

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

2-(4-Methoxyphenyl)-1H-indene

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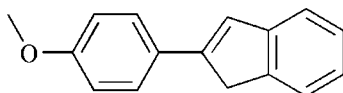
Received 27 April 2008; accepted 28 June 2008

Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.051; wR factor = 0.136; data-to-parameter ratio = 17.6.

Excluding four H atoms, the molecule of the title compound, $\text{C}_{16}\text{H}_{14}\text{O}$, is almost planar, with an r.m.s. deviation of 0.0801 (2) Å. Due to p - π conjugation, the lengths of the two single bonds attached to the O atom are significantly different.

Related literature

For related literature, see: Rayabarapu *et al.* (2003); Senanayake *et al.* (1995).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{14}\text{O}$
 $M_r = 222.27$
 Monoclinic, $P2_1/c$
 $a = 5.8347$ (8) Å
 $b = 7.5584$ (10) Å
 $c = 26.135$ (4) Å
 $\beta = 92.772$ (11)°

$V = 1151.3$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 113$ (2) K
 $0.34 \times 0.32 \times 0.12$ mm

Data collection

Rigaku Saturn diffractometer
 Absorption correction: multi-scan
 (*CrystalClear*; Molecular
 Structure Corporation & Rigaku,
 1999)
 $T_{\min} = 0.974$, $T_{\max} = 0.991$

10940 measured reflections
 2724 independent reflections
 2360 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.135$
 $S = 1.10$
 2724 reflections

155 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.32$ e Å⁻³
 $\Delta\rho_{\min} = -0.39$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

O1—C1	1.3724 (16)	O1—C16	1.4301 (19)
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Data collection: *CrystalClear* (Molecular Structure Corporation & Rigaku, 1999); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

The project was supported by the Fund for Doctorates of Henan University of Science and Technology.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WK2085).

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supplementary materials

Acta Cryst. (2008). E64, o1406 [doi:10.1107/S1600536808019776]

2-(4-Methoxyphenyl)-1*H*-indene

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Comment

Indene ring frameworks are present in a large number of biologically active compounds, and their metallocene complexes are able to catalyze olefin polymerization (Senanayake *et al.*, 1995; Rayabarapu *et al.*, 2003). Some derivatives have shown analgesic and myorelaxation activity, and others are used as valuable intermediates for the synthesis of indenyl chrysanthemates that possess insecticidal properties. So in the recent three decades, many chemists have been attracted by the synthesis of indenenes. In this context, we report the synthesis and crystal structure of the title compound, (I), namely 2-(4-methoxyphenyl)-1*H*-indene.

The title compound was obtained as colourless plate-like crystals in the monoclinic space group $P 1 21/c 1$. A view of the molecular structure of (I) with the numbering scheme is shown in Fig. 1. The whole molecular structure is almost planar with an r.m.s. deviation of 0.0801 (2) Å. Due to the p - π conjugation of atom O1 and benzene ring, the single-bond distance of the O1—C1 [1.3724 (16) Å] is significantly shorter than that of O1—C16 [1.4301 (19) Å].

Experimental

o-Bromobenzyl zinc bromide (3.5 mmol, 3.5 equiv) in 3.5 ml CH₂Cl₂ was added to a degassed refluxing CH₂Cl₂ solution (8 ml) of 1-ethynyl-4-methoxybenzene (1.0 mmol, 1.0 equiv) and Ni(PPh₃)₂I₂ (0.1 mmol, 0.1 equiv). After being stirred at 313 K for 6 h, the solution was cooled to room temperature. The resultant solution was diluted with 50 ml ethyl acetate. The organic layer was washed with 10 ml aqueous HCl solution, saturated NaCl. The aqueous layer was back-extracted with Ethyl acetate. The combined organic layer was dried over anhydrous Na₂SO₄. After filtration, the solvent was removed under reduced pressure and the residue was purified *via* flash chromatography (SiO₂) to afford the compound. Single crystal suitable for X-ray analysis were obtained by slow evaporation at 298 K of a CH₂Cl₂ solution.

Refinement

H atoms were positioned geometrically and refined as riding with C—H = 0.95–0.99 Å. For the CH and CH₂ groups, $U_{\text{iso}}(\text{H})$ values are set equal to 1.2 U_{eq} (carrier atom) and for the methyl groups they are set equal to 1.5 U_{eq} (carrier atom).

Figures

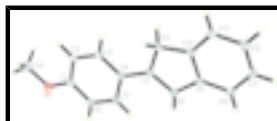


Fig. 1. View of the molecule of (I) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 35% probability level.

2-(4-Methoxyphenyl)-1H-indene

Crystal data

$C_{16}H_{14}O$	$F_{000} = 472$
$M_r = 222.27$	$D_x = 1.282 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
Hall symbol: -P 2ybc	$\lambda = 0.71070 \text{ \AA}$
$a = 5.8347 (8) \text{ \AA}$	Cell parameters from 2782 reflections
$b = 7.5584 (10) \text{ \AA}$	$\theta = 2.8\text{--}27.9^\circ$
$c = 26.135 (4) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 92.772 (11)^\circ$	$T = 113 (2) \text{ K}$
$V = 1151.3 (3) \text{ \AA}^3$	Plate, colourless
$Z = 4$	$0.34 \times 0.32 \times 0.12 \text{ mm}$

Data collection

Rigaku Saturn diffractometer	2724 independent reflections
Radiation source: rotating anode	2360 reflections with $I > 2\sigma(I)$
Monochromator: confocal	$R_{\text{int}} = 0.038$
Detector resolution: $7.31 \text{ pixels mm}^{-1}$	$\theta_{\text{max}} = 27.9^\circ$
$T = 113(2) \text{ K}$	$\theta_{\text{min}} = 2.8^\circ$
ω scans	$h = -7 \rightarrow 7$
Absorption correction: multi-scan (CrystalClear; Molecular Structure Corporation & Rigaku, 1999)	$k = -9 \rightarrow 9$
$T_{\text{min}} = 0.974$, $T_{\text{max}} = 0.991$	$l = -34 \rightarrow 34$
10940 measured reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.051$	H-atom parameters constrained
$wR(F^2) = 0.135$	$w = 1/[\sigma^2(F_o^2) + (0.0654P)^2 + 0.4321P]$
$S = 1.10$	where $P = (F_o^2 + 2F_c^2)/3$
2724 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
155 parameters	$\Delta\rho_{\text{max}} = 0.32 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\text{min}} = -0.39 \text{ e \AA}^{-3}$
	Extinction correction: none

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.06468 (18)	0.38400 (14)	0.43568 (4)	0.0246 (3)
C1	0.0999 (2)	0.37824 (18)	0.38417 (5)	0.0196 (3)
C2	-0.0458 (2)	0.29441 (18)	0.34780 (5)	0.0209 (3)
H2	-0.1824	0.2384	0.3578	0.025*
C3	0.0118 (2)	0.29399 (18)	0.29665 (5)	0.0205 (3)
H3	-0.0875	0.2366	0.2720	0.025*
C4	0.2110 (2)	0.37525 (17)	0.28040 (5)	0.0188 (3)
C5	0.3523 (2)	0.46050 (18)	0.31800 (5)	0.0207 (3)
H5	0.4884	0.5177	0.3082	0.025*
C6	0.2972 (2)	0.46277 (18)	0.36889 (5)	0.0217 (3)
H6	0.3944	0.5223	0.3935	0.026*
C7	0.2705 (2)	0.37168 (17)	0.22661 (5)	0.0191 (3)
C8	0.4701 (2)	0.44705 (18)	0.20655 (5)	0.0203 (3)
H8	0.5885	0.5082	0.2255	0.024*
C9	0.4594 (2)	0.41280 (18)	0.15079 (5)	0.0193 (3)
C10	0.6102 (3)	0.45594 (19)	0.11311 (5)	0.0234 (3)
H10	0.7468	0.5203	0.1215	0.028*
C11	0.5577 (3)	0.40313 (19)	0.06279 (6)	0.0251 (3)
H11	0.6602	0.4309	0.0368	0.030*
C12	0.3566 (3)	0.31006 (19)	0.05014 (5)	0.0246 (3)
H12	0.3239	0.2747	0.0157	0.030*
C13	0.2030 (3)	0.26833 (18)	0.08766 (5)	0.0220 (3)
H13	0.0648	0.2064	0.0790	0.026*
C14	0.2559 (2)	0.31909 (17)	0.13805 (5)	0.0188 (3)
C15	0.1310 (2)	0.29032 (18)	0.18543 (5)	0.0197 (3)
H15A	-0.0226	0.3461	0.1824	0.024*
H15B	0.1118	0.1623	0.1919	0.024*
C16	-0.1393 (3)	0.3029 (2)	0.45282 (6)	0.0311 (4)
H16A	-0.1495	0.3227	0.4897	0.047*
H16B	-0.1347	0.1755	0.4460	0.047*
H16C	-0.2736	0.3548	0.4345	0.047*

supplementary materials

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0271 (6)	0.0282 (6)	0.0184 (5)	-0.0015 (4)	0.0008 (4)	0.0018 (4)
C1	0.0221 (7)	0.0180 (6)	0.0187 (6)	0.0040 (5)	-0.0003 (5)	0.0017 (5)
C2	0.0195 (7)	0.0196 (7)	0.0237 (7)	-0.0020 (5)	0.0006 (5)	0.0014 (5)
C3	0.0205 (7)	0.0188 (6)	0.0219 (7)	-0.0012 (5)	-0.0019 (5)	-0.0009 (5)
C4	0.0197 (7)	0.0141 (6)	0.0224 (7)	0.0037 (5)	0.0004 (5)	0.0003 (5)
C5	0.0179 (7)	0.0195 (7)	0.0247 (7)	-0.0006 (5)	0.0011 (5)	-0.0011 (5)
C6	0.0205 (7)	0.0204 (7)	0.0240 (7)	0.0004 (5)	-0.0025 (5)	-0.0020 (5)
C7	0.0216 (7)	0.0139 (6)	0.0220 (7)	0.0030 (5)	0.0012 (5)	-0.0006 (5)
C8	0.0181 (7)	0.0212 (7)	0.0215 (7)	-0.0014 (5)	-0.0001 (5)	-0.0021 (5)
C9	0.0193 (7)	0.0158 (6)	0.0225 (7)	0.0011 (5)	-0.0001 (5)	-0.0001 (5)
C10	0.0229 (7)	0.0209 (7)	0.0267 (7)	-0.0024 (6)	0.0026 (6)	0.0015 (5)
C11	0.0280 (8)	0.0241 (7)	0.0235 (7)	0.0019 (6)	0.0060 (6)	0.0053 (6)
C12	0.0311 (8)	0.0230 (7)	0.0198 (7)	0.0021 (6)	0.0006 (6)	0.0011 (5)
C13	0.0242 (8)	0.0200 (7)	0.0216 (7)	-0.0013 (6)	-0.0018 (5)	0.0005 (5)
C14	0.0191 (7)	0.0157 (6)	0.0216 (7)	0.0016 (5)	0.0002 (5)	0.0011 (5)
C15	0.0192 (7)	0.0185 (6)	0.0211 (6)	0.0008 (5)	-0.0008 (5)	-0.0004 (5)
C16	0.0332 (9)	0.0371 (9)	0.0234 (7)	-0.0048 (7)	0.0043 (6)	0.0049 (6)

Geometric parameters (\AA , $^\circ$)

O1—C1	1.3724 (16)	C9—C10	1.391 (2)
O1—C16	1.4301 (19)	C9—C14	1.409 (2)
C1—C6	1.392 (2)	C10—C11	1.394 (2)
C1—C2	1.3963 (19)	C10—H10	0.9500
C2—C3	1.394 (2)	C11—C12	1.394 (2)
C2—H2	0.9500	C11—H11	0.9500
C3—C4	1.399 (2)	C12—C13	1.397 (2)
C3—H3	0.9500	C12—H12	0.9500
C4—C5	1.4078 (19)	C13—C14	1.3919 (19)
C4—C7	1.4645 (19)	C13—H13	0.9500
C5—C6	1.384 (2)	C14—C15	1.4829 (19)
C5—H5	0.9500	C15—H15A	0.9900
C6—H6	0.9500	C15—H15B	0.9900
C7—C8	1.419 (2)	C16—H16A	0.9800
C7—C15	1.4542 (19)	C16—H16B	0.9800
C8—C9	1.4785 (19)	C16—H16C	0.9800
C8—H8	0.9500		
C1—O1—C16	117.40 (12)	C9—C10—C11	118.88 (14)
O1—C1—C6	115.60 (12)	C9—C10—H10	120.6
O1—C1—C2	124.55 (13)	C11—C10—H10	120.6
C6—C1—C2	119.85 (13)	C12—C11—C10	120.83 (14)
C3—C2—C1	119.13 (13)	C12—C11—H11	119.6
C3—C2—H2	120.4	C10—C11—H11	119.6
C1—C2—H2	120.4	C11—C12—C13	120.59 (13)

C2—C3—C4	122.18 (13)	C11—C12—H12	119.7
C2—C3—H3	118.9	C13—C12—H12	119.7
C4—C3—H3	118.9	C14—C13—C12	118.78 (14)
C3—C4—C5	117.14 (13)	C14—C13—H13	120.6
C3—C4—C7	121.50 (12)	C12—C13—H13	120.6
C5—C4—C7	121.36 (13)	C13—C14—C9	120.54 (13)
C6—C5—C4	121.43 (13)	C13—C14—C15	130.85 (13)
C6—C5—H5	119.3	C9—C14—C15	108.60 (12)
C4—C5—H5	119.3	C7—C15—C14	106.02 (12)
C5—C6—C1	120.26 (13)	C7—C15—H15A	110.5
C5—C6—H6	119.9	C14—C15—H15A	110.5
C1—C6—H6	119.9	C7—C15—H15B	110.5
C8—C7—C15	109.64 (12)	C14—C15—H15B	110.5
C8—C7—C4	125.75 (12)	H15A—C15—H15B	108.7
C15—C7—C4	124.61 (13)	O1—C16—H16A	109.5
C7—C8—C9	107.34 (12)	O1—C16—H16B	109.5
C7—C8—H8	126.3	H16A—C16—H16B	109.5
C9—C8—H8	126.3	O1—C16—H16C	109.5
C10—C9—C14	120.36 (13)	H16A—C16—H16C	109.5
C10—C9—C8	131.24 (13)	H16B—C16—H16C	109.5
C14—C9—C8	108.40 (12)		
C16—O1—C1—C6	178.34 (12)	C7—C8—C9—C10	-179.34 (14)
C16—O1—C1—C2	-2.0 (2)	C7—C8—C9—C14	0.24 (15)
O1—C1—C2—C3	-178.41 (13)	C14—C9—C10—C11	-0.7 (2)
C6—C1—C2—C3	1.3 (2)	C8—C9—C10—C11	178.81 (14)
C1—C2—C3—C4	-0.1 (2)	C9—C10—C11—C12	0.6 (2)
C2—C3—C4—C5	-0.7 (2)	C10—C11—C12—C13	0.3 (2)
C2—C3—C4—C7	179.21 (13)	C11—C12—C13—C14	-0.9 (2)
C3—C4—C5—C6	0.4 (2)	C12—C13—C14—C9	0.8 (2)
C7—C4—C5—C6	-179.47 (13)	C12—C13—C14—C15	-178.47 (14)
C4—C5—C6—C1	0.7 (2)	C10—C9—C14—C13	0.0 (2)
O1—C1—C6—C5	178.17 (12)	C8—C9—C14—C13	-179.60 (12)
C2—C1—C6—C5	-1.5 (2)	C10—C9—C14—C15	179.45 (12)
C3—C4—C7—C8	-178.77 (13)	C8—C9—C14—C15	-0.18 (15)
C5—C4—C7—C8	1.1 (2)	C8—C7—C15—C14	0.09 (15)
C3—C4—C7—C15	1.7 (2)	C4—C7—C15—C14	179.70 (12)
C5—C4—C7—C15	-178.42 (13)	C13—C14—C15—C7	179.39 (14)
C15—C7—C8—C9	-0.20 (15)	C9—C14—C15—C7	0.06 (15)
C4—C7—C8—C9	-179.80 (12)		

Fig. 1

