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Key indicators

Single-crystal X-ray study
 $T = 173$ K
 Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.022
 wR factor = 0.057
 Data-to-parameter ratio = 15.9

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

2,6-Dibromo-4-chlorobenzonitrile

2,6-Dibromo-4-chlorobenzonitrile, $\text{C}_7\text{H}_2\text{Br}_2\text{ClN}$, forms layers in the crystal structure, with $\text{Br} \cdots \text{N}$ contacts of 3.049 (2) Å the strongest intermolecular interactions. The crystal structure is isomorphous with 2,4,6-tribromobenzonitrile.

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Comment

Previously, the structure of 2,6-dibromo-4-chlorobenzonitrile (Britton *et al.*, 2002) was found to be isomorphous with 2,4,6-tribromobenzonitrile (Carter & Britton, 1972), based on the similarities between the cell dimensions. The structure has been determined in full to obtain a precise measurement of the short $\text{Br} \cdots \text{N}$ distance that was expected in the structure.

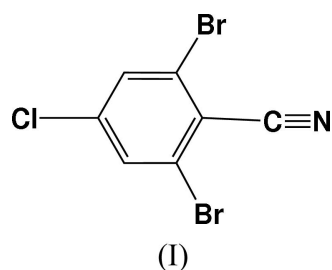


Fig. 1 shows the labeling and the anisotropic displacement ellipsoids. The bond lengths and angles are normal.

Fig. 2 shows the packing. The molecules form layers that are virtually identical with those in the tribromo analog. The $\text{Br} \cdots \text{N}$ contacts are 3.049 (2) Å, with $\text{C}-\text{Br} \cdots \text{N} = 166.0$ (2)° and $\text{Br} \cdots \text{N}-\text{C} = 133.2$ (3)°. This is consistent with other structures in which Br acts as a Lewis acid and N acts as a bifurcated Lewis base. There are also $\text{H} \cdots \text{Cl}$ contacts, with $\text{H} \cdots \text{Cl} = 3.08$ (3) Å, $\text{C}-\text{H} \cdots \text{Cl} = 16$ (2)° and $\text{H} \cdots \text{Cl}-\text{C} = 100$ (2)°; the angles are consistent with a $\text{C}-\text{H} \cdots \text{Cl}$ hydrogen

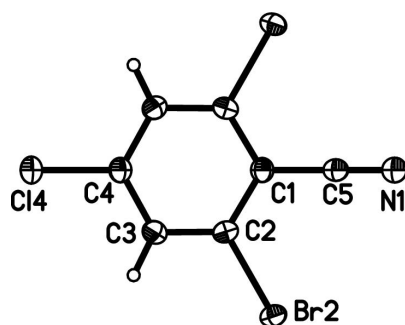


Figure 1

View of the title compound, with displacement ellipsoids shown at the 50% probability level. Unlabeled atoms are related to labeled atoms by $(x, \frac{1}{2} - y, z)$.

bond although the distance is slightly larger than the usual van der Waals distance for $\text{H} \cdots \text{Cl}$ (Bondi, 1964).

The layers are parallel to $(20\bar{1})$ with the molecules tilted by $1.4 (1)^\circ$ out of the layer. Adjacent layers are $3.318 (1) \text{ \AA}$ apart.

Experimental

The compound was prepared from the corresponding aniline *via* the Sandmeyer reaction. [For details of the preparation of the tribromo isomorph, see Carter & Britton (1972).]

Crystal data

$\text{C}_7\text{H}_2\text{Br}_2\text{ClN}$	$D_x = 2.346 \text{ Mg m}^{-3}$
$M_r = 295.37$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/m$	Cell parameters from 2165 reflections
$a = 8.666 (2) \text{ \AA}$	$\theta = 2.4\text{--}27.4^\circ$
$b = 10.125 (3) \text{ \AA}$	$\mu = 9.94 \text{ mm}^{-1}$
$c = 4.7776 (12) \text{ \AA}$	$T = 173 (2) \text{ K}$
$\beta = 93.98 (1)^\circ$	Needle, colorless
$V = 418.19 (19) \text{ \AA}^3$	$0.45 \times 0.15 \times 0.10 \text{ mm}$
$Z = 2$	

Data collection

Bruker SMART 1K CCD area-detector diffractometer	1001 independent reflections
ω scans	930 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996; Blessing, 1995)	$R_{\text{int}} = 0.024$
$T_{\text{min}} = 0.19$, $T_{\text{max}} = 0.37$	$\theta_{\text{max}} = 27.5^\circ$
2839 measured reflections	$h = -11 \rightarrow 5$
	$k = -13 \rightarrow 13$
	$l = -6 \rightarrow 6$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.033P)^2 + 0.25P]$
$R[F^2 > 2\sigma(F^2)] = 0.022$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.057$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.09$	$\Delta\rho_{\text{max}} = 0.43 \text{ e \AA}^{-3}$
1001 reflections	$\Delta\rho_{\text{min}} = -0.39 \text{ e \AA}^{-3}$
63 parameters	Extinction correction: <i>SHELXTL</i>
All H-atom parameters refined	Extinction coefficient: $0.047 (3)$

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine

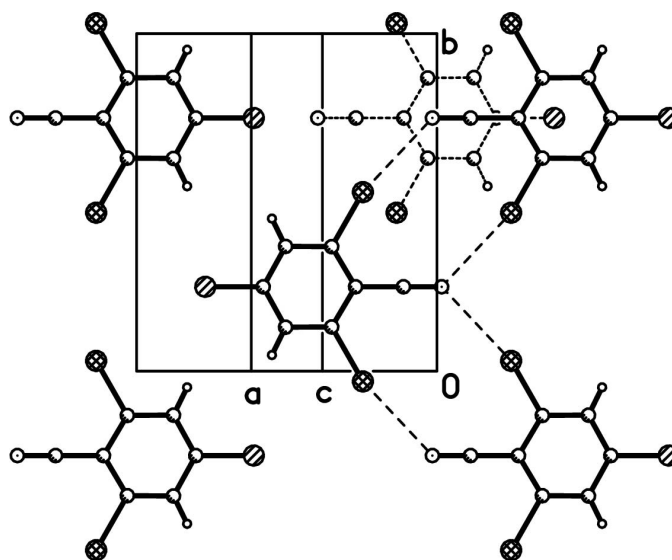


Figure 2

The packing in 2,6-dibromo-4-chlorobenzonitrile, viewed normal to $(20\bar{1})$. The molecules in one layer are shown with heavy bonds and the $\text{Br} \cdots \text{N}$ contacts are shown with dashed lines. One molecule in an adjacent layer is shown with dashed bonds.

structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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References

- Blessing, R. H. (1995). *Acta Cryst.* **A51**, 33–38.
 Bondi, A. (1964). *J. Phys. Chem.* **68**, 441–451.
 Britton, D., Noland, W. E. & Henke, T. A. (2002). *Acta Cryst.* **E58**, o185–o187.
 Bruker (2002). *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
 Carter, V. B. & Britton, D. (1972). *Acta Cryst.* **B28**, 945–950.
 Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
 Sheldrick, G. M. (1997). *SHELXTL*. Version 5.1. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.