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3,4-Bis(4-bromophenyl)-*N*-phenylmaleimideYong Liu,^a Shuai Zheng,^b Jing Zhou,^b Dongmei Gao^b and Zhen-Ting Du^{b*}

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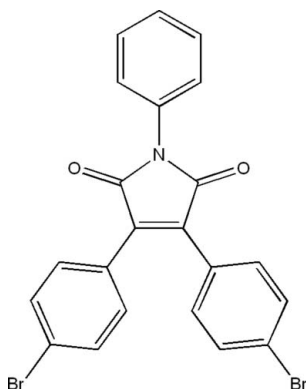
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.011$ Å; R factor = 0.050; wR factor = 0.133; data-to-parameter ratio = 13.9.

In the title molecule, $\text{C}_{22}\text{H}_{13}\text{Br}_2\text{NO}_2$, the three benzene rings are arranged in a propeller-like fashion around the central maleimide ring, making dihedral angles of 48.2 (4), 30.2 (4) and 34.8 (4)° with the maleimide ring. The C—C single-bond lengths connecting benzene groups and maleimide are significantly shorter [C—C = 1.468 (9) and 1.478 (9) Å] than a typical $\text{Csp}^3-\text{Csp}^3$ single bond, indicating partial conjugation between the benzene and the maleimide. A weak nonclassical C—H...O hydrogen bond helps to stabilize the crystal structure.

Related literature

For general background to 3,4-diaryl-substituted maleimide derivatives, see: Fujii *et al.* (2001); Onimura *et al.* (2010); Shorunov *et al.* (2006).



Experimental

Crystal data

$\text{C}_{22}\text{H}_{13}\text{Br}_2\text{NO}_2$
 $M_r = 483.13$
Monoclinic, $P2_1/c$
 $a = 10.844$ (5) Å
 $b = 18.594$ (9) Å
 $c = 9.602$ (5) Å
 $\beta = 102.760$ (6)°

$V = 1888.3$ (16) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 4.31$ mm⁻¹
 $T = 296$ K
 $0.32 \times 0.30 \times 0.24$ mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1998)
 $T_{\min} = 0.339$, $T_{\max} = 0.424$

8594 measured reflections
3382 independent reflections
1427 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.117$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.133$
 $S = 0.94$
3382 reflections

244 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.51$ e Å⁻³
 $\Delta\rho_{\min} = -0.71$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C21—H21...O1 ⁱ	0.93	2.41	3.267 (9)	153

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

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2272).

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supporting information

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3,4-Bis(4-bromophenyl)-*N*-phenylmaleimide

Yong Liu, Shuai Zheng, Jing Zhou, Dongmei Gao and Zhen-Ting Du

S1. Comment

3,4-Diaryl-substituted maleimide is a conjugated unit which has interesting optical and electronic properties. A number of 3,4-diaryl-substituted maleimide derivatives have been designed and synthesized to be used as monomer of some electro-optic polymers (Shorunov *et al.*, 2006; Onimura *et al.*, 2010). In the course of exploring new electro-optic compounds, we obtained an intermediate compound 3,4-bis(4-bromophenyl)-*N*-phenylmaleimide **I**. Here we report the structure and synthesis of title compound.

The molecule holds four rings. The maleimide ring locates the core position, and the other three benzene rings are arranged in a propeller-like fashion around the central maleimide 5-membering ring with the dihedral angles being 48.2 (4)° (C1–C6), 30.2 (4)° (C9–C14) and 34.8 (4)° (C17–C22). The C—C single bond lengths connecting benzene groups and maleimide unit are respectively 1.468 (9) Å (C4–C7) and 1.478 (9) Å (C17–C18), which are obviously shorter than typical Csp^3 — Csp^3 single bond. This means that the bonding between the benzenes and the maleimide is quite conjugated.

The molecules of **I** are crystalized in $P2_1/c$ space group which is different from that of *N*-3,4-triphenylmaleimide (*Pbca*, Fujii *et al.*, 2001). There are no classic hydrogen bonds in this crystal structure. However, the weak intermolecular interaction C21—H21 $\cdots$ O1ⁱ is helpful to the stabilization of the packing (Fig. 2). This intermolecular non-classical H-bond is characterized by the parameters: bond lengths of 0.93 Å (C21—H21), 2.41 Å (H21 $\cdots$ O1ⁱ), 3.267 (9) Å (C21 $\cdots$ O1ⁱ) and angle 153° (C21—H21 $\cdots$ O1ⁱ). Symmetry code: (i) 1-*x*, -1/2+*y*, 1/2-*z*.

S2. Experimental

3,4-Bis(4-bromophenyl)maleic anhydride (0.60 g, 1.47 mmol), 4-methylbenzenesulfonic acid (0.30 g, 3.26 mmol) and H₂SO₄ (2 ml) were dissolved in *N,N*-dimethylformamide (DMF, 1.0 ml) and toluene (20 ml) mixed solvent. After heating the mixture to refluxing, a toluene solution of anilin (0.20 g, 2.17 mmol) was slowly added. After stirring for 2 h, the mixture was cooled to 333 K and poured into 8% Na₂CO₃ solution and further stirred for 10 min. The solution was extracted with toluene and dried over Na₂SO₄. After removing the solvent, the crude product was purified by recrystallization from ethanol, affording the title compound, **I**, (0.51 g, 72%). Then the compound **I** was dissolved in THF, and yellow crystals were formed on slow evaporation at room temperature over one week.

S3. Refinement

All H atoms were placed in geometrically calculated positions and refined using a riding model with C—H = 0.93 Å (for aromatic H) with $U_{iso}(H) = 1.2U_{eq}(C)$.

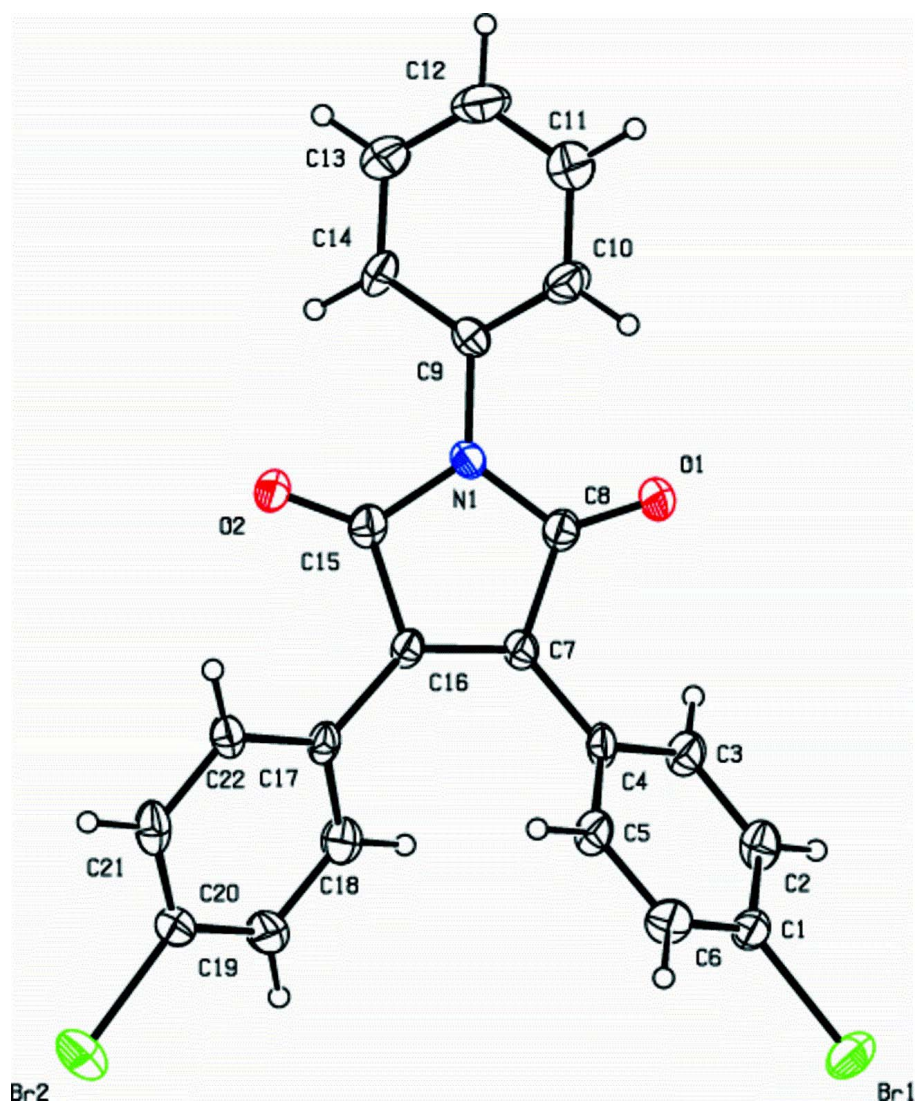


Figure 1

The molecular structure of **I** with the atom numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are presented as a small spheres of arbitrary radius.

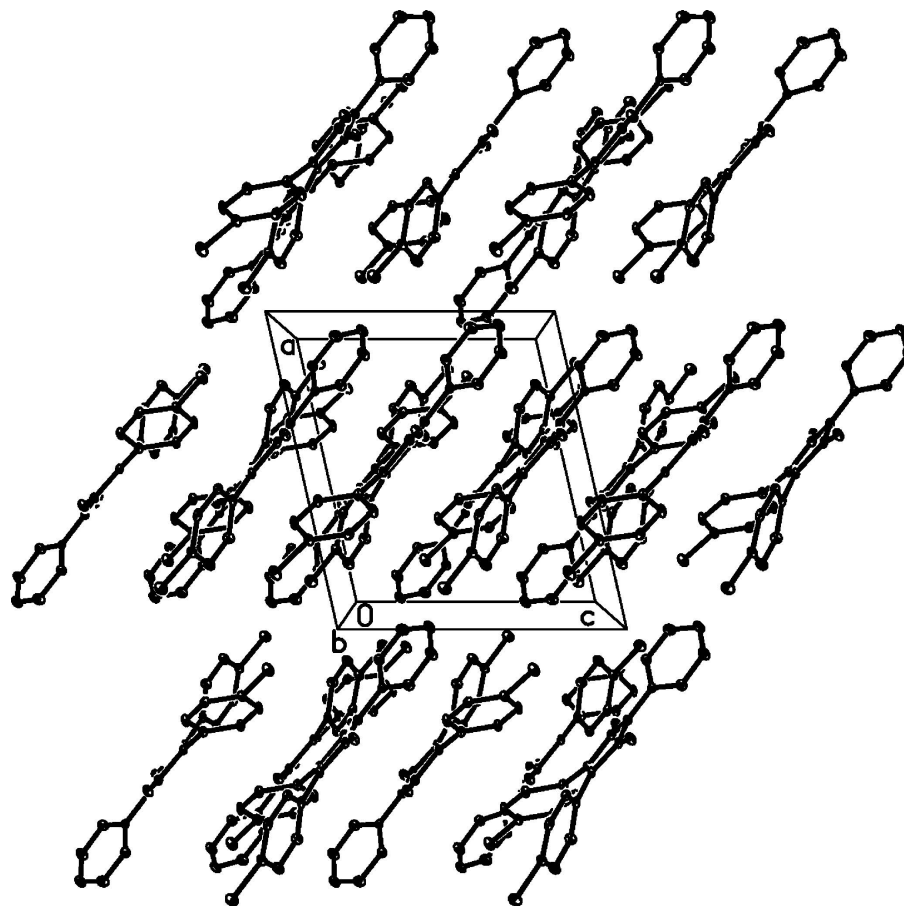


Figure 2

The molecular packing of **I** along *b* axis.

3,4-bis(4-bromophenyl)-1-phenyl-2,5-dihydro-1H-pyrrole-2,5-dione

Crystal data

$\text{C}_{22}\text{H}_{13}\text{Br}_2\text{NO}_2$

$M_r = 483.13$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 10.844 (5) \text{ \AA}$

$b = 18.594 (9) \text{ \AA}$

$c = 9.602 (5) \text{ \AA}$

$\beta = 102.760 (6)^\circ$

$V = 1888.3 (16) \text{ \AA}^3$

$Z = 4$

$F(000) = 952$

$D_x = 1.699 \text{ Mg m}^{-3}$

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

Cell parameters from 1226 reflections

$\theta = 2.2\text{--}20.8^\circ$

$\mu = 4.31 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Block, yellow

$0.32 \times 0.30 \times 0.24 \text{ mm}$

Data collection

Bruker APEXII CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1998)

$T_{\min} = 0.339$, $T_{\max} = 0.424$

8594 measured reflections

3382 independent reflections

1427 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.117$
 $\theta_{\text{max}} = 25.2^\circ$, $\theta_{\text{min}} = 2.2^\circ$
 $h = -12 \rightarrow 12$

$k = -22 \rightarrow 21$
 $l = -9 \rightarrow 11$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.133$
 $S = 0.94$
 3382 reflections
 244 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
 $w = 1/[\sigma^2(F_o^2) + (0.0468P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.51 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.71 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s 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 $F^2 > \sigma(F^2)$ 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 ($\text{\AA}^2$)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.18239 (9)	1.08051 (4)	−0.16015 (9)	0.0659 (4)
Br2	0.12170 (9)	0.59744 (5)	−0.09998 (9)	0.0681 (4)
C1	0.2818 (7)	1.0177 (4)	−0.0272 (8)	0.042 (2)
C2	0.3009 (7)	1.0324 (4)	0.1166 (8)	0.044 (2)
H2	0.2661	1.0735	0.1474	0.052*
C3	0.3718 (7)	0.9859 (4)	0.2152 (7)	0.046 (2)
H3	0.3854	0.9965	0.3121	0.055*
C4	0.4223 (7)	0.9244 (3)	0.1720 (7)	0.037 (2)
C5	0.4019 (7)	0.9103 (4)	0.0241 (7)	0.041 (2)
H5	0.4353	0.8689	−0.0080	0.050*
C6	0.3333 (8)	0.9573 (4)	−0.0718 (7)	0.049 (2)
H6	0.3214	0.9480	−0.1690	0.058*
C7	0.4950 (7)	0.8740 (4)	0.2765 (7)	0.0317 (19)
C8	0.5979 (7)	0.8995 (4)	0.3949 (7)	0.037 (2)
C9	0.7668 (7)	0.8405 (4)	0.5799 (7)	0.0324 (19)
C10	0.8549 (8)	0.8948 (4)	0.5883 (8)	0.049 (2)
H10	0.8449	0.9299	0.5176	0.059*
C11	0.9570 (8)	0.8967 (5)	0.7012 (9)	0.064 (3)
H11	1.0169	0.9329	0.7067	0.077*
C12	0.9715 (9)	0.8455 (5)	0.8061 (8)	0.066 (3)
H12	1.0385	0.8483	0.8852	0.079*
C13	0.8877 (9)	0.7907 (5)	0.7936 (9)	0.067 (3)

H13	0.9013	0.7541	0.8612	0.080*
C14	0.7840 (8)	0.7881 (4)	0.6845 (8)	0.049 (2)
H14	0.7252	0.7513	0.6802	0.059*
C15	0.5930 (7)	0.7768 (4)	0.4054 (7)	0.038 (2)
C16	0.4904 (7)	0.8021 (4)	0.2827 (7)	0.0321 (19)
C17	0.3999 (8)	0.7505 (4)	0.1988 (7)	0.035 (2)
C18	0.2750 (8)	0.7690 (4)	0.1499 (8)	0.050 (2)
H18	0.2470	0.8136	0.1741	0.060*
C19	0.1900 (8)	0.7226 (4)	0.0653 (7)	0.052 (2)
H19	0.1053	0.7352	0.0353	0.063*
C20	0.2331 (8)	0.6571 (4)	0.0260 (7)	0.043 (2)
C21	0.3549 (8)	0.6361 (4)	0.0806 (7)	0.046 (2)
H21	0.3816	0.5905	0.0607	0.055*
C22	0.4388 (7)	0.6829 (4)	0.1659 (7)	0.036 (2)
H22	0.5221	0.6687	0.2014	0.044*
N1	0.6588 (6)	0.8394 (3)	0.4625 (6)	0.0341 (16)
O1	0.6229 (5)	0.9610 (3)	0.4283 (5)	0.0494 (15)
O2	0.6167 (5)	0.7165 (3)	0.4467 (5)	0.0480 (15)

Atomic displacement parameters ($\text{\AA}^2$)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0834 (8)	0.0516 (6)	0.0525 (6)	0.0141 (5)	−0.0068 (5)	0.0128 (4)
Br2	0.0699 (7)	0.0590 (6)	0.0689 (7)	−0.0182 (5)	0.0015 (5)	−0.0198 (5)
C1	0.056 (6)	0.032 (5)	0.034 (5)	0.004 (4)	0.002 (4)	0.000 (4)
C2	0.058 (6)	0.034 (5)	0.043 (5)	0.007 (4)	0.021 (5)	−0.006 (4)
C3	0.073 (7)	0.030 (5)	0.031 (5)	0.008 (5)	0.005 (4)	−0.006 (4)
C4	0.055 (6)	0.018 (4)	0.037 (5)	−0.003 (4)	0.011 (4)	−0.002 (4)
C5	0.064 (6)	0.027 (5)	0.033 (4)	0.012 (4)	0.011 (4)	−0.008 (4)
C6	0.078 (7)	0.048 (5)	0.019 (4)	0.003 (5)	0.008 (4)	−0.007 (4)
C7	0.043 (6)	0.025 (4)	0.027 (4)	0.002 (4)	0.009 (4)	−0.006 (3)
C8	0.047 (6)	0.031 (5)	0.034 (5)	0.000 (4)	0.009 (4)	0.002 (4)
C9	0.039 (6)	0.030 (4)	0.029 (5)	0.000 (4)	0.010 (4)	−0.011 (4)
C10	0.050 (6)	0.054 (6)	0.041 (5)	0.002 (5)	0.004 (5)	0.014 (4)
C11	0.060 (7)	0.055 (6)	0.072 (6)	−0.007 (5)	0.003 (5)	0.008 (5)
C12	0.060 (7)	0.080 (7)	0.047 (6)	0.011 (6)	−0.012 (5)	0.003 (5)
C13	0.072 (8)	0.064 (7)	0.054 (6)	−0.008 (6)	−0.010 (6)	0.021 (5)
C14	0.056 (7)	0.035 (5)	0.052 (6)	0.003 (4)	0.004 (5)	0.019 (4)
C15	0.055 (6)	0.035 (5)	0.025 (5)	−0.003 (4)	0.014 (4)	0.002 (4)
C16	0.051 (6)	0.023 (4)	0.022 (4)	0.004 (4)	0.007 (4)	−0.001 (3)
C17	0.058 (7)	0.024 (5)	0.023 (4)	−0.001 (4)	0.011 (4)	0.003 (3)
C18	0.054 (7)	0.034 (5)	0.059 (6)	0.008 (5)	0.007 (5)	−0.006 (4)
C19	0.058 (7)	0.038 (5)	0.053 (5)	0.003 (5)	−0.004 (5)	−0.010 (4)
C20	0.049 (7)	0.038 (5)	0.038 (5)	−0.005 (4)	0.000 (4)	−0.003 (4)
C21	0.075 (8)	0.029 (5)	0.041 (5)	−0.006 (5)	0.027 (5)	−0.004 (4)
C22	0.047 (6)	0.027 (5)	0.037 (5)	−0.005 (4)	0.012 (4)	−0.003 (4)
N1	0.039 (4)	0.030 (4)	0.031 (4)	−0.008 (3)	0.004 (3)	−0.001 (3)
O1	0.067 (4)	0.028 (3)	0.046 (3)	−0.003 (3)	−0.005 (3)	−0.005 (3)

O2	0.066 (4)	0.028 (3)	0.043 (3)	0.000 (3)	-0.004 (3)	0.004 (2)
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Geometric parameters (Å, °)

Br1—C1	1.883 (7)	C11—C12	1.369 (10)
Br2—C20	1.873 (7)	C11—H11	0.9300
C1—C6	1.365 (9)	C12—C13	1.352 (11)
C1—C2	1.378 (8)	C12—H12	0.9300
C2—C3	1.384 (9)	C13—C14	1.357 (9)
C2—H2	0.9300	C13—H13	0.9300
C3—C4	1.371 (9)	C14—H14	0.9300
C3—H3	0.9300	C15—O2	1.198 (7)
C4—C5	1.414 (8)	C15—N1	1.412 (8)
C4—C7	1.468 (9)	C15—C16	1.506 (9)
C5—C6	1.365 (9)	C16—C17	1.478 (9)
C5—H5	0.9300	C17—C18	1.375 (10)
C6—H6	0.9300	C17—C22	1.385 (9)
C7—C16	1.339 (9)	C18—C19	1.387 (9)
C7—C8	1.484 (9)	C18—H18	0.9300
C8—O1	1.202 (7)	C19—C20	1.386 (9)
C8—N1	1.385 (8)	C19—H19	0.9300
C9—C10	1.380 (9)	C20—C21	1.366 (10)
C9—C14	1.383 (8)	C21—C22	1.388 (9)
C9—N1	1.435 (8)	C21—H21	0.9300
C10—C11	1.369 (10)	C22—H22	0.9300
C10—H10	0.9300		
C6—C1—C2	119.7 (6)	C11—C12—H12	120.3
C6—C1—Br1	120.7 (6)	C12—C13—C14	121.4 (8)
C2—C1—Br1	119.6 (6)	C12—C13—H13	119.3
C1—C2—C3	120.0 (7)	C14—C13—H13	119.3
C1—C2—H2	120.0	C13—C14—C9	119.4 (8)
C3—C2—H2	120.0	C13—C14—H14	120.3
C4—C3—C2	120.9 (7)	C9—C14—H14	120.3
C4—C3—H3	119.6	O2—C15—N1	126.1 (7)
C2—C3—H3	119.6	O2—C15—C16	128.3 (7)
C3—C4—C5	118.3 (6)	N1—C15—C16	105.6 (6)
C3—C4—C7	121.0 (6)	C7—C16—C17	130.6 (7)
C5—C4—C7	120.6 (6)	C7—C16—C15	108.6 (6)
C6—C5—C4	120.0 (7)	C17—C16—C15	120.7 (6)
C6—C5—H5	120.0	C18—C17—C22	118.5 (7)
C4—C5—H5	120.0	C18—C17—C16	120.6 (7)
C5—C6—C1	121.1 (7)	C22—C17—C16	121.0 (7)
C5—C6—H6	119.5	C17—C18—C19	121.4 (7)
C1—C6—H6	119.5	C17—C18—H18	119.3
C16—C7—C4	130.3 (6)	C19—C18—H18	119.3
C16—C7—C8	108.3 (6)	C18—C19—C20	119.0 (8)
C4—C7—C8	121.3 (6)	C18—C19—H19	120.5

O1—C8—N1	125.9 (7)	C20—C19—H19	120.5
O1—C8—C7	126.6 (7)	C21—C20—C19	120.1 (7)
N1—C8—C7	107.5 (6)	C21—C20—Br2	120.7 (6)
C10—C9—C14	119.5 (7)	C19—C20—Br2	119.2 (6)
C10—C9—N1	119.4 (6)	C20—C21—C22	120.0 (7)
C14—C9—N1	121.1 (7)	C20—C21—H21	120.0
C11—C10—C9	119.6 (7)	C22—C21—H21	120.0
C11—C10—H10	120.2	C17—C22—C21	120.6 (7)
C9—C10—H10	120.2	C17—C22—H22	119.7
C10—C11—C12	120.4 (9)	C21—C22—H22	119.7
C10—C11—H11	119.8	C8—N1—C15	109.6 (6)
C12—C11—H11	119.8	C8—N1—C9	125.3 (6)
C13—C12—C11	119.5 (8)	C15—N1—C9	124.9 (6)
C13—C12—H12	120.3		
C6—C1—C2—C3	−0.1 (12)	O2—C15—C16—C7	178.1 (8)
Br1—C1—C2—C3	178.8 (6)	N1—C15—C16—C7	−2.8 (8)
C1—C2—C3—C4	−1.0 (12)	O2—C15—C16—C17	2.3 (12)
C2—C3—C4—C5	1.0 (11)	N1—C15—C16—C17	−178.6 (6)
C2—C3—C4—C7	−178.5 (7)	C7—C16—C17—C18	−32.3 (12)
C3—C4—C5—C6	−0.1 (11)	C15—C16—C17—C18	142.5 (7)
C7—C4—C5—C6	179.5 (7)	C7—C16—C17—C22	146.8 (8)
C4—C5—C6—C1	−1.0 (12)	C15—C16—C17—C22	−38.4 (10)
C2—C1—C6—C5	1.1 (12)	C22—C17—C18—C19	−2.3 (11)
Br1—C1—C6—C5	−177.8 (6)	C16—C17—C18—C19	176.9 (7)
C3—C4—C7—C16	135.2 (9)	C17—C18—C19—C20	−1.9 (12)
C5—C4—C7—C16	−44.3 (12)	C18—C19—C20—C21	5.8 (12)
C3—C4—C7—C8	−48.7 (10)	C18—C19—C20—Br2	−175.2 (6)
C5—C4—C7—C8	131.7 (7)	C19—C20—C21—C22	−5.5 (12)
C16—C7—C8—O1	−174.1 (8)	Br2—C20—C21—C22	175.5 (5)
C4—C7—C8—O1	9.1 (12)	C18—C17—C22—C21	2.7 (10)
C16—C7—C8—N1	4.7 (8)	C16—C17—C22—C21	−176.5 (6)
C4—C7—C8—N1	−172.1 (6)	C20—C21—C22—C17	1.2 (11)
C14—C9—C10—C11	−1.1 (11)	O1—C8—N1—C15	172.3 (7)
N1—C9—C10—C11	179.0 (7)	C7—C8—N1—C15	−6.5 (8)
C9—C10—C11—C12	−0.5 (13)	O1—C8—N1—C9	−3.4 (12)
C10—C11—C12—C13	3.5 (14)	C7—C8—N1—C9	177.8 (6)
C11—C12—C13—C14	−4.8 (15)	O2—C15—N1—C8	−175.1 (7)
C12—C13—C14—C9	3.2 (14)	C16—C15—N1—C8	5.8 (8)
C10—C9—C14—C13	−0.2 (11)	O2—C15—N1—C9	0.6 (12)
N1—C9—C14—C13	179.7 (7)	C16—C15—N1—C9	−178.5 (6)
C4—C7—C16—C17	−9.3 (14)	C10—C9—N1—C8	−33.2 (10)
C8—C7—C16—C17	174.2 (7)	C14—C9—N1—C8	146.9 (7)
C4—C7—C16—C15	175.3 (7)	C10—C9—N1—C15	151.7 (7)
C8—C7—C16—C15	−1.1 (8)	C14—C9—N1—C15	−28.2 (10)

Hydrogen-bond geometry (Å, °)

<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>
C21—H21 $\cdots$ O1 ⁱ	0.93	2.41	3.267 (9)	153

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