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4,4',6,6'-Tetrabromo-2,2'-(2,8-diazonia-5-azanona-1,8-diene-1,9-diyl)diphenolate

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.005 Å; disorder in main residue; R factor = 0.038; wR factor = 0.095; data-to-parameter ratio = 18.3.

In the zwitterionic title compound, $C_{18}H_{17}Br_4N_3O_2$, the two salicylaldimine groups form a dihedral angle of 51.94 (2)° and the dihedral angle between the aromatic ring planes is 51.14 (2)°. One of the C atoms adjacent to the aza N atom is disordered over two positions; the site-occupancy factors are 0.51 (1) and 0.49 (1). There are two strong intramolecular N-H···O hydrogen bonds in the molecule.

Related literature

For general background on the use of Schiff bases in metal complexes, see: Vigato *et al.* (2007).



Experimental

Crystal data

Data collection

Bruker SMART APEXII diffractometer Absorption correction: multi-scan (SADABS; Sheldrick, 1997) $T_{min} = 0.149, T_{max} = 0.227$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$	
$wR(F^2) = 0.095$	
S = 1.05	
4693 reflections	
256 parameters	
6 restraints	

17118 measured reflections 4693 independent reflections 3747 reflections with $I > 2\sigma(I)$ $R_{int} = 0.040$

H atoms treated by a mixture of independent and constrained refinement $\Delta \rho_{max} = 2.07 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{min} = -0.94 \text{ e } \text{\AA}^{-3}$

Table 1 Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1−H1 <i>A</i> ···O1	0.97 (6)	1.70 (6)	2.553 (5)	144 (5)
N3−H3 <i>A</i> ···O2	0.87 (6)	1.84 (6)	2.597 (4)	144 (5)

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *XP* in *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BQ2108).

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4,4',6,6'-Tetrabromo-2,2'-(2,8-diazonia-5-azanona-1,8-diene-1,9-diyl)diphenolate

Z.-J. Chen and K.-W. Lei

Comment

The Schiff bases are widely employed as ligands in coordination chemistry. These ligands are readily available, versatile and, depending on the nature of the starting materials (primary amines and carbonyl precursors), they exhibit various denticities and functionalities. Moreover, the number, the nature, and the relative position of the donor atoms of a Schiff base ligand allow a good control over the stereochemistry of the metallic centers, as well as over the number of the metal ions within homo- and heteropolynuclear complexes. All these advantages make Schiff bases very good candidates in the effort to synthesize metal complexes of interest in bioinorganic chemistry, catalysis, encapsulation, transport and separation processes, magnetochemistry (Vigato *et al.*, 2007). So we report here the crystal structure of the new Schiff base ligand, 4,4',6,6'-Tetrabromo-2,2'-[3-azapentane-1,5-diylbis(nitrilomethylidyne)]diphenol(I).

The molecular structure of (I) is illustrated in Fig. 1. The two pendant moieties in a *cis* conformation attach to the ends of the C—C—N—C—C backbone. The N2 atom exhibits tetrahedral sp^3 hybridization, whereas the two amide N atoms display planar sp^2 hybridization. There is no H atom attached to O1 and O2 atoms. Instead these H atoms are attached to the N1 and N3 atoms. The double-bonds C7—N1 (1.295 (6) Å) and C12—N3 (1.296 (6) Å) show the typical character of Schiff base. The dihedral angle between the salicylaldimine groups is 51.94 (2)°. The crystal structure of (I) is stabilized by intramolecular N—H···O hydrogen bonding. The C10 atom is disorder over two positions with the site-occupancy factors of 0.51 (1) and 0.49 (1). The larger than normal range of thermal motion is mostly due to the difference between the disordered group and the other atoms which are not disordered.

Experimental

N-(2-aminoethyl)ethane-1,2-diamine (0.01 mol, 1.03 g) and 2-hydroxy-3,5-dibromobenzaldehyde(0.02 mol, 5.60 g) were dissolved in 20 ml e thanol and the solution was stirred for 3 h. After filtration and evaporation, a pure yellow product was recrystallized from ethanol. Yield: 81.7%. Calcd. for $C_{18}H_{17}Br_4N_3O_2$: C, 34.48; H, 2.73; N, 6.70; Found: C, 34.59; H, 2.62; N, 6.81%.

Refinement

All H atoms except the N attached H1A and H3A which refined freely were placed in geometrically idealized positions and constrained to ride on their parent atoms (C—H = 0.93%A, 0.97%A; N—H = 0.86 Å; and U_{iso} (H) values equal to 1.2 U_{eq} C.

Figures



Fig. 1. The structure of (I), showing 50% probability displacement ellipsoids and the atomnumbering scheme. Dashed lines show H-bondings. Only the major component is shown.

4,4',6,6'-Tetrabromo-2,2'-(2,8-diazonia-5-azanona-1,8-diene-1,9-diyl)diphenolate

Crystal data	
$C_{18}H_{17}Br_4N_3O_2$	$F_{000} = 1208$
$M_r = 626.99$	$D_{\rm x} = 2.052 \ {\rm Mg \ m}^{-3}$
Monoclinic, $P2_1/n$	Mo K α radiation $\lambda = 0.71073$ Å
Hall symbol: -P 2yn	Cell parameters from 5793 reflections
a = 9.4506 (11) Å	$\theta = 1.0-27.6^{\circ}$
b = 9.1242 (11) Å	$\mu = 7.95 \text{ mm}^{-1}$
c = 23.618 (3) Å	T = 293 (2) K
$\beta = 94.774 \ (2)^{\circ}$	BLOCK, yellow
V = 2029.5 (4) Å ³	$0.26 \times 0.21 \times 0.19 \text{ mm}$
Z = 4	

Data collection

Bruker SMART APEXII diffractometer	4693 independent reflections
Radiation source: fine-focus sealed tube	3747 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\rm int} = 0.040$
T = 293(2) K	$\theta_{\text{max}} = 27.6^{\circ}$
ϕ and ω scans	$\theta_{\min} = 1.7^{\circ}$
Absorption correction: multi-scan (SADABS; Sheldrick, 1997)	$h = -12 \rightarrow 12$
$T_{\min} = 0.149, \ T_{\max} = 0.227$	$k = -11 \rightarrow 10$
17118 measured reflections	$l = -30 \rightarrow 30$

Refinement

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
$w = 1/[\sigma^2(F_0^2) + (0.0444P)^2 + 3.6177P]$

	where $P = (F_0^2 + 2F_c^2)/3$
<i>S</i> = 1.05	$(\Delta/\sigma)_{max} = 0.001$
4693 reflections	$\Delta\rho_{max} = 2.07 \text{ e} \text{ Å}^{-3}$
256 parameters	$\Delta \rho_{min} = -0.94 \text{ e } \text{\AA}^{-3}$
6 restraints	Extinction correction: none
Primary atom site location: structure-invariant direct	

methods

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 (A^2)

	x	У	z	$U_{\rm iso}*/U_{\rm eq}$	Occ. (<1)
C1	1.1639 (4)	1.1651 (4)	0.03605 (16)	0.0255 (8)	
C2	1.2507 (4)	1.0675 (4)	0.00631 (15)	0.0244 (8)	
C3	1.2518 (4)	0.9186 (4)	0.01388 (16)	0.0260 (8)	
Н3	1.3097	0.8596	-0.0065	0.031*	
C4	1.1645 (4)	0.8557 (4)	0.05281 (18)	0.0280 (9)	
C5	1.0778 (4)	0.9402 (5)	0.08246 (17)	0.0277 (8)	
Н5	1.0197	0.8968	0.1076	0.033*	
C6	1.0763 (4)	1.0945 (5)	0.07503 (16)	0.0252 (8)	
C7	0.9904 (4)	1.1819 (5)	0.10911 (17)	0.0286 (9)	
H7	0.9341	1.1363	0.1344	0.034*	
C8	0.9087 (5)	1.4217 (5)	0.13912 (18)	0.0362 (10)	
H8A	0.9734	1.4858	0.1615	0.043*	
H8B	0.8566	1.3647	0.1651	0.043*	
C9	0.8055 (5)	1.5135 (5)	0.10130 (19)	0.0367 (10)	
H9A	0.7693	1.5925	0.1235	0.044*	
H9B	0.8551	1.5567	0.0711	0.044*	
C10	0.5604 (11)	1.4631 (12)	0.1098 (5)	0.0500 (18)	0.501 (9)
H10A	0.5928	1.4777	0.1495	0.060*	0.501 (9)
H10B	0.5188	1.5544	0.0955	0.060*	0.501 (9)
C10'	0.5427 (11)	1.4357 (13)	0.0666 (5)	0.0500 (18)	0.499 (9)
H10C	0.5168	1.3988	0.0286	0.060*	0.499 (9)
H10D	0.5160	1.5383	0.0670	0.060*	0.499 (9)
C11	0.4590 (6)	1.3573 (6)	0.1061 (3)	0.0619 (17)	
H11A	0.4238	1.3449	0.0666	0.074*	
H11B	0.3801	1.3881	0.1270	0.074*	

C12	0.4349 (4)	1.1126 (5)	0.14599 (17)	0.0303 (9)
H12	0.3368	1.1242	0.1417	0.036*
C13	0.4899 (4)	0.9821 (4)	0.17117 (15)	0.0235 (8)
C14	0.6423 (4)	0.9637 (4)	0.18085 (15)	0.0228 (8)
C15	0.6853 (4)	0.8296 (4)	0.20880 (15)	0.0227 (8)
C16	0.5925 (4)	0.7256 (4)	0.22511 (16)	0.0259 (8)
H16	0.6260	0.6405	0.2433	0.031*
C17	0.4452 (4)	0.7494 (4)	0.21399 (17)	0.0277 (8)
C18	0.3950 (4)	0.8739 (5)	0.18751 (16)	0.0266 (8)
H18	0.2977	0.8875	0.1802	0.032*
Br1	1.37074 (5)	1.15131 (5)	-0.045130 (17)	0.03273 (12)
Br2	1.17523 (5)	0.64919 (5)	0.06248 (2)	0.04388 (14)
Br3	0.88370 (4)	0.79812 (5)	0.222670 (17)	0.03065 (12)
Br4	0.31565 (5)	0.60611 (6)	0.23712 (2)	0.04384 (14)
N1	0.9899 (4)	1.3230 (4)	0.10528 (15)	0.0310 (8)
N2	0.6880 (4)	1.4267 (5)	0.07679 (18)	0.0447 (10)
H2	0.6896	1.3661	0.0490	0.054*
N3	0.5128 (4)	1.2170 (4)	0.12852 (16)	0.0342 (8)
01	1.1653 (3)	1.3038 (3)	0.02905 (13)	0.0365 (7)
O2	0.7294 (3)	1.0603 (3)	0.16669 (11)	0.0265 (6)
H1A	1.050 (6)	1.358 (6)	0.077 (2)	0.052 (15)*
H3A	0.603 (6)	1.199 (6)	0.137 (2)	0.052 (16)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0248 (19)	0.026 (2)	0.0259 (19)	0.0012 (16)	0.0033 (14)	0.0018 (15)
C2	0.0241 (19)	0.026 (2)	0.0231 (18)	-0.0036 (15)	0.0034 (14)	0.0021 (15)
C3	0.0238 (19)	0.024 (2)	0.0295 (19)	0.0033 (15)	-0.0026 (15)	-0.0002 (16)
C4	0.025 (2)	0.021 (2)	0.037 (2)	-0.0015 (16)	-0.0051 (16)	0.0068 (17)
C5	0.0217 (19)	0.030 (2)	0.031 (2)	-0.0039 (16)	-0.0001 (15)	0.0095 (17)
C6	0.0200 (18)	0.030 (2)	0.0259 (19)	0.0013 (16)	0.0020 (14)	0.0043 (16)
C7	0.024 (2)	0.033 (2)	0.029 (2)	0.0017 (17)	0.0043 (15)	0.0062 (17)
C8	0.040 (2)	0.038 (3)	0.032 (2)	0.009 (2)	0.0082 (18)	-0.0018 (19)
C9	0.047 (3)	0.028 (2)	0.037 (2)	0.009 (2)	0.0102 (19)	-0.0003 (18)
C10	0.050 (4)	0.050 (4)	0.050 (4)	0.000 (3)	0.004 (4)	0.000 (4)
C10'	0.050 (4)	0.050 (4)	0.050 (4)	0.000 (3)	0.004 (4)	0.000 (4)
C11	0.046 (3)	0.049 (3)	0.094 (5)	0.021 (3)	0.026 (3)	0.042 (3)
C12	0.0202 (19)	0.041 (3)	0.030 (2)	0.0029 (17)	0.0028 (15)	0.0027 (18)
C13	0.0210 (18)	0.029 (2)	0.0210 (17)	-0.0010 (15)	0.0009 (14)	-0.0005 (15)
C14	0.0226 (18)	0.027 (2)	0.0194 (17)	-0.0007 (16)	0.0040 (14)	-0.0058 (15)
C15	0.0234 (18)	0.024 (2)	0.0205 (17)	0.0015 (15)	0.0031 (14)	-0.0046 (15)
C16	0.034 (2)	0.022 (2)	0.0232 (18)	0.0001 (16)	0.0073 (15)	-0.0017 (15)
C17	0.030 (2)	0.027 (2)	0.0271 (19)	-0.0079 (17)	0.0100 (15)	-0.0036 (16)
C18	0.0217 (19)	0.033 (2)	0.0249 (18)	0.0004 (16)	0.0036 (14)	-0.0046 (16)
Br1	0.0407 (2)	0.0288 (2)	0.0309 (2)	0.00118 (18)	0.01642 (17)	0.00159 (16)
Br2	0.0412 (3)	0.0226 (2)	0.0680 (3)	0.00063 (19)	0.0055 (2)	0.0134 (2)
Br3	0.0242 (2)	0.0332 (2)	0.0343 (2)	0.00332 (17)	0.00105 (15)	0.00357 (17)

Br4	0.0378(3)	0.0422(3)	0.0531 (3)	-0.0142(2)	0.0130(2)	0.0056 (2)
N1	0.0311(19)	0.0122(3) 0.031(2)	0.0321(18)	0.0112(2) 0.0054(15)	0.0130(2)	0.0000(2) 0.0043(15)
N2	0.040 (2)	0.031(2) 0.042(3)	0.052(2)	-0.0019(19)	0.0068 (18)	0.0108 (19)
N3	0.0275(19)	0.035(2)	0.040(2)	0.0112 (16)	0.0045 (15)	0.0110 (17)
01	0.0476 (19)	0.0202 (15)	0.0448 (17)	0.0028 (14)	0.0220 (14)	0.0030 (13)
02	0.0227 (13)	0.0244 (15)	0.0325 (14)	-0.0013 (11)	0.0037 (11)	0.0024 (11)
						()
Geometric param	neters (Å, °)					
C1—O1		1.277 (5)	C10'—	N2	1.376	(11)
C1—C2		1.434 (6)	C10'—	C11	1.459	(12)
C1—C6		1.440 (5)	C10'—	H10C	0.970	0
C2—C3		1.370 (6)	C10'—	H10D	0.970	0
C2—Br1		1.892 (4)	C11—1	N3	1.460	(6)
C3—C4		1.408 (6)	C11—I	H11A	0.970	0
С3—Н3		0.9300	C11—I	H11B	0.970	0
C4—C5		1.360 (6)	C12—1	N3	1.293	(6)
C4—Br2		1.900 (4)	C12—0	213	1.411	(6)
C5—C6		1.419 (6)	C12—I	H12	0.930	0
С5—Н5		0.9300	C13—0	C18	1.409	(6)
C6—C7		1.433 (6)	C13—0	C14	1.449	(5)
C7—N1		1.290 (6)	C14—0	02	1.270	(5)
С7—Н7		0.9300	C14—0	215	1.433	(5)
C8—N1		1.463 (5)	C15—0	216	1.369	(6)
С8—С9		1.518 (6)	C15—I	Br3	1.898	(4)
C8—H8A		0.9700	C16—0	C17	1.412	(6)
C8—H8B		0.9700	C16—I	416	0.930	0
C9—N2		1.445 (6)	C17—0	C18	1.363	(6)
С9—Н9А		0.9700	C17—I	Br4	1.902	(4)
С9—Н9В		0.9700	C18—I	118	0.930	0
C10-C11		1.359 (12)	N1—H	1A	0.97 (6)
C10—N2		1.526 (11)	N2—H	2	0.860	0
C10—H10A		0.9700	N3—H	3A	0.87 (6)
C10—H10B		0.9700				
O1—C1—C2		122.7 (4)	H10C-	-C10'-H10D	107.3	
O1—C1—C6		122.6 (4)	C10—0	C11—C10'	43.6 (6)
C2—C1—C6		114.7 (4)	C10—0	C11—N3	112.1	(7)
C3—C2—C1		123.4 (4)	C10'—	C11—N3	118.1	(6)
C3—C2—Br1		119.1 (3)	C10—0	C11—H11A	109.2	
C1—C2—Br1		117.5 (3)	C10'—	C11—H11A	66.7	
C2—C3—C4		119.4 (4)	N3—C	11—H11A	109.2	
С2—С3—Н3		120.3	C10—0	C11—H11B	109.2	
С4—С3—Н3		120.3	C10'—	С11—Н11В	131.6	
C5—C4—C3		121.0 (4)	N3—C	11—H11B	109.2	
C5—C4—Br2		121.9 (3)	H11A-	C11H11B	107.9	
C3—C4—Br2		117.0 (3)	N3—C	12—C13	123.8	(4)
C4—C5—C6		119.9 (4)	N3—C	12—H12	118.1	
C4—C5—H5		120.0	C13—0	С12—Н12	118.1	
С6—С5—Н5		120.0	C18—0	C13—C12	119.1	(3)

C5—C6—C7	118.9 (4)	C18—C13—C14	121.5 (4)
C5—C6—C1	121.5 (4)	C12—C13—C14	119.4 (4)
C7—C6—C1	119.5 (4)	O2—C14—C15	123.3 (3)
N1—C7—C6	121.0 (4)	O2—C14—C13	122.4 (4)
N1—C7—H7	119.5	C15-C14-C13	114.3 (3)
С6—С7—Н7	119.5	C16—C15—C14	123.9 (4)
N1—C8—C9	111.0 (4)	C16—C15—Br3	119.6 (3)
N1—C8—H8A	109.4	C14—C15—Br3	116.5 (3)
С9—С8—Н8А	109.4	C15—C16—C17	119.1 (4)
N1—C8—H8B	109.4	C15-C16-H16	120.5
С9—С8—Н8В	109.4	С17—С16—Н16	120.5
H8A—C8—H8B	108.0	C18—C17—C16	120.9 (4)
N2—C9—C8	111.6 (4)	C18—C17—Br4	119.8 (3)
N2—C9—H9A	109.3	C16—C17—Br4	119.3 (3)
С8—С9—Н9А	109.3	C17—C18—C13	120.3 (4)
N2—C9—H9B	109.3	C17—C18—H18	119.8
С8—С9—Н9В	109.3	C13—C18—H18	119.8
H9A—C9—H9B	108.0	C7—N1—C8	125.1 (4)
C11—C10—N2	113.3 (8)	C7—N1—H1A	112 (3)
C11-C10-H10A	108.9	C8—N1—H1A	123 (3)
N2-C10-H10A	108.9	C10'—N2—C9	139.4 (6)
C11—C10—H10B	108.9	C10'—N2—C10	42.1 (6)
N2-C10-H10B	108.9	C9—N2—C10	106.8 (5)
H10A—C10—H10B	107.7	C10'—N2—H2	89.2
N2—C10'—C11	116.4 (8)	C9—N2—H2	126.6
N2—C10'—H10C	108.2	C10—N2—H2	126.6
C11—C10'—H10C	108.2	C12—N3—C11	124.8 (4)
N2—C10'—H10D	108.2	C12—N3—H3A	111 (4)
C11—C10'—H10D	108.2	C11—N3—H3A	123 (4)
O1—C1—C2—C3	179.1 (4)	C12—C13—C14—O2	-1.1 (6)
C6—C1—C2—C3	-0.1 (6)	C18—C13—C14—C15	-1.0 (5)
O1-C1-C2-Br1	-0.1 (5)	C12—C13—C14—C15	177.5 (3)
C6—C1—C2—Br1	-179.3 (3)	O2-C14-C15-C16	178.9 (4)
C1—C2—C3—C4	-0.3 (6)	C13-C14-C15-C16	0.3 (5)
Br1—C2—C3—C4	178.9 (3)	O2—C14—C15—Br3	-1.6 (5)
C2—C3—C4—C5	0.8 (6)	C13—C14—C15—Br3	179.8 (3)
C2—C3—C4—Br2	-178.4 (3)	C14—C15—C16—C17	0.2 (6)
C3—C4—C5—C6	-1.0 (6)	Br3-C15-C16-C17	-179.3 (3)
Br2C4C5C6	178.2 (3)	C15—C16—C17—C18	-0.1 (6)
C4—C5—C6—C7	-176.5 (4)	C15-C16-C17-Br4	-179.5 (3)
C4—C5—C6—C1	0.7 (6)	C16-C17-C18-C13	-0.6 (6)
O1—C1—C6—C5	-179.3 (4)	Br4—C17—C18—C13	178.8 (3)
C2—C1—C6—C5	-0.1 (5)	C12-C13-C18-C17	-177.4 (4)
O1—C1—C6—C7	-2.2 (6)	C14—C13—C18—C17	1.2 (6)
C2-C1-C6-C7	177.0 (3)	C6—C7—N1—C8	-178.4 (4)
C5—C6—C7—N1	177.4 (4)	C9—C8—N1—C7	-120.1 (5)
C1—C6—C7—N1	0.2 (6)	C11—C10'—N2—C9	-101.3 (11)
N1	72.0 (5)	C11—C10'—N2—C10	-50.4 (9)
N2-C10-C11-C10'	-47.6 (8)	C8—C9—N2—C10'	136.5 (8)

N2-C10-C11-N3	60.4 (10)	C8—C9—N2—C10	103.6 (6)
N2-C10'-C11-C10	57.2 (10)	C11—C10—N2—C10'	53.8 (10)
N2-C10'-C11-N3	-36.2 (12)	C11—C10—N2—C9	-158.1 (7)
N3-C12-C13-C18	-178.6 (4)	C13-C12-N3-C11	-175.4 (5)
N3-C12-C13-C14	2.9 (6)	C10-C11-N3-C12	158.2 (7)
C18—C13—C14—O2	-179.6 (3)	C10'-C11-N3-C12	-153.8 (7)

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	H…A	$D \cdots A$	D—H··· A
N1—H1A···O1	0.97 (6)	1.70 (6)	2.553 (5)	144 (5)
N3—H3A…O2	0.87 (6)	1.84 (6)	2.597 (4)	144 (5)



