

2,4-Bis(2-bromophenyl)-3-azabicyclo-[3.3.1]nonan-9-one

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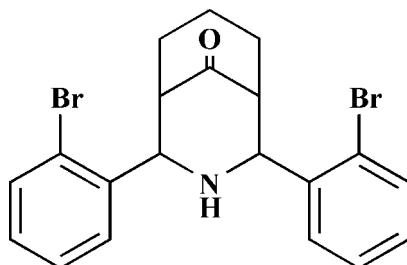
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.027; wR factor = 0.060; data-to-parameter ratio = 18.5.

In the molecular structure of the title compound, $\text{C}_{20}\text{H}_{19}\text{Br}_2\text{NO}$, the fused six-membered heterocyclic and cyclohexane rings adopt a twin-chair conformation with equatorial orientations of all the substituents. Both the *ortho*-bromo substituents of the benzene rings are oriented towards the carbonyl group; the dihedral angle between the ring planes is $29.13(3)^\circ$. In the crystal structure, the N—H group does not participate in any hydrogen bonds.

Related literature

For 3-azabicyclononan-9-ones and their significance as bioactive molecules, see: Barker *et al.* (2005); Jeyaraman & Avila (1981). For puckering parameters, see: Cremer & Pople (1975); Web & Becker (1967). For a similar structure see; Parthiban *et al.* (2008).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{19}\text{Br}_2\text{NO}$	$\gamma = 97.399(1)^\circ$
$M_r = 449.18$	$V = 885.94(5)\text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 2$
$a = 7.8389(3)\text{ \AA}$	Mo $K\alpha$ radiation
$b = 10.5770(3)\text{ \AA}$	$\mu = 4.58\text{ mm}^{-1}$
$c = 11.0274(3)\text{ \AA}$	$T = 298(2)\text{ K}$
$\alpha = 101.099(2)^\circ$	$0.45 \times 0.38 \times 0.35\text{ mm}$
$\beta = 93.725(2)^\circ$	

Data collection

Bruker APEXII CCD	10959 measured reflections
diffractometer	4098 independent reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 1999)	3266 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.232$, $T_{\max} = 0.297$	$R_{\text{int}} = 0.017$
(expected range = 0.157–0.201)	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.027$	H atoms treated by a mixture of
$wR(F^2) = 0.060$	independent and constrained
$S = 1.00$	refinement
4098 reflections	$\Delta\rho_{\max} = 0.58\text{ e \AA}^{-3}$
221 parameters	$\Delta\rho_{\min} = -0.56\text{ e \AA}^{-3}$

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT-Plus* (Bruker, 2004); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2843).

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supporting information

Acta Cryst. (2008). E64, o2385 [doi:10.1107/S1600536808037501]

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S1. Comment

3-Azabicyclononan-9-ones are important class of compounds due to their significance as bio-active molecules (Jeyaraman & Avila, 1981; Barker et al., 2005).

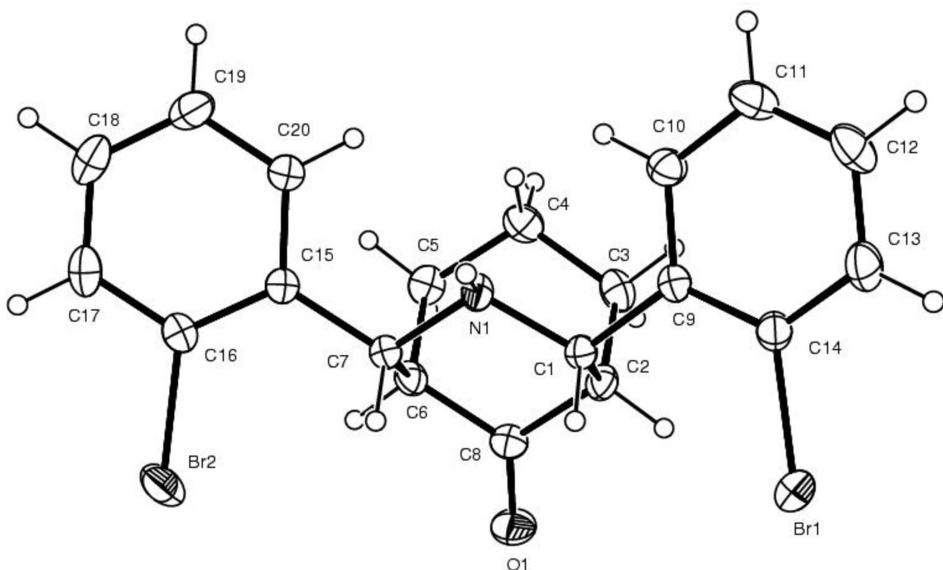
The title compound, (I), exists in a chair-chair conformation with equatorial orientations of the ortho bromo-phenyl groups on each side of the secondary amino group with the torsion angles of C8—C2—C1—C9 and C8—C6—C7—C15 being 177.88 (4) and 179.42 (6) $^{\circ}$, respectively. In both aryl groups, the bromo substituents point towards the carbonyl group and the dihedral angle between the ring planes is 29.13 (3) $^{\circ}$. The piperidine ring adopts near ideal chair conformation with the deviation of ring atoms N1 and C8 from the C1/C2/C6/C7 plane by -0.635 (3) and 0.705 (3) \AA , respectively, $Q_T = 0.599$ (2) \AA , $q(2)=0.047$ (2) \AA , $q(3)=0.597$ (2) \AA , $\theta = 4.71$ (19) $^{\circ}$ Cremer & Pople, 1975; Web & Becker, 1967), whereas the cyclohexane ring atoms C4 and C8 deviate from the C2/C3/C5/C6 plane by -0.539 (4) and 0.725 (3) \AA , respectively, $Q_T = 0.565$ (2) \AA , $q(2)=0.141$ (2) \AA , $q(3)=0.548$ (2) \AA , $\theta = 14.4$ (2) $^{\circ}$, indicating a deviation from the ideal chair conformation of the cyclohexane part in the title compound. The crystal structure is stabilized by the intermolecular van der Waals interactions.

S2. Experimental

A mixture of cyclohexanone (0.05 mol) and *ortho* bromobenzaldehyde (0.1 mol) was added to a warm solution of ammonium acetate (0.075 mol) in 50 ml of absolute ethanol. The mixture was gently warmed on a hot plate until a yellow colour was formed and then cooled to room temperature. Then, 50 ml of ether was added and allowed to stir over night at room temperature. At the end, the crude azabicyclic ketone was separated by filtration and washed with 1:5 v/v ethanol–ether mixture till the solid became colourless. Recrystallization of the compound from acetone gave colourless blocks of (I).

S3. Refinement

The nitrogen-bound H atom was located in a difference map and refined isotropically. The other hydrogen atoms were fixed geometrically ($C—H = 0.93$ – 0.98\AA) and refined as riding with $U_{iso}(H) = 1.2U_{eq}(C)$.

**Figure 1**

The molecular structure of (I) with non-hydrogen atoms represented as 30% probability ellipsoids.

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Crystal data

$C_{20}H_{19}Br_2NO$
 $M_r = 449.18$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 7.8389 (3)$ Å
 $b = 10.5770 (3)$ Å
 $c = 11.0274 (3)$ Å
 $\alpha = 101.099 (2)^\circ$
 $\beta = 93.725 (2)^\circ$
 $\gamma = 97.399 (1)^\circ$
 $V = 885.94 (5)$ Å³

$Z = 2$
 $F(000) = 448$
 $D_x = 1.684 \text{ Mg m}^{-3}$
 $Mo K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 5642 reflections
 $\theta = 2.5\text{--}28.2^\circ$
 $\mu = 4.58 \text{ mm}^{-1}$
 $T = 298$ K
Block, colourless
 $0.45 \times 0.38 \times 0.35$ mm

Data collection

Bruker APEXII CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 1999)
 $T_{\min} = 0.232$, $T_{\max} = 0.297$

10959 measured reflections
4098 independent reflections
3266 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 1.9^\circ$
 $h = -10 \rightarrow 10$
 $k = -13 \rightarrow 13$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.060$
 $S = 1.00$
4098 reflections

221 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 atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0207P)^2 + 0.5638P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.002$$

$$\Delta\rho_{\max} = 0.58 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.56 \text{ e \AA}^{-3}$$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	-0.00438 (3)	0.47719 (2)	0.77681 (2)	0.05065 (8)
Br2	0.16949 (3)	1.23193 (3)	1.01448 (2)	0.06070 (9)
C1	0.2124 (2)	0.74756 (18)	0.75331 (18)	0.0301 (4)
H1	0.1437	0.7370	0.8228	0.036*
C2	0.0889 (2)	0.76537 (19)	0.64395 (19)	0.0344 (4)
H2	-0.0049	0.6918	0.6254	0.041*
C3	0.1735 (3)	0.7773 (2)	0.5238 (2)	0.0428 (5)
H3A	0.2294	0.7011	0.4980	0.051*
H3B	0.0839	0.7778	0.4591	0.051*
C4	0.3064 (3)	0.8987 (2)	0.5360 (2)	0.0432 (5)
H4A	0.4119	0.8865	0.5806	0.052*
H4B	0.3333	0.9102	0.4539	0.052*
C5	0.2434 (3)	1.0214 (2)	0.60396 (19)	0.0392 (5)
H5A	0.1647	1.0500	0.5468	0.047*
H5B	0.3418	1.0895	0.6280	0.047*
C6	0.1515 (2)	1.00528 (19)	0.72040 (18)	0.0331 (4)
H6	0.0967	1.0828	0.7479	0.040*
C7	0.2692 (2)	0.98341 (18)	0.83118 (17)	0.0294 (4)
H7	0.1987	0.9747	0.9001	0.035*
C8	0.0132 (2)	0.8886 (2)	0.68529 (18)	0.0342 (4)
C9	0.3014 (2)	0.62834 (18)	0.72007 (17)	0.0304 (4)
C10	0.4692 (3)	0.6383 (2)	0.6847 (2)	0.0389 (5)
H10	0.5265	0.7194	0.6796	0.047*
C11	0.5525 (3)	0.5302 (2)	0.6570 (2)	0.0495 (6)
H11	0.6641	0.5392	0.6328	0.059*
C12	0.4711 (3)	0.4095 (2)	0.6651 (2)	0.0556 (6)

H12	0.5280	0.3371	0.6472	0.067*
C13	0.3055 (3)	0.3958 (2)	0.6995 (2)	0.0482 (6)
H13	0.2496	0.3142	0.7047	0.058*
C14	0.2228 (3)	0.50413 (19)	0.72632 (19)	0.0350 (4)
C15	0.4125 (2)	1.09742 (18)	0.87430 (17)	0.0290 (4)
C16	0.3860 (3)	1.21258 (19)	0.95107 (18)	0.0337 (4)
C17	0.5159 (3)	1.3181 (2)	0.9863 (2)	0.0436 (5)
H17	0.4946	1.3938	1.0380	0.052*
C18	0.6757 (3)	1.3104 (2)	0.9446 (2)	0.0499 (6)
H18	0.7629	1.3812	0.9670	0.060*
C19	0.7071 (3)	1.1976 (2)	0.8696 (2)	0.0482 (6)
H19	0.8161	1.1920	0.8421	0.058*
C20	0.5772 (3)	1.0923 (2)	0.8347 (2)	0.0381 (5)
H20	0.6003	1.0166	0.7839	0.046*
N1	0.3422 (2)	0.86343 (15)	0.79409 (16)	0.0304 (4)
O1	-0.13950 (19)	0.89419 (17)	0.68681 (16)	0.0521 (4)
H1A	0.407 (3)	0.851 (2)	0.850 (2)	0.040 (7)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.04366 (13)	0.03775 (13)	0.06740 (17)	-0.00469 (9)	0.01281 (11)	0.00737 (11)
Br2	0.05389 (15)	0.06429 (18)	0.05949 (17)	0.02181 (12)	0.01497 (12)	-0.01039 (13)
C1	0.0296 (9)	0.0256 (10)	0.0349 (10)	0.0042 (7)	0.0038 (8)	0.0056 (8)
C2	0.0286 (9)	0.0300 (10)	0.0415 (11)	0.0019 (8)	-0.0038 (8)	0.0035 (9)
C3	0.0511 (13)	0.0398 (12)	0.0351 (11)	0.0125 (10)	-0.0022 (10)	-0.0001 (9)
C4	0.0505 (13)	0.0492 (13)	0.0325 (11)	0.0104 (10)	0.0097 (9)	0.0107 (10)
C5	0.0441 (11)	0.0369 (12)	0.0386 (12)	0.0063 (9)	-0.0008 (9)	0.0138 (9)
C6	0.0326 (10)	0.0299 (10)	0.0371 (11)	0.0102 (8)	0.0012 (8)	0.0046 (8)
C7	0.0300 (9)	0.0276 (10)	0.0302 (10)	0.0044 (7)	0.0033 (7)	0.0049 (8)
C8	0.0310 (9)	0.0405 (12)	0.0327 (11)	0.0088 (8)	-0.0003 (8)	0.0100 (9)
C9	0.0335 (9)	0.0278 (10)	0.0295 (10)	0.0060 (8)	0.0003 (8)	0.0046 (8)
C10	0.0357 (10)	0.0380 (12)	0.0448 (12)	0.0081 (9)	0.0059 (9)	0.0104 (9)
C11	0.0417 (12)	0.0553 (15)	0.0575 (15)	0.0220 (11)	0.0123 (11)	0.0135 (12)
C12	0.0657 (16)	0.0446 (14)	0.0637 (16)	0.0311 (12)	0.0140 (13)	0.0109 (12)
C13	0.0608 (15)	0.0297 (12)	0.0551 (14)	0.0101 (10)	0.0065 (11)	0.0083 (10)
C14	0.0371 (10)	0.0304 (10)	0.0362 (11)	0.0041 (8)	0.0025 (8)	0.0038 (8)
C15	0.0314 (9)	0.0280 (10)	0.0284 (10)	0.0053 (8)	0.0005 (7)	0.0079 (8)
C16	0.0403 (10)	0.0322 (11)	0.0297 (10)	0.0100 (8)	0.0010 (8)	0.0062 (8)
C17	0.0646 (15)	0.0278 (11)	0.0353 (12)	0.0034 (10)	-0.0054 (10)	0.0047 (9)
C18	0.0547 (14)	0.0427 (13)	0.0455 (13)	-0.0154 (11)	-0.0072 (11)	0.0108 (11)
C19	0.0346 (11)	0.0573 (15)	0.0500 (14)	-0.0043 (10)	0.0028 (10)	0.0117 (12)
C20	0.0337 (10)	0.0375 (11)	0.0411 (12)	0.0048 (9)	0.0047 (9)	0.0029 (9)
N1	0.0292 (8)	0.0249 (8)	0.0362 (9)	0.0055 (6)	-0.0050 (7)	0.0053 (7)
O1	0.0292 (7)	0.0588 (11)	0.0693 (11)	0.0119 (7)	0.0028 (7)	0.0120 (9)

Geometric parameters (\AA , $\text{^{\circ}}$)

Br1—C14	1.903 (2)	C7—H7	0.9800
Br2—C16	1.897 (2)	C8—O1	1.207 (2)
C1—N1	1.465 (2)	C9—C10	1.393 (3)
C1—C9	1.515 (3)	C9—C14	1.394 (3)
C1—C2	1.552 (3)	C10—C11	1.382 (3)
C1—H1	0.9800	C10—H10	0.9300
C2—C8	1.505 (3)	C11—C12	1.373 (4)
C2—C3	1.539 (3)	C11—H11	0.9300
C2—H2	0.9800	C12—C13	1.373 (3)
C3—C4	1.524 (3)	C12—H12	0.9300
C3—H3A	0.9700	C13—C14	1.381 (3)
C3—H3B	0.9700	C13—H13	0.9300
C4—C5	1.525 (3)	C15—C20	1.393 (3)
C4—H4A	0.9700	C15—C16	1.391 (3)
C4—H4B	0.9700	C16—C17	1.387 (3)
C5—C6	1.539 (3)	C17—C18	1.369 (3)
C5—H5A	0.9700	C17—H17	0.9300
C5—H5B	0.9700	C18—C19	1.375 (4)
C6—C8	1.506 (3)	C18—H18	0.9300
C6—C7	1.554 (3)	C19—C20	1.385 (3)
C6—H6	0.9800	C19—H19	0.9300
C7—N1	1.457 (2)	C20—H20	0.9300
C7—C15	1.518 (2)	N1—H1A	0.81 (2)
N1—C1—C9	109.63 (15)	O1—C8—C2	124.48 (19)
N1—C1—C2	110.38 (16)	O1—C8—C6	123.98 (19)
C9—C1—C2	112.32 (16)	C2—C8—C6	111.51 (16)
N1—C1—H1	108.1	C10—C9—C14	116.51 (18)
C9—C1—H1	108.1	C10—C9—C1	121.26 (17)
C2—C1—H1	108.1	C14—C9—C1	122.21 (17)
C8—C2—C3	107.21 (17)	C11—C10—C9	121.5 (2)
C8—C2—C1	107.83 (16)	C11—C10—H10	119.3
C3—C2—C1	115.42 (16)	C9—C10—H10	119.3
C8—C2—H2	108.7	C12—C11—C10	120.3 (2)
C3—C2—H2	108.7	C12—C11—H11	119.9
C1—C2—H2	108.7	C10—C11—H11	119.9
C4—C3—C2	114.03 (17)	C13—C12—C11	119.9 (2)
C4—C3—H3A	108.7	C13—C12—H12	120.1
C2—C3—H3A	108.7	C11—C12—H12	120.1
C4—C3—H3B	108.7	C12—C13—C14	119.5 (2)
C2—C3—H3B	108.7	C12—C13—H13	120.3
H3A—C3—H3B	107.6	C14—C13—H13	120.3
C5—C4—C3	112.67 (18)	C13—C14—C9	122.3 (2)
C5—C4—H4A	109.1	C13—C14—Br1	116.76 (16)
C3—C4—H4A	109.1	C9—C14—Br1	120.89 (15)
C5—C4—H4B	109.1	C20—C15—C16	116.80 (18)

C3—C4—H4B	109.1	C20—C15—C7	120.81 (17)
H4A—C4—H4B	107.8	C16—C15—C7	122.36 (17)
C4—C5—C6	114.82 (17)	C17—C16—C15	122.0 (2)
C4—C5—H5A	108.6	C17—C16—Br2	116.64 (16)
C6—C5—H5A	108.6	C15—C16—Br2	121.32 (15)
C4—C5—H5B	108.6	C18—C17—C16	119.7 (2)
C6—C5—H5B	108.6	C18—C17—H17	120.2
H5A—C5—H5B	107.5	C16—C17—H17	120.2
C8—C6—C5	108.13 (16)	C17—C18—C19	119.9 (2)
C8—C6—C7	107.18 (16)	C17—C18—H18	120.0
C5—C6—C7	115.25 (16)	C19—C18—H18	120.0
C8—C6—H6	108.7	C18—C19—C20	120.3 (2)
C5—C6—H6	108.7	C18—C19—H19	119.9
C7—C6—H6	108.7	C20—C19—H19	119.9
N1—C7—C15	110.21 (15)	C19—C20—C15	121.3 (2)
N1—C7—C6	109.31 (15)	C19—C20—H20	119.3
C15—C7—C6	111.09 (15)	C15—C20—H20	119.3
N1—C7—H7	108.7	C7—N1—C1	113.89 (15)
C15—C7—H7	108.7	C7—N1—H1A	111.0 (16)
C6—C7—H7	108.7	C1—N1—H1A	108.6 (16)
N1—C1—C2—C8	-55.2 (2)	C9—C10—C11—C12	-0.6 (4)
C9—C1—C2—C8	-177.89 (16)	C10—C11—C12—C13	0.7 (4)
N1—C1—C2—C3	64.6 (2)	C11—C12—C13—C14	-0.3 (4)
C9—C1—C2—C3	-58.1 (2)	C12—C13—C14—C9	-0.2 (3)
C8—C2—C3—C4	55.1 (2)	C12—C13—C14—Br1	-178.65 (19)
C1—C2—C3—C4	-65.0 (2)	C10—C9—C14—C13	0.2 (3)
C2—C3—C4—C5	-45.4 (3)	C1—C9—C14—C13	-178.10 (19)
C3—C4—C5—C6	43.4 (3)	C10—C9—C14—Br1	178.65 (15)
C4—C5—C6—C8	-51.3 (2)	C1—C9—C14—Br1	0.3 (3)
C4—C5—C6—C7	68.6 (2)	N1—C7—C15—C20	23.6 (2)
C8—C6—C7—N1	58.74 (19)	C6—C7—C15—C20	-97.7 (2)
C5—C6—C7—N1	-61.6 (2)	N1—C7—C15—C16	-158.48 (18)
C8—C6—C7—C15	-179.43 (15)	C6—C7—C15—C16	80.2 (2)
C5—C6—C7—C15	60.2 (2)	C20—C15—C16—C17	0.5 (3)
C3—C2—C8—O1	113.0 (2)	C7—C15—C16—C17	-177.53 (18)
C1—C2—C8—O1	-122.1 (2)	C20—C15—C16—Br2	-178.76 (15)
C3—C2—C8—C6	-64.9 (2)	C7—C15—C16—Br2	3.2 (3)
C1—C2—C8—C6	59.9 (2)	C15—C16—C17—C18	0.3 (3)
C5—C6—C8—O1	-114.9 (2)	Br2—C16—C17—C18	179.52 (17)
C7—C6—C8—O1	120.3 (2)	C16—C17—C18—C19	-0.9 (3)
C5—C6—C8—C2	63.1 (2)	C17—C18—C19—C20	0.8 (4)
C7—C6—C8—C2	-61.7 (2)	C18—C19—C20—C15	0.0 (3)
N1—C1—C9—C10	-24.4 (3)	C16—C15—C20—C19	-0.6 (3)
C2—C1—C9—C10	98.7 (2)	C7—C15—C20—C19	177.4 (2)
N1—C1—C9—C14	153.82 (18)	C15—C7—N1—C1	178.86 (15)
C2—C1—C9—C14	-83.1 (2)	C6—C7—N1—C1	-58.8 (2)
C14—C9—C10—C11	0.2 (3)	C9—C1—N1—C7	-178.59 (16)

supporting information

C1—C9—C10—C11

178.5 (2)

C2—C1—N1—C7

57.2 (2)
