

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

4-(Furan-2-carbonyl)piperazin-1-ium
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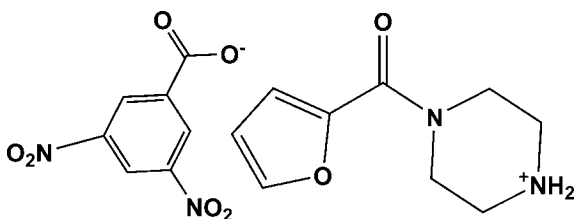
Received 14 May 2014; accepted 15 May 2014 Edited by E. R. T. Tiekink, University of Malaya, Malaysia

Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; disorder in main residue; R factor = 0.043; wR factor = 0.123; data-to-parameter ratio = 12.4.

In the cation of the title salt, $\text{C}_9\text{H}_{13}\text{N}_2\text{O}_2^+ \cdot \text{C}_7\text{H}_3\text{N}_2\text{O}_6^-$, the piperazine ring adopts a slightly distorted chair conformation. Twofold rotational disorder is exhibited by the furan ring in a 0.430 (4):0.570 (4) ratio. In the crystal, $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds link the ions into chains along [010]. Additional weak $\text{C}-\text{H} \cdots \text{O}$ interactions are observed, leading to a supra-molecular layer parallel to (011).

Related literature

For the synthesis of the drug Prazosin [systematic name: 2-[4-(2-furoyl)piperazin-1-yl]-6,7-dimethoxyquinazolin-4-amine], see: Honkanen *et al.* (1980). For the drug 1(2-furoyl)piperazine, used in the treatment of high blood pressure and anxiety, see: Brogden *et al.* (1977). For therapeutic uses of piperazines, see: Brockunier *et al.* (2004); Bogatcheva *et al.* (2006). For the use of the piperazine moiety in the construction of bioactive molecules, see: Choudhary *et al.* (2006). For a related structure, see: Dayananda *et al.* (2012). For puckering parameters, see: Cremer & Pople (1975).



Experimental

Crystal data

 $\text{C}_9\text{H}_{13}\text{N}_2\text{O}_2^+ \cdot \text{C}_7\text{H}_3\text{N}_2\text{O}_6^-$
 $M_r = 392.33$ Orthorhombic, $Pbca$
 $a = 9.6060$ (2) Å $b = 10.4572$ (2) Å
 $c = 33.8766$ (7) Å
 $V = 3402.97$ (13) Å³
 $Z = 8$ Cu $K\alpha$ radiation
 $\mu = 1.08$ mm⁻¹
 $T = 173$ K
 $0.28 \times 0.22 \times 0.18$ mm

Data collection

Agilent Eos Gemini diffractometer
Absorption correction: multi-scan
(*CrysAlis PRO* and *CrysAlis RED*; Agilent, 2012)
 $T_{\min} = 0.863$, $T_{\max} = 1.000$ 21195 measured reflections
3352 independent reflections
2915 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.079$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.123$
 $S = 1.02$
3352 reflections
270 parameters10 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.26$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻³**Table 1**
Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{N2B}-\text{H2BA} \cdots \text{O1A}^i$	0.99	2.51	3.1607 (16)	123
$\text{N2B}-\text{H2BA} \cdots \text{O2A}^i$	0.99	1.72	2.7093 (16)	176
$\text{N2B}-\text{H2BB} \cdots \text{O1A}^{ii}$	0.99	1.77	2.7424 (16)	166
$\text{C5A}-\text{H5A} \cdots \text{O2A}^{iii}$	0.95	2.47	3.3170 (18)	148
$\text{C9B}-\text{H9B} \cdots \text{O6A}^{iv}$	0.95	2.44	3.183 (6)	134
$\text{C8BB}-\text{H8BB} \cdots \text{O3A}^v$	0.95	2.50	3.395 (5)	158
$\text{C2B}-\text{H2BC} \cdots \text{O1B}^i$	0.99	2.59	3.2300 (19)	122

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

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis RED* (Agilent, 2012); program(s) used to solve structure: *SUPERFLIP* (Palatinus & Chapuis, 2007); program(s) used to refine structure: *SHELXL2012* (Sheldrick, 2008); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2*.

CNK thanks the University of Mysore for research facilities and is also grateful to the Principal, Maharani's Science College for Women, Mysore, for giving permission to undertake research. JPJ acknowledges the NSF-MRI program (grant No. CHE-1039027) for funds to purchase the X-ray diffractometer.

Supporting information for this paper is available from the IUCr electronic archives (Reference: TK5314).

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supporting information

Acta Cryst. (2014). E70, o700–o701 [doi:10.1107/S160053681401126X]

4-(Furan-2-carbonyl)piperazin-1-ium 3,5-dinitrobenzoate

Channappa N. Kavitha, Manpreet Kaur, Jerry P. Jasinski, Ray J. Butcher and H.S. Yathirajan

S1. Structural commentary

1(2-Furoyl)piperazine, used to synthesise the drug Prazosin (Honkanen *et al.*, 1980), is the first of a new class of anti-hypertensives. It is a sympatholytic drug used to treat high blood pressure and anxiety (Brogden *et al.*, 1977). Piperazines are found in biologically active compounds across a number of different therapeutic areas (Brockunier *et al.*, 2004; Bogatcheva *et al.*, 2006). The piperazine moiety is extensively employed to construct various bioactive molecules with anti-bacterial and anti-malarial activity, and as anti-psychotic agents (Choudhary *et al.*, 2006). The crystal structures of a similar salt *viz.*, cinnarizinium 3,5-dinitrosalicylate (Dayananda *et al.*, 2012) has been reported. In view of the above importance of piperazines, this paper reports the crystal structure of the title salt, (I) $C_9H_{13}N_2O_2^+ \cdot C_7H_3N_2O_6^-$.

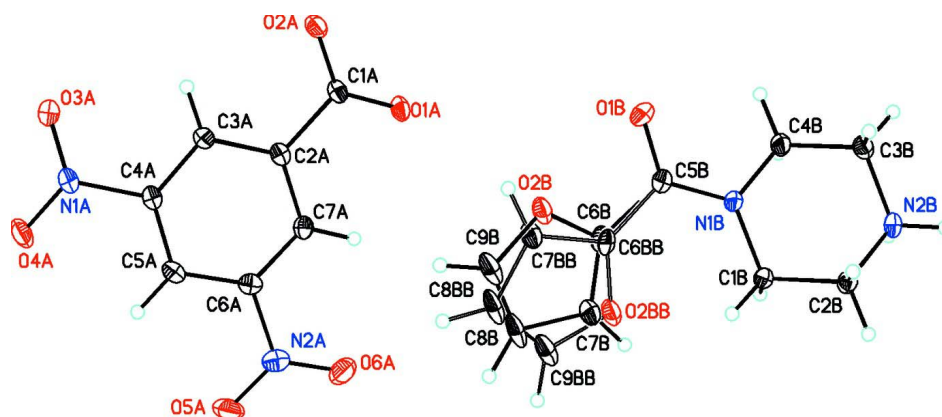
The title compound, (I), crystallizes with one independent piperazinium cation and a 3,5-dinitrobenzoate anion in the asymmetric unit (Fig. 1). In the cation, the piperazine ring adopts a slightly distorted chair conformation with puckering parameters Q , θ , and $\varphi = 0.5552(15)\text{\AA}$, $173.13(14)^\circ$ and $4.2(14)^\circ$, respectively (Cremer & Pople, 1975). Two-fold rotational disorder is exhibited by the furan ring in a 0.430(4):0.570(4) ratio represents two different conformations of the molecule that exist in the same crystal form. N—H \cdots O intermolecular hydrogen bonds link the cations and anions into infinite 1-D chains along [0 1 0] (Fig. 2). Additional weak C—H \cdots O intermolecular interactions are observed (Table 1) forming chains along [0 0 1] resulting in a 2-D supramolecular network structure.

S2. Synthesis and crystallization

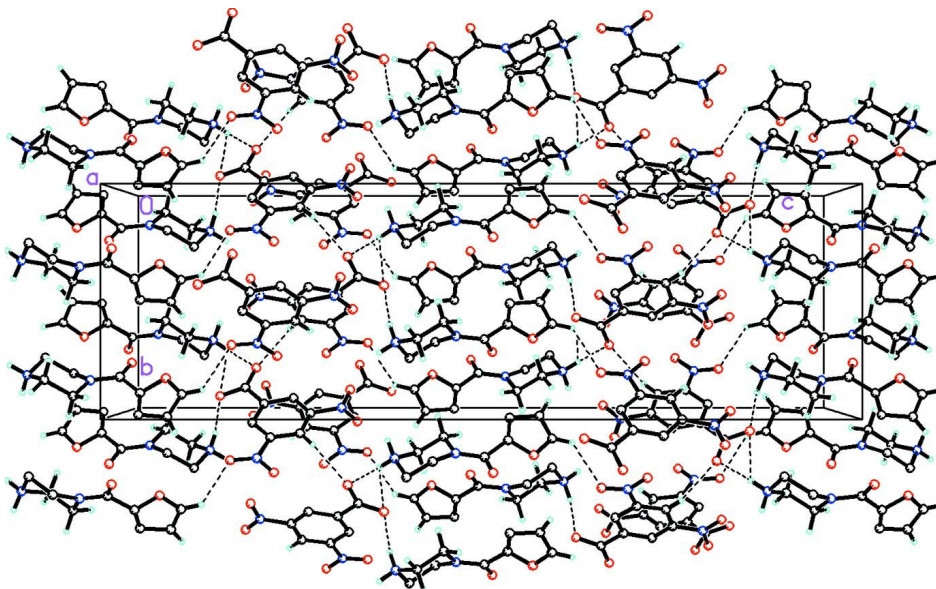
1(2-Furoyl)piperazine (0.9 g, 0.005 mol) and 3,5-dinitrobenzoic acid (1.0 g, 0.005 mol) were dissolved in *N,N*-dimethylformamide and stirred for 10 minutes at 333 K. The resulting solution was allowed to cool slowly at room temperature. The crystals of salt (I) (M.pt: 453–459 K) appeared after a few days.

S3. Refinement

All of the H atoms were placed in their calculated positions and then refined using the riding model with atom—H lengths of 0.95 Å (CH); 0.99 Å (CH₂) and 0.92 Å (NH₂), and with $U_{iso} = 1.2 \times U_{eq}(\text{parent atom})$. Owing to poor agreement, the following reflections were omitted from the final cycles of refinement: (2 2 2), (1 2 3), (1 0 6), (0 2 2), (0 4 1), (2 3 0), (1 2 4), (2 1 2), (1 2 1), (2 3 2) and (1 1 13).

**Figure 1**

ORTEP drawing of (I) ($C_9H_{13}N_2O_2^+$, $C_7H_3N_2O_6^-$) showing the labeling scheme with 30% probability displacement ellipsoids. Two-fold rotational disorder exhibited by the furan ring in a 0.430 (4):0.570 (4) ratio is displayed.

**Figure 2**

Molecular packing for (I) viewed along the *a* axis. Dashed lines indicate N—H \cdots O intermolecular hydrogen bonds forming infinite chain along the *b* axis and weak C—H \cdots O intermolecular interactions. Only the major disordered component [0.570 (4)] of the furan ring is displayed. H atoms not involved in hydrogen bonding have been removed for clarity.

4-(Furan-2-carbonyl)piperazin-1-ium 3,5-dinitrobenzoate

Crystal data

$C_9H_{13}N_2O_2^+ \cdot C_7H_3N_2O_6^-$

$M_r = 392.33$

Orthorhombic, *Pbca*

$a = 9.6060$ (2) Å

$b = 10.4572$ (2) Å

$c = 33.8766$ (7) Å

$V = 3402.97$ (13) Å³

$Z = 8$

$F(000) = 1632$

$D_x = 1.532$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 7148 reflections

$\theta = 3.9\text{--}72.3^\circ$

$\mu = 1.08$ mm⁻¹

$T = 173$ K
Irregular, colourless

$0.28 \times 0.22 \times 0.18$ mm

Data collection

Agilent Eos Gemini
diffractometer
Detector resolution: 16.0416 pixels mm^{-1}
 ω scans
Absorption correction: multi-scan
(*CrysAlis PRO* and *CrysAlis RED*; Agilent,
2012)
 $T_{\min} = 0.863$, $T_{\max} = 1.000$

21195 measured reflections
3352 independent reflections
2915 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.079$
 $\theta_{\max} = 72.4^\circ$, $\theta_{\min} = 5.2^\circ$
 $h = -10 \rightarrow 11$
 $k = -12 \rightarrow 12$
 $l = -30 \rightarrow 41$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.123$
 $S = 1.02$
3352 reflections
270 parameters
10 restraints
Primary atom site location: structure-invariant
direct methods

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0723P)^2 + 0.8847P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.26 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL2012* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0015 (3)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1A	0.36870 (12)	0.57178 (10)	0.63310 (3)	0.0284 (3)	
O2A	0.28703 (12)	0.69735 (10)	0.68122 (3)	0.0317 (3)	
O3A	0.42159 (14)	0.60237 (15)	0.81693 (3)	0.0463 (4)	
O4A	0.58504 (14)	0.46802 (13)	0.82985 (3)	0.0415 (3)	
O5A	0.82825 (13)	0.26144 (13)	0.72229 (4)	0.0456 (3)	
O6A	0.77657 (15)	0.31036 (13)	0.66197 (4)	0.0468 (4)	
N1A	0.51100 (13)	0.52552 (13)	0.80671 (4)	0.0281 (3)	
N2A	0.75838 (14)	0.31806 (13)	0.69759 (4)	0.0326 (3)	
C1A	0.36297 (15)	0.60956 (14)	0.66819 (4)	0.0239 (3)	
C2A	0.45703 (14)	0.54250 (13)	0.69778 (4)	0.0224 (3)	
C3A	0.44092 (15)	0.56475 (13)	0.73799 (4)	0.0223 (3)	
H3A	0.3713	0.6217	0.7473	0.027*	
C4A	0.52856 (15)	0.50207 (14)	0.76415 (4)	0.0233 (3)	
C5A	0.63161 (15)	0.41803 (14)	0.75234 (4)	0.0252 (3)	
H5A	0.6891	0.3744	0.7708	0.030*	
C6A	0.64582 (15)	0.40151 (14)	0.71212 (4)	0.0254 (3)	
C7A	0.56135 (15)	0.46153 (14)	0.68461 (4)	0.0241 (3)	

H7A	0.5748	0.4474	0.6572	0.029*	
O1B	0.49213 (15)	0.72567 (16)	0.51522 (4)	0.0557 (4)	
O2B	0.6605 (4)	0.6627 (4)	0.57013 (17)	0.0347 (6)	0.430 (4)
C6B	0.672 (3)	0.602 (3)	0.5349 (8)	0.023 (2)	0.430 (4)
C7B	0.7724 (9)	0.5082 (8)	0.53851 (19)	0.0311 (8)	0.430 (4)
H7B	0.8108	0.4585	0.5177	0.037*	0.430 (4)
C8B	0.8067 (7)	0.4998 (7)	0.5786 (2)	0.0392 (15)	0.430 (4)
H8B	0.8656	0.4388	0.5910	0.047*	0.430 (4)
C9B	0.7390 (6)	0.5956 (6)	0.59554 (18)	0.0405 (9)	0.430 (4)
H9B	0.7454	0.6149	0.6229	0.049*	0.430 (4)
O2BB	0.7728 (4)	0.5083 (4)	0.52700 (8)	0.0347 (6)	0.570 (4)
C6BB	0.6943 (19)	0.613 (2)	0.5349 (6)	0.023 (2)	0.570 (4)
C7BB	0.6920 (4)	0.6323 (4)	0.57507 (17)	0.0311 (8)	0.570 (4)
H7BB	0.6370	0.6936	0.5887	0.037*	0.570 (4)
C8BB	0.7856 (5)	0.5455 (5)	0.59227 (13)	0.0392 (15)	0.570 (4)
H8BB	0.8115	0.5390	0.6193	0.047*	0.570 (4)
C9BB	0.8307 (4)	0.4732 (4)	0.56174 (14)	0.0405 (9)	0.570 (4)
H9BB	0.8958	0.4054	0.5644	0.049*	0.570 (4)
N1B	0.62643 (12)	0.65199 (13)	0.46545 (3)	0.0251 (3)	
N2B	0.69338 (13)	0.67832 (11)	0.38373 (3)	0.0233 (3)	
H2BA	0.7311	0.7252	0.3607	0.028*	
H2BB	0.6603	0.5938	0.3745	0.028*	
C1B	0.75009 (15)	0.58881 (15)	0.44944 (4)	0.0248 (3)	
H1BA	0.7264	0.5001	0.4417	0.030*	
H1BB	0.8233	0.5849	0.4700	0.030*	
C2B	0.80476 (15)	0.66087 (15)	0.41378 (4)	0.0262 (3)	
H2BC	0.8400	0.7456	0.4222	0.031*	
H2BD	0.8832	0.6128	0.4020	0.031*	
C3B	0.57552 (16)	0.75100 (15)	0.40107 (4)	0.0273 (3)	
H3BA	0.5016	0.7624	0.3810	0.033*	
H3BB	0.6079	0.8367	0.4094	0.033*	
C4B	0.51769 (15)	0.67938 (17)	0.43631 (4)	0.0301 (4)	
H4BA	0.4435	0.7311	0.4488	0.036*	
H4BB	0.4756	0.5980	0.4273	0.036*	
C5B	0.59507 (16)	0.66629 (16)	0.50423 (4)	0.0299 (4)	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1A	0.0419 (6)	0.0242 (5)	0.0189 (5)	0.0007 (4)	-0.0049 (4)	-0.0006 (4)
O2A	0.0430 (6)	0.0275 (6)	0.0246 (5)	0.0102 (5)	-0.0106 (4)	-0.0032 (4)
O3A	0.0509 (8)	0.0651 (9)	0.0228 (6)	0.0221 (7)	0.0003 (5)	-0.0018 (5)
O4A	0.0528 (8)	0.0454 (7)	0.0262 (6)	0.0075 (6)	-0.0137 (5)	0.0072 (5)
O5A	0.0365 (7)	0.0445 (7)	0.0560 (8)	0.0169 (6)	-0.0095 (6)	-0.0018 (6)
O6A	0.0505 (8)	0.0472 (8)	0.0427 (7)	0.0127 (6)	0.0179 (6)	0.0016 (6)
N1A	0.0314 (6)	0.0315 (7)	0.0216 (6)	-0.0008 (5)	-0.0046 (5)	0.0036 (5)
N2A	0.0269 (6)	0.0275 (7)	0.0433 (8)	0.0004 (5)	0.0028 (6)	0.0001 (6)
C1A	0.0305 (7)	0.0198 (7)	0.0214 (7)	-0.0021 (6)	-0.0054 (6)	0.0016 (5)

C2A	0.0249 (7)	0.0206 (7)	0.0218 (7)	-0.0045 (5)	-0.0035 (5)	0.0024 (5)
C3A	0.0233 (7)	0.0204 (7)	0.0233 (7)	-0.0013 (5)	-0.0028 (5)	0.0001 (5)
C4A	0.0252 (7)	0.0240 (7)	0.0207 (6)	-0.0041 (6)	-0.0027 (5)	0.0023 (5)
C5A	0.0234 (7)	0.0235 (7)	0.0287 (7)	-0.0017 (5)	-0.0057 (6)	0.0043 (6)
C6A	0.0223 (7)	0.0224 (7)	0.0316 (7)	-0.0014 (5)	-0.0004 (6)	0.0009 (6)
C7A	0.0268 (7)	0.0238 (7)	0.0215 (7)	-0.0038 (6)	-0.0004 (5)	-0.0007 (5)
O1B	0.0552 (8)	0.0842 (11)	0.0278 (6)	0.0375 (8)	0.0117 (6)	0.0020 (6)
O2B	0.0447 (11)	0.0406 (11)	0.0188 (11)	0.0075 (8)	-0.0056 (12)	0.0038 (13)
C6B	0.019 (5)	0.029 (3)	0.0200 (7)	-0.010 (4)	0.003 (3)	0.0016 (14)
C7B	0.0344 (14)	0.0367 (15)	0.022 (2)	0.0009 (11)	-0.0054 (17)	0.0024 (18)
C8B	0.044 (2)	0.049 (3)	0.024 (2)	-0.016 (3)	-0.017 (2)	0.015 (2)
C9B	0.0447 (16)	0.0482 (18)	0.0284 (14)	-0.0067 (13)	-0.0148 (15)	0.0141 (16)
O2BB	0.0447 (11)	0.0406 (11)	0.0188 (11)	0.0075 (8)	-0.0056 (12)	0.0038 (13)
C6BB	0.019 (5)	0.029 (3)	0.0200 (7)	-0.010 (4)	0.003 (3)	0.0016 (14)
C7BB	0.0344 (14)	0.0367 (15)	0.022 (2)	0.0009 (11)	-0.0054 (17)	0.0024 (18)
C8BB	0.044 (2)	0.049 (3)	0.024 (2)	-0.016 (3)	-0.017 (2)	0.015 (2)
C9BB	0.0447 (16)	0.0482 (18)	0.0284 (14)	-0.0067 (13)	-0.0148 (15)	0.0141 (16)
N1B	0.0236 (6)	0.0343 (7)	0.0174 (6)	0.0045 (5)	0.0012 (4)	0.0017 (5)
N2B	0.0292 (6)	0.0245 (6)	0.0163 (5)	-0.0049 (5)	0.0008 (5)	0.0003 (4)
C1B	0.0267 (7)	0.0294 (7)	0.0184 (6)	0.0052 (6)	0.0015 (5)	-0.0009 (5)
C2B	0.0242 (7)	0.0351 (8)	0.0192 (7)	-0.0006 (6)	0.0010 (5)	-0.0008 (6)
C3B	0.0303 (7)	0.0289 (7)	0.0227 (7)	0.0020 (6)	-0.0026 (6)	0.0030 (6)
C4B	0.0233 (7)	0.0444 (9)	0.0227 (7)	0.0024 (6)	0.0003 (6)	0.0059 (6)
C5B	0.0330 (8)	0.0354 (8)	0.0212 (7)	0.0056 (7)	0.0042 (6)	0.0007 (6)

Geometric parameters (Å, °)

O1A—C1A	1.2539 (17)	C9B—H9B	0.9500
O2A—C1A	1.2529 (18)	O2BB—C6BB	1.35 (3)
O3A—N1A	1.2261 (18)	O2BB—C9BB	1.352 (5)
O4A—N1A	1.2173 (17)	C6BB—C7BB	1.38 (2)
O5A—N2A	1.2251 (19)	C6BB—C5B	1.517 (9)
O6A—N2A	1.2220 (19)	C7BB—H7BB	0.9500
N1A—C4A	1.4722 (18)	C7BB—C8BB	1.405 (6)
N2A—C6A	1.4741 (19)	C8BB—H8BB	0.9500
C1A—C2A	1.5208 (19)	C8BB—C9BB	1.352 (6)
C2A—C3A	1.3905 (19)	C9BB—H9BB	0.9500
C2A—C7A	1.386 (2)	N1B—C1B	1.4634 (18)
C3A—H3A	0.9500	N1B—C4B	1.4656 (18)
C3A—C4A	1.387 (2)	N1B—C5B	1.3563 (19)
C4A—C5A	1.383 (2)	N2B—H2BA	0.9900
C5A—H5A	0.9500	N2B—H2BB	0.9900
C5A—C6A	1.380 (2)	N2B—C2B	1.4880 (18)
C6A—C7A	1.386 (2)	N2B—C3B	1.4848 (19)
C7A—H7A	0.9500	C1B—H1BA	0.9900
O1B—C5B	1.226 (2)	C1B—H1BB	0.9900
O2B—C6B	1.36 (3)	C1B—C2B	1.5178 (19)
O2B—C9B	1.342 (6)	C2B—H2BC	0.9900

C6B—C7B	1.38 (3)	C2B—H2BD	0.9900
C6B—C5B	1.444 (12)	C3B—H3BA	0.9900
C7B—H7B	0.9500	C3B—H3BB	0.9900
C7B—C8B	1.401 (8)	C3B—C4B	1.515 (2)
C8B—H8B	0.9500	C4B—H4BA	0.9900
C8B—C9B	1.325 (9)	C4B—H4BB	0.9900
O3A—N1A—C4A	117.76 (12)	C8BB—C7BB—H7BB	126.2
O4A—N1A—O3A	123.46 (13)	C7BB—C8BB—H8BB	127.8
O4A—N1A—C4A	118.78 (13)	C9BB—C8BB—C7BB	104.4 (4)
O5A—N2A—C6A	117.39 (14)	C9BB—C8BB—H8BB	127.8
O6A—N2A—O5A	124.34 (14)	O2BB—C9BB—H9BB	123.7
O6A—N2A—C6A	118.27 (13)	C8BB—C9BB—O2BB	112.5 (4)
O1A—C1A—C2A	116.97 (13)	C8BB—C9BB—H9BB	123.7
O2A—C1A—O1A	126.19 (13)	C1B—N1B—C4B	114.66 (11)
O2A—C1A—C2A	116.84 (12)	C5B—N1B—C1B	126.09 (12)
C3A—C2A—C1A	120.16 (13)	C5B—N1B—C4B	118.21 (12)
C7A—C2A—C1A	119.94 (13)	H2BA—N2B—H2BB	108.2
C7A—C2A—C3A	119.88 (13)	C2B—N2B—H2BA	109.7
C2A—C3A—H3A	120.7	C2B—N2B—H2BB	109.7
C4A—C3A—C2A	118.65 (13)	C3B—N2B—H2BA	109.7
C4A—C3A—H3A	120.7	C3B—N2B—H2BB	109.7
C3A—C4A—N1A	118.53 (13)	C3B—N2B—C2B	109.90 (11)
C5A—C4A—N1A	118.11 (12)	N1B—C1B—H1BA	109.5
C5A—C4A—C3A	123.36 (13)	N1B—C1B—H1BB	109.5
C4A—C5A—H5A	122.1	N1B—C1B—C2B	110.59 (12)
C6A—C5A—C4A	115.85 (13)	H1BA—C1B—H1BB	108.1
C6A—C5A—H5A	122.1	C2B—C1B—H1BA	109.5
C5A—C6A—N2A	118.44 (13)	C2B—C1B—H1BB	109.5
C5A—C6A—C7A	123.33 (14)	N2B—C2B—C1B	110.90 (12)
C7A—C6A—N2A	118.22 (13)	N2B—C2B—H2BC	109.5
C2A—C7A—C6A	118.91 (13)	N2B—C2B—H2BD	109.5
C2A—C7A—H7A	120.5	C1B—C2B—H2BC	109.5
C6A—C7A—H7A	120.5	C1B—C2B—H2BD	109.5
C9B—O2B—C6B	105.8 (9)	H2BC—C2B—H2BD	108.0
O2B—C6B—C7B	108.3 (9)	N2B—C3B—H3BA	109.7
O2B—C6B—C5B	112 (2)	N2B—C3B—H3BB	109.7
C7B—C6B—C5B	139 (3)	N2B—C3B—C4B	109.78 (12)
C6B—C7B—H7B	126.5	H3BA—C3B—H3BB	108.2
C6B—C7B—C8B	107.1 (13)	C4B—C3B—H3BA	109.7
C8B—C7B—H7B	126.5	C4B—C3B—H3BB	109.7
C7B—C8B—H8B	127.6	N1B—C4B—C3B	111.47 (12)
C9B—C8B—C7B	104.9 (5)	N1B—C4B—H4BA	109.3
C9B—C8B—H8B	127.6	N1B—C4B—H4BB	109.3
O2B—C9B—H9B	123.4	C3B—C4B—H4BA	109.3
C8B—C9B—O2B	113.2 (5)	C3B—C4B—H4BB	109.3
C8B—C9B—H9B	123.4	H4BA—C4B—H4BB	108.0
C6BB—O2BB—C9BB	106.0 (7)	O1B—C5B—C6B	115.6 (15)

O2BB—C6BB—C7BB	109.0 (7)	O1B—C5B—C6BB	119.1 (10)
O2BB—C6BB—C5B	120.9 (17)	O1B—C5B—N1B	121.96 (15)
C7BB—C6BB—C5B	127.6 (18)	N1B—C5B—C6B	122.1 (15)
C6BB—C7BB—H7BB	126.2	N1B—C5B—C6BB	118.8 (10)
C6BB—C7BB—C8BB	107.6 (9)		
O1A—C1A—C2A—C3A	-169.82 (13)	C7B—C6B—C5B—N1B	-10 (4)
O1A—C1A—C2A—C7A	11.8 (2)	C7B—C8B—C9B—O2B	2.0 (7)
O2A—C1A—C2A—C3A	10.2 (2)	C9B—O2B—C6B—C7B	-8 (2)
O2A—C1A—C2A—C7A	-168.20 (14)	C9B—O2B—C6B—C5B	-178.6 (14)
O3A—N1A—C4A—C3A	-1.9 (2)	O2BB—C6BB—C7BB—C8BB	6.9 (14)
O3A—N1A—C4A—C5A	178.35 (15)	O2BB—C6BB—C5B—O1B	153.1 (11)
O4A—N1A—C4A—C3A	177.96 (14)	O2BB—C6BB—C5B—C6B	85 (17)
O4A—N1A—C4A—C5A	-1.8 (2)	O2BB—C6BB—C5B—N1B	-29.8 (18)
O5A—N2A—C6A—C5A	4.4 (2)	C6BB—O2BB—C9BB—C8BB	4.1 (10)
O5A—N2A—C6A—C7A	-176.94 (14)	C6BB—C7BB—C8BB—C9BB	-4.2 (10)
O6A—N2A—C6A—C5A	-175.20 (14)	C7BB—C6BB—C5B—O1B	-7 (2)
O6A—N2A—C6A—C7A	3.5 (2)	C7BB—C6BB—C5B—C6B	-75 (17)
N1A—C4A—C5A—C6A	-178.63 (12)	C7BB—C6BB—C5B—N1B	170.0 (13)
N2A—C6A—C7A—C2A	-178.59 (13)	C7BB—C8BB—C9BB—O2BB	0.1 (5)
C1A—C2A—C3A—C4A	-179.95 (12)	C9BB—O2BB—C6BB—C7BB	-6.7 (14)
C1A—C2A—C7A—C6A	179.97 (13)	C9BB—O2BB—C6BB—C5B	-170.1 (11)
C2A—C3A—C4A—N1A	-179.82 (12)	N1B—C1B—C2B—N2B	-54.25 (16)
C2A—C3A—C4A—C5A	-0.1 (2)	N2B—C3B—C4B—N1B	55.11 (17)
C3A—C2A—C7A—C6A	1.6 (2)	C1B—N1B—C4B—C3B	-52.13 (18)
C3A—C4A—C5A—C6A	1.6 (2)	C1B—N1B—C5B—O1B	176.66 (17)
C4A—C5A—C6A—N2A	177.02 (13)	C1B—N1B—C5B—C6B	-9.7 (15)
C4A—C5A—C6A—C7A	-1.6 (2)	C1B—N1B—C5B—C6BB	-0.3 (10)
C5A—C6A—C7A—C2A	0.0 (2)	C2B—N2B—C3B—C4B	-59.34 (15)
C7A—C2A—C3A—C4A	-1.6 (2)	C3B—N2B—C2B—C1B	59.48 (15)
O2B—C6B—C7B—C8B	9 (2)	C4B—N1B—C1B—C2B	51.18 (17)
O2B—C6B—C5B—O1B	-29 (2)	C4B—N1B—C5B—O1B	-15.7 (3)
O2B—C6B—C5B—C6BB	87 (17)	C4B—N1B—C5B—C6B	157.9 (14)
O2B—C6B—C5B—N1B	157.2 (12)	C4B—N1B—C5B—C6BB	167.3 (10)
C6B—O2B—C9B—C8B	3.4 (14)	C5B—C6B—C7B—C8B	176 (3)
C6B—C7B—C8B—C9B	-6.6 (14)	C5B—C6BB—C7BB—C8BB	168.9 (14)
C7B—C6B—C5B—O1B	164 (3)	C5B—N1B—C1B—C2B	-140.79 (15)
C7B—C6B—C5B—C6BB	-80 (17)	C5B—N1B—C4B—C3B	138.83 (15)

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2B—H2BA \cdots O1A ⁱ	0.99	2.51	3.1607 (16)	123
N2B—H2BA \cdots O2A ⁱ	0.99	1.72	2.7093 (16)	176
N2B—H2BB \cdots O1A ⁱⁱ	0.99	1.77	2.7424 (16)	166
C5A—H5A \cdots O2A ⁱⁱⁱ	0.95	2.47	3.3170 (18)	148
C9B—H9B \cdots O6A ^{iv}	0.95	2.44	3.183 (6)	134

<i>C8BB—H8BB···O3A^v</i>	0.95	2.50	3.395 (5)	158
<i>C2B—H2BC···O1Bⁱ</i>	0.99	2.59	3.2300 (19)	122

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