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1,1'-Bicyclohexyl-1,1'-diyl 2,2'-bipyridine-3,3'-dicarboxylate

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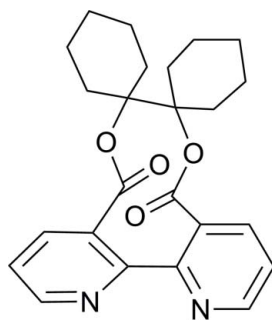
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.001$ Å; R factor = 0.043; wR factor = 0.119; data-to-parameter ratio = 26.3.

The title compound, $\text{C}_{24}\text{H}_{26}\text{N}_2\text{O}_4$, lies about a crystallographic twofold rotation axis. The cyclohexane rings adopts a chair conformation. The two pyridine rings form a dihedral angle of 41.02 (4)°. In the crystal, molecules are linked *via* $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds into a layer parallel to the bc plane.

Related literature

For the background to this study, see the first paper in this series: Fun, Quah, Wu & Zhang (2012). For a related structure, see: Fun, Quah & Wu (2012). For the stability of the temperature controller used in the the data collection, see: Cosier & Glazer (1986). For standard bond-length data, see: Allen *et al.* (1987). For ring conformations, see: Cremer & Pople (1975). For the preparation, see: Wu *et al.* (2012).



Experimental

Crystal data

 $\text{C}_{24}\text{H}_{26}\text{N}_2\text{O}_4$
 $M_r = 406.47$

 Monoclinic, $C2/c$
 $a = 16.7647$ (3) Å
 $b = 10.2618$ (2) Å
 $c = 11.5755$ (2) Å
 $\beta = 99.810$ (1)°
 $V = 1962.28$ (6) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 100$ K
 $0.53 \times 0.24 \times 0.12$ mm

Data collection

 Bruker SMART APEXII CCD
 area-detector diffractometer
 Absorption correction: multi-scan
 (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.952$, $T_{\max} = 0.989$

 11734 measured reflections
 3578 independent reflections
 2972 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.119$
 $S = 1.04$
 3578 reflections

 136 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.50$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.27$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C3}-\text{H3A}\cdots\text{O2}^i$	0.95	2.60	3.5319 (12)	168
$\text{C12}-\text{H12A}\cdots\text{N1}^{ii}$	0.99	2.54	3.4641 (12)	156

 Symmetry codes: (i) $x, -y + 1, z + \frac{1}{2}$; (ii) $x, y + 1, z$.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS5124).

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§ Thomson Reuters ResearcherID: A-5525-2009.

supplementary materials

Acta Cryst. (2012). E68, o1629 [doi:10.1107/S1600536812018508]

1,1'-Bicyclohexyl-1,1'-diyl 2,2'-bipyridine-3,3'-dicarboxylate**Hoong-Kun Fun, Ming Yeng Lim, Ching Kheng Quah and Dongdong Wu****Comment**

The title compound is a ten-membered bislactone with biaryl moiety fused. Easy preparation of this compound could be achieved through a concise photochemical method (Wu *et al.*, 2012). The title compound, Fig. 1, lies about a crystallographic twofold axis generated by the symmetry code $-x, y, -z + 1/2$. The cyclohexane ring (C7–C12) adopts a chair conformation with puckering parameters (Cremer & Pople, 1975) $Q = 0.5773$ (10) Å, $\Theta = 177.88$ (10)° and $\varphi = 256$ (3)°. The two pyridine rings (N1/C1–C5 & N1A/C1A–C5A) are essentially planar [maximum deviation of 0.018 (1) Å at atom C4/C4A] and form a dihedral angle of 41.02 (4)°. Bond lengths (Allen *et al.*, 1987) and angles are within normal ranges and are comparable to related structures (Fun, Quah, Wu & Zhang, 2012; Fun, Quah & Wu, 2012). In the crystal (Fig. 2), molecules are linked *via* intermolecular C3—H3A...O2 and C12—H12A...N1 hydrogen bonds (Table 1) into layers parallel to the (100) plane.

Experimental

The title compound was the product from the photo-oxidation between 2,3-dispirohexyl-2,3-dihydro-[1,4]dioxino[2,3-f][1,10]phenanthroline and oxygen. The compound was purified by flash column chromatography with ethyl acetate/petroleum ether (1:10) as eluents. X-ray quality crystals of the title compound (*m.p.* 180–183 °C), were obtained from slow evaporation of an acetone and petroleum ether solution (1:10).

Refinement

All H atoms were positioned geometrically and refined using a riding model with C—H = 0.95 or 0.99 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

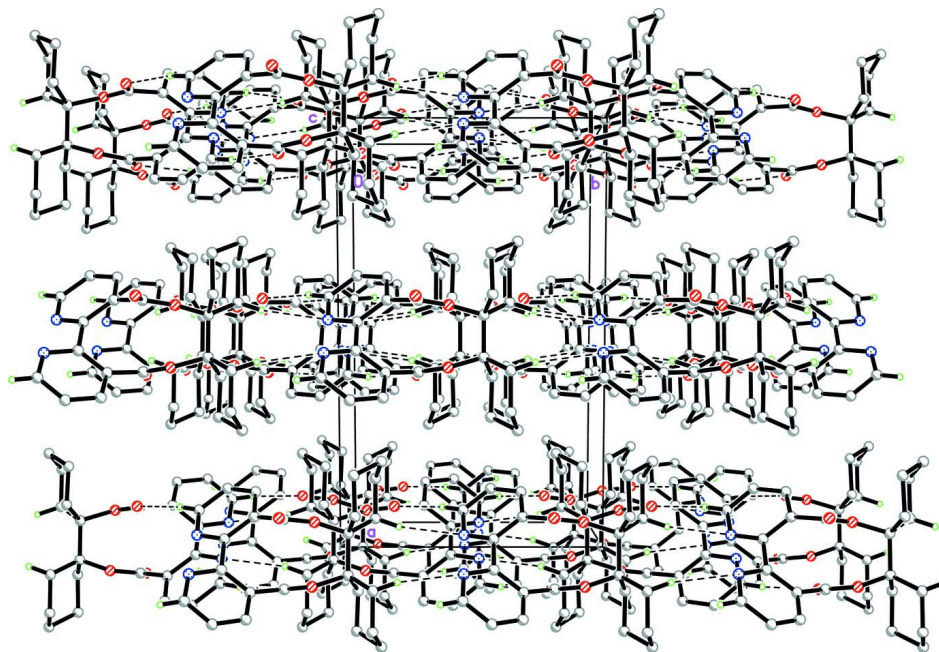
Computing details

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT* (Bruker, 2009); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009).



Figure 1

The molecular structure of the title compound, showing 50% probability displacement ellipsoids for non-H atoms. Atoms with suffix A have been generated by the symmetry code $-x, y, -z + 1/2$.

**Figure 2**

A packing diagram of the title compound, viewed along the c axis. H atoms not involved in hydrogen bonds (dashed lines) have been omitted for clarity.

1,1'-Bicyclohexyl-1,1'-diyl 2,2'-bipyridine-3,3'-dicarboxylate

Crystal data

$C_{24}H_{26}N_2O_4$

$M_r = 406.47$

Monoclinic, $C2/c$

Hall symbol: $-C 2yc$

$a = 16.7647 (3) \text{ \AA}$

$b = 10.2618 (2) \text{ \AA}$

$c = 11.5755 (2) \text{ \AA}$

$\beta = 99.810 (1)^\circ$

$V = 1962.28 (6) \text{ \AA}^3$

$Z = 4$

$F(000) = 864$

$D_x = 1.376 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 4821 reflections

$\theta = 2.3\text{--}32.4^\circ$

$\mu = 0.09 \text{ mm}^{-1}$

$T = 100 \text{ K}$

Block, colourless

$0.53 \times 0.24 \times 0.12 \text{ mm}$

Data collection

Bruker SMART APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2009)

$T_{\min} = 0.952$, $T_{\max} = 0.989$

11734 measured reflections

3578 independent reflections

2972 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\max} = 32.7^\circ$, $\theta_{\min} = 2.3^\circ$

$h = -24 \rightarrow 25$

$k = -15 \rightarrow 15$

$l = -14 \rightarrow 17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.119$

$S = 1.03$

3578 reflections

136 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0602P)^2 + 1.2358P]$

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

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

$\Delta\rho_{\max} = 0.50 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F , and R-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.07035 (4)	0.91569 (6)	0.32195 (5)	0.01138 (14)
O2	0.08919 (4)	0.79994 (7)	0.15933 (6)	0.01480 (15)
N1	0.03750 (5)	0.47163 (8)	0.36731 (7)	0.01641 (17)
C1	0.13527 (6)	0.68507 (10)	0.44475 (8)	0.01575 (18)
H1A	0.1678	0.7583	0.4717	0.019*
C2	0.13993 (6)	0.57190 (10)	0.51086 (8)	0.01792 (19)
H2A	0.1757	0.5658	0.5837	0.022*
C3	0.09090 (6)	0.46750 (10)	0.46771 (8)	0.01813 (19)
H3A	0.0955	0.3889	0.5117	0.022*
C4	0.03211 (5)	0.58187 (9)	0.30384 (8)	0.01315 (17)
C5	0.08212 (5)	0.68998 (9)	0.33802 (8)	0.01229 (16)
C6	0.08110 (5)	0.80731 (9)	0.26097 (8)	0.01163 (16)
C7	0.04781 (5)	1.04426 (9)	0.26757 (7)	0.01063 (16)
C8	0.09239 (5)	1.07527 (9)	0.16519 (8)	0.01365 (17)
H8A	0.0760	1.0116	0.1013	0.016*
H8B	0.0764	1.1631	0.1343	0.016*
C9	0.18467 (6)	1.07060 (10)	0.20256 (9)	0.01769 (19)
H9A	0.2013	0.9814	0.2289	0.021*
H9B	0.2110	1.0921	0.1346	0.021*
C10	0.21245 (6)	1.16711 (11)	0.30191 (9)	0.0207 (2)
H10A	0.2716	1.1591	0.3280	0.025*
H10B	0.2007	1.2572	0.2734	0.025*
C11	0.16867 (6)	1.13963 (10)	0.40464 (8)	0.01758 (19)
H11A	0.1842	1.2065	0.4660	0.021*

H11B	0.1860	1.0537	0.4388	0.021*
C12	0.07650 (5)	1.14012 (9)	0.36753 (8)	0.01346 (17)
H12A	0.0585	1.2291	0.3422	0.016*
H12B	0.0510	1.1171	0.4359	0.016*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0130 (3)	0.0087 (3)	0.0120 (3)	0.0012 (2)	0.0009 (2)	0.0006 (2)
O2	0.0162 (3)	0.0141 (3)	0.0145 (3)	0.0007 (2)	0.0040 (2)	-0.0013 (2)
N1	0.0203 (4)	0.0111 (4)	0.0178 (3)	0.0014 (3)	0.0030 (3)	0.0021 (3)
C1	0.0156 (4)	0.0141 (4)	0.0164 (4)	0.0025 (3)	-0.0004 (3)	-0.0002 (3)
C2	0.0196 (4)	0.0177 (5)	0.0154 (4)	0.0054 (4)	-0.0002 (3)	0.0016 (3)
C3	0.0223 (4)	0.0140 (4)	0.0180 (4)	0.0052 (4)	0.0032 (3)	0.0039 (3)
C4	0.0156 (4)	0.0099 (4)	0.0138 (4)	0.0012 (3)	0.0022 (3)	0.0001 (3)
C5	0.0132 (3)	0.0097 (4)	0.0139 (3)	0.0020 (3)	0.0020 (3)	0.0005 (3)
C6	0.0094 (3)	0.0098 (4)	0.0151 (4)	-0.0001 (3)	0.0003 (3)	-0.0002 (3)
C7	0.0118 (3)	0.0076 (3)	0.0119 (3)	-0.0002 (3)	0.0003 (3)	0.0009 (3)
C8	0.0132 (4)	0.0137 (4)	0.0140 (4)	-0.0011 (3)	0.0023 (3)	0.0021 (3)
C9	0.0134 (4)	0.0198 (5)	0.0203 (4)	-0.0020 (3)	0.0040 (3)	0.0032 (4)
C10	0.0143 (4)	0.0193 (5)	0.0271 (5)	-0.0050 (4)	-0.0001 (3)	0.0018 (4)
C11	0.0141 (4)	0.0167 (4)	0.0200 (4)	-0.0014 (3)	-0.0028 (3)	-0.0025 (4)
C12	0.0142 (4)	0.0101 (4)	0.0148 (4)	-0.0005 (3)	-0.0011 (3)	-0.0023 (3)

Geometric parameters (\AA , $^\circ$)

O1—C6	1.3458 (11)	C7—C7 ⁱ	1.5854 (17)
O1—C7	1.4828 (11)	C8—C9	1.5344 (13)
O2—C6	1.2094 (11)	C8—H8A	0.9900
N1—C3	1.3414 (12)	C8—H8B	0.9900
N1—C4	1.3435 (12)	C9—C10	1.5290 (15)
C1—C2	1.3856 (14)	C9—H9A	0.9900
C1—C5	1.3956 (12)	C9—H9B	0.9900
C1—H1A	0.9500	C10—C11	1.5264 (15)
C2—C3	1.3903 (15)	C10—H10A	0.9900
C2—H2A	0.9500	C10—H10B	0.9900
C3—H3A	0.9500	C11—C12	1.5317 (13)
C4—C5	1.4061 (13)	C11—H11A	0.9900
C4—C4 ⁱ	1.5018 (18)	C11—H11B	0.9900
C5—C6	1.4968 (12)	C12—H12A	0.9900
C7—C12	1.5324 (12)	C12—H12B	0.9900
C7—C8	1.5382 (12)		
C6—O1—C7	124.05 (6)	C7—C8—H8A	109.2
C3—N1—C4	118.24 (9)	C9—C8—H8B	109.2
C2—C1—C5	119.12 (9)	C7—C8—H8B	109.2
C2—C1—H1A	120.4	H8A—C8—H8B	107.9
C5—C1—H1A	120.4	C10—C9—C8	110.80 (8)
C1—C2—C3	118.24 (9)	C10—C9—H9A	109.5
C1—C2—H2A	120.9	C8—C9—H9A	109.5

C3—C2—H2A	120.9	C10—C9—H9B	109.5
N1—C3—C2	123.60 (9)	C8—C9—H9B	109.5
N1—C3—H3A	118.2	H9A—C9—H9B	108.1
C2—C3—H3A	118.2	C11—C10—C9	109.94 (8)
N1—C4—C5	121.94 (8)	C11—C10—H10A	109.7
N1—C4—C4 ⁱ	115.20 (6)	C9—C10—H10A	109.7
C5—C4—C4 ⁱ	122.84 (6)	C11—C10—H10B	109.7
C1—C5—C4	118.74 (8)	C9—C10—H10B	109.7
C1—C5—C6	119.82 (8)	H10A—C10—H10B	108.2
C4—C5—C6	121.42 (8)	C10—C11—C12	112.15 (8)
O2—C6—O1	127.53 (8)	C10—C11—H11A	109.2
O2—C6—C5	122.52 (8)	C12—C11—H11A	109.2
O1—C6—C5	109.95 (7)	C10—C11—H11B	109.2
O1—C7—C12	103.07 (6)	C12—C11—H11B	109.2
O1—C7—C8	112.90 (7)	H11A—C11—H11B	107.9
C12—C7—C8	108.53 (7)	C11—C12—C7	112.42 (8)
O1—C7—C7 ⁱ	106.40 (5)	C11—C12—H12A	109.1
C12—C7—C7 ⁱ	111.54 (7)	C7—C12—H12A	109.1
C8—C7—C7 ⁱ	113.91 (8)	C11—C12—H12B	109.1
C9—C8—C7	112.04 (7)	C7—C12—H12B	109.1
C9—C8—H8A	109.2	H12A—C12—H12B	107.9
C5—C1—C2—C3	-0.15 (14)	C1—C5—C6—O1	-53.79 (11)
C4—N1—C3—C2	1.26 (15)	C4—C5—C6—O1	128.11 (9)
C1—C2—C3—N1	-2.13 (16)	C6—O1—C7—C12	-157.78 (7)
C3—N1—C4—C5	1.90 (14)	C6—O1—C7—C8	-40.90 (10)
C3—N1—C4—C4 ⁱ	-176.74 (10)	C6—O1—C7—C7 ⁱ	84.75 (10)
C2—C1—C5—C4	3.07 (14)	O1—C7—C8—C9	-56.99 (10)
C2—C1—C5—C6	-175.08 (8)	C12—C7—C8—C9	56.61 (10)
N1—C4—C5—C1	-4.06 (14)	C7 ⁱ —C7—C8—C9	-178.48 (6)
C4 ⁱ —C4—C5—C1	174.47 (10)	C7—C8—C9—C10	-58.49 (11)
N1—C4—C5—C6	174.05 (8)	C8—C9—C10—C11	56.01 (11)
C4 ⁱ —C4—C5—C6	-7.41 (15)	C9—C10—C11—C12	-54.88 (11)
C7—O1—C6—O2	14.66 (13)	C10—C11—C12—C7	55.67 (11)
C7—O1—C6—C5	-165.46 (7)	O1—C7—C12—C11	65.03 (9)
C1—C5—C6—O2	126.10 (10)	C8—C7—C12—C11	-54.90 (10)
C4—C5—C6—O2	-52.00 (13)	C7 ⁱ —C7—C12—C11	178.81 (7)

Symmetry code: (i) $-x, y, -z+1/2$.

Hydrogen-bond geometry (\AA , $^\circ$)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C3—H3A \cdots O2 ⁱⁱ	0.95	2.60	3.5319 (12)	168
C12—H12A \cdots N1 ⁱⁱⁱ	0.99	2.54	3.4641 (12)	156

Symmetry codes: (ii) $x, -y+1, z+1/2$; (iii) $x, y+1, z$.