

The unusual (*syn*-/*anti*-)₂ conformation of a dimethoxypyrimidyl-based tennimide

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Received 25 May 2023

Accepted 3 August 2023

Edited by J. Ellena, Universidade de São Paulo, Brazil

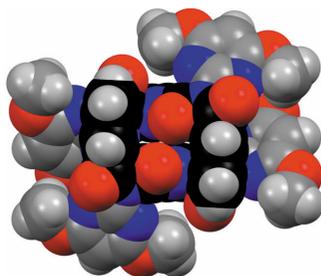
Keywords: conformation; macrocycle; methoxy; pyridine; pyrimidine; tennimide; tetramer; crystal structure.**CCDC reference:** 2286597**Supporting information:** this article has supporting information at journals.iucr.org/e

The tennimide macrocycle, (**I**) (C₅₂H₄₀N₁₆O₁₆·0.167H₂O), was synthesized from 2-amino-4,6-dimethoxypyrimidine and pyridine-2,6-dicarbonyl dichloride. Compound (**I**) represents the first tennimide incorporating pyridine rings in the macrocycle scaffold. In the macrocycle ring, the carbonyl groups at each successive dicarbonyl(pyridine) moiety adopt the (*syn/anti*)₂ conformation. This contrasts with all previously reported tetraimide macrocycles, which exhibit the (*syn*)₄ conformation. The effect is to close any potential cavity or *niche* by having two of the central pyridine C₅N rings aligned close to each other [with closest pyridine C_g··C_g ring centroid separations of 3.5775 (19) Å; closest C··C = 3.467 (5) Å]. A partial occupancy water molecule (with s.o.f. = 0.167), resides with its oxygen atom on a twofold axis at hydrogen-bonding distances to the carbonyl O atom, in a molecular *niche* between two pyridine rings. Macrocycles of (**I**) have all six C=O groups and all eight methoxy O atoms present on the macrocycle surface. However, all twelve N atoms are effectively shielded on steric grounds from any potential intermolecular interactions. The remaining two C=O O atoms interact with the partial occupancy water molecule *via* two O—H··O=C hydrogen bonds. Macrocycles of (**I**) stack as one-dimensional chains along the *b*-axis direction with primary intermolecular interactions involving weak C—H··O=C/OCH₃/H₂O contacts. Chains interlock weakly *via* methoxy–methoxy C—H··O interactions into two-dimensional sheets.

1. Chemical context

Developments in macrocyclic chemistry continue with an emphasis on structure, function and developing new architectures with pendant functional groups. An aim is to achieve new applications in coordination chemistry, nanoscience, natural products, medicinal chemistry and applied sciences (Böhmer, 1995; Vicens & Harrowfield, 2007). Macrocyclic science now spans several scientific fields and finds applications across a vast range of chemical, physical and biomedical sciences (Gloe, 2005; Davis & Higson, 2011).

Macrocycles usually contain donor atoms such as O, N, S and P and are utilized with a wide variety of aliphatic groups (*e.g.* in crown ethers) and/or aromatic rings (*e.g.* in calixarenes, porphyrins) (Böhmer, 1995; Gloe, 2005). An objective is that particular functional groups are incorporated onto a scaffold to accommodate a wide range of metals, their oxidation states and coordination chemistry geometries. This originates from the initial crown ethers, through macrocycles such as calix[*n*]arenes (Böhmer, 1995) to macromolecular macrocycles (Davis & Higson, 2011). Constituent functional groups now span a relatively large range of types as incorporated into many macrocycles in the form of amides, pyridines, imides *etc.* (Pappalardo *et al.*, 1992*a,b*; Böhmer, 1995; Vicens &



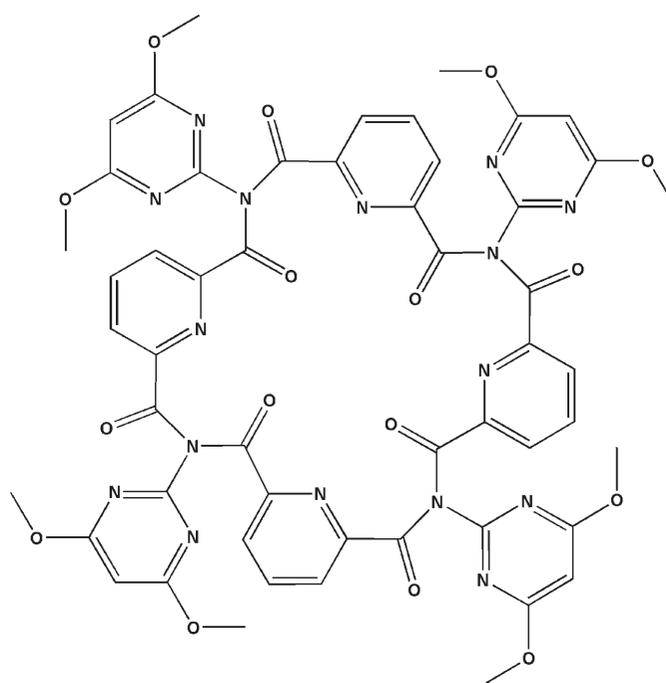
Harrowfield, 2007). Recently, considerable effort has been made to incorporate biological moieties (*e.g.* peptide chains) for a range of applications including artificial ion channels and transport (Xin *et al.*, 2015; Legrand & Barboiu, 2013).

In terms of macrocyclic conformations, there are many examples where a macrocycle is isolated in a defined, stable geometry and subsequently shown to adopt alternate conformations. Usually these can be structurally characterized and the different conformations may or may not be interconvertible in solution or even in the solid state. For example, the cone conformation of calix[4]arenes has been well described (Andreotti *et al.*, 1979; Gutsche, 1983). In addition, the partial cone, *syn*-distal and *syn*-proximal conformations have been studied in many different calix[4]arene derivatives (Gutsche *et al.*, 1983; Ferguson *et al.*, 1992, 1993; Pappalardo *et al.*, 1992*a,b*; Shinkai, 1993). Even in tennimide chemistry, we were fortunate to isolate a tennimide known as (26IO)₄ with three distinct solid-state geometries (Mocilac & Gallagher, 2013), though these are interconvertible in solution. The three (26IO)₄ conformations differ in terms of the size and apertures of the molecular cavity. For the smaller trezimide (trimer) systems, two conformations have been isolated as the distinct *P* and *R* conformations (Mocilac & Gallagher, 2013). Therefore, isolation and characterization of a new macrocyclic conformation in a class of imide-based macrocycles is of interest to researchers studying imide-based and related macrocycles. Furthermore, researchers have continued to advance conformational analysis and especially with respect to non-rigid macrocycles (Bohle & Grimme, 2022). Their study utilized the automated generation of macrocyclic conformers using computational methods as applied to ¹³C-NMR data of flexible cycloalkanes. With such developments, new macrocyclic conformations will be postulated for a range of macrocycle types in tandem with synthetic experimental studies. In addition, it has recently been shown using synthetic strategies, the engineering of molecular topology in pseudo-peptidic macrocycles (Sharma *et al.*, 2017).

Of particular note is that Balakrishna and co-workers have studied phosphorus-based systems using >PCH₃ and >PC₆H₅ as linker groups between the isophthaloyl moieties (Balakrishna, 2018; Kashid *et al.*, 2017). This contrasts with the >N(pyridine) and >N(pyrimidine) linkers that we have studied to date. Structural examples of the P-based crystal structures as VAWVIB and VAWVOH are available on the CSD (Groom *et al.*, 2016). In addition, the highly constrained butterfly structures as ‘dimers’ have been reported as RAYFII and ZAFJAV (Saunders *et al.*, 2012; Pearce & Crossley, 2020). He and co-workers in their structures NUKZIF, NULHOU, NULHUA (Wang *et al.*, 2020) noted that such diimides and polyimides are still relatively rare (Wang *et al.*, 2020). These systems are more closely related to the well-explored planar diimides such as naphthalene diimide (Takenaka, 2021).

From previous benzamide studies (Donnelly *et al.*, 2008), we reported macrocyclic trimers (trezimides) and tetramers (tennimides) based on the isophthaloyl residue and imide linker group (Evans & Gale, 2004; Mocilac & Gallagher, 2013, 2014, 2016; Gallagher & Mocilac, 2021). These macrocycles

with pendant pyridine, pyrimidine and pyridinyl ester groups are attached to the central scaffold and have potential to bind to metal complexes. Trimers and tetramers are typically synthesized and isolated in modest yields, together with oligomers and polymers from which they have to be carefully separated by column chromatography. We herein report a new pyridine-based tennimide macrocycle derived from 2-amino-4,6-dimethoxypyrimidine and pyridine-2,6-dicarbonyldichloride (Fig. 1) with an unusual (*syn/anti*)₂ tennimide conformation (Fig. 2). The isolation of a new conformation using a pyridine-based scaffold demonstrates that these macrocycles can be investigated to exploit this new molecular conformation.



2. Structural commentary

The macrocycle (**I**), as synthesized from the condensation reaction of 2-amino-4,6-dimethoxypyrimidine with pyridine-2,6-dicarbonyldichloride, adopts a different (*syn/anti*)₂ conformation to the (*syn*)₄ seen in all reported tennimides (Evans & Gale, 2004; Mocilac & Gallagher, 2013, 2014, 2016) (Figs. 1, 2, 3). (**I**) represents the first tennimide synthesized using pyridine-2,6-dicarbonyl dichloride (in comparison to benzene dicarbonyldichloride) and thus has four pyridines incorporated into the scaffold. It comprises four pyridine N donor, eight pyrimidine N, eight carbonyl O donor atoms, together with eight methoxy groups and four aliphatic imide N atoms. However, the orientation of most aromatic N donors is not suitable for coordination due to shielding from the eight methoxy groups on the external surface of the macrocycle. There is no inner macrocyclic cavity because of the spatial arrangement of the pyridine rings in the (*syn/anti*)₂ conformation. However, a partial occupancy water molecule is

observed in a macrocyclic niche at hydrogen-bonding distances to two imide carbonyl oxygen atoms (Fig. 2).

In all tennimides (Evans & Gale, 2004; Mocilac & Gallagher, 2013, 2014, 2016; Gallagher & Mocilac, 2021) the carbonyl group conformations with respect to the isophthaloyl residues is noted as *syn*-. Therefore, these macrocycles are classed as having a $(syn)_4$ conformation (Mocilac & Gallagher, 2013, 2014, 2016; Gallagher & Mocilac, 2021). In (**I**), the pyridine conformation with respect to the carbonyl groups alternates as *syn*- and *anti*- and is defined as $(syn/anti)_2$. This $(syn/anti)_2$ conformation has no internal cavity available to incorporate an ion or small molecule (although a small niche is present). The geometric details are: two of the pyridine C₅N rings are positioned close to each other [with closest pyridine ring centroid C_g··C_g separations = 3.5775 (19) Å; with closest C24··C24_a = 3.467 (5) Å across the twofold axis; where $a = -x, y, \frac{1}{2} - z$]. In the macrocycle, the imide '(O=C)N' hinge O=C··C=O torsion angles are oriented at -92.5 (4) and -91.59 (5)° for O1=C1··C2=O2 and O3=C3··C4=O4, respectively, which are typical of imide conformations in tennimides. However, the pyridine dicarbonyl moiety torsion angles differ with 87.0 (5)° for the *syn*-conformation (in O1=C1··C3=O3) and -144.89 (5)° for the *anti*-conformation (in O2=C2··C4=O4). The *syn*-torsion data is close to the average isophthaloyl (O=C)C₆H₄(C=O) torsion angle noted previously (Evans & Gale, 2004; Mocilac & Gallagher, 2013, 2014, 2016; Gallagher & Mocilac, 2021). In the four dimethoxypyrimidine rings, the methoxy groups all adopt similar conformations, as noted previously (Gallagher *et al.*, 2001). The proximally related pyridine rings are almost orthogonal to one another at 87.00 (10), 87.09 (10)° and with the opposite (distal-related) pyridine rings almost parallel at

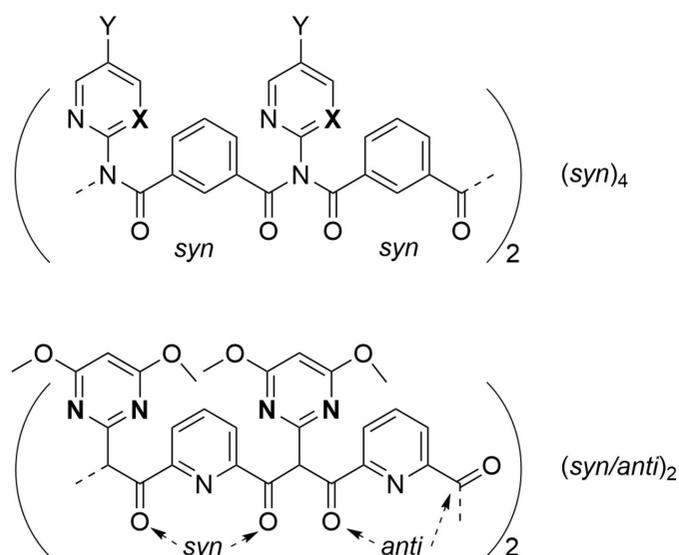


Figure 1
Schematic diagrams of (top) the tennimide (**I**) with the relative geometries depicted to minimize atom/group overlap and (bottom) the relative differences between the $(syn)_4$ and $(syn/anti)_2$ -conformations. The X and Y labels refer to: X = C or N and Y = H, CH₃ or halogen atom (F, Cl, Br).

Table 1
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>
C15—H15···O3 ⁱ	0.93	2.47	3.148 (4)	130
C15—H15···O4 ⁱⁱ	0.93	2.42	3.084 (4)	129
C18A—H18C···O23A	0.96	2.47	3.393 (6)	160
C24—H24···O1W ⁱⁱⁱ	0.93	2.49	3.14 (2)	127
C26—H26···N22A	0.93	2.62	3.325 (4)	133
C27A—H27B···O23A ^{iv}	0.96	2.57	3.144 (4)	119
O1W—H1W···O2	0.85	2.32	2.97 (3)	134 (5)

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

9.25 (11), 0.97 (11)°. For symmetry-related pyridine rings, the C=O groups are positioned *anti*- with respect to one another.

A partial occupancy water molecule O1W (site occupancy of 0.167) occupies a niche between two distal pyridines (with *syn*-related C=O groups), separated by *ca* 6 Å. The water molecule O1W forms O—H···O=C hydrogen bonds [O···O = 2.97 (3) Å] with two symmetry-related carbonyl O2 atoms and two weaker C—H···O complete the hydrogen bonding (H···O = 2.49 Å).

3. Supramolecular features

The primary interactions involving the macrocycle are a range of rather weak aliphatic/aromatic C—H···O contacts (Table 1) in the absence of strong hydrogen-bond donors. This arises in a system with a vast excess of potentially strong acceptor groups on the tennimide surface. Molecules of (**I**) aggregate as 1D chains along the *b*-axis direction with primary intermolecular interactions involving weak C—H···O=C/OCH₃/H₂O contacts (Fig. 4). Chains interlock weakly *via* methoxy···methoxy C—H···O interactions into 2D sheets. This type of 1D aggregation is quite common for this class of

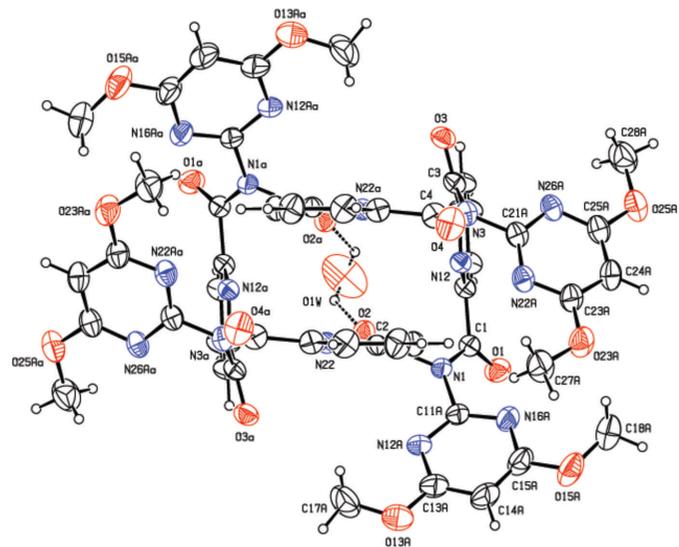


Figure 2
A view of (**I**) with the atomic-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level for clarity. The partial occupancy water molecule O1W is depicted with dashed lines representing the hydrogen bonding to (**I**). Symmetry code: (a) $-x, y, -z + \frac{1}{2}$.

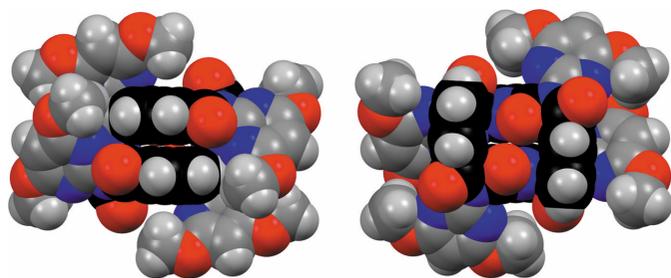


Figure 3
Two views of (**I**) with all atoms drawn as their van der Waals spheres and the central pyridine backbone drawn in black using the *Mercury* program (Macrae *et al.*, 2020).

tennimide macrocycle. It was noted in the 1D macrocyclic stacking driven by $C-Br \cdots O=C/N_{\text{pyridine}}$ halogen bonding between molecules in brominated tennimides (Mocilac & Gallagher, 2013).

4. Synthesis and crystallization

Synthetic reaction conditions:

Pyridine-2,6-dicarbonyl dichloride (2.041 g, 10 mmol, 1 eq.) was dissolved in 100 ml of dry CH_2Cl_2 (DCM) and a catalytic quantity of 14.4 mg 4-dimethylaminopyridine (DMAP) with 4 ml (29.4 mmol, 3 eq.) of Et_3N were added to the solution under N_2 . This solution was cooled to 255 K and 1.64 g (10 mmol, 1 eq.) of 4,6-dimethoxyppyrimidin-2-amine dissolved

in 40 ml dry of DCM were added. The solution was stirred for 72 h and thin layer chromatography (TLC: CH_2Cl_2 /ethyl acetate, 4:1) indicated that conversion was incomplete (the related aminopyridine reactants show full conversion within 24 h). The solution was then diluted to 250 ml with technical grade CH_2Cl_2 , washed four times with 100 ml of NH_4Cl solution (pH 4), dried using MgSO_4 and the solvent mixture removed at reduced pressure. Filtration through silica gel (CH_2Cl_2 /ethyl acetate, 2:1) was performed before final purification was attempted by column chromatography. The expected products were the acyclic (2:1), (3:2) and (4:3) mixed imide benzamides as well as longer chain oligomers and some polymeric materials. TLC indicated four bands (one of which was the pyrimidin-2-amine starting material) in addition to polymeric material at the bottom of the TLC plate. The polymer was mostly removed by preliminary filtration. Further attempts at purification again involved multiple steps of column chromatography. The first column (using *n*-hexane/ethyl acetate, 2:1) resulted in poor separation and merely eliminated the remaining starting material. A subsequent column (CH_2Cl_2 /ethyl acetate, 10:1) gave some separation and two products were isolated. The first product is the (2:1) pyridine-2,6-dicarboxamide and the second was shown to be a (4:4) macrocyclic tennimide (**I**) isolated in a relatively low yield of *ca* 5%. The final major product could not be isolated and purified.

N^2,N^6 -bis(4,6-dimethoxyppyrimidin-2-yl)pyridine-2,6-dicarboxamide (2:1 product) $^1\text{H-NMR}$ (400 MHz, CDCl_3): δ = 10.24 (*s*, 2H, NH), 8.50 (*d*, J = 7.8 Hz, 2H, Ar-H), 8.12 (*t*, J = 7.8 Hz, 1H, Ar-H), 5.81 (*s*, 2H, Ar-H), 3.94 (*s*, 2H, O- CH_3). $^{13}\text{C-NMR}$ (100 MHz, CDCl_3): δ = 172.1 (4C, Ar- C_q), 160.5 (2C, Carbonyl- C_q), 155.8 (2C, Ar- C_q), 148.5 (2C, Ar- C_q), 139.8 (1C, Ar- C-H), 126.4 (2C, Ar- C-H), 86.0 (2C, Ar- C-H), 54.2 (4C, O- CH_3). **IR**: 3460, 3386, 3245, 3101, 3052, 2996, 2924, 2854, 2580, 2163, 1708, 1598, 1573, 1509, 1478, 1455, 1424, 1365, 1300, 1256, 1217, 1190, 1162, 1141, 1094, 1063, 1041, 1000, 987, 958, 909, 879, 856, 834, 777, 754, 696, 663 cm^{-1} .

Tennimide macrocycle [4:4 product or (I**)]** $^1\text{H-NMR}$ (400 MHz, CDCl_3): δ = 8.13 (*m*, 2H, Ar-H), 8.03 (*m*, 2H, Ar-H), 7.96 (*s*, 4H, Ar-H), 7.80 (*m*, 2H, Ar-H), 7.72 (*m*, 2H, Ar-H), 5.67 (*s*, 2H, Ar-H), 5.63 (*s*, 2H, Ar-H), 3.58 (*s*, 12H, O- CH_3), 3.50 (*s*, 12H, O- CH_3). **IR**: 3100, 2957, 2924, 2853, 1733, 1711, 1586, 1557, 1467, 1402, 1363, 1309, 1282, 1192, 1157, 1086, 1077, 1054, 995, 937, 923, 842, 824, 778, 741, 709, 652 cm^{-1} .

5. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. H atoms attached to C atoms were treated as riding using the *SHELXL14* (Sheldrick, 2015b) defaults at 294 (1) K with $C-H$ = 0.93 Å (aromatic) and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ (aromatic); the methyl $C-H$ = 0.96 Å (aliphatic) and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$. The H atoms of the partial occupancy water molecule were treated using three DFIX restraints at chemically sensible positions and directed towards the closest $O=C$ acceptor groups. The presence of

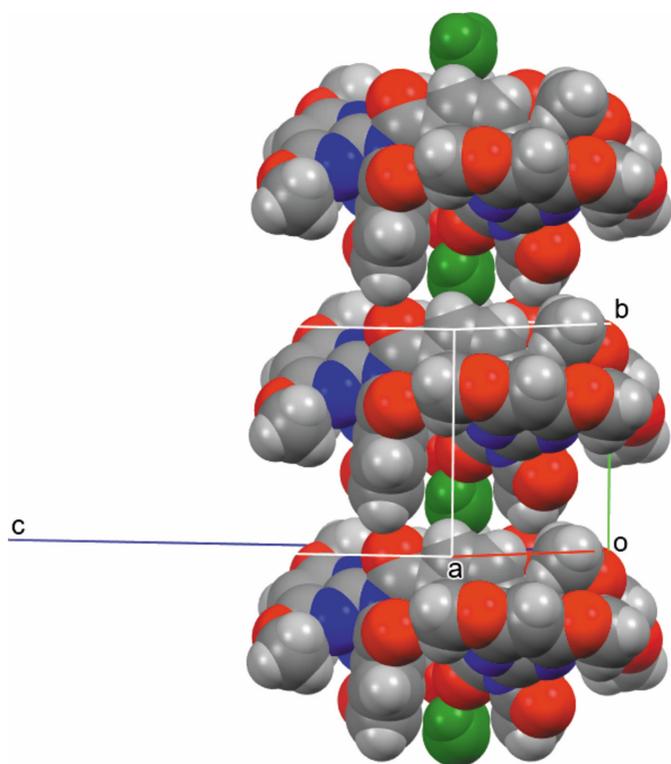


Figure 4
A view of the primary stacking in the crystal structure of (**I**) along the *b*-axis direction with atoms drawn as their van der Waals spheres and partial occupancy water O1W in green.

the water in this location is similar to that noted in a 26(BrIO)₄ structure (XOCHUU; Mocilac & Gallagher, 2014) where a hemihydrate spans a pyridine N atom and a carbonyl O=C by intermolecular hydrogen bonding at the macrocycle cavity entrance.

Structural analysis in the penultimate stages of refinement demonstrates that by omitting the partial occupancy water molecule, the *R*-factor increases from 0.054 to 0.056. The residual electron density increases from +0.19 to 0.58 e Å⁻³, resulting in a single peak of residual electron density on the twofold axis. This is where the partial occupancy water molecule is located. The WGHT card increases from 0.067 to 0.081. There is no other atom or group disorder in the structure of (I).

Acknowledgements

JFG thanks Dublin City University for grants in aid of chemical research. The research was part-funded under the Programme for Research in Third Level Institutions (PRTL) Cycle 4 (Ireland). It was co-funded through the European Regional Development Fund (ERDF), part of the European Union Structural Funds Programme (ESF) 2007–2013.

Funding information

Funding for this research was provided by: Dublin City University.

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Table 2

Experimental details.

Crystal data	
Chemical formula	C ₅₂ H ₄₀ N ₁₆ O ₁₆ ·0.167H ₂ O
<i>M_r</i>	1148.02
Crystal system, space group	Orthorhombic, <i>Pbcn</i>
Temperature (K)	294
<i>a</i> , <i>b</i> , <i>c</i> (Å)	18.8065 (10), 10.0745 (6), 28.847 (3)
<i>V</i> (Å ³)	5465.6 (7)
<i>Z</i>	4
Radiation type	Cu Kα
μ (mm ⁻¹)	0.91
Crystal size (mm)	0.39 × 0.30 × 0.04
Data collection	
Diffractometer	Xcalibur, Sapphire3, Gemini Ultra
Absorption correction	Analytical (ABSFAC; Clark & Reid, 1998)
<i>T_{min}</i> , <i>T_{max}</i>	0.778, 0.973
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	14675, 4439, 3036
<i>R_{int}</i>	0.038
(sin θ/λ) _{max} (Å ⁻¹)	0.581
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.054, 0.159, 1.03
No. of reflections	4439
No. of parameters	392
No. of restraints	3
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
Δρ _{max} , Δρ _{min} (e Å ⁻³)	0.19, -0.18

Computer programs: *CrysAlis PRO* (Rigaku OD, 2015), *SHELXT14/6* (Sheldrick, 2015a), *SHELXL14/6* (Sheldrick, 2015b), *PLATON* (Spek, 2020) and *Mercury* (Macrae *et al.*, 2020).

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supporting information

Acta Cryst. (2023). E79, 837-841 [https://doi.org/10.1107/S2056989023006837]

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Computing details

Data collection: *CrysAlis PRO* (Rigaku OD, 2015); cell refinement: *CrysAlis PRO* (Rigaku OD, 2015); data reduction: *CrysAlis PRO* (Rigaku OD, 2015); program(s) used to solve structure: *SHELXT14/6* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL14/6* (Sheldrick, 2015b); molecular graphics: *PLATON* (Spek, 2020) and *Mercury* (Macrae *et al.*, 2020); software used to prepare material for publication: *SHELXL14/6* (Sheldrick, 2015b).

3,11,19,27-Tetrakis(4,6-dimethoxypyrimidin-2-yl)-3,11,19,27,33,34,35,36-octaazapentacyclo[27.3.1.1^{5,9}.1^{13,17}.1^{21,25}]hexatriaconta-1(33),5,7,9(36),13(35),14,16,21(34),22,24,29,31-dodecaen-2,4,10,12,18,20,26,28-octone 0.167-hydrate

Crystal data

C₅₂H₄₀N₁₆O₁₆·0.167H₂O

M_r = 1148.02

Orthorhombic, *Pbcn*

a = 18.8065 (10) Å

b = 10.0745 (6) Å

c = 28.847 (3) Å

V = 5465.6 (7) Å³

Z = 4

F(000) = 2374.7

D_x = 1.395 Mg m⁻³

Cu *Kα* radiation, λ = 1.54184 Å

Cell parameters from 2330 reflections

θ = 2.8–63.6°

μ = 0.91 mm⁻¹

T = 294 K

Plate, colourless

0.39 × 0.30 × 0.04 mm

Data collection

Xcalibur, Sapphire3, Gemini Ultra diffractometer

Radiation source: Enhance Ultra (Cu) X-ray Source

Mirror monochromator

ω scans

Absorption correction: analytical (*ABSFAC*; Clark & Reid, 1998)

T_{min} = 0.778, *T_{max}* = 0.973

14675 measured reflections

4439 independent reflections

3036 reflections with *I* > 2σ(*I*)

R_{int} = 0.038

θ_{max} = 63.6°, θ_{min} = 3.1°

h = -19→21

k = -6→11

l = -33→30

2330 standard reflections every 60 min

intensity decay: 1%

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.054

wR(*F*²) = 0.159

S = 1.03

4439 reflections

392 parameters

3 restraints

Primary atom site location: structure-invariant direct methods

Secondary atom site location: inferred from neighbouring sites

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0686P)^2 + 1.3472P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.19 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$$

Extinction correction: SHELXL,

$$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$$

Extinction coefficient: 0.00055 (7)

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	-0.05721 (12)	0.3212 (2)	0.40663 (8)	0.0750 (6)	
C1	-0.03715 (15)	0.3931 (3)	0.37594 (10)	0.0543 (7)	
N1	-0.07837 (12)	0.4970 (2)	0.35800 (8)	0.0544 (6)	
O2	-0.06477 (11)	0.43970 (19)	0.28185 (7)	0.0598 (5)	
C2	-0.07302 (13)	0.5274 (3)	0.30973 (10)	0.0504 (7)	
O3	0.22174 (11)	0.6319 (2)	0.29594 (7)	0.0652 (6)	
C3	0.16807 (15)	0.6223 (3)	0.31815 (10)	0.0527 (7)	
N3	0.13350 (13)	0.7385 (2)	0.33702 (8)	0.0548 (6)	
O4	0.11289 (15)	0.9585 (2)	0.32596 (8)	0.0876 (8)	
C4	0.11694 (17)	0.8484 (3)	0.31003 (11)	0.0619 (8)	
C11	0.13402 (14)	0.4934 (3)	0.33012 (10)	0.0499 (6)	
N12	0.06713 (11)	0.4984 (2)	0.34536 (7)	0.0487 (5)	
C13	0.03634 (14)	0.3830 (3)	0.35609 (9)	0.0487 (6)	
C14	0.07002 (16)	0.2620 (3)	0.35217 (11)	0.0621 (8)	
H14	0.0469	0.1839	0.3605	0.074*	
C15	0.13824 (17)	0.2597 (3)	0.33575 (13)	0.0738 (10)	
H15	0.1622	0.1794	0.3324	0.089*	
C16	0.17103 (16)	0.3770 (3)	0.32431 (12)	0.0655 (8)	
H16	0.2173	0.3776	0.3129	0.079*	
C11A	-0.13160 (16)	0.5612 (3)	0.38435 (10)	0.0612 (8)	
N12A	-0.18786 (13)	0.5977 (3)	0.36019 (9)	0.0684 (7)	
C13A	-0.2347 (2)	0.6711 (4)	0.38399 (14)	0.0897 (12)	
C14A	-0.2263 (2)	0.6991 (5)	0.43042 (14)	0.1027 (14)	
H14A	-0.2600	0.7475	0.4469	0.123*	
C15A	-0.1659 (2)	0.6523 (4)	0.45107 (12)	0.0858 (11)	
N16A	-0.11615 (15)	0.5832 (3)	0.42871 (9)	0.0714 (8)	
O13A	-0.29127 (15)	0.7190 (4)	0.36166 (12)	0.1228 (11)	
C17A	-0.2926 (3)	0.7082 (6)	0.31264 (19)	0.1275 (18)	
H17A	-0.2552	0.7613	0.2996	0.191*	
H17B	-0.3377	0.7387	0.3012	0.191*	
H17C	-0.2858	0.6171	0.3039	0.191*	
O15A	-0.15514 (19)	0.6812 (3)	0.49625 (9)	0.1107 (10)	

C18A	-0.0908 (3)	0.6365 (5)	0.51825 (14)	0.1134 (15)	
H18A	-0.0829	0.5449	0.5108	0.170*	
H18B	-0.0952	0.6462	0.5512	0.170*	
H18C	-0.0515	0.6887	0.5074	0.170*	
C21	-0.07603 (14)	0.6707 (3)	0.29661 (9)	0.0504 (7)	
N22	-0.09433 (12)	0.6954 (2)	0.25293 (8)	0.0502 (6)	
C23	-0.09991 (17)	0.8212 (3)	0.23984 (10)	0.0581 (7)	
C24	-0.0874 (2)	0.9276 (3)	0.26911 (13)	0.0771 (10)	
H24	-0.0937	1.0146	0.2591	0.093*	
C25	-0.0652 (2)	0.9002 (3)	0.31365 (13)	0.0816 (11)	
H25	-0.0547	0.9689	0.3341	0.098*	
C26	-0.05880 (17)	0.7711 (3)	0.32738 (11)	0.0670 (8)	
H26	-0.0430	0.7508	0.3571	0.080*	
C21A	0.12724 (17)	0.7424 (3)	0.38590 (10)	0.0580 (8)	
N22A	0.07257 (14)	0.8089 (2)	0.40337 (9)	0.0644 (7)	
C23A	0.06838 (19)	0.8052 (3)	0.44966 (11)	0.0704 (9)	
C24A	0.1147 (2)	0.7351 (3)	0.47695 (12)	0.0773 (10)	
H24A	0.1097	0.7312	0.5090	0.093*	
C25A	0.1687 (2)	0.6709 (3)	0.45415 (12)	0.0705 (9)	
N26A	0.17692 (14)	0.6744 (2)	0.40795 (9)	0.0630 (7)	
O23A	0.01530 (15)	0.8719 (2)	0.47037 (8)	0.0903 (8)	
C27A	-0.0336 (2)	0.9460 (4)	0.44195 (14)	0.1050 (14)	
H27A	-0.0076	1.0071	0.4228	0.158*	
H27B	-0.0659	0.9944	0.4614	0.158*	
H27C	-0.0599	0.8860	0.4226	0.158*	
O25A	0.21549 (15)	0.6015 (3)	0.47933 (8)	0.0899 (8)	
C28A	0.2726 (2)	0.5372 (5)	0.45500 (16)	0.1113 (15)	
H28A	0.2535	0.4717	0.4343	0.167*	
H28B	0.3037	0.4950	0.4769	0.167*	
H28C	0.2988	0.6019	0.4376	0.167*	
O1W	0.0000	0.187 (3)	0.2500	0.180 (15)	0.167
H1W	-0.0358 (4)	0.238 (3)	0.248 (3)	0.100*	0.167

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0701 (14)	0.0827 (15)	0.0721 (14)	-0.0008 (12)	0.0112 (12)	0.0203 (12)
C1	0.0507 (17)	0.0590 (17)	0.0533 (17)	-0.0002 (14)	-0.0006 (13)	-0.0020 (14)
N1	0.0467 (13)	0.0683 (15)	0.0483 (13)	0.0126 (12)	0.0028 (10)	0.0008 (11)
O2	0.0654 (13)	0.0597 (12)	0.0544 (12)	0.0076 (10)	-0.0001 (10)	-0.0033 (10)
C2	0.0391 (15)	0.0606 (17)	0.0515 (16)	0.0028 (13)	-0.0011 (12)	-0.0036 (14)
O3	0.0487 (12)	0.0717 (14)	0.0752 (14)	-0.0099 (10)	-0.0037 (10)	-0.0028 (10)
C3	0.0463 (16)	0.0493 (16)	0.0625 (18)	-0.0019 (13)	-0.0076 (14)	-0.0046 (13)
N3	0.0650 (15)	0.0426 (12)	0.0567 (14)	-0.0054 (11)	-0.0084 (12)	-0.0050 (10)
O4	0.129 (2)	0.0410 (12)	0.0928 (17)	-0.0117 (13)	-0.0023 (15)	-0.0104 (11)
C4	0.072 (2)	0.0402 (16)	0.073 (2)	-0.0086 (14)	0.0004 (16)	-0.0058 (14)
C11	0.0437 (15)	0.0464 (15)	0.0596 (16)	-0.0008 (13)	-0.0039 (13)	-0.0072 (12)
N12	0.0469 (13)	0.0459 (12)	0.0532 (13)	0.0028 (11)	-0.0017 (10)	-0.0055 (10)

C13	0.0459 (15)	0.0459 (15)	0.0541 (16)	-0.0008 (12)	-0.0036 (12)	0.0004 (12)
C14	0.0559 (18)	0.0419 (15)	0.088 (2)	-0.0023 (14)	0.0000 (16)	-0.0003 (14)
C15	0.0575 (19)	0.0411 (16)	0.123 (3)	0.0085 (15)	0.0102 (19)	-0.0078 (17)
C16	0.0487 (17)	0.0519 (17)	0.096 (2)	0.0026 (14)	0.0069 (16)	-0.0086 (15)
C11A	0.0531 (18)	0.072 (2)	0.0581 (19)	0.0104 (16)	0.0079 (14)	-0.0033 (15)
N12A	0.0498 (15)	0.0873 (19)	0.0681 (17)	0.0140 (14)	0.0091 (12)	-0.0036 (14)
C13A	0.068 (2)	0.110 (3)	0.091 (3)	0.029 (2)	0.015 (2)	-0.007 (2)
C14A	0.095 (3)	0.133 (4)	0.080 (3)	0.038 (3)	0.025 (2)	-0.019 (2)
C15A	0.101 (3)	0.095 (3)	0.061 (2)	0.016 (2)	0.015 (2)	-0.0103 (19)
N16A	0.0756 (18)	0.0832 (19)	0.0553 (16)	0.0099 (15)	0.0056 (13)	-0.0074 (13)
O13A	0.0773 (19)	0.177 (3)	0.114 (2)	0.062 (2)	0.0076 (17)	-0.011 (2)
C17A	0.092 (3)	0.164 (5)	0.126 (4)	0.048 (3)	-0.036 (3)	-0.026 (4)
O15A	0.147 (3)	0.124 (2)	0.0608 (16)	0.022 (2)	0.0187 (17)	-0.0204 (15)
C18A	0.145 (4)	0.128 (4)	0.067 (3)	0.002 (3)	-0.006 (3)	-0.008 (2)
C21	0.0460 (15)	0.0531 (16)	0.0520 (16)	-0.0005 (13)	0.0010 (12)	-0.0060 (13)
N22	0.0504 (13)	0.0455 (13)	0.0547 (14)	0.0051 (10)	-0.0002 (10)	-0.0015 (10)
C23	0.0652 (19)	0.0432 (16)	0.0660 (19)	0.0025 (14)	0.0029 (15)	-0.0047 (14)
C24	0.097 (3)	0.0496 (18)	0.085 (3)	0.0003 (18)	0.004 (2)	-0.0084 (16)
C25	0.105 (3)	0.064 (2)	0.076 (2)	-0.013 (2)	-0.003 (2)	-0.0195 (18)
C26	0.069 (2)	0.071 (2)	0.0607 (19)	-0.0050 (17)	-0.0029 (16)	-0.0110 (15)
C21A	0.067 (2)	0.0444 (15)	0.0626 (19)	-0.0084 (15)	-0.0097 (16)	-0.0120 (13)
N22A	0.0752 (18)	0.0561 (15)	0.0617 (16)	-0.0021 (13)	-0.0107 (13)	-0.0155 (12)
C23A	0.086 (2)	0.0623 (19)	0.063 (2)	-0.0023 (18)	-0.0076 (18)	-0.0193 (16)
C24A	0.100 (3)	0.074 (2)	0.058 (2)	0.000 (2)	-0.0156 (19)	-0.0111 (17)
C25A	0.082 (2)	0.0599 (19)	0.070 (2)	-0.0039 (17)	-0.0193 (19)	-0.0057 (16)
N26A	0.0690 (17)	0.0546 (14)	0.0654 (16)	-0.0052 (13)	-0.0150 (13)	-0.0086 (12)
O23A	0.110 (2)	0.0919 (17)	0.0689 (15)	0.0239 (16)	-0.0025 (14)	-0.0257 (13)
C27A	0.112 (3)	0.111 (3)	0.091 (3)	0.039 (3)	-0.015 (3)	-0.039 (2)
O25A	0.0992 (19)	0.0954 (18)	0.0752 (16)	0.0111 (15)	-0.0286 (15)	-0.0032 (13)
C28A	0.103 (3)	0.121 (4)	0.110 (3)	0.033 (3)	-0.026 (3)	-0.004 (3)
O1W	0.17 (3)	0.085 (19)	0.28 (4)	0.000	-0.06 (3)	0.000

Geometric parameters (Å, °)

O1—C1	1.205 (3)	C17A—H17B	0.9600
C1—N1	1.402 (3)	C17A—H17C	0.9600
C1—C13	1.499 (4)	O15A—C18A	1.438 (5)
N1—C11A	1.414 (3)	C18A—H18A	0.9600
N1—C2	1.429 (3)	C18A—H18B	0.9600
O2—C2	1.205 (3)	C18A—H18C	0.9600
C2—C21	1.494 (4)	C21—N22	1.330 (3)
O3—C3	1.200 (3)	C21—C26	1.384 (4)
C3—N3	1.446 (4)	N22—C23	1.327 (3)
C3—C11	1.488 (4)	C23—C24	1.385 (4)
N3—C4	1.389 (4)	C23—C4 ⁱ	1.499 (4)
N3—C21A	1.416 (4)	C24—C25	1.379 (5)
O4—C4	1.203 (3)	C24—H24	0.9300
C4—C23 ⁱ	1.499 (4)	C25—C26	1.365 (5)

C11—N12	1.333 (3)	C25—H25	0.9300
C11—C16	1.374 (4)	C26—H26	0.9300
N12—C13	1.335 (3)	C21A—N26A	1.322 (4)
C13—C14	1.378 (4)	C21A—N22A	1.326 (4)
C14—C15	1.368 (4)	N22A—C23A	1.338 (4)
C14—H14	0.9300	C23A—O23A	1.343 (4)
C15—C16	1.373 (4)	C23A—C24A	1.370 (5)
C15—H15	0.9300	C24A—C25A	1.373 (5)
C16—H16	0.9300	C24A—H24A	0.9300
C11A—N12A	1.319 (4)	C25A—O25A	1.338 (4)
C11A—N16A	1.331 (4)	C25A—N26A	1.342 (4)
N12A—C13A	1.339 (4)	O23A—C27A	1.441 (4)
C13A—O13A	1.334 (4)	C27A—H27A	0.9600
C13A—C14A	1.378 (5)	C27A—H27B	0.9600
C14A—C15A	1.367 (5)	C27A—H27C	0.9600
C14A—H14A	0.9300	O25A—C28A	1.438 (5)
C15A—N16A	1.332 (4)	C28A—H28A	0.9600
C15A—O15A	1.351 (4)	C28A—H28B	0.9600
O13A—C17A	1.418 (5)	C28A—H28C	0.9600
C17A—H17A	0.9600	O1W—H1W	0.850 (5)
O1—C1—N1	123.2 (3)	H17A—C17A—H17C	109.5
O1—C1—C13	121.9 (3)	H17B—C17A—H17C	109.5
N1—C1—C13	114.8 (2)	C15A—O15A—C18A	119.0 (3)
C1—N1—C11A	122.3 (2)	O15A—C18A—H18A	109.5
C1—N1—C2	118.7 (2)	O15A—C18A—H18B	109.5
C11A—N1—C2	118.4 (2)	H18A—C18A—H18B	109.5
O2—C2—N1	120.1 (3)	O15A—C18A—H18C	109.5
O2—C2—C21	123.0 (3)	H18A—C18A—H18C	109.5
N1—C2—C21	116.8 (2)	H18B—C18A—H18C	109.5
O3—C3—N3	120.9 (3)	N22—C21—C26	122.1 (3)
O3—C3—C11	123.8 (3)	N22—C21—C2	115.5 (2)
N3—C3—C11	115.2 (3)	C26—C21—C2	122.3 (3)
C4—N3—C21A	121.2 (2)	C23—N22—C21	117.9 (2)
C4—N3—C3	122.4 (2)	N22—C23—C24	123.6 (3)
C21A—N3—C3	115.8 (2)	N22—C23—C4 ⁱ	117.7 (2)
O4—C4—N3	122.3 (3)	C24—C23—C4 ⁱ	118.7 (3)
O4—C4—C23 ⁱ	121.4 (3)	C25—C24—C23	117.7 (3)
N3—C4—C23 ⁱ	116.1 (2)	C25—C24—H24	121.2
N12—C11—C16	123.4 (3)	C23—C24—H24	121.2
N12—C11—C3	116.7 (2)	C26—C25—C24	119.2 (3)
C16—C11—C3	119.9 (3)	C26—C25—H25	120.4
C11—N12—C13	116.9 (2)	C24—C25—H25	120.4
N12—C13—C14	123.5 (3)	C25—C26—C21	119.3 (3)
N12—C13—C1	115.4 (2)	C25—C26—H26	120.3
C14—C13—C1	121.0 (3)	C21—C26—H26	120.3
C15—C14—C13	118.4 (3)	N26A—C21A—N22A	128.8 (3)
C15—C14—H14	120.8	N26A—C21A—N3	114.0 (3)

C13—C14—H14	120.8	N22A—C21A—N3	117.2 (3)
C14—C15—C16	119.3 (3)	C21A—N22A—C23A	114.3 (3)
C14—C15—H15	120.4	N22A—C23A—O23A	118.3 (3)
C16—C15—H15	120.4	N22A—C23A—C24A	123.4 (3)
C15—C16—C11	118.5 (3)	O23A—C23A—C24A	118.3 (3)
C15—C16—H16	120.7	C23A—C24A—C25A	116.0 (3)
C11—C16—H16	120.7	C23A—C24A—H24A	122.0
N12A—C11A—N16A	129.5 (3)	C25A—C24A—H24A	122.0
N12A—C11A—N1	114.3 (3)	O25A—C25A—N26A	118.5 (3)
N16A—C11A—N1	116.0 (3)	O25A—C25A—C24A	118.2 (3)
C11A—N12A—C13A	114.2 (3)	N26A—C25A—C24A	123.3 (3)
O13A—C13A—N12A	118.5 (4)	C21A—N26A—C25A	114.2 (3)
O13A—C13A—C14A	119.1 (3)	C23A—O23A—C27A	118.7 (3)
N12A—C13A—C14A	122.4 (4)	O23A—C27A—H27A	109.5
C15A—C14A—C13A	116.6 (3)	O23A—C27A—H27B	109.5
C15A—C14A—H14A	121.7	H27A—C27A—H27B	109.5
C13A—C14A—H14A	121.7	O23A—C27A—H27C	109.5
N16A—C15A—O15A	118.3 (4)	H27A—C27A—H27C	109.5
N16A—C15A—C14A	123.6 (3)	H27B—C27A—H27C	109.5
O15A—C15A—C14A	118.1 (3)	C25A—O25A—C28A	117.5 (3)
C11A—N16A—C15A	113.5 (3)	O25A—C28A—H28A	109.5
C13A—O13A—C17A	117.9 (3)	O25A—C28A—H28B	109.5
O13A—C17A—H17A	109.5	H28A—C28A—H28B	109.5
O13A—C17A—H17B	109.5	O25A—C28A—H28C	109.5
H17A—C17A—H17B	109.5	H28A—C28A—H28C	109.5
O13A—C17A—H17C	109.5	H28B—C28A—H28C	109.5
O1—C1—N1—C11A	26.2 (4)	C13A—C14A—C15A—N16A	0.4 (7)
C13—C1—N1—C11A	-150.1 (3)	C13A—C14A—C15A—O15A	178.3 (4)
O1—C1—N1—C2	-145.4 (3)	N12A—C11A—N16A—C15A	0.1 (5)
C13—C1—N1—C2	38.4 (3)	N1—C11A—N16A—C15A	175.5 (3)
C1—N1—C2—O2	35.9 (4)	O15A—C15A—N16A—C11A	-179.4 (3)
C11A—N1—C2—O2	-136.0 (3)	C14A—C15A—N16A—C11A	-1.5 (6)
C1—N1—C2—C21	-142.4 (3)	N12A—C13A—O13A—C17A	-10.2 (6)
C11A—N1—C2—C21	45.7 (3)	C14A—C13A—O13A—C17A	169.5 (5)
O3—C3—N3—C4	51.6 (4)	N16A—C15A—O15A—C18A	0.0 (6)
C11—C3—N3—C4	-131.0 (3)	C14A—C15A—O15A—C18A	-178.0 (4)
O3—C3—N3—C21A	-119.2 (3)	O2—C2—C21—N22	23.0 (4)
C11—C3—N3—C21A	58.3 (3)	N1—C2—C21—N22	-158.8 (2)
C21A—N3—C4—O4	17.8 (5)	O2—C2—C21—C26	-155.0 (3)
C3—N3—C4—O4	-152.5 (3)	N1—C2—C21—C26	23.2 (4)
C21A—N3—C4—C23 ⁱ	-158.6 (3)	C26—C21—N22—C23	-3.9 (4)
C3—N3—C4—C23 ⁱ	31.1 (4)	C2—C21—N22—C23	178.0 (2)
O3—C3—C11—N12	-165.0 (3)	C21—N22—C23—C24	0.3 (5)
N3—C3—C11—N12	17.6 (4)	C21—N22—C23—C4 ⁱ	177.0 (3)
O3—C3—C11—C16	13.7 (4)	N22—C23—C24—C25	2.7 (5)
N3—C3—C11—C16	-163.7 (3)	C4 ⁱ —C23—C24—C25	-174.0 (3)
C16—C11—N12—C13	1.3 (4)	C23—C24—C25—C26	-2.1 (6)

C3—C11—N12—C13	179.9 (2)	C24—C25—C26—C21	-1.3 (5)
C11—N12—C13—C14	0.0 (4)	N22—C21—C26—C25	4.5 (5)
C11—N12—C13—C1	176.3 (2)	C2—C21—C26—C25	-177.6 (3)
O1—C1—C13—N12	-141.8 (3)	C4—N3—C21A—N26A	-143.9 (3)
N1—C1—C13—N12	34.4 (3)	C3—N3—C21A—N26A	27.0 (3)
O1—C1—C13—C14	34.5 (4)	C4—N3—C21A—N22A	38.0 (4)
N1—C1—C13—C14	-149.2 (3)	C3—N3—C21A—N22A	-151.2 (2)
N12—C13—C14—C15	-1.1 (5)	N26A—C21A—N22A—C23A	0.1 (4)
C1—C13—C14—C15	-177.1 (3)	N3—C21A—N22A—C23A	178.0 (3)
C13—C14—C15—C16	0.8 (5)	C21A—N22A—C23A—O23A	178.8 (3)
C14—C15—C16—C11	0.4 (5)	C21A—N22A—C23A—C24A	-2.4 (5)
N12—C11—C16—C15	-1.5 (5)	N22A—C23A—C24A—C25A	2.4 (5)
C3—C11—C16—C15	179.9 (3)	O23A—C23A—C24A—C25A	-178.8 (3)
C1—N1—C11A—N12A	-144.5 (3)	C23A—C24A—C25A—O25A	179.7 (3)
C2—N1—C11A—N12A	27.1 (4)	C23A—C24A—C25A—N26A	-0.3 (5)
C1—N1—C11A—N16A	39.5 (4)	N22A—C21A—N26A—C25A	1.8 (4)
C2—N1—C11A—N16A	-148.9 (3)	N3—C21A—N26A—C25A	-176.1 (2)
N16A—C11A—N12A—C13A	2.3 (5)	O25A—C25A—N26A—C21A	178.3 (3)
N1—C11A—N12A—C13A	-173.1 (3)	C24A—C25A—N26A—C21A	-1.7 (5)
C11A—N12A—C13A—O13A	176.2 (4)	N22A—C23A—O23A—C27A	-1.2 (5)
C11A—N12A—C13A—C14A	-3.5 (6)	C24A—C23A—O23A—C27A	179.9 (3)
O13A—C13A—C14A—C15A	-177.3 (4)	N26A—C25A—O25A—C28A	1.3 (5)
N12A—C13A—C14A—C15A	2.3 (7)	C24A—C25A—O25A—C28A	-178.7 (3)

Symmetry code: (i) $-x, y, -z+1/2$.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C15—H15 \cdots O3 ⁱⁱ	0.93	2.47	3.148 (4)	130
C15—H15 \cdots O4 ⁱⁱⁱ	0.93	2.42	3.084 (4)	129
C18A—H18C \cdots O23A	0.96	2.47	3.393 (6)	160
C24—H24 \cdots O1 ^{iv}	0.93	2.49	3.14 (2)	127
C26—H26 \cdots N22A	0.93	2.62	3.325 (4)	133
C27A—H27B \cdots O23A ^v	0.96	2.57	3.144 (4)	119

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