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**(E)-tert-Butyl 2-(5-[4-(dimethylamino)-phenyl]diazenyl)-2,6-dioxo-1H-pyrimidin-3-yl)acetate dichloromethane monosolvate**

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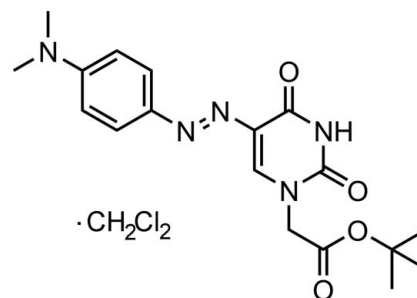
Received 11 March 2014; accepted 3 April 2014

Key indicators: single-crystal X-ray study;  $T = 150$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å; disorder in solvent or counterion;  $R$  factor = 0.051;  $wR$  factor = 0.128; data-to-parameter ratio = 15.9.

In the title compound,  $\text{C}_{18}\text{H}_{23}\text{N}_5\text{O}_4 \cdot \text{CH}_2\text{Cl}_2$ , the dichloromethane solvent molecule is disordered over two sets of sites in a 0.630 (13):0.370 (13) ratio. The dihedral angle between the uracil and phenyl rings is  $30.2$  (1)°. In the crystal, the principal interactions are  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds, which link uracil units across centres of symmetry, forming eight-membered rings with an  $R_2^2(8)$  graph-set motif. The structure also displays  $\text{C}-\text{H} \cdots \text{O}$  and  $\text{C}-\text{H} \cdots \text{Cl}$  hydrogen bonds. Intramolecular  $\text{C}-\text{H} \cdots \text{O}$  short contacts are also observed.

**Related literature**

As part of our program in the synthesis of modified nucleobases that possess intrinsic fluorescence while maintaining an unadulterated base-pairing face, we have prepared an asymmetrical azo compound as a hybrid between a nucleobase and the known fluorescence quencher 4-((4-(dimethylamino)phenyl)azo)benzoic acid (DABCYL), see: Dodd & Hudson (2009); Tyagi & Kramer (1996). For an azo-based fluorescence quencher in peptide nucleic acid, see: Moustafa & Hudson (2011). For an example of photoisomerization of azo groups in peptide nucleic acid, see: Yue *et al.* (2009), and in DNA, see: Asanuma *et al.* (1999). The title compound was prepared following standard procedures, see: Thurber & Townsend (1972), Tsupak *et al.* (2002) and Moustafa (2011).

**Experimental***Crystal data*

$\text{C}_{18}\text{H}_{23}\text{N}_5\text{O}_4 \cdot \text{CH}_2\text{Cl}_2$   
 $M_r = 458.34$   
 Monoclinic,  $P2_1/n$   
 $a = 13.208$  (9) Å  
 $b = 10.783$  (6) Å  
 $c = 17.255$  (11) Å  
 $\beta = 112.33$  (2)°

$V = 2273$  (3) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.32$  mm<sup>-1</sup>  
 $T = 150$  K  
 $0.20 \times 0.18 \times 0.15$  mm

*Data collection*

Nonius KappaCCD diffractometer  
 Absorption correction: multi-scan  
 (SADABS; Bruker, 2008)  
 $T_{\min} = 0.689$ ,  $T_{\max} = 0.746$

34970 measured reflections  
 5944 independent reflections  
 3567 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.054$

*Refinement*

$R[F^2 > 2\sigma(F^2)] = 0.051$   
 $wR(F^2) = 0.128$   
 $S = 1.03$   
 5944 reflections  
 374 parameters  
 12 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\text{max}} = 0.53$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.59$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{N4}-\text{H4A} \cdots \text{O2}^{\text{i}}$	0.84 (3)	2.02 (3)	2.851 (2)	171 (2)
$\text{C2}-\text{H2C} \cdots \text{O1}^{\text{ii}}$	1.03 (3)	2.62 (3)	3.189 (4)	115 (2)
$\text{C4}-\text{H4} \cdots \text{Cl2X}^{\text{iii}}$	0.94 (2)	2.98 (2)	3.784 (4)	143.5 (17)
$\text{C12}-\text{H12} \cdots \text{O1}^{\text{iv}}$	0.99 (2)	2.34 (2)	3.214 (3)	145.8 (18)
$\text{C13}-\text{H13B} \cdots \text{Cl2X}^{\text{v}}$	0.96 (2)	2.99 (2)	3.895 (4)	158.6 (16)
$\text{C16}-\text{H16A} \cdots \text{O3}$	0.98 (3)	2.51 (3)	3.042 (3)	114 (2)
$\text{C17}-\text{H17C} \cdots \text{O3}$	0.96 (3)	2.48 (3)	2.994 (4)	113 (2)
$\text{C1X}-\text{H1X1} \cdots \text{O3}^{\text{vi}}$	0.99	2.52	3.346 (4)	141
$\text{C1X}-\text{H1X2} \cdots \text{O1}^{\text{ii}}$	0.99	2.48	3.256 (4)	135
$\text{C1Y}-\text{H1Y1} \cdots \text{O3}^{\text{vi}}$	0.99	2.50	3.346 (4)	143

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

Data collection: COLLECT (Nonius, 1999); cell refinement: SAINT (Bruker, 2008); data reduction: SAINT; program(s) used to solve structure: SHELXS2014 (Sheldrick, 2008); program(s) used to refine structure: SHELXL2014 (Sheldrick, 2008); molecular graphics: NRCVAX (Gabe *et al.*, 1989); software used to prepare material for publication: cif2tables.py (Boyle, 2008).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: ZP2013).

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## supporting information

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**(*E*)-*tert*-Butyl 2-(5-{[4-(dimethylamino)phenyl]diazenyl}-2,6-dioxo-1*H*-pyrimidin-3-yl)acetate dichloromethane monosolvate**

**Robert H. E. Hudson, Mohamed E. Moustafa and Paul D. Boyle**

### S1. Introduction

As part of our program in the synthesis of modified nucleobases that possess intrinsic fluorescence while maintaining an unadulterated base-pairing face, we have prepared an asymmetrical azo compound as a hybrid between a nucleobase and the known fluorescence quencher 4-((4-(dimethylamino)phenyl)azo)benzoic acid (DABCYL) see: Dodd & Hudson (2009) and Tyagi & Kramer (1996).

### S2. Experimental

The title compound was prepared following standard procedures see: Thurber & Townsend (1972), Tsupak *et al.* (2002) and Moustafa (2011).

#### S2.1. Synthesis and crystallization

The title compound was crystallized by slow diffusion of hexanes into a solution of dichloromethane. Orange plates of suitable quality for diffraction were obtained.

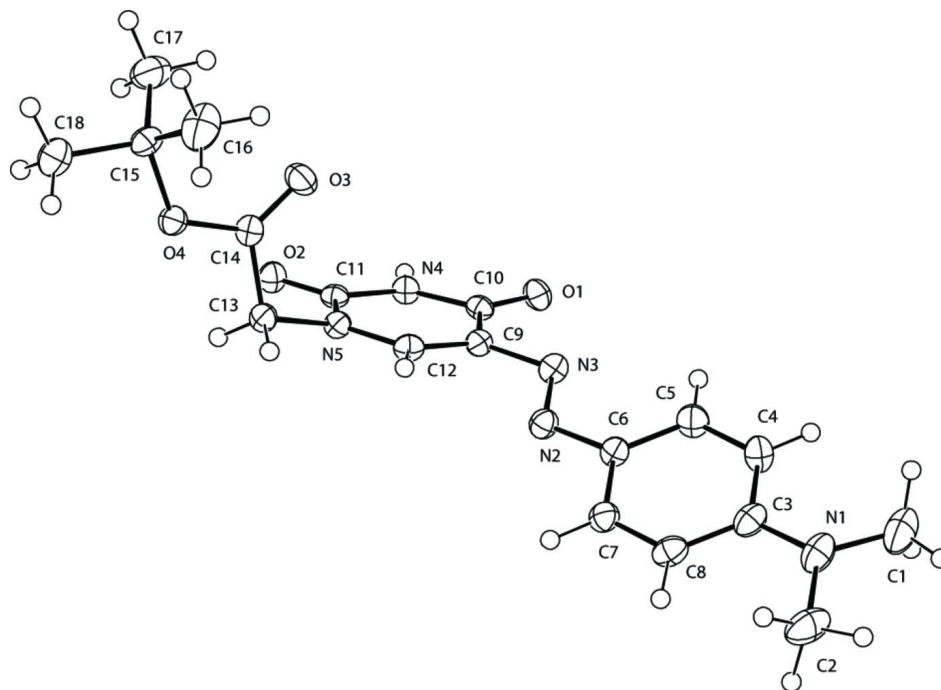
#### S2.2. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 1.

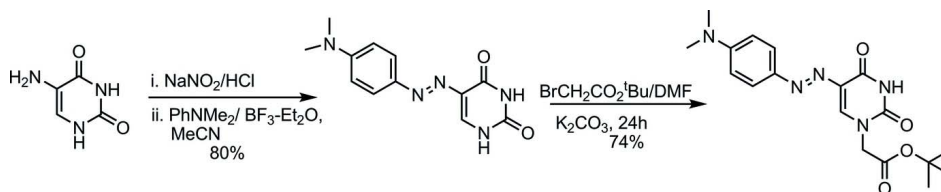
### S3. Results and discussion

The molecule resides at a general position in the lattice as does a disordered methylene chloride of solvation. Selected intermolecular hydrogen bonding interactions are mentioned briefly below. All potential hydrogen bonding interactions are listed in Table 1.

The principal intermolecular interactions are N—H $\cdots$ O hydrogen bonds which join uracil moieties across a centre of symmetry forming an eight membered ring, designated as R<sub>2</sub><sup>2</sup>(8) ring in graph set notation. This hydrogen bond consists of N4—H4A $\cdots$ O2 with an H $\cdots$ A distance of 2.02 (3) Å. The other uracil carbonyl oxygen, O1, participates in two intermolecular C—H $\cdots$ O hydrogen bonds. The shorter of the two interactions arises with the atoms C12—H12 acting as the donor group. The hydrogen bond pattern forms a C(5) chain. The H12 $\cdots$ O1 distance is 2.34 (2) Å. The longer interaction is with the atom H1X1 from the CH<sub>2</sub>Cl<sub>2</sub> of solvation. The H1X2 $\cdots$ O1 distance is 2.48 Å.


**Figure 1**

ORTEP representation of the title compound showing the atom-numbering. The asymmetric unit contains an enclathrated  $\text{CH}_2\text{Cl}_2$  also.


**Figure 2**

Reaction scheme.

**(*E*)-*tert*-Butyl 2-(5-[[4-(dimethylamino)phenyl]diazenyl]-2,6-dioxo-1*H*-pyrimidin-3-yl)acetate dichloromethane monosolvate**

*Crystal data*

$\text{C}_{18}\text{H}_{23}\text{N}_5\text{O}_4 \cdot \text{CH}_2\text{Cl}_2$

$M_r = 458.34$

Monoclinic,  $P2_1/n$

$a = 13.208$  (9) Å

$b = 10.783$  (6) Å

$c = 17.255$  (11) Å

$\beta = 112.33$  (2)°

$V = 2273$  (3) Å<sup>3</sup>

$Z = 4$

$F(000) = 960$

$D_x = 1.339$  Mg m<sup>-3</sup>

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

Cell parameters from 9939 reflections

$\theta = 2.3$ – $28.1$ °

$\mu = 0.32$  mm<sup>-1</sup>

$T = 150$  K

Prism, colourless

$0.20 \times 0.18 \times 0.15$  mm

*Data collection*

Nonius KappaCCD  
diffractometer

Radiation source: sealed tube  
phi and  $\omega$  scans

Absorption correction: multi-scan  
(SADABS; Bruker, 2008)

$T_{\min} = 0.689$ ,  $T_{\max} = 0.746$

34970 measured reflections

5944 independent reflections

3567 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.054$

$\theta_{\max} = 29.6^\circ$ ,  $\theta_{\min} = 2.3^\circ$

$h = -18 \rightarrow 17$

$k = -14 \rightarrow 14$

$l = -22 \rightarrow 23$

*Refinement*

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.051$

$wR(F^2) = 0.128$

$S = 1.03$

5944 reflections

374 parameters

12 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

$w = 1/[\sigma^2(F_o^2) + (0.0446P)^2 + 1.4761P]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.53 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.59 \text{ e } \text{\AA}^{-3}$

*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.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.9052 (2)	0.7150 (3)	1.05913 (18)	0.0474 (7)	
H1A	0.873 (2)	0.777 (2)	1.0869 (16)	0.045 (7)*	
H1B	0.929 (3)	0.766 (3)	1.020 (2)	0.084 (11)*	
H1C	0.969 (3)	0.675 (3)	1.101 (2)	0.083 (11)*	
C2	0.8704 (2)	0.4912 (3)	1.02020 (18)	0.0443 (7)	
H2A	0.939 (3)	0.490 (3)	1.069 (2)	0.071 (9)*	
H2B	0.891 (2)	0.465 (2)	0.9674 (18)	0.056 (8)*	
H2C	0.817 (3)	0.427 (3)	1.0273 (19)	0.071 (10)*	
N1	0.82913 (15)	0.61752 (19)	1.01530 (12)	0.0372 (5)	
C3	0.72814 (16)	0.6469 (2)	0.95471 (13)	0.0279 (5)	
C4	0.68748 (18)	0.7695 (2)	0.94533 (14)	0.0305 (5)	
H4	0.7300 (18)	0.833 (2)	0.9799 (14)	0.028 (6)*	
C5	0.58531 (18)	0.7985 (2)	0.88637 (14)	0.0293 (5)	
H5	0.5580 (19)	0.877 (2)	0.8811 (15)	0.039 (7)*	
C6	0.51979 (16)	0.70655 (18)	0.83333 (12)	0.0232 (4)	
C7	0.56023 (17)	0.58499 (19)	0.84203 (14)	0.0261 (5)	
H7	0.5167 (17)	0.524 (2)	0.8034 (14)	0.029 (6)*	
C8	0.66189 (17)	0.5549 (2)	0.90144 (14)	0.0281 (5)	
H8	0.6874 (18)	0.469 (2)	0.9054 (14)	0.034 (6)*	

N2	0.41469 (13)	0.72338 (16)	0.77085 (10)	0.0250 (4)	
N3	0.37871 (13)	0.83448 (15)	0.76006 (10)	0.0245 (4)	
C9	0.26949 (15)	0.83883 (18)	0.69941 (12)	0.0213 (4)	
C10	0.23176 (16)	0.95536 (18)	0.65479 (12)	0.0210 (4)	
O1	0.28351 (11)	1.05269 (13)	0.66708 (9)	0.0269 (3)	
N4	0.12641 (13)	0.94789 (16)	0.59289 (10)	0.0216 (4)	
H4A	0.1030 (19)	1.012 (2)	0.5641 (16)	0.037 (7)*	
C11	0.05549 (15)	0.84931 (18)	0.57493 (12)	0.0199 (4)	
O2	-0.03756 (11)	0.85176 (13)	0.52046 (8)	0.0254 (3)	
N5	0.09479 (13)	0.74509 (14)	0.62430 (10)	0.0207 (4)	
C12	0.19932 (16)	0.74158 (19)	0.68434 (12)	0.0226 (4)	
H12	0.2182 (18)	0.663 (2)	0.7162 (14)	0.033 (6)*	
C13	0.01573 (17)	0.64735 (19)	0.61964 (14)	0.0229 (4)	
H13A	0.0566 (19)	0.576 (2)	0.6456 (15)	0.036 (6)*	
H13B	-0.0272 (17)	0.6270 (19)	0.5626 (14)	0.023 (5)*	
C14	-0.05649 (16)	0.68765 (18)	0.66563 (12)	0.0220 (4)	
O3	-0.03756 (12)	0.77674 (14)	0.71118 (9)	0.0318 (4)	
O4	-0.14075 (11)	0.60916 (13)	0.64849 (8)	0.0245 (3)	
C15	-0.21469 (16)	0.6153 (2)	0.69607 (13)	0.0271 (5)	
C16	-0.1464 (2)	0.5914 (3)	0.78781 (16)	0.0456 (7)	
H16A	-0.095 (2)	0.659 (3)	0.8133 (19)	0.066 (9)*	
H16B	-0.104 (3)	0.516 (3)	0.7938 (19)	0.070 (10)*	
H16C	-0.193 (3)	0.584 (3)	0.819 (2)	0.072 (10)*	
C17	-0.2744 (2)	0.7388 (3)	0.6801 (2)	0.0440 (6)	
H17A	-0.328 (2)	0.738 (3)	0.7089 (17)	0.057 (8)*	
H17B	-0.311 (2)	0.755 (3)	0.6188 (19)	0.056 (8)*	
H17C	-0.227 (3)	0.807 (3)	0.7051 (19)	0.065 (9)*	
C18	-0.2945 (2)	0.5097 (3)	0.65820 (18)	0.0382 (6)	
H18A	-0.256 (2)	0.432 (2)	0.6654 (15)	0.040 (7)*	
H18B	-0.337 (2)	0.524 (3)	0.595 (2)	0.067 (9)*	
H18C	-0.347 (2)	0.506 (3)	0.6847 (17)	0.054 (8)*	
C1X	0.8190 (3)	0.1657 (3)	1.1140 (2)	0.0728 (10)	0.630 (13)
H1X1	0.8564	0.1666	1.1758	0.087*	0.630 (13)
H1X2	0.7807	0.2460	1.0970	0.087*	0.630 (13)
C11X	0.91767 (6)	0.15439 (9)	1.07048 (5)	0.0687 (3)	0.630 (13)
C12X	0.72201 (17)	0.0480 (2)	1.08651 (16)	0.0511 (7)	0.630 (13)
C1Y	0.8190 (3)	0.1657 (3)	1.1140 (2)	0.0728 (10)	0.370 (13)
H1Y1	0.8550	0.1888	1.1739	0.087*	0.370 (13)
H1Y2	0.7669	0.2326	1.0853	0.087*	0.370 (13)
C11Y	0.91767 (6)	0.15439 (9)	1.07048 (5)	0.0687 (3)	0.370 (13)
C12Y	0.7489 (11)	0.0313 (6)	1.1063 (8)	0.114 (2)	0.370 (13)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0272 (13)	0.072 (2)	0.0370 (15)	-0.0029 (13)	0.0051 (12)	0.0058 (15)
C2	0.0346 (14)	0.0589 (18)	0.0386 (15)	0.0208 (13)	0.0128 (12)	0.0140 (13)
N1	0.0243 (9)	0.0461 (12)	0.0349 (11)	0.0039 (9)	0.0042 (8)	0.0078 (9)

C3	0.0231 (10)	0.0355 (12)	0.0267 (11)	0.0020 (9)	0.0112 (9)	0.0078 (9)
C4	0.0291 (11)	0.0286 (12)	0.0293 (11)	-0.0049 (10)	0.0061 (10)	-0.0004 (10)
C5	0.0305 (11)	0.0224 (11)	0.0322 (12)	0.0019 (9)	0.0087 (10)	0.0025 (9)
C6	0.0223 (10)	0.0211 (10)	0.0257 (10)	0.0003 (8)	0.0083 (8)	0.0028 (8)
C7	0.0241 (10)	0.0226 (11)	0.0331 (12)	-0.0018 (9)	0.0126 (9)	0.0001 (9)
C8	0.0274 (11)	0.0238 (11)	0.0362 (12)	0.0046 (9)	0.0154 (10)	0.0069 (9)
N2	0.0231 (9)	0.0244 (9)	0.0264 (9)	0.0010 (7)	0.0083 (7)	0.0023 (7)
N3	0.0241 (9)	0.0250 (9)	0.0238 (9)	0.0010 (7)	0.0085 (7)	0.0040 (7)
C9	0.0211 (9)	0.0227 (10)	0.0199 (10)	0.0007 (8)	0.0076 (8)	0.0011 (8)
C10	0.0250 (10)	0.0207 (10)	0.0196 (9)	0.0000 (8)	0.0109 (8)	-0.0008 (8)
O1	0.0308 (8)	0.0201 (7)	0.0284 (8)	-0.0036 (6)	0.0097 (6)	-0.0010 (6)
N4	0.0235 (8)	0.0181 (9)	0.0221 (9)	0.0009 (7)	0.0073 (7)	0.0038 (7)
C11	0.0233 (10)	0.0190 (10)	0.0206 (10)	0.0012 (8)	0.0117 (8)	0.0008 (8)
O2	0.0236 (7)	0.0233 (7)	0.0257 (7)	0.0011 (6)	0.0052 (6)	0.0032 (6)
N5	0.0211 (8)	0.0177 (8)	0.0237 (8)	-0.0002 (7)	0.0088 (7)	0.0026 (7)
C12	0.0251 (10)	0.0207 (11)	0.0226 (10)	0.0040 (8)	0.0096 (9)	0.0038 (8)
C13	0.0229 (10)	0.0170 (10)	0.0287 (11)	-0.0010 (8)	0.0097 (9)	0.0017 (9)
C14	0.0228 (10)	0.0193 (10)	0.0224 (10)	-0.0006 (8)	0.0068 (8)	0.0053 (8)
O3	0.0373 (9)	0.0270 (8)	0.0348 (8)	-0.0071 (7)	0.0179 (7)	-0.0087 (7)
O4	0.0248 (7)	0.0231 (7)	0.0287 (8)	-0.0038 (6)	0.0138 (6)	-0.0004 (6)
C15	0.0244 (10)	0.0315 (12)	0.0297 (11)	-0.0011 (9)	0.0151 (9)	0.0013 (9)
C16	0.0399 (15)	0.069 (2)	0.0298 (13)	-0.0125 (15)	0.0155 (12)	0.0091 (13)
C17	0.0367 (14)	0.0416 (16)	0.0602 (19)	0.0039 (12)	0.0255 (14)	-0.0042 (14)
C18	0.0325 (13)	0.0400 (15)	0.0457 (15)	-0.0086 (11)	0.0189 (12)	-0.0017 (12)
C1X	0.085 (2)	0.0521 (19)	0.105 (3)	-0.0259 (17)	0.063 (2)	-0.0354 (18)
C11X	0.0585 (5)	0.0981 (7)	0.0553 (5)	-0.0121 (4)	0.0282 (4)	-0.0193 (4)
C12X	0.0499 (14)	0.0513 (13)	0.0461 (11)	-0.0163 (7)	0.0114 (11)	-0.0014 (7)
C1Y	0.085 (2)	0.0521 (19)	0.105 (3)	-0.0259 (17)	0.063 (2)	-0.0354 (18)
C11Y	0.0585 (5)	0.0981 (7)	0.0553 (5)	-0.0121 (4)	0.0282 (4)	-0.0193 (4)
C12Y	0.232 (6)	0.0284 (17)	0.153 (5)	-0.024 (3)	0.155 (5)	-0.013 (3)

*Geometric parameters (Å, °)*

C1—N1	1.452 (3)	C11—O2	1.232 (2)
C1—H1A	1.00 (3)	C11—N5	1.386 (2)
C1—H1B	1.01 (4)	N5—C12	1.376 (3)
C1—H1C	0.98 (4)	N5—C13	1.464 (3)
C2—N1	1.458 (3)	C12—H12	0.99 (2)
C2—H2A	0.98 (3)	C13—C14	1.518 (3)
C2—H2B	1.08 (3)	C13—H13A	0.95 (2)
C2—H2C	1.03 (3)	C13—H13B	0.96 (2)
N1—C3	1.383 (3)	C14—O3	1.206 (2)
C3—C8	1.408 (3)	C14—O4	1.339 (2)
C3—C4	1.414 (3)	O4—C15	1.497 (3)
C4—C5	1.382 (3)	C15—C16	1.517 (3)
C4—H4	0.94 (2)	C15—C17	1.518 (3)
C5—C6	1.403 (3)	C15—C18	1.521 (3)
C5—H5	0.91 (3)	C16—H16A	0.98 (3)

C6—C7	1.402 (3)	C16—H16B	0.97 (3)
C6—N2	1.409 (3)	C16—H16C	0.95 (3)
C7—C8	1.383 (3)	C17—H17A	1.00 (3)
C7—H7	0.95 (2)	C17—H17B	1.00 (3)
C8—H8	0.98 (2)	C17—H17C	0.96 (3)
N2—N3	1.276 (2)	C18—H18A	0.97 (3)
N3—C9	1.425 (3)	C18—H18B	1.03 (3)
C9—C12	1.358 (3)	C18—H18C	0.97 (3)
C9—C10	1.459 (3)	C1X—C12X	1.737 (3)
C10—O1	1.226 (2)	C1X—C11X	1.738 (3)
C10—N4	1.398 (3)	C1X—H1X1	0.9900
N4—C11	1.372 (3)	C1X—H1X2	0.9900
N4—H4A	0.84 (3)		
N1—C1—H1A	113.5 (15)	C12—N5—C11	121.27 (16)
N1—C1—H1B	112.0 (19)	C12—N5—C13	120.81 (16)
H1A—C1—H1B	104 (2)	C11—N5—C13	117.19 (16)
N1—C1—H1C	108 (2)	C9—C12—N5	122.66 (18)
H1A—C1—H1C	110 (2)	C9—C12—H12	123.2 (13)
H1B—C1—H1C	109 (3)	N5—C12—H12	114.2 (13)
N1—C2—H2A	105.4 (19)	N5—C13—C14	110.01 (17)
N1—C2—H2B	113.6 (14)	N5—C13—H13A	107.0 (14)
H2A—C2—H2B	106 (2)	C14—C13—H13A	110.0 (14)
N1—C2—H2C	112.1 (17)	N5—C13—H13B	110.7 (13)
H2A—C2—H2C	110 (2)	C14—C13—H13B	111.1 (13)
H2B—C2—H2C	109 (2)	H13A—C13—H13B	107.9 (19)
C3—N1—C1	120.4 (2)	O3—C14—O4	126.64 (19)
C3—N1—C2	119.3 (2)	O3—C14—C13	123.57 (18)
C1—N1—C2	118.5 (2)	O4—C14—C13	109.78 (17)
N1—C3—C8	121.0 (2)	C14—O4—C15	120.76 (16)
N1—C3—C4	121.1 (2)	O4—C15—C16	108.16 (19)
C8—C3—C4	117.98 (19)	O4—C15—C17	109.97 (18)
C5—C4—C3	121.2 (2)	C16—C15—C17	113.6 (2)
C5—C4—H4	118.8 (13)	O4—C15—C18	102.75 (18)
C3—C4—H4	120.0 (13)	C16—C15—C18	111.2 (2)
C4—C5—C6	120.6 (2)	C17—C15—C18	110.6 (2)
C4—C5—H5	121.4 (15)	C15—C16—H16A	112.1 (18)
C6—C5—H5	118.0 (15)	C15—C16—H16B	110.1 (19)
C7—C6—C5	118.39 (19)	H16A—C16—H16B	108 (2)
C7—C6—N2	115.27 (18)	C15—C16—H16C	109.9 (19)
C5—C6—N2	126.34 (19)	H16A—C16—H16C	107 (3)
C8—C7—C6	121.4 (2)	H16B—C16—H16C	110 (3)
C8—C7—H7	121.1 (13)	C15—C17—H17A	108.7 (16)
C6—C7—H7	117.4 (13)	C15—C17—H17B	110.5 (17)
C7—C8—C3	120.4 (2)	H17A—C17—H17B	112 (2)
C7—C8—H8	119.0 (13)	C15—C17—H17C	112.6 (18)
C3—C8—H8	120.6 (13)	H17A—C17—H17C	104 (2)
N3—N2—C6	115.89 (16)	H17B—C17—H17C	108 (2)



N2—N3—C9	110.80 (16)	C15—C18—H18A	110.5 (15)
C12—C9—N3	122.83 (18)	C15—C18—H18B	110.8 (16)
C12—C9—C10	119.52 (18)	H18A—C18—H18B	109 (2)
N3—C9—C10	117.61 (17)	C15—C18—H18C	109.2 (17)
O1—C10—N4	120.61 (18)	H18A—C18—H18C	110 (2)
O1—C10—C9	126.13 (18)	H18B—C18—H18C	107 (2)
N4—C10—C9	113.26 (17)	Cl2X—C1X—Cl1X	115.3 (2)
C11—N4—C10	127.82 (17)	Cl2X—C1X—H1X1	108.5
C11—N4—H4A	116.1 (17)	Cl1X—C1X—H1X1	108.5
C10—N4—H4A	116.0 (17)	Cl2X—C1X—H1X2	108.5
O2—C11—N4	123.60 (18)	Cl1X—C1X—H1X2	108.5
O2—C11—N5	121.23 (17)	H1X1—C1X—H1X2	107.5
N4—C11—N5	115.15 (17)		
C1—N1—C3—C8	168.5 (2)	N3—C9—C10—N4	175.99 (16)
C2—N1—C3—C8	4.1 (3)	O1—C10—N4—C11	-175.49 (18)
C1—N1—C3—C4	-12.6 (3)	C9—C10—N4—C11	4.9 (3)
C2—N1—C3—C4	-177.0 (2)	C10—N4—C11—O2	177.93 (19)
N1—C3—C4—C5	-178.2 (2)	C10—N4—C11—N5	-0.6 (3)
C8—C3—C4—C5	0.7 (3)	O2—C11—N5—C12	179.03 (18)
C3—C4—C5—C6	-0.9 (3)	N4—C11—N5—C12	-2.4 (3)
C4—C5—C6—C7	0.4 (3)	O2—C11—N5—C13	-10.7 (3)
C4—C5—C6—N2	-180.0 (2)	N4—C11—N5—C13	167.88 (17)
C5—C6—C7—C8	0.3 (3)	N3—C9—C12—N5	-178.38 (18)
N2—C6—C7—C8	-179.39 (19)	C10—C9—C12—N5	4.2 (3)
C6—C7—C8—C3	-0.5 (3)	C11—N5—C12—C9	0.5 (3)
N1—C3—C8—C7	178.9 (2)	C13—N5—C12—C9	-169.43 (19)
C4—C3—C8—C7	0.0 (3)	C12—N5—C13—C14	96.3 (2)
C7—C6—N2—N3	-177.35 (18)	C11—N5—C13—C14	-74.0 (2)
C5—C6—N2—N3	3.0 (3)	N5—C13—C14—O3	-12.2 (3)
C6—N2—N3—C9	-176.25 (16)	N5—C13—C14—O4	168.70 (15)
N2—N3—C9—C12	26.9 (3)	O3—C14—O4—C15	-8.9 (3)
N2—N3—C9—C10	-155.63 (17)	C13—C14—O4—C15	170.22 (16)
C12—C9—C10—O1	173.92 (19)	C14—O4—C15—C16	-60.9 (2)
N3—C9—C10—O1	-3.6 (3)	C14—O4—C15—C17	63.7 (2)
C12—C9—C10—N4	-6.5 (3)	C14—O4—C15—C18	-178.54 (18)

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>
N4—H4A...O2 <sup>i</sup>	0.84 (3)	2.02 (3)	2.851 (2)	171 (2)
C2—H2C...O1 <sup>ii</sup>	1.03 (3)	2.62 (3)	3.189 (4)	115 (2)
C4—H4...Cl2X <sup>iii</sup>	0.94 (2)	2.98 (2)	3.784 (4)	143.5 (17)
C12—H12...O1 <sup>iv</sup>	0.99 (2)	2.34 (2)	3.214 (3)	145.8 (18)
C13—H13B...Cl2X <sup>v</sup>	0.96 (2)	2.99 (2)	3.895 (4)	158.6 (16)
C16—H16A...O3	0.98 (3)	2.51 (3)	3.042 (3)	114 (2)
C17—H17C...O3	0.96 (3)	2.48 (3)	2.994 (4)	113 (2)
C1X—H1X1...O3 <sup>vi</sup>	0.99	2.52	3.346 (4)	141

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C1X—H1X2...O1 <sup>ii</sup>	0.99	2.48	3.256 (4)	135
C1Y—H1Y1...O3 <sup>vi</sup>	0.99	2.50	3.346 (4)	143

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