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## 2-[(4-Methoxyanilino)methyl]phenol

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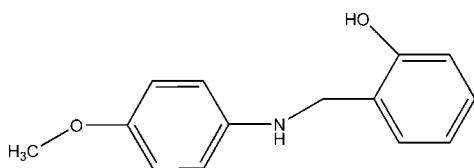
Received 2 August 2011; accepted 5 August 2011

Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.056;  $wR$  factor = 0.128; data-to-parameter ratio = 16.5.

In the title compound,  $\text{C}_{14}\text{H}_{15}\text{NO}_2$ , the dihedral angle between the two benzene rings is  $71.10(5)^\circ$ . In the crystal, molecules are linked by intermolecular  $\text{N}-\text{H}\cdots\text{O}$ , and  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bonds into a chain running parallel to the  $b$  axis.

## Related literature

For the synthesis of the title compound, see: Noda (1959). For other related structures, see: Liu *et al.* (2007); Qu *et al.* (2007).



## Experimental

## Crystal data

 $\text{C}_{14}\text{H}_{15}\text{NO}_2$  $M_r = 229.27$ Monoclinic,  $P2_1/c$  $a = 7.8132(16)$  Å $b = 5.7947(12)$  Å $c = 26.175(5)$  Å $\beta = 95.02(3)^\circ$  $V = 1180.5(4)$  Å<sup>3</sup> $Z = 4$ Mo  $K\alpha$  radiation $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 298$  K $0.40 \times 0.30 \times 0.20$  mm

## Data collection

Rigaku SCXmini diffractometer  
Absorption correction: multi-scan  
(*CrystalClear*; Rigaku, 2005)  
 $T_{\min} = 0.970$ ,  $T_{\max} = 0.983$ 10971 measured reflections  
2693 independent reflections  
1692 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.056$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.056$   
 $wR(F^2) = 0.128$   
 $S = 1.03$   
2693 reflections  
163 parameters  
2 restraintsH atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.17$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.22$  e Å<sup>-3</sup>

**Table 1**  
Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.86 (1)	2.22 (1)	3.058 (2)	163 (2)
$\text{O2}-\text{H2}\cdots\text{N1}^{\text{ii}}$	0.86 (1)	1.89 (1)	2.741 (2)	172 (2)

Symmetry codes: (i)  $x, y - 1, z$ ; (ii)  $-x, -y + 1, -z + 1$ .

Data collection: *CrystalClear* (Rigaku, 2005); 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: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2441).

## References

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Qu, Y., Tian, L.-J. & Dong, J. (2007). *Acta Cryst.* **E63**, o4832.  
Rigaku (2005). *CrystalClear*. Rigaku Corporation, Tokyo, Japan.  
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## supporting information

*Acta Cryst.* (2011). E67, o2304 [doi:10.1107/S1600536811031679]

## 2-[(4-Methoxyanilino)methyl]phenol

Hong Shu, Ning-Shu Yu, Guo-Lan Xie and Li-Zhuang Chen

### S1. Comment

Recently, the crystal structures of compounds closely related to the title molecule, *e.g.*, 2-[(4-chlorophenyl)amino-methyl]-6-methoxyphenol (Liu *et al.*, 2007) and 2-(anilinomethyl)phenol (Qu *et al.*, 2007) have been reported. We report here the crystal structure of a new member of this family of compounds.

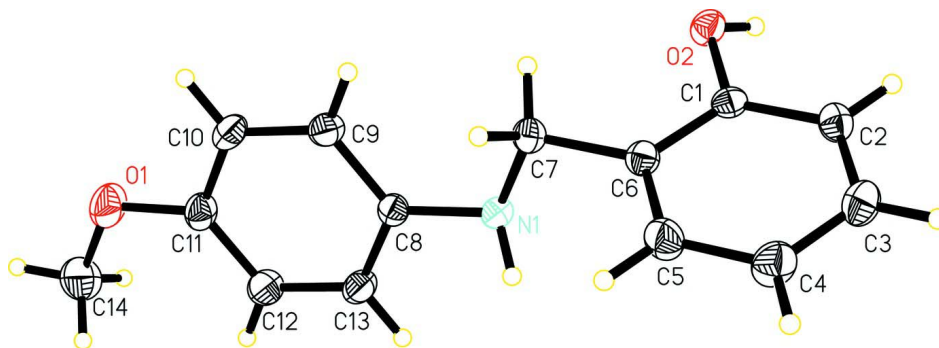
In the title compound (Fig. 1), the dihedral angle between the two benzene ring planes is 71.10 (5)°. In the crystal structure, the molecules are linked by intermolecular N—H···O, and O—H···N hydrogen bonds into a one-dimensional chain lying parallel to the *b*-axis (Fig. 2).

### S2. Experimental

The title compound was synthesized by the reaction of 2-((4-methoxyphenylimino)-methyl)phenol (2.76 g, 10 mmol) with NaBH<sub>4</sub> (0.38 g, 10 mmol) in methanol (50 ml) according to the reported method (Noda, 1959). Crystals were obtained from an ethanolic (95%) solution by slow evaporation at room temperature.

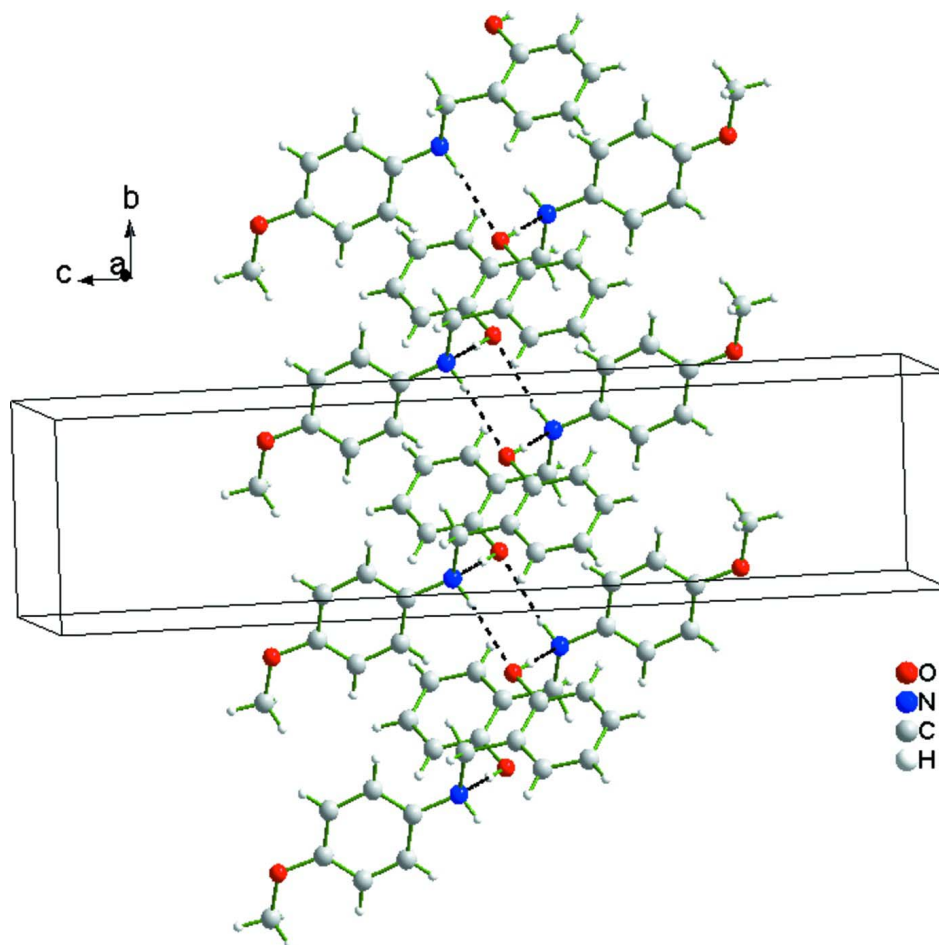
### S3. Refinement

H atoms were placed at calculated positions and were included in the refinement in the riding-model approximation, with C—H = 0.93, 0.96 and 0.97 Å, for aryl, methyl and methylene H-atoms, respectively, with  $U_{\text{iso}}(\text{H}) = 1.2$  (or 1.5 for methyl)  $U_{\text{eq}}(\text{C})$ . The hydrogen atoms bonded to O and N were included in the positions obtained from a difference map and were allowed to refine with distances constrained at N—H and O—H = 0.86 (1) Å.



**Figure 1**

The asymmetric unit of the title compound with atom labels. Displacement ellipsoids are drawn at the 30% probability level.

**Figure 2**

The unit cell packing of the title compound viewed along the *b*-axis. Hydrogen bonds are drawn as dashed lines.

### 2-[(4-Methoxyanilino)methyl]phenol

#### Crystal data

$C_{14}H_{15}NO_2$

$M_r = 229.27$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P\ 2_1/c$

$a = 7.8132\ (16)\ \text{\AA}$

$b = 5.7947\ (12)\ \text{\AA}$

$c = 26.175\ (5)\ \text{\AA}$

$\beta = 95.02\ (3)^\circ$

$V = 1180.5\ (4)\ \text{\AA}^3$

$Z = 4$

$F(000) = 488$

$D_x = 1.290\ \text{Mg m}^{-3}$

Mo  $K\alpha$  radiation,  $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 8657 reflections

$\theta = 3.1\text{--}27.5^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 298\ \text{K}$

Prism, colorless

$0.40 \times 0.30 \times 0.20\ \text{mm}$

#### Data collection

Rigaku SCXmini  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution:  $13.6612\ \text{pixels mm}^{-1}$

CCD\_Profile\_fitting scans

Absorption correction: multi-scan  
(*CrystalClear*; Rigaku, 2005)

$T_{\min} = 0.970$ ,  $T_{\max} = 0.983$

10971 measured reflections

2693 independent reflections

1692 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.056$   
 $\theta_{\text{max}} = 27.5^\circ$ ,  $\theta_{\text{min}} = 3.1^\circ$   
 $h = -9 \rightarrow 10$

$k = -7 \rightarrow 7$   
 $l = -33 \rightarrow 33$

### Refinement

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.056$   
 $wR(F^2) = 0.128$   
 $S = 1.03$   
 2693 reflections  
 163 parameters  
 2 restraints  
 Primary atom site location: structure-invariant  
 direct methods

Secondary atom site location: difference Fourier  
 map  
 Hydrogen site location: inferred from  
 neighbouring sites  
 H atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0527P)^2 + 0.1931P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.17 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.22 \text{ e } \text{\AA}^{-3}$

### 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 ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O2	0.12511 (17)	0.7060 (2)	0.48819 (5)	0.0369 (4)
N1	0.1955 (2)	0.1501 (3)	0.54954 (6)	0.0322 (4)
C8	0.2161 (2)	0.0685 (3)	0.60126 (6)	0.0301 (4)
C7	0.3137 (2)	0.3330 (3)	0.53561 (7)	0.0350 (5)
H7A	0.4309	0.2888	0.5467	0.042*
H7B	0.2883	0.4746	0.5532	0.042*
C11	0.2449 (3)	-0.1048 (4)	0.70119 (7)	0.0395 (5)
C6	0.2976 (2)	0.3745 (3)	0.47851 (6)	0.0317 (4)
C1	0.2033 (2)	0.5582 (3)	0.45649 (6)	0.0295 (4)
C9	0.3148 (2)	0.1824 (3)	0.64015 (7)	0.0370 (5)
H9	0.3717	0.3180	0.6330	0.044*
O1	0.2685 (2)	-0.1765 (3)	0.75158 (5)	0.0600 (5)
C2	0.1942 (2)	0.5952 (3)	0.40425 (7)	0.0365 (5)
H2A	0.1322	0.7199	0.3900	0.044*
C10	0.3290 (3)	0.0950 (3)	0.68955 (7)	0.0410 (5)
H10	0.3961	0.1721	0.7152	0.049*
C13	0.1307 (3)	-0.1295 (3)	0.61362 (7)	0.0397 (5)
H13	0.0618	-0.2056	0.5882	0.048*
C5	0.3797 (3)	0.2304 (3)	0.44614 (8)	0.0442 (5)
H5	0.4435	0.1067	0.4601	0.053*
C12	0.1450 (3)	-0.2177 (4)	0.66294 (7)	0.0430 (5)

H12	0.0874	-0.3525	0.6703	0.052*
C4	0.3699 (3)	0.2647 (4)	0.39393 (8)	0.0488 (6)
H4	0.4257	0.1649	0.3731	0.059*
C3	0.2769 (3)	0.4478 (4)	0.37300 (7)	0.0437 (5)
H3	0.2696	0.4726	0.3378	0.052*
C14	0.1791 (4)	-0.3750 (4)	0.76569 (8)	0.0698 (8)
H14A	0.0576	-0.3478	0.7603	0.105*
H14B	0.2096	-0.4089	0.8012	0.105*
H14C	0.2092	-0.5034	0.7451	0.105*
H1	0.195 (2)	0.032 (2)	0.5294 (6)	0.046 (6)*
H2	0.0288 (19)	0.756 (4)	0.4738 (9)	0.083 (9)*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O2	0.0365 (8)	0.0393 (8)	0.0342 (7)	0.0069 (7)	-0.0009 (6)	-0.0054 (6)
N1	0.0345 (9)	0.0315 (9)	0.0301 (9)	-0.0004 (7)	-0.0006 (7)	-0.0026 (7)
C8	0.0285 (10)	0.0322 (10)	0.0295 (9)	0.0057 (8)	0.0021 (8)	-0.0013 (8)
C7	0.0339 (11)	0.0342 (11)	0.0358 (10)	0.0013 (9)	-0.0029 (8)	0.0008 (8)
C11	0.0444 (12)	0.0466 (12)	0.0278 (10)	0.0049 (10)	0.0054 (9)	0.0003 (9)
C6	0.0299 (10)	0.0330 (10)	0.0319 (10)	-0.0002 (8)	0.0016 (8)	0.0000 (8)
C1	0.0267 (9)	0.0296 (10)	0.0325 (10)	-0.0044 (8)	0.0037 (8)	-0.0040 (8)
C9	0.0397 (11)	0.0355 (11)	0.0360 (11)	-0.0041 (9)	0.0036 (9)	-0.0029 (8)
O1	0.0814 (12)	0.0676 (11)	0.0304 (8)	-0.0100 (9)	0.0011 (8)	0.0072 (7)
C2	0.0355 (11)	0.0379 (11)	0.0355 (10)	0.0016 (9)	0.0005 (9)	0.0046 (9)
C10	0.0441 (12)	0.0468 (12)	0.0309 (10)	-0.0020 (10)	-0.0029 (9)	-0.0083 (9)
C13	0.0436 (12)	0.0420 (12)	0.0326 (10)	-0.0072 (9)	-0.0015 (9)	-0.0043 (9)
C5	0.0486 (13)	0.0406 (12)	0.0434 (12)	0.0136 (10)	0.0038 (10)	0.0013 (9)
C12	0.0496 (13)	0.0424 (12)	0.0372 (11)	-0.0101 (10)	0.0057 (10)	0.0023 (9)
C4	0.0520 (13)	0.0541 (14)	0.0415 (12)	0.0128 (11)	0.0106 (10)	-0.0083 (10)
C3	0.0455 (12)	0.0547 (13)	0.0317 (10)	-0.0011 (11)	0.0073 (9)	0.0000 (9)
C14	0.111 (2)	0.0580 (16)	0.0417 (13)	-0.0056 (15)	0.0122 (14)	0.0121 (11)

*Geometric parameters (Å, °)*

O2—C1	1.372 (2)	C9—H9	0.9300
O2—H2	0.862 (10)	O1—C14	1.412 (3)
N1—C8	1.430 (2)	C2—C3	1.382 (3)
N1—C7	1.472 (2)	C2—H2A	0.9300
N1—H1	0.864 (9)	C10—H10	0.9300
C8—C13	1.380 (3)	C13—C12	1.384 (3)
C8—C9	1.389 (2)	C13—H13	0.9300
C7—C6	1.508 (2)	C5—C4	1.376 (3)
C7—H7A	0.9700	C5—H5	0.9300
C7—H7B	0.9700	C12—H12	0.9300
C11—C10	1.378 (3)	C4—C3	1.372 (3)
C11—C12	1.379 (3)	C4—H4	0.9300
C11—O1	1.380 (2)	C3—H3	0.9300

C6—C5	1.386 (3)	C14—H14A	0.9600
C6—C1	1.390 (2)	C14—H14B	0.9600
C1—C2	1.380 (2)	C14—H14C	0.9600
C9—C10	1.384 (3)		
C1—O2—H2	111.4 (17)	C1—C2—H2A	119.8
C8—N1—C7	116.84 (14)	C3—C2—H2A	119.8
C8—N1—H1	108.1 (13)	C11—C10—C9	120.88 (18)
C7—N1—H1	112.8 (13)	C11—C10—H10	119.6
C13—C8—C9	118.17 (17)	C9—C10—H10	119.6
C13—C8—N1	118.71 (16)	C8—C13—C12	121.63 (18)
C9—C8—N1	123.10 (17)	C8—C13—H13	119.2
N1—C7—C6	111.14 (15)	C12—C13—H13	119.2
N1—C7—H7A	109.4	C4—C5—C6	122.09 (19)
C6—C7—H7A	109.4	C4—C5—H5	119.0
N1—C7—H7B	109.4	C6—C5—H5	119.0
C6—C7—H7B	109.4	C11—C12—C13	119.77 (19)
H7A—C7—H7B	108.0	C11—C12—H12	120.1
C10—C11—C12	119.22 (18)	C13—C12—H12	120.1
C10—C11—O1	115.96 (18)	C3—C4—C5	119.31 (19)
C12—C11—O1	124.81 (19)	C3—C4—H4	120.3
C5—C6—C1	117.69 (17)	C5—C4—H4	120.3
C5—C6—C7	120.43 (17)	C4—C3—C2	120.01 (18)
C1—C6—C7	121.88 (16)	C4—C3—H3	120.0
O2—C1—C2	121.01 (16)	C2—C3—H3	120.0
O2—C1—C6	118.35 (15)	O1—C14—H14A	109.5
C2—C1—C6	120.59 (16)	O1—C14—H14B	109.5
C10—C9—C8	120.31 (18)	H14A—C14—H14B	109.5
C10—C9—H9	119.8	O1—C14—H14C	109.5
C8—C9—H9	119.8	H14A—C14—H14C	109.5
C11—O1—C14	117.88 (17)	H14B—C14—H14C	109.5
C1—C2—C3	120.31 (18)		
C7—N1—C8—C13	-168.27 (16)	C6—C1—C2—C3	0.9 (3)
C7—N1—C8—C9	13.4 (2)	C12—C11—C10—C9	-0.3 (3)
C8—N1—C7—C6	170.60 (14)	O1—C11—C10—C9	179.82 (18)
N1—C7—C6—C5	-80.6 (2)	C8—C9—C10—C11	-0.5 (3)
N1—C7—C6—C1	100.3 (2)	C9—C8—C13—C12	-1.6 (3)
C5—C6—C1—O2	-178.27 (16)	N1—C8—C13—C12	-179.97 (17)
C7—C6—C1—O2	0.8 (3)	C1—C6—C5—C4	-0.1 (3)
C5—C6—C1—C2	-0.6 (3)	C7—C6—C5—C4	-179.12 (19)
C7—C6—C1—C2	178.44 (17)	C10—C11—C12—C13	0.1 (3)
C13—C8—C9—C10	1.3 (3)	O1—C11—C12—C13	179.99 (18)
N1—C8—C9—C10	179.69 (17)	C8—C13—C12—C11	0.8 (3)
C10—C11—O1—C14	177.40 (19)	C6—C5—C4—C3	0.4 (3)
C12—C11—O1—C14	-2.5 (3)	C5—C4—C3—C2	-0.1 (3)
O2—C1—C2—C3	178.53 (17)	C1—C2—C3—C4	-0.6 (3)

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
N1—H1 $\cdots$ O2 <sup>i</sup>	0.86 (1)	2.22 (1)	3.058 (2)	163 (2)
O2—H2 $\cdots$ N1 <sup>ii</sup>	0.86 (1)	1.89 (1)	2.741 (2)	172 (2)

Symmetry codes: (i)  $x, y-1, z$ ; (ii)  $-x, -y+1, -z+1$ .