

## 2,2'-(Piperazine-1,4-diyl)diethanaminium dibenzoate

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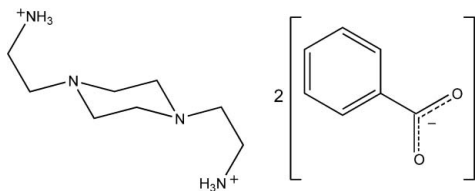
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Key indicators: single-crystal X-ray study;  $T = 180$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.040;  $wR$  factor = 0.110; data-to-parameter ratio = 17.9.

The asymmetric unit of the title salt  $\text{C}_8\text{H}_{22}\text{N}_4^{2+} \cdot 2\text{C}_7\text{H}_5\text{O}_2^-$ , comprises two independent pairs of half a 2,2'-(piperazine-1,4-diyl)diethanaminium dication plus a benzoate anion. The dications are symmetrical and lie across crystallographic centres of inversion. The crystal structure was refined as a two-component pseudo-merohedral twin using the twin law 001 010 100 [the domain fractions are 0.8645 (8) and 0.1355 (8)]. The anions and cations are linked by  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds and weak  $\text{N}-\text{H} \cdots \text{O}$  intermolecular interactions to form infinite two-dimensional networks parallel to [101]. The conformation adopted by the cation in the crystal structure is very similar to that adopted by the same cation in the structures of the 2-hydroxybenzoate [Cukrowski *et al.* (2012). *Acta Cryst.* E68, o2387], the nitrate and the tetrahydrogen pentaborate salts.

### Related literature

For the structures of the 2-hydroxybenzoate, the nitrate and the tetrahydrogen pentaborate salts of the 1,4-di(2-ammonioethyl)piperazine cation, see: Cukrowski *et al.* (2012); Junk & Smith (2005); Jiang *et al.* (2009), respectively.



### Experimental

#### Crystal data

$\text{C}_8\text{H}_{22}\text{N}_4^{2+} \cdot 2\text{C}_7\text{H}_5\text{O}_2^-$   
 $M_r = 416.52$   
 Monoclinic,  $P2_1/n$   
 $a = 19.5300$  (4) Å  
 $b = 6.6694$  (2) Å

$c = 19.6178$  (4) Å  
 $\beta = 115.989$  (1)°  
 $V = 2296.89$  (10) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation

$\mu = 0.08$  mm<sup>-1</sup>  
 $T = 180$  K

0.28 × 0.23 × 0.12 mm

#### Data collection

Nonius KappaCCD diffractometer  
 Absorption correction: multi-scan  
 (SORTAV; Blessing, 1995)  
 $T_{\min} = 0.910$ ,  $T_{\max} = 0.991$

20137 measured reflections  
 5194 independent reflections  
 3970 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.031$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$   
 $wR(F^2) = 0.110$   
 $S = 1.02$   
 5194 reflections  
 290 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\text{max}} = 0.16$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.18$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{N1}-\text{H1A} \cdots \text{O15}^i$	0.945 (18)	1.833 (19)	2.7739 (17)	173.0 (15)
$\text{N1}-\text{H1B} \cdots \text{O15}$	0.902 (16)	1.895 (17)	2.7836 (15)	167.8 (14)
$\text{N1}-\text{H1B} \cdots \text{O14}$	0.902 (16)	2.632 (16)	3.2580 (16)	127.2 (13)
$\text{N1}-\text{H1C} \cdots \text{O14}^{ii}$	0.925 (18)	1.887 (18)	2.7660 (16)	157.9 (14)
$\text{N1}'-\text{H1}'\text{A} \cdots \text{O14}'$	0.882 (19)	1.857 (19)	2.7355 (17)	174.2 (15)
$\text{N1}'-\text{H1}'\text{B} \cdots \text{O14}^{iii}$	0.897 (17)	1.908 (17)	2.7836 (16)	164.8 (15)
$\text{N1}'-\text{H1}'\text{B} \cdots \text{O15}^{iii}$	0.897 (17)	2.533 (16)	3.1858 (16)	130.1 (13)
$\text{N1}'-\text{H1}'\text{C} \cdots \text{O15}^{iv}$	0.916 (18)	1.934 (18)	2.7585 (16)	148.8 (14)

Symmetry codes: (i)  $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (ii)  $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$ ; (iii)  $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iv)  $x, y + 1, z$ .

Data collection: COLLECT (Nonius, 1998); cell refinement: SCALEPACK (Otwinowski & Minor, 1997); data reduction: DENZO (Otwinowski & Minor, 1997), SCALEPACK and SORTAV (Blessing, 1995); program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997), POV-RAY (Cason, 2004) and Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: JJ2134).

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## supplementary materials

*Acta Cryst.* (2012). E68, o2389 [doi:10.1107/S1600536812030115]

**2,2'-(Piperazine-1,4-diyl)diethanaminium dibenzoate**

Ignacy Cukrowski, Adedapo S. Adeyinka and David C. Liles

**Comment**

The title compound [ $C_8H_{22}N_4^{2+} 2(C_7H_5O_2^-)$ ] (**1**) was obtained as an unintended product during an attempt to prepare a benzoate salt of a singly protonated *N,N'*-di(2-aminoethyl)-2-aminoethane-1-ammonium ion ( $C_6H_{19}N_4^+ C_7H_5O_2^-$ ). This occurred because the starting material, instead of being pure *N,N'*-di(2-aminoethyl)-ethane-1,2-diamine, ( $C_6H_{18}N_4$ ), was a mixture of that compound and 1,4-di(2-aminoethyl)piperazine ( $C_8H_{22}N_4$ ) see Cukrowski, *et al.* (2012).

The asymmetric unit of the title compound,  $C_8H_{22}N_4^{2+}, 2(C_7H_5O_2^-)$ , **1**, is a salt with two independent pairs of half a  $C_8H_{22}N_4^{2+}$  cation plus a  $C_7H_5O_2^-$  anion. The  $C_8H_{22}N_4^{2+}$  cations are symmetrical and lie across crystallographic centres of inversion (Fig. 1). The crystal structure was refined as a two-component pseudo-merohedral twin using the twin law 0 0 1 0 - 1 0 1 0 0. The fractional contribution refined to 0.1355 (8).

All three H atoms of each ammonium group in the cations of **1** are hydrogen bonded to the O atoms of the carboxylate groups of the anions. For each ammonium group, one H atom forms a bifurcated hydrogen bond to both of the O atoms of the carboxylate group of one anion, whereas the other two H atoms each form single hydrogen bonds to one O atom of the carboxylate group of each of two additional anions (Fig. 2). Thus both the O atoms of each carboxylate group are each acceptors for two hydrogen bonds. N—H $\cdots$ O hydrogen bonds and weak N—H $\cdots$ O intermolecular interactions link the cations and anions to form a two-dimensional network with layers parallel to the [101] plane (Fig. 2). Each of the two independent cation-anion pairs form the content of alternate network layers.

The conformation adopted by the  $C_8H_{22}N_4^{2+}$  cation in the crystal structure of **1** is very similar to the conformations adopted by the same cation in the crystal structures of the 2-hydroxybenzoate (Cukrowski, *et al.*, 2012), the nitrate (Junk & Smith, 2005) and the tetrahydrogenpentaborate (Jiang, *et al.*, 2009) salts despite the differences in the size and shape of the anions in the various structures.

**Experimental**

2 ml of a 3.32 M aqueous solution of what was claimed by the supplier (QinHuangDao JinLei Chemical Co.Ltd) to be *N,N'*-di(2-aminoethyl)-ethane-1,2-diamine, but which turned out to be a mixture of that compound ( $C_6H_{18}N_4$ , 6.64 mmol) and 1,4-di(2-aminoethyl)piperazine ( $C_8H_{20}N_4$ , 5.57(1-*n*) mmol) was added to 0.78 g of benzoic acid (6.96 mmol), resulting in a clear colourless solution. 0.2 ml of ethanol was added to the solution and the mixture was heated for 3 h at 70 °C. The solution was cooled to room temperature and then left covered for six days and then allowed to slowly evaporate by covering the container with perforated aluminium foil. Yellow crystals were obtained after four days of slow evaporation.

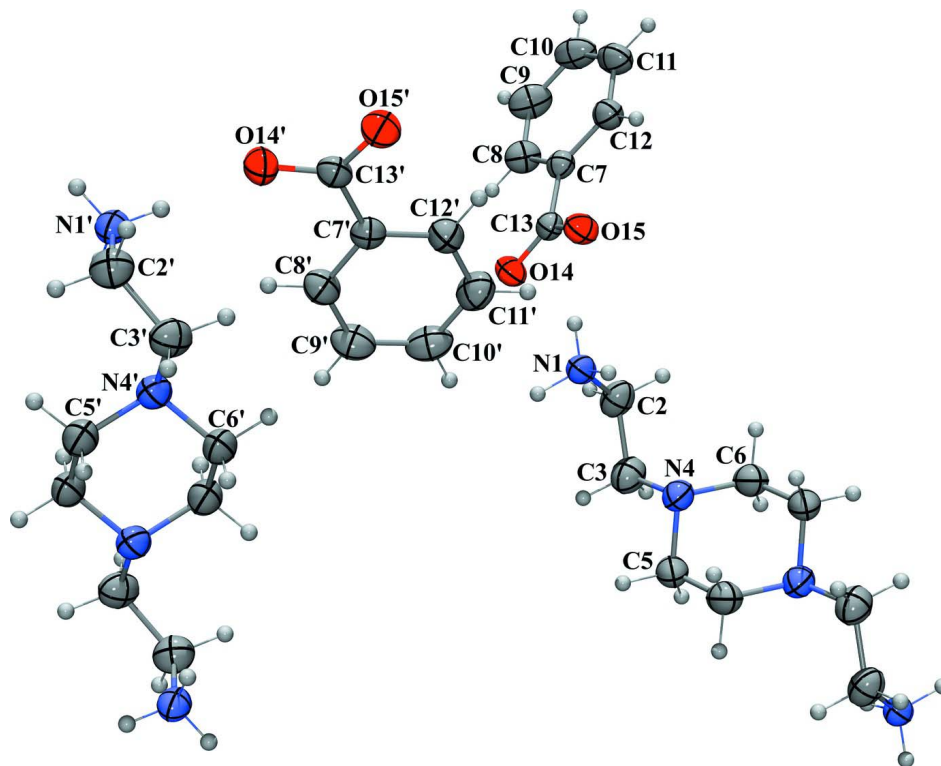
**Refinement**

The crystal structure was refined as a two component pseudo-merohedral twin using the twin law 0 0 1 0 - 1 0 1 0 0. The fractional contribution refined to 0.1355 (8). H1A, H1B and H1C were located by a difference map and their coordinates

were refined. All of the remaining H atoms were placed in their calculated positions and then refined using the riding model with Atom—H lengths of 0.95 Å, (CH) or 0.99 Å (CH<sub>2</sub>). Isotropic displacement parameters for all hydrogen atoms were set to 1.20 times  $U_{eq}$  of the parent atom.

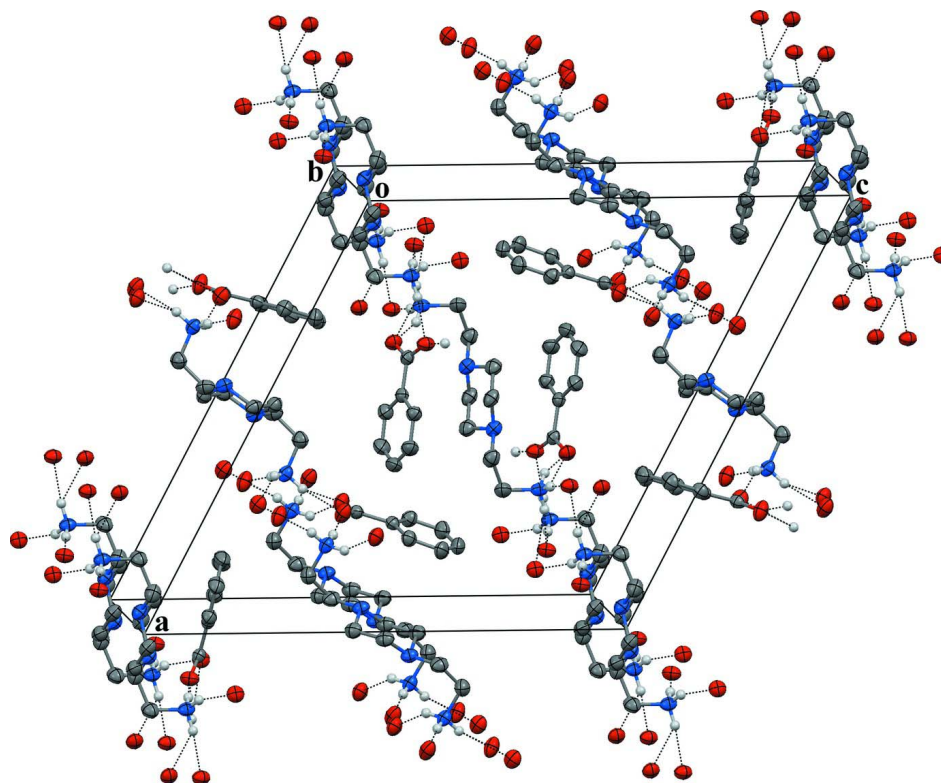
### Computing details

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997), *SCALEPACK* and *SORTAV* (Blessing, 1995); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997), *POV-RAY* (Cason, 2004) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008) and *PLATON* (Spek, 2009).



**Figure 1**

Molecular structure of the title compound showing the atom labeling scheme and 50° probability displacement ellipsoids.

**Figure 2**

Packing diagram of the title compound viewed offset from along the *b* axis. Dashed lines indicate N—H $\cdots$ O and O—H $\cdots$ O hydrogen bonds. The intermolecular N—H $\cdots$ O hydrogen bonds form a two-dimensional network.

### 2,2'-(Piperazine-1,4-diyl)diethanaminium dibenzoate

#### Crystal data

$C_8H_{22}N_4^{2+} \cdot 2C_7H_5O_2^-$   
 $M_r = 416.52$   
 Monoclinic,  $P2_1/n$   
 Hall symbol: -P 2yn  
 $a = 19.5300(4) \text{ \AA}$   
 $b = 6.6694(2) \text{ \AA}$   
 $c = 19.6178(4) \text{ \AA}$   
 $\beta = 115.989(1)^\circ$   
 $V = 2296.89(10) \text{ \AA}^3$   
 $Z = 4$

$F(000) = 896$   
 $D_x = 1.204 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation,  $\lambda = 0.71070 \text{ \AA}$   
 Cell parameters from 12295 reflections  
 $\theta = 1.0\text{--}27.5^\circ$   
 $\mu = 0.08 \text{ mm}^{-1}$   
 $T = 180 \text{ K}$   
 Block, yellow  
 $0.28 \times 0.23 \times 0.12 \text{ mm}$

#### Data collection

Nonius KappaCCD  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 Thin slice  $\omega$  and  $\varphi$  scans  
 Absorption correction: multi-scan  
 (SORTAV; Blessing, 1995)  
 $T_{\min} = 0.910$ ,  $T_{\max} = 0.991$

20137 measured reflections  
 5194 independent reflections  
 3970 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.031$   
 $\theta_{\text{max}} = 27.5^\circ$ ,  $\theta_{\text{min}} = 3.6^\circ$   
 $h = -25 \rightarrow 22$   
 $k = -7 \rightarrow 8$   
 $l = -25 \rightarrow 25$

Refinement

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.040$   
 $wR(F^2) = 0.110$   
 $S = 1.02$   
 5194 reflections  
 290 parameters  
 0 restraints  
 Primary atom site location: structure-invariant  
 direct methods

Secondary atom site location: difference Fourier  
 map  
 Hydrogen site location: difference Fourier map  
 H atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0702P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.16 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.19472 (7)	0.2447 (2)	0.19552 (7)	0.0375 (3)
H1A	0.1776 (9)	0.374 (3)	0.2008 (8)	0.045*
H1B	0.2431 (9)	0.223 (2)	0.2302 (9)	0.045*
H1C	0.1652 (9)	0.142 (3)	0.2002 (8)	0.045*
C2	0.19529 (8)	0.2437 (3)	0.12037 (8)	0.0491 (4)
H2A	0.2127	0.1110	0.1116	0.059*
H2B	0.2319	0.3457	0.1198	0.059*
C3	0.11811 (9)	0.2872 (3)	0.05773 (8)	0.0512 (4)
H3A	0.0994	0.4156	0.0686	0.061*
H3B	0.1226	0.3034	0.0097	0.061*
N4	0.06208 (7)	0.12951 (18)	0.04783 (6)	0.0418 (3)
C5	-0.01461 (8)	0.2057 (2)	0.00103 (8)	0.0458 (4)
H5A	-0.0182	0.2489	-0.0487	0.055*
H5B	-0.0242	0.3241	0.0260	0.055*
C6	0.07429 (8)	-0.0487 (2)	0.01107 (8)	0.0483 (4)
H6A	0.1254	-0.1048	0.0430	0.058*
H6B	0.0726	-0.0106	-0.0384	0.058*
C7	0.46941 (7)	0.2132 (2)	0.30262 (7)	0.0347 (3)
C8	0.51949 (8)	0.3595 (3)	0.30237 (8)	0.0470 (4)
H8	0.5013	0.4909	0.2852	0.056*
C9	0.59622 (9)	0.3155 (3)	0.32714 (9)	0.0583 (5)
H9	0.6304	0.4175	0.3276	0.070*
C10	0.62302 (9)	0.1248 (3)	0.35110 (9)	0.0566 (5)
H10	0.6755	0.0952	0.3680	0.068*
C11	0.57335 (9)	-0.0231 (3)	0.35052 (8)	0.0512 (4)
H11	0.5915	-0.1552	0.3664	0.061*

C12	0.49691 (8)	0.0210 (2)	0.32676 (8)	0.0415 (3)
H12	0.4631	-0.0809	0.3270	0.050*
C13	0.38687 (7)	0.2669 (2)	0.27915 (7)	0.0340 (3)
O14	0.36433 (5)	0.43634 (15)	0.25215 (5)	0.0429 (2)
O15	0.34467 (5)	0.13526 (15)	0.28820 (6)	0.0435 (3)
N1'	0.18677 (7)	0.8130 (2)	0.69616 (7)	0.0387 (3)
H1'A	0.2019 (9)	0.704 (3)	0.6812 (9)	0.046*
H1'B	0.2193 (9)	0.845 (2)	0.7439 (10)	0.046*
H1'C	0.1828 (9)	0.917 (3)	0.6640 (9)	0.046*
C2'	0.11280 (9)	0.7645 (3)	0.69548 (8)	0.0490 (4)
H2'A	0.0946	0.8814	0.7141	0.059*
H2'B	0.1193	0.6506	0.7301	0.059*
C3'	0.05450 (8)	0.7108 (3)	0.61682 (8)	0.0484 (4)
H3'A	0.0703	0.5854	0.6006	0.058*
H3'B	0.0049	0.6857	0.6177	0.058*
N4'	0.04519 (6)	0.86872 (18)	0.56193 (6)	0.0409 (3)
C5'	0.00761 (8)	1.0480 (2)	0.57298 (8)	0.0458 (4)
H5'A	0.0379	1.1020	0.6246	0.055*
H5'B	-0.0434	1.0114	0.5683	0.055*
C6'	0.00043 (8)	0.7944 (2)	0.48472 (8)	0.0441 (4)
H6'A	-0.0507	0.7539	0.4785	0.053*
H6'B	0.0256	0.6747	0.4762	0.053*
C7'	0.19900 (7)	0.3513 (2)	0.53125 (7)	0.0351 (3)
C8'	0.17823 (8)	0.5420 (2)	0.50088 (8)	0.0425 (3)
H8'	0.1815	0.6515	0.5332	0.051*
C9'	0.15272 (9)	0.5741 (3)	0.42383 (9)	0.0515 (4)
H9'	0.1377	0.7044	0.4032	0.062*
C10'	0.14927 (9)	0.4151 (3)	0.37717 (9)	0.0533 (4)
H10'	0.1315	0.4365	0.3243	0.064*
C11'	0.17134 (9)	0.2262 (3)	0.40677 (8)	0.0518 (4)
H11'	0.1699	0.1181	0.3746	0.062*
C12'	0.19571 (8)	0.1943 (2)	0.48389 (8)	0.0440 (3)
H12'	0.2102	0.0634	0.5043	0.053*
C13'	0.22236 (7)	0.3122 (2)	0.61444 (7)	0.0355 (3)
O14'	0.23245 (6)	0.46100 (17)	0.65735 (5)	0.0521 (3)
O15'	0.23097 (6)	0.13489 (16)	0.63648 (6)	0.0478 (3)

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0311 (6)	0.0325 (7)	0.0436 (6)	-0.0018 (5)	0.0115 (5)	-0.0006 (5)
C2	0.0417 (8)	0.0570 (10)	0.0495 (8)	-0.0097 (7)	0.0207 (6)	-0.0086 (7)
C3	0.0576 (9)	0.0497 (10)	0.0432 (8)	-0.0072 (8)	0.0193 (7)	0.0004 (7)
N4	0.0427 (7)	0.0410 (7)	0.0360 (6)	-0.0020 (5)	0.0119 (5)	-0.0013 (5)
C5	0.0519 (9)	0.0402 (8)	0.0360 (7)	0.0059 (7)	0.0107 (6)	-0.0001 (6)
C6	0.0448 (8)	0.0519 (10)	0.0428 (7)	0.0073 (7)	0.0144 (6)	-0.0012 (7)
C7	0.0362 (7)	0.0386 (8)	0.0291 (6)	0.0007 (6)	0.0141 (5)	-0.0019 (5)
C8	0.0428 (8)	0.0523 (10)	0.0458 (8)	-0.0024 (7)	0.0195 (6)	0.0051 (7)
C9	0.0424 (9)	0.0790 (14)	0.0564 (9)	-0.0083 (9)	0.0242 (7)	0.0073 (8)
C10	0.0378 (8)	0.0872 (14)	0.0471 (8)	0.0120 (9)	0.0209 (6)	0.0019 (8)

C11	0.0468 (8)	0.0586 (11)	0.0473 (8)	0.0164 (8)	0.0198 (6)	0.0009 (7)
C12	0.0405 (8)	0.0430 (9)	0.0401 (7)	0.0045 (6)	0.0169 (6)	-0.0012 (6)
C13	0.0375 (7)	0.0331 (8)	0.0299 (6)	0.0004 (6)	0.0133 (5)	-0.0028 (5)
O14	0.0447 (5)	0.0350 (6)	0.0472 (5)	0.0067 (4)	0.0184 (4)	0.0049 (4)
O15	0.0346 (5)	0.0389 (6)	0.0540 (6)	-0.0003 (4)	0.0167 (4)	0.0055 (4)
N1'	0.0444 (7)	0.0342 (7)	0.0334 (6)	-0.0010 (5)	0.0132 (5)	0.0019 (5)
C2'	0.0499 (9)	0.0546 (10)	0.0447 (8)	-0.0003 (7)	0.0227 (7)	0.0058 (7)
C3'	0.0402 (8)	0.0472 (10)	0.0529 (9)	-0.0060 (7)	0.0159 (6)	0.0023 (7)
N4'	0.0353 (6)	0.0389 (7)	0.0423 (6)	-0.0011 (5)	0.0113 (5)	-0.0021 (5)
C5'	0.0419 (8)	0.0485 (9)	0.0432 (8)	-0.0010 (7)	0.0152 (6)	-0.0082 (7)
C6'	0.0376 (7)	0.0406 (9)	0.0476 (8)	-0.0017 (6)	0.0127 (6)	-0.0088 (6)
C7'	0.0308 (6)	0.0379 (8)	0.0375 (7)	-0.0013 (6)	0.0159 (5)	0.0026 (6)
C8'	0.0444 (8)	0.0390 (9)	0.0466 (8)	0.0006 (6)	0.0223 (6)	0.0053 (6)
C9'	0.0560 (9)	0.0530 (10)	0.0492 (8)	0.0069 (8)	0.0265 (7)	0.0185 (8)
C10'	0.0478 (8)	0.0758 (13)	0.0375 (7)	0.0002 (8)	0.0198 (6)	0.0123 (8)
C11'	0.0554 (9)	0.0616 (11)	0.0402 (8)	-0.0002 (8)	0.0226 (7)	-0.0054 (7)
C12'	0.0472 (8)	0.0404 (8)	0.0425 (8)	0.0024 (7)	0.0179 (6)	0.0008 (6)
C13'	0.0312 (7)	0.0365 (8)	0.0388 (7)	0.0026 (6)	0.0154 (5)	0.0036 (6)
O14'	0.0690 (7)	0.0428 (7)	0.0385 (5)	0.0114 (5)	0.0181 (5)	-0.0018 (5)
O15'	0.0582 (7)	0.0390 (6)	0.0432 (5)	-0.0003 (5)	0.0195 (5)	0.0086 (4)

*Geometric parameters (Å, °)*

N1—C2	1.4792 (19)	N1'—C2'	1.475 (2)
N1—H1A	0.945 (18)	N1'—H1'A	0.882 (19)
N1—H1B	0.902 (16)	N1'—H1'B	0.897 (17)
N1—H1C	0.925 (18)	N1'—H1'C	0.916 (18)
C2—C3	1.498 (2)	C2'—C3'	1.505 (2)
C2—H2A	0.9900	C2'—H2'A	0.9900
C2—H2B	0.9900	C2'—H2'B	0.9900
C3—N4	1.468 (2)	C3'—N4'	1.460 (2)
C3—H3A	0.9900	C3'—H3'A	0.9900
C3—H3B	0.9900	C3'—H3'B	0.9900
N4—C5	1.4625 (19)	N4'—C6'	1.4640 (18)
N4—C6	1.463 (2)	N4'—C5'	1.468 (2)
C5—C6 <sup>i</sup>	1.506 (2)	C5'—C6' <sup>ii</sup>	1.502 (2)
C5—H5A	0.9900	C5'—H5'A	0.9900
C5—H5B	0.9900	C5'—H5'B	0.9900
C6—C5 <sup>i</sup>	1.506 (2)	C6'—C5' <sup>ii</sup>	1.502 (2)
C6—H6A	0.9900	C6'—H6'A	0.9900
C6—H6B	0.9900	C6'—H6'B	0.9900
C7—C8	1.383 (2)	C7'—C12'	1.383 (2)
C7—C12	1.390 (2)	C7'—C8'	1.388 (2)
C7—C13	1.5127 (18)	C7'—C13'	1.5125 (18)
C8—C9	1.389 (2)	C8'—C9'	1.385 (2)
C8—H8	0.9500	C8'—H8'	0.9500
C9—C10	1.378 (3)	C9'—C10'	1.383 (3)
C9—H9	0.9500	C9'—H9'	0.9500
C10—C11	1.380 (3)	C10'—C11'	1.375 (3)
C10—H10	0.9500	C10'—H10'	0.9500

C11—C12	1.387 (2)	C11'—C12'	1.389 (2)
C11—H11	0.9500	C11'—H11'	0.9500
C12—H12	0.9500	C12'—H12'	0.9500
C13—O14	1.2436 (17)	C13'—O15'	1.2447 (17)
C13—O15	1.2690 (17)	C13'—O14'	1.2599 (17)
C2—N1—H1A	105.8 (9)	C2'—N1'—H1'A	106.7 (11)
C2—N1—H1B	106.8 (9)	C2'—N1'—H1'B	107.8 (10)
H1A—N1—H1B	111.6 (14)	H1'A—N1'—H1'B	110.8 (14)
C2—N1—H1C	111.9 (10)	C2'—N1'—H1'C	111.8 (10)
H1A—N1—H1C	113.4 (14)	H1'A—N1'—H1'C	109.5 (14)
H1B—N1—H1C	107.2 (14)	H1'B—N1'—H1'C	110.2 (15)
N1—C2—C3	111.87 (13)	N1'—C2'—C3'	111.07 (12)
N1—C2—H2A	109.2	N1'—C2'—H2'A	109.4
C3—C2—H2A	109.2	C3'—C2'—H2'A	109.4
N1—C2—H2B	109.2	N1'—C2'—H2'B	109.4
C3—C2—H2B	109.2	C3'—C2'—H2'B	109.4
H2A—C2—H2B	107.9	H2'A—C2'—H2'B	108.0
N4—C3—C2	113.21 (13)	N4'—C3'—C2'	112.22 (13)
N4—C3—H3A	108.9	N4'—C3'—H3'A	109.2
C2—C3—H3A	108.9	C2'—C3'—H3'A	109.2
N4—C3—H3B	108.9	N4'—C3'—H3'B	109.2
C2—C3—H3B	108.9	C2'—C3'—H3'B	109.2
H3A—C3—H3B	107.8	H3'A—C3'—H3'B	107.9
C5—N4—C6	108.42 (11)	C3'—N4'—C6'	110.11 (12)
C5—N4—C3	109.47 (12)	C3'—N4'—C5'	112.79 (12)
C6—N4—C3	111.96 (12)	C6'—N4'—C5'	108.54 (11)
N4—C5—C6 <sup>i</sup>	111.48 (13)	N4'—C5'—C6' <sup>ii</sup>	110.65 (12)
N4—C5—H5A	109.3	N4'—C5'—H5'A	109.5
C6 <sup>i</sup> —C5—H5A	109.3	C6' <sup>ii</sup> —C5'—H5'A	109.5
N4—C5—H5B	109.3	N4'—C5'—H5'B	109.5
C6 <sup>i</sup> —C5—H5B	109.3	C6' <sup>ii</sup> —C5'—H5'B	109.5
H5A—C5—H5B	108.0	H5'A—C5'—H5'B	108.1
N4—C6—C5 <sup>i</sup>	111.13 (12)	N4'—C6'—C5' <sup>ii</sup>	111.14 (12)
N4—C6—H6A	109.4	N4'—C6'—H6'A	109.4
C5 <sup>i</sup> —C6—H6A	109.4	C5' <sup>ii</sup> —C6'—H6'A	109.4
N4—C6—H6B	109.4	N4'—C6'—H6'B	109.4
C5 <sup>i</sup> —C6—H6B	109.4	C5' <sup>ii</sup> —C6'—H6'B	109.4
H6A—C6—H6B	108.0	H6'A—C6'—H6'B	108.0
C8—C7—C12	118.90 (13)	C12'—C7'—C8'	119.15 (12)
C8—C7—C13	119.50 (13)	C12'—C7'—C13'	119.82 (13)
C12—C7—C13	121.57 (12)	C8'—C7'—C13'	120.99 (13)
C7—C8—C9	120.39 (16)	C9'—C8'—C7'	120.50 (14)
C7—C8—H8	119.8	C9'—C8'—H8'	119.7
C9—C8—H8	119.8	C7'—C8'—H8'	119.7
C10—C9—C8	120.31 (16)	C10'—C9'—C8'	119.60 (15)
C10—C9—H9	119.8	C10'—C9'—H9'	120.2
C8—C9—H9	119.8	C8'—C9'—H9'	120.2
C9—C10—C11	119.78 (15)	C11'—C10'—C9'	120.50 (14)



C9—C10—H10	120.1	C11'—C10'—H10'	119.7
C11—C10—H10	120.1	C9'—C10'—H10'	119.7
C10—C11—C12	120.04 (16)	C10'—C11'—C12'	119.65 (16)
C10—C11—H11	120.0	C10'—C11'—H11'	120.2
C12—C11—H11	120.0	C12'—C11'—H11'	120.2
C11—C12—C7	120.57 (14)	C7'—C12'—C11'	120.56 (15)
C11—C12—H12	119.7	C7'—C12'—H12'	119.7
C7—C12—H12	119.7	C11'—C12'—H12'	119.7
O14—C13—O15	123.91 (12)	O15'—C13'—O14'	123.96 (12)
O14—C13—C7	118.44 (12)	O15'—C13'—C7'	118.02 (12)
O15—C13—C7	117.65 (12)	O14'—C13'—C7'	118.01 (12)
N1—C2—C3—N4	-66.32 (18)	N1'—C2'—C3'—N4'	-56.03 (18)
C2—C3—N4—C5	164.89 (13)	C2'—C3'—N4'—C6'	168.40 (13)
C2—C3—N4—C6	-74.84 (16)	C2'—C3'—N4'—C5'	-70.19 (16)
C6—N4—C5—C6 <sup>i</sup>	57.36 (17)	C3'—N4'—C5'—C6' <sup>ii</sup>	179.90 (11)
C3—N4—C5—C6 <sup>i</sup>	179.76 (12)	C6'—N4'—C5'—C6' <sup>ii</sup>	-57.80 (16)
C5—N4—C6—C5 <sup>i</sup>	-57.14 (17)	C3'—N4'—C6'—C5' <sup>ii</sup>	-177.99 (12)
C3—N4—C6—C5 <sup>i</sup>	-178.01 (11)	C5'—N4'—C6'—C5' <sup>ii</sup>	58.10 (16)
C12—C7—C8—C9	-1.1 (2)	C12'—C7'—C8'—C9'	1.6 (2)
C13—C7—C8—C9	176.94 (13)	C13'—C7'—C8'—C9'	-176.24 (13)
C7—C8—C9—C10	1.0 (2)	C7'—C8'—C9'—C10'	-1.2 (2)
C8—C9—C10—C11	-0.1 (2)	C8'—C9'—C10'—C11'	-0.3 (2)
C9—C10—C11—C12	-0.8 (2)	C9'—C10'—C11'—C12'	1.4 (2)
C10—C11—C12—C7	0.7 (2)	C8'—C7'—C12'—C11'	-0.5 (2)
C8—C7—C12—C11	0.28 (19)	C13'—C7'—C12'—C11'	177.32 (13)
C13—C7—C12—C11	-177.74 (12)	C10'—C11'—C12'—C7'	-0.9 (2)
C8—C7—C13—O14	7.89 (17)	C12'—C7'—C13'—O15'	-6.79 (19)
C12—C7—C13—O14	-174.10 (12)	C8'—C7'—C13'—O15'	171.02 (13)
C8—C7—C13—O15	-172.22 (12)	C12'—C7'—C13'—O14'	172.71 (13)
C12—C7—C13—O15	5.79 (17)	C8'—C7'—C13'—O14'	-9.48 (19)

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

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A $\cdots$ O15 <sup>iii</sup>	0.945 (18)	1.833 (19)	2.7739 (17)	173.0 (15)
N1—H1B $\cdots$ O15	0.902 (16)	1.895 (17)	2.7836 (15)	167.8 (14)
N1—H1B $\cdots$ O14	0.902 (16)	2.632 (16)	3.2580 (16)	127.2 (13)
N1—H1C $\cdots$ O14 <sup>iv</sup>	0.925 (18)	1.887 (18)	2.7660 (16)	157.9 (14)
N1'—H1'A $\cdots$ O14'	0.882 (19)	1.857 (19)	2.7355 (17)	174.2 (15)
N1'—H1'B $\cdots$ O14 <sup>v</sup>	0.897 (17)	1.908 (17)	2.7836 (16)	164.8 (15)
N1'—H1'B $\cdots$ O15 <sup>v</sup>	0.897 (17)	2.533 (16)	3.1858 (16)	130.1 (13)
N1'—H1'C $\cdots$ O15 <sup>vi</sup>	0.916 (18)	1.934 (18)	2.7585 (16)	148.8 (14)

Symmetry codes: (iii)  $-x+1/2, y+1/2, -z+1/2$ ; (iv)  $-x+1/2, y-1/2, -z+1/2$ ; (v)  $-x+1/2, y+1/2, -z+3/2$ ; (vi)  $x, y+1, z$ .