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(3-Aminophenyl)methanol

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Key indicators: single-crystal X-ray study; T = 200 K; mean σ (C–C) = 0.002 Å; R factor = 0.033; wR factor = 0.084; data-to-parameter ratio = 10.2.

In the title compound, C_7H_9NO , a derivative of benzyl alcohol, the endocyclic C–C–C angles are in the range 119.50 (12)– 121.04 (12)°. In the crystal, molecules are linked by N–H···O hydrogen-bond interactions, forming an extended two-dimensional framework parallel to *ab*. O–H···N interactions are also observed.

Related literature

For the crystal structure of (3-(hydroxymethyl)phenyl)-bis-(diphenylphosphinomethyl)amine, see: Hursthouse *et al.* (2003). For the crystal structure of 3-nitrobenzyl alcohol as a co-crystal with platinum-containing coordination compounds, see: Oskui *et al.* (1999). For graph-set analysis of hydrogen bonds, see: Etter *et al.* (1990); Bernstein *et al.* (1995). For the use of chelating ligands in coordination chemistry, see: Gade (1998).



Experimental

a = 4.7977 (4) Å
b = 6.2954 (6) Å
c = 21.6341 (18)

 $V = 653.42 (10) \text{ Å}^3$ Z = 4Mo *K*\alpha radiation

Data collection

Bruker APEXII CCD diffractometer Absorption correction: multi-scan (SADABS; Bruker, 2008) $T_{min} = 0.869, T_{max} = 1.000$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.033$ $wR(F^2) = 0.084$ S = 1.11961 reflections 94 parameters

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O1 - H81 \cdots N1^{i}$ $N1 - H71 \cdots O1^{ii}$ $N1 - H72 \cdots O1^{iii}$	0.85 (3) 0.87 (2) 0.904 (19)	2.02 (3) 2.19 (2) 2.24 (2)	2.8620 (18) 3.0588 (16) 3.1204 (16)	171 (2) 177 (2) 165.3 (16)

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

 $0.53 \times 0.47 \times 0.19 \text{ mm}$

6010 measured reflections

961 independent reflections

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

H atoms treated by a mixture of

independent and constrained

T = 200 K

 $R_{\rm int}=0.020$

refinement

 $\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.16 \text{ e } \text{\AA}^{-3}$

Symmetry codes: (i) x, y - 1, z; (ii) x + 1, y + 1, z; (iii) $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *APEX2* (Bruker, 2010); cell refinement: *SAINT* (Bruker, 2010); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BX2363).

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supporting information

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S1. Comment

Chelate ligands have found widespread use in coordination chemistry due to the enhanced thermodynamic stability of resultant coordination compounds in relation to coordination compounds exclusively applying comparable monodentate ligands (Gade, 1998). Combining two different donor atoms, a molecular set-up to accomodate a large variety of metal centers of variable Lewis acidity is at hand. In this aspect, 3-aminobenzyl alcohol seemed of interest due to its possible use as a strictly neutral or, depending on the pH value, as an anionic or cationic ligand. In addition, due to the set-up of its functional groups, it may act as mono- or bidentate ligand offering the possibility to create seven-membered chelate rings. To enable comparative studies in terms of bond lengths and angles in envisioned coordination compounds, we determined the molecular and crystal structure of the title compound. Information about the crystal structure of (3-(Hy-droxymethyl)phenyl)-bis(diphenylphosphinomethyl)amine (Hursthouse *et al.*, 2003) as well as 3-nitrobenzyl alcohol as a co-crystallizate with platinum-containing coordination compounds (Oskui *et al.*, 1999) is available in the literature.

The hydroxymethyl group is not in plane with the aromatic system, the respective dihedral angle was found at 33.0 (2) °. Endocyclic C–C–C angles hardly deviate from the expected ideal values of 120 ° and range from 119.50 (12)–121.04 (12) °. The biggest angle is found on the C atom bearing the amino group (Fig. 1).

In the crystal structure, a cooperative set of hydrogen bonds involving all nitrogen- and oxygen-bonded hydrogen atoms is present. While the oxygen atom acts as twofold acceptor for hydrogen bonds exclusively stemming from the H atoms of the amino group, the nitrogen atom of the amino group serves as acceptor for a hydrogen bond originating from the hydroxyl group's O atom. In terms of graph-set analysis (Etter *et al.*, 1990; Bernstein *et al.*, 1995), the descriptor for this hydrogen bonding system on the unitary level is $C_1^1(7)C_1^1(7)C_1^1(7)$. In the crystal the molecules are linked by N— H···O hydrogen-bond interactions, forming an extended two-dimensional framework parallel to the ab, Fig. 2. π -Stacking is not a prominent feature of the crystal structure with the shortest intercentroid distance between two aromatic systems measured at 5.5741 (10) Å .(Fig. 2).

S2. Experimental

The compound was obtained commercially (Aldrich). Crystals suitable for the X-ray diffraction study were obtained upon free evaporation of a solution of the compound in acetonitrile at room temperature.

S3. Refinement

Carbon-bound H atoms were placed in calculated positions (C—H 0.99-0.95 Å) and were included in the refinement in the riding model approximation, with U(H) set to $1.2U_{eq}(C)$. The hydrogen atoms of the hydroxyl group as well as of the amino group were located on a difference Fourier map and refined freely.



Figure 1

The molecular structure of the title compound, with atom labels and anisotropic displacement ellipsoids (drawn at 50% probability level).



Figure 2

Intermolecular contacts, viewed along [-1 0 0]. Symmetry operators: ⁱ x, y - 1, z; ⁱⁱ x - 1, y - 1, z; ⁱⁱⁱ -x + 1, y - 1/2, -z + 1/2; ^{iv} -x + 1, y + 1, z; ^{iv} -x + 1, y + 1/2, -z + 1/2; ^v x, y + 1, z; ^{vi} x + 1, y + 1, z.

(3-Aminophenyl)methanol

Crystal data	
C ₇ H ₉ NO	F(000) = 264
$M_r = 123.15$	$D_{\rm x} = 1.252 {\rm ~Mg} {\rm ~m}^{-3}$
Orthorhombic, $P2_12_12_1$	Mo <i>K</i> α radiation, $\lambda = 0.71073$ Å
Hall symbol: P 2ac 2ab	Cell parameters from 5183 reflections
a = 4.7977 (4) Å	$\theta = 3.4 - 28.1^{\circ}$
b = 6.2954 (6) Å	$\mu=0.09~\mathrm{mm^{-1}}$
c = 21.6341 (18) Å	T = 200 K
$V = 653.42 (10) Å^3$	Platelet, brown
Z = 4	$0.53\times0.47\times0.19~mm$

Data collection

Bruker APEXII CCD diffractometer Radiation source: fine-focus sealed tube Graphite monochromator φ and ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2008) $T_{\min} = 0.869, T_{\max} = 1.000$ <i>Refinement</i>	6010 measured reflections 961 independent reflections 924 reflections with $I > 2\sigma(I)$ $R_{int} = 0.020$ $\theta_{max} = 28.0^{\circ}, \ \theta_{min} = 3.4^{\circ}$ $h = -6 \rightarrow 6$ $k = -7 \rightarrow 8$ $l = -23 \rightarrow 28$
Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.033$ $wR(F^2) = 0.084$ S = 1.11 961 reflections 94 parameters 0 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 $w = 1/[\sigma^2(F_o^2) + (0.0484P)^2 + 0.1019P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.19 \text{ e } \text{Å}^{-3}$ $\Delta\rho_{min} = -0.16 \text{ e } \text{Å}^{-3}$

Special details

Refinement. Due to the absence of a strong anomalous scatterer, the Flack parameter is meaningless. Thus, Friedel opposites (600 pairs) have been merged and the item was removed from the CIF.

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
01	0.3733 (2)	-0.37165 (17)	0.18291 (4)	0.0277 (3)	
H81	0.524 (6)	-0.439 (4)	0.1893 (9)	0.049 (6)*	
N1	0.8480 (3)	0.35877 (19)	0.19920 (6)	0.0287 (3)	
H71	1.002 (5)	0.432 (3)	0.1955 (9)	0.040 (5)*	
H72	0.811 (4)	0.303 (3)	0.2368 (9)	0.035 (5)*	
C1	0.3760 (3)	-0.2875 (2)	0.12190 (6)	0.0313 (3)	
H1A	0.1858	-0.2383	0.1114	0.038*	
H1B	0.4250	-0.4028	0.0927	0.038*	
C2	0.5769 (3)	-0.1056 (2)	0.11274 (6)	0.0236 (3)	
C3	0.6316 (3)	0.0359 (2)	0.16033 (6)	0.0252 (3)	
H3	0.5471	0.0147	0.1996	0.030*	
C4	0.8087 (3)	0.2089 (2)	0.15151 (6)	0.0237 (3)	
C5	0.9317 (3)	0.2381 (2)	0.09359 (7)	0.0303 (3)	
H5	1.0527	0.3552	0.0867	0.036*	
C6	0.8774 (3)	0.0967 (2)	0.04623 (7)	0.0325 (3)	
H6	0.9615	0.1177	0.0070	0.039*	
C7	0.7016 (3)	-0.0758 (2)	0.05529 (6)	0.0278 (3)	
H7	0.6667	-0.1727	0.0225	0.033*	

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01	0.0305 (5)	0.0253 (5)	0.0271 (5)	-0.0050 (5)	0.0030 (4)	0.0028 (4)
N1	0.0328 (6)	0.0248 (6)	0.0284 (6)	-0.0090 (5)	0.0021 (5)	-0.0024 (5)
C1	0.0372 (7)	0.0310 (7)	0.0259 (6)	-0.0156 (7)	-0.0035 (6)	0.0025 (5)
C2	0.0230 (6)	0.0219 (6)	0.0259 (6)	-0.0044 (5)	-0.0029 (5)	0.0024 (5)
C3	0.0274 (6)	0.0257 (6)	0.0225 (5)	-0.0066 (6)	0.0022 (5)	0.0013 (5)
C4	0.0239 (6)	0.0209 (6)	0.0265 (6)	-0.0022 (6)	-0.0011 (5)	0.0004 (5)
C5	0.0320 (7)	0.0275 (7)	0.0313 (7)	-0.0102 (6)	0.0051 (6)	0.0015 (6)
C6	0.0359 (7)	0.0359 (7)	0.0257 (6)	-0.0078 (7)	0.0071 (6)	0.0011 (6)
C7	0.0306 (6)	0.0281 (7)	0.0248 (6)	-0.0043 (6)	-0.0002 (6)	-0.0024 (5)

Atomic displacement parameters $(Å^2)$

Geometric parameters (Å, °)

01—C1	1.4222 (16)	С2—С7	1.3922 (19)
O1—H81	0.85 (3)	C3—C4	1.3945 (19)
N1—C4	1.4105 (17)	С3—Н3	0.9500
N1—H71	0.87 (2)	C4—C5	1.3972 (19)
N1—H72	0.904 (19)	C5—C6	1.382 (2)
C1—C2	1.5100 (19)	С5—Н5	0.9500
C1—H1A	0.9900	C6—C7	1.389 (2)
C1—H1B	0.9900	С6—Н6	0.9500
C2—C3	1.3865 (19)	С7—Н7	0.9500
C1—O1—H81	109.2 (14)	С2—С3—Н3	119.5
C4—N1—H71	113.4 (13)	С4—С3—Н3	119.5
C4—N1—H72	111.9 (12)	C3—C4—C5	118.85 (12)
H71—N1—H72	117.0 (16)	C3—C4—N1	120.24 (12)
O1—C1—C2	114.20 (11)	C5—C4—N1	120.78 (12)
O1—C1—H1A	108.7	C6—C5—C4	120.04 (13)
C2C1H1A	108.7	C6—C5—H5	120.0
O1—C1—H1B	108.7	C4—C5—H5	120.0
C2C1H1B	108.7	C5—C6—C7	120.90 (13)
H1A—C1—H1B	107.6	С5—С6—Н6	119.6
C3—C2—C7	119.66 (12)	С7—С6—Н6	119.6
C3—C2—C1	120.72 (12)	C6—C7—C2	119.50 (12)
C7—C2—C1	119.58 (12)	С6—С7—Н7	120.3
C2—C3—C4	121.04 (12)	С2—С7—Н7	120.3
O1—C1—C2—C3	-33.0 (2)	C3—C4—C5—C6	-0.1 (2)
O1—C1—C2—C7	149.06 (13)	N1-C4-C5-C6	-176.05 (14)
C7—C2—C3—C4	0.4 (2)	C4—C5—C6—C7	-0.1 (2)
C1—C2—C3—C4	-177.54 (13)	C5—C6—C7—C2	0.5 (2)
C2—C3—C4—C5	0.0 (2)	C3—C2—C7—C6	-0.6 (2)
C2—C3—C4—N1	175.95 (13)	C1—C2—C7—C6	177.36 (13)

D—H···A	<i>D</i> —Н	H···A	$D \cdots A$	D—H···A
01—H81…N1 ⁱ	0.85 (3)	2.02 (3)	2.8620 (18)	171 (2)
N1—H71····O1 ⁱⁱ	0.87 (2)	2.19 (2)	3.0588 (16)	177 (2)
N1—H72…O1 ⁱⁱⁱ	0.904 (19)	2.24 (2)	3.1204 (16)	165.3 (16)

Hydrogen-bond geometry (Å, °)

Symmetry codes: (i) *x*, *y*-1, *z*; (ii) *x*+1, *y*+1, *z*; (iii) -*x*+1, *y*+1/2, -*z*+1/2.