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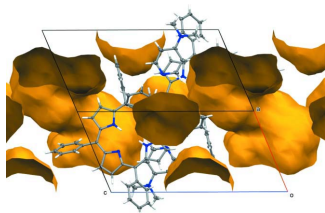
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Crystal structure of *cis*-7,8-dihydroxy-5,10,15,20-tetraphenylchlorin and its zinc(II)–ethylenediamine complex

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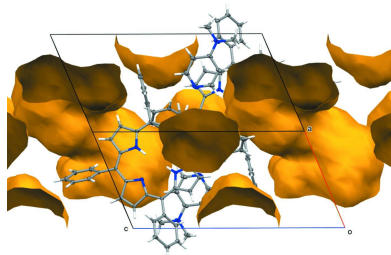
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The title chlorin, **2^{Ph}H₂**, hydrogen-bonded to dimethylaminopyridine (DMAP), C₄₄H₃₂N₄O₂·C₇H₁₀N₂, and its corresponding zinc(II) complex, **2^{Ph}Zn**, axially coordinated to ethylenediamine (EDA), [Zn(C₄₄H₃₀N₄O₂)]·C₂H₈N₂, were isolated and crystallized by adventitious reduction of the corresponding osmate esters by DMAP and EDA, respectively. Known since 1996 and, *inter alia*, used for the preparation of a wide range of (planar and non-planar) chlorin analogues (so-called pyrrole-modified porphyrins), their conformational analyses in the solid state are important benchmarks. Both macrocycles are only modestly distorted from planarity and both are slightly more non-planar than the corresponding dimethoxy-derivative, but less planar than a free-base *meso*-pentafluorophenyl-based osmate ester. NSD analyses provide quantitative and qualitative analyses of the distortion modes. One origin of the non-planarity is presumably the avoidance of the eclipsed configuration of the two *vic-cis* diols on the pyrroline moiety; the resulting deformation of the pyrroline translates in some cases into the macrocycle. The structure of **2^{Ph}H₂** features voids making up *ca* 26% of the unit-cell volume filled with highly disordered solvate molecules (chloroform and hexanes). **2^{Ph}Zn** crystallized with a 13.6 (4)% occupied solvate methanol molecule.

1. Chemical context

The study of synthetic chlorins as functional, spectroscopic, or structural models for nature's premiere light-harvesting pigment chlorophyll is one of the central aspects in contemporary porphyrinoid chemistry (Flitsch, 1988; Liu *et al.*, 2018; Taniguchi & Lindsey, 2017; Lindsey, 2015). Because of the facility of the synthesis of a wide range of *meso*-tetraarylporphyrins, their conversion to chlorins has been widely studied (Flitsch, 1988; Taniguchi & Lindsey, 2017).

We contributed to the field the description of the OsO₄-mediated dihydroxylation of *meso*-tetraarylporphyrins **1^{Ar}M**, generating the corresponding chlorin diols **2^{Ar}M** (Fig. 1) (Brückner & Dolphin, 1995*a*; Brückner *et al.*, 1998). Depending on the stoichiometric ratio of OsO₄ used and whether the porphyrin metal complex or free base is used, the reaction may also lead to the regioselective formation of tetrahydroxymetalloisobacteriochlorins or tetrahydroxybacteriochlorins, respectively (Brückner & Dolphin, 1995*b*; Samankumara *et al.*, 2010; Hyland *et al.*, 2012; Bruhn & Brückner, 2015). Chlorin diols **2^{Ar}H₂** have shown efficacy as photosensitizers in photodynamic therapy (Macalpine *et al.*, 2002) or are substrates toward their oxidation to the corresponding diones (Starnes *et al.*, 2000, 2001; Daniell *et al.*, 2003).



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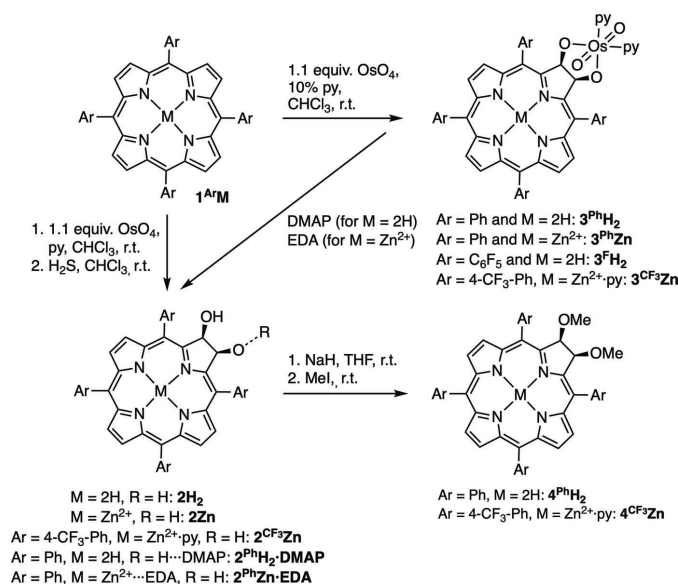


Figure 1
Synthetic pathways towards $2^{\text{Ph}}\text{H}_2\cdot\text{DMAP}$ and $2^{\text{Ph}}\text{Zn}\cdot\text{EDA}$ and their methoxy ethers.

Importantly, chlorin diols 2^{Ar}H_2 are the starting materials for the generation of a wide range of planar and non-planar chlorin analogues (so-called pyrrole-modified porphyrins) (Brückner, 2016; Sharma *et al.*, 2017; Hewage *et al.*, 2019; Brückner *et al.*, 2020; Luciano *et al.*, 2020; Wu *et al.*, 2020), whereby the parent chlorin diols 2^{Ph}H_2 and 2^{Ph}Zn generally serve as spectroscopic benchmarks. Since the conformation of a porphyrinic macrocycle greatly influences its electronic structure, the structural characterization of the benchmark compounds 2^{Ph}H_2 and 2^{Ph}Zn is important. Curiously, however, even though these fundamental compounds are known since 1996, crystals suitable for single X-ray crystal structure analyses could not be grown to date. However, related derivatives, such as osmate ester 3^{F}H_2 (Hewage *et al.*, 2019), a number of tetrahydroxybacteriochlorins and isobacteriochlorins (Samankumara *et al.*, 2010), and a number of

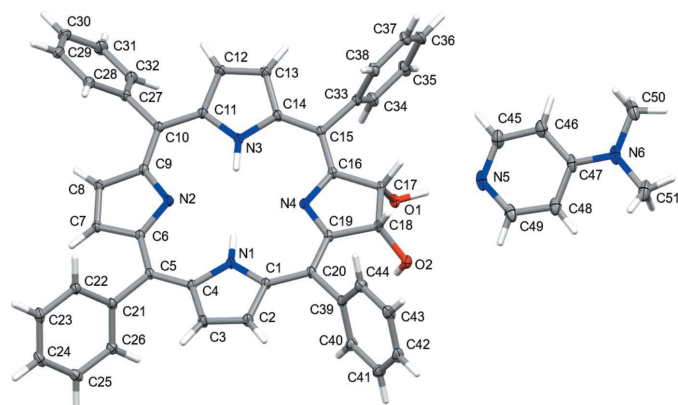
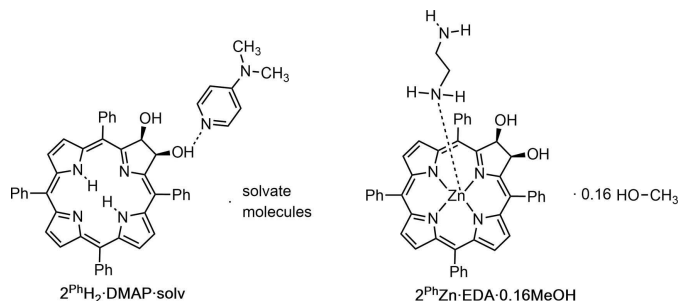


Figure 2
X-ray structure of $2^{\text{Ph}}\text{H}_2\cdot\text{DMAP}$ with the atom-labeling scheme for non-H atoms. 50% probability ellipsoids.

alkylated diol free base and metal complexes 4^{Ar}M (M = 2H, Ni, Cu, Zn, Pd) (Samankumara *et al.*, 2010; Sharma *et al.*, 2017) could be structurally characterized.



In due course of working with the intermediate osmate esters and attempts to form crystals of the amine adducts, we inadvertently reduced the osmate ester and the long-sought parent free base *meso*-phenyl chlorin diol 2^{Ph}H_2 , as $2^{\text{Ph}}\text{H}_2\cdot\text{DMAP}$ hydrogen-bonded to DMAP (4-dimethylaminopyridine) and the zinc(II) complex 2^{Ph}Zn , in the form $2^{\text{Ph}}\text{Zn}\cdot\text{EDA}$ in which the metal is axially coordinated to ethylenediamine (EDA), crystallized in single-crystal X-ray diffraction quality.

2. Structural commentary

The structures of both $2^{\text{Ph}}\text{H}_2\cdot\text{DMAP}$ and $2^{\text{Ph}}\text{Zn}\cdot\text{EDA}$ confirm the *cis-vic* stereochemistry of the diol functionality and the near-perpendicular arrangement of the *meso*-phenyl groups – structural features well known for these types of *meso*-arylchlorin diols (Hewage *et al.*, 2019; Samankumara *et al.*, 2010; Sharma *et al.*, 2017) or *meso*-arylporphyrinoids, in general (Senge, 2000) (Figs. 2 and 3).

Importantly, the structures allow the determination of the conformation of their chromophores. The dissection of the

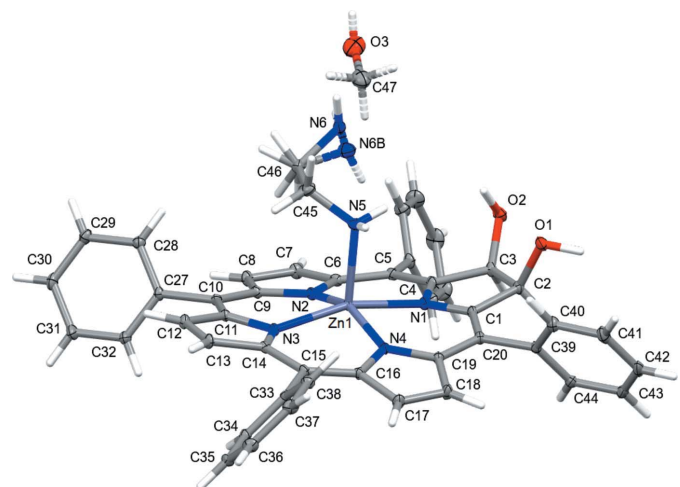


Figure 3
X-ray structure of the zinc(II) complex $2^{\text{Ph}}\text{Zn}\cdot\text{EDA}$, with the atom-labeling scheme for non-H atoms. 50% probability ellipsoids. Dashed bonds indicate the minor disordered amine [11.8 (12)% occupancy], and the partially occupied MeOH solvate [13.6 (4)% occupancy]. Atom labels for the backwards pointing phenyl ring (C21–C26) are omitted for clarity.

conformation of **2^{Ph}H₂·DMAP** using a normal mode structural decomposition (NSD) analysis (Kingsbury & Senge, 2021; Shelnutt *et al.*, 1998) shows that its chromophore exhibits a considerable saddling distortion. In comparison, the dimethoxy derivative **4^{Ph}H₂** (Samankumara *et al.*, 2010) is more planar, with only very modest distortions evenly spread over a number of distortion modes (Fig. 4a). In **4^{Ph}H₂**, both methoxy substituents point toward the outside, whereas the corresponding hydroxy groups in **2^{Ph}H₂·DMAP** point in opposite directions, with only the hydrogen-bonded (to DMAP) hydroxy group pointing outwards. A slight deformation of the pyrroline moiety in **2^{Ph}H₂·DMAP** alleviates the steric interactions between the two hydroxy groups [26.65 (13)° O—C—C—O torsion angle] that would be otherwise forced to be eclipsed. The corresponding torsion angle in **4^{Ph}H₂** is slightly smaller [17.23 (17)°; Samankumara *et al.*, 2010]. This *vic-cis*-substituents-induced pyrroline deformation was also observed previously (Sharma *et al.*, 2017; Hewage *et al.*, 2019).

The out-of-plane plots (Kingsbury & Senge, 2021) of the two free-base chlorins **2^{Ph}H₂·DMAP** and **4^{Ph}H₂** also illustrate the qualitative and quantitative differences in the conformations of the two (Fig. 5a).

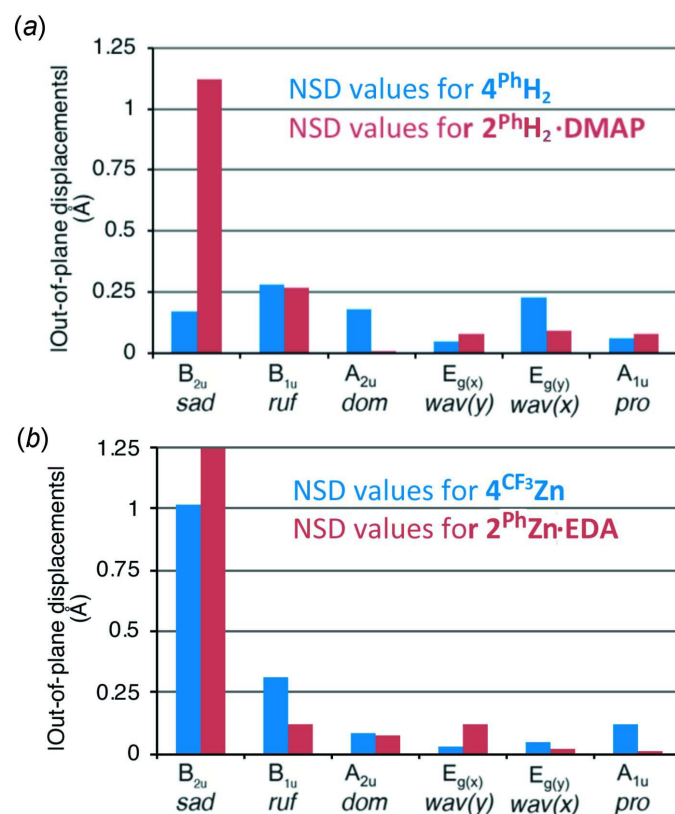


Figure 4
Normal mode Structural Decomposition (NSD) analysis (Kingsbury & Senge, 2021) of (a), the chromophore conformations of dihydroxychlorin **2^{Ph}H₂·DMAP** (hydrogen-bonded to DMAP) in comparison to the conformation of the chromophore of dimethoxychlorin **4^{Ph}H₂** (Samankumara *et al.*, 2010), and (b), the equivalent chromophore conformation analysis of **2^{Ph}Zn-EDA** in comparison to the closely related dimethoxy derivative **4^{CF3}Zn** (Sharma *et al.*, 2017).

The saddling deformation is more pronounced in the corresponding zinc(II) complexes but the deformation modes observed in either of the complexes are very similar (Fig. 4b and 5b). This (small) B_{2u} deformation mode is typical for penta-coordinated, square-pyramidal porphyrinoid zinc(II) complexes (Kingsbury & Senge, 2021). The differences in conformation quality and quantity is only minimal between the parent compound **2^{Ph}Zn-EDA** and its *p*-aryl-substituted and methylated analogue **4^{CF3}Zn-py**. In addition, both molecules carry their axial ligand on the same hemisphere defined by the macrocycle the diol/dimethoxy moieties are located. Nonetheless, there are differences. For instance, a smaller O—C—C—O torsion angle was observed in the diol zinc complex **2^{Ph}Zn-EDA** [O—C_β—C_β—O dihedral angle = 7.86 (17)°], whereas the corresponding angle in the dimethoxy derivative **4^{CF3}Zn** is much larger at 28.1 (4)° (Sharma *et al.*, 2017).

In neither the free base nor the zinc complex of the diol chlorins are any significant in-plane deformations observed. The change in the macrocycle conformation upon methylation and/or hydrogen bonding to an amine acceptor reiterates the conformational malleability of the chlorin chromophore (Kratky *et al.*, 1985), as previously also shown in the varying conformations of a range of transition-metal complexes (Sharma *et al.*, 2017).

3. Supramolecular features

The dominant supramolecular interactions in both **2^{Ph}H₂·DMAP** and **2^{Ph}Zn-EDA** are hydrogen-bonding inter-

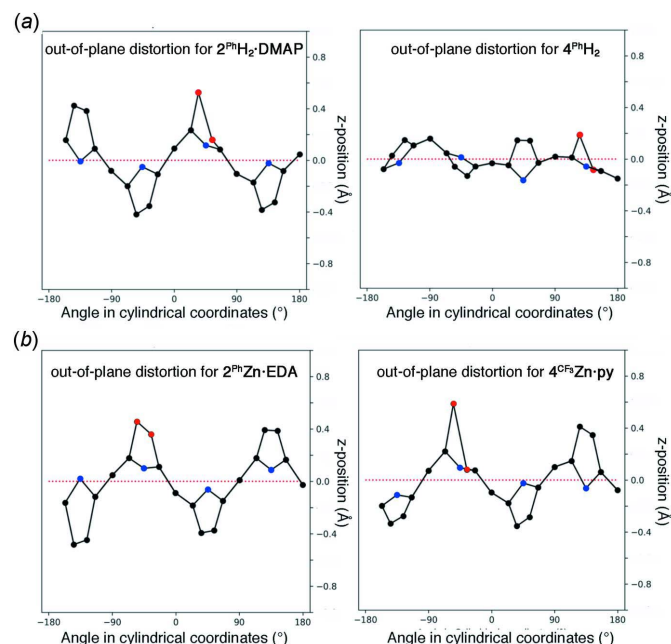


Figure 5
Out-of-plane plots (Kingsbury & Senge, 2021) of the chromophore conformations of (a), dihydroxychlorin **2^{Ph}H₂·DMAP** and dimethoxychlorin **4^{Ph}H₂** (Samankumara *et al.*, 2010), and (b), the equivalent plots of **2^{Ph}Zn-EDA** and **4^{CF3}Zn-py** (Sharma *et al.*, 2017). The atoms indicated in red are the pyrroline β -carbons carrying the *cis*-hydroxy or methoxy groups.

Table 1
Hydrogen-bond geometry (Å, °) for **2^{Ph}H₂**.

<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—H1O...N5	0.973 (17)	1.727 (17)	2.6968 (14)	174.1 (14)
O2—H2O...O1 ⁱ	0.927 (17)	1.882 (17)	2.7798 (12)	162.5 (14)
N1—H1N...N2	0.925 (15)	2.346 (15)	2.9064 (13)	118.7 (11)
N1—H1N...N4	0.925 (15)	2.383 (15)	2.9518 (13)	119.6 (11)
N3—H3N...N2	0.915 (16)	2.292 (16)	2.8868 (13)	122.3 (12)
N3—H3N...N4	0.915 (16)	2.458 (15)	2.9766 (14)	116.1 (12)
C37—H37...O2 ⁱⁱ	0.95	2.51	3.3840 (16)	153
C38—H38...C48 ⁱⁱ	0.95	2.77	3.6779 (19)	161
C50—H50B...N4 ⁱⁱ	0.98	2.57	3.544 (2)	171

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

actions between the hydroxyl functions of the chlorin molecules, and the DMAP and EDA bases incorporated into the crystal structure.

In **2^{Ph}H₂·DMAP** one of the hydroxyl groups acts as a donor towards the DMAP with O1—H1O...N5 = 2.6968 (14) Å. O1 in turn acts as acceptor for an O—H...O bond originating from O2 of a neighboring molecule. A symmetry-equivalent interaction (by inversion) connects the other two oxygen atoms of the same two molecules with each other, creating an inversion-symmetric dimer (Fig. 6). A number of additional interactions that augment the strong hydrogen bonds, among them C—H...O, C—H...N and C—H... π interactions, are listed in the hydrogen-bonding Table 1.

The structure of **2^{Ph}H₂·DMAP** also contains 647 Å³ (ca 26% of the unit-cell volume) of solvent-accessible voids occupied by highly disordered solvent molecules that could not be properly modeled or refined (Fig. 7). The content of these voids, presumably chloroform and hexane, the crystallization solvents, were instead included in the model *via* reverse-Fourier-transform methods using the SQUEEZE routine (van der Sluis & Spek, 1990; Spek, 2015) as imple-

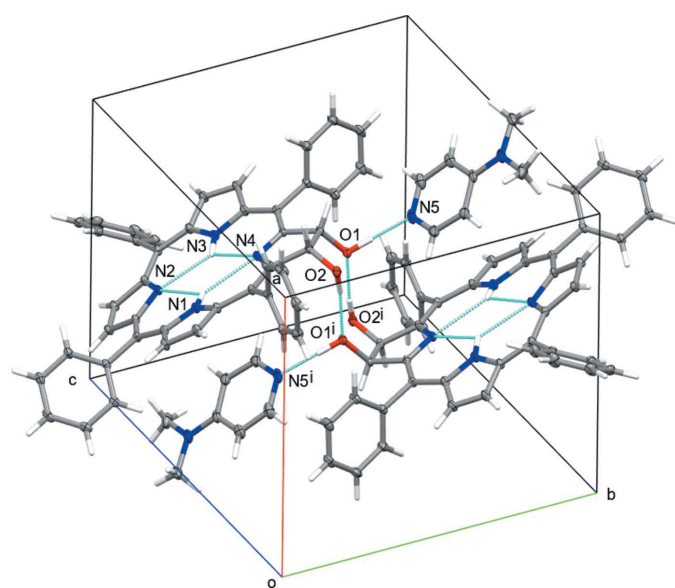


Figure 6
Hydrogen bonding and packing of **2^{Ph}H₂·DMAP**. 50% probability ellipsoids. Symmetry code: (i) $1 - x, 1 - y, 1 - z$.

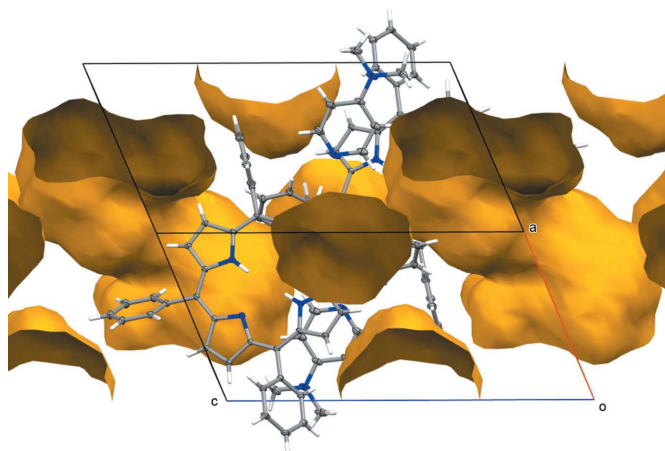


Figure 7
Solvent-accessible voids in **2^{Ph}H₂·DMAP**. The void volume is 647 Å³, or ca 26% of the unit-cell volume.

mented in the program *PLATON* (Spek, 2020), and added as additional not-model-based structure-factor contributions. The procedure corrected for 162 electrons within the solvent-accessible voids.

Hydrogen bonding in **2^{Ph}Zn·EDA** is similar to that of **2^{Ph}H₂·DMAP**, but more complex. In contrast to the DMAP molecule in **2^{Ph}H₂·DMAP**, the amino NH₂ groups of the ethylene diamine in **2^{Ph}Zn·EDA** can act as both hydrogen-bond acceptors as well as hydrogen-bond donors. One of the two amine moieties of the EDA base is axially coordinated to the zinc center of the chlorin complex, and is thus not available as a hydrogen-bond acceptor. The partially occupied methanol molecule also takes part in hydrogen-bonding interactions, and the disorder of the not-metal-coordinated amino group further complicates the hydrogen-bonding network of **2^{Ph}Zn·EDA**.

The two hydroxyl groups again both act as hydrogen-bond donors, and similar to in **2^{Ph}H₂·DMAP** they form an inversion-symmetric dimer (Fig. 8). O1 again acts as a hydrogen-bond donor towards the base, here the disordered amino group, of the other molecule of the dimer. Different from the DMAP molecule, which lacks acidic H atoms, the amines also act as hydrogen-bond donors. The metal-coordinated amine

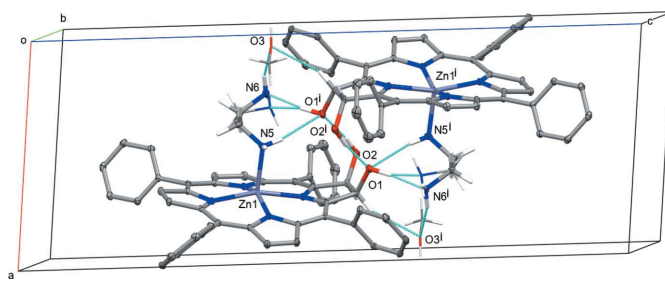


Figure 8
Hydrogen bonding and packing of **2^{Ph}Zn·EDA**. 50% probability ellipsoids. Symmetry code: (i) $1 - x, 1 - y, 1 - z$. 50% ellipsoids for fully occupied and major occupancy non-H atoms. Others in capped stick mode. Phenyl and pyrrole H atoms are omitted for clarity.

Table 2
Hydrogen-bond geometry (Å, °) for **2^{Ph}Zn**.

<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—H1...N6 ⁱ	0.99	1.73	2.710 (3)	168
O1—H1...N6B ⁱ	0.99	1.54	2.510 (17)	165
O2—H2A...O1 ⁱ	0.99	1.82	2.8056 (18)	171
C2—H2...O3 ⁱ	1.00	2.53	3.460 (14)	155
N5—H5A...O1 ⁱ	0.88 (2)	2.38 (2)	3.2442 (18)	166 (2)
C46—H46A...N2	0.99	2.49	3.368 (2)	148
N6—H6A...O3	0.90 (2)	2.08 (2)	2.932 (14)	159 (3)
C46B—H46C...N2	0.99	2.68	3.368 (2)	126
O3—H3O...N4 ⁱⁱ	0.84	2.20	2.992 (14)	157

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

creates an N—H...O bond that provides an additional connection within the dimer to create a 3D hydrogen-bonding network between the two molecules (Fig. 8).

Several 'terminal' hydrogen bonds or hydrogen-bond-like interactions cap off the not yet used acidic and basic atoms, which are listed in the hydrogen-bonding Table 2 (interactions not shown). The second amine H atom of the metal-coordinated NH₂ group is engaged in an N—H... π interaction towards the π -density of C29 of the phenyl ring of a neighboring molecule. The major moiety of the disordered amino group hydrogen bonds with the partially occupied methanol molecule. However, this interaction is not always present, as the occupancy of the MeOH molecule is only 13.6 (4)%, while that of the amino group is 88.2 (12)%. The second amino H atom is not involved in any directional interactions. One of the H atoms of the minor amino moiety might be engaged in another N—H... π interaction towards the π -density of C43 and C43 of a phenyl ring of the second dimer molecule, but the exact positions of the amino H atoms are not determined accurately given the low occupancy of the amino fragment [11.8 (12)%]. The same is true for the position of the methanol hydroxyl H atom, which appears to be engaged in a weak O—H... π interaction with the porphyrinic π -system of a molecule at $-1+x, y, z$. O3, the methanol oxygen atom, acts as acceptor for a C—H...O interaction originating from a phenyl C atom of a molecule not part of the dimer. The H...O distance is unusually short for a C—H...O interaction, 2.53 Å, which could be an artifact of the low occupancy of the methanol molecule.

4. Database survey

A search of the Cambridge Structural Database (CSD Version 5.43, Nov 2021; Groom *et al.*, 2016) for *meso*-tetraarylchlorins or their metal(II) complexes revealed in excess of 75 structures, but few are directly comparable to the title compounds: Most examples contain a variety of bulky substituents or annulated rings at the pyrroline positions [the closest being an imidazolone-annulated dihydroxychlorin, TAKDUI (Luciano *et al.* 2020)] or contain other (sterically encumbering) substituents at the pyrrolic β -positions or on the *meso*-aryl groups. Most metallochlorins contain also a different metal than zinc(II). Only a few compounds are structurally closely related

to **2^{Ph}H₂-DMAP** or **2^{Ph}Zn-EDA**. Among them is the parent non-hydroxylated chlorin zinc chelate [5,10,15,20-tetraphenylchlorinato]zinc(II)-pyridine complex (HPORZN10; Spaulding *et al.*, 1977), the bis- β -*n*-butylated free base and zinc(II) chlorins (QAKLUJ and QAKMAQ, respectively; Senge *et al.*, 2000), free base 5,10,15,20-tetraphenyl-7-hydroxychlorin (SAZSAP; Samankumara *et al.*, 2010), the β -nitrated analogue of **2^{Ph}H₂** (TIPBIF; Worlinsky *et al.*, 2013), dimethoxy derivatives **4^{Ph}H₂** (SAZROC; Samankumara *et al.*, 2010) and **4^{CF₃}Zn-py** (PEDKER; Sharma *et al.*, 2017), osmate ester **3^FH₂** (SIZFUF; Hewage *et al.*, 2019), and *trans*-7,8-diol-7,8-dimethyltetraphenylchlorin (ZAZNIZ; Banerjee *et al.*, 2012).

5. Synthesis and crystallization

The OsO₄-mediated dihydroxylation of porphyrin **1H₂** is a two-step sequence: the formation of the osmate ester **3^AH₂** in the first step is followed by the reduction of the osmate ester to the target dihydroxychlorin **2^AH₂** (often performed as a two-step, one-pot process) (Brückner & Dolphin, 1995b; Samankumara *et al.*, 2010; Hyland *et al.*, 2012). Here, we prepared the intermediate *meso*-tetraphenyl-2,3-*vic*-dihydroxychlorin osmate ester according to the established oxidation of *meso*-tetraphenylporphyrins **1^{Ph}H₂** (Brückner *et al.*, 1998). Metalation of the free base **1^{Ph}H₂** using Zn(OAc)₂·2H₂O under standard conditions (Buchler, 1978) (refluxing CHCl₃/MeOH for 35–40 min) formed the corresponding Zn^{II} osmate ester **3^{Ph}Zn**.

While crystallizing the osmate esters in CH₂Cl₂ and layering with the non-solvent hexane in the presence of DMAP (for **3^{Ph}H₂**) or by allowing a solution of the ester in CH₂Cl₂/MeOH to slowly evaporate in the presence of EDA (for **3^{Ph}Zn**), both osmate esters adventitiously reduced and diols **2^{Ph}H₂-DMAP** and **2^{Ph}Zn-EDA** crystallized, respectively. The spectroscopic data of both known chromophores are as described previously (Brückner *et al.*, 1998).

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. C—H bond distances were constrained to 0.95 Å for aromatic and alkene C—H groups, and to 1.00, 0.99 and 0.98 Å for aliphatic C—H, CH₂ and CH₃ groups, respectively. Positions of N—H and NH₂ hydrogen atoms were refined. N—H distances within NH₂ groups in **2^{Ph}Zn-EDA** were restrained to 0.88 (2) Å and H—N—H and H—N—C angles were restrained to be similar to each other. Methyl CH₃ and hydroxyl H atoms were allowed to rotate but not to tip to best fit the experimental electron density. The hydroxyl H atom of the partially occupied methanol molecule in **2^{Ph}Zn-EDA** was restrained to hydrogen bond to a porphyrin N atom of a neighboring complex. *U*_{iso}(H) values were set to a multiple of *U*_{eq}(C/O/N) with 1.5 for CH₃ and OH, and 1.2 for C—H, CH₂, N—H and NH₂ units, respectively.

In the structure of **2^{Ph}Zn-EDA**, disorder of the not-metal-coordinated amino group of the ethylene diamine molecule is

Table 3
Experimental details.

	2^{Pb}H₂	2^{Pb}Zn
Crystal data		
Chemical formula	C ₄₄ H ₃₂ N ₄ O ₂ ·C ₇ H ₁₀ N ₂ ·[+solvent]	[Zn(C ₄₄ H ₃₀ N ₄ O ₂)]·C ₂ H ₈ N ₂ ·0.136CH ₄ O
<i>M_r</i>	770.90	776.57
Crystal system, space group	Triclinic, <i>P</i> $\bar{1}$	Monoclinic, <i>P</i> ₂ /c
Temperature (K)	150	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	10.0193 (4), 15.2554 (8), 17.7983 (10)	10.1249 (3), 13.5400 (4), 27.0447 (8)
α , β , γ (°)	69.918 (2), 74.926 (2), 84.140 (2)	90, 95.1464 (11), 90
<i>V</i> (Å ³)	2466.9 (2)	3692.64 (19)
<i>Z</i>	2	4
Radiation type	Mo <i>K</i> α	Cu <i>K</i> α
μ (mm ⁻¹)	0.06	1.32
Crystal size (mm)	0.33 × 0.21 × 0.19	0.27 × 0.25 × 0.18
Data collection		
Diffractometer	Bruker AXS D8 Quest diffractometer with PhotonII charge-integrating pixel array detector (CPAD)	Bruker AXS D8 Quest diffractometer with PhotonIII-C14 charge-integrating and photon counting pixel array detector
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)
<i>T</i> _{min} , <i>T</i> _{max}	0.665, 0.746	0.606, 0.754
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	48645, 14738, 9891	21319, 7551, 7037
<i>R</i> _{int}	0.060	0.024
(sin θ /λ) _{max} (Å ⁻¹)	0.714	0.638
Refinement		
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.048, 0.133, 1.04	0.031, 0.088, 1.04
No. of reflections	14738	7551
No. of parameters	549	549
No. of restraints	0	17
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³)	0.45, -0.21	0.31, -0.44

Computer programs: *APEX4* (Bruker, 2021), *APEX3* and *SAINT* (Bruker, 2019), *SHELXT* (Sheldrick, 2015a), *SHELXS97* (Sheldrick, 2008), *SHELXL2018/3* (Sheldrick, 2015b), *ShelXle* (Hübschle *et al.*, 2011), *Mercury* (Macrae *et al.*, 2020), and *publCIF* (Westrip, 2010).

observed and a methanol solvate molecule is partially occupied. The C—N bonds were restrained to be similar in length. A partially occupied methanol molecule is located nearby the major disordered amino group and hydrogen-bonded to it. The hydroxyl H atom was restrained to hydrogen bond to a porphyrin N atom of a neighboring complex. Subject to these conditions, the occupancy ratio for the amino groups refined to 0.882 (12): 0.118 (12), and the occupancy rate for the methanol molecule refined to 0.136 (4). The occupancy of the methanol molecule is not correlated with the disorder of the amino group (the major 88% occupied amino group is hydrogen-bonded to the 14% occupied methanol molecule).

The structure of **2^{Pb}H₂·DMAP** contains 647 Å³ of solvent-accessible voids occupied by highly disordered solvate molecules (presumably chloroform and hexane, the crystallization solvents). The residual electron-density peaks are not arranged in an interpretable pattern and no unambiguous disorder model could be developed. The structure factors were instead augmented *via* reverse-Fourier-transform methods using the SQUEEZE routine (van Sluis & Spek, 1990; Spek, 2015), as implemented in the program *PLATON* (Spek, 2020). The resultant .fab file containing the structure-factor contribution from the electron content of the void space was used in together with the original hkl file in the further

refinement. The SQUEEZE procedure accounted for 162 electrons within the solvent-accessible voids.

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supporting information

Crystal structure of *cis*-7,8-*vic*-dihydroxy-5,10,15,20-tetraphenylchlorin and its zinc(II)–ethylenediamine complex

Nivedita Chaudhri, Christian Brückner* and Matthias Zeller

Computing details

Data collection: *APEX4* (Bruker, 2021) for 2PhH2; *APEX3* (Bruker, 2019) for 2PhZn. For both structures, cell refinement: *SAINT* (Bruker, 2019); data reduction: *SAINT* (Bruker, 2019). Program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a) for 2PhH2; *SHELXS97* (Sheldrick, 2008) for 2PhZn. For both structures, program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b), *ShelXle* (Hübschle *et al.*, 2011); molecular graphics: *Mercury* (Macrae *et al.*, 2020); software used to prepare material for publication: *publCIF* (Westrip, 2010).

cis-7,8-*vic*-Dihydroxy-5,10,15,20-tetraphenylchlorin dimethylaminopyridine monosolvate (2PhH2)

Crystal data

C₄₄H₃₂N₄O₂·C₇H₁₀N₂·[+solvent]*M_r* = 770.90Triclinic, *P*1*a* = 10.0193 (4) Å*b* = 15.2554 (8) Å*c* = 17.7983 (10) Å*α* = 69.918 (2)°*β* = 74.926 (2)°*γ* = 84.140 (2)°*V* = 2466.9 (2) Å³*Z* = 2*F*(000) = 812*D_x* = 1.038 Mg m⁻³Mo *Kα* radiation, *λ* = 0.71073 Å

Cell parameters from 9960 reflections

θ = 2.4–31.9°*μ* = 0.06 mm⁻¹*T* = 150 K

Fragment, black

0.33 × 0.21 × 0.19 mm

Data collection

Bruker AXS D8 Quest
diffractometer with PhotonII charge-integrating
pixel array detector (CPAD)

Radiation source: fine focus sealed tube X-ray
source

Triumph curved graphite crystal
monochromator

Detector resolution: 7.4074 pixels mm⁻¹*ω* and *φ* scans

Absorption correction: multi-scan
(*SADABS*; Krause *et al.*, 2015)

T_{min} = 0.665, *T_{max}* = 0.746

48645 measured reflections

14738 independent reflections

9891 reflections with *I* > 2σ(*I*)*R_{int}* = 0.060*θ_{max}* = 30.5°, *θ_{min}* = 2.2°*h* = −14→14*k* = −21→21*l* = −25→25

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.048*wR*(*F*²) = 0.133*S* = 1.04

14738 reflections

549 parameters

0 restraints

Primary atom site location: dual

Secondary atom site location: difference Fourier
map

Hydrogen site location: mixed

- 36 H atoms treated by a mixture of independent
and constrained refinement $(\Delta/\sigma)_{\max} = 0.001$
37 $w = 1/[\sigma^2(F_o^2) + (0.0604P)^2 + 0.2687P]$ $\Delta\rho_{\max} = 0.45 \text{ e } \text{\AA}^{-3}$
where $P = (F_o^2 + 2F_c^2)/3$ $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$

38 *Special details*

- 39 **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.
- 40 **Refinement.** The structure contains 647 Å ng₃ of solvent accessible voids occupied by highly disordered solvate molecules (presumably chloroform and hexane, the crystallization solvents). The residual electron density peaks are not arranged in an interpretable pattern and no unambiguous disorder model could be developed. The structure factors were instead augmented *via* reverse Fourier transform methods using the SQUEEZE routine (P. van der Sluis & A.L. Spek (1990). Acta Cryst. A46, 194–201) as implemented in the program *PLATON*. The resultant FAB file containing the structure factor contribution from the electron content of the void space was used in together with the original *hkl* file in the further refinement. (The FAB file with details of the Squeeze results is appended to this cif file). The Squeeze procedure corrected for 162 electrons within the solvent accessible voids.

41 *Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)*

42		x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
43	O1	0.62197 (8)	0.52783 (6)	0.53614 (5)	0.03032 (18)
44	H1O	0.6812 (16)	0.5788 (11)	0.4973 (10)	0.045*
45	O2	0.65897 (9)	0.42954 (6)	0.42876 (5)	0.03175 (18)
46	H2O	0.5704 (17)	0.4513 (11)	0.4458 (10)	0.048*
47	N1	0.43218 (10)	0.14254 (6)	0.68820 (6)	0.02595 (19)
48	H1N	0.4495 (15)	0.1795 (10)	0.7162 (9)	0.039*
49	N2	0.37812 (10)	0.12770 (6)	0.86064 (6)	0.02649 (19)
50	N3	0.55460 (10)	0.28307 (7)	0.82259 (6)	0.0283 (2)
51	H3N	0.5129 (16)	0.2577 (11)	0.7949 (10)	0.042*
52	N4	0.59891 (9)	0.30953 (6)	0.64424 (6)	0.02597 (19)
53	N5	0.79133 (13)	0.66232 (8)	0.42197 (8)	0.0513 (3)
54	N6	1.09618 (14)	0.82611 (9)	0.22875 (8)	0.0542 (3)
55	C1	0.48576 (12)	0.15869 (8)	0.60568 (7)	0.0272 (2)
56	C2	0.43926 (14)	0.08451 (8)	0.58748 (8)	0.0344 (3)
57	H2	0.461662	0.076127	0.535118	0.041*
58	C3	0.35751 (13)	0.02790 (8)	0.65773 (7)	0.0329 (3)
59	H3	0.312720	−0.026520	0.662782	0.039*
60	C4	0.35068 (11)	0.06429 (7)	0.72235 (7)	0.0261 (2)
61	C5	0.27900 (11)	0.02833 (7)	0.80485 (7)	0.0258 (2)
62	C6	0.28682 (11)	0.06218 (8)	0.86799 (7)	0.0258 (2)
63	C7	0.20051 (12)	0.02932 (8)	0.95097 (7)	0.0292 (2)
64	H7	0.128291	−0.014687	0.970852	0.035*
65	C8	0.24279 (12)	0.07350 (8)	0.99458 (7)	0.0300 (2)
66	H8	0.206040	0.066657	1.051130	0.036*
67	C9	0.35477 (11)	0.13318 (8)	0.93846 (7)	0.0270 (2)
68	C10	0.42982 (12)	0.18860 (8)	0.96202 (7)	0.0282 (2)
69	C11	0.52658 (12)	0.25542 (8)	0.90752 (7)	0.0293 (2)
70	C12	0.60953 (13)	0.31160 (9)	0.92638 (8)	0.0349 (3)

71	H12	0.614596	0.307363	0.980081	0.042*
72	C13	0.68056 (13)	0.37261 (9)	0.85413 (8)	0.0344 (3)
73	H13	0.742769	0.418500	0.848983	0.041*
74	C14	0.64544 (12)	0.35568 (8)	0.78779 (7)	0.0290 (2)
75	C15	0.69121 (11)	0.40654 (7)	0.70330 (7)	0.0266 (2)
76	C16	0.66552 (11)	0.38612 (7)	0.63837 (7)	0.0256 (2)
77	C17	0.71366 (11)	0.44961 (8)	0.54998 (7)	0.0269 (2)
78	H17	0.810994	0.469724	0.538272	0.032*
79	C18	0.70357 (11)	0.38542 (8)	0.50137 (7)	0.0271 (2)
80	H18	0.798257	0.359268	0.485428	0.033*
81	C19	0.61540 (11)	0.30561 (7)	0.56736 (7)	0.0258 (2)
82	C20	0.56811 (11)	0.23380 (8)	0.54904 (7)	0.0267 (2)
83	C21	0.18885 (12)	-0.05389 (8)	0.82851 (7)	0.0277 (2)
84	C22	0.21160 (13)	-0.13703 (8)	0.88883 (8)	0.0340 (3)
85	H22	0.286768	-0.142088	0.913483	0.041*
86	C23	0.12514 (15)	-0.21243 (9)	0.91308 (9)	0.0418 (3)
87	H23	0.141490	-0.268761	0.954185	0.050*
88	C24	0.01525 (14)	-0.20589 (10)	0.87763 (9)	0.0428 (3)
89	H24	-0.044311	-0.257364	0.894713	0.051*
90	C25	-0.00731 (13)	-0.12454 (10)	0.81758 (9)	0.0392 (3)
91	H25	-0.082307	-0.120161	0.792940	0.047*
92	C26	0.07881 (12)	-0.04862 (9)	0.79264 (8)	0.0321 (2)
93	H26	0.062527	0.007135	0.750918	0.039*
94	C27	0.40379 (12)	0.17673 (9)	1.05123 (7)	0.0306 (2)
95	C28	0.42357 (13)	0.09033 (9)	1.10829 (8)	0.0350 (3)
96	H28	0.453285	0.038015	1.090474	0.042*
97	C29	0.39991 (14)	0.08048 (11)	1.19136 (8)	0.0438 (3)
98	H29	0.413317	0.021333	1.230008	0.053*
99	C30	0.35704 (17)	0.15613 (13)	1.21802 (9)	0.0518 (4)
100	H30	0.341547	0.149010	1.274766	0.062*
101	C31	0.33677 (18)	0.24209 (12)	1.16204 (9)	0.0532 (4)
102	H31	0.307152	0.294134	1.180274	0.064*
103	C32	0.35962 (15)	0.25240 (10)	1.07935 (8)	0.0409 (3)
104	H32	0.345099	0.311659	1.041196	0.049*
105	C33	0.77693 (12)	0.48962 (8)	0.68495 (7)	0.0277 (2)
106	C34	0.71456 (13)	0.57598 (9)	0.68111 (8)	0.0357 (3)
107	H34	0.617350	0.583199	0.687579	0.043*
108	C35	0.79380 (15)	0.65217 (10)	0.66780 (9)	0.0435 (3)
109	H35	0.750570	0.711286	0.664713	0.052*
110	C36	0.93499 (15)	0.64211 (10)	0.65908 (9)	0.0443 (3)
111	H36	0.988404	0.693901	0.651294	0.053*
112	C37	0.99855 (14)	0.55681 (11)	0.66166 (9)	0.0444 (3)
113	H37	1.095810	0.549926	0.655054	0.053*
114	C38	0.91967 (13)	0.48109 (9)	0.67397 (9)	0.0379 (3)
115	H38	0.963808	0.422699	0.674888	0.045*
116	C39	0.61114 (12)	0.23285 (8)	0.46200 (7)	0.0285 (2)
117	C40	0.51769 (14)	0.25180 (9)	0.41304 (8)	0.0353 (3)
118	H40	0.424591	0.267589	0.434044	0.042*

119	C41	0.55892 (15)	0.24792 (10)	0.33351 (8)	0.0408 (3)
120	H41	0.494024	0.261266	0.300357	0.049*
121	C42	0.69418 (15)	0.22468 (9)	0.30227 (8)	0.0405 (3)
122	H42	0.722094	0.221702	0.247937	0.049*
123	C43	0.78742 (14)	0.20604 (10)	0.35014 (9)	0.0424 (3)
124	H43	0.880424	0.190326	0.328850	0.051*
125	C44	0.74671 (13)	0.20998 (9)	0.42957 (8)	0.0369 (3)
126	H44	0.812284	0.196878	0.462273	0.044*
127	C45	0.89678 (18)	0.69732 (11)	0.43549 (10)	0.0546 (4)
128	H45	0.901928	0.682842	0.491021	0.065*
129	C46	0.99821 (17)	0.75277 (10)	0.37465 (10)	0.0504 (4)
130	H46	1.069780	0.775754	0.388603	0.060*
131	C47	0.99487 (15)	0.77500 (9)	0.29192 (9)	0.0435 (3)
132	C48	0.88343 (15)	0.73985 (10)	0.27753 (10)	0.0484 (4)
133	H48	0.873990	0.753889	0.222851	0.058*
134	C49	0.78732 (15)	0.68461 (10)	0.34342 (11)	0.0504 (4)
135	H49	0.713560	0.661011	0.331852	0.060*
136	C50	1.20258 (17)	0.86988 (12)	0.24494 (12)	0.0637 (5)
137	H50A	1.267695	0.901455	0.192839	0.096*
138	H50B	1.252073	0.822161	0.281093	0.096*
139	H50C	1.159873	0.915591	0.271836	0.096*
140	C51	1.09541 (19)	0.84148 (13)	0.14456 (10)	0.0656 (5)
141	H51A	1.184094	0.867545	0.108436	0.098*
142	H51B	1.020473	0.885300	0.129652	0.098*
143	H51C	1.081150	0.782071	0.138165	0.098*

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

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
146	O1	0.0308 (4)	0.0229 (4)	0.0316 (4)	−0.0034 (3)	−0.0017 (3)
147	O2	0.0346 (4)	0.0310 (4)	0.0244 (4)	−0.0029 (4)	−0.0022 (3)
148	N1	0.0303 (5)	0.0227 (4)	0.0226 (4)	−0.0058 (4)	−0.0006 (4)
149	N2	0.0297 (5)	0.0251 (4)	0.0228 (5)	−0.0050 (4)	−0.0019 (4)
150	N3	0.0328 (5)	0.0262 (5)	0.0241 (5)	−0.0088 (4)	−0.0024 (4)
151	N4	0.0278 (4)	0.0240 (4)	0.0237 (5)	−0.0044 (4)	−0.0022 (4)
152	N5	0.0458 (7)	0.0300 (6)	0.0536 (8)	−0.0030 (5)	0.0108 (6)
153	N6	0.0464 (7)	0.0388 (6)	0.0517 (8)	−0.0073 (6)	0.0050 (6)
154	C1	0.0315 (5)	0.0248 (5)	0.0230 (5)	−0.0046 (4)	−0.0009 (4)
155	C2	0.0451 (7)	0.0297 (6)	0.0270 (6)	−0.0103 (5)	0.0010 (5)
156	C3	0.0430 (6)	0.0273 (5)	0.0275 (6)	−0.0102 (5)	−0.0015 (5)
157	C4	0.0301 (5)	0.0214 (5)	0.0244 (5)	−0.0051 (4)	−0.0025 (4)
158	C5	0.0265 (5)	0.0230 (5)	0.0256 (5)	−0.0041 (4)	−0.0028 (4)
159	C6	0.0267 (5)	0.0248 (5)	0.0231 (5)	−0.0036 (4)	−0.0025 (4)
160	C7	0.0281 (5)	0.0315 (6)	0.0243 (5)	−0.0077 (5)	−0.0002 (4)
161	C8	0.0309 (5)	0.0335 (6)	0.0222 (5)	−0.0056 (5)	0.0002 (4)
162	C9	0.0287 (5)	0.0258 (5)	0.0240 (5)	−0.0027 (4)	−0.0026 (4)
163	C10	0.0323 (5)	0.0263 (5)	0.0240 (5)	−0.0040 (4)	−0.0032 (4)
164	C11	0.0339 (6)	0.0281 (5)	0.0250 (5)	−0.0048 (5)	−0.0044 (5)

165	C12	0.0417 (7)	0.0355 (6)	0.0280 (6)	−0.0111 (5)	−0.0069 (5)	−0.0091 (5)
166	C13	0.0386 (6)	0.0351 (6)	0.0308 (6)	−0.0118 (5)	−0.0062 (5)	−0.0107 (5)
167	C14	0.0324 (5)	0.0251 (5)	0.0285 (6)	−0.0063 (4)	−0.0035 (5)	−0.0087 (4)
168	C15	0.0277 (5)	0.0227 (5)	0.0270 (5)	−0.0051 (4)	−0.0025 (4)	−0.0069 (4)
169	C16	0.0252 (5)	0.0229 (5)	0.0250 (5)	−0.0037 (4)	−0.0020 (4)	−0.0053 (4)
170	C17	0.0256 (5)	0.0242 (5)	0.0256 (5)	−0.0048 (4)	−0.0006 (4)	−0.0044 (4)
171	C18	0.0264 (5)	0.0264 (5)	0.0238 (5)	−0.0039 (4)	0.0002 (4)	−0.0059 (4)
172	C19	0.0256 (5)	0.0238 (5)	0.0236 (5)	−0.0023 (4)	−0.0004 (4)	−0.0059 (4)
173	C20	0.0290 (5)	0.0251 (5)	0.0227 (5)	−0.0032 (4)	−0.0009 (4)	−0.0071 (4)
174	C21	0.0289 (5)	0.0266 (5)	0.0256 (5)	−0.0062 (4)	0.0013 (4)	−0.0102 (4)
175	C22	0.0375 (6)	0.0303 (6)	0.0303 (6)	−0.0087 (5)	−0.0032 (5)	−0.0063 (5)
176	C23	0.0463 (7)	0.0291 (6)	0.0406 (7)	−0.0117 (5)	0.0022 (6)	−0.0059 (5)
177	C24	0.0380 (7)	0.0378 (7)	0.0496 (8)	−0.0173 (6)	0.0082 (6)	−0.0198 (6)
178	C25	0.0302 (6)	0.0460 (7)	0.0463 (8)	−0.0084 (5)	−0.0011 (6)	−0.0252 (6)
179	C26	0.0308 (6)	0.0334 (6)	0.0328 (6)	−0.0036 (5)	−0.0025 (5)	−0.0146 (5)
180	C27	0.0324 (6)	0.0347 (6)	0.0242 (5)	−0.0088 (5)	−0.0044 (5)	−0.0084 (5)
181	C28	0.0312 (6)	0.0395 (7)	0.0312 (6)	−0.0064 (5)	−0.0076 (5)	−0.0061 (5)
182	C29	0.0402 (7)	0.0565 (9)	0.0297 (7)	−0.0134 (6)	−0.0135 (6)	−0.0006 (6)
183	C30	0.0581 (9)	0.0730 (11)	0.0286 (7)	−0.0243 (8)	−0.0086 (7)	−0.0167 (7)
184	C31	0.0709 (10)	0.0573 (9)	0.0370 (8)	−0.0205 (8)	−0.0013 (7)	−0.0256 (7)
185	C32	0.0537 (8)	0.0374 (7)	0.0320 (7)	−0.0110 (6)	−0.0038 (6)	−0.0139 (5)
186	C33	0.0295 (5)	0.0276 (5)	0.0239 (5)	−0.0075 (4)	−0.0009 (4)	−0.0079 (4)
187	C34	0.0327 (6)	0.0314 (6)	0.0412 (7)	−0.0070 (5)	0.0012 (5)	−0.0153 (5)
188	C35	0.0499 (8)	0.0330 (6)	0.0470 (8)	−0.0116 (6)	0.0020 (6)	−0.0193 (6)
189	C36	0.0509 (8)	0.0459 (8)	0.0363 (7)	−0.0257 (7)	−0.0007 (6)	−0.0140 (6)
190	C37	0.0332 (6)	0.0545 (8)	0.0392 (7)	−0.0189 (6)	−0.0025 (6)	−0.0072 (6)
191	C38	0.0301 (6)	0.0364 (6)	0.0413 (7)	−0.0054 (5)	−0.0036 (5)	−0.0076 (6)
192	C39	0.0349 (6)	0.0239 (5)	0.0237 (5)	−0.0080 (4)	0.0015 (5)	−0.0081 (4)
193	C40	0.0376 (6)	0.0368 (6)	0.0296 (6)	−0.0028 (5)	−0.0023 (5)	−0.0121 (5)
194	C41	0.0533 (8)	0.0402 (7)	0.0308 (7)	−0.0056 (6)	−0.0090 (6)	−0.0134 (6)
195	C42	0.0548 (8)	0.0362 (7)	0.0292 (6)	−0.0141 (6)	0.0052 (6)	−0.0165 (5)
196	C43	0.0393 (7)	0.0483 (8)	0.0418 (7)	−0.0106 (6)	0.0071 (6)	−0.0272 (7)
197	C44	0.0352 (6)	0.0403 (7)	0.0374 (7)	−0.0039 (5)	−0.0009 (5)	−0.0204 (6)
198	C45	0.0650 (10)	0.0410 (8)	0.0440 (8)	−0.0082 (7)	0.0077 (8)	−0.0101 (7)
199	C46	0.0550 (9)	0.0366 (7)	0.0508 (9)	−0.0091 (6)	0.0013 (7)	−0.0117 (6)
200	C47	0.0415 (7)	0.0234 (6)	0.0466 (8)	−0.0005 (5)	0.0051 (6)	0.0006 (5)
201	C48	0.0431 (8)	0.0367 (7)	0.0486 (8)	0.0037 (6)	−0.0048 (6)	0.0013 (6)
202	C49	0.0368 (7)	0.0340 (7)	0.0629 (10)	0.0013 (6)	−0.0042 (7)	−0.0005 (7)
203	C50	0.0418 (8)	0.0402 (8)	0.0829 (13)	−0.0060 (7)	−0.0022 (8)	0.0046 (8)
204	C51	0.0553 (10)	0.0607 (10)	0.0467 (9)	0.0059 (8)	0.0061 (8)	0.0095 (8)

205 *Geometric parameters (Å, °)*

206	O1—C17	1.4214 (14)	C23—C24	1.384 (2)
207	O1—H1O	0.973 (17)	C23—H23	0.9500
208	O2—C18	1.4016 (14)	C24—C25	1.375 (2)
209	O2—H2O	0.927 (17)	C24—H24	0.9500
210	N1—C1	1.3692 (14)	C25—C26	1.3909 (17)

211	N1—C4	1.3797 (14)	C25—H25	0.9500
212	N1—H1N	0.925 (15)	C26—H26	0.9500
213	N2—C9	1.3737 (14)	C27—C28	1.3929 (18)
214	N2—C6	1.3740 (14)	C27—C32	1.3986 (17)
215	N3—C14	1.3714 (14)	C28—C29	1.3916 (18)
216	N3—C11	1.3811 (15)	C28—H28	0.9500
217	N3—H3N	0.915 (16)	C29—C30	1.382 (2)
218	N4—C19	1.3565 (14)	C29—H29	0.9500
219	N4—C16	1.3639 (14)	C30—C31	1.381 (2)
220	N5—C49	1.331 (2)	C30—H30	0.9500
221	N5—C45	1.340 (2)	C31—C32	1.3843 (19)
222	N6—C47	1.3669 (18)	C31—H31	0.9500
223	N6—C51	1.437 (2)	C32—H32	0.9500
224	N6—C50	1.450 (2)	C33—C34	1.3867 (17)
225	C1—C20	1.4062 (15)	C33—C38	1.3910 (17)
226	C1—C2	1.4284 (16)	C34—C35	1.3924 (17)
227	C2—C3	1.3602 (17)	C34—H34	0.9500
228	C2—H2	0.9500	C35—C36	1.381 (2)
229	C3—C4	1.4232 (16)	C35—H35	0.9500
230	C3—H3	0.9500	C36—C37	1.381 (2)
231	C4—C5	1.3970 (15)	C36—H36	0.9500
232	C5—C6	1.4101 (16)	C37—C38	1.3891 (19)
233	C5—C21	1.4958 (15)	C37—H37	0.9500
234	C6—C7	1.4478 (16)	C38—H38	0.9500
235	C7—C8	1.3508 (16)	C39—C40	1.3840 (18)
236	C7—H7	0.9500	C39—C44	1.3905 (17)
237	C8—C9	1.4445 (16)	C40—C41	1.3879 (17)
238	C8—H8	0.9500	C40—H40	0.9500
239	C9—C10	1.4119 (16)	C41—C42	1.384 (2)
240	C10—C11	1.3977 (16)	C41—H41	0.9500
241	C10—C27	1.4895 (16)	C42—C43	1.368 (2)
242	C11—C12	1.4233 (16)	C42—H42	0.9500
243	C12—C13	1.3651 (18)	C43—C44	1.3864 (18)
244	C12—H12	0.9500	C43—H43	0.9500
245	C13—C14	1.4243 (17)	C44—H44	0.9500
246	C13—H13	0.9500	C45—C46	1.377 (2)
247	C14—C15	1.4094 (16)	C45—H45	0.9500
248	C15—C16	1.3846 (16)	C46—C47	1.402 (2)
249	C15—C33	1.4977 (15)	C46—H46	0.9500
250	C16—C17	1.5166 (16)	C47—C48	1.402 (2)
251	C17—C18	1.5380 (16)	C48—C49	1.383 (2)
252	C17—H17	1.0000	C48—H48	0.9500
253	C18—C19	1.5298 (15)	C49—H49	0.9500
254	C18—H18	1.0000	C50—H50A	0.9800
255	C19—C20	1.4008 (15)	C50—H50B	0.9800
256	C20—C39	1.5008 (15)	C50—H50C	0.9800
257	C21—C26	1.3937 (17)	C51—H51A	0.9800
258	C21—C22	1.3951 (17)	C51—H51B	0.9800

259	C22—C23	1.3884 (17)	C51—H51C	0.9800
260	C22—H22	0.9500		
261				
262	C17—O1—H1O	103.8 (9)	C25—C24—H24	120.2
263	C18—O2—H2O	105.0 (10)	C23—C24—H24	120.2
264	C1—N1—C4	110.47 (9)	C24—C25—C26	120.52 (13)
265	C1—N1—H1N	123.8 (9)	C24—C25—H25	119.7
266	C4—N1—H1N	125.7 (9)	C26—C25—H25	119.7
267	C9—N2—C6	104.78 (9)	C25—C26—C21	120.38 (12)
268	C14—N3—C11	110.47 (10)	C25—C26—H26	119.8
269	C14—N3—H3N	126.2 (10)	C21—C26—H26	119.8
270	C11—N3—H3N	123.2 (10)	C28—C27—C32	118.61 (12)
271	C19—N4—C16	108.59 (9)	C28—C27—C10	120.80 (11)
272	C49—N5—C45	115.86 (13)	C32—C27—C10	120.58 (11)
273	C47—N6—C51	120.56 (15)	C29—C28—C27	120.15 (13)
274	C47—N6—C50	120.85 (15)	C29—C28—H28	119.9
275	C51—N6—C50	118.52 (14)	C27—C28—H28	119.9
276	N1—C1—C20	127.44 (10)	C30—C29—C28	120.47 (14)
277	N1—C1—C2	106.31 (10)	C30—C29—H29	119.8
278	C20—C1—C2	126.24 (10)	C28—C29—H29	119.8
279	C3—C2—C1	108.53 (10)	C31—C30—C29	119.90 (13)
280	C3—C2—H2	125.7	C31—C30—H30	120.1
281	C1—C2—H2	125.7	C29—C30—H30	120.1
282	C2—C3—C4	108.18 (10)	C30—C31—C32	119.99 (14)
283	C2—C3—H3	125.9	C30—C31—H31	120.0
284	C4—C3—H3	125.9	C32—C31—H31	120.0
285	N1—C4—C5	125.72 (10)	C31—C32—C27	120.88 (14)
286	N1—C4—C3	106.47 (10)	C31—C32—H32	119.6
287	C5—C4—C3	127.81 (10)	C27—C32—H32	119.6
288	C4—C5—C6	125.26 (10)	C34—C33—C38	118.83 (11)
289	C4—C5—C21	117.43 (10)	C34—C33—C15	120.39 (10)
290	C6—C5—C21	117.30 (10)	C38—C33—C15	120.75 (11)
291	N2—C6—C5	125.47 (10)	C33—C34—C35	120.29 (12)
292	N2—C6—C7	110.91 (9)	C33—C34—H34	119.9
293	C5—C6—C7	123.59 (10)	C35—C34—H34	119.9
294	C8—C7—C6	106.56 (10)	C36—C35—C34	120.24 (13)
295	C8—C7—H7	126.7	C36—C35—H35	119.9
296	C6—C7—H7	126.7	C34—C35—H35	119.9
297	C7—C8—C9	106.62 (10)	C37—C36—C35	120.02 (12)
298	C7—C8—H8	126.7	C37—C36—H36	120.0
299	C9—C8—H8	126.7	C35—C36—H36	120.0
300	N2—C9—C10	125.32 (10)	C36—C37—C38	119.72 (13)
301	N2—C9—C8	111.05 (10)	C36—C37—H37	120.1
302	C10—C9—C8	123.63 (10)	C38—C37—H37	120.1
303	C11—C10—C9	124.84 (11)	C37—C38—C33	120.87 (13)
304	C11—C10—C27	116.86 (10)	C37—C38—H38	119.6
305	C9—C10—C27	118.30 (10)	C33—C38—H38	119.6
306	N3—C11—C10	125.39 (10)	C40—C39—C44	118.43 (11)

307	N3—C11—C12	106.27 (10)	C40—C39—C20	121.53 (10)
308	C10—C11—C12	128.27 (11)	C44—C39—C20	120.02 (11)
309	C13—C12—C11	108.35 (11)	C39—C40—C41	120.53 (12)
310	C13—C12—H12	125.8	C39—C40—H40	119.7
311	C11—C12—H12	125.8	C41—C40—H40	119.7
312	C12—C13—C14	108.32 (11)	C42—C41—C40	120.31 (13)
313	C12—C13—H13	125.8	C42—C41—H41	119.8
314	C14—C13—H13	125.8	C40—C41—H41	119.8
315	N3—C14—C15	127.08 (10)	C43—C42—C41	119.60 (12)
316	N3—C14—C13	106.53 (10)	C43—C42—H42	120.2
317	C15—C14—C13	126.34 (10)	C41—C42—H42	120.2
318	C16—C15—C14	126.44 (10)	C42—C43—C44	120.28 (12)
319	C16—C15—C33	118.94 (10)	C42—C43—H43	119.9
320	C14—C15—C33	114.62 (10)	C44—C43—H43	119.9
321	N4—C16—C15	126.10 (10)	C43—C44—C39	120.86 (13)
322	N4—C16—C17	112.44 (9)	C43—C44—H44	119.6
323	C15—C16—C17	121.45 (10)	C39—C44—H44	119.6
324	O1—C17—C16	108.57 (9)	N5—C45—C46	124.66 (17)
325	O1—C17—C18	112.64 (9)	N5—C45—H45	117.7
326	C16—C17—C18	101.86 (9)	C46—C45—H45	117.7
327	O1—C17—H17	111.1	C45—C46—C47	119.36 (16)
328	C16—C17—H17	111.1	C45—C46—H46	120.3
329	C18—C17—H17	111.1	C47—C46—H46	120.3
330	O2—C18—C19	117.07 (9)	N6—C47—C46	121.97 (15)
331	O2—C18—C17	115.00 (9)	N6—C47—C48	121.83 (15)
332	C19—C18—C17	102.30 (9)	C46—C47—C48	116.18 (13)
333	O2—C18—H18	107.3	C49—C48—C47	119.59 (16)
334	C19—C18—H18	107.3	C49—C48—H48	120.2
335	C17—C18—H18	107.3	C47—C48—H48	120.2
336	N4—C19—C20	125.15 (10)	N5—C49—C48	124.33 (16)
337	N4—C19—C18	112.16 (9)	N5—C49—H49	117.8
338	C20—C19—C18	122.51 (10)	C48—C49—H49	117.8
339	C19—C20—C1	126.08 (10)	N6—C50—H50A	109.5
340	C19—C20—C39	118.64 (10)	N6—C50—H50B	109.5
341	C1—C20—C39	115.24 (10)	H50A—C50—H50B	109.5
342	C26—C21—C22	118.64 (11)	N6—C50—H50C	109.5
343	C26—C21—C5	121.07 (11)	H50A—C50—H50C	109.5
344	C22—C21—C5	120.27 (11)	H50B—C50—H50C	109.5
345	C23—C22—C21	120.46 (12)	N6—C51—H51A	109.5
346	C23—C22—H22	119.8	N6—C51—H51B	109.5
347	C21—C22—H22	119.8	H51A—C51—H51B	109.5
348	C24—C23—C22	120.31 (13)	N6—C51—H51C	109.5
349	C24—C23—H23	119.8	H51A—C51—H51C	109.5
350	C22—C23—H23	119.8	H51B—C51—H51C	109.5
351	C25—C24—C23	119.69 (12)		
352				
353	C4—N1—C1—C20	176.83 (11)	C18—C19—C20—C1	−179.31 (11)
354	C4—N1—C1—C2	−2.11 (13)	N4—C19—C20—C39	−171.75 (10)

355	N1—C1—C2—C3	1.49 (14)	C18—C19—C20—C39	2.90 (16)
356	C20—C1—C2—C3	−177.46 (12)	N1—C1—C20—C19	2.4 (2)
357	C1—C2—C3—C4	−0.35 (15)	C2—C1—C20—C19	−178.88 (12)
358	C1—N1—C4—C5	−179.30 (11)	N1—C1—C20—C39	−179.76 (11)
359	C1—N1—C4—C3	1.91 (13)	C2—C1—C20—C39	−1.02 (18)
360	C2—C3—C4—N1	−0.92 (14)	C4—C5—C21—C26	−60.11 (15)
361	C2—C3—C4—C5	−179.68 (12)	C6—C5—C21—C26	121.08 (12)
362	N1—C4—C5—C6	−5.13 (19)	C4—C5—C21—C22	121.51 (12)
363	C3—C4—C5—C6	173.41 (12)	C6—C5—C21—C22	−57.30 (15)
364	N1—C4—C5—C21	176.17 (10)	C26—C21—C22—C23	−0.72 (18)
365	C3—C4—C5—C21	−5.30 (18)	C5—C21—C22—C23	177.70 (11)
366	C9—N2—C6—C5	−175.53 (11)	C21—C22—C23—C24	0.0 (2)
367	C9—N2—C6—C7	2.71 (12)	C22—C23—C24—C25	0.6 (2)
368	C4—C5—C6—N2	−8.48 (18)	C23—C24—C25—C26	−0.5 (2)
369	C21—C5—C6—N2	170.23 (10)	C24—C25—C26—C21	−0.30 (18)
370	C4—C5—C6—C7	173.50 (11)	C22—C21—C26—C25	0.88 (17)
371	C21—C5—C6—C7	−7.80 (16)	C5—C21—C26—C25	−177.52 (10)
372	N2—C6—C7—C8	−1.69 (13)	C11—C10—C27—C28	−122.22 (13)
373	C5—C6—C7—C8	176.58 (11)	C9—C10—C27—C28	58.57 (16)
374	C6—C7—C8—C9	−0.05 (13)	C11—C10—C27—C32	57.43 (16)
375	C6—N2—C9—C10	176.68 (11)	C9—C10—C27—C32	−121.78 (13)
376	C6—N2—C9—C8	−2.75 (12)	C32—C27—C28—C29	−0.18 (18)
377	C7—C8—C9—N2	1.78 (13)	C10—C27—C28—C29	179.47 (11)
378	C7—C8—C9—C10	−177.66 (11)	C27—C28—C29—C30	−0.2 (2)
379	N2—C9—C10—C11	9.00 (19)	C28—C29—C30—C31	0.3 (2)
380	C8—C9—C10—C11	−171.64 (11)	C29—C30—C31—C32	−0.1 (2)
381	N2—C9—C10—C27	−171.86 (11)	C30—C31—C32—C27	−0.3 (2)
382	C8—C9—C10—C27	7.50 (17)	C28—C27—C32—C31	0.4 (2)
383	C14—N3—C11—C10	174.45 (11)	C10—C27—C32—C31	−179.25 (13)
384	C14—N3—C11—C12	−2.57 (13)	C16—C15—C33—C34	91.51 (14)
385	C9—C10—C11—N3	5.97 (19)	C14—C15—C33—C34	−89.12 (14)
386	C27—C10—C11—N3	−173.17 (11)	C16—C15—C33—C38	−90.45 (15)
387	C9—C10—C11—C12	−177.67 (12)	C14—C15—C33—C38	88.92 (14)
388	C27—C10—C11—C12	3.18 (19)	C38—C33—C34—C35	−1.03 (19)
389	N3—C11—C12—C13	1.97 (14)	C15—C33—C34—C35	177.04 (12)
390	C10—C11—C12—C13	−174.93 (12)	C33—C34—C35—C36	−0.6 (2)
391	C11—C12—C13—C14	−0.69 (15)	C34—C35—C36—C37	1.5 (2)
392	C11—N3—C14—C15	−175.45 (11)	C35—C36—C37—C38	−0.7 (2)
393	C11—N3—C14—C13	2.16 (13)	C36—C37—C38—C33	−0.9 (2)
394	C12—C13—C14—N3	−0.87 (14)	C34—C33—C38—C37	1.8 (2)
395	C12—C13—C14—C15	176.76 (12)	C15—C33—C38—C37	−176.26 (12)
396	N3—C14—C15—C16	−7.9 (2)	C19—C20—C39—C40	−110.08 (13)
397	C13—C14—C15—C16	174.91 (12)	C1—C20—C39—C40	71.89 (14)
398	N3—C14—C15—C33	172.75 (11)	C19—C20—C39—C44	71.77 (15)
399	C13—C14—C15—C33	−4.41 (17)	C1—C20—C39—C44	−106.26 (13)
400	C19—N4—C16—C15	−169.25 (11)	C44—C39—C40—C41	0.02 (18)
401	C19—N4—C16—C17	9.77 (12)	C20—C39—C40—C41	−178.16 (11)
402	C14—C15—C16—N4	−4.65 (19)	C39—C40—C41—C42	0.2 (2)

403	C33—C15—C16—N4	174.64 (10)	C40—C41—C42—C43	−0.4 (2)
404	C14—C15—C16—C17	176.41 (11)	C41—C42—C43—C44	0.3 (2)
405	C33—C15—C16—C17	−4.31 (16)	C42—C43—C44—C39	0.0 (2)
406	N4—C16—C17—O1	103.05 (10)	C40—C39—C44—C43	−0.12 (19)
407	C15—C16—C17—O1	−77.88 (13)	C20—C39—C44—C43	178.09 (12)
408	N4—C16—C17—C18	−16.00 (12)	C49—N5—C45—C46	−0.4 (2)
409	C15—C16—C17—C18	163.07 (10)	N5—C45—C46—C47	−0.6 (2)
410	O1—C17—C18—O2	26.65 (13)	C51—N6—C47—C46	175.09 (14)
411	C16—C17—C18—O2	142.76 (9)	C50—N6—C47—C46	−8.1 (2)
412	O1—C17—C18—C19	−101.33 (10)	C51—N6—C47—C48	−3.3 (2)
413	C16—C17—C18—C19	14.78 (10)	C50—N6—C47—C48	173.57 (13)
414	C16—N4—C19—C20	176.25 (10)	C45—C46—C47—N6	−176.90 (14)
415	C16—N4—C19—C18	1.12 (12)	C45—C46—C47—C48	1.6 (2)
416	O2—C18—C19—N4	−137.49 (10)	N6—C47—C48—C49	176.77 (13)
417	C17—C18—C19—N4	−10.83 (12)	C46—C47—C48—C49	−1.7 (2)
418	O2—C18—C19—C20	47.23 (15)	C45—N5—C49—C48	0.2 (2)
419	C17—C18—C19—C20	173.89 (10)	C47—C48—C49—N5	0.9 (2)
420	N4—C19—C20—C1	6.04 (19)		

421 *Hydrogen-bond geometry (Å, °)*

422	<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>
423	O1—H1O $\cdots$ N5	0.973 (17)	1.727 (17)	2.6968 (14)	174.1 (14)
424	O2—H2O $\cdots$ O1 ⁱ	0.927 (17)	1.882 (17)	2.7798 (12)	162.5 (14)
425	N1—H1N $\cdots$ N2	0.925 (15)	2.346 (15)	2.9064 (13)	118.7 (11)
426	N1—H1N $\cdots$ N4	0.925 (15)	2.383 (15)	2.9518 (13)	119.6 (11)
427	N3—H3N $\cdots$ N2	0.915 (16)	2.292 (16)	2.8868 (13)	122.3 (12)
428	N3—H3N $\cdots$ N4	0.915 (16)	2.458 (15)	2.9766 (14)	116.1 (12)
429	C37—H37 $\cdots$ O2 ⁱⁱ	0.95	2.51	3.3840 (16)	153
430	C38—H38 $\cdots$ C48 ⁱⁱ	0.95	2.77	3.6779 (19)	161
431	C50—H50B $\cdots$ N4 ⁱⁱ	0.98	2.57	3.544 (2)	171

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

433 **[*cis*-7,8-*vic*-Dihydroxy-5,10,15,20-tetraphenylchlorinato(2-)]zinc(II)–ethylenediamine–methanol (1/1/0.136)**

434 **(2PhZn)**

435 *Crystal data*436 [Zn(C₄₄H₃₀N₄O₂)]·C₂H₈N₂·0.136CH₄O437 *M_r* = 776.57438 Monoclinic, *P*2₁/*c*439 *a* = 10.1249 (3) Å440 *b* = 13.5400 (4) Å441 *c* = 27.0447 (8) Å442 β = 95.1464 (11)°443 *V* = 3692.64 (19) Å³444 *Z* = 4*F*(000) = 1618*D_x* = 1.397 Mg m^{−3}Cu *K*α radiation, λ = 1.54178 Å

Cell parameters from 9950 reflections

 θ = 3.3–79.4° μ = 1.32 mm^{−1}*T* = 150 K

Block, black

0.27 × 0.25 × 0.18 mm

445 *Data collection*

446 Bruker AXS D8 Quest
diffractometer with PhotonIII-C14 charge-
integrating and photon counting pixel array
detector

447 Radiation source: I-mu-S microsource X-ray
tube

448 Laterally graded multilayer (Goebel) mirror
monochromator

449 Detector resolution: 7.4074 pixels mm⁻¹

450 ω and ϕ scans

Absorption correction: multi-scan
(*SADABS*; Krause *et al.*, 2015)

$T_{\min} = 0.606$, $T_{\max} = 0.754$

21319 measured reflections

7551 independent reflections

7037 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.024$

$\theta_{\max} = 79.5^\circ$, $\theta_{\min} = 3.3^\circ$

$h = -12 \rightarrow 11$

$k = -16 \rightarrow 15$

$l = -29 \rightarrow 34$

451 *Refinement*

452 Refinement on F^2

453 Least-squares matrix: full

454 $R[F^2 > 2\sigma(F^2)] = 0.031$

455 $wR(F^2) = 0.088$

456 $S = 1.04$

457 7551 reflections

458 549 parameters

459 17 restraints

460 Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: mixed

H atoms treated by a mixture of independent
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0454P)^2 + 1.8191P]$

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

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

$\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$

461 *Special details*

462 **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.

463 **Refinement.** The not metal coordinated amino group of an ethylene diamine ligand was refined as disordered. The C—N
bonds were restrained to be similar in length. Amine H atom positions were refined and N—H distances were restrained
to 0.88 (2) Angstrom. Equivalent H···H and C···H distances were restrained to be similar to each other. Subject to these
conditions the occupancy ratio refined to 0.882 (12) to 0.118 (12).

A partially occupied methanol molecule is located nearby the major disordered amino group and H-bonded to it. The
hydroxyl H atom was restrained to hydrogen bond to a porphyrin N atom of a neighboring complex. Subject to these
conditions the occupancy rate refined to 0.136 (4).

464 *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)
465 Zn1	0.70722 (2)	0.44915 (2)	0.36168 (2)	0.01832 (7)	
466 O1	0.62030 (12)	0.47166 (9)	0.54271 (4)	0.0308 (2)	
467 H1	0.6482 (10)	0.4528 (15)	0.5775 (9)	0.046*	
468 O2	0.56022 (12)	0.65470 (9)	0.51056 (4)	0.0328 (3)	
469 H2A	0.4922 (17)	0.6092 (15)	0.4949 (8)	0.049*	
470 N1	0.73694 (12)	0.51455 (9)	0.43385 (4)	0.0204 (2)	
471 N2	0.68821 (12)	0.58909 (9)	0.33244 (4)	0.0202 (2)	
472 N3	0.73798 (12)	0.39779 (9)	0.29110 (4)	0.0192 (2)	
473 N4	0.80823 (12)	0.32619 (9)	0.39012 (4)	0.0201 (2)	
474 C1	0.76495 (14)	0.46245 (11)	0.47647 (5)	0.0211 (3)	
475 C2	0.72768 (15)	0.51995 (12)	0.52170 (5)	0.0247 (3)	

477	H2	0.805841	0.525874	0.546886	0.030*
478	C3	0.68812 (16)	0.62230 (12)	0.49999 (5)	0.0256 (3)
479	H3	0.755257	0.671994	0.513274	0.031*
480	C4	0.70045 (14)	0.60845 (11)	0.44448 (5)	0.0217 (3)
481	C5	0.67805 (14)	0.68419 (11)	0.41046 (5)	0.0221 (3)
482	C6	0.67411 (14)	0.67471 (11)	0.35816 (5)	0.0214 (3)
483	C7	0.65058 (16)	0.75574 (11)	0.32390 (6)	0.0273 (3)
484	H7	0.638690	0.823086	0.332289	0.033*
485	C8	0.64860 (16)	0.71772 (11)	0.27743 (6)	0.0274 (3)
486	H8	0.635401	0.753563	0.247193	0.033*
487	C9	0.67021 (14)	0.61307 (11)	0.28244 (5)	0.0216 (3)
488	C10	0.66810 (14)	0.54655 (11)	0.24303 (5)	0.0212 (3)
489	C11	0.69457 (14)	0.44448 (11)	0.24758 (5)	0.0207 (3)
490	C12	0.68755 (15)	0.37502 (12)	0.20741 (5)	0.0255 (3)
491	H12	0.658048	0.388023	0.173700	0.031*
492	C13	0.73118 (15)	0.28702 (11)	0.22677 (5)	0.0246 (3)
493	H13	0.737737	0.226813	0.209116	0.030*
494	C14	0.76550 (14)	0.30210 (11)	0.27891 (5)	0.0200 (3)
495	C15	0.82584 (14)	0.23173 (11)	0.31236 (5)	0.0205 (3)
496	C16	0.85130 (14)	0.24664 (11)	0.36358 (5)	0.0221 (3)
497	C17	0.92352 (17)	0.17956 (12)	0.39728 (6)	0.0295 (3)
498	H17	0.965688	0.120098	0.388676	0.035*
499	C18	0.92010 (17)	0.21707 (13)	0.44358 (6)	0.0304 (3)
500	H18	0.958959	0.188554	0.473521	0.036*
501	C19	0.84659 (14)	0.30825 (11)	0.43899 (5)	0.0227 (3)
502	C20	0.81842 (14)	0.36856 (11)	0.47976 (5)	0.0222 (3)
503	C21	0.66512 (16)	0.78725 (11)	0.42976 (5)	0.0257 (3)
504	C22	0.54419 (18)	0.83574 (13)	0.42621 (7)	0.0343 (4)
505	H22	0.466488	0.801861	0.413263	0.041*
506	C23	0.5353 (2)	0.93293 (15)	0.44131 (8)	0.0464 (5)
507	H23	0.451780	0.965418	0.438633	0.056*
508	C24	0.6477 (2)	0.98287 (14)	0.46029 (8)	0.0492 (5)
509	H24	0.641627	1.049810	0.470365	0.059*
510	C25	0.7680 (2)	0.93559 (14)	0.46454 (8)	0.0478 (5)
511	H25	0.845182	0.969597	0.477894	0.057*
512	C26	0.77712 (19)	0.83764 (13)	0.44927 (7)	0.0370 (4)
513	H26	0.860658	0.805211	0.452263	0.044*
514	C27	0.63996 (15)	0.58672 (11)	0.19152 (5)	0.0222 (3)
515	C28	0.51396 (15)	0.62173 (12)	0.17514 (6)	0.0271 (3)
516	H28	0.445274	0.620404	0.196869	0.033*
517	C29	0.48819 (17)	0.65857 (12)	0.12724 (6)	0.0303 (3)
518	H29	0.401410	0.680622	0.116268	0.036*
519	C30	0.58747 (18)	0.66344 (12)	0.09542 (6)	0.0317 (3)
520	H30	0.569838	0.689923	0.062973	0.038*
521	C31	0.71357 (18)	0.62906 (15)	0.11152 (6)	0.0376 (4)
522	H31	0.782543	0.632158	0.089955	0.045*
523	C32	0.73909 (16)	0.59025 (14)	0.15896 (6)	0.0323 (4)
524	H32	0.825091	0.565819	0.169340	0.039*

525	C33	0.86940 (14)	0.13604 (11)	0.29145 (5)	0.0213 (3)	
526	C34	0.95353 (15)	0.13496 (11)	0.25326 (5)	0.0233 (3)	
527	H34	0.982623	0.195715	0.240408	0.028*	
528	C35	0.99553 (16)	0.04640 (12)	0.23368 (6)	0.0274 (3)	
529	H35	1.052341	0.047091	0.207570	0.033*	
530	C36	0.95451 (17)	−0.04261 (12)	0.25227 (7)	0.0313 (3)	
531	H36	0.982821	−0.103132	0.238932	0.038*	
532	C37	0.87209 (17)	−0.04298 (12)	0.29037 (7)	0.0309 (3)	
533	H37	0.844492	−0.104022	0.303371	0.037*	
534	C38	0.82924 (16)	0.04546 (11)	0.30982 (6)	0.0263 (3)	
535	H38	0.772140	0.044189	0.335840	0.032*	
536	C39	0.85297 (15)	0.32409 (12)	0.53006 (5)	0.0237 (3)	
537	C40	0.78919 (17)	0.23834 (13)	0.54385 (6)	0.0313 (3)	
538	H40	0.720053	0.210821	0.522050	0.038*	
539	C41	0.8254 (2)	0.19268 (15)	0.58903 (7)	0.0394 (4)	
540	H41	0.782257	0.133703	0.597723	0.047*	
541	C42	0.92487 (19)	0.23337 (16)	0.62148 (6)	0.0419 (5)	
542	H42	0.949622	0.202435	0.652456	0.050*	
543	C43	0.98729 (17)	0.31836 (16)	0.60868 (6)	0.0381 (4)	
544	H43	1.054483	0.346547	0.631109	0.046*	
545	C44	0.95294 (15)	0.36380 (13)	0.56300 (6)	0.0291 (3)	
546	H44	0.997849	0.422012	0.554307	0.035*	
547	N5	0.50664 (13)	0.40837 (10)	0.36913 (5)	0.0271 (3)	
548	H5A	0.480 (2)	0.4337 (14)	0.3967 (7)	0.041*	
549	H5B	0.509 (2)	0.3447 (11)	0.3761 (8)	0.041*	
550	C45	0.40746 (17)	0.42549 (15)	0.32609 (7)	0.0385 (4)	
551	H45A	0.329199	0.383047	0.329539	0.046*	
552	H45B	0.446349	0.405890	0.295254	0.046*	
553	C46	0.36344 (16)	0.53096 (14)	0.32166 (6)	0.0326 (4)	0.882 (12)
554	H46A	0.442264	0.574411	0.322569	0.039*	0.882 (12)
555	H46B	0.311072	0.540812	0.289330	0.039*	0.882 (12)
556	N6	0.2830 (4)	0.5585 (2)	0.36180 (8)	0.0361 (8)	0.882 (12)
557	H6A	0.2073 (19)	0.5235 (18)	0.3589 (9)	0.054*	0.882 (12)
558	H6B	0.257 (3)	0.6185 (13)	0.3606 (9)	0.054*	0.882 (12)
559	C46B	0.36344 (16)	0.53096 (14)	0.32166 (6)	0.0326 (4)	0.118 (12)
560	H46C	0.428985	0.567714	0.303754	0.039*	0.118 (12)
561	H46D	0.277978	0.532993	0.300800	0.039*	0.118 (12)
562	N6B	0.346 (3)	0.5839 (15)	0.3685 (6)	0.046 (5)	0.118 (12)
563	H6C	0.305 (16)	0.640 (6)	0.3622 (17)	0.068*	0.118 (12)
564	H6D	0.425 (5)	0.604 (12)	0.382 (4)	0.068*	0.118 (12)
565	O3	0.0708 (14)	0.4152 (11)	0.3732 (6)	0.070 (4)	0.136 (4)
566	H3O	−0.010618	0.402558	0.371916	0.084*	0.136 (4)
567	C47	0.1402 (18)	0.3301 (15)	0.3759 (7)	0.060 (5)	0.136 (4)
568	H47A	0.117547	0.291039	0.345850	0.072*	0.136 (4)
569	H47B	0.118247	0.292480	0.405043	0.072*	0.136 (4)
570	H47C	0.235366	0.344884	0.378874	0.072*	0.136 (4)

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

572		U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
573	Zn1	0.02239 (11)	0.01794 (11)	0.01494 (10)	0.00134 (6)	0.00332 (7)	−0.00039 (6)
574	O1	0.0343 (6)	0.0355 (6)	0.0240 (6)	0.0011 (5)	0.0097 (4)	0.0021 (5)
575	O2	0.0390 (6)	0.0310 (6)	0.0299 (6)	0.0063 (5)	0.0107 (5)	−0.0054 (5)
576	N1	0.0240 (6)	0.0202 (6)	0.0175 (6)	0.0000 (4)	0.0033 (4)	−0.0014 (4)
577	N2	0.0242 (6)	0.0190 (6)	0.0179 (6)	0.0007 (4)	0.0041 (4)	−0.0005 (4)
578	N3	0.0241 (6)	0.0187 (6)	0.0153 (5)	0.0017 (4)	0.0038 (4)	0.0000 (4)
579	N4	0.0235 (6)	0.0220 (6)	0.0148 (5)	0.0035 (4)	0.0027 (4)	−0.0006 (4)
580	C1	0.0224 (7)	0.0249 (7)	0.0161 (6)	−0.0013 (5)	0.0029 (5)	−0.0024 (5)
581	C2	0.0277 (7)	0.0281 (8)	0.0182 (7)	0.0010 (6)	0.0019 (5)	−0.0036 (6)
582	C3	0.0332 (8)	0.0245 (7)	0.0191 (7)	−0.0008 (6)	0.0021 (6)	−0.0044 (6)
583	C4	0.0229 (6)	0.0229 (7)	0.0196 (7)	−0.0010 (5)	0.0032 (5)	−0.0048 (5)
584	C5	0.0238 (7)	0.0202 (7)	0.0226 (7)	0.0005 (5)	0.0040 (5)	−0.0036 (5)
585	C6	0.0230 (7)	0.0184 (7)	0.0233 (7)	−0.0002 (5)	0.0044 (5)	−0.0012 (5)
586	C7	0.0372 (8)	0.0170 (7)	0.0282 (8)	0.0009 (6)	0.0059 (6)	0.0012 (6)
587	C8	0.0368 (8)	0.0211 (7)	0.0249 (7)	0.0012 (6)	0.0055 (6)	0.0042 (6)
588	C9	0.0251 (7)	0.0203 (7)	0.0198 (7)	0.0012 (5)	0.0045 (5)	0.0031 (5)
589	C10	0.0229 (7)	0.0230 (7)	0.0182 (7)	0.0014 (5)	0.0038 (5)	0.0024 (5)
590	C11	0.0239 (7)	0.0220 (7)	0.0164 (6)	0.0013 (5)	0.0027 (5)	0.0008 (5)
591	C12	0.0329 (8)	0.0270 (8)	0.0163 (6)	0.0017 (6)	0.0003 (5)	−0.0013 (6)
592	C13	0.0322 (8)	0.0224 (7)	0.0190 (7)	0.0008 (6)	0.0008 (6)	−0.0042 (5)
593	C14	0.0229 (6)	0.0203 (7)	0.0172 (6)	−0.0002 (5)	0.0044 (5)	−0.0020 (5)
594	C15	0.0223 (6)	0.0203 (7)	0.0193 (7)	0.0012 (5)	0.0040 (5)	−0.0015 (5)
595	C16	0.0244 (7)	0.0225 (7)	0.0198 (7)	0.0047 (5)	0.0038 (5)	−0.0003 (5)
596	C17	0.0366 (8)	0.0292 (8)	0.0223 (7)	0.0136 (6)	0.0008 (6)	−0.0007 (6)
597	C18	0.0377 (8)	0.0332 (9)	0.0197 (7)	0.0148 (7)	−0.0009 (6)	0.0018 (6)
598	C19	0.0254 (7)	0.0247 (7)	0.0179 (7)	0.0034 (6)	0.0015 (5)	0.0000 (5)
599	C20	0.0241 (7)	0.0262 (7)	0.0162 (6)	0.0008 (5)	0.0021 (5)	−0.0002 (5)
600	C21	0.0355 (8)	0.0209 (7)	0.0210 (7)	0.0014 (6)	0.0048 (6)	−0.0030 (5)
601	C22	0.0375 (9)	0.0282 (9)	0.0377 (9)	0.0053 (7)	0.0059 (7)	−0.0034 (7)
602	C23	0.0566 (12)	0.0319 (10)	0.0513 (12)	0.0176 (9)	0.0074 (9)	−0.0057 (8)
603	C24	0.0779 (15)	0.0224 (9)	0.0467 (11)	0.0083 (9)	0.0025 (10)	−0.0114 (8)
604	C25	0.0609 (13)	0.0278 (9)	0.0529 (12)	−0.0050 (8)	−0.0042 (10)	−0.0136 (8)
605	C26	0.0414 (9)	0.0290 (9)	0.0397 (10)	0.0009 (7)	−0.0015 (7)	−0.0101 (7)
606	C27	0.0290 (7)	0.0188 (7)	0.0188 (7)	0.0006 (5)	0.0023 (5)	0.0021 (5)
607	C28	0.0280 (7)	0.0274 (8)	0.0261 (7)	0.0034 (6)	0.0027 (6)	0.0002 (6)
608	C29	0.0342 (8)	0.0270 (8)	0.0286 (8)	0.0055 (6)	−0.0041 (6)	0.0016 (6)
609	C30	0.0430 (9)	0.0290 (8)	0.0219 (7)	−0.0030 (7)	−0.0037 (6)	0.0071 (6)
610	C31	0.0352 (9)	0.0543 (11)	0.0239 (8)	−0.0044 (8)	0.0059 (6)	0.0111 (7)
611	C32	0.0269 (8)	0.0463 (10)	0.0240 (8)	0.0030 (7)	0.0039 (6)	0.0084 (7)
612	C33	0.0238 (7)	0.0219 (7)	0.0179 (6)	0.0036 (5)	−0.0002 (5)	−0.0017 (5)
613	C34	0.0266 (7)	0.0226 (7)	0.0207 (7)	0.0018 (5)	0.0032 (5)	−0.0014 (5)
614	C35	0.0271 (7)	0.0319 (8)	0.0236 (7)	0.0058 (6)	0.0044 (6)	−0.0044 (6)
615	C36	0.0329 (8)	0.0237 (8)	0.0369 (9)	0.0081 (6)	0.0015 (7)	−0.0072 (6)
616	C37	0.0329 (8)	0.0208 (8)	0.0390 (9)	0.0026 (6)	0.0036 (7)	0.0029 (6)
617	C38	0.0275 (7)	0.0250 (8)	0.0270 (8)	0.0037 (6)	0.0058 (6)	0.0022 (6)

618	C39	0.0260 (7)	0.0289 (8)	0.0164 (6)	0.0067 (6)	0.0038 (5)	−0.0004 (6)
619	C40	0.0390 (9)	0.0319 (9)	0.0230 (7)	0.0024 (7)	0.0037 (6)	0.0014 (6)
620	C41	0.0496 (10)	0.0411 (10)	0.0290 (8)	0.0103 (8)	0.0120 (7)	0.0109 (7)
621	C42	0.0438 (10)	0.0636 (13)	0.0191 (8)	0.0266 (9)	0.0067 (7)	0.0104 (8)
622	C43	0.0298 (8)	0.0623 (12)	0.0212 (8)	0.0162 (8)	−0.0031 (6)	−0.0050 (8)
623	C44	0.0247 (7)	0.0396 (9)	0.0232 (7)	0.0067 (6)	0.0024 (6)	−0.0034 (6)
624	N5	0.0246 (6)	0.0233 (7)	0.0341 (7)	0.0007 (5)	0.0062 (5)	0.0004 (5)
625	C45	0.0281 (8)	0.0444 (10)	0.0419 (10)	−0.0002 (7)	−0.0037 (7)	−0.0157 (8)
626	C46	0.0270 (8)	0.0478 (10)	0.0229 (8)	0.0033 (7)	0.0011 (6)	0.0014 (7)
627	N6	0.0404 (17)	0.0460 (14)	0.0224 (9)	0.0156 (12)	0.0061 (10)	0.0046 (8)
628	C46B	0.0270 (8)	0.0478 (10)	0.0229 (8)	0.0033 (7)	0.0011 (6)	0.0014 (7)
629	N6B	0.035 (12)	0.049 (10)	0.052 (10)	−0.002 (8)	0.002 (8)	−0.006 (7)
630	O3	0.064 (8)	0.067 (9)	0.078 (10)	−0.017 (7)	−0.003 (7)	0.016 (7)
631	C47	0.054 (10)	0.072 (12)	0.056 (10)	−0.017 (9)	0.011 (8)	−0.011 (9)

632 *Geometric parameters (Å, °)*

633	Zn1—N2	2.0556 (12)	C25—H25	0.9500
634	Zn1—N4	2.0660 (12)	C26—H26	0.9500
635	Zn1—N3	2.0812 (11)	C27—C32	1.394 (2)
636	Zn1—N5	2.1315 (13)	C27—C28	1.395 (2)
637	Zn1—N1	2.1399 (12)	C28—C29	1.391 (2)
638	O1—C2	1.4294 (19)	C28—H28	0.9500
639	O1—H1	0.99 (2)	C29—C30	1.382 (3)
640	O2—C3	1.4204 (19)	C29—H29	0.9500
641	O2—H2A	0.99 (3)	C30—C31	1.392 (3)
642	N1—C1	1.3593 (19)	C30—H30	0.9500
643	N1—C4	1.3618 (19)	C31—C32	1.389 (2)
644	N2—C6	1.3661 (18)	C31—H31	0.9500
645	N2—C9	1.3865 (18)	C32—H32	0.9500
646	N3—C14	1.3716 (18)	C33—C34	1.397 (2)
647	N3—C11	1.3731 (18)	C33—C38	1.397 (2)
648	N4—C19	1.3654 (18)	C34—C35	1.393 (2)
649	N4—C16	1.3863 (18)	C34—H34	0.9500
650	C1—C20	1.382 (2)	C35—C36	1.384 (2)
651	C1—C2	1.5257 (19)	C35—H35	0.9500
652	C2—C3	1.544 (2)	C36—C37	1.383 (3)
653	C2—H2	1.0000	C36—H36	0.9500
654	C3—C4	1.5292 (19)	C37—C38	1.393 (2)
655	C3—H3	1.0000	C37—H37	0.9500
656	C4—C5	1.383 (2)	C38—H38	0.9500
657	C5—C6	1.417 (2)	C39—C44	1.395 (2)
658	C5—C21	1.500 (2)	C39—C40	1.395 (2)
659	C6—C7	1.442 (2)	C40—C41	1.389 (2)
660	C7—C8	1.356 (2)	C40—H40	0.9500
661	C7—H7	0.9500	C41—C42	1.389 (3)
662	C8—C9	1.438 (2)	C41—H41	0.9500
663	C8—H8	0.9500	C42—C43	1.372 (3)

664	C9—C10	1.394 (2)	C42—H42	0.9500
665	C10—C11	1.411 (2)	C43—C44	1.396 (2)
666	C10—C27	1.4990 (19)	C43—H43	0.9500
667	C11—C12	1.434 (2)	C44—H44	0.9500
668	C12—C13	1.359 (2)	N5—C45	1.486 (2)
669	C12—H12	0.9500	N5—H5A	0.884 (15)
670	C13—C14	1.4364 (19)	N5—H5B	0.882 (15)
671	C13—H13	0.9500	C45—C46B	1.498 (3)
672	C14—C15	1.414 (2)	C45—C46	1.498 (3)
673	C15—C16	1.401 (2)	C45—H45A	0.9900
674	C15—C33	1.4960 (19)	C45—H45B	0.9900
675	C16—C17	1.439 (2)	C46—N6	1.463 (2)
676	C17—C18	1.354 (2)	C46—H46A	0.9900
677	C17—H17	0.9500	C46—H46B	0.9900
678	C18—C19	1.441 (2)	N6—H6A	0.898 (16)
679	C18—H18	0.9500	N6—H6B	0.853 (16)
680	C19—C20	1.421 (2)	C46B—N6B	1.479 (14)
681	C20—C39	1.5001 (19)	C46B—H46C	0.9900
682	C21—C22	1.385 (2)	C46B—H46D	0.9900
683	C21—C26	1.387 (2)	N6B—H6C	0.88 (2)
684	C22—C23	1.383 (3)	N6B—H6D	0.89 (2)
685	C22—H22	0.9500	O3—C47	1.35 (2)
686	C23—C24	1.382 (3)	O3—H3O	0.8400
687	C23—H23	0.9500	C47—H47A	0.9800
688	C24—C25	1.372 (3)	C47—H47B	0.9800
689	C24—H24	0.9500	C47—H47C	0.9800
690	C25—C26	1.394 (3)		
691				
692	N2—Zn1—N4	155.81 (5)	C25—C24—H24	120.1
693	N2—Zn1—N3	88.38 (5)	C23—C24—H24	120.1
694	N4—Zn1—N3	87.85 (5)	C24—C25—C26	120.1 (2)
695	N2—Zn1—N5	102.59 (5)	C24—C25—H25	120.0
696	N4—Zn1—N5	101.54 (5)	C26—C25—H25	120.0
697	N3—Zn1—N5	102.84 (5)	C21—C26—C25	120.47 (18)
698	N2—Zn1—N1	88.29 (5)	C21—C26—H26	119.8
699	N4—Zn1—N1	88.25 (5)	C25—C26—H26	119.8
700	N3—Zn1—N1	162.66 (5)	C32—C27—C28	118.52 (14)
701	N5—Zn1—N1	94.49 (5)	C32—C27—C10	120.79 (13)
702	C2—O1—H1	109.5	C28—C27—C10	120.69 (13)
703	C3—O2—H2A	109.5	C29—C28—C27	120.48 (15)
704	C1—N1—C4	110.22 (12)	C29—C28—H28	119.8
705	C1—N1—Zn1	124.07 (10)	C27—C28—H28	119.8
706	C4—N1—Zn1	124.07 (9)	C30—C29—C28	120.72 (15)
707	C6—N2—C9	106.70 (12)	C30—C29—H29	119.6
708	C6—N2—Zn1	126.63 (10)	C28—C29—H29	119.6
709	C9—N2—Zn1	126.20 (10)	C29—C30—C31	119.17 (15)
710	C14—N3—C11	106.58 (11)	C29—C30—H30	120.4
711	C14—N3—Zn1	125.90 (9)	C31—C30—H30	120.4

712	C11—N3—Zn1	124.72 (9)	C32—C31—C30	120.32 (16)
713	C19—N4—C16	106.72 (12)	C32—C31—H31	119.8
714	C19—N4—Zn1	126.30 (10)	C30—C31—H31	119.8
715	C16—N4—Zn1	126.97 (9)	C31—C32—C27	120.76 (15)
716	N1—C1—C20	125.66 (13)	C31—C32—H32	119.6
717	N1—C1—C2	111.54 (12)	C27—C32—H32	119.6
718	C20—C1—C2	122.79 (13)	C34—C33—C38	118.04 (13)
719	O1—C2—C1	109.69 (12)	C34—C33—C15	120.58 (13)
720	O1—C2—C3	112.41 (12)	C38—C33—C15	121.37 (13)
721	C1—C2—C3	103.15 (12)	C35—C34—C33	121.16 (14)
722	O1—C2—H2	110.5	C35—C34—H34	119.4
723	C1—C2—H2	110.5	C33—C34—H34	119.4
724	C3—C2—H2	110.5	C36—C35—C34	119.99 (15)
725	O2—C3—C4	113.06 (12)	C36—C35—H35	120.0
726	O2—C3—C2	114.25 (13)	C34—C35—H35	120.0
727	C4—C3—C2	102.83 (12)	C37—C36—C35	119.65 (14)
728	O2—C3—H3	108.8	C37—C36—H36	120.2
729	C4—C3—H3	108.8	C35—C36—H36	120.2
730	C2—C3—H3	108.8	C36—C37—C38	120.50 (15)
731	N1—C4—C5	125.69 (13)	C36—C37—H37	119.7
732	N1—C4—C3	111.67 (12)	C38—C37—H37	119.7
733	C5—C4—C3	122.63 (13)	C37—C38—C33	120.65 (15)
734	C4—C5—C6	125.81 (13)	C37—C38—H38	119.7
735	C4—C5—C21	118.21 (13)	C33—C38—H38	119.7
736	C6—C5—C21	115.86 (13)	C44—C39—C40	118.47 (14)
737	N2—C6—C5	126.18 (13)	C44—C39—C20	121.42 (14)
738	N2—C6—C7	109.70 (12)	C40—C39—C20	120.03 (14)
739	C5—C6—C7	124.10 (13)	C41—C40—C39	120.89 (17)
740	C8—C7—C6	107.13 (13)	C41—C40—H40	119.6
741	C8—C7—H7	126.4	C39—C40—H40	119.6
742	C6—C7—H7	126.4	C42—C41—C40	119.91 (18)
743	C7—C8—C9	107.28 (13)	C42—C41—H41	120.0
744	C7—C8—H8	126.4	C40—C41—H41	120.0
745	C9—C8—H8	126.4	C43—C42—C41	119.83 (16)
746	N2—C9—C10	125.87 (13)	C43—C42—H42	120.1
747	N2—C9—C8	109.15 (13)	C41—C42—H42	120.1
748	C10—C9—C8	124.93 (13)	C42—C43—C44	120.60 (17)
749	C9—C10—C11	125.20 (13)	C42—C43—H43	119.7
750	C9—C10—C27	117.69 (13)	C44—C43—H43	119.7
751	C11—C10—C27	117.08 (13)	C39—C44—C43	120.28 (17)
752	N3—C11—C10	124.71 (13)	C39—C44—H44	119.9
753	N3—C11—C12	109.73 (12)	C43—C44—H44	119.9
754	C10—C11—C12	125.48 (13)	C45—N5—Zn1	117.93 (11)
755	C13—C12—C11	106.91 (13)	C45—N5—H5A	111.4 (15)
756	C13—C12—H12	126.5	Zn1—N5—H5A	110.1 (15)
757	C11—C12—H12	126.5	C45—N5—H5B	108.8 (14)
758	C12—C13—C14	107.17 (13)	Zn1—N5—H5B	105.3 (14)
759	C12—C13—H13	126.4	H5A—N5—H5B	101.8 (18)

760	C14—C13—H13	126.4	N5—C45—C46B	112.77 (14)
761	N3—C14—C15	124.61 (12)	N5—C45—C46	112.77 (14)
762	N3—C14—C13	109.49 (12)	N5—C45—H45A	109.0
763	C15—C14—C13	125.79 (13)	C46—C45—H45A	109.0
764	C16—C15—C14	124.48 (13)	N5—C45—H45B	109.0
765	C16—C15—C33	117.67 (13)	C46—C45—H45B	109.0
766	C14—C15—C33	117.82 (12)	H45A—C45—H45B	107.8
767	N4—C16—C15	125.85 (13)	N6—C46—C45	111.44 (17)
768	N4—C16—C17	109.14 (12)	N6—C46—H46A	109.3
769	C15—C16—C17	124.99 (13)	C45—C46—H46A	109.3
770	C18—C17—C16	107.17 (13)	N6—C46—H46B	109.3
771	C18—C17—H17	126.4	C45—C46—H46B	109.3
772	C16—C17—H17	126.4	H46A—C46—H46B	108.0
773	C17—C18—C19	107.31 (13)	C46—N6—H6A	109.2 (15)
774	C17—C18—H18	126.3	C46—N6—H6B	114.0 (16)
775	C19—C18—H18	126.3	H6A—N6—H6B	104 (2)
776	N4—C19—C20	126.10 (13)	N6B—C46B—C45	116.8 (7)
777	N4—C19—C18	109.61 (13)	N6B—C46B—H46C	108.1
778	C20—C19—C18	124.29 (13)	C45—C46B—H46C	108.1
779	C1—C20—C19	125.69 (13)	N6B—C46B—H46D	108.1
780	C1—C20—C39	119.08 (13)	C45—C46B—H46D	108.1
781	C19—C20—C39	115.23 (13)	H46C—C46B—H46D	107.3
782	C22—C21—C26	118.74 (15)	C46B—N6B—H6C	110 (3)
783	C22—C21—C5	121.41 (15)	C46B—N6B—H6D	109 (3)
784	C26—C21—C5	119.76 (14)	H6C—N6B—H6D	102 (4)
785	C23—C22—C21	120.74 (18)	C47—O3—H3O	109.5
786	C23—C22—H22	119.6	O3—C47—H47A	109.5
787	C21—C22—H22	119.6	O3—C47—H47B	109.5
788	C24—C23—C22	120.11 (19)	H47A—C47—H47B	109.5
789	C24—C23—H23	119.9	O3—C47—H47C	109.5
790	C22—C23—H23	119.9	H47A—C47—H47C	109.5
791	C25—C24—C23	119.88 (17)	H47B—C47—H47C	109.5
792				
793	C4—N1—C1—C20	−172.69 (14)	C14—C15—C16—C17	−173.90 (15)
794	Zn1—N1—C1—C20	21.5 (2)	C33—C15—C16—C17	4.1 (2)
795	C4—N1—C1—C2	8.29 (16)	N4—C16—C17—C18	1.83 (19)
796	Zn1—N1—C1—C2	−157.56 (10)	C15—C16—C17—C18	−176.45 (15)
797	N1—C1—C2—O1	113.02 (14)	C16—C17—C18—C19	−0.5 (2)
798	C20—C1—C2—O1	−66.03 (18)	C16—N4—C19—C20	−177.20 (14)
799	N1—C1—C2—C3	−6.94 (16)	Zn1—N4—C19—C20	3.4 (2)
800	C20—C1—C2—C3	174.00 (14)	C16—N4—C19—C18	2.20 (17)
801	O1—C2—C3—O2	7.86 (17)	Zn1—N4—C19—C18	−177.24 (11)
802	C1—C2—C3—O2	125.94 (13)	C17—C18—C19—N4	−1.09 (19)
803	O1—C2—C3—C4	−115.04 (13)	C17—C18—C19—C20	178.32 (15)
804	C1—C2—C3—C4	3.04 (15)	N1—C1—C20—C19	−3.6 (2)
805	C1—N1—C4—C5	173.14 (14)	C2—C1—C20—C19	175.28 (14)
806	Zn1—N1—C4—C5	−21.0 (2)	N1—C1—C20—C39	175.45 (13)
807	C1—N1—C4—C3	−6.09 (17)	C2—C1—C20—C39	−5.6 (2)

808	Zn1—N1—C4—C3	159.75 (10)	N4—C19—C20—C1	−10.3 (2)
809	O2—C3—C4—N1	−122.20 (14)	C18—C19—C20—C1	170.36 (15)
810	C2—C3—C4—N1	1.50 (16)	N4—C19—C20—C39	170.55 (14)
811	O2—C3—C4—C5	58.54 (19)	C18—C19—C20—C39	−8.8 (2)
812	C2—C3—C4—C5	−177.76 (13)	C4—C5—C21—C22	−108.81 (18)
813	N1—C4—C5—C6	7.8 (2)	C6—C5—C21—C22	74.97 (19)
814	C3—C4—C5—C6	−173.08 (14)	C4—C5—C21—C26	74.7 (2)
815	N1—C4—C5—C21	−168.03 (14)	C6—C5—C21—C26	−101.57 (18)
816	C3—C4—C5—C21	11.1 (2)	C26—C21—C22—C23	0.7 (3)
817	C9—N2—C6—C5	176.68 (14)	C5—C21—C22—C23	−175.87 (17)
818	Zn1—N2—C6—C5	4.2 (2)	C21—C22—C23—C24	−0.1 (3)
819	C9—N2—C6—C7	−1.77 (16)	C22—C23—C24—C25	−0.6 (3)
820	Zn1—N2—C6—C7	−174.29 (10)	C23—C24—C25—C26	0.7 (4)
821	C4—C5—C6—N2	1.8 (2)	C22—C21—C26—C25	−0.6 (3)
822	C21—C5—C6—N2	177.74 (14)	C5—C21—C26—C25	176.01 (18)
823	C4—C5—C6—C7	−179.92 (15)	C24—C25—C26—C21	−0.1 (3)
824	C21—C5—C6—C7	−4.0 (2)	C9—C10—C27—C32	110.01 (18)
825	N2—C6—C7—C8	0.94 (18)	C11—C10—C27—C32	−68.0 (2)
826	C5—C6—C7—C8	−177.54 (15)	C9—C10—C27—C28	−69.78 (19)
827	C6—C7—C8—C9	0.26 (18)	C11—C10—C27—C28	112.19 (17)
828	C6—N2—C9—C10	−175.80 (14)	C32—C27—C28—C29	0.5 (2)
829	Zn1—N2—C9—C10	−3.2 (2)	C10—C27—C28—C29	−179.74 (14)
830	C6—N2—C9—C8	1.93 (16)	C27—C28—C29—C30	−1.6 (2)
831	Zn1—N2—C9—C8	174.49 (10)	C28—C29—C30—C31	1.3 (3)
832	C7—C8—C9—N2	−1.36 (18)	C29—C30—C31—C32	0.1 (3)
833	C7—C8—C9—C10	176.39 (15)	C30—C31—C32—C27	−1.2 (3)
834	N2—C9—C10—C11	−5.4 (2)	C28—C27—C32—C31	0.9 (3)
835	C8—C9—C10—C11	177.25 (15)	C10—C27—C32—C31	−178.87 (16)
836	N2—C9—C10—C27	176.78 (13)	C16—C15—C33—C34	−123.99 (15)
837	C8—C9—C10—C27	−0.6 (2)	C14—C15—C33—C34	54.12 (19)
838	C14—N3—C11—C10	−173.52 (14)	C16—C15—C33—C38	55.03 (19)
839	Zn1—N3—C11—C10	24.5 (2)	C14—C15—C33—C38	−126.86 (15)
840	C14—N3—C11—C12	3.29 (16)	C38—C33—C34—C35	0.5 (2)
841	Zn1—N3—C11—C12	−158.64 (10)	C15—C33—C34—C35	179.57 (14)
842	C9—C10—C11—N3	−6.2 (2)	C33—C34—C35—C36	−0.4 (2)
843	C27—C10—C11—N3	171.62 (13)	C34—C35—C36—C37	−0.2 (3)
844	C9—C10—C11—C12	177.44 (15)	C35—C36—C37—C38	0.6 (3)
845	C27—C10—C11—C12	−4.7 (2)	C36—C37—C38—C33	−0.4 (3)
846	N3—C11—C12—C13	−1.97 (17)	C34—C33—C38—C37	−0.1 (2)
847	C10—C11—C12—C13	174.82 (15)	C15—C33—C38—C37	−179.16 (14)
848	C11—C12—C13—C14	−0.14 (17)	C1—C20—C39—C44	−65.50 (19)
849	C11—N3—C14—C15	173.12 (13)	C19—C20—C39—C44	113.68 (16)
850	Zn1—N3—C14—C15	−25.2 (2)	C1—C20—C39—C40	117.69 (17)
851	C11—N3—C14—C13	−3.37 (16)	C19—C20—C39—C40	−63.13 (19)
852	Zn1—N3—C14—C13	158.29 (10)	C44—C39—C40—C41	−0.9 (2)
853	C12—C13—C14—N3	2.20 (17)	C20—C39—C40—C41	176.05 (15)
854	C12—C13—C14—C15	−174.24 (14)	C39—C40—C41—C42	1.1 (3)
855	N3—C14—C15—C16	7.9 (2)	C40—C41—C42—C43	−0.2 (3)

856	C13—C14—C15—C16	−176.19 (14)	C41—C42—C43—C44	−0.8 (3)
857	N3—C14—C15—C33	−170.08 (13)	C40—C39—C44—C43	−0.2 (2)
858	C13—C14—C15—C33	5.8 (2)	C20—C39—C44—C43	−177.06 (14)
859	C19—N4—C16—C15	175.79 (14)	C42—C43—C44—C39	1.0 (2)
860	Zn1—N4—C16—C15	−4.8 (2)	Zn1—N5—C45—C46B	−78.92 (16)
861	C19—N4—C16—C17	−2.47 (17)	Zn1—N5—C45—C46	−78.92 (16)
862	Zn1—N4—C16—C17	176.96 (11)	N5—C45—C46—N6	−69.5 (3)
863	C14—C15—C16—N4	8.1 (2)	N5—C45—C46B—N6B	−38.3 (15)
864	C33—C15—C16—N4	−173.93 (13)		

865 *Hydrogen-bond geometry (Å, °)*

866	<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>
867	O1—H1 $\cdots$ N6 ⁱ	0.99	1.73	2.710 (3)	168
868	O1—H1 $\cdots$ N6B ⁱ	0.99	1.54	2.510 (17)	165
869	O2—H2A $\cdots$ O1 ⁱ	0.99	1.82	2.8056 (18)	171
870	C2—H2 $\cdots$ O3 ⁱ	1.00	2.53	3.460 (14)	155
871	N5—H5A $\cdots$ O1 ⁱ	0.88 (2)	2.38 (2)	3.2442 (18)	166 (2)
872	C46—H46A $\cdots$ N2	0.99	2.49	3.368 (2)	148
873	N6—H6A $\cdots$ O3	0.90 (2)	2.08 (2)	2.932 (14)	159 (3)
874	C46B—H46C $\cdots$ N2	0.99	2.68	3.368 (2)	126
875	O3—H3O $\cdots$ N4 ⁱⁱ	0.84	2.20	2.992 (14)	157

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