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1-(4-Amino-3,5-dichlorophenyl)ethanol

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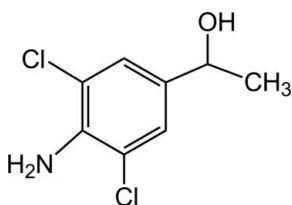
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 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.032; wR factor = 0.128; data-to-parameter ratio = 11.8.

The asymmetric unit of the title compound, $\text{C}_8\text{H}_9\text{Cl}_2\text{NO}$, contains two crystallographically independent molecules which are connected *via* an $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond. There is aromatic $\pi-\pi$ stacking in the crystal, with a centroid-centroid distance between benzene rings of 3.48 (2) Å. The crystal packing is stabilized by intermolecular hydrogen bonds.

Related literature

For the synthetic use of the title compound and related compounds, see: Judkins *et al.* (1991); Ehrhardt (1990); Kelsner (2007); Lu (2001); Pri-Bar *et al.* (1990); Shukrallah *et al.* (2004).



Experimental

Crystal data

$\text{C}_8\text{H}_9\text{Cl}_2\text{NO}$
 $M_r = 206.06$
 Monoclinic, $C2/c$
 $a = 16.472$ (2) Å
 $b = 16.110$ (2) Å
 $c = 14.6756$ (19) Å
 $\beta = 107.049$ (2)°

$V = 3723.2$ (8) Å³
 $Z = 16$
 Mo $K\alpha$ radiation
 $\mu = 0.65$ mm⁻¹
 $T = 296$ K
 $0.15 \times 0.13 \times 0.08$ mm

Data collection

Bruker APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2007)
 $T_{\min} = 0.909$, $T_{\max} = 0.950$

9273 measured reflections
 3279 independent reflections
 2855 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.128$
 $S = 1.06$
 3279 reflections
 277 parameters
 1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.42$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.46$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H13}\cdots\text{O2}^i$	0.82 (3)	2.17 (3)	2.927 (3)	154 (3)
$\text{O1}-\text{H16}\cdots\text{N1}^{ii}$	0.83 (3)	2.12 (3)	2.941 (3)	169 (3)
$\text{O2}-\text{H18}\cdots\text{N2}^{iii}$	0.79 (3)	2.13 (3)	2.912 (3)	172 (2)

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

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); 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.*, 2006); software used to prepare material for publication: publCIF (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZK2012).

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supporting information

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1-(4-Amino-3,5-dichlorophenyl)ethanol

Peng Liu, Jian-Feng Wu, Ping-An Wang, Wei He and Hai-Bo Wang

S1. Comment

We report here the crystal structure of the title compound 1-(4-amino-3,5-dichlorophenyl) ethanol (I), a typical secondary aromatic alcohol and an important organic building block (Fig. 1). Bond lengths and angles are within normal ranges. As part of our ongoing studies of the secondary aromatic alcohol, the title compound was synthesized and characterized by X-ray diffraction. The asymmetric unit of the title compound (I) contains two crystallographically independent molecules in which both molecules are connected *via* N—H \cdots O hydrogen bonds. The dihedral angle between the two benzene rings is 60.49°. The packing of molecules in the crystal structure is stabilized by intermolecular O—H \cdots N hydrogen bonds.

S2. Experimental

1-(4-amino-3,5-dichlorophenyl)ethanone (1.03 g, 5 mmol) was dissolved in anhydrous methanol (25 ml) at room temperature, followed by addition of NaBH₄ (0.227 g, 6 mmol). The mixture was stirred vigorously at room temperature until TLC showed no ethanone. The solvent was evaporated to dryness under reduced pressure to obtain a crude product, which was purified by a flash column chromatography (n-hexane/EtOAc 8:1) to afford pure colorless compound in 89.6% yield. Then, the title compound (40 mg, 0.19 mmol) was dissolved in ethyl acetate/n-hexane (7:3, 15 ml). Colorless crystals were isolated after several days.

S3. Refinement

In both structures all the H atoms were discernible in the difference Fourier maps. However, they were constrained by riding model approximation. C—H_{methyl} = 0.96 Å; C—H_{aryl} = 0.93 Å; $U_{\text{iso}}(\text{H}_{\text{methyl}})$ and $U_{\text{iso}}(\text{H}_{\text{aryl}})$ are 1.5 $U_{\text{eq}}(\text{C})$ and 1.2 $U_{\text{eq}}(\text{C})$, respectively.

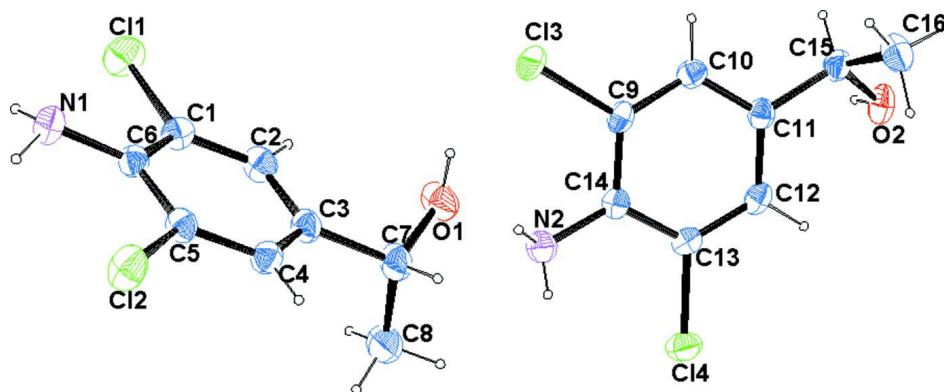


Figure 1

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

1-(4-Amino-3,5-dichlorophenyl)ethanol

Crystal data

$C_8H_9Cl_2NO$
 $M_r = 206.06$
 Monoclinic, $C2/c$
 $a = 16.472(2) \text{ \AA}$
 $b = 16.110(2) \text{ \AA}$
 $c = 14.6756(19) \text{ \AA}$
 $\beta = 107.049(2)^\circ$
 $V = 3723.2(8) \text{ \AA}^3$
 $Z = 16$

$F(000) = 1696$
 $D_x = 1.470 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 5355 reflections
 $\theta = 2.6\text{--}28.3^\circ$
 $\mu = 0.65 \text{ mm}^{-1}$
 $T = 296 \text{ K}$
 Block, colorless
 $0.15 \times 0.13 \times 0.08 \text{ mm}$

Data collection

Bruker APEXII CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (*SADABS*; Bruker, 2007)
 $T_{\min} = 0.909$, $T_{\max} = 0.950$

9273 measured reflections
 3279 independent reflections
 2855 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 1.8^\circ$
 $h = -19 \rightarrow 15$
 $k = -19 \rightarrow 17$
 $l = -17 \rightarrow 16$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.128$
 $S = 1.06$
 3279 reflections
 277 parameters
 1 restraint
 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
 $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.42 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.46 \text{ e \AA}^{-3}$

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	1.10844 (4)	0.04952 (4)	0.86690 (4)	0.0620 (2)
C12	1.09417 (4)	0.16958 (4)	1.20328 (4)	0.0598 (2)
C13	0.70719 (3)	-0.07065 (4)	0.67509 (4)	0.0557 (2)
C14	0.44673 (4)	0.14455 (3)	0.61899 (4)	0.0527 (2)
O1	0.78216 (11)	0.06516 (11)	0.85101 (13)	0.0617 (5)
O2	0.31701 (9)	-0.14705 (11)	0.48677 (12)	0.0519 (4)
N1	1.18071 (11)	0.10254 (12)	1.06839 (15)	0.0468 (4)
N2	0.63069 (12)	0.09603 (13)	0.67790 (13)	0.0440 (4)
C1	1.04879 (13)	0.08619 (12)	0.93915 (14)	0.0416 (5)
C2	0.96140 (14)	0.08983 (14)	0.90409 (15)	0.0458 (5)
C3	0.91343 (12)	0.11733 (12)	0.96181 (14)	0.0414 (5)
C4	0.95554 (13)	0.14186 (12)	1.05372 (15)	0.0408 (5)
C5	1.04275 (12)	0.13832 (12)	1.08668 (13)	0.0389 (4)
C6	1.09322 (12)	0.11026 (11)	1.03173 (13)	0.0375 (4)
C7	0.81676 (14)	0.12321 (15)	0.92535 (18)	0.0517 (5)
C8	0.78811 (16)	0.20783 (16)	0.8855 (2)	0.0712 (8)
H8A	0.7272	0.2094	0.8632	0.107*
H8B	0.8106	0.2193	0.8334	0.107*
H8C	0.8082	0.2489	0.9344	0.107*
C9	0.59864 (12)	-0.05077 (12)	0.64539 (13)	0.0377 (4)
C10	0.54210 (12)	-0.11507 (13)	0.61455 (14)	0.0388 (4)
C11	0.45556 (12)	-0.10121 (12)	0.58610 (13)	0.0380 (4)
C12	0.42693 (12)	-0.02043 (13)	0.58929 (14)	0.0406 (5)
C13	0.48443 (12)	0.04327 (12)	0.62044 (13)	0.0366 (4)
C14	0.57235 (11)	0.03095 (12)	0.65081 (12)	0.0354 (4)
C15	0.39451 (13)	-0.17286 (14)	0.55311 (16)	0.0458 (5)
C16	0.37270 (17)	-0.21407 (17)	0.6356 (2)	0.0578 (6)
H1	0.3668 (13)	-0.0092 (11)	0.5668 (14)	0.035 (5)*
H2	0.5640 (15)	-0.1665 (14)	0.6168 (17)	0.048 (6)*
H3	0.9238 (15)	0.1595 (14)	1.0923 (18)	0.048 (6)*
H4	0.4188 (13)	-0.2126 (14)	0.5260 (15)	0.045 (6)*
H5	0.9332 (14)	0.0707 (13)	0.8489 (16)	0.045 (6)*
H6	0.3307 (18)	-0.2551 (16)	0.612 (2)	0.069 (7)*
H8	0.6703 (16)	0.0815 (14)	0.7248 (17)	0.045 (6)*
H9	0.3513 (16)	-0.1676 (16)	0.6741 (18)	0.058 (7)*

H10	0.4207 (17)	-0.2342 (17)	0.6731 (19)	0.062 (7)*
H11	0.6117 (18)	0.1357 (18)	0.6943 (19)	0.061 (9)*
H12	0.7895 (15)	0.1125 (15)	0.9809 (17)	0.051 (6)*
H13	1.2082 (18)	0.1076 (17)	1.030 (2)	0.062 (8)*
H15	1.2046 (18)	0.1337 (17)	1.118 (2)	0.062 (8)*
H16	0.794 (2)	0.016 (2)	0.867 (2)	0.084 (10)*
H18	0.3276 (19)	-0.1363 (17)	0.4388 (16)	0.070 (9)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0664 (4)	0.0764 (4)	0.0496 (4)	0.0058 (3)	0.0267 (3)	-0.0065 (3)
C12	0.0441 (3)	0.0892 (5)	0.0391 (3)	0.0033 (3)	0.0011 (2)	-0.0144 (3)
C13	0.0275 (3)	0.0586 (4)	0.0742 (4)	0.0044 (2)	0.0044 (2)	-0.0059 (3)
C14	0.0508 (4)	0.0457 (3)	0.0588 (4)	0.0116 (2)	0.0116 (3)	-0.0049 (2)
O1	0.0475 (9)	0.0491 (10)	0.0669 (11)	-0.0050 (7)	-0.0171 (8)	-0.0017 (8)
O2	0.0332 (8)	0.0783 (11)	0.0422 (9)	-0.0106 (7)	0.0082 (6)	-0.0022 (7)
N1	0.0341 (9)	0.0582 (12)	0.0472 (11)	-0.0009 (8)	0.0106 (8)	-0.0004 (9)
N2	0.0387 (10)	0.0460 (10)	0.0430 (10)	-0.0016 (8)	0.0053 (8)	-0.0039 (8)
C1	0.0448 (11)	0.0432 (11)	0.0381 (10)	0.0021 (9)	0.0142 (9)	0.0011 (8)
C2	0.0463 (12)	0.0497 (12)	0.0345 (11)	-0.0036 (9)	0.0010 (9)	-0.0027 (9)
C3	0.0345 (10)	0.0410 (10)	0.0432 (11)	-0.0028 (8)	0.0027 (8)	0.0011 (8)
C4	0.0359 (11)	0.0431 (11)	0.0418 (11)	0.0021 (8)	0.0089 (9)	-0.0019 (8)
C5	0.0369 (10)	0.0421 (11)	0.0333 (10)	-0.0026 (8)	0.0033 (8)	-0.0008 (8)
C6	0.0337 (10)	0.0377 (10)	0.0395 (10)	0.0011 (7)	0.0082 (8)	0.0054 (8)
C7	0.0368 (11)	0.0531 (12)	0.0561 (13)	-0.0047 (9)	-0.0009 (10)	0.0000 (10)
C8	0.0454 (14)	0.0526 (14)	0.101 (2)	0.0025 (11)	-0.0017 (13)	0.0039 (13)
C9	0.0269 (9)	0.0488 (11)	0.0348 (10)	0.0023 (8)	0.0050 (7)	0.0024 (8)
C10	0.0365 (10)	0.0404 (11)	0.0379 (10)	0.0041 (8)	0.0086 (8)	0.0008 (8)
C11	0.0341 (10)	0.0448 (11)	0.0345 (10)	-0.0027 (8)	0.0094 (8)	-0.0022 (8)
C12	0.0282 (10)	0.0541 (12)	0.0383 (10)	0.0026 (8)	0.0081 (8)	-0.0010 (9)
C13	0.0363 (10)	0.0413 (10)	0.0323 (9)	0.0073 (8)	0.0101 (8)	0.0007 (8)
C14	0.0331 (10)	0.0432 (10)	0.0283 (9)	-0.0013 (8)	0.0068 (7)	0.0001 (8)
C15	0.0352 (11)	0.0500 (12)	0.0525 (13)	-0.0036 (9)	0.0132 (9)	-0.0109 (10)
C16	0.0518 (15)	0.0547 (14)	0.0610 (15)	-0.0106 (12)	0.0073 (12)	0.0077 (12)

Geometric parameters (Å, °)

C11—C1	1.7463 (19)	C4—H3	0.92 (2)
C12—C5	1.7462 (19)	C5—C6	1.392 (3)
C13—C9	1.7415 (19)	C7—C8	1.504 (3)
C14—C13	1.7437 (19)	C7—H12	1.05 (2)
O1—C7	1.423 (3)	C8—H8A	0.9600
O1—H16	0.84 (3)	C8—H8B	0.9600
O2—C15	1.422 (3)	C8—H8C	0.9600
O2—H18	0.792 (17)	C9—C10	1.378 (3)
N1—C6	1.388 (3)	C9—C14	1.395 (3)
N1—H13	0.82 (3)	C10—C11	1.381 (3)

N1—H15	0.87 (3)	C10—H2	0.90 (2)
N2—C14	1.399 (3)	C11—C12	1.390 (3)
N2—H8	0.83 (3)	C11—C15	1.514 (3)
N2—H11	0.78 (3)	C12—C13	1.380 (3)
C1—C2	1.381 (3)	C12—H1	0.96 (2)
C1—C6	1.396 (3)	C13—C14	1.399 (3)
C2—C3	1.389 (3)	C15—C16	1.513 (3)
C2—H5	0.86 (2)	C15—H4	0.91 (2)
C3—C4	1.381 (3)	C16—H6	0.95 (3)
C3—C7	1.527 (3)	C16—H9	1.06 (3)
C4—C5	1.376 (3)	C16—H10	0.88 (3)
C7—O1—H16	113 (2)	H8A—C8—H8B	109.5
C15—O2—H18	106 (2)	C7—C8—H8C	109.5
C6—N1—H13	116 (2)	H8A—C8—H8C	109.5
C6—N1—H15	115.7 (18)	H8B—C8—H8C	109.5
H13—N1—H15	108 (3)	C10—C9—C14	122.46 (17)
C14—N2—H8	109.3 (16)	C10—C9—C13	119.32 (15)
C14—N2—H11	113 (2)	C14—C9—C13	118.18 (14)
H8—N2—H11	105 (3)	C9—C10—C11	121.09 (19)
C2—C1—C6	122.87 (18)	C9—C10—H2	117.1 (15)
C2—C1—C11	119.93 (16)	C11—C10—H2	121.8 (15)
C6—C1—C11	117.20 (15)	C10—C11—C12	118.14 (18)
C1—C2—C3	120.23 (19)	C10—C11—C15	120.22 (18)
C1—C2—H5	123.4 (15)	C12—C11—C15	121.64 (17)
C3—C2—H5	116.1 (15)	C13—C12—C11	120.04 (17)
C4—C3—C2	118.29 (18)	C13—C12—H1	120.7 (11)
C4—C3—C7	119.86 (19)	C11—C12—H1	119.2 (11)
C2—C3—C7	121.82 (18)	C12—C13—C14	123.08 (17)
C5—C4—C3	120.33 (19)	C12—C13—C14	118.96 (15)
C5—C4—H3	121.3 (15)	C14—C13—C14	117.93 (15)
C3—C4—H3	118.4 (15)	C9—C14—C13	115.17 (16)
C4—C5—C6	123.30 (18)	C9—C14—N2	121.63 (17)
C4—C5—C12	119.27 (15)	C13—C14—N2	123.02 (18)
C6—C5—C12	117.44 (15)	O2—C15—C16	107.45 (18)
N1—C6—C5	122.21 (18)	O2—C15—C11	112.01 (18)
N1—C6—C1	122.73 (18)	C16—C15—C11	111.70 (19)
C5—C6—C1	114.97 (17)	O2—C15—H4	109.2 (14)
O1—C7—C8	106.7 (2)	C16—C15—H4	106.3 (13)
O1—C7—C3	111.71 (19)	C11—C15—H4	109.9 (14)
C8—C7—C3	111.63 (19)	C15—C16—H6	109.3 (17)
O1—C7—H12	108.6 (13)	C15—C16—H9	108.2 (13)
C8—C7—H12	107.1 (13)	H6—C16—H9	112 (2)
C3—C7—H12	110.9 (13)	C15—C16—H10	106.3 (16)
C7—C8—H8A	109.5	H6—C16—H10	113 (2)
C7—C8—H8B	109.5	H9—C16—H10	108 (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>
N1—H13···O2 ⁱ	0.82 (3)	2.17 (3)	2.927 (3)	154 (3)
O1—H16···N1 ⁱⁱ	0.83 (3)	2.12 (3)	2.941 (3)	169 (3)
O2—H18···N2 ⁱⁱⁱ	0.79 (3)	2.13 (3)	2.912 (3)	172 (2)

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