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Dimethyl *trans*-3-(4-bromophenyl)-2-methylisoxazolidine-4,5-dicarboxylate

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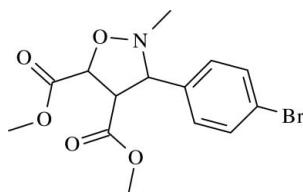
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 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.044; wR factor = 0.101; data-to-parameter ratio = 16.7.

In the title compound, $\text{C}_{14}\text{H}_{16}\text{BrNO}_5$, the isoxazolidine ring adopts an envelope conformation, with the N atom at the flap. In the crystal, intermolecular $\text{C}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds generate $R_3^2(18)$ ring motifs which are fused into a ribbon-like structure extending along the b axis.

Related literature

For general background, see: Confalone & Huie (1988); Torssell (1988); Frederickson (1997); Gothelf & Jorgensen (1998); Chiacchio *et al.* (2003); Padwa *et al.* (1981, 1984); Ochiai *et al.* (1967); Baldwin & Aube (1987); Heaney *et al.* (2001). For hydrogen-bond motifs, see: Bernstein *et al.* (1995); Etter (1990). For ring conformations, see: Cremer & Pople (1975).



Experimental

Crystal data

$\text{C}_{14}\text{H}_{16}\text{BrNO}_5$
 $M_r = 358.19$
 Monoclinic, $P2_1/c$
 $a = 10.9020$ (4) Å
 $b = 8.1780$ (3) Å

$c = 17.8127$ (8) Å
 $\beta = 101.622$ (3)°
 $V = 1555.56$ (11) Å³
 $Z = 4$
 Mo $K\alpha$ radiation

$\mu = 2.67$ mm⁻¹
 $T = 296$ K

0.71 × 0.60 × 0.45 mm

Data collection

Stoe IPDS II diffractometer
 Absorption correction: integration
 (*X-RED32*; Stoe & Cie, 2002)
 $T_{\min} = 0.327$, $T_{\max} = 0.480$

15678 measured reflections
 3232 independent reflections
 2696 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.043$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.101$
 $S = 1.12$
 3232 reflections

193 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.49$ e Å⁻³
 $\Delta\rho_{\min} = -0.93$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C3}-\text{H3}\cdots\text{N1}^i$	0.93	2.56	3.492 (4)	179
$\text{C12}-\text{H12C}\cdots\text{O1}^{ii}$	0.96	2.52	3.434 (5)	158

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

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The authors acknowledge the Faculty of Arts and Sciences, Ondokuz Mayıs University, Turkey, for the use of the Stoe IPDS II diffractometer (purchased under grant F.279 of the University Research Fund).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2876).

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supporting information

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Dimethyl *trans*-3-(4-bromophenyl)-2-methylisoxazolidine-4,5-dicarboxylate**Orhan Büyükgüngör, Serkan Yavuz, Mustafa Odabaşoğlu, Hamdi Özkan, Özgür Pamir and Yılmaz Yıldırım****S1. Comment**

The 1,3-dipolar cycloaddition of nitrones and alkenes is a powerful synthetic device that allows up to three new stereogenic centers to be assembled in a stereospecific manner in a single step (Confalone & Huie, 1988; Torrsell, 1988; Frederickson, 1997; Gothelf & Jorgensen, 1998). Among these N and O containing five-membered heterocycles, isoxazolidines and isoxazoline derivatives have emerged as important candidates and have been shown to display useful anticancer and antiviral properties (Chiacchio *et al.*, 2003).

The syntheses of isoxazolidine derivatives is an important subject in organic chemistry because they are found in the structure of most natural compounds and drugs. In recent years, isoxazolidine derivatives have been synthesized in high yield *via* intermolecular cycloaddition of *N*-methylnitron with disubstituted olefins and are employed for biological evaluation.

These isoxazolidines are used in the syntheses of β -lactams (Padwa *et al.*, 1981) which are of value in the treatment of bacterial infections (Ochiai *et al.*, 1967), occur as natural products (Baldwin & Aube, 1987), serve as versatile synthetic intermediates (Padwa *et al.*, 1984), and are biologically interesting compounds. In view of the interest shown in these compounds, we report herein the crystal structure of the title compound, (I).

The overall view and atom-labelling of the molecule of (I) are displayed in Fig. 1. The isoxazolidine ring (O1/N1/C7-C9) adopts an envelope conformation, with atom N1 displaced by 0.326 (2) Å from the plane of the other ring atoms (Cremer & Pople, 1975).

The crystal packing is stabilized by intermolecular C—H \cdots N and C—H \cdots O hydrogen bonds (Table 1). As shown in Fig. 2, these hydrogen bonds form $R_3^3(18)$ motifs which are fused to form ribbon-like structure extending along the *b* axis.

S2. Experimental

N-Methyl-*C*-(4-bromophenyl)nitron was prepared from 4-bromo benzaldehyde, *N*-methyl-hydroxylamine hydrochloride and sodium carbonate in CH₂Cl₂ according to the literature method (Heaney *et al.*, 2001). For the preparation of the title compound, *N*-methyl-*C*-(4-bromophenyl) nitron (453 mg, 3 mmol) and dimethylmaleate (475 mg, 3,3 mmol) were dissolved in benzene (50 ml). The reaction mixture was refluxed for 9 h, and monitored by TLC. After evaporation of the solvent, the reaction mixture was separated by column chromatography, using a mixture of hexane-ethyl acetate (1:2) as the eluent. The *trans*-isomer was recrystallized from methanol in 3 d (m.p. 354–355 K).

S3. Refinement

H atoms were positioned geometrically (C-H = 0.93–0.98 Å) and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{methyl C})$. A rotating-group model was used for the methyl groups.

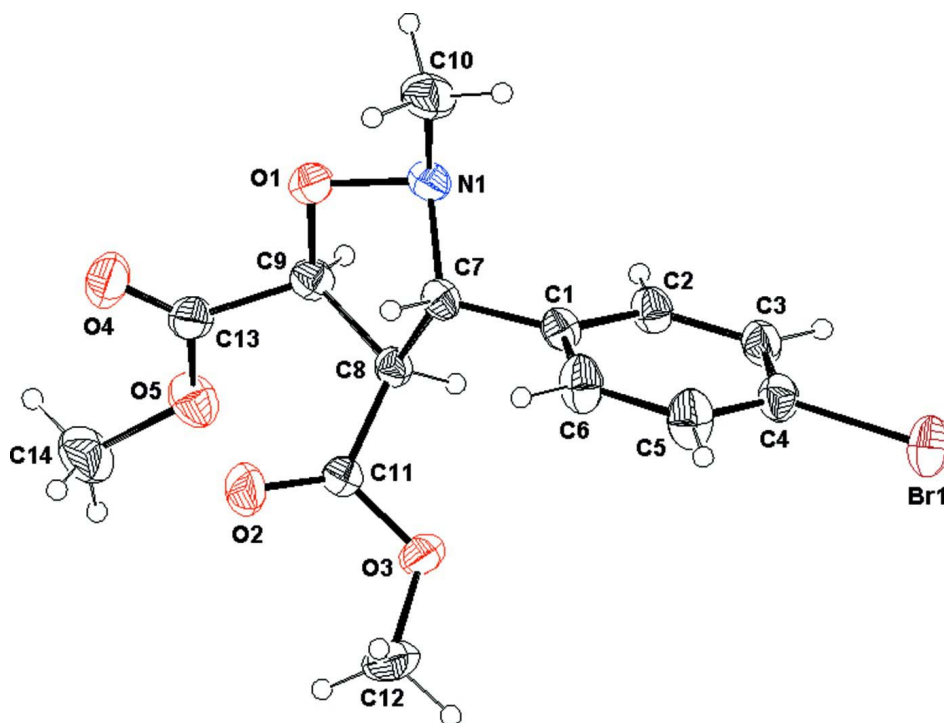


Figure 1

A view of (I) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

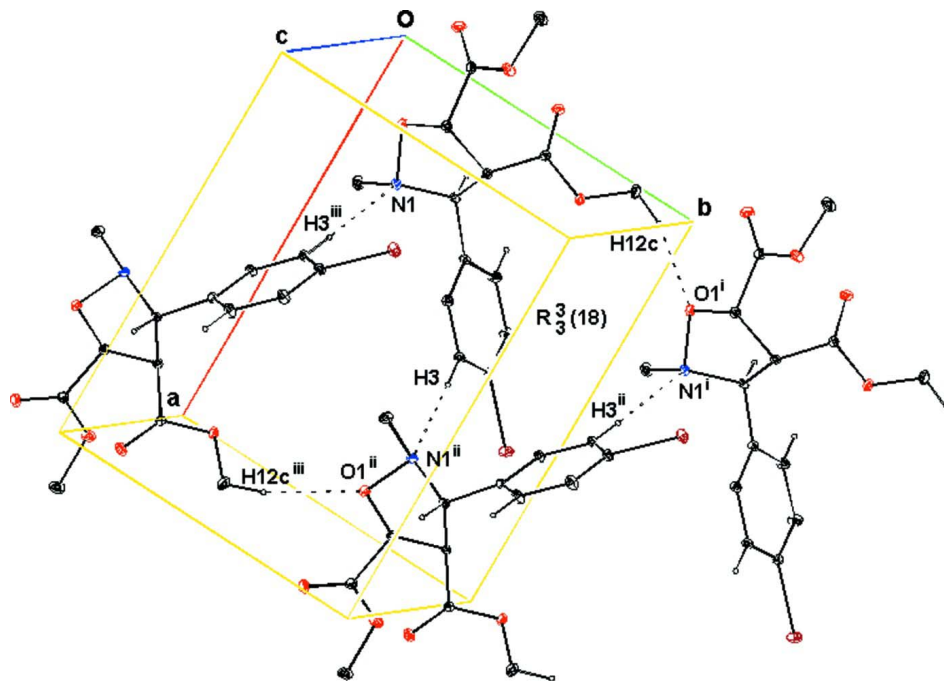
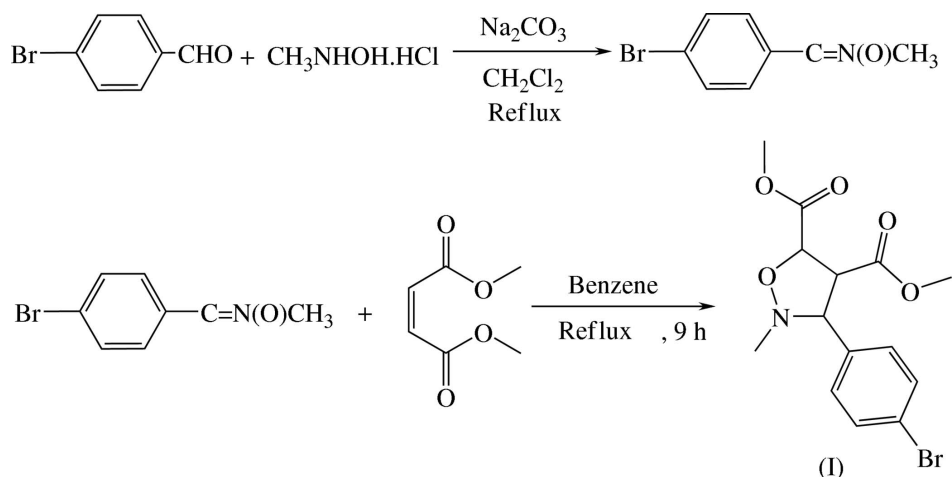


Figure 2

Part of the crystal structure of (I), showing the formation of hydrogen-bonded $R_3^3(18)$ motifs. H atoms not involved in hydrogen bonds have been omitted for clarity. Dashed lines indicate hydrogen bonds. [Symmetry codes: (i) $x, 1 + y, z$; (ii) $1 - x, 1/2 + y, 1/2 - z$; (iii) $1 - x, y - 1/2, 1/2 - z$].


Figure 3

Preparation of the title compound.

(3*R*,4*S*,5*R*)-Dimethyl *trans*-3-(4-bromophenyl)-2-methylisoxazolidine-4,5-dicarboxylate
Crystal data
 $C_{14}H_{16}BrNO_5$
 $M_r = 358.19$

 Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

 $a = 10.9020$ (4) Å

 $b = 8.1780$ (3) Å

 $c = 17.8127$ (8) Å

 $\beta = 101.622$ (3)°

 $V = 1555.56$ (11) Å³
 $Z = 4$
 $F(000) = 728$
 $D_x = 1.529$ Mg m⁻³

 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 15678 reflections

 $\theta = 1.9$ – 28.0 °

 $\mu = 2.67$ mm⁻¹
 $T = 296$ K

Block, colourless

 $0.71 \times 0.60 \times 0.45$ mm

Data collection

Stoe IPDS II

diffractometer

Radiation source: sealed X-ray tube, 12 x 0.4 mm long-fine focus

Plane graphite monochromator

 Detector resolution: 6.67 pixels mm⁻¹
 ω -scan rotation method

Absorption correction: integration

 (*X-RED32*; Stoe & Cie, 2002)

 $T_{\min} = 0.327$, $T_{\max} = 0.480$

15678 measured reflections

3232 independent reflections

 2696 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.043$
 $\theta_{\max} = 26.5$ °, $\theta_{\min} = 1.9$ °

 $h = -13$ → 13
 $k = -10$ → 10
 $l = -22$ → 22
Refinement

 Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.101$
 $S = 1.12$

3232 reflections

193 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

 $w = 1/[\sigma^2(F_o^2) + (0.0399P)^2 + 1.0321P]$

 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.49$ e Å⁻³
 $\Delta\rho_{\min} = -0.93$ e Å⁻³

Special details

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C2	0.4201 (3)	0.5503 (4)	0.18030 (16)	0.0460 (7)
H2	0.4149	0.4978	0.2259	0.055*
C8	0.1314 (2)	0.4572 (3)	0.16330 (15)	0.0371 (5)
H8	0.1754	0.5207	0.2072	0.044*
C5	0.4354 (3)	0.7096 (5)	0.04589 (18)	0.0617 (9)
H5	0.4399	0.7642	0.0007	0.074*
C12	0.0010 (4)	0.8290 (4)	0.0637 (3)	0.0793 (12)
H12C	0.0311	0.9380	0.0761	0.095*
H12B	-0.0850	0.8216	0.0685	0.095*
H12A	0.0074	0.8037	0.0120	0.095*
C11	0.0347 (3)	0.5632 (3)	0.11402 (16)	0.0403 (6)
O3	0.0753 (2)	0.7144 (3)	0.11549 (15)	0.0625 (6)
C10	0.3202 (3)	0.1297 (5)	0.1037 (2)	0.0643 (9)
H10C	0.3990	0.1723	0.0965	0.077*
H10B	0.2642	0.1192	0.0549	0.077*
H10A	0.3329	0.0243	0.1278	0.077*
C4	0.5204 (3)	0.7411 (4)	0.11187 (18)	0.0473 (7)
C6	0.3420 (3)	0.5947 (5)	0.04732 (18)	0.0575 (8)
H6	0.2844	0.5715	0.0025	0.069*
N1	0.2659 (2)	0.2410 (3)	0.15238 (14)	0.0448 (5)
O1	0.14367 (19)	0.1690 (2)	0.15537 (13)	0.0506 (5)
O2	-0.0615 (2)	0.5161 (3)	0.07465 (15)	0.0670 (7)
C13	-0.0534 (3)	0.2653 (4)	0.17647 (19)	0.0497 (7)
C14	-0.2378 (3)	0.3749 (6)	0.2055 (3)	0.0783 (12)
H14C	-0.2637	0.4566	0.2378	0.094*
H14B	-0.2667	0.2696	0.2183	0.094*
H14A	-0.2728	0.3994	0.1529	0.094*
O4	-0.1095 (2)	0.1619 (3)	0.1377 (2)	0.0846 (9)
O5	-0.1029 (2)	0.3738 (3)	0.21707 (14)	0.0628 (6)
Br1	0.64870 (3)	0.89981 (5)	0.11195 (2)	0.06568 (15)
C3	0.5147 (3)	0.6630 (4)	0.17970 (17)	0.0462 (7)
H3	0.5732	0.6857	0.2242	0.055*
C7	0.2260 (2)	0.3988 (3)	0.11494 (15)	0.0385 (6)
H7	0.1817	0.3788	0.0622	0.046*
C9	0.0865 (3)	0.2939 (4)	0.19170 (16)	0.0432 (6)

H9	0.1185	0.2863	0.2471	0.052*
C1	0.3332 (2)	0.5142 (4)	0.11427 (15)	0.0400 (6)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C2	0.0426 (14)	0.0554 (18)	0.0379 (14)	-0.0027 (13)	0.0028 (12)	0.0067 (13)
C8	0.0342 (12)	0.0394 (14)	0.0357 (13)	-0.0002 (11)	0.0024 (11)	-0.0037 (11)
C5	0.0592 (19)	0.082 (3)	0.0438 (17)	-0.0175 (18)	0.0108 (15)	0.0108 (17)
C12	0.089 (3)	0.0383 (18)	0.097 (3)	0.0084 (18)	-0.013 (2)	0.005 (2)
C11	0.0381 (13)	0.0392 (15)	0.0430 (14)	0.0014 (11)	0.0070 (12)	-0.0011 (12)
O3	0.0635 (14)	0.0341 (11)	0.0769 (15)	-0.0017 (10)	-0.0164 (12)	-0.0014 (11)
C10	0.060 (2)	0.061 (2)	0.075 (2)	0.0151 (17)	0.0198 (18)	-0.0072 (18)
C4	0.0387 (14)	0.0532 (17)	0.0517 (17)	-0.0047 (13)	0.0135 (13)	-0.0030 (14)
C6	0.0530 (17)	0.081 (2)	0.0361 (15)	-0.0153 (17)	0.0021 (13)	0.0038 (16)
N1	0.0378 (11)	0.0448 (14)	0.0520 (14)	0.0067 (10)	0.0094 (11)	0.0022 (11)
O1	0.0472 (11)	0.0391 (11)	0.0675 (14)	0.0010 (9)	0.0162 (10)	0.0016 (10)
O2	0.0493 (12)	0.0642 (16)	0.0749 (16)	-0.0133 (11)	-0.0174 (12)	0.0165 (13)
C13	0.0463 (16)	0.0454 (17)	0.0604 (18)	0.0012 (14)	0.0176 (15)	0.0068 (15)
C14	0.0518 (19)	0.094 (3)	0.095 (3)	0.012 (2)	0.030 (2)	0.002 (2)
O4	0.0544 (14)	0.0595 (16)	0.142 (3)	-0.0144 (12)	0.0256 (17)	-0.0311 (18)
O5	0.0482 (12)	0.0809 (17)	0.0622 (14)	0.0056 (11)	0.0182 (11)	-0.0075 (12)
Br1	0.0557 (2)	0.0767 (3)	0.0680 (2)	-0.02186 (17)	0.02031 (16)	0.00019 (19)
C3	0.0383 (14)	0.0518 (17)	0.0452 (15)	-0.0021 (12)	0.0007 (12)	-0.0004 (14)
C7	0.0352 (12)	0.0441 (15)	0.0345 (13)	0.0012 (11)	0.0026 (10)	0.0000 (12)
C9	0.0434 (14)	0.0454 (16)	0.0414 (14)	0.0032 (12)	0.0102 (12)	0.0049 (13)
C1	0.0352 (13)	0.0477 (16)	0.0361 (13)	0.0005 (12)	0.0046 (11)	0.0011 (12)

Geometric parameters (Å, °)

C2—C3	1.385 (4)	C10—H10A	0.96
C2—C1	1.386 (4)	C4—C3	1.379 (4)
C2—H2	0.93	C4—Br1	1.908 (3)
C8—C11	1.503 (4)	C6—C1	1.382 (4)
C8—C9	1.542 (4)	C6—H6	0.93
C8—C7	1.547 (3)	N1—O1	1.468 (3)
C8—H8	0.98	N1—C7	1.478 (4)
C5—C4	1.367 (5)	O1—C9	1.419 (3)
C5—C6	1.390 (5)	C13—O4	1.181 (4)
C5—H5	0.93	C13—O5	1.326 (4)
C12—O3	1.444 (4)	C13—C9	1.512 (4)
C12—H12C	0.96	C14—O5	1.444 (4)
C12—H12B	0.96	C14—H14C	0.96
C12—H12A	0.96	C14—H14B	0.96
C11—O2	1.202 (4)	C14—H14A	0.96
C11—O3	1.312 (4)	C3—H3	0.93
C10—N1	1.462 (4)	C7—C1	1.505 (4)
C10—H10C	0.96	C7—H7	0.98

C10—H10B	0.96	C9—H9	0.98
C3—C2—C1	121.2 (3)	C5—C6—H6	119.5
C3—C2—H2	119.4	C10—N1—O1	104.5 (2)
C1—C2—H2	119.4	C10—N1—C7	113.2 (2)
C11—C8—C9	117.3 (2)	O1—N1—C7	100.31 (19)
C11—C8—C7	108.7 (2)	C9—O1—N1	102.4 (2)
C9—C8—C7	101.9 (2)	O4—C13—O5	125.3 (3)
C11—C8—H8	109.5	O4—C13—C9	126.9 (3)
C9—C8—H8	109.5	O5—C13—C9	107.7 (3)
C7—C8—H8	109.5	O5—C14—H14C	109.5
C4—C5—C6	119.0 (3)	O5—C14—H14B	109.5
C4—C5—H5	120.5	H14C—C14—H14B	109.5
C6—C5—H5	120.5	O5—C14—H14A	109.5
O3—C12—H12C	109.5	H14C—C14—H14A	109.5
O3—C12—H12B	109.5	H14B—C14—H14A	109.5
H12C—C12—H12B	109.5	C13—O5—C14	116.2 (3)
O3—C12—H12A	109.5	C4—C3—C2	118.7 (3)
H12C—C12—H12A	109.5	C4—C3—H3	120.6
H12B—C12—H12A	109.5	C2—C3—H3	120.6
O2—C11—O3	124.4 (3)	N1—C7—C1	113.1 (2)
O2—C11—C8	125.7 (3)	N1—C7—C8	100.8 (2)
O3—C11—C8	109.8 (2)	C1—C7—C8	114.6 (2)
C11—O3—C12	117.1 (3)	N1—C7—H7	109.3
N1—C10—H10C	109.5	C1—C7—H7	109.3
N1—C10—H10B	109.5	C8—C7—H7	109.3
H10C—C10—H10B	109.5	O1—C9—C13	109.1 (2)
N1—C10—H10A	109.5	O1—C9—C8	106.0 (2)
H10C—C10—H10A	109.5	C13—C9—C8	116.8 (2)
H10B—C10—H10A	109.5	O1—C9—H9	108.2
C5—C4—C3	121.5 (3)	C13—C9—H9	108.2
C5—C4—Br1	120.0 (2)	C8—C9—H9	108.2
C3—C4—Br1	118.5 (2)	C6—C1—C2	118.5 (3)
C1—C6—C5	121.1 (3)	C6—C1—C7	119.6 (3)
C1—C6—H6	119.5	C2—C1—C7	121.8 (2)
C9—C8—C11—O2	-25.5 (4)	C9—C8—C7—N1	-26.7 (2)
C7—C8—C11—O2	89.3 (3)	C11—C8—C7—C1	87.0 (3)
C9—C8—C11—O3	159.4 (2)	C9—C8—C7—C1	-148.5 (2)
C7—C8—C11—O3	-85.8 (3)	N1—O1—C9—C13	161.6 (2)
O2—C11—O3—C12	-2.7 (5)	N1—O1—C9—C8	35.1 (3)
C8—C11—O3—C12	172.4 (3)	O4—C13—C9—O1	-3.5 (5)
C6—C5—C4—C3	-0.8 (5)	O5—C13—C9—O1	174.0 (2)
C6—C5—C4—Br1	-179.4 (3)	O4—C13—C9—C8	116.5 (4)
C4—C5—C6—C1	0.8 (6)	O5—C13—C9—C8	-65.9 (3)
C10—N1—O1—C9	-170.5 (2)	C11—C8—C9—O1	113.7 (3)
C7—N1—O1—C9	-53.0 (2)	C7—C8—C9—O1	-4.9 (3)
O4—C13—O5—C14	-7.2 (5)	C11—C8—C9—C13	-8.0 (4)

C9—C13—O5—C14	175.2 (3)	C7—C8—C9—C13	-126.6 (3)
C5—C4—C3—C2	0.0 (5)	C5—C6—C1—C2	-0.2 (5)
Br1—C4—C3—C2	178.6 (2)	C5—C6—C1—C7	176.0 (3)
C1—C2—C3—C4	0.7 (5)	C3—C2—C1—C6	-0.6 (5)
C10—N1—C7—C1	-77.9 (3)	C3—C2—C1—C7	-176.7 (3)
O1—N1—C7—C1	171.2 (2)	N1—C7—C1—C6	133.1 (3)
C10—N1—C7—C8	159.2 (2)	C8—C7—C1—C6	-112.1 (3)
O1—N1—C7—C8	48.3 (2)	N1—C7—C1—C2	-50.9 (4)
C11—C8—C7—N1	-151.2 (2)	C8—C7—C1—C2	63.9 (4)

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
C3—H3...N1 ⁱ	0.93	2.56	3.492 (4)	179
C12—H12C...O1 ⁱⁱ	0.96	2.52	3.434 (5)	158

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