

Wen-Bing Yuan, Lan Yan and
Ru-Dong Yang*Department of Chemistry, Lanzhou University,
Lanzhou 730000, People's Republic of China

Correspondence e-mail: lzuanwb@163.com

Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.008$ Å
 R factor = 0.071
 wR factor = 0.255
Data-to-parameter ratio = 11.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

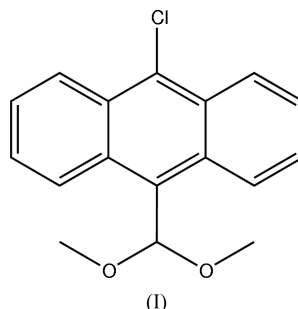
9-Chloro-10-(dimethoxymethyl)anthracene

In the title compound, $\text{C}_{17}\text{H}_{15}\text{ClO}_2$, the anthracene ring system is planar and the bond lengths and angles are unexceptional. In the crystal structure, the molecules stack along the a axis and there are no significantly short intermolecular contacts.

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Comment

In the course of complex syntheses, for instance, in the total syntheses of natural products, carbonyl groups must often be protected against nucleophilic attack, *e.g.* by organometallic compounds, strong bases, reduction and sometimes also oxidation. To achieve this, the carbonyl groups are generally transformed into appropriate acetals, hydrazones, oximes and cyanohydrins. Dialkyl acetals and ketals can easily be formed from carbonyl compounds with alcohols under acidic conditions.



We report here the structure of a new acetal, (I) (Fig. 1), formed by reacting 10-chloro-9-anthraldehyde with methanol in the presence of CuCl_2 . The bond lengths and angles lie within normal ranges (You & Zhu, 2004). The dihedral angles between the anthracene plane and the groups $\text{C}1-\text{O}1-\text{C}16$ and $\text{C}1-\text{O}2-\text{C}17$ are $62.1(4)$ and $83.5(4)^\circ$, respectively; the dihedral angle between planes $\text{C}1-\text{O}1-\text{C}16$ and $\text{C}1-\text{O}2-\text{C}17$ is $79.1(4)^\circ$. In the crystal structure, the molecules stack along the a axis and there are no significantly short intermolecular contacts.

Experimental

10-Chloro-9-anthraldehyde, methanol, ethanol and CuCl_2 are available commercially and were used without further purification. 10-Chloro-9-anthraldehyde (0.5 mmol, 120.3 mg) and CuCl_2 (0.1 mmol, 13.4 mg) were dissolved in MeOH–EtOH (1:1 *v/v*, 5 ml). The mixture was stirred for 2 h at room temperature, filtered, and the filtrate was left to evaporate. After 4 weeks, some crystals suitable for X-ray crystallographic analysis were formed; these were washed three times with the same ratio mixture of solvent (yield 63%). Analysis found: C 71.1, H 5.3%; calculated for $\text{C}_{17}\text{H}_{15}\text{ClO}_2$: C 71.2, H 5.2%.

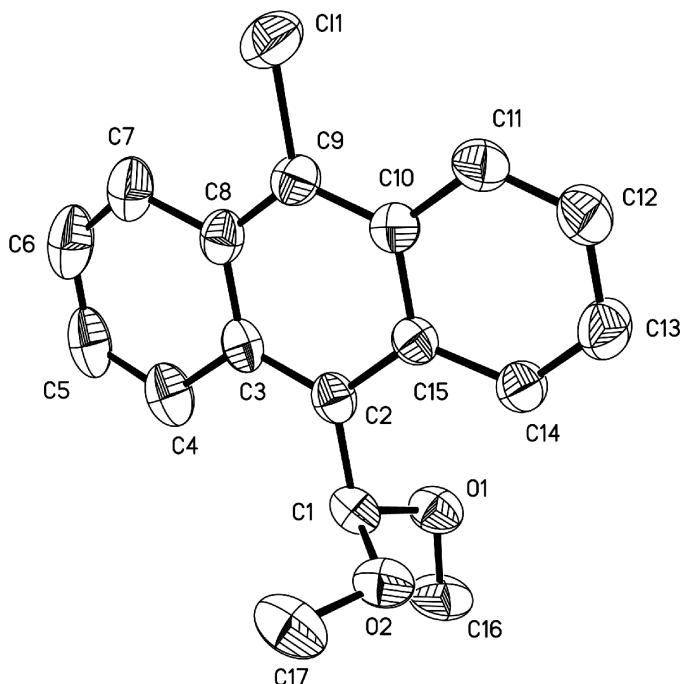


Figure 1

The structure of the title compound, showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted.

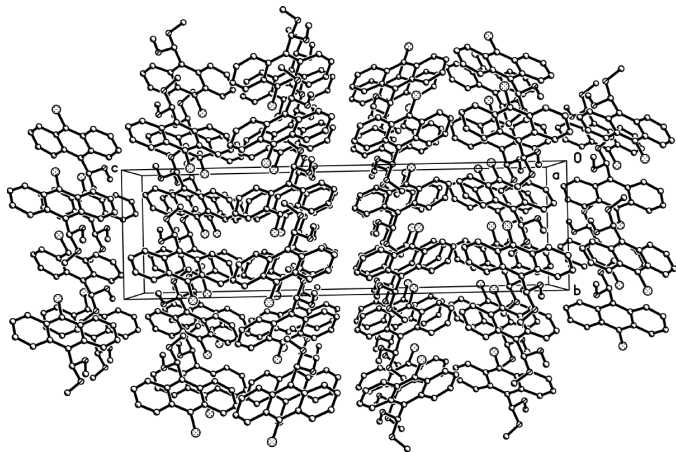


Figure 2

The crystal packing of the title compound, viewed along the *a* axis. H atoms have been omitted.

Crystal data

$C_{17}H_{15}ClO_2$
 $M_r = 286.74$
 Orthorhombic, *Pbca*
 $a = 10.553 (12) \text{ \AA}$
 $b = 8.948 (10) \text{ \AA}$
 $c = 30.84 (3) \text{ \AA}$
 $V = 2912 (6) \text{ \AA}^3$
 $Z = 8$
 $D_x = 1.308 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation
 Cell parameters from 1879 reflections
 $\theta = 2.3\text{--}20.1^\circ$
 $\mu = 0.26 \text{ mm}^{-1}$
 $T = 293 (2) \text{ K}$
 Block, colourless
 $0.42 \times 0.35 \times 0.28 \text{ mm}$

Data collection

Bruker SMART 1000 CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.899$, $T_{\max} = 0.931$
 13634 measured reflections

2550 independent reflections
 1069 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.087$
 $\theta_{\max} = 25.0^\circ$
 $h = -12 \rightarrow 12$
 $k = -10 \rightarrow 10$
 $l = -36 \rightarrow 25$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.071$
 $wR(F^2) = 0.255$
 $S = 0.95$
 2550 reflections
 226 parameters

All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.141P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.78 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.24 \text{ e \AA}^{-3}$

All H atoms were refined isotropically, giving C–H distances in the range $0.93 (2)\text{--}0.99 (2) \text{ \AA}$; $U_{\text{iso}}(\text{H})$ values were fixed at 0.08 \AA^2 .

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); 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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