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Methyl 2-[(carbamoylamino)imino]-2-(3-[1-[(carbamoylamino)imino]-2-methoxy-2-oxoethyl]phenyl)acetate ethanol monosolvate monohydrate

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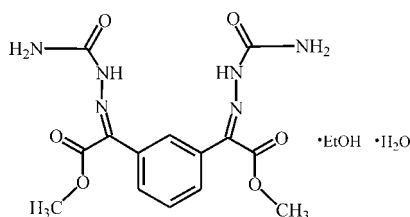
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Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.043; wR factor = 0.125; data-to-parameter ratio = 14.5.

In the title compound, $\text{C}_{14}\text{H}_{16}\text{N}_6\text{O}_6 \cdot \text{C}_2\text{H}_6\text{O} \cdot \text{H}_2\text{O}$, both substituents of the benzene ring are approximately planar with maximum deviations from the mean plane of 0.0561 (12) (an imine N atom) and 0.1419 (11) Å (a methoxy O atom). The substituents are tilted out of the plane of the benzene ring by 64.48 (4) and 70.08 (5)°, respectively. In the crystal, molecules form centrosymmetric dimers associated *via* pairs of N—H...O hydrogen bonds. The dimers are linked *via* the water and ethanol molecules, forming two-dimensional hydrogen-bond networks lying parallel to (100).

Related literature

For details of the synthesis of 1,4-benzodiketodicarboxylic acid and its derivatives, see: Ismatov *et al.* (1991). For the synthesis and biological activity of 1,3-benzodiketodicarboxylic acid and its derivatives, see: Ismatov *et al.* (1998, 2001).



Experimental

Crystal data

 $\text{C}_{14}\text{H}_{16}\text{N}_6\text{O}_6 \cdot \text{C}_2\text{H}_6\text{O} \cdot \text{H}_2\text{O}$ $M_r = 428.41$

Monoclinic, $P2_1/c$
 $a = 7.5810$ (2) Å
 $b = 12.1216$ (3) Å
 $c = 23.0379$ (7) Å
 $\beta = 97.254$ (3)°
 $V = 2100.11$ (10) Å³

$Z = 4$
 Cu $K\alpha$ radiation
 $\mu = 0.94$ mm⁻¹
 $T = 294$ K
 $0.41 \times 0.32 \times 0.26$ mm

Data collection

Oxford Diffraction Xcalibur diffractometer
 Absorption correction: multi-scan (ABSPACK in *CrysAlis PRO*; Oxford Diffraction, 2009)
 $T_{\min} = 0.700$, $T_{\max} = 1.000$

15863 measured reflections
 4317 independent reflections
 2927 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.125$
 $S = 0.98$
 4317 reflections
 298 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.27$ e Å⁻³
 $\Delta\rho_{\min} = -0.18$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O1W—H1WA...O6	0.85	2.05	2.879 (3)	166
O1W—H1WB...O5 ⁱ	0.85	2.16	2.941 (3)	152
N3—H3A...O7	0.86 (4)	2.13 (4)	2.980 (2)	169 (3)
N3—H3B...O5 ⁱⁱ	0.85 (3)	2.61 (2)	3.082 (2)	116.7 (18)
N5—H5A...O3 ⁱⁱⁱ	0.86	2.19	3.010 (2)	160
N6—H6A...O1W ^{iv}	0.84 (3)	2.34 (3)	3.137 (3)	158 (3)
N6—H6B...O7 ^v	0.90 (3)	1.99 (3)	2.871 (3)	166 (2)
O7—H7...O2	0.88 (3)	1.96 (3)	2.801 (2)	158 (3)
O7—H7...N1	0.88 (3)	2.52 (3)	3.0677 (19)	121 (2)
C4—H4...O6 ^{vi}	0.93	2.57	3.493 (2)	173
C13—H13C...N6 ^{vii}	0.96	2.62	3.405 (3)	139

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2009); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FY2032).

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supporting information

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Methyl 2-[(carbamoylamino)imino]-2-(3-{1-[(carbamoylamino)imino]-2-methoxy-2-oxoethyl}phenyl)acetate ethanol monosolvate monohydrate

Dilmurot Ismatov, Umarkhon Azizov, Jamshid Ashurov and Samat Talipov

S1. Comment

It has been reported previously that arylketodicarbonic acid derivatives possess antiinflammatory activity (Ismatov *et al.*, 1991, Ismatov *et al.*, 1998, Ismatov *et al.*, 2001). We report herein the X-ray crystallographic study of a disemicarbazodimethylether-1,3-benzenediketodicarbonic acid monohydrate ethanol solvate.

The asymmetric unit of the title compound, $C_{14}H_{16}N_6O_6 \cdot C_2H_6O \cdot H_2O$, contains a disemicarbazodimethylether-1,3-benzenediketodicarbonic acid (DBA), an ethanol and a water molecule as shown in Fig. 1. In the DBA molecule the planes of the substituent fragments (C1/C7/C8/N1/N2/O1/O2 and C3/C11/C12/N4/N5/O4/O5) are tilted out of the mean plane of the benzene ring by $64.48(4)^\circ$ and $70.08(5)^\circ$, respectively.

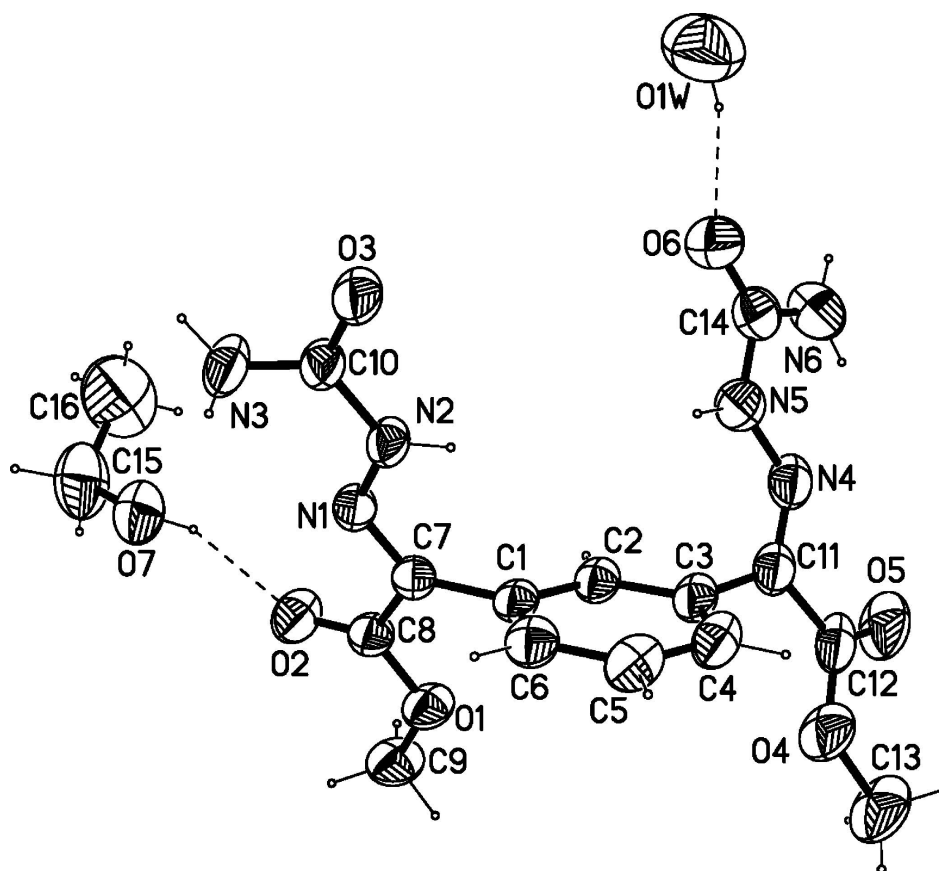
In the crystal structure (Fig. 2), DBA molecules form centrosymmetric pair associates *via* $N2-H \cdots O3$ and $N5-H \cdots O3$ H-bonds and the associates are linked *via* H-bonds by water ($O1W-H1 \cdots O5$, $O1W-H2 \cdots O6$, $N6-H \cdots O1W$) and ethanol ($N6-H \cdots O7$, $N3-H3 \cdots O7$, $O7-H7 \cdots O2$) molecules forming two-dimensional networks parallel to the (100) plane (Table 1).

S2. Experimental

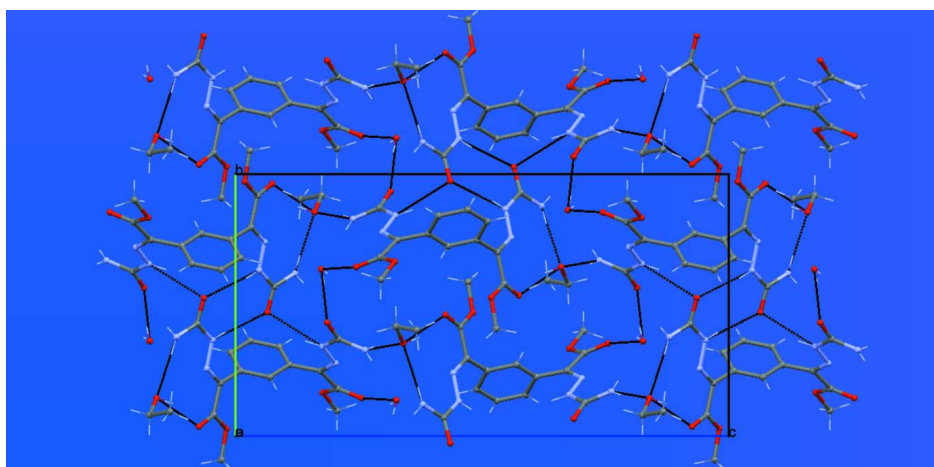
In a round bottom flask, dimethylether-1,3-benzodiketodicarboxylic acid (0.02 mol, 0.51 g) was dissolved in 5 ml methanol. Drops of semicarbazide chloride (0.04 mol, 0.44 g) dissolved in ethanol (5 ml) were then added. The reaction mixture was refluxed for 30 min. The precipitate (0.44 g) formed after 24 h was filtered. The precipitate was then dissolved in ethanol at room temperature. After few days, colorless crystals (m.p. $120-121^\circ\text{C}$) were formed by slow evaporation.

S3. Refinement

C-bound H atoms were placed in calculated positions with C—H 0.93 Å for aromatic, 0.97 Å for CH_2 and 0.96 Å for CH_3 hydrogens and were refined as riding on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ or $1.5 U_{\text{eq}}(\text{C})$ for methyl H atoms. NH H atoms were placed in calculated positions with N—H 0.86 Å and refined in riding mode with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{N})$. The H-atoms bonded to the O atom of the water molecule were found from difference Fourier map and their coordinates were refined independently with $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O})$. The H-atom bonded to the O atom of the ethanol molecule and the NH_2 H atoms were refined freely.

**Figure 1**

The molecular structure of title compound, with the atom numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

Packing diagram for title compound.

Methyl 2-[(carbamoylamino)imino]-2-(3-{1-[(carbamoylamino)imino]-2-methoxy-2-oxoethyl}phenyl)acetate ethanol monosolvate monohydrate

Crystal data

C₁₄H₁₆N₆O₆·C₂H₆O·H₂O

M_r = 428.41

Monoclinic, *P*2₁/*c*

Hall symbol: -P 2ybc

a = 7.5810 (2) Å

b = 12.1216 (3) Å

c = 23.0379 (7) Å

β = 97.254 (3)°

V = 2100.11 (10) Å³

Z = 4

F(000) = 904

D_x = 1.355 Mg m⁻³

Cu *K*α radiation, λ = 1.54184 Å

θ = 3.6–75.7°

μ = 0.94 mm⁻¹

T = 294 K

Block, colourless

0.41 × 0.32 × 0.26 mm

Data collection

Oxford Diffraction Xcalibur diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 10.2576 pixels mm⁻¹

ω scans

Absorption correction: multi-scan

(ABSPACK in *CrysAlis PRO*; Oxford Diffraction, 2009)

T_{min} = 0.700, *T_{max}* = 1.000

15863 measured reflections

4317 independent reflections

2927 reflections with *I* > 2σ(*I*)

R_{int} = 0.035

θ_{max} = 75.9°, θ_{min} = 3.9°

h = -8→9

k = -14→15

l = -27→28

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.043

wR(*F*²) = 0.125

S = 0.98

4317 reflections

298 parameters

0 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/[σ²(*F_o*²) + (0.0775*P*)²]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.002

Δρ_{max} = 0.27 e Å⁻³

Δρ_{min} = -0.18 e Å⁻³

Extinction correction: *SHELXL97* (Sheldrick, 2008), *F_c** = *kF_c*[1 + 0.001*xF_c*²λ³/sin(2θ)]^{-1/4}

Extinction coefficient: 0.0014 (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 > σ(*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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U_{iso}</i> */ <i>U_{eq}</i>
C1	0.2929 (2)	0.25039 (12)	0.51014 (7)	0.0377 (4)

C2	0.3041 (2)	0.27386 (13)	0.56962 (7)	0.0400 (4)
H2	0.2219	0.3212	0.5831	0.048*
C3	0.4375 (2)	0.22701 (13)	0.60902 (7)	0.0427 (4)
C4	0.5626 (2)	0.15889 (14)	0.58852 (8)	0.0501 (4)
H4	0.6537	0.1288	0.6145	0.060*
C5	0.5520 (2)	0.13560 (14)	0.52931 (9)	0.0526 (4)
H5	0.6362	0.0897	0.5157	0.063*
C6	0.4171 (2)	0.18025 (13)	0.49020 (8)	0.0457 (4)
H6	0.4096	0.1633	0.4506	0.055*
C7	0.1457 (2)	0.29742 (13)	0.46895 (7)	0.0385 (4)
C8	0.1357 (2)	0.41831 (13)	0.45552 (7)	0.0412 (4)
C9	0.2828 (3)	0.58940 (15)	0.47140 (10)	0.0582 (5)
H9A	0.2586	0.6042	0.4302	0.087*
H9B	0.3987	0.6170	0.4860	0.087*
H9C	0.1951	0.6251	0.4915	0.087*
C10	-0.1273 (2)	0.07098 (14)	0.42197 (7)	0.0455 (4)
C11	0.4352 (2)	0.25050 (14)	0.67237 (7)	0.0475 (4)
C12	0.5777 (3)	0.31779 (15)	0.70685 (8)	0.0553 (5)
C13	0.8581 (3)	0.4016 (2)	0.70538 (13)	0.0876 (8)
H13A	0.9592	0.3958	0.6843	0.131*
H13B	0.8902	0.3768	0.7449	0.131*
H13C	0.8199	0.4771	0.7056	0.131*
C14	0.0394 (3)	0.12867 (16)	0.70835 (9)	0.0583 (5)
N1	0.01612 (18)	0.24187 (11)	0.44138 (6)	0.0408 (3)
N2	0.00539 (19)	0.13309 (10)	0.45248 (6)	0.0446 (3)
H2A	0.0812	0.1026	0.4785	0.053*
N3	-0.2322 (3)	0.12126 (15)	0.37973 (8)	0.0628 (5)
N4	0.3108 (2)	0.22112 (12)	0.70208 (7)	0.0528 (4)
N5	0.1754 (2)	0.15785 (13)	0.67639 (7)	0.0560 (4)
H5A	0.1739	0.1361	0.6408	0.067*
N6	0.0482 (3)	0.17095 (18)	0.76207 (9)	0.0713 (5)
O1	0.27699 (16)	0.47111 (9)	0.48126 (6)	0.0489 (3)
O2	0.01418 (18)	0.46321 (10)	0.42578 (6)	0.0585 (4)
O3	-0.14134 (19)	-0.02689 (10)	0.43545 (6)	0.0583 (4)
O4	0.7144 (2)	0.33357 (12)	0.67710 (7)	0.0670 (4)
O5	0.5688 (2)	0.35432 (13)	0.75522 (6)	0.0767 (5)
O6	-0.0770 (2)	0.06677 (13)	0.68590 (7)	0.0725 (4)
C15	-0.3706 (4)	0.4190 (2)	0.32573 (11)	0.0836 (8)
H15A	-0.4259	0.3992	0.2869	0.100*
H15B	-0.3366	0.4961	0.3249	0.100*
C16	-0.4991 (4)	0.4054 (3)	0.36663 (16)	0.1227 (12)
H16A	-0.6073	0.4428	0.3520	0.184*
H16B	-0.4523	0.4361	0.4038	0.184*
H16C	-0.5229	0.3283	0.3712	0.184*
O7	-0.2166 (2)	0.35452 (12)	0.33981 (7)	0.0649 (4)
O1W	-0.2726 (4)	-0.13725 (18)	0.67387 (11)	0.1195 (8)
H1WA	-0.2003	-0.0837	0.6802	0.179*
H1WB	-0.3699	-0.1212	0.6866	0.179*

H3A	-0.212 (4)	0.188 (3)	0.3697 (13)	0.106 (10)*
H3B	-0.317 (3)	0.084 (2)	0.3617 (10)	0.073 (7)*
H6A	0.132 (4)	0.212 (2)	0.7761 (12)	0.095 (10)*
H6B	-0.046 (4)	0.158 (2)	0.7812 (11)	0.081 (8)*
H7	-0.156 (4)	0.376 (2)	0.3730 (13)	0.111 (10)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0379 (8)	0.0285 (7)	0.0444 (8)	-0.0087 (6)	-0.0032 (6)	0.0021 (6)
C2	0.0412 (9)	0.0310 (8)	0.0458 (9)	-0.0047 (6)	-0.0024 (7)	-0.0002 (6)
C3	0.0459 (9)	0.0318 (8)	0.0466 (9)	-0.0088 (7)	-0.0094 (7)	0.0010 (6)
C4	0.0473 (10)	0.0358 (8)	0.0623 (11)	-0.0004 (7)	-0.0120 (8)	0.0012 (8)
C5	0.0463 (10)	0.0410 (9)	0.0688 (12)	0.0020 (7)	0.0005 (9)	-0.0042 (8)
C6	0.0502 (10)	0.0364 (8)	0.0495 (9)	-0.0061 (7)	0.0019 (8)	-0.0041 (7)
C7	0.0408 (8)	0.0352 (8)	0.0385 (8)	-0.0059 (6)	0.0007 (7)	0.0027 (6)
C8	0.0417 (9)	0.0356 (8)	0.0455 (9)	-0.0049 (7)	0.0022 (7)	0.0032 (7)
C9	0.0573 (11)	0.0344 (9)	0.0817 (14)	-0.0099 (8)	0.0041 (10)	0.0042 (9)
C10	0.0529 (10)	0.0399 (9)	0.0402 (8)	-0.0106 (7)	-0.0078 (7)	0.0024 (7)
C11	0.0530 (10)	0.0392 (9)	0.0459 (9)	-0.0043 (8)	-0.0110 (8)	0.0042 (7)
C12	0.0653 (13)	0.0402 (9)	0.0524 (11)	-0.0038 (8)	-0.0231 (9)	0.0046 (8)
C13	0.0760 (16)	0.0672 (14)	0.1084 (19)	-0.0257 (12)	-0.0323 (14)	-0.0023 (14)
C14	0.0591 (12)	0.0493 (11)	0.0636 (12)	0.0003 (9)	-0.0030 (10)	0.0152 (9)
N1	0.0460 (8)	0.0343 (7)	0.0397 (7)	-0.0063 (6)	-0.0043 (6)	0.0042 (5)
N2	0.0496 (8)	0.0340 (7)	0.0452 (7)	-0.0086 (6)	-0.0131 (6)	0.0074 (6)
N3	0.0705 (11)	0.0482 (10)	0.0599 (10)	-0.0145 (8)	-0.0302 (8)	0.0064 (8)
N4	0.0620 (10)	0.0447 (8)	0.0470 (8)	-0.0047 (7)	-0.0112 (7)	0.0032 (6)
N5	0.0623 (10)	0.0561 (10)	0.0474 (8)	-0.0137 (8)	-0.0021 (7)	-0.0009 (7)
N6	0.0813 (15)	0.0677 (12)	0.0659 (12)	-0.0023 (11)	0.0131 (11)	0.0088 (10)
O1	0.0457 (7)	0.0326 (6)	0.0661 (8)	-0.0088 (5)	-0.0019 (6)	0.0034 (5)
O2	0.0556 (8)	0.0396 (7)	0.0740 (9)	-0.0047 (6)	-0.0157 (7)	0.0123 (6)
O3	0.0735 (9)	0.0405 (7)	0.0540 (7)	-0.0196 (6)	-0.0186 (6)	0.0065 (5)
O4	0.0638 (9)	0.0579 (8)	0.0729 (9)	-0.0211 (7)	-0.0159 (7)	-0.0047 (7)
O5	0.0873 (11)	0.0771 (10)	0.0574 (9)	-0.0112 (8)	-0.0233 (8)	-0.0126 (7)
O6	0.0637 (9)	0.0672 (9)	0.0839 (11)	-0.0132 (7)	-0.0017 (8)	0.0137 (8)
C15	0.0932 (19)	0.0754 (16)	0.0761 (15)	0.0252 (13)	-0.0131 (14)	0.0127 (12)
C16	0.095 (2)	0.151 (3)	0.125 (3)	0.047 (2)	0.026 (2)	0.030 (2)
O7	0.0647 (9)	0.0672 (9)	0.0588 (8)	0.0062 (7)	-0.0071 (7)	0.0032 (7)
O1W	0.154 (2)	0.1015 (16)	0.1034 (16)	-0.0471 (15)	0.0175 (16)	0.0006 (12)

Geometric parameters (Å, °)

C1—C6	1.389 (2)	C12—O4	1.326 (3)
C1—C2	1.392 (2)	C13—O4	1.454 (2)
C1—C7	1.484 (2)	C13—H13A	0.9600
C2—C3	1.392 (2)	C13—H13B	0.9600
C2—H2	0.9300	C13—H13C	0.9600
C3—C4	1.385 (3)	C14—O6	1.222 (2)

C3—C11	1.489 (2)	C14—N6	1.333 (3)
C4—C5	1.385 (3)	C14—N5	1.387 (3)
C4—H4	0.9300	N1—N2	1.3476 (17)
C5—C6	1.385 (2)	N2—H2A	0.8600
C5—H5	0.9300	N3—H3A	0.86 (3)
C6—H6	0.9300	N3—H3B	0.85 (3)
C7—N1	1.290 (2)	N4—N5	1.356 (2)
C7—C8	1.498 (2)	N5—H5A	0.8600
C8—O2	1.206 (2)	N6—H6A	0.84 (3)
C8—O1	1.3224 (19)	N6—H6B	0.90 (3)
C9—O1	1.453 (2)	C15—O7	1.408 (3)
C9—H9A	0.9600	C15—C16	1.448 (4)
C9—H9B	0.9600	C15—H15A	0.9700
C9—H9C	0.9600	C15—H15B	0.9700
C10—O3	1.234 (2)	C16—H16A	0.9600
C10—N3	1.325 (2)	C16—H16B	0.9600
C10—N2	1.377 (2)	C16—H16C	0.9600
C11—N4	1.283 (2)	O7—H7	0.88 (3)
C11—C12	1.499 (2)	O1W—H1WA	0.8501
C12—O5	1.209 (2)	O1W—H1WB	0.8495
C6—C1—C2	119.41 (15)	O4—C13—H13A	109.5
C6—C1—C7	120.66 (15)	O4—C13—H13B	109.5
C2—C1—C7	119.90 (15)	H13A—C13—H13B	109.5
C1—C2—C3	120.53 (16)	O4—C13—H13C	109.5
C1—C2—H2	119.7	H13A—C13—H13C	109.5
C3—C2—H2	119.7	H13B—C13—H13C	109.5
C4—C3—C2	119.55 (16)	O6—C14—N6	125.3 (2)
C4—C3—C11	122.79 (15)	O6—C14—N5	118.5 (2)
C2—C3—C11	117.63 (16)	N6—C14—N5	116.2 (2)
C3—C4—C5	120.03 (16)	C7—N1—N2	118.53 (13)
C3—C4—H4	120.0	N1—N2—C10	119.73 (13)
C5—C4—H4	120.0	N1—N2—H2A	120.1
C6—C5—C4	120.47 (18)	C10—N2—H2A	120.1
C6—C5—H5	119.8	C10—N3—H3A	121 (2)
C4—C5—H5	119.8	C10—N3—H3B	117.5 (16)
C5—C6—C1	119.98 (17)	H3A—N3—H3B	121 (2)
C5—C6—H6	120.0	C11—N4—N5	119.39 (15)
C1—C6—H6	120.0	N4—N5—C14	119.03 (16)
N1—C7—C1	125.35 (14)	N4—N5—H5A	120.5
N1—C7—C8	113.16 (14)	C14—N5—H5A	120.5
C1—C7—C8	121.49 (13)	C14—N6—H6A	122 (2)
O2—C8—O1	123.61 (15)	C14—N6—H6B	115.9 (16)
O2—C8—C7	125.05 (15)	H6A—N6—H6B	122 (3)
O1—C8—C7	111.33 (14)	C8—O1—C9	116.52 (14)
O1—C9—H9A	109.5	C12—O4—C13	116.15 (18)
O1—C9—H9B	109.5	O7—C15—C16	113.2 (2)
H9A—C9—H9B	109.5	O7—C15—H15A	108.9

O1—C9—H9C	109.5	C16—C15—H15A	108.9
H9A—C9—H9C	109.5	O7—C15—H15B	108.9
H9B—C9—H9C	109.5	C16—C15—H15B	108.9
O3—C10—N3	124.35 (16)	H15A—C15—H15B	107.8
O3—C10—N2	118.54 (15)	C15—C16—H16A	109.5
N3—C10—N2	117.12 (15)	C15—C16—H16B	109.5
N4—C11—C3	124.74 (15)	H16A—C16—H16B	109.5
N4—C11—C12	113.41 (16)	C15—C16—H16C	109.5
C3—C11—C12	121.77 (17)	H16A—C16—H16C	109.5
O5—C12—O4	124.16 (18)	H16B—C16—H16C	109.5
O5—C12—C11	124.6 (2)	C15—O7—H7	111 (2)
O4—C12—C11	111.25 (17)	H1WA—O1W—H1WB	109.6

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>
O1 <i>W</i> —H1 <i>WA</i> ...O6	0.85	2.05	2.879 (3)	166
O1 <i>W</i> —H1 <i>WB</i> ...O5 ⁱ	0.85	2.16	2.941 (3)	152
N3—H3 <i>A</i> ...O7	0.86 (4)	2.13 (4)	2.980 (2)	169 (3)
N3—H3 <i>B</i> ...O5 ⁱⁱ	0.85 (3)	2.61 (2)	3.082 (2)	116.7 (18)
N5—H5 <i>A</i> ...O3 ⁱⁱⁱ	0.86	2.19	3.010 (2)	160
N6—H6 <i>A</i> ...O1 <i>W</i> ^{iv}	0.84 (3)	2.34 (3)	3.137 (3)	158 (3)
N6—H6 <i>B</i> ...O7 ^v	0.90 (3)	1.99 (3)	2.871 (3)	166 (2)
O7—H7...O2	0.88 (3)	1.96 (3)	2.801 (2)	158 (3)
O7—H7...N1	0.88 (3)	2.52 (3)	3.0677 (19)	121 (2)
C4—H4...O6 ^{vi}	0.93	2.57	3.493 (2)	173
C13—H13 <i>C</i> ...N6 ^{vii}	0.96	2.62	3.405 (3)	139

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