

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

4,4'-Bipyridyl-4,4'-(hydroxymethylene)-dibenzoic acid (1/1)

Lan Qin, Lan-Ping Xu and Lei Han*

 Faculty of Materials Science & Chemical Engineering, Ningbo University, Ningbo, Zhejiang 315211, People's Republic of China
 Correspondence e-mail: hanlei@nbu.edu.cn

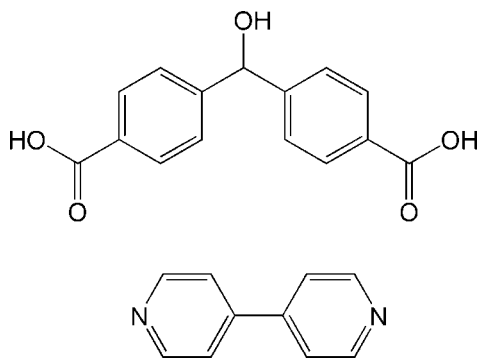
Received 19 March 2012; accepted 20 March 2012

 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.077; wR factor = 0.210; data-to-parameter ratio = 12.8.

In the title 1:1 co-crystal, $\text{C}_{10}\text{H}_8\text{N}_2 \cdot \text{C}_{15}\text{H}_{12}\text{O}_5$, strong intermolecular $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds link alternating molecules of 4,4'-(hydroxymethylene)dibenzoic acid and 4,4'-bipyridyl into zigzag chains in [501]. The crystal packing also exhibits $\pi-\pi$ interactions between the 4,4'-bipyridyl rings of neighbouring chains [centroid-centroid distance = 3.608 (3) Å] and weak $\text{C}-\text{H} \cdots \text{O}$ hydrogen bonds.

Related literature

For background to supramolecular crystal engineering, see: Simon & Bassoul (2000). For aromatic carboxylic acids as supramolecular synthons, see: Desiraju (1995). For studies of bent arenedicarboxylate ligands, see: Koichi *et al.* (2011); Xu *et al.* (2011).



Experimental

Crystal data

 $\text{C}_{10}\text{H}_8\text{N}_2 \cdot \text{C}_{15}\text{H}_{12}\text{O}_5$
 $M_r = 428.43$

 Monoclinic, $P2_1/n$
 $a = 8.0528$ (16) Å
 $b = 11.683$ (2) Å
 $c = 21.922$ (4) Å
 $\beta = 96.66$ (3)°
 $V = 2048.6$ (7) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹
 $T = 298$ K
 $0.31 \times 0.14 \times 0.12$ mm

Data collection

 Rigaku R-AXIS RAPID
 diffractometer
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.984$, $T_{\max} = 0.988$

 16322 measured reflections
 3800 independent reflections
 1903 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.089$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.077$
 $wR(F^2) = 0.210$
 $S = 0.99$
 3800 reflections
 298 parameters

 H atoms treated by a mixture of
 independent and constrained
 refinement
 $\Delta\rho_{\text{max}} = 0.68$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{O2}-\text{H2} \cdots \text{N2}^{\text{i}}$	0.84 (5)	1.82 (5)	2.660 (5)	173 (5)
$\text{O4}-\text{H1} \cdots \text{N1}^{\text{ii}}$	0.87 (6)	1.76 (6)	2.605 (5)	166 (5)
$\text{C19}-\text{H19A} \cdots \text{O3}^{\text{iii}}$	0.93	2.40	3.321 (5)	171
$\text{C17}-\text{H17A} \cdots \text{O1}^{\text{iv}}$	0.93	2.57	3.212 (5)	126
$\text{C8}-\text{H8A} \cdots \text{O3}^{\text{iv}}$	0.98	2.51	3.376 (6)	148

 Symmetry codes: (i) $-x + \frac{5}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x, -y, -z$; (iii) $-x + 1, -y, -z$; (iv) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

This work was supported by the National Natural Science Foundation of China (grant No. 21071087) and the K. C. Wong Magna Fund in Ningbo University.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5265).

References

- Desiraju, G. R. (1995). *Angew. Chem. Int. Ed.* **34**, 2311–2327.
 Higashi, T. (1995). *ABSCOR*. Rigaku Corporation, Tokyo, Japan.
 Koichi, K., Eriko, S. & Takuji, H. (2011). *Chem. Eur. J.* **17**, 11527–11534.
 Rigaku (1998). *RAPID-AUTO*. Rigaku Corporation, Tokyo, Japan.
 Rigaku/MS (2004). *CrystalStructure*. Rigaku/MS, The Woodlands, Texas, USA.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Simon, J. & Bassoul, P. (2000). In *Design of Molecular Materials: Supramolecular Engineering*. Berlin: Wiley-VCH.
 Xu, L.-P., Zhao, W.-N. & Han, L. (2011). *Acta Cryst.* **E67**, o1971.

supporting information

Acta Cryst. (2012). E68, o1175 [https://doi.org/10.1107/S1600536812011956]

4,4'-Bipyridyl–4,4'-(hydroxymethylene)dibenzoic acid (1/1)**Lan Qin, Lan-Ping Xu and Lei Han****S1. Comment**

Supramolecular crystal engineering has attracted growing interest over the past few decades because of their importance in biological system and molecular recognition (Simon *et al.*, 2000). Aromatic carboxylic acid is one of the most important supramolecular synthons to construct novel organic networks by hydrogen bonds and π – π interactions (Desiraju, 1995). Recently, interest has been devoted to the assembly of extended solids from the long and bent arene-dicarboxylate ligands (Koichi *et al.*, 2011; Xu *et al.*, 2011). We have employed 4,4'-(hydroxymethylene)dibenzoic acid as an excellent candidate for the construction of targeted supramolecular structures. A new organic cocrystal compound, $C_{15}H_{12}O_5 \cdot C_{10}H_8N_2$, has been synthesized by reacting 4,4'-(hydroxymethylene)dibenzoic acid and 4,4'-bipyridyl under hydrothermal conditions, and its crystal structure is reported here.

The asymmetric unit of the title compound consists of one 4,4'-(hydroxymethylene)dibenzoic acid and one 4,4'-bipyridyl molecule (Figure 1). The dihedral angle formed by two pyridine rings in 4,4'-bipyridyl is $24.74(1)^\circ$, and the dihedral angle between the two benzene rings in bent 4,4'-(hydroxymethylene)dibenzoic acid ligand is $85.95(3)^\circ$. In the 1:1 cocrystal, strong intermolecular O—H \cdots N hydrogen bonds link the alternating molecules of 4,4'-(hydroxymethylene)dibenzoic acid and 4,4'-bipyridyl into zigzag chains in [501], as shown in Figure 2. Furthermore, the crystal packing exhibits also π – π interactions between the rings of 4,4'-bipyridyl from the neighbouring chains [centroid–centroid distance of $3.608(3)$ Å] and weak C—H \cdots O hydrogen bonds.

S2. Experimental

A mixture of 4,4'-(hydroxymethylene)dibenzoic acid (26.4 mg, 0.1 mmol) and 4,4'-bipyridyl (15.1 mg, 0.1 mmol) in H₂O (8 ml) was sealed in a 25 ml Teflon-lined stainless steel reactor and heated at 447 K for 3 d. Colorless single crystals of the title compound was obtained after cooling the solution to room temperature. Block-shaped crystals were collected and washed with distilled water. The yield was approximately 70% based on 4,4'-(hydroxymethylene)dibenzoic acid.

S3. Refinement

H atoms attached to O2 and O4 were located in difference maps and refined isotropically. C-bound H atoms were positioned geometrically and allowed to ride on their respective parent atoms at distances of C—H = 0.93 Å, with $U_{iso}(H) = 1.2U_{eq}(C)$.

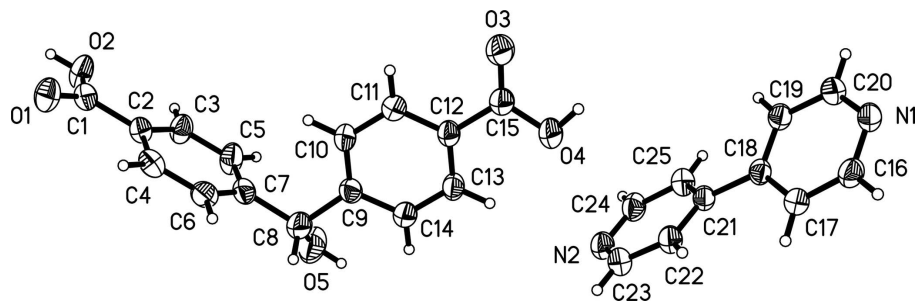


Figure 1

A content of asymmetric unit of the title compound showing the atomic numbering and 30% probability displacement ellipsoids.

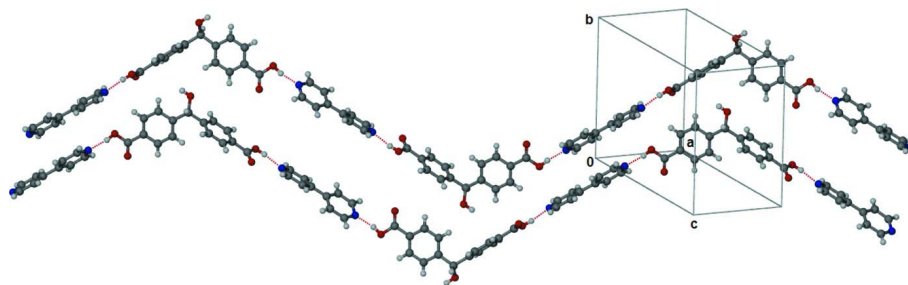


Figure 2

A portion of the crystal packing showing O—H...N hydrogen bonds as dashed lines.

4,4'-Bipyridyl-4,4'-(hydroxymethylene)dibenzoic acid (1/1)

Crystal data

$C_{10}H_8N_2 \cdot C_{15}H_{12}O_5$

$M_r = 428.43$

Monoclinic, $P2_1/n$

Hall symbol: $-P 2_1n$

$a = 8.0528 (16) \text{ \AA}$

$b = 11.683 (2) \text{ \AA}$

$c = 21.922 (4) \text{ \AA}$

$\beta = 96.66 (3)^\circ$

$V = 2048.6 (7) \text{ \AA}^3$

$Z = 4$

$F(000) = 896$

$D_x = 1.389 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 2687 reflections

$\theta = 3.1\text{--}25.5^\circ$

$\mu = 0.10 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Block, colourless

$0.31 \times 0.14 \times 0.12 \text{ mm}$

Data collection

Rigaku R-AXIS RAPID

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 0 pixels mm^{-1}

ω scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.984$, $T_{\max} = 0.988$

16322 measured reflections

3800 independent reflections

1903 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.089$

$\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 3.1^\circ$

$h = -9 \rightarrow 9$

$k = -14 \rightarrow 14$

$l = -26 \rightarrow 24$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.077$ $wR(F^2) = 0.210$ $S = 0.99$

3800 reflections

298 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sitesH atoms treated by a mixture of independent
and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0824P)^2 + 1.4919P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.68 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001xF_c^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0067 (18)

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.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F , and R-factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.7007 (4)	-0.0589 (3)	0.44854 (15)	0.0828 (10)
O2	1.8293 (4)	-0.0048 (3)	0.36999 (15)	0.0844 (11)
O3	0.6425 (4)	-0.0831 (3)	0.12982 (16)	0.0831 (10)
O4	0.5017 (4)	0.0793 (3)	0.11948 (17)	0.0863 (12)
O5	1.1910 (4)	0.3660 (3)	0.2676 (2)	0.1155 (15)
H5A	1.1098	0.4045	0.2542	0.173*
N1	-0.2656 (4)	0.0472 (3)	-0.06295 (16)	0.0628 (10)
N2	0.4192 (4)	0.3635 (3)	0.08219 (19)	0.0694 (10)
C1	1.7018 (5)	-0.0067 (4)	0.4015 (2)	0.0610 (11)
C2	1.5595 (5)	0.0650 (4)	0.37421 (18)	0.0557 (11)
C3	1.5610 (5)	0.1201 (4)	0.31932 (19)	0.0686 (13)
H3A	1.6546	0.1139	0.2984	0.082*
C4	1.4188 (5)	0.0754 (4)	0.40432 (19)	0.0633 (12)
H4A	1.4142	0.0385	0.4417	0.076*
C5	1.4258 (5)	0.1850 (5)	0.2941 (2)	0.0780 (15)
H5B	1.4302	0.2220	0.2568	0.094*
C6	1.2844 (5)	0.1406 (4)	0.3790 (2)	0.0668 (12)
H6A	1.1908	0.1475	0.3999	0.080*
C7	1.2865 (5)	0.1952 (4)	0.3238 (2)	0.0653 (12)
C8	1.1347 (5)	0.2650 (4)	0.2987 (2)	0.0784 (15)
H8A	1.0768	0.2904	0.3332	0.094*
C9	1.0110 (4)	0.1983 (4)	0.25372 (18)	0.0586 (11)
C10	1.0175 (5)	0.0821 (4)	0.24747 (19)	0.0660 (12)

H10A	1.1054	0.0415	0.2686	0.079*
C11	0.8944 (5)	0.0235 (4)	0.20995 (19)	0.0613 (11)
H11A	0.9000	-0.0557	0.2064	0.074*
C12	0.7646 (4)	0.0829 (4)	0.17821 (17)	0.0526 (10)
C13	0.7589 (5)	0.1998 (4)	0.18338 (18)	0.0608 (11)
H13A	0.6723	0.2404	0.1614	0.073*
C14	0.8806 (5)	0.2575 (4)	0.22089 (19)	0.0618 (11)
H14A	0.8752	0.3368	0.2242	0.074*
C15	0.6312 (5)	0.0179 (4)	0.1404 (2)	0.0616 (11)
C16	-0.2813 (5)	0.1129 (4)	-0.0146 (2)	0.0671 (12)
H16A	-0.3849	0.1154	0.0003	0.081*
C17	-0.1537 (5)	0.1778 (4)	0.01501 (19)	0.0594 (11)
H17A	-0.1712	0.2220	0.0489	0.071*
C18	0.0024 (4)	0.1760 (3)	-0.00674 (18)	0.0515 (10)
C19	0.0182 (5)	0.1095 (3)	-0.05812 (18)	0.0578 (11)
H19A	0.1191	0.1070	-0.0748	0.069*
C20	-0.1179 (5)	0.0466 (4)	-0.0844 (2)	0.0641 (12)
H20A	-0.1051	0.0020	-0.1187	0.077*
C21	0.1463 (5)	0.2406 (3)	0.02397 (18)	0.0526 (10)
C22	0.1556 (5)	0.2692 (4)	0.0859 (2)	0.0649 (12)
H22A	0.0711	0.2478	0.1091	0.078*
C23	0.2930 (5)	0.3303 (4)	0.1123 (2)	0.0725 (13)
H23A	0.2973	0.3492	0.1536	0.087*
C24	0.4097 (5)	0.3339 (4)	0.0235 (2)	0.0679 (13)
H24A	0.4970	0.3555	0.0017	0.082*
C25	0.2790 (5)	0.2731 (4)	-0.0073 (2)	0.0631 (12)
H25A	0.2798	0.2542	-0.0484	0.076*
H2	1.915 (6)	-0.042 (5)	0.385 (2)	0.100 (19)*
H1	0.436 (7)	0.031 (5)	0.099 (2)	0.11 (2)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.084 (2)	0.091 (3)	0.071 (2)	0.0073 (18)	0.0029 (17)	0.0258 (19)
O2	0.061 (2)	0.109 (3)	0.082 (2)	0.023 (2)	0.0036 (18)	0.026 (2)
O3	0.0630 (19)	0.078 (3)	0.105 (3)	-0.0035 (17)	-0.0046 (17)	-0.008 (2)
O4	0.062 (2)	0.083 (3)	0.106 (3)	0.0024 (18)	-0.0254 (19)	-0.017 (2)
O5	0.089 (3)	0.090 (3)	0.156 (4)	-0.001 (2)	-0.033 (2)	0.012 (3)
N1	0.053 (2)	0.064 (2)	0.068 (2)	0.0002 (17)	-0.0067 (18)	-0.0012 (19)
N2	0.054 (2)	0.068 (3)	0.083 (3)	-0.0012 (18)	-0.007 (2)	-0.005 (2)
C1	0.057 (3)	0.068 (3)	0.057 (3)	-0.005 (2)	-0.002 (2)	-0.001 (2)
C2	0.051 (2)	0.064 (3)	0.051 (2)	-0.0029 (19)	0.0005 (19)	0.000 (2)
C3	0.053 (2)	0.095 (4)	0.058 (3)	0.014 (2)	0.009 (2)	0.012 (3)
C4	0.066 (3)	0.073 (3)	0.050 (2)	-0.013 (2)	0.007 (2)	-0.004 (2)
C5	0.064 (3)	0.109 (4)	0.060 (3)	0.015 (3)	0.002 (2)	0.015 (3)
C6	0.047 (2)	0.085 (3)	0.070 (3)	-0.001 (2)	0.010 (2)	-0.020 (3)
C7	0.052 (2)	0.079 (3)	0.063 (3)	0.002 (2)	-0.007 (2)	-0.015 (2)
C8	0.063 (3)	0.074 (4)	0.094 (4)	0.007 (2)	-0.011 (3)	-0.014 (3)

C9	0.045 (2)	0.071 (3)	0.059 (2)	0.002 (2)	0.0015 (19)	-0.012 (2)
C10	0.051 (2)	0.077 (3)	0.065 (3)	0.010 (2)	-0.011 (2)	-0.007 (2)
C11	0.058 (2)	0.059 (3)	0.064 (3)	0.005 (2)	0.000 (2)	-0.002 (2)
C12	0.044 (2)	0.064 (3)	0.049 (2)	-0.0003 (19)	0.0033 (17)	0.001 (2)
C13	0.048 (2)	0.074 (3)	0.058 (2)	0.006 (2)	-0.0040 (19)	0.000 (2)
C14	0.056 (2)	0.061 (3)	0.066 (3)	0.003 (2)	-0.001 (2)	-0.003 (2)
C15	0.055 (3)	0.064 (3)	0.065 (3)	0.001 (2)	0.001 (2)	-0.001 (2)
C16	0.047 (2)	0.068 (3)	0.086 (3)	0.001 (2)	0.005 (2)	-0.002 (3)
C17	0.054 (2)	0.061 (3)	0.062 (3)	0.003 (2)	0.001 (2)	-0.002 (2)
C18	0.048 (2)	0.051 (3)	0.053 (2)	0.0009 (18)	-0.0034 (18)	0.005 (2)
C19	0.053 (2)	0.059 (3)	0.059 (3)	0.003 (2)	-0.001 (2)	0.002 (2)
C20	0.063 (3)	0.066 (3)	0.060 (3)	0.003 (2)	-0.005 (2)	-0.003 (2)
C21	0.049 (2)	0.051 (3)	0.055 (2)	0.0005 (18)	-0.0044 (19)	0.0004 (19)
C22	0.056 (2)	0.065 (3)	0.071 (3)	0.000 (2)	-0.006 (2)	-0.004 (2)
C23	0.066 (3)	0.076 (3)	0.071 (3)	0.006 (2)	-0.012 (2)	-0.009 (3)
C24	0.052 (2)	0.070 (3)	0.080 (3)	-0.003 (2)	-0.002 (2)	0.004 (3)
C25	0.057 (2)	0.069 (3)	0.062 (3)	-0.003 (2)	0.000 (2)	0.003 (2)

Geometric parameters (Å, °)

O1—C1	1.199 (5)	C9—C14	1.387 (5)
O2—C1	1.302 (5)	C10—C11	1.392 (5)
O2—H2	0.84 (5)	C10—H10A	0.9300
O3—C15	1.208 (5)	C11—C12	1.375 (5)
O4—C15	1.304 (5)	C11—H11A	0.9300
O4—H1	0.87 (6)	C12—C13	1.371 (6)
O5—C8	1.460 (6)	C12—C15	1.487 (6)
O5—H5A	0.8200	C13—C14	1.380 (5)
N1—C16	1.327 (5)	C13—H13A	0.9300
N1—C20	1.329 (5)	C14—H14A	0.9300
N2—C24	1.325 (6)	C16—C17	1.377 (5)
N2—C23	1.332 (6)	C16—H16A	0.9300
C1—C2	1.488 (6)	C17—C18	1.395 (5)
C2—C3	1.365 (5)	C17—H17A	0.9300
C2—C4	1.381 (6)	C18—C19	1.386 (5)
C3—C5	1.388 (6)	C18—C21	1.478 (5)
C3—H3A	0.9300	C19—C20	1.387 (5)
C4—C6	1.386 (6)	C19—H19A	0.9300
C4—H4A	0.9300	C20—H20A	0.9300
C5—C7	1.365 (6)	C21—C25	1.387 (6)
C5—H5B	0.9300	C21—C22	1.391 (5)
C6—C7	1.370 (6)	C22—C23	1.385 (6)
C6—H6A	0.9300	C22—H22A	0.9300
C7—C8	1.518 (6)	C23—H23A	0.9300
C8—C9	1.531 (6)	C24—C25	1.379 (6)
C8—H8A	0.9800	C24—H24A	0.9300
C9—C10	1.367 (6)	C25—H25A	0.9300

C1—O2—H2	116 (4)	C13—C12—C15	121.7 (4)
C15—O4—H1	104 (4)	C11—C12—C15	118.8 (4)
C8—O5—H5A	109.5	C12—C13—C14	120.5 (4)
C16—N1—C20	117.2 (4)	C12—C13—H13A	119.7
C24—N2—C23	116.3 (4)	C14—C13—H13A	119.7
O1—C1—O2	123.3 (4)	C13—C14—C9	120.5 (4)
O1—C1—C2	123.3 (4)	C13—C14—H14A	119.7
O2—C1—C2	113.3 (4)	C9—C14—H14A	119.7
C3—C2—C4	118.1 (4)	O3—C15—O4	123.0 (4)
C3—C2—C1	122.3 (4)	O3—C15—C12	122.7 (4)
C4—C2—C1	119.6 (4)	O4—C15—C12	114.3 (4)
C2—C3—C5	121.4 (4)	N1—C16—C17	124.0 (4)
C2—C3—H3A	119.3	N1—C16—H16A	118.0
C5—C3—H3A	119.3	C17—C16—H16A	118.0
C2—C4—C6	120.2 (4)	C16—C17—C18	118.9 (4)
C2—C4—H4A	119.9	C16—C17—H17A	120.5
C6—C4—H4A	119.9	C18—C17—H17A	120.5
C7—C5—C3	120.6 (4)	C19—C18—C17	117.2 (4)
C7—C5—H5B	119.7	C19—C18—C21	121.0 (4)
C3—C5—H5B	119.7	C17—C18—C21	121.8 (4)
C7—C6—C4	121.4 (4)	C18—C19—C20	119.5 (4)
C7—C6—H6A	119.3	C18—C19—H19A	120.3
C4—C6—H6A	119.3	C20—C19—H19A	120.3
C5—C7—C6	118.3 (4)	N1—C20—C19	123.1 (4)
C5—C7—C8	123.1 (5)	N1—C20—H20A	118.4
C6—C7—C8	118.5 (4)	C19—C20—H20A	118.4
O5—C8—C7	108.8 (4)	C25—C21—C22	117.3 (4)
O5—C8—C9	108.9 (4)	C25—C21—C18	121.6 (4)
C7—C8—C9	113.3 (4)	C22—C21—C18	121.2 (4)
O5—C8—H8A	108.6	C23—C22—C21	118.8 (4)
C7—C8—H8A	108.6	C23—C22—H22A	120.6
C9—C8—H8A	108.6	C21—C22—H22A	120.6
C10—C9—C14	118.6 (4)	N2—C23—C22	124.1 (5)
C10—C9—C8	122.7 (4)	N2—C23—H23A	117.9
C14—C9—C8	118.5 (4)	C22—C23—H23A	117.9
C9—C10—C11	121.0 (4)	N2—C24—C25	124.3 (4)
C9—C10—H10A	119.5	N2—C24—H24A	117.8
C11—C10—H10A	119.5	C25—C24—H24A	117.8
C12—C11—C10	119.9 (4)	C24—C25—C21	119.2 (4)
C12—C11—H11A	120.1	C24—C25—H25A	120.4
C10—C11—H11A	120.1	C21—C25—H25A	120.4
C13—C12—C11	119.5 (4)		

Hydrogen-bond geometry (Å, °)

<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>
O2—H2 \cdots N2 ⁱ	0.84 (5)	1.82 (5)	2.660 (5)	173 (5)
O4—H1 \cdots N1 ⁱⁱ	0.87 (6)	1.76 (6)	2.605 (5)	166 (5)

C19—H19A···O3 ⁱⁱⁱ	0.93	2.40	3.321 (5)	171
C17—H17A···O1 ^{iv}	0.93	2.57	3.212 (5)	126
C8—H8A···O3 ^{iv}	0.98	2.51	3.376 (6)	148

Symmetry codes: (i) $-x+5/2, y-1/2, -z+1/2$; (ii) $-x, -y, -z$; (iii) $-x+1, -y, -z$; (iv) $-x+3/2, y+1/2, -z+1/2$.