Acta Crystallographica Section C Crystal Structure Communications

ISSN 0108-2701

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Electronic paper

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Redetermination of *cis***-4-cyclohexene**-1,2-dicarboxylic acid at 173 K

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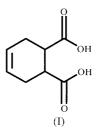
Received 22 May 2000 Accepted 20 June 2000

Data validation number: IUC0000164

The structure of the title compound, $C_8H_{10}O_4$, previously determined by Küppers & Kim [Acta Cryst. (1993), C49, 1218-1220] has been redetermined at 173 K. The cyclohexene ring exhibits a half-chair conformation. The molecular geometry and the crystal packing, which is stabilized by two hydrogen bonds, agree well with the room-temperature structure determination.

Comment

The crystal structure of the title compound, (I), was determined in order to establish unambiguously the nature of a reaction product. The molecular geometry and the crystal structure of the title compound agree well with the structure determined at room temperature (Küppers & Kim, 1993). A least-squares fit of all non-H atoms shows an r.m.s. deviation of 0.014 Å. The cyclohexene ring exhibits a half-chair conformation, with atoms C3, C4, C5 and C6 in a common plane ($\sigma = 0.003$ Å), and C1 0.326 (5) Å above and C2 0.400 (5) Å below that plane. The cell axes are a little shorter at 173 K, and the anisotropic displacement parameters are, as expected, smaller. However, there are no significant differences between the structures at different temperatures. The crystal packing is stabilized by two hydrogen bonds (Table 1).



Experimental

The title compound was synthesized according to the method of Hünig et al. (1979) in two steps. A solution of freshly sublimed maleic anhydride (19.6 g, 200 mmol) in dried benzene (50 ml) was heated to approximately 350 K. Butadiene was then passed into the solution for 30 min with continuous stirring. The reaction product, 4-cyclohexene1,2-dicarboxylic acid anhydride, was washed with petroleum ether and recrystallized from diethyl ether. The second step comprised the hydrolysis of the anhydride. 4-Cyclohexene-1,2-dicarboxylic acid anhydride (1 g) was dissolved in water (10 ml) and refluxed under a take-off condenser for 2 h. When the solution was cooled down to room temperature the product precipitated. Recrystallization from ethyl acetate yielded crystals suitable for an X-ray structure analysis.

Crystal data

 $C_8H_{10}O_4$ Z = 2 $M_r = 170.16$ $D_x = 1.400 \text{ Mg m}^{-3}$ Triclinic, P1 Mo $K\alpha$ radiation a = 6.308 (4) ÅCell parameters from 2610 b = 6.892 (4) Åreflections c = 10.875(7) Å $\theta = 0-25^{\circ}$ $\mu = 0.113 \text{ mm}^{-1}$ $\alpha = 74.57 (1)^{\circ}$ $\beta = 73.78 \ (1)^{\circ}$ T = 173 KBlock, colourless $\nu = 64.41 \ (1)^{\circ}$ $V = 403.6 (4) \text{ Å}^3$ $0.25\,\times\,0.15\,\times\,0.15$ mm Data collection

> $R_{\rm int} = 0.043$ $\theta_{\text{max}} = 26.37^{\circ}$ $h = -7 \rightarrow 7$

 $k = -8 \rightarrow 8$

 $l = -13 \rightarrow 13$

503 standard reflections

frequency: 1100 min

intensity decay: none

Siemens CCD three-circle diffractometer ω scans Absorption correction: empirical (SADABS; Sheldrick, 1996) $T_{\min} = 0.972, \ T_{\max} = 0.983$ 4809 measured reflections 1645 independent reflections

1151 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0414P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.052$	+ 0.3350P]
$wR(F^2) = 0.127$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.078	$(\Delta/\sigma)_{\rm max} < 0.001$
1645 reflections	$\Delta \rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3}$
112 parameters	$\Delta \rho_{\rm min} = -0.23 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
	Extinction coefficient: 0.068 (11)

Table 1

H	yc	lrogen-	bonding	g geome	try	(A, °)).
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
012-H12···011 ⁱ	0.84	1.83	2.667 (3)	172
$O22-H22\cdots O21^{ii}$	0.84	1.81	2.645 (3)	176
Commentation and and (i) 1		(;;) 1 1	. 1 .	

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

All H atoms were located by difference Fourier synthesis and refined with $U(H) = 1.2U_{eq}(C)$ or $U(H) = 1.2U_{eq}(O)$ using a riding model with Csp^2 -H = 0.95, C-H(secondary) = 0.99, C-H(tertiary) = 1.00 and O-H = 0.84 Å. The OH groups were allowed to rotate about the C-O axis.

Data collection: SMART (Siemens, 1995); cell refinement: SMART; data reduction: SAINT (Siemens, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997).

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