1626 reflections with $I > 2\sigma(I)$	frequency: 60 min
	intensity decay: none

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0888P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.151$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.10$	$\Delta\rho_{\text{max}} = 0.34 \text{ e \AA}^{-3}$
2171 reflections	$\Delta\rho_{\text{min}} = -0.33 \text{ e \AA}^{-3}$
241 parameters	Extinction correction: <i>SHELXL97</i>
H atoms treated by a mixture of independent and constrained refinement	(Sheldrick, 1997)
	Extinction coefficient: 0.084 (12)

Table 1

Selected geometric parameters (\AA , $^\circ$) for (I).

O1—C5	1.231 (4)	N1—C18	1.408 (5)
O2—C22	1.219 (4)	C10—C15	1.350 (4)
N1—C22	1.354 (4)		
C15—C10—C14	118.5 (3)	C17—C16—C21	119.0 (3)
C15—C10—C13	125.0 (3)	C17—C16—C15	122.5 (3)
C14—C10—C15—C16	−176.1 (4)	C10—C15—C16—C21	−127.7 (4)
C13—C10—C15—C16	2.9 (6)	C22—N1—C18—C19	−22.1 (6)
C10—C15—C16—C17	55.5 (6)	C22—N1—C18—C17	159.2 (4)

H atoms were positioned geometrically, with C—H distances set to 0.96 \AA for methyl H atoms and 0.93 \AA for the remainder, and refined

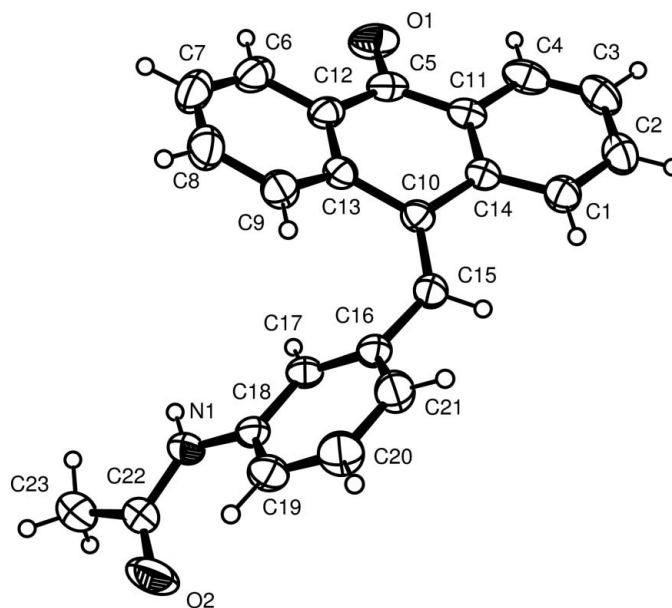


Figure 1

The structure of (I), with 30% probability displacement ellipsoids.

using the riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2$ (or 1.5 for methyl H atoms) times $U_{\text{eq}}(\text{parent atom})$. In the absence of significant anomalous scattering, Friedel pairs were merged before the final refinement.

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD* (McArdle, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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