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3-Carboxypyrazino[2,3-*f*][1,10]-phenanthrolin-9-ium-2-carboxylateXiao-Ning Zhang,^a Feng Fu,^{a*} Dong-Sheng Li^b and Cai-Xia Meng^a

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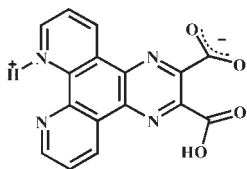
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.062; wR factor = 0.177; data-to-parameter ratio = 10.1.

In the title zwitterionic compound, $\text{C}_{16}\text{H}_8\text{N}_4\text{O}_4$, the dihedral angle between the carboxyl and carboxylate groups is $72.14(2)^\circ$. In the crystal, molecules are linked by strong intermolecular $\text{O}-\text{H}\cdots\text{O}^-$ and $\text{N}^+-\text{H}\cdots\text{O}^-$ hydrogen bonds into double chains extended along [001]. These chains are additionally stabilized by $\pi-\pi$ stacking interactions between the pyridine and benzene rings [centroid-centroid distance = $3.5542(8)$ Å].

Related literature

For coordination compounds of the title ligand, see: Weng *et al.* (2009).



Experimental

Crystal data

$\text{C}_{16}\text{H}_8\text{N}_4\text{O}_4$
 $M_r = 320.26$
Triclinic, $P\bar{1}$

$a = 7.302(2)$ Å
 $b = 9.662(3)$ Å
 $c = 10.726(3)$ Å

$\alpha = 63.564(3)^\circ$
 $\beta = 71.386(3)^\circ$
 $\gamma = 78.283(4)^\circ$
 $V = 640.5(3)$ Å³
 $Z = 2$

Mo $K\alpha$ radiation
 $\mu = 0.12$ mm⁻¹
 $T = 293$ K
 $0.32 \times 0.21 \times 0.15$ mm

Data collection

Bruker SMART CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.969$, $T_{\max} = 0.982$

4652 measured reflections
2243 independent reflections
1268 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.037$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.177$
 $S = 1.05$
2243 reflections
221 parameters
1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.38$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.31$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O3}-\text{H3}\cdots\text{O1}^{\text{i}}$	0.82	1.82	2.638 (4)	171
$\text{N2}-\text{H2}\cdots\text{O2}^{\text{ii}}$	0.81 (2)	1.87 (3)	2.638 (4)	159 (5)

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

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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2240).

References

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Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
Weng, Z.-H., Liu, D.-C., Chen, Z.-L., Zou, H.-H., Qin, S.-N. & Liang, F.-P. (2009). *Cryst. Growth Des.* **9**, 4163–4170.

supporting information

Acta Cryst. (2010). E66, o740 [doi:10.1107/S1600536810007361]

3-Carboxypyrazino[2,3-*f*][1,10]phenanthroline-9-ium-2-carboxylate

Xiao-Ning Zhang, Feng Fu, Dong-Sheng Li and Cai-Xia Meng

S1. Comment

[2,3-*f*]Pyrazino[1,10]phenanthroline-2,3-dicarboxylic acid (H₂ppdb) is a multidentate O/N-donor ligand, which was rarely used up to date (Weng *et al.*, 2009). Owing to the presence of two carboxylic groups and a large conjugated π system, H₂ppdb hoards the recognition information for the formation of interesting supramolecular structures. We attempted to synthesize a Tm^{III} complex with the H₂ppdb ligand in hydrothermal synthesis conditions however we were unsuccessful. Instead of crystals of the complex, single crystals of the H₂ppdb were isolated and the structure of H₂ppdb is reported here.

The molecular structure of title compound is shown in Fig. 1. One of the carboxyl groups is deprotonated and the proton is transferred to the phenanthroline fragment. Intermolecular hydrogen-bonding and π - π stacking interactions are the most important features of the title compound. O—H \cdots O and N—H \cdots O hydrogen bonds (Table 1), and π - π stacking interactions between the pyridine rings and phenyl rings from the neighboring H₂ppdb molecules [centroid–centroid distances = 3.5542 (8) Å] link the molecules into a double chain extending along the *c* axis. The double chains are further connected by weak C—H \cdots O hydrogen bonds involving the carboxyl oxygen and the C—H groups adjacent to the pyridine rings and π - π stacking interactions as indicated by a center-to-center distance of 3.7402 (7) Å, forming a three-dimensional supramolecular network (Fig.2).

S2. Experimental

All chemicals were of analytical reagent grade, commercially available and used without further purification. A mixture of H₂ppdb (0.1 mmol, 0.0320 g), Tm₂O₃ (0.05 mmol, 0.0193 g) was adjusted to pH 1.5 with HNO₃ (0.2 mol/L). It was then sealed in a 25 ml Teflon-lined stainless steel reactor together with water (10 ml), heated at 140 °C for 4 days, and then cooled slowly to room temperature, to furnish yellow crystals.

S3. Refinement

The H atom attached to N atom was located in a difference Fourier map and refined with the N—H distance restrained to 0.82 (2) Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{N})$. The other H atoms were placed in idealized positions, with C—H = 0.93 and O—H = 0.82 Å, and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

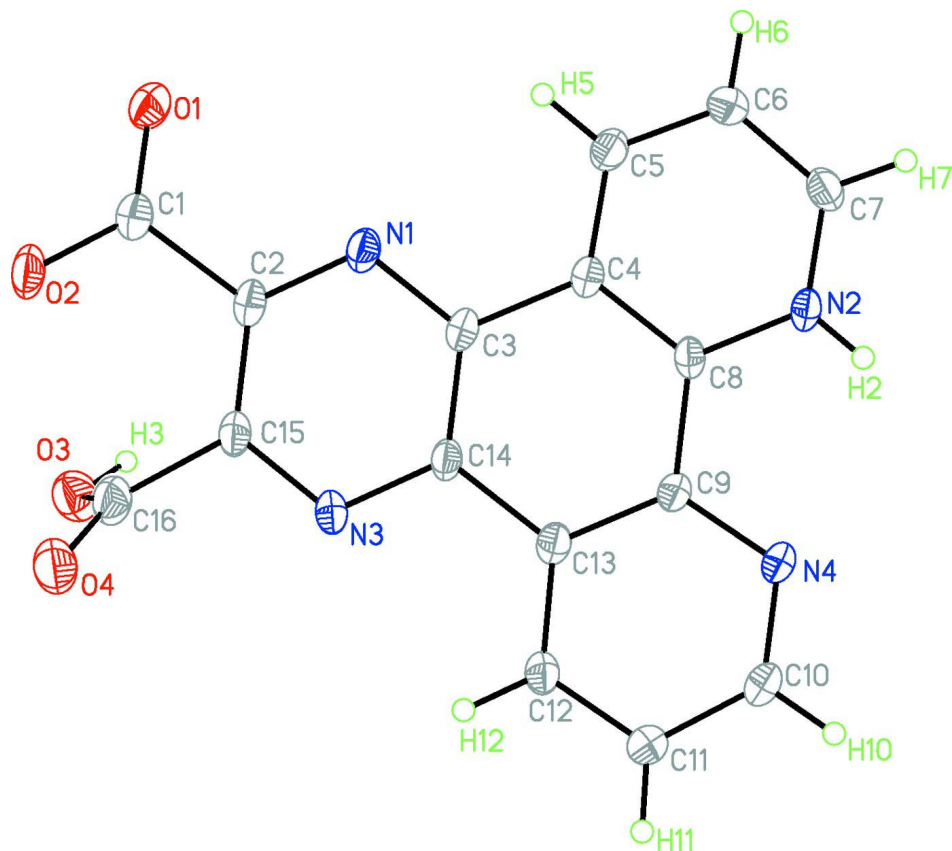


Figure 1

The structure of the title compound with the atom-numbering scheme showing displacement ellipsoids at the 50% probability level.

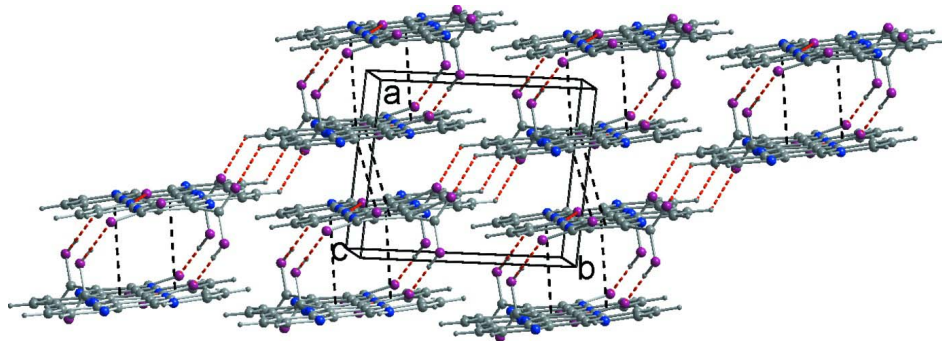


Figure 2

Three-dimensional supramolecular network formed by hydrogen bonds and π - π stacking interactions.

3-Carboxypyrazino[2,3-*f*][1,10]phenanthroline-9-ium-2-carboxylate

Crystal data

$C_{16}H_8N_4O_4$

$M_r = 320.26$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 7.302 (2) \text{ \AA}$

$b = 9.662 (3) \text{ \AA}$

$c = 10.726 (3) \text{ \AA}$

$\alpha = 63.564 (3)^\circ$

$\beta = 71.386 (3)^\circ$
 $\gamma = 78.283 (4)^\circ$
 $V = 640.5 (3) \text{ \AA}^3$
 $Z = 2$
 $F(000) = 328$
 $D_x = 1.661 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1426 reflections
 $\theta = 2.5\text{--}25.0^\circ$
 $\mu = 0.12 \text{ mm}^{-1}$
 $T = 293 \text{ K}$
 Prism, colorless
 $0.32 \times 0.21 \times 0.15 \text{ mm}$

Data collection

Bruker SMART CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 phi and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.969$, $T_{\max} = 0.982$

4652 measured reflections
 2243 independent reflections
 1268 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.037$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.5^\circ$
 $h = -8 \rightarrow 8$
 $k = -11 \rightarrow 11$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.177$
 $S = 1.05$
 2243 reflections
 221 parameters
 1 restraint
 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
 $w = 1/[\sigma^2(F_o^2) + (0.0802P)^2 + 0.2P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.38 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.31 \text{ e \AA}^{-3}$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.1681 (4)	-0.2077 (3)	0.4832 (3)	0.0468 (8)
O2	0.2929 (4)	-0.0009 (3)	0.2840 (3)	0.0473 (8)
O3	0.0845 (4)	0.3107 (4)	0.2623 (3)	0.0529 (8)
H3	0.0010	0.2742	0.3381	0.079*
O4	0.4003 (5)	0.3285 (4)	0.1966 (3)	0.0551 (9)
N1	0.2349 (4)	-0.0766 (4)	0.6468 (3)	0.0324 (8)
N3	0.2887 (5)	0.2420 (4)	0.5116 (3)	0.0374 (8)
N2	0.2403 (5)	-0.0825 (4)	1.0943 (3)	0.0317 (8)
N4	0.3104 (4)	0.2237 (4)	0.9626 (3)	0.0336 (8)

C4	0.2334 (5)	-0.0835 (4)	0.8748 (4)	0.0285 (9)
C9	0.2928 (5)	0.1575 (4)	0.8803 (4)	0.0285 (9)
C14	0.2812 (5)	0.1585 (4)	0.6532 (4)	0.0291 (9)
C15	0.2689 (5)	0.1654 (5)	0.4411 (4)	0.0348 (10)
C6	0.1733 (5)	-0.3119 (4)	1.0965 (4)	0.0352 (10)
H6	0.1440	-0.4150	1.1477	0.042*
C8	0.2571 (5)	-0.0055 (4)	0.9502 (4)	0.0275 (9)
C1	0.2339 (5)	-0.0776 (5)	0.4193 (4)	0.0359 (10)
C3	0.2491 (5)	0.0009 (4)	0.7207 (4)	0.0284 (9)
C13	0.3009 (5)	0.2401 (4)	0.7332 (4)	0.0302 (9)
C12	0.3264 (6)	0.3985 (5)	0.6710 (4)	0.0364 (10)
H12	0.3301	0.4574	0.5742	0.044*
C7	0.2011 (5)	-0.2294 (5)	1.1653 (4)	0.0357 (10)
H7	0.1920	-0.2783	1.2637	0.043*
C10	0.3366 (6)	0.3728 (5)	0.8992 (4)	0.0392 (10)
H10	0.3499	0.4194	0.9550	0.047*
C5	0.1900 (5)	-0.2382 (4)	0.9509 (4)	0.0343 (9)
H5	0.1722	-0.2920	0.9027	0.041*
C2	0.2468 (5)	0.0059 (5)	0.5071 (4)	0.0319 (9)
C11	0.3458 (6)	0.4662 (5)	0.7535 (4)	0.0419 (11)
H11	0.3647	0.5712	0.7141	0.050*
C16	0.2599 (7)	0.2711 (5)	0.2886 (5)	0.0442 (11)
H2	0.250 (7)	-0.035 (5)	1.138 (4)	0.066*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.062 (2)	0.0476 (19)	0.0407 (17)	-0.0151 (16)	-0.0143 (14)	-0.0215 (15)
O2	0.063 (2)	0.062 (2)	0.0262 (16)	-0.0189 (16)	-0.0099 (14)	-0.0215 (14)
O3	0.052 (2)	0.057 (2)	0.0469 (18)	-0.0093 (16)	-0.0152 (15)	-0.0159 (16)
O4	0.066 (2)	0.066 (2)	0.0367 (17)	-0.0204 (18)	-0.0114 (16)	-0.0185 (16)
N1	0.0352 (19)	0.040 (2)	0.0284 (18)	-0.0048 (15)	-0.0108 (14)	-0.0173 (15)
N3	0.051 (2)	0.040 (2)	0.0215 (17)	-0.0109 (16)	-0.0091 (15)	-0.0102 (15)
N2	0.039 (2)	0.038 (2)	0.0243 (18)	-0.0015 (16)	-0.0131 (15)	-0.0153 (15)
N4	0.042 (2)	0.036 (2)	0.0307 (17)	-0.0063 (15)	-0.0120 (15)	-0.0178 (16)
C4	0.028 (2)	0.036 (2)	0.0259 (19)	-0.0023 (17)	-0.0081 (16)	-0.0156 (17)
C9	0.032 (2)	0.032 (2)	0.026 (2)	-0.0039 (17)	-0.0074 (16)	-0.0151 (17)
C14	0.031 (2)	0.038 (2)	0.0234 (19)	-0.0035 (17)	-0.0099 (16)	-0.0140 (17)
C15	0.043 (2)	0.043 (3)	0.025 (2)	-0.006 (2)	-0.0110 (18)	-0.0165 (19)
C6	0.041 (2)	0.030 (2)	0.031 (2)	-0.0048 (18)	-0.0075 (18)	-0.0106 (18)
C8	0.028 (2)	0.035 (2)	0.0211 (19)	-0.0038 (17)	-0.0079 (15)	-0.0110 (17)
C1	0.036 (2)	0.044 (3)	0.033 (2)	-0.002 (2)	-0.0114 (19)	-0.019 (2)
C3	0.028 (2)	0.036 (2)	0.028 (2)	-0.0012 (17)	-0.0086 (16)	-0.0179 (18)
C13	0.030 (2)	0.036 (2)	0.027 (2)	-0.0042 (17)	-0.0063 (16)	-0.0151 (18)
C12	0.044 (2)	0.043 (3)	0.024 (2)	-0.012 (2)	-0.0076 (17)	-0.0122 (19)
C7	0.042 (2)	0.034 (2)	0.026 (2)	-0.0012 (19)	-0.0110 (18)	-0.0072 (18)
C10	0.050 (3)	0.040 (3)	0.039 (2)	-0.006 (2)	-0.017 (2)	-0.021 (2)
C5	0.038 (2)	0.038 (2)	0.033 (2)	-0.0047 (18)	-0.0085 (18)	-0.0193 (19)

C2	0.030 (2)	0.046 (3)	0.026 (2)	-0.0032 (18)	-0.0076 (17)	-0.0194 (19)
C11	0.056 (3)	0.038 (2)	0.036 (2)	-0.014 (2)	-0.012 (2)	-0.014 (2)
C16	0.046 (3)	0.057 (3)	0.035 (2)	-0.016 (2)	-0.010 (2)	-0.020 (2)

Geometric parameters (Å, °)

O1—C1	1.244 (4)	C9—C8	1.449 (5)
O2—C1	1.270 (4)	C14—C3	1.398 (5)
O3—C16	1.341 (5)	C14—C13	1.450 (5)
O3—H3	0.8200	C15—C2	1.400 (5)
O4—C16	1.196 (5)	C15—C16	1.510 (5)
N1—C2	1.330 (4)	C6—C5	1.373 (5)
N1—C3	1.346 (4)	C6—C7	1.380 (5)
N3—C15	1.323 (5)	C6—H6	0.9300
N3—C14	1.354 (4)	C1—C2	1.520 (5)
N2—C7	1.319 (5)	C13—C12	1.397 (5)
N2—C8	1.360 (4)	C12—C11	1.365 (5)
N2—H2	0.808 (19)	C12—H12	0.9300
N4—C10	1.316 (5)	C7—H7	0.9300
N4—C9	1.347 (4)	C10—C11	1.402 (5)
C4—C5	1.393 (5)	C10—H10	0.9300
C4—C8	1.392 (5)	C5—H5	0.9300
C4—C3	1.458 (5)	C11—H11	0.9300
C9—C13	1.404 (5)		
C16—O3—H3	109.5	N1—C3—C14	121.7 (3)
C2—N1—C3	116.5 (3)	N1—C3—C4	118.7 (3)
C15—N3—C14	116.4 (3)	C14—C3—C4	119.6 (3)
C7—N2—C8	121.8 (3)	C12—C13—C9	117.8 (3)
C7—N2—H2	120 (3)	C12—C13—C14	123.1 (3)
C8—N2—H2	118 (3)	C9—C13—C14	119.1 (3)
C10—N4—C9	117.1 (3)	C11—C12—C13	119.4 (4)
C5—C4—C8	118.4 (3)	C11—C12—H12	120.3
C5—C4—C3	122.8 (3)	C13—C12—H12	120.3
C8—C4—C3	118.8 (3)	N2—C7—C6	121.3 (4)
N4—C9—C13	123.0 (3)	N2—C7—H7	119.3
N4—C9—C8	117.6 (3)	C6—C7—H7	119.3
C13—C9—C8	119.4 (3)	N4—C10—C11	124.5 (4)
N3—C14—C3	121.2 (3)	N4—C10—H10	117.7
N3—C14—C13	117.5 (3)	C11—C10—H10	117.7
C3—C14—C13	121.3 (3)	C6—C5—C4	120.6 (3)
N3—C15—C2	122.3 (3)	C6—C5—H5	119.7
N3—C15—C16	112.3 (3)	C4—C5—H5	119.7
C2—C15—C16	125.2 (3)	N1—C2—C15	121.7 (3)
C5—C6—C7	118.5 (4)	N1—C2—C1	118.1 (3)
C5—C6—H6	120.8	C15—C2—C1	120.1 (3)
C7—C6—H6	120.8	C12—C11—C10	118.1 (4)
N2—C8—C4	119.4 (3)	C12—C11—H11	121.0

N2—C8—C9	118.8 (3)	C10—C11—H11	121.0
C4—C8—C9	121.8 (3)	O4—C16—O3	120.5 (4)
O1—C1—O2	127.5 (4)	O4—C16—C15	121.8 (4)
O1—C1—C2	119.1 (3)	O3—C16—C15	117.4 (4)
O2—C1—C2	113.4 (4)		
C10—N4—C9—C13	-0.1 (5)	C8—C9—C13—C14	-2.5 (5)
C10—N4—C9—C8	-177.9 (3)	N3—C14—C13—C12	0.1 (5)
C15—N3—C14—C3	-1.6 (5)	C3—C14—C13—C12	-177.7 (3)
C15—N3—C14—C13	-179.5 (3)	N3—C14—C13—C9	179.6 (3)
C14—N3—C15—C2	-1.7 (6)	C3—C14—C13—C9	1.8 (5)
C14—N3—C15—C16	174.5 (3)	C9—C13—C12—C11	1.2 (6)
C7—N2—C8—C4	-0.4 (5)	C14—C13—C12—C11	-179.3 (3)
C7—N2—C8—C9	177.6 (3)	C8—N2—C7—C6	-0.6 (6)
C5—C4—C8—N2	1.0 (5)	C5—C6—C7—N2	0.9 (6)
C3—C4—C8—N2	179.2 (3)	C9—N4—C10—C11	0.5 (6)
C5—C4—C8—C9	-176.9 (3)	C7—C6—C5—C4	-0.2 (6)
C3—C4—C8—C9	1.3 (5)	C8—C4—C5—C6	-0.7 (5)
N4—C9—C8—N2	1.0 (5)	C3—C4—C5—C6	-178.8 (3)
C13—C9—C8—N2	-176.9 (3)	C3—N1—C2—C15	-1.2 (5)
N4—C9—C8—C4	178.8 (3)	C3—N1—C2—C1	179.3 (3)
C13—C9—C8—C4	1.0 (5)	N3—C15—C2—N1	3.3 (6)
C2—N1—C3—C14	-2.1 (5)	C16—C15—C2—N1	-172.4 (4)
C2—N1—C3—C4	178.9 (3)	N3—C15—C2—C1	-177.2 (3)
N3—C14—C3—N1	3.7 (6)	C16—C15—C2—C1	7.1 (6)
C13—C14—C3—N1	-178.5 (3)	O1—C1—C2—N1	16.9 (5)
N3—C14—C3—C4	-177.3 (3)	O2—C1—C2—N1	-163.8 (3)
C13—C14—C3—C4	0.5 (5)	O1—C1—C2—C15	-162.6 (4)
C5—C4—C3—N1	-4.9 (5)	O2—C1—C2—C15	16.8 (5)
C8—C4—C3—N1	177.0 (3)	C13—C12—C11—C10	-0.9 (6)
C5—C4—C3—C14	176.1 (3)	N4—C10—C11—C12	0.0 (6)
C8—C4—C3—C14	-2.0 (5)	N3—C15—C16—O4	73.0 (5)
N4—C9—C13—C12	-0.7 (6)	C2—C15—C16—O4	-110.9 (5)
C8—C9—C13—C12	177.0 (3)	N3—C15—C16—O3	-100.7 (4)
N4—C9—C13—C14	179.8 (3)	C2—C15—C16—O3	75.3 (5)

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3 \cdots O1 ⁱ	0.82	1.82	2.638 (4)	171
N2—H2 \cdots O2 ⁱⁱ	0.81 (2)	1.87 (3)	2.638 (4)	159 (5)

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