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## Structure Reports

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## Peter G. Jones, ${ }^{\text {a }}$ *

Peter Bubenitschek, ${ }^{\text {b }}$
Henning Hopf ${ }^{\text {b }}$ and Bernhard Witulski ${ }^{\text {b }}$
${ }^{\text {a }}$ Institut für Anorganische und Analytische Chemie, Technische Universität Braunschweig, Postfach 3329, 38023 Braunschweig, Germany, and ${ }^{\mathbf{b}}$ Institut für Organische Chemie, Technische Universität Braunschweig, Postfach 3329, 38023 Braunschweig, Germany

Correspondence e-mail: p.jones@tu-bs.de

## Key indicators

Single-crystal X-ray study
$T=178 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.046$
$w R$ factor $=0.133$
Data-to-parameter ratio $=16.6$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## [2.2](2,5)Furanoparacyclophane-12,13dicarbaldehyde

In the title compound, $\mathrm{C}_{16} \mathrm{H}_{14} \mathrm{O}_{3}$, the furan ring makes an angle of $22.0(1)^{\circ}$ with the plane of the four non-bridgehead C atoms of the six-membered ring. The molecules are linked by a short $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interaction to form helical chains parallel to the $b$ axis.

## Comment

In our studies of the addition of highly reactive triple-bond dienophiles to strained aromatic compounds, we prepared the dinitrile (1) by the addition of cyanoacetylene to [2.2](2,5)furanoparacyclophane (Hopf \& Witulski, 1995; Witulski, 1992). To utilize this adduct for further transformations, we have reduced the nitrile to the bis-aldehyde, (2), using diisobutyl aluminium hydride (DIBAH) (Witulski, 1992).


The structure of (2) is shown in Fig. 1. The six-membered ring shows the flattened boat form typical of paracyclophanes, with atom C3 lying 0.160 (3) $\AA$ and C6 0.170 (3) $\AA$ out of the plane of the other four atoms. The angle between the plane of these four atoms and the pane of the furan ring (r.m.s. deviation $0.02 \AA$ ) is $22.0(1)^{\circ}$.

The packing is determined by a short $\mathrm{C} 1-\mathrm{H} 1 b \cdots \mathrm{O} 3$ interaction (Table 1), which links adjacent molecules related by the $2_{1}$ screw axis, to form helical chains with an overall direction parallel to the $b$ axis (Fig. 2). There are two such chains per unit cell.

## Experimental

The title compound was prepared according to Witulski (1992) and was recrystallized from chloroform/pentane.

Crystal data

$$
\begin{aligned}
& \mathrm{C}_{16} \mathrm{H}_{14} \mathrm{O}_{3} \\
& M_{r}=254.27 \\
& \text { Monoclinic, } P 2_{\mathrm{A}} / c \\
& a=13.813(3) \AA \\
& b=8.909(3) \AA \\
& c=10.658(3) \AA \\
& \beta=107.44(2)^{\circ}{ }^{\circ} \AA^{3} \\
& V=1251.3(6) \AA^{3} \\
& Z=4
\end{aligned}
$$

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Figure 1
The molecule of compound (2) in the crystal. Ellipsoids are drawn at the $30 \%$ probability level and H -atom radii are arbitrary.

## Data collection

Nicolet $R 3$ diffractometer $\omega$ scans
Absorption correction: none 3018 measured reflections 2857 independent reflections 1807 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.020$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.046$
$w R\left(F^{2}\right)=0.133$
$S=1.02$
2857 reflections
172 parameters
H -atom parameters constrained

$$
\begin{aligned}
& \theta_{\max }=27.6^{\circ} \\
& h=-17 \rightarrow 17 \\
& k=-1 \rightarrow 11 \\
& l=0 \rightarrow 13 \\
& 3 \text { standard reflections } \\
& \quad \text { every } 147 \text { reflections } \\
& \quad \text { intensity decay: none }
\end{aligned}
$$

Table 1
Hydrogen-bonding geometry $\left(\AA^{\circ},^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 1-\mathrm{H} 1 B \cdots \mathrm{OB}^{\mathrm{i}}$ | 0.99 | 2.45 | $3.393(3)$ | 160 |

Symmetry code: (i) $1-x, \frac{1}{2}+y, \frac{3}{2}-z$.


Figure 2
Packing diagram of compound (2). Hydrogen bonds are indicated by dashed lines. H atoms not involved in hydrogen bonding have been omitted. The view direction is perpendicular to (101).

H atoms were included using a riding model, with fixed $\mathrm{C}-\mathrm{H}$ bond lengths ( $s p^{2} \mathrm{C}-\mathrm{H}=0.95 \AA$ and $\mathrm{CH}_{2}=0.98 \AA$ ); $U_{\text {iso }}(\mathrm{H})$ values were fixed at 1.2 times the $U_{\text {eq }}$ values of the parent atom.

Data collection: P3 (Nicolet, 1987); cell refinement: P3; data reduction: XDISK (Nicolet, 1987); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP (Siemens, 1994); software used to prepare material for publication: SHELXL97.

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