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1,10-Phenanthroline-1-ium 2-carboxy-4,5-dichlorobenzoate

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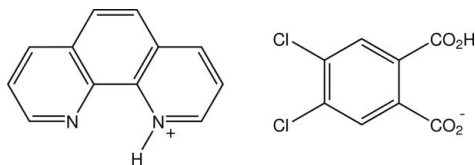
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 Key indicators: single-crystal X-ray study; $T = 130$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.032; wR factor = 0.085; data-to-parameter ratio = 14.3.

In the structure of the 1:1 proton-transfer compound of 1,10-phenanthroline with 4,5-dichlorophthalic acid, $\text{C}_{12}\text{H}_9\text{N}_2^+ \cdot \text{C}_8\text{H}_3\text{Cl}_2\text{O}_4^-$, determined at 130 K, the 1,10-phenanthrolium cation and the hydrogen 4,5-dichlorophthalate anion associate through a single $\text{N}-\text{H} \cdots \text{O}_{\text{carboxyl}}$ hydrogen bond giving discrete units which have no extension except through a number of weak cation $\text{C}-\text{H} \cdots \text{O}_{\text{anion}}$ associations and weak cation-anion aromatic ring $\pi-\pi$ interactions [minimum centroid-centroid separation = 3.6815 (12) Å]. The anions are essentially planar [maximum deviation 0.214 (1) Å (a carboxyl O)] with the *syn*-related H atom of the carboxyl group, forming a short intramolecular $\text{O}-\text{H} \cdots \text{O}_{\text{carboxyl}}$ hydrogen bond.

Related literature

For the structures of other hydrogen 4,5-dichlorophthalate salts, see: Mallinson *et al.* (2003); Bozkurt *et al.* (2006); Smith *et al.* (2007, 2008*a,b*, 2009*a,b*). For hydrogen-bond motifs, see: Etter *et al.* (1990).



Experimental

Crystal data

 $\text{C}_{12}\text{H}_9\text{N}_2^+ \cdot \text{C}_8\text{H}_3\text{Cl}_2\text{O}_4^-$
 $M_r = 415.22$
 Monoclinic, $P2_1$
 $a = 6.4598$ (11) Å
 $b = 7.3696$ (12) Å

 $c = 18.302$ (3) Å
 $\beta = 94.978$ (3)°
 $V = 868.0$ (2) Å³
 $Z = 2$
 Mo $K\alpha$ radiation

 $\mu = 0.41$ mm⁻¹
 $T = 130$ K

 $0.55 \times 0.45 \times 0.05$ mm

Data collection

 Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\text{min}} = 0.81$, $T_{\text{max}} = 0.98$

 5464 measured reflections
 3734 independent reflections
 3629 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.085$
 $S = 1.04$
 3734 reflections
 261 parameters
 1 restraint

 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.30$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³
 Absolute structure: Flack (1983), 1564 Friedel pairs
 Flack parameter: 0.00 (4)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{N1A}-\text{H1A} \cdots \text{O22}$	0.90 (2)	1.83 (2)	2.6926 (19)	158 (2)
$\text{N1A}-\text{H1A} \cdots \text{N10A}$	0.90 (2)	2.38 (2)	2.749 (2)	104.3 (15)
$\text{O12}-\text{H12} \cdots \text{O21}$	0.98 (3)	1.43 (3)	2.4054 (19)	179 (4)
$\text{C2A}-\text{H2A} \cdots \text{O21}$	0.93	2.52	3.279 (2)	140
$\text{C3}-\text{H3} \cdots \text{O22}$	0.93	2.26	2.647 (2)	104
$\text{C3A}-\text{H3A} \cdots \text{O11}^{\text{i}}$	0.93	2.44	3.355 (2)	168
$\text{C4A}-\text{H4A} \cdots \text{O21}^{\text{ii}}$	0.93	2.49	3.252 (2)	139
$\text{C6}-\text{H6} \cdots \text{O11}$	0.93	2.29	2.668 (2)	103
$\text{C6A}-\text{H6A} \cdots \text{O11}^{\text{iii}}$	0.93	2.59	3.270 (2)	130

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

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 1999); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FL2261).

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supporting information

Acta Cryst. (2009). E65, o2333 [doi:10.1107/S1600536809034448]

1,10-Phenanthroline-1-ium 2-carboxy-4,5-dichlorobenzoate

Graham Smith, Urs D. Wermuth and Jonathan M. White

S1. Comment

The 1:1 proton-transfer compounds of 4,5-dichlorophthalic acid (DCPA) with the aromatic nitrogen Lewis bases commonly have low-dimensional hydrogen-bonded structures (Smith *et al.*, 2007, 2008a, 2008b, 2009a, 2009b; Bozkurt *et al.*, 2006; Mallinson *et al.*, 2003). In the two-dimensional examples the DCPA anions assume non-planar conformations and form into sheet substructures which in the case of the compounds with the *meta*- and *para*-amino-benzoic acids (Smith *et al.*, 2008b) are extended into three-dimensional frameworks through peripheral cyclic head-to-head carboxylic acid hydrogen-bonding associations. However, with the majority of the structures, e.g. the brucinium salt (Smith *et al.*, 2007), the DCPA anions are essentially planar with short intramolecular carboxylic acid O–H \cdots O_{carboxyl} hydrogen bonds. These features were therefore expected and found in the 1:1 proton-transfer compound of DCPA with 1,10-phenanthroline, (I), reported here.

In (I), a single N⁺–H \cdots O_{carboxyl} hydrogen bond links the phenanthroline cation and the DCPA anion (Fig. 1). A weak aromatic ring C–H \cdots O_{carboxyl} interaction (Table 1) completes an asymmetric R²₂(7) cyclic association (Etter *et al.*, 1990). Three additional anion C–H \cdots O interactions represent the only structure extensions present. Some overlap is present between the anion aromatic ring (C1–C6) and one six-membered ring of the cation (N10A, C9A, C8A, C7A, C6A, C14A) [minimum ring centroid separation, 3.6815 (12) Å] (Fig. 2), giving weak π – π stacking interactions (Fig. 3). The DCPA anion is essentially planar [torsion angles C2–C1–C11–O11, -168.30 (16)°; C1–C2–C21–O22, -179.53 (16)°], and exhibits a short intramolecular O–H \cdots O_{carboxyl} hydrogen bond [2.4054 (19) Å]. Associated with this bond is a significant distortion of the *exo*-C1 and C2 bond angles [C1–C2–C21, 128.55 (15) ° and C2–C1–C11, 129.18 (16) °]. This and a lengthening of the C1–C11 and C2–C21 bonds [1.538 (3) and 1.536 (3) Å] is common to the planar DCPA anions in the series of 1:1 proton-transfer compounds [angle range: 127.88 (16)° in the nicotinamide salt (Smith *et al.*, 2009a) to 129.27 (14)° in the 8-aminoquinoline salt (Smith *et al.*, 2008a); bond length range: 1.523 (3)–1.535 (3) Å, both in the brucinium salt (Smith *et al.*, 2007).

S2. Experimental

The title compound (I) was synthesized by heating 1 mmol quantities of 1,10-phenanthroline and 4,5-dichlorophthalic acid in 50 ml of 95% ethanol for 10 min under reflux. After concentration to *ca.* 30 ml, partial room-temperature evaporation of the hot-filtered solution gave colourless plates (m.p. 464–465 K) suitable for data collection.

S3. Refinement

Hydrogen atoms potentially involved in hydrogen-bonding interactions were located by difference methods and their positional and isotropic displacement parameters were refined. Other H atoms were included in the refinement at calculated positions [C–H, 0.93 Å] and treated as riding models with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

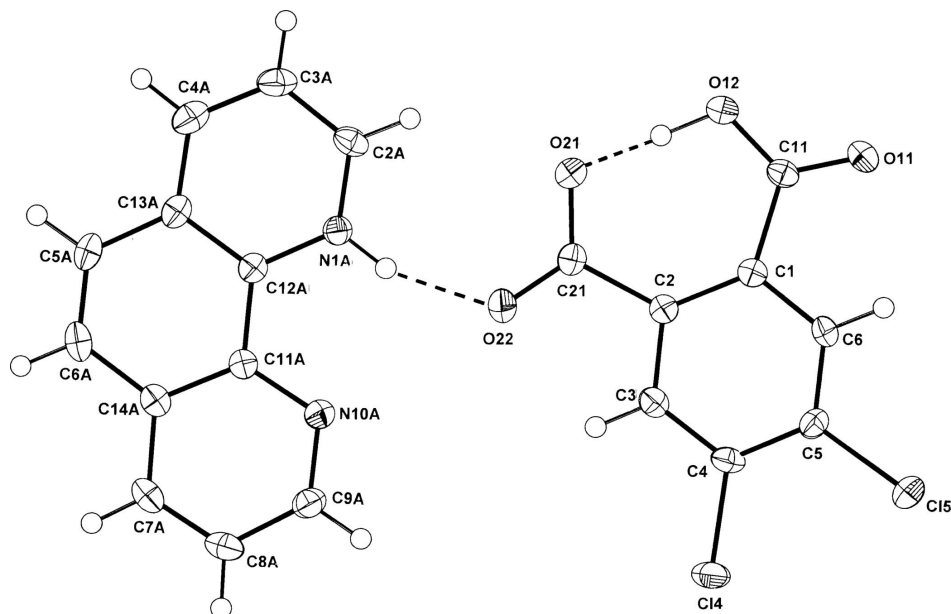


Figure 1

Molecular configuration and atom numbering scheme for the 1,10-phenanthroline-1-ium cation and the hydrogen 4,5-dichlorophthalate anion in (I). Non-H atoms are shown as 50% probability displacement ellipsoids. The inter-species hydrogen bond is shown as a dashed line.

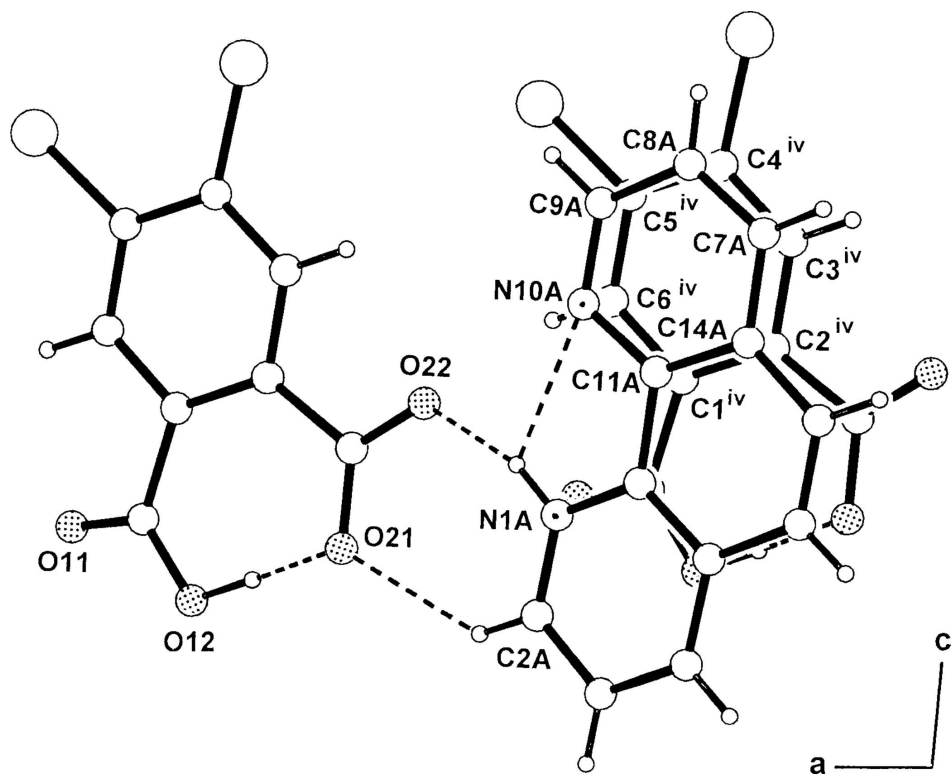


Figure 2

Cation-anion aromatic ring overlap in (I) viewed down the approximate *b* direction in the unit cell. For symmetry code (iv): $x + 1, y - 1, z$.

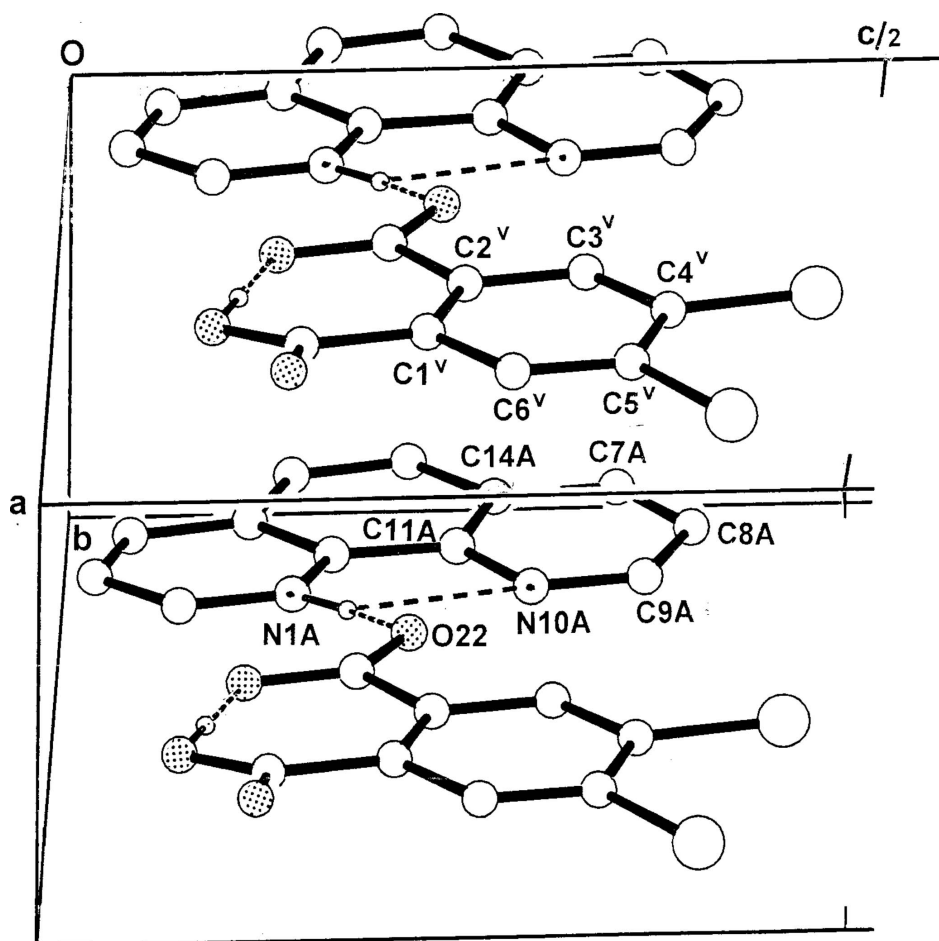


Figure 3

Aromatic ring π - π interactions in a perspective view of part of the unit cell. Non-interactive H atoms are omitted. For symmetry code (v): $x - 1, y, z$.

1,10-Phenanthroline-1-ium 2-carboxy-4,5-dichlorobenzoate

Crystal data

$C_{12}H_9N_2^+ \cdot C_8H_3Cl_2O_4^-$

$M_r = 415.22$

Monoclinic, $P2_1$

Hall symbol: $P\ 2y_b$

$a = 6.4598$ (11) Å

$b = 7.3696$ (12) Å

$c = 18.302$ (3) Å

$\beta = 94.978$ (3)°

$V = 868.0$ (2) Å³

$Z = 2$

$F(000) = 424$

$D_x = 1.589$ Mg m⁻³

Melting point = 464–465 K

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

Cell parameters from 3630 reflections

$\theta = 2.2$ – 27.5 °

$\mu = 0.41$ mm⁻¹

$T = 130$ K

Plate, colourless

$0.55 \times 0.45 \times 0.05$ mm

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.81$, $T_{\max} = 0.98$

5464 measured reflections
 3734 independent reflections
 3629 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

$\theta_{\text{max}} = 27.6^\circ$, $\theta_{\text{min}} = 1.1^\circ$
 $h = -5 \rightarrow 8$
 $k = -9 \rightarrow 9$
 $l = -23 \rightarrow 21$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.085$
 $S = 1.04$
 3734 reflections
 261 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
 $w = 1/[\sigma^2(F_o^2) + (0.053P)^2 + 0.0522P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.30 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.19 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 1564 Friedel
 pairs
 Absolute structure parameter: 0.00 (4)

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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	$U_{\text{iso}}^*/U_{\text{eq}}$
N1A	0.4667 (2)	0.7304 (2)	0.14863 (8)	0.0192 (4)
N10A	0.4177 (2)	0.7715 (2)	0.29542 (8)	0.0212 (4)
C2A	0.5046 (3)	0.7149 (3)	0.07881 (10)	0.0232 (5)
C3A	0.3645 (3)	0.7828 (3)	0.02322 (10)	0.0277 (5)
C4A	0.1845 (3)	0.8641 (3)	0.04213 (10)	0.0268 (5)
C5A	-0.0459 (3)	0.9562 (3)	0.13825 (11)	0.0233 (5)
C6A	-0.0804 (3)	0.9634 (2)	0.20961 (11)	0.0246 (5)
C7A	0.0442 (3)	0.9113 (2)	0.34093 (10)	0.0259 (6)
C8A	0.2013 (3)	0.8540 (3)	0.39058 (10)	0.0286 (5)
C9A	0.3857 (3)	0.7854 (3)	0.36523 (10)	0.0253 (5)
C11A	0.2617 (3)	0.8274 (2)	0.24641 (9)	0.0184 (4)
C12A	0.2916 (3)	0.8115 (2)	0.16952 (10)	0.0178 (4)
C13A	0.1421 (3)	0.8783 (2)	0.11565 (10)	0.0207 (4)
C14A	0.0717 (3)	0.9002 (2)	0.26564 (10)	0.0207 (5)
Cl4	1.15820 (7)	0.37080 (8)	0.46649 (2)	0.0332 (1)
Cl5	1.59276 (7)	0.22144 (8)	0.42219 (2)	0.0316 (1)
O11	1.5052 (2)	0.18688 (18)	0.14604 (7)	0.0251 (4)
O12	1.2404 (2)	0.34157 (19)	0.09355 (7)	0.0265 (4)
O21	0.9277 (2)	0.4836 (2)	0.12771 (7)	0.0271 (4)
O22	0.76248 (19)	0.53666 (18)	0.22655 (7)	0.0254 (4)

C1	1.2820 (3)	0.3211 (2)	0.22748 (9)	0.0179 (5)
C2	1.0928 (3)	0.3971 (2)	0.24707 (9)	0.0178 (4)
C3	1.0603 (3)	0.4092 (2)	0.32117 (10)	0.0206 (5)
C4	1.2096 (3)	0.3536 (3)	0.37559 (9)	0.0215 (5)
C5	1.3983 (3)	0.2862 (2)	0.35636 (10)	0.0207 (5)
C6	1.4314 (3)	0.2693 (2)	0.28333 (10)	0.0196 (4)
C11	1.3502 (3)	0.2788 (2)	0.15083 (9)	0.0195 (5)
C21	0.9127 (3)	0.4764 (2)	0.19673 (9)	0.0193 (4)
H1A	0.556 (3)	0.683 (3)	0.1842 (12)	0.021 (5)*
H2A	0.62580	0.65840	0.06680	0.0280*
H3A	0.39190	0.77330	-0.02570	0.0330*
H4A	0.09010	0.91030	0.00560	0.0320*
H5A	-0.14490	1.00210	0.10310	0.0280*
H6A	-0.20540	1.01040	0.22290	0.0300*
H7A	-0.07860	0.95670	0.35670	0.0310*
H8A	0.18650	0.86030	0.44060	0.0340*
H9A	0.49080	0.74780	0.39980	0.0300*
H3	0.93520	0.45580	0.33450	0.0250*
H6	1.55700	0.22200	0.27080	0.0230*
H12	1.113 (5)	0.400 (4)	0.1072 (17)	0.035 (9)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1A	0.0167 (6)	0.0212 (7)	0.0195 (7)	-0.0015 (6)	0.0013 (5)	0.0001 (6)
N10A	0.0217 (7)	0.0224 (7)	0.0193 (7)	0.0012 (6)	0.0005 (6)	-0.0004 (6)
C2A	0.0222 (8)	0.0261 (8)	0.0222 (8)	-0.0047 (8)	0.0078 (7)	-0.0028 (7)
C3A	0.0292 (10)	0.0360 (10)	0.0182 (8)	-0.0077 (8)	0.0036 (7)	0.0005 (7)
C4A	0.0268 (9)	0.0298 (9)	0.0230 (8)	-0.0069 (8)	-0.0029 (7)	0.0060 (8)
C5A	0.0176 (8)	0.0211 (8)	0.0302 (9)	-0.0004 (7)	-0.0040 (7)	0.0041 (7)
C6A	0.0164 (8)	0.0202 (9)	0.0375 (10)	0.0014 (7)	0.0033 (7)	-0.0017 (8)
C7A	0.0242 (9)	0.0253 (10)	0.0295 (10)	0.0011 (8)	0.0104 (7)	-0.0027 (7)
C8A	0.0358 (10)	0.0308 (10)	0.0203 (8)	0.0001 (9)	0.0096 (7)	-0.0023 (8)
C9A	0.0295 (9)	0.0263 (9)	0.0197 (8)	0.0025 (8)	-0.0007 (7)	0.0020 (7)
C11A	0.0182 (8)	0.0164 (7)	0.0207 (8)	-0.0009 (6)	0.0026 (6)	0.0009 (6)
C12A	0.0163 (7)	0.0166 (7)	0.0204 (8)	-0.0040 (6)	0.0005 (6)	0.0001 (6)
C13A	0.0186 (7)	0.0188 (7)	0.0241 (8)	-0.0039 (7)	-0.0020 (6)	0.0027 (7)
C14A	0.0195 (8)	0.0181 (8)	0.0248 (8)	-0.0031 (7)	0.0039 (6)	-0.0021 (7)
C14	0.0291 (2)	0.0541 (3)	0.0169 (2)	0.0014 (2)	0.0045 (2)	-0.0034 (2)
C15	0.0251 (2)	0.0463 (3)	0.0221 (2)	0.0067 (2)	-0.0046 (2)	-0.0001 (2)
O11	0.0227 (6)	0.0297 (7)	0.0236 (6)	0.0044 (6)	0.0058 (5)	-0.0029 (5)
O12	0.0260 (6)	0.0354 (7)	0.0187 (6)	0.0056 (6)	0.0052 (5)	0.0018 (5)
O21	0.0264 (7)	0.0343 (7)	0.0204 (6)	0.0068 (6)	0.0009 (5)	0.0030 (5)
O22	0.0200 (6)	0.0284 (7)	0.0277 (7)	0.0044 (5)	0.0023 (5)	0.0012 (5)
C1	0.0189 (8)	0.0157 (8)	0.0192 (8)	-0.0027 (6)	0.0029 (6)	0.0006 (6)
C2	0.0188 (8)	0.0149 (7)	0.0197 (8)	-0.0012 (6)	0.0024 (6)	0.0007 (6)
C3	0.0189 (8)	0.0215 (9)	0.0215 (8)	-0.0012 (7)	0.0032 (7)	-0.0034 (6)
C4	0.0229 (8)	0.0263 (9)	0.0161 (7)	-0.0035 (7)	0.0057 (6)	-0.0019 (7)

C5	0.0188 (8)	0.0231 (8)	0.0196 (8)	-0.0005 (7)	-0.0019 (6)	0.0019 (7)
C6	0.0155 (7)	0.0207 (8)	0.0229 (8)	0.0010 (6)	0.0040 (6)	-0.0014 (7)
C11	0.0202 (8)	0.0197 (8)	0.0192 (8)	-0.0041 (7)	0.0055 (6)	-0.0012 (6)
C21	0.0168 (7)	0.0173 (8)	0.0233 (8)	-0.0023 (6)	-0.0005 (6)	-0.0004 (6)

Geometric parameters (Å, °)

C14—C4	1.7291 (17)	C11A—C12A	1.442 (2)
C15—C5	1.7305 (19)	C11A—C14A	1.412 (3)
O11—C11	1.218 (2)	C12A—C13A	1.408 (3)
O12—C11	1.299 (2)	C2A—H2A	0.9300
O21—C21	1.276 (2)	C3A—H3A	0.9300
O22—C21	1.236 (2)	C4A—H4A	0.9300
O12—H12	0.98 (3)	C5A—H5A	0.9300
N1A—C12A	1.363 (2)	C6A—H6A	0.9300
N1A—C2A	1.327 (2)	C7A—H7A	0.9300
N10A—C11A	1.354 (2)	C8A—H8A	0.9300
N10A—C9A	1.316 (2)	C9A—H9A	0.9300
N1A—H1A	0.90 (2)	C1—C2	1.419 (3)
C2A—C3A	1.395 (3)	C1—C11	1.538 (2)
C3A—C4A	1.379 (3)	C1—C6	1.397 (3)
C4A—C13A	1.400 (3)	C2—C21	1.536 (3)
C5A—C6A	1.345 (3)	C2—C3	1.393 (2)
C5A—C13A	1.436 (3)	C3—C4	1.387 (3)
C6A—C14A	1.435 (3)	C4—C5	1.390 (3)
C7A—C14A	1.407 (3)	C5—C6	1.377 (3)
C7A—C8A	1.369 (3)	C3—H3	0.9300
C8A—C9A	1.409 (3)	C6—H6	0.9300
C11—O12—H12	111 (2)	C14A—C6A—H6A	119.00
C2A—N1A—C12A	122.34 (16)	C5A—C6A—H6A	119.00
C9A—N10A—C11A	116.68 (15)	C8A—C7A—H7A	121.00
C12A—N1A—H1A	117.5 (13)	C14A—C7A—H7A	121.00
C2A—N1A—H1A	120.1 (13)	C9A—C8A—H8A	120.00
N1A—C2A—C3A	120.63 (18)	C7A—C8A—H8A	120.00
C2A—C3A—C4A	118.74 (17)	C8A—C9A—H9A	118.00
C3A—C4A—C13A	120.89 (17)	N10A—C9A—H9A	118.00
C6A—C5A—C13A	120.73 (18)	C2—C1—C11	129.18 (16)
C5A—C6A—C14A	121.35 (18)	C2—C1—C6	118.60 (15)
C8A—C7A—C14A	118.87 (17)	C6—C1—C11	112.19 (16)
C7A—C8A—C9A	119.43 (17)	C1—C2—C21	128.55 (15)
N10A—C9A—C8A	123.79 (17)	C1—C2—C3	118.51 (16)
N10A—C11A—C14A	124.31 (15)	C3—C2—C21	112.92 (16)
C12A—C11A—C14A	117.85 (16)	C2—C3—C4	121.74 (17)
N10A—C11A—C12A	117.83 (16)	C14—C4—C5	121.05 (14)
C11A—C12A—C13A	120.89 (17)	C3—C4—C5	119.63 (16)
N1A—C12A—C11A	119.61 (16)	C14—C4—C3	119.32 (15)
N1A—C12A—C13A	119.50 (16)	C4—C5—C6	119.44 (17)

C5A—C13A—C12A	118.94 (17)	C15—C5—C4	121.47 (14)
C4A—C13A—C5A	123.19 (17)	C15—C5—C6	119.09 (15)
C4A—C13A—C12A	117.87 (17)	C1—C6—C5	122.00 (17)
C6A—C14A—C7A	122.97 (17)	O11—C11—C1	118.72 (15)
C7A—C14A—C11A	116.91 (16)	O12—C11—C1	118.97 (15)
C6A—C14A—C11A	120.09 (16)	O11—C11—O12	122.32 (16)
C3A—C2A—H2A	120.00	O22—C21—C2	117.04 (15)
N1A—C2A—H2A	120.00	O21—C21—O22	123.49 (17)
C2A—C3A—H3A	121.00	O21—C21—C2	119.41 (16)
C4A—C3A—H3A	121.00	C2—C3—H3	119.00
C3A—C4A—H4A	120.00	C4—C3—H3	119.00
C13A—C4A—H4A	120.00	C1—C6—H6	119.00
C6A—C5A—H5A	120.00	C5—C6—H6	119.00
C13A—C5A—H5A	120.00		
C12A—N1A—C2A—C3A	0.2 (3)	N1A—C12A—C13A—C5A	177.27 (16)
C2A—N1A—C12A—C11A	-178.71 (17)	C11A—C12A—C13A—C4A	177.79 (16)
C2A—N1A—C12A—C13A	1.4 (3)	C11A—C12A—C13A—C5A	-2.6 (2)
C11A—N10A—C9A—C8A	0.0 (3)	C6—C1—C2—C3	-2.9 (2)
C9A—N10A—C11A—C12A	-179.22 (16)	C6—C1—C2—C21	175.21 (15)
C9A—N10A—C11A—C14A	0.8 (2)	C11—C1—C2—C3	174.90 (15)
N1A—C2A—C3A—C4A	-0.9 (3)	C11—C1—C2—C21	-7.0 (3)
C2A—C3A—C4A—C13A	-0.1 (3)	C2—C1—C6—C5	1.5 (2)
C3A—C4A—C13A—C5A	-177.89 (19)	C11—C1—C6—C5	-176.68 (14)
C3A—C4A—C13A—C12A	1.7 (3)	C2—C1—C11—O11	-168.30 (16)
C13A—C5A—C6A—C14A	2.4 (3)	C2—C1—C11—O12	11.4 (3)
C6A—C5A—C13A—C4A	178.76 (18)	C6—C1—C11—O11	9.6 (2)
C6A—C5A—C13A—C12A	-0.8 (3)	C6—C1—C11—O12	-170.67 (15)
C5A—C6A—C14A—C7A	177.74 (17)	C1—C2—C3—C4	1.7 (2)
C5A—C6A—C14A—C11A	-0.5 (2)	C21—C2—C3—C4	-176.64 (16)
C14A—C7A—C8A—C9A	-0.1 (3)	C1—C2—C21—O21	-2.2 (3)
C8A—C7A—C14A—C6A	-177.47 (17)	C1—C2—C21—O22	-179.53 (16)
C8A—C7A—C14A—C11A	0.8 (2)	C3—C2—C21—O21	175.96 (15)
C7A—C8A—C9A—N10A	-0.3 (3)	C3—C2—C21—O22	-1.4 (2)
N10A—C11A—C12A—N1A	4.5 (2)	C2—C3—C4—C14	-179.50 (14)
N10A—C11A—C12A—C13A	-175.62 (15)	C2—C3—C4—C5	0.9 (3)
C14A—C11A—C12A—N1A	-175.47 (15)	C14—C4—C5—C15	-1.6 (2)
C14A—C11A—C12A—C13A	4.4 (2)	C14—C4—C5—C6	178.04 (14)
N10A—C11A—C14A—C6A	177.13 (15)	C3—C4—C5—C15	178.02 (14)
N10A—C11A—C14A—C7A	-1.2 (2)	C3—C4—C5—C6	-2.4 (3)
C12A—C11A—C14A—C6A	-2.9 (2)	C15—C5—C6—C1	-179.20 (12)
C12A—C11A—C14A—C7A	178.83 (14)	C4—C5—C6—C1	1.2 (2)
N1A—C12A—C13A—C4A	-2.4 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1A—H1A...O22	0.90 (2)	1.83 (2)	2.6926 (19)	158 (2)

N1A—H1A...N10A	0.90 (2)	2.38 (2)	2.749 (2)	104.3 (15)
O12—H12...O21	0.98 (3)	1.43 (3)	2.4054 (19)	179 (4)
C2A—H2A...O21	0.93	2.52	3.279 (2)	140
C3—H3...O22	0.93	2.26	2.647 (2)	104
C3A—H3A...O11 ⁱ	0.93	2.44	3.355 (2)	168
C4A—H4A...O21 ⁱⁱ	0.93	2.49	3.252 (2)	139
C6—H6...O11	0.93	2.29	2.668 (2)	103
C6A—H6A...O11 ⁱⁱⁱ	0.93	2.59	3.270 (2)	130

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