

## 2-[(*E*)-4-Quinolylmethyleneamino]-phenol

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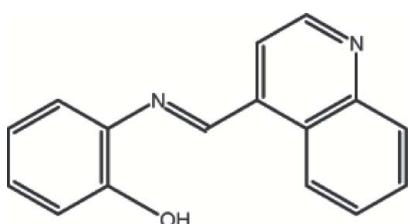
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Key indicators: single-crystal X-ray study;  $T = 150\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$ ;  $R$  factor = 0.042;  $wR$  factor = 0.105; data-to-parameter ratio = 13.6.

In the title compound,  $\text{C}_{16}\text{H}_{12}\text{N}_2\text{O}$ , the dihedral angle between the two aromatic ring systems is  $68.54(5)^\circ$ . The molecular packing is stabilized by intra- and intermolecular  $\text{O}-\text{H}\cdots\text{N}$  and intramolecular  $\text{C}-\text{H}\cdots\text{N}$  hydrogen-bond interactions.

## Related literature

For bond-length data, see: Allen *et al.* (1987). For general background, see: Gao *et al.* (2005); Hagen *et al.* (1983); Lozytska *et al.* (2004); Sessler *et al.* (2004); Kuz'min *et al.* (2000). For related structures, see: Räisänen, Elo *et al.* (2007); Räisänen Leskelä & Repo (2007). For experimental procedures, see: Gümüş (2002).



## Experimental

### Crystal data

$\text{C}_{16}\text{H}_{12}\text{N}_2\text{O}$   
 $M_r = 248.28$   
Monoclinic,  $P2_1/n$   
 $a = 6.7174(6)\text{ \AA}$   
 $b = 23.931(2)\text{ \AA}$   
 $c = 7.7495(6)\text{ \AA}$   
 $\beta = 105.521(1)^\circ$   
 $V = 1200.33(17)\text{ \AA}^3$   
 $Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 0.09\text{ mm}^{-1}$   
 $T = 150(2)\text{ K}$   
 $0.25 \times 0.20 \times 0.10\text{ mm}$

### Data collection

Bruker APEXII CCD diffractometer  
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003)  
 $T_{\min} = 0.978$ ,  $T_{\max} = 0.991$   
12294 measured reflections  
2991 independent reflections  
2338 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.037$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.105$   
 $S = 1.05$   
2991 reflections  
220 parameters  
H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.25\text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.24\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1 $\cdots$ N1	0.95 (2)	2.34 (2)	2.7766 (16)	107.5 (15)
O1—H1 $\cdots$ N2 <sup>i</sup>	0.95 (2)	1.91 (2)	2.8085 (16)	156.4 (19)
C13—H13 $\cdots$ N1	0.975 (15)	2.356 (14)	2.9776 (17)	121.0 (11)

Symmetry code: (i)  $-x + 2, -y, -z + 2$ .

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2807).

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## **supplementary materials**

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## 2-[*(E*)-4-Quinolylmethylideneamino]phenol

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### Comment

2- and 4-(*N*-arylimino)quinolines have been investigated to their clinical efficacy for use as an analgetic drug (Hagen *et al.*, 1983). A number of azomethines and their complexes possess high biological activity, including antineoplastic (Kuz'min *et al.*, 2000; Gao *et al.*, 2005) and antiviral (Lozytska *et al.*, 2004). It has been recently shown that the macrocyclic Schiff bases can selectively bind different anions acting as artificial anionic receptors (Sessler *et al.*, 2004).

In the title compound, (I), (Fig. 1), all bond lengths and angles are within normal ranges (Räisänen, Elo *et al.*, 2007; Räisänen, Leskelä, & Repo 2007; Allen *et al.*, 1987). The molecule has a non-planar conformation. The dihedral angle between its two aromatic ring systems (C1–C6) and (N2/C8–C16) is 68.54 (5)°. The C6—N1—C7—C8 torsion angle is -179.98 (13)° and the N1—C6—C1—O1 angle deviates from zero only by -7.27 (18)°.

The molecular packing of the compound is stabilized by O—H···N and C—H···N hydrogen bonds (Table 1, Fig. 2).

### Experimental

Quinoline-4-carboxaldehyde (1.0 mmol) was dissolved in hot absolute ethanol (5 ml) and an equimolar amount of 2-aminophenol dissolved in hot absolute ethanol (10 ml) was added. The mixture was refluxed at reaction temperature for 2 h. The progress of the reaction was monitored by TLC using n-hexane/ethylacetate (1:1 v/v) as eluent. Upon completion of the reaction, the crude product which precipitated on cooling was collected by filtration. Further purification was accomplished by recrystallization from ethanol to yield yellow blocks of (I). [yield 80%, m.p. 475–476 K]. UV (CHCl<sub>3</sub>):  $\lambda_{\text{max}}$  248, 343, 377 nm. IR (KBr): 3029, 2851, 1625, 1574 and 1472, 1293–1165, 830 and 757 cm<sup>-1</sup>. <sup>1</sup>H NMR (CDCl<sub>3</sub>, 200 MHz): δ (p.p.m.) 6.95–9.08 (m, ArH and CH, 11H), 9.41 (s, OH, 1H). MS: m/z 249 (*M*+1), 248 (*M*), 247 (*M*-1), 128 (*M*-120), 120 (*M*-128). Analysis calculated for C<sub>16</sub>H<sub>12</sub>N<sub>2</sub>O: C 77.40, H 4.87, N 11.28%; found: C 77.62, H 4.96, N 11.12% (Gümüş, 2002).

### Refinement

The H atoms were found from a difference map and refined freely.

### Figures

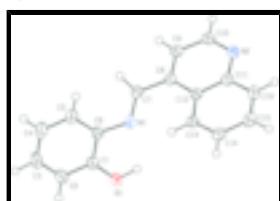


Fig. 1. View of (I): displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

# supplementary materials

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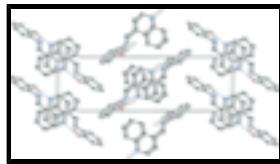


Fig. 2. The hydrogen bonding interactions of (I) viewed down the  $a$  axis. For clarity, H atoms not involved in hydrogen bonding have been omitted.

## 2-[**(E**)-4-Quinolylmethylenamino]phenol

### Crystal data

C <sub>16</sub> H <sub>12</sub> N <sub>2</sub> O	$F_{000} = 520$
$M_r = 248.28$	$D_x = 1.374 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
Hall symbol: -P 2yn	$\lambda = 0.71073 \text{ \AA}$
$a = 6.7174 (6) \text{ \AA}$	Cell parameters from 2762 reflections
$b = 23.931 (2) \text{ \AA}$	$\theta = 2.9\text{--}26.4^\circ$
$c = 7.7495 (6) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 105.521 (1)^\circ$	$T = 150 (2) \text{ K}$
$V = 1200.33 (17) \text{ \AA}^3$	Block, yellow
$Z = 4$	$0.25 \times 0.20 \times 0.10 \text{ mm}$

### Data collection

Bruker APEXII CCD diffractometer	2991 independent reflections
Radiation source: sealed tube	2338 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.037$
$T = 150(2) \text{ K}$	$\theta_{\text{max}} = 28.4^\circ$
$\varphi$ and $\omega$ scans	$\theta_{\text{min}} = 1.7^\circ$
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	$h = -8 \rightarrow 8$
$T_{\text{min}} = 0.978$ , $T_{\text{max}} = 0.991$	$k = -31 \rightarrow 31$
12294 measured reflections	$l = -10 \rightarrow 10$

### Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: difmap and geom
$R[F^2 > 2\sigma(F^2)] = 0.042$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.105$	$w = 1/\sigma^2(F_o^2) + (0.0424P)^2 + 0.3303P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2991 reflections	$\Delta\rho_{\text{max}} = 0.25 \text{ e \AA}^{-3}$
220 parameters	$\Delta\rho_{\text{min}} = -0.24 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction correction: none

*Special details*

**Geometry.** Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement on  $F^2$  for ALL reflections except those flagged by the user for potential systematic errors. Weighted  $R$ -factors  $wR$  and all goodnesses 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 observed criterion of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factor-obs *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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.06305 (16)	0.13183 (4)	0.54890 (14)	0.0340 (3)
N1	0.76446 (16)	0.10349 (4)	0.72156 (14)	0.0243 (3)
N2	0.80975 (16)	-0.03956 (5)	1.22446 (15)	0.0258 (3)
C1	0.8930 (2)	0.16414 (5)	0.52971 (17)	0.0264 (4)
C2	0.8697 (3)	0.20982 (6)	0.41486 (19)	0.0340 (4)
C3	0.6997 (3)	0.24448 (6)	0.3910 (2)	0.0372 (5)
C4	0.5529 (3)	0.23493 (6)	0.4839 (2)	0.0370 (5)
C5	0.5745 (2)	0.18955 (6)	0.59862 (19)	0.0313 (4)
C6	0.7433 (2)	0.15362 (5)	0.62205 (17)	0.0242 (4)
C7	0.72483 (19)	0.10243 (5)	0.87303 (17)	0.0235 (3)
C8	0.74355 (18)	0.05168 (5)	0.98390 (16)	0.0215 (3)
C9	0.78135 (19)	0.05904 (6)	1.16557 (17)	0.0236 (3)
C10	0.81518 (19)	0.01260 (6)	1.28017 (17)	0.0260 (4)
C11	0.76319 (18)	-0.04841 (5)	1.04285 (17)	0.0233 (3)
C12	0.72937 (18)	-0.00427 (5)	0.91566 (16)	0.0212 (3)
C13	0.67854 (19)	-0.01837 (6)	0.73121 (17)	0.0240 (4)
C14	0.6654 (2)	-0.07303 (6)	0.67751 (18)	0.0278 (4)
C15	0.7014 (2)	-0.11657 (6)	0.8042 (2)	0.0308 (4)
C16	0.7487 (2)	-0.10454 (5)	0.98290 (19)	0.0282 (4)
H1	1.068 (3)	0.1013 (9)	0.628 (3)	0.072 (6)*
H2	0.976 (3)	0.2161 (7)	0.350 (2)	0.044 (5)*
H3	0.686 (3)	0.2753 (7)	0.307 (2)	0.046 (5)*
H4	0.433 (3)	0.2587 (7)	0.469 (2)	0.042 (5)*
H5	0.468 (2)	0.1808 (6)	0.657 (2)	0.033 (4)*
H7	0.689 (2)	0.1376 (6)	0.929 (2)	0.031 (4)*
H9	0.788 (2)	0.0966 (6)	1.214 (2)	0.031 (4)*
H10	0.844 (2)	0.0186 (6)	1.409 (2)	0.024 (4)*
H13	0.653 (2)	0.0114 (6)	0.642 (2)	0.025 (4)*
H14	0.632 (2)	-0.0822 (6)	0.550 (2)	0.032 (4)*
H15	0.693 (2)	-0.1566 (7)	0.765 (2)	0.035 (4)*
H16	0.772 (2)	-0.1340 (7)	1.074 (2)	0.033 (4)*

## supplementary materials

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### Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0385 (6)	0.0322 (5)	0.0359 (6)	0.0058 (4)	0.0181 (4)	0.0097 (4)
N1	0.0246 (5)	0.0223 (5)	0.0254 (6)	0.0006 (4)	0.0055 (4)	0.0023 (4)
N2	0.0220 (5)	0.0295 (6)	0.0259 (6)	0.0025 (4)	0.0062 (4)	0.0040 (5)
C1	0.0343 (7)	0.0221 (6)	0.0214 (6)	-0.0012 (5)	0.0049 (5)	-0.0020 (5)
C2	0.0475 (9)	0.0274 (7)	0.0272 (7)	-0.0040 (6)	0.0100 (6)	0.0022 (6)
C3	0.0549 (10)	0.0214 (7)	0.0308 (8)	-0.0020 (6)	0.0037 (7)	0.0037 (6)
C4	0.0436 (9)	0.0238 (7)	0.0383 (8)	0.0074 (6)	0.0018 (7)	0.0010 (6)
C5	0.0335 (7)	0.0257 (7)	0.0328 (7)	0.0021 (6)	0.0057 (6)	0.0002 (6)
C6	0.0303 (7)	0.0185 (6)	0.0217 (6)	-0.0016 (5)	0.0036 (5)	-0.0009 (5)
C7	0.0226 (6)	0.0221 (6)	0.0249 (6)	0.0003 (5)	0.0050 (5)	-0.0017 (5)
C8	0.0168 (6)	0.0238 (6)	0.0238 (6)	-0.0003 (5)	0.0052 (4)	0.0004 (5)
C9	0.0211 (6)	0.0251 (6)	0.0248 (6)	-0.0007 (5)	0.0064 (5)	-0.0023 (5)
C10	0.0227 (6)	0.0345 (7)	0.0210 (6)	0.0009 (5)	0.0060 (5)	0.0005 (5)
C11	0.0173 (6)	0.0259 (6)	0.0269 (6)	0.0004 (5)	0.0062 (5)	0.0017 (5)
C12	0.0166 (6)	0.0234 (6)	0.0241 (6)	0.0000 (4)	0.0064 (4)	-0.0004 (5)
C13	0.0212 (6)	0.0266 (7)	0.0247 (6)	0.0006 (5)	0.0070 (5)	0.0001 (5)
C14	0.0249 (6)	0.0297 (7)	0.0285 (7)	-0.0002 (5)	0.0068 (5)	-0.0056 (5)
C15	0.0282 (7)	0.0240 (7)	0.0388 (8)	0.0006 (5)	0.0064 (6)	-0.0036 (6)
C16	0.0257 (7)	0.0224 (6)	0.0358 (7)	0.0012 (5)	0.0071 (5)	0.0036 (6)

### Geometric parameters ( $\text{\AA}$ , $^\circ$ )

O1—C1	1.3542 (17)	C11—C12	1.4210 (17)
O1—H1	0.95 (2)	C12—C13	1.4186 (18)
N1—C6	1.4126 (16)	C13—C14	1.368 (2)
N1—C7	1.2715 (17)	C14—C15	1.408 (2)
N2—C11	1.3740 (17)	C15—C16	1.366 (2)
N2—C10	1.3181 (19)	C2—H2	0.989 (19)
C1—C2	1.3918 (19)	C3—H3	0.972 (16)
C1—C6	1.4042 (19)	C4—H4	0.967 (19)
C2—C3	1.383 (3)	C5—H5	0.966 (14)
C3—C4	1.387 (3)	C7—H7	1.005 (15)
C4—C5	1.386 (2)	C9—H9	0.971 (15)
C5—C6	1.3957 (19)	C10—H10	0.975 (15)
C7—C8	1.4735 (17)	C13—H13	0.975 (15)
C8—C9	1.3734 (18)	C14—H14	0.978 (15)
C8—C12	1.4334 (17)	C15—H15	1.002 (17)
C9—C10	1.403 (2)	C16—H16	0.980 (16)
C11—C16	1.4161 (17)		
C1—O1—H1	113.2 (13)	C13—C14—C15	120.70 (13)
C6—N1—C7	120.37 (11)	C14—C15—C16	120.07 (13)
C10—N2—C11	117.40 (11)	C11—C16—C15	120.62 (12)
O1—C1—C2	117.77 (13)	C1—C2—H2	118.1 (10)
C2—C1—C6	119.56 (13)	C3—C2—H2	121.6 (10)

O1—C1—C6	122.67 (11)	C2—C3—H3	118.1 (12)
C1—C2—C3	120.27 (16)	C4—C3—H3	121.3 (12)
C2—C3—C4	120.54 (14)	C3—C4—H4	121.7 (10)
C3—C4—C5	119.68 (16)	C5—C4—H4	118.6 (10)
C4—C5—C6	120.53 (14)	C4—C5—H5	120.4 (9)
N1—C6—C1	116.70 (11)	C6—C5—H5	118.9 (9)
C1—C6—C5	119.40 (12)	N1—C7—H7	121.0 (8)
N1—C6—C5	123.63 (12)	C8—C7—H7	115.6 (8)
N1—C7—C8	123.26 (11)	C8—C9—H9	119.5 (9)
C7—C8—C12	124.61 (11)	C10—C9—H9	120.4 (9)
C9—C8—C12	118.28 (11)	N2—C10—H10	117.1 (9)
C7—C8—C9	117.10 (11)	C9—C10—H10	119.1 (9)
C8—C9—C10	120.11 (13)	C12—C13—H13	119.3 (9)
N2—C10—C9	123.87 (12)	C14—C13—H13	119.9 (9)
N2—C11—C16	117.33 (11)	C13—C14—H14	120.0 (9)
C12—C11—C16	119.56 (12)	C15—C14—H14	119.3 (9)
N2—C11—C12	123.11 (11)	C14—C15—H15	120.7 (9)
C8—C12—C13	124.63 (11)	C16—C15—H15	119.2 (9)
C11—C12—C13	118.22 (11)	C11—C16—H16	117.5 (9)
C8—C12—C11	117.13 (11)	C15—C16—H16	121.8 (9)
C12—C13—C14	120.82 (12)		
C7—N1—C6—C1	141.39 (13)	C7—C8—C9—C10	-175.57 (12)
C7—N1—C6—C5	-44.64 (19)	C12—C8—C9—C10	3.15 (19)
C6—N1—C7—C8	-179.98 (13)	C7—C8—C12—C11	176.39 (12)
C11—N2—C10—C9	-1.61 (19)	C7—C8—C12—C13	-5.5 (2)
C10—N2—C11—C12	2.52 (19)	C9—C8—C12—C11	-2.23 (18)
C10—N2—C11—C16	-177.36 (12)	C9—C8—C12—C13	175.91 (13)
O1—C1—C2—C3	-179.58 (13)	C8—C9—C10—N2	-1.3 (2)
C6—C1—C2—C3	-0.4 (2)	N2—C11—C12—C8	-0.62 (19)
O1—C1—C6—N1	-7.27 (18)	N2—C11—C12—C13	-178.87 (12)
O1—C1—C6—C5	178.49 (12)	C16—C11—C12—C8	179.26 (12)
C2—C1—C6—N1	173.54 (12)	C16—C11—C12—C13	1.00 (18)
C2—C1—C6—C5	-0.70 (19)	N2—C11—C16—C15	179.50 (13)
C1—C2—C3—C4	1.4 (2)	C12—C11—C16—C15	-0.4 (2)
C2—C3—C4—C5	-1.4 (2)	C8—C12—C13—C14	-179.03 (13)
C3—C4—C5—C6	0.4 (2)	C11—C12—C13—C14	-0.92 (19)
C4—C5—C6—N1	-173.12 (13)	C12—C13—C14—C15	0.2 (2)
C4—C5—C6—C1	0.7 (2)	C13—C14—C15—C16	0.4 (2)
N1—C7—C8—C9	153.65 (13)	C14—C15—C16—C11	-0.3 (2)
N1—C7—C8—C12	-25.0 (2)		

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O1—H1···N1	0.95 (2)	2.34 (2)	2.7766 (16)	107.5 (15)
O1—H1···N2 <sup>i</sup>	0.95 (2)	1.91 (2)	2.8085 (16)	156.4 (19)
C13—H13···N1	0.975 (15)	2.356 (14)	2.9776 (17)	121.0 (11)

Symmetry codes: (i)  $-x+2, -y, -z+2$ .

## supplementary materials

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Fig. 1

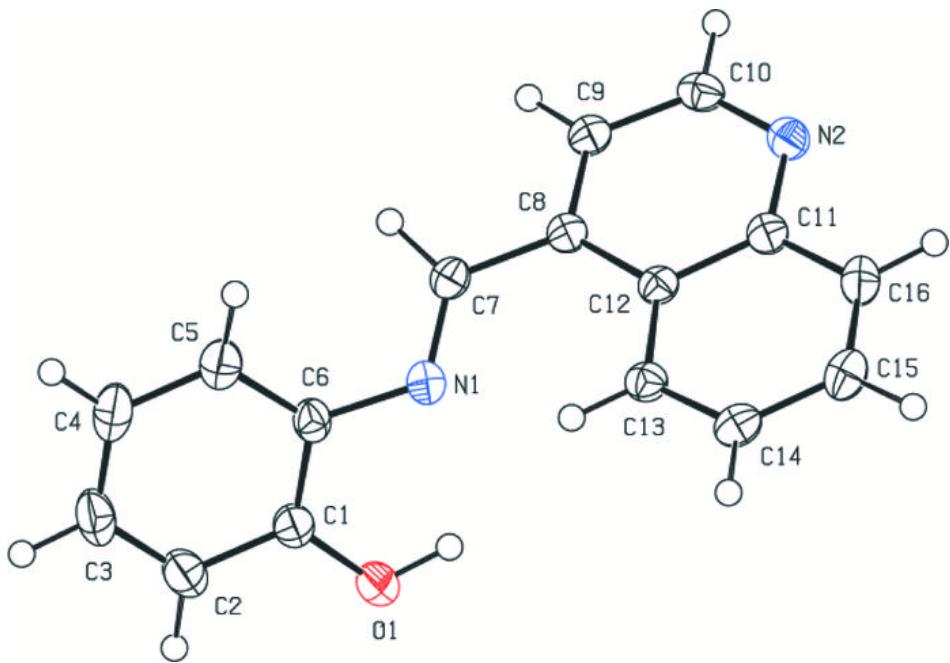


Fig. 2

