

***trans*-1,4-Dibromocyclohexane-1,4-dicarboxylic acid**

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Key indicators

Single-crystal X-ray study
 $T = 110\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.026
 wR factor = 0.077
Data-to-parameter ratio = 14.6

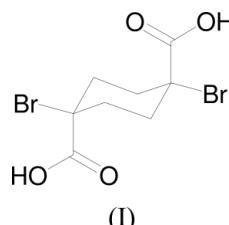
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Molecules of the title compound, $\text{C}_8\text{H}_{10}\text{Br}_2\text{O}_4$, located on symmetry centers, are in a rigid chair conformation, with the COOH and Br substituents axial and equatorial, respectively. The carboxylic acid groups form hydrogen bonds, in a cyclic motif, leading to infinite chains along the b axis.

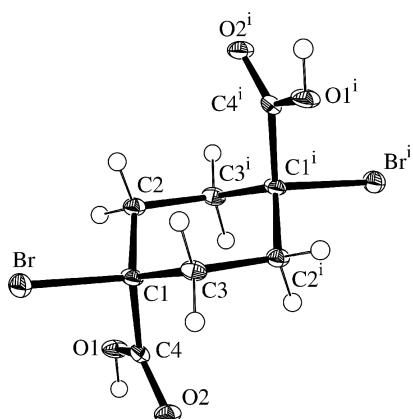
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Comment

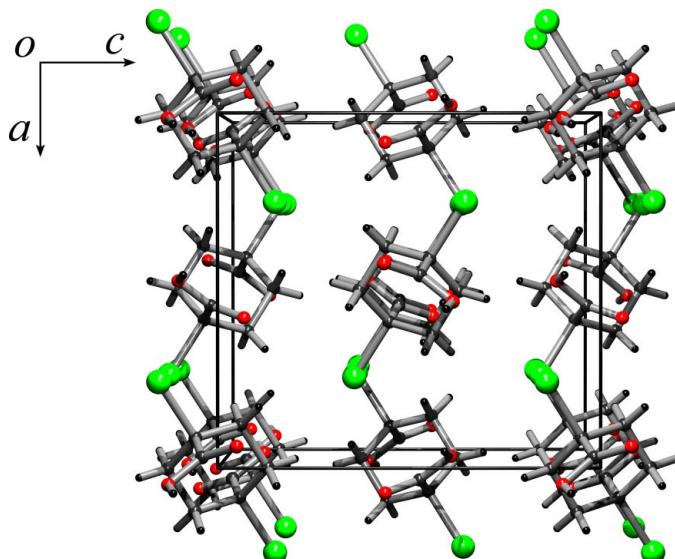
This paper is part of a study of the molecular conformation and crystal polarity of *trans*-1,4-di- and tetrasubstituted symmetrical cyclohexanes (Echeverría *et al.*, 2000, 1995*a,b*).



The intramolecular bond lengths and the endocyclic bond angles that describe the molecular conformation of the title compound, (I), are given in Table 1. The mean C–C ring bond length and the mean endocyclic angle are $1.532(3)\text{ \AA}$ and $110.7(2)^\circ$, respectively; the latter value is close to the value for the ideal cyclohexane ring, C–C–C 111.1° (Bucourt & Hainaut, 1965) and to the observed mean value in the parent compound *trans*-1,4-cyclohexanedicarboxylic acid, (II) [$111.4(4)^\circ$; Dunitz & Strickler, 1966; Von Luger *et al.*, 1972]. The cyclohexane ring, as described by the puckering parameters (Cremer & Pople, 1975), $Q_T = 0.585(2)\text{ \AA}$, $\theta = 0.0(1)^\circ$ and φ undefined, is not distorted. The torsion angle C1–C2–C3ⁱ–C1ⁱ in (I), $56.2(2)^\circ$, can be compared with $53.4(3)^\circ$ in (II), $54.8(8)^\circ$ in *trans*-1,4-dibromocyclohexane-1,4-dicarbonitrile (Echeverría *et al.*, 1995*a*), and $57.1(3)^\circ$ in *trans*-cyclohexane-1,4-dicarbonitrile (Echeverría *et al.*, 1995*b*). In contrast to observations in other studies, where flatter chairs were found as the size of substituents was increased (Juaristi, 1995; Echeverría *et al.*, 2000), the molecule of (I) exhibits a more pronounced chair conformation. The exocyclic angle involving the substituents (C4–C1–Br1) is $102.2(1)^\circ$. This departs from the experimental $H_{\text{eq}}-\text{C}-H_{\text{ax}}$ value, 106.6° , obtained from isotopomers of cyclohexane selectively substituted with deuterium and ^{13}C , using pulsed microwave Fourier transform spectroscopy (Dommen *et al.*, 1990). The $R_{\text{eq}}-\text{C}-R_{\text{ax}}$ angle in (I) is also smaller than the values obtained in other heavily substituted cyclohexanes that include halogens and/or carboxymethyl groups, *e.g.* $105.7(1)^\circ$ in *trans*-dimethyl

**Figure 1**

A view of (I), with displacement ellipsoids drawn at the 50% probability level. Symmetry code: (i) $1 - x, -y, 1 - z$.

**Figure 2**

A view, along b , of the unit-cell contents for (I).

1,4-bis(difluoromethyl)cyclohexane-1,4-dicarboxylate (Swenson *et al.*, 1996) and 106.7 (2) $^{\circ}$ in 1H a :2He:4Ha:5He-octa-fluorocyclohexane (Goodhand & Hamor, 1978). In spite of the equatorial position of Br, the C–Br bond length, 1.988 (2) Å, is longer than the C–Br bond length, 1.834 Å, observed in *trans*-1,4-dibromocyclohexane (Hassel & Hadler Vihovde, 1953). This is consistent with the elongation predicted by HF calculations in overcrowded, perhalogenated cyclohexanes (Ciolowski *et al.*, 1995).

The three-dimensional structure consists of infinite chains extending in the **b** direction. These chains result from hydrogen bonds, in a cyclic motif, formed by the carboxylic acid groups (Table 2). The chains are linked by weak C–H \cdots O bonds (Steiner, 2002) and weak C–H \cdots Br contacts.

Experimental

The title compound was synthesized according to Barón *et al.* (1975) and crystallized by slow evaporation from acetone.

Crystal data



$M_r = 329.98$

Orthorhombic, $Pbca$

$a = 10.5125 (8)$ Å

$b = 8.3836 (5)$ Å

$c = 11.2817 (9)$ Å

$V = 994.29 (12)$ Å³

$Z = 4$

$D_x = 2.204$ Mg m⁻³

Mo $K\alpha$ radiation

Cell parameters from 914 reflections

$\theta = 11.5\text{--}26.3^{\circ}$

$\mu = 8.14$ mm⁻¹

$T = 110 (2)$ K

Prism, colourless

$0.20 \times 0.13 \times 0.10$ mm

Data collection

Bruker SMART 1K CCD diffractometer

ω scans

Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.244$, $T_{\max} = 0.443$

10 606 measured reflections

1224 independent reflections
975 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$

$\theta_{\max} = 28.2^{\circ}$

$h = -13 \rightarrow 13$

$k = -11 \rightarrow 11$

$l = -13 \rightarrow 15$

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.027$

$wR(F^2) = 0.077$

$S = 1.06$

1224 reflections

84 parameters

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0491P)^2 + 0.3512P]$$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 1.04$ e Å⁻³

$\Delta\rho_{\min} = -0.56$ e Å⁻³

Table 1
Selected geometric parameters (Å, °).

Br1–C1	1.988 (2)	C3–C1	1.525 (3)
O1–C4	1.304 (3)	C3–C2 ⁱ	1.545 (3)
O1–H1	0.82	C2–C1	1.525 (3)
O2–C4	1.235 (3)	C1–C4	1.536 (3)
C1–C3–C2 ⁱ	109.97 (18)	C2–C1–Br1	107.70 (16)
C1–C2–C3 ⁱ	110.57 (18)	C4–C1–Br1	102.23 (13)
C3–C1–C2	111.75 (18)	O2–C4–O1	125.0 (2)
C3–C1–C4	111.99 (18)	O2–C4–C1	120.60 (19)
C2–C1–C4	113.69 (18)	O1–C4–C1	114.39 (19)
C3–C1–Br1	108.88 (15)		
C1–C2–C3 ⁱ –C1 ⁱ	−56.1 (2)	C3–C1–C4–O2	25.0 (3)
C2 ⁱ –C3–C1–C2	−56.8 (2)	C2–C1–C4–O2	152.8 (2)
C2 ⁱ –C3–C1–C4	72.1 (2)	Br1–C1–C4–O2	−91.4 (2)
C2 ⁱ –C3–C1–Br1	−175.6 (1)	C3–C1–C4–O1	−156.2 (2)
C3 ⁱ –C2–C1–C3	57.1 (2)	C2–C1–C4–O1	−28.4 (3)
C3 ⁱ –C2–C1–C4	−70.8 (2)	Br1–C1–C4–O1	87.4 (2)
C3 ⁱ –C2–C1–Br1	176.7 (1)		

Symmetry code: (i) $1 - x, -y, 1 - z$.

Table 2
Hydrogen-bonding geometry (Å, °).

D–H \cdots A	D–H	H \cdots A	D \cdots A	D–H \cdots A
O1–H1 \cdots O2 ⁱⁱ	0.82	1.82	2.639 (2)	176
C2–H21 \cdots O1 ⁱⁱⁱ	0.97 (3)	2.73 (2)	3.274 (3)	116 (2)
C2–H22 \cdots O2 ^{iv}	0.96 (2)	2.93 (2)	3.719 (3)	140 (2)
C3–H31 \cdots O2 ^v	1.01 (3)	2.71 (3)	3.431 (3)	129 (2)
C3–H32 \cdots O1 ^{vi}	0.96 (3)	3.18 (3)	3.889 (3)	132 (2)
C2–H21 \cdots Br1 ^v	0.96 (3)	3.17 (3)	4.081 (2)	158 (2)
C2–H22 \cdots Br1 ^{vii}	0.96 (2)	3.03 (2)	3.813 (2)	139 (2)
C3–H31 \cdots Br1 ^v	1.01 (3)	3.22 (2)	4.137 (2)	152 (2)

Symmetry codes: (ii) $-x, -y, -z$; (iii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$; (iv) $x, \frac{1}{2} - y, z - \frac{1}{2}$; (v) $\frac{3}{2} - x, y - \frac{1}{2}, z$; (vi) $\frac{1}{2} + x, \frac{1}{2} - y, 1 - z$; (vii) $x - \frac{1}{2}, y, \frac{1}{2} - z$.

H atoms bound to C atom were located in difference Fourier syntheses and were refined freely in isotropic mode [C—H distances are in the range 0.96 (2)–1.01 (3) Å]. The H atom of the hydroxyl group was stereochemically positioned and allowed to ride on the bound O atom. The O—H bond was also allowed to rotate about the C—O bond. The largest difference electron-density peak is 0.94 Å from Br1.

Data collection: *SMART-NT* (Bruker, 1998); cell refinement: *SMART-NT*; data reduction: *SAINT-NT* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 1990), *PARST* (Nardelli, 1983, 1995) and *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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