

## 1,4,10,13-Tetraoxa-7,16-diazoniacyclo-octadecane bis(1*H*-pyrrole-2-carboxylate)

Fanglei Zeng and Zhenming Yin\*

Tianjin Key Laboratory of Structure and Performance for Functional Molecules, Key Laboratory of Inorganic-Organic Hybrid Functional Material Chemistry, Ministry of Education, College of Chemistry, Tianjin Normal University, Tianjin 300387, People's Republic of China

Correspondence e-mail: tjyinzm@aliyun.com

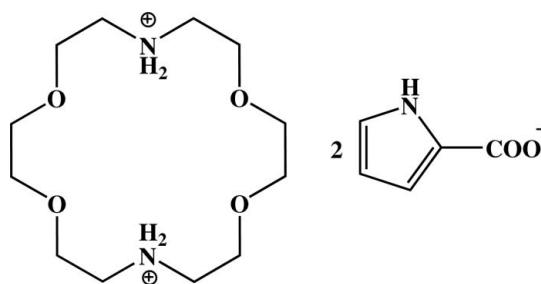
Received 24 May 2013; accepted 11 June 2013

Key indicators: single-crystal X-ray study;  $T = 294\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$ ;  $R$  factor = 0.042;  $wR$  factor = 0.113; data-to-parameter ratio = 14.9.

In the title salt,  $\text{C}_{12}\text{H}_{28}\text{N}_2\text{O}_4^{2+} \cdot 2\text{C}_5\text{H}_4\text{NO}_2^-$ , the 1,4,10,13-tetraoxa-7,16-diazacyclooctadecane dication possesses inversion symmetry. In the crystal, the pyrrole-carboxylate anions are linked via pairs of  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds, forming inversion dimers. These dimers are linked by the dications, via  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds, forming chains propagating along [110].

### Related literature

For background to hydrogen-bonded supramolecular assemblies, see: Burrows (2004). For the hydrogen-bonded assemblies of pyrrole-based structures, see: Wang & Yin (2007); Yin & Li (2006); Cui *et al.* (2009); Li *et al.* (2012).



### Experimental

#### Crystal data

$\text{C}_{12}\text{H}_{28}\text{N}_2\text{O}_4^{2+} \cdot 2\text{C}_5\text{H}_4\text{NO}_2^-$	$\gamma = 77.824 (4)^\circ$
$M_r = 484.55$	$V = 614.8 (3)\text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 1$
$a = 7.8963 (19)\text{ \AA}$	Mo $K\alpha$ radiation
$b = 9.164 (2)\text{ \AA}$	$\mu = 0.10\text{ mm}^{-1}$
$c = 9.244 (2)\text{ \AA}$	$T = 294\text{ K}$
$\alpha = 73.028 (4)^\circ$	$0.24 \times 0.22 \times 0.18\text{ mm}$
$\beta = 76.547 (4)^\circ$	

#### Data collection

Bruker SMART CCD area-detector diffractometer	3484 measured reflections
Absorption correction: multi-scan ( <i>SADABS</i> ; Bruker, 1997)	2471 independent reflections
$T_{\min} = 0.970$ , $T_{\max} = 0.982$	1695 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.017$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.113$	$\Delta\rho_{\text{max}} = 0.14\text{ e \AA}^{-3}$
$S = 1.03$	$\Delta\rho_{\text{min}} = -0.26\text{ e \AA}^{-3}$
2471 reflections	
166 parameters	

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{N}2-\text{H}2 \cdots \text{O}4^{\text{i}}$	0.877 (19)	1.94 (2)	2.7741 (19)	158.0 (17)
$\text{N}1-\text{H}1B \cdots \text{O}3^{\text{ii}}$	0.91 (2)	2.01 (2)	2.8167 (19)	147.4 (16)
$\text{N}1-\text{H}1A \cdots \text{O}4$	0.94 (2)	2.489 (19)	3.137 (2)	125.9 (14)
$\text{N}1-\text{H}1A \cdots \text{O}3$	0.94 (2)	1.81 (2)	2.7452 (19)	171.1 (17)

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

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); 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*.

We sincerely thank the Natural Science Foundation of China for financial support (NSFC No. 21172174).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FF2108).

### References

- Bruker (1997). *SADABS, SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Burrows, A. D. (2004). *Struct. Bond.* **108**, 55–96.
- Cui, Y., Yin, Z., Dong, L. & He, J. (2009). *J. Mol. Struct.* **938**, 322–327.
- Li, C., Zhang, G. & Yin, Z. (2012). *Acta Cryst. E* **68**, m323.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Wang, W.-Y. & Yin, Z.-M. (2007). *Acta Cryst. E* **63**, o2737–o2738.
- Yin, Z. & Li, Z. (2006). *Tetrahedron Lett.* **47**, 7875–7879.

# supporting information

*Acta Cryst.* (2013). E69, o1108 [https://doi.org/10.1107/S1600536813016176]

## 1,4,10,13-Tetraoxa-7,16-diaza-cyclooctadecane bis(1*H*-pyrrole-2-carboxylate)

Fanglei Zeng and Zhenming Yin

### S1. Comment

Hydrogen-bond-mediated self-assembly represents an area of considerable current interest (Burrows, 2004). It has recently been found that pyrrole-based entities are also capable of undergoing self-assembly through hydrogen bonds, especially in the solid state. In our previous works, we have reported the hydrogen-bonded assemblies of 4-pyridylmethyl 1*H*-pyrrole-2-carboxylate (Wang & Yin, 2007) and some other pyrrole-based compounds (Yin & Li, 2006; Cui *et al.* 2009; Li *et al.* 2012) in the solid state. Here we report the self-assembly of the title compound, (I), *via* conventional N—H···O hydrogen bonds.

The molecular structure of (I) is shown in Fig.1. In the solid state, the compound adopts central symmetrical conformation. Each pyrrole-2-carboxylate group is planar and interact with protonated amino group through two charge assisted N—H···O hydrogen bonds.

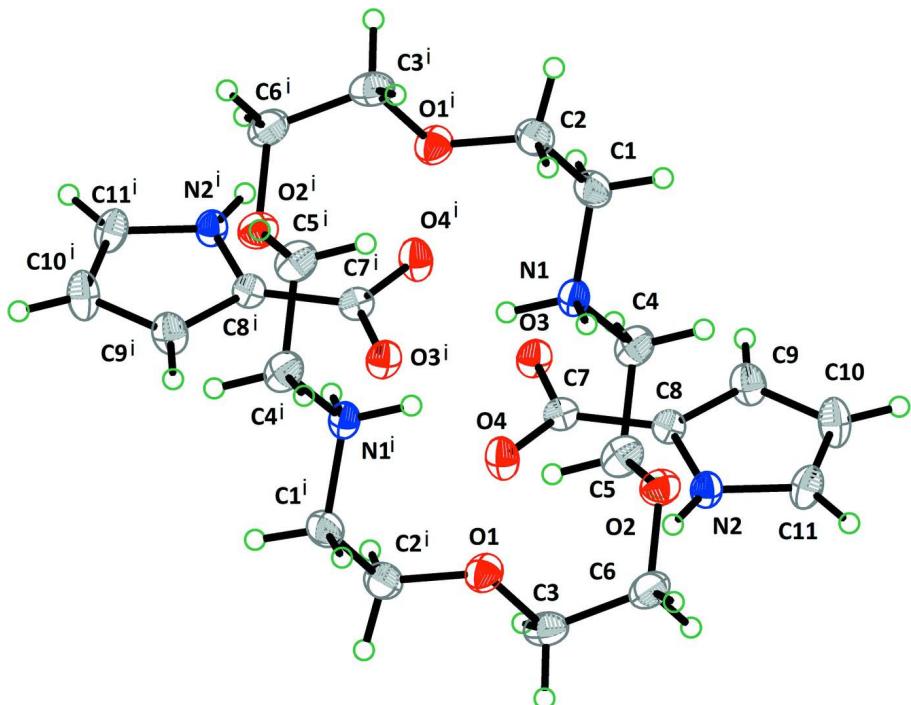
In the crystal structure, the molecules of (I) are held together by a pair of N—H···O hydrogen bonds between the pyrrole and carbonyl groups (Fig.2). Consequently, the molecules of (I) form a one-dimensional infinite chain structure.

### S2. Experimental

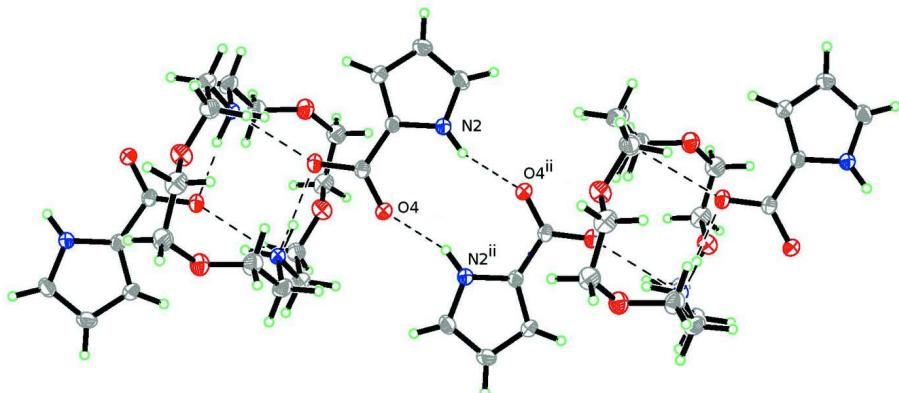
1,4,10,13-Tetraoxa-7,16-diaza-cyclooctadecane (262 mg, 100 mmol), 1*H*-pyrrole-2-carboxylic acid (222 mg, 200 mmol) were added to alcohol (20 ml), and the mixture was stirred in room temperature for 1 h. The solution was then evaporated and afforded the title compound (colorless crystals, 387 mg, 70%).

### S3. Refinement

The N-bound H atoms were located in a difference map and refined freely. Other H atoms were positioned geometrically (C—H = 0.93 or 0.97 Å) and refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

**Figure 1**

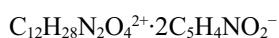
The molecular structure of (I), with the numbering scheme and 30% probability displacement ellipsoids. [Symmetry code: (i) 1-x, 1-y, 1-z]

**Figure 2**

The dimer of molecules of (I) connected by N—H···O hydrogen bonds (dashed lines). [Symmetry code: (ii) -x + 2, -y, -z + 1]

### 1,4,10,13-Tetraoxa-7,16-diazoniacyclooctadecane bis(1*H*-pyrrole-2-carboxylate)

#### Crystal data



$$M_r = 484.55$$

Triclinic,  $P\bar{1}$

Hall symbol: -P 1

$$a = 7.8963 (19) \text{ \AA}$$

$$b = 9.164 (2) \text{ \AA}$$

$$c = 9.244 (2) \text{ \AA}$$

$$\alpha = 73.028 (4)^\circ$$

$$\beta = 76.547 (4)^\circ$$

$$\gamma = 77.824 (4)^\circ$$

$$V = 614.8 (3) \text{ \AA}^3$$

$$Z = 1$$

$F(000) = 260$   
 $D_x = 1.309 \text{ Mg m}^{-3}$   
Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$   
Cell parameters from 1288 reflections  
 $\theta = 2.7\text{--}25.6^\circ$

$\mu = 0.10 \text{ mm}^{-1}$   
 $T = 294 \text{ K}$   
Block, colourless  
 $0.24 \times 0.22 \times 0.18 \text{ mm}$

#### Data collection

Bruker SMART CCD area-detector  
diffractometer  
Radiation source: fine-focus sealed tube  
Graphite monochromator  
phi and  $\omega$  scans  
Absorption correction: multi-scan  
(SADABS; Bruker, 1997)  
 $T_{\min} = 0.970$ ,  $T_{\max} = 0.982$

3484 measured reflections  
2471 independent reflections  
1695 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.017$   
 $\theta_{\max} = 26.3^\circ$ ,  $\theta_{\min} = 2.3^\circ$   
 $h = -9 \rightarrow 9$   
 $k = -11 \rightarrow 7$   
 $l = -11 \rightarrow 11$

#### Refinement

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.042$   
 $wR(F^2) = 0.113$   
 $S = 1.03$   
2471 reflections  
166 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0587P)^2 + 0.0353P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.14 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.26 \text{ e \AA}^{-3}$

#### 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 ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.5212 (2)	0.36442 (16)	0.71663 (16)	0.0346 (3)
H1A	0.592 (3)	0.369 (2)	0.618 (2)	0.051 (5)*
H1B	0.434 (3)	0.447 (2)	0.706 (2)	0.049 (5)*
N2	0.99820 (18)	0.10017 (16)	0.27227 (16)	0.0346 (3)
H2	1.027 (2)	0.022 (2)	0.348 (2)	0.048 (5)*
O1	0.41888 (15)	0.19373 (14)	0.53943 (14)	0.0462 (3)
O2	0.55409 (16)	0.64120 (13)	0.77534 (14)	0.0462 (3)
O3	0.72871 (15)	0.40741 (12)	0.42918 (12)	0.0391 (3)
O4	0.85227 (16)	0.17742 (13)	0.55299 (13)	0.0439 (3)
C1	0.4410 (3)	0.2209 (2)	0.7790 (2)	0.0453 (5)
H1C	0.3761	0.2187	0.8823	0.054*

H1D	0.5342	0.1326	0.7857	0.054*
C2	0.3194 (2)	0.2067 (2)	0.6832 (2)	0.0457 (5)
H2A	0.2624	0.1163	0.7332	0.055*
H2B	0.2290	0.2970	0.6705	0.055*
C3	0.3199 (3)	0.1628 (2)	0.4441 (2)	0.0495 (5)
H3A	0.2031	0.2233	0.4538	0.059*
H3B	0.3075	0.0544	0.4754	0.059*
C4	0.6376 (3)	0.3752 (2)	0.8169 (2)	0.0482 (5)
H4A	0.7369	0.2923	0.8172	0.058*
H4B	0.5729	0.3636	0.9216	0.058*
C5	0.7031 (2)	0.5272 (2)	0.7613 (2)	0.0486 (5)
H5A	0.7851	0.5313	0.8230	0.058*
H5B	0.7627	0.5427	0.6548	0.058*
C6	0.5862 (3)	0.7961 (2)	0.7188 (2)	0.0507 (5)
H6A	0.6552	0.8153	0.7836	0.061*
H6B	0.4742	0.8641	0.7278	0.061*
C7	0.8236 (2)	0.27649 (18)	0.43185 (18)	0.0318 (4)
C8	0.9026 (2)	0.24102 (18)	0.28271 (18)	0.0313 (4)
C9	0.8985 (2)	0.3296 (2)	0.13557 (19)	0.0432 (4)
H9	0.8426	0.4309	0.1075	0.052*
C10	0.9934 (3)	0.2405 (2)	0.0351 (2)	0.0505 (5)
H10	1.0120	0.2716	-0.0718	0.061*
C11	1.0535 (2)	0.0996 (2)	0.1224 (2)	0.0437 (5)
H11	1.1209	0.0174	0.0854	0.052*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0400 (8)	0.0321 (8)	0.0289 (8)	0.0031 (7)	-0.0085 (7)	-0.0080 (6)
N2	0.0376 (8)	0.0328 (8)	0.0310 (8)	0.0017 (6)	-0.0068 (6)	-0.0091 (6)
O1	0.0407 (7)	0.0546 (8)	0.0490 (8)	-0.0112 (6)	-0.0072 (6)	-0.0200 (6)
O2	0.0475 (7)	0.0390 (7)	0.0528 (8)	-0.0051 (6)	-0.0082 (6)	-0.0147 (6)
O3	0.0416 (7)	0.0323 (6)	0.0401 (7)	0.0072 (5)	-0.0095 (5)	-0.0115 (5)
O4	0.0509 (8)	0.0387 (7)	0.0324 (6)	0.0094 (6)	-0.0086 (6)	-0.0049 (5)
C1	0.0548 (12)	0.0358 (10)	0.0393 (10)	-0.0065 (8)	-0.0051 (9)	-0.0034 (8)
C2	0.0458 (11)	0.0414 (10)	0.0467 (11)	-0.0092 (8)	-0.0022 (9)	-0.0096 (8)
C3	0.0525 (12)	0.0437 (11)	0.0602 (12)	-0.0163 (9)	-0.0190 (10)	-0.0121 (9)
C4	0.0646 (13)	0.0408 (11)	0.0425 (10)	0.0004 (9)	-0.0267 (9)	-0.0086 (8)
C5	0.0481 (11)	0.0492 (11)	0.0541 (12)	-0.0011 (9)	-0.0225 (9)	-0.0157 (9)
C6	0.0632 (13)	0.0426 (11)	0.0542 (12)	-0.0095 (9)	-0.0155 (10)	-0.0196 (9)
C7	0.0281 (9)	0.0310 (9)	0.0370 (9)	-0.0019 (7)	-0.0086 (7)	-0.0092 (7)
C8	0.0283 (8)	0.0309 (8)	0.0337 (9)	-0.0002 (7)	-0.0077 (7)	-0.0082 (7)
C9	0.0443 (10)	0.0405 (10)	0.0363 (10)	0.0057 (8)	-0.0095 (8)	-0.0040 (8)
C10	0.0497 (11)	0.0651 (13)	0.0304 (9)	0.0053 (10)	-0.0089 (8)	-0.0115 (9)
C11	0.0425 (10)	0.0499 (11)	0.0404 (10)	0.0052 (8)	-0.0076 (8)	-0.0225 (9)

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

N1—C1	1.483 (2)	C3—C6 <sup>i</sup>	1.492 (3)
N1—C4	1.484 (2)	C3—H3A	0.9700
N1—H1A	0.94 (2)	C3—H3B	0.9700
N1—H1B	0.91 (2)	C4—C5	1.495 (3)
N2—C11	1.354 (2)	C4—H4A	0.9700
N2—C8	1.369 (2)	C4—H4B	0.9700
N2—H2	0.877 (19)	C5—H5A	0.9700
O1—C2	1.402 (2)	C5—H5B	0.9700
O1—C3	1.424 (2)	C6—C3 <sup>i</sup>	1.492 (3)
O2—C5	1.408 (2)	C6—H6A	0.9700
O2—C6	1.417 (2)	C6—H6B	0.9700
O3—C7	1.2701 (18)	C7—C8	1.474 (2)
O4—C7	1.2519 (18)	C8—C9	1.369 (2)
C1—C2	1.496 (2)	C9—C10	1.397 (2)
C1—H1C	0.9700	C9—H9	0.9300
C1—H1D	0.9700	C10—C11	1.361 (3)
C2—H2A	0.9700	C10—H10	0.9300
C2—H2B	0.9700	C11—H11	0.9300
C1—N1—C4	111.26 (13)	C5—C4—H4A	109.5
C1—N1—H1A	112.0 (11)	N1—C4—H4B	109.5
C4—N1—H1A	106.5 (11)	C5—C4—H4B	109.5
C1—N1—H1B	109.1 (12)	H4A—C4—H4B	108.1
C4—N1—H1B	111.2 (12)	O2—C5—C4	106.59 (16)
H1A—N1—H1B	106.7 (16)	O2—C5—H5A	110.4
C11—N2—C8	109.53 (15)	C4—C5—H5A	110.4
C11—N2—H2	122.7 (12)	O2—C5—H5B	110.4
C8—N2—H2	127.7 (12)	C4—C5—H5B	110.4
C2—O1—C3	113.17 (14)	H5A—C5—H5B	108.6
C5—O2—C6	115.71 (15)	O2—C6—C3 <sup>i</sup>	114.91 (15)
N1—C1—C2	113.14 (14)	O2—C6—H6A	108.5
N1—C1—H1C	109.0	C3 <sup>i</sup> —C6—H6A	108.5
C2—C1—H1C	109.0	O2—C6—H6B	108.5
N1—C1—H1D	109.0	C3 <sup>i</sup> —C6—H6B	108.5
C2—C1—H1D	109.0	H6A—C6—H6B	107.5
H1C—C1—H1D	107.8	O4—C7—O3	123.76 (15)
O1—C2—C1	108.26 (15)	O4—C7—C8	118.96 (13)
O1—C2—H2A	110.0	O3—C7—C8	117.28 (14)
C1—C2—H2A	110.0	N2—C8—C9	107.10 (14)
O1—C2—H2B	110.0	N2—C8—C7	122.19 (14)
C1—C2—H2B	110.0	C9—C8—C7	130.70 (15)
H2A—C2—H2B	108.4	C8—C9—C10	107.78 (16)
O1—C3—C6 <sup>i</sup>	108.78 (15)	C8—C9—H9	126.1
O1—C3—H3A	109.9	C10—C9—H9	126.1
C6 <sup>i</sup> —C3—H3A	109.9	C11—C10—C9	107.46 (15)
O1—C3—H3B	109.9	C11—C10—H10	126.3

C6 <sup>i</sup> —C3—H3B	109.9	C9—C10—H10	126.3
H3A—C3—H3B	108.3	N2—C11—C10	108.13 (15)
N1—C4—C5	110.60 (14)	N2—C11—H11	125.9
N1—C4—H4A	109.5	C10—C11—H11	125.9
C4—N1—C1—C2	178.78 (15)	O4—C7—C8—N2	3.8 (2)
C3—O1—C2—C1	173.64 (14)	O3—C7—C8—N2	-175.99 (14)
N1—C1—C2—O1	64.80 (19)	O4—C7—C8—C9	-176.92 (17)
C2—O1—C3—C6 <sup>i</sup>	160.84 (15)	O3—C7—C8—C9	3.3 (3)
C1—N1—C4—C5	-174.38 (16)	N2—C8—C9—C10	-0.1 (2)
C6—O2—C5—C4	-176.05 (14)	C7—C8—C9—C10	-179.46 (16)
N1—C4—C5—O2	63.76 (19)	C8—C9—C10—C11	0.0 (2)
C5—O2—C6—C3 <sup>i</sup>	54.4 (2)	C8—N2—C11—C10	-0.1 (2)
C11—N2—C8—C9	0.12 (19)	C9—C10—C11—N2	0.1 (2)
C11—N2—C8—C7	179.58 (15)		

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

#### Hydrogen-bond geometry ( $\text{\AA}$ , $^\circ$ )

$D\text{—H}\cdots A$	$D\text{—H}$	$\text{H}\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N2—H2 $\cdots$ O4 <sup>ii</sup>	0.877 (19)	1.94 (2)	2.7741 (19)	158.0 (17)
N1—H1B $\cdots$ O3 <sup>i</sup>	0.91 (2)	2.01 (2)	2.8167 (19)	147.4 (16)
N1—H1A $\cdots$ O4	0.94 (2)	2.489 (19)	3.137 (2)	125.9 (14)
N1—H1A $\cdots$ O3	0.94 (2)	1.81 (2)	2.7452 (19)	171.1 (17)

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