organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.013 Å R factor = 0.058 wR factor = 0.198 Data-to-parameter ratio = 15.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

4-Bromo-2,6-dichloroaniline

The title compound, $C_6H_4BrCl_2N$, crystallizes in the space group $P2_1/c$ with two molecules in the asymmetric unit. Molecules related by an *a*-axis translation are stacked over each other, bound by $\pi-\pi$ interactions. Molecules in adjacent stacks are linked to each other through weak $N-H\cdots N$ hydrogen bonds.

Comment

The title compound, (I), was synthesized for a study of the electrochemical reductive cleavage of carbon-halogen bonds (Arun Prasad & Sangaranarayanan, 2004). It crystallizes with two molecules in the asymmetric unit. The average C-Br and C-Cl bond lengths of 1.895 and 1.743 Å are in excellent agreement with the standard values of 1.899 (12) and 1.739 (10) Å, respectively (Allen *et al.*, 1987).



Molecules related by an *a*-axis translation are stacked over each other and the perpendicular distance between the stacking planes is 3.596 Å (Fig. 2). It is, therefore, inferred that the stacked molecules are bound to each other by π - π interactions (Hunter & Sanders, 1990). The stacked columns are weakly linked by N-H···N hydrogen bonds (Table 1).

Experimental

2,6-Dichloroaniline (21 g) was added to 6 M hydrochloric acid (120 ml) with continuous stirring. After complete dissolution of 2,6-





Received 28 January 2005 Accepted 1 February 2005 Online 26 February 2005

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved dichloroaniline, bromine vapour was passed into the solution, during which the reaction flask was cooled with ice. When the solution attained a yellow colour, the passage of bromine vapour was stopped and the final product was a dark pasty mass. Crude 4-bromo-2,6dichloroaniline was filtered off, washed well with water and thoroughly dried. It was then purified by column chromatography (silica gel) using hexane as eluent; slow solvent evaporation produced colourless crystals of (I).

> $D_x = 2.011 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 25 reflections $\theta = 10-15^{\circ}$

 $\mu = 5.75 \text{ mm}^{-1}$ T = 293 (2) KBlock, colourless $0.3 \times 0.2 \times 0.2 \text{ mm}$

 $R_{\rm int} = 0.061$

 $\theta_{\rm max} = 25.0^{\circ}$

 $h = 0 \rightarrow 4$

 $k = 0 \rightarrow 18$

 $l=-30\rightarrow 30$

2 standard reflections

frequency: 60 min

intensity decay: none

Crystal data

C ₆ H ₄ BrCl ₂ N
$M_r = 240.91$
Monoclinic, $P2_1/c$
a = 4.015 (3) Å
b = 15.41 (1) Å
c = 25.730(5) Å
$\beta = 91.22 \ (7)^{\circ}$
$V = 1591.6 (16) \text{ Å}^3$
Z = 8

Data collection

Enraf-Nonius CAD-4 diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan (North *et al.*, 1968) $T_{\min} = 0.187$, $T_{\max} = 0.316$ 3225 measured reflections 2796 independent reflections 1287 reflections with $I > 2\sigma(I)$

Refinement

 Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.1152P)^2]$
 $R[F^2 > 2\sigma(F^2)] = 0.059$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.198$ $(\Delta/\sigma)_{max} < 0.001$

 S = 1.00 $\Delta\rho_{max} = 0.66 \text{ e Å}^{-3}$

 2796 reflections
 $\Delta\rho_{min} = -0.64 \text{ e Å}^{-3}$

 182 parameters
 Extinction correction: SHELXL97

 H-atom parameters constrained
 Extinction coefficient: 0.0043 (11)

Table 1

Hydrogen-bonding geometry (Å, °).

D-II···A	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$N1-H1A\cdots N2^{i}$	0.86	2.39	3.148 (11)	147
$N2-H2A\cdots N1^{ii}$	0.86	2.41	3.199 (11)	152

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

All H atoms were positioned geometrically and were refined using the riding-model approximation, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm carrier \ atom})$ and C–H = 0.93 Å or N–H = 0.86 Å.



Figure 2 A packing diagram of the title compound, viewed down [100].

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

MAP thanks the CSIR, India, for the award of a Senior Research Fellowship. The authors gratefully acknowledge the instrumentation facility provided by SAIF, IIT Madras.

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