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3-Aminobenzoic acid–1,2-bis(4-pyridyl)ethane (1/1)

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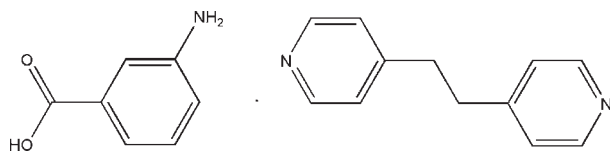
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.042; wR factor = 0.108; data-to-parameter ratio = 13.1.

The asymmetric unit of the title compound, $\text{C}_{12}\text{H}_{12}\text{N}_2 \cdot \text{C}_7\text{H}_7\text{NO}_2$, contains two 3-aminobenzoic acid molecules and two 1,2-bis(4-pyridyl)ethane molecules. In the two 1,2-bis(4-pyridyl)ethane molecules, the dihedral angles between the pyridyl rings are 2.99 (9) and 46.78 (8)°. In the crystal, the molecules associate through amine and carboxyl group $\text{N}-\text{H} \cdots \text{O}=\text{C}$ interactions between one of the 3-aminobenzoic acid molecules and one of the 1,2-bis(4-pyridyl)ethane molecules, generating $R_2^2(14)$ dimers, which are extended head-to-tail *via* amine and pyridine $\text{N}-\text{H} \cdots \text{N}$ hydrogen bonds. Intermolecular $\text{O}-\text{H} \cdots \text{N}$, $\text{N}-\text{H} \cdots \text{O}$, $\text{N}-\text{H} \cdots \text{N}$ and $\text{C}-\text{H} \cdots \text{O}$ hydrogen bonding are observed in the crystal structure. $\text{C}-\text{H} \cdots \pi$ and $\pi-\pi$ stacking interactions [centroid-centroid distance = 3.9985 (10) Å] are also present.

Related literature

For applications of 3-aminobenzoic acid, see: Lynch & McClenaghan (2001); Smith (2005). For related structures, see: Smith *et al.* (1995); Lynch *et al.* (1998). For a similar dimeric $R_2^2(14)$ structure, see: Etter *et al.* (1990).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_2 \cdot \text{C}_7\text{H}_7\text{NO}_2$
 $M_r = 321.37$
 Triclinic, $P\bar{1}$
 $a = 9.0430$ (3) Å
 $b = 13.0565$ (5) Å

$c = 14.6300$ (5) Å
 $\alpha = 88.172$ (3)°
 $\beta = 79.366$ (3)°
 $\gamma = 74.506$ (3)°
 $V = 1635.72$ (10) Å³

$Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹

$T = 100$ K
 $0.54 \times 0.18 \times 0.15$ mm

Data collection

Oxford Diffraction Gemini-S CCD diffractometer
 Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2009)
 $T_{\min} = 0.997$, $T_{\max} = 1.000$
 12385 measured reflections
 5971 independent reflections
 4125 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.108$
 $S = 0.99$
 5971 reflections
 455 parameters
 2 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.57$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.35$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg5 is the centroid of the C2–C7 ring.

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{O1}-\text{H1C} \cdots \text{N2}$	0.84 (2)	1.79 (2)	2.6294 (19)	176 (2)
$\text{O3}-\text{H3A} \cdots \text{N6}^i$	0.84 (1)	1.75 (1)	2.5790 (19)	171 (2)
$\text{N1}-\text{H1A} \cdots \text{O4}^{ii}$	0.89 (2)	2.21 (2)	3.061 (2)	158.6 (17)
$\text{N1}-\text{H1B} \cdots \text{N3}^{iii}$	0.882 (19)	2.17 (2)	3.048 (2)	177.8 (19)
$\text{N4}-\text{H4A} \cdots \text{O2}^{ii}$	0.89 (2)	2.19 (2)	3.035 (2)	157.6 (18)
$\text{N4}-\text{H4B} \cdots \text{N5}^{iv}$	0.89 (2)	2.13 (2)	3.017 (2)	171.9 (17)
$\text{C3}-\text{H3C} \cdots \text{O4}^{ii}$	0.95	2.56	3.353 (2)	141
$\text{C22}-\text{H22A} \cdots \text{O2}^{ii}$	0.95	2.54	3.327 (2)	141
$\text{C28}-\text{H28A} \cdots \text{O2}^v$	0.95	2.55	3.492 (2)	172
$\text{C38}-\text{H38A} \cdots \text{O4}^{vi}$	0.95	2.50	3.448 (2)	177
$\text{C12}-\text{H12A} \cdots \text{Cg5}^v$	0.95	2.67	3.5510 (18)	154

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

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

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU2742).

References

- Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). *Acta Cryst.* **B46**, 256–262.
 Lynch, D. E. & McClenaghan, I. (2001). *Acta Cryst.* **C57**, 830–832.
 Lynch, D. E., Smith, G., Byriel, K. A. & Kennard, C. H. L. (1998). *Aust. J. Chem.* **51**, 587–592.
 Oxford Diffraction (2008). *CrysAlis CCD* and *CrysAlis RED*. Oxford Diffraction Ltd, Yarnton, England.
 Oxford Diffraction (2009). *CrysAlis PRO*. Oxford Diffraction Ltd, Yarnton, England.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Smith, G. (2005). *Acta Cryst.* **E61**, o3398–o3400.
 Smith, G., Gentner, J. M., Lynch, D. E., Byriel, K. A. & Kennard, C. H. L. (1995). *Aust. J. Chem.* **48**, 1151–1166.
 Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supporting information

Acta Cryst. (2010). E66, o1138 [https://doi.org/10.1107/S1600536810014261]

3-Aminobenzoic acid–1,2-bis(4-pyridyl)ethane (1/1)**Fwu Ming Shen and Shie Fu Lush****S1. Comment**

The structures of a number of 1:1 adduct compounds of 3-aminobenzoic acid with Lewis bases have been reported. These include compounds with 2-amino-pyrimidine (Smith *et al.*, 1995) and 2-aminobenzothiazole (Lynch *et al.*, 1998). Not only 3-aminobenzoic acid is more effective for the promotion of hydrogen-bonding extensions, but also forms various weak noncovalent interactions, such as π - π stacking and C—H \cdots π interactions.

Our research program aims to gain hydrogen-bonding networks involving 3-aminobenzoic acid with 1,2-bis(4-pyridyl)ethane, which can develop well defined noncovalent supramolecular architectures and form multiple hydrogen bonds containing components of complementary arrays of hydrogen-bonding sites. The resulting crystal X-Ray structure (Fig. 1) consists entirely of neutral 3-aminobenzoic acid molecules and neutral 1,2-bis(4-pyridyl)ethane units, with no proton transfer (Lynch & McClenaghan, 2001; Smith, 2005). The title compound comprises non-planar molecules, similar to other analogous compounds (Smith *et al.*, 1995; Lynch *et al.*, 1998) that associate 3-aminobenzoic acid via amine and carboxylic group N—H \cdots O [N \cdots O 3.061 (2) and 3.035 (2) Å] dimer $R_2^2(14)$ (Etter *et al.*, 1990) and form linear hydrogen-bonded via amine and pyridine N—H \cdots N [N \cdots N 3.048 (2) and 3.017 (2) Å].

The title compound's supramolecular structure can be readily analyzed in terms of carboxyl atom O1 and amino group N1 act as hydrogen-bond donors to pyridyl atoms N2 and N3. Similarly, N4, C22, C28, N1, C3, C38 act as hydrogen-bond donors to carboxyl atoms O2, O4, respectively (Table 1 and Fig. 2).

This layer is consolidated by C—H \cdots π interaction (C12—H12A \cdots Cg5; full details and symmetry code are given in Table 1). Furthermore, π - π ring stacking interactions are between neighboring complexes in the structure. The distance between Cg1 (N2/C8—C12) \cdots Cg4 (N6/C34—C38) is 3.9985 (10) Å and dihedral angle between two rings is 9.86 (9) °.

S2. Experimental

The 3-aminobenzoic acid (171.0 mg, 1.0 mmol) and 1,2-bis(4-pyridyl)ethane (184 mg, 1.0 mmol) were dissolved in 20 ml 50% methanol-water, the solution was refluxed for 30 min. The filtered solution was transferred to a 25 ml tube after one week at room temperature, and colorless transparent crystals formed (yield 59.36%). Elemental analysis calcd(%) for C₁₉H₁₉N₃O₂ (Mr=321.37): C, 55.83; H, 5.96; N, 13.08; Found: C, 56.03; H, 5.92; N, 12.96.

S3. Refinement

Amino H atoms were located in a difference Fourier map and were refined isotropically. Water H atoms were located in a difference Fourier map and refined with the distances constraints of O—H = 0.84 Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. Other H atoms were positioned geometrically with C—H = 0.95 (aromatic) and 0.99 Å (methylene), and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

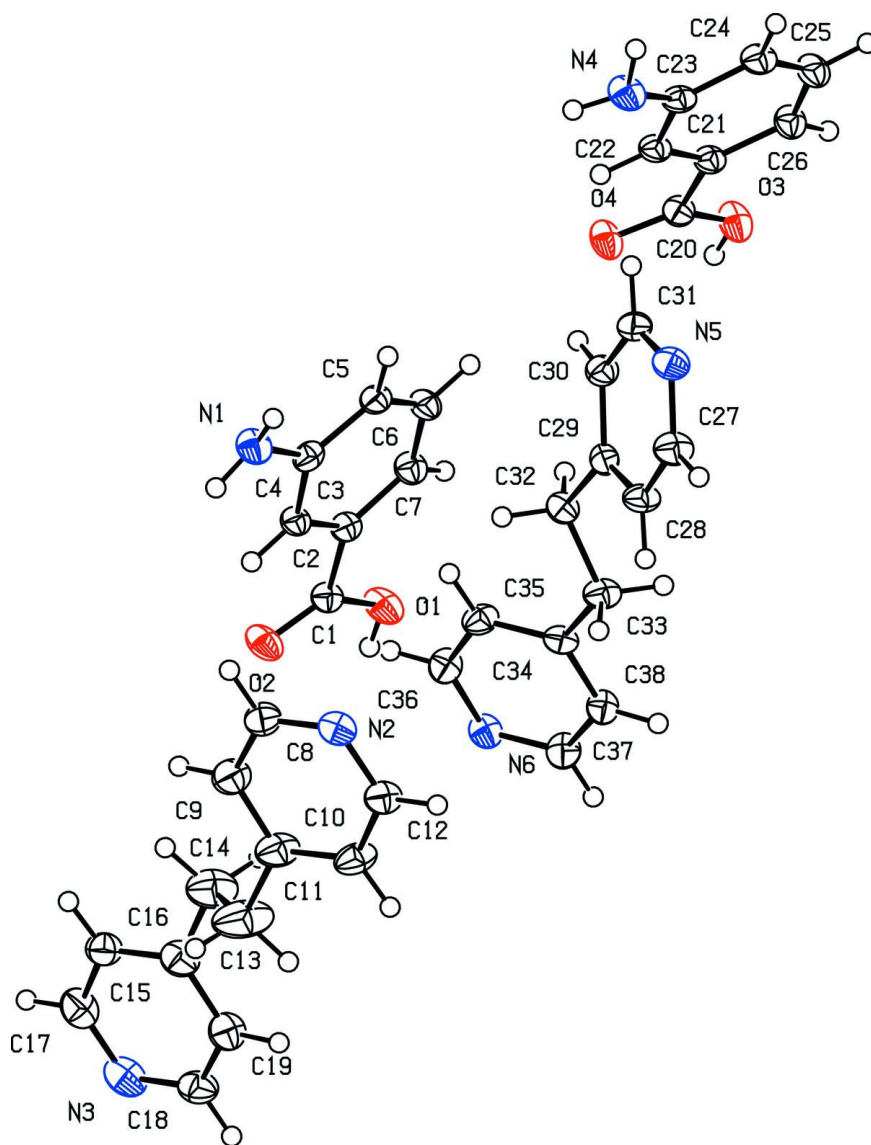


Figure 1

View of the title compound with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

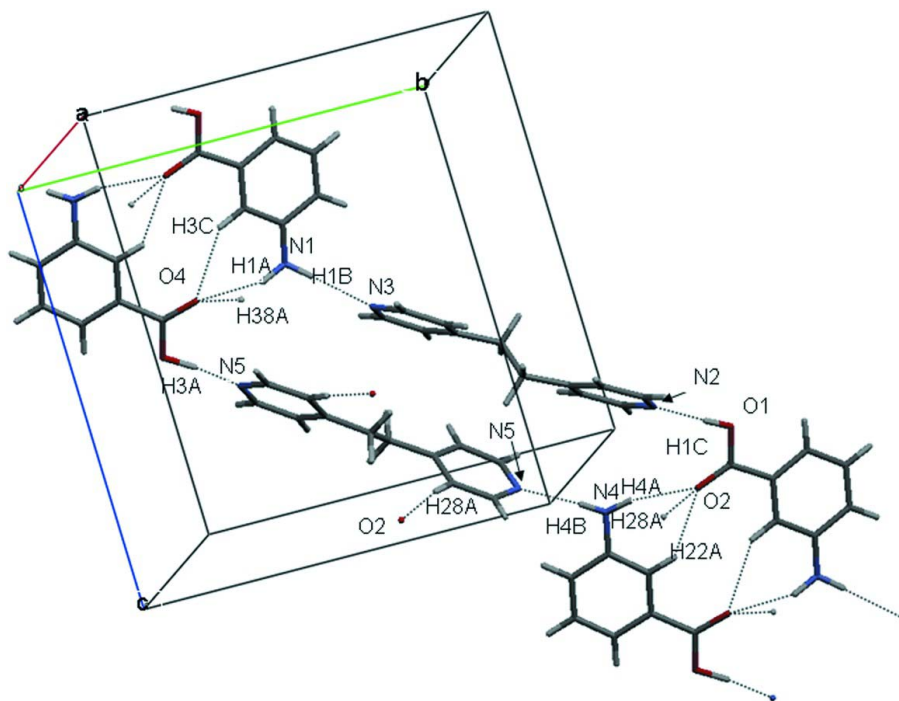


Figure 2

The molecular packing for the title compound. Hydrogen-bonding associations are shown as dotted lines.

3-Aminobenzoic acid-1,2-bis(4-pyridyl)ethane (1/1)

Crystal data

$C_{12}H_{12}N_2 \cdot C_7H_7NO_2$

$M_r = 321.37$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 9.0430$ (3) Å

$b = 13.0565$ (5) Å

$c = 14.6300$ (5) Å

$\alpha = 88.172$ (3)°

$\beta = 79.366$ (3)°

$\gamma = 74.506$ (3)°

$V = 1635.72$ (10) Å³

$Z = 4$

$F(000) = 680$

$D_x = 1.305$ Mg m⁻³

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

Cell parameters from 6302 reflections

$\theta = 2.5\text{--}29.2^\circ$

$\mu = 0.09$ mm⁻¹

$T = 100$ K

Parallelepiped, colorless

$0.54 \times 0.18 \times 0.15$ mm

Data collection

Oxford Diffraction Gemini-S CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*CrysAlis PRO*; Oxford Diffraction, 2009)

$T_{\min} = 0.997$, $T_{\max} = 1.000$

12385 measured reflections

5971 independent reflections

4125 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.020$

$\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 2.5^\circ$

$h = -10 \rightarrow 10$

$k = -13 \rightarrow 15$

$l = -17 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.108$
 $S = 0.99$
 5971 reflections
 455 parameters
 2 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0632P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.57 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.35 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.76671 (15)	0.58027 (10)	0.03742 (8)	0.0334 (3)
O2	0.77042 (15)	0.69496 (9)	-0.07886 (8)	0.0328 (3)
O3	0.26995 (15)	0.16295 (9)	0.48714 (9)	0.0323 (3)
O4	0.23015 (14)	0.28364 (9)	0.37667 (8)	0.0312 (3)
N1	0.69956 (18)	0.49939 (14)	-0.35303 (11)	0.0306 (4)
N2	0.81019 (17)	0.73452 (11)	0.13170 (10)	0.0281 (3)
N3	0.7482 (2)	1.31954 (12)	0.50754 (11)	0.0374 (4)
N4	0.28039 (18)	0.06579 (14)	0.08726 (11)	0.0305 (4)
N5	0.77629 (17)	0.10653 (11)	0.02344 (10)	0.0299 (4)
N6	0.76549 (16)	0.67214 (11)	0.41027 (10)	0.0261 (3)
C1	0.75649 (18)	0.60925 (13)	-0.04865 (12)	0.0230 (4)
C2	0.72593 (18)	0.52804 (13)	-0.10732 (12)	0.0225 (4)
C3	0.73369 (18)	0.54740 (13)	-0.20172 (12)	0.0226 (4)
H3C	0.7591	0.6100	-0.2264	0.027*
C4	0.70458 (18)	0.47598 (13)	-0.26121 (12)	0.0238 (4)
C5	0.67203 (19)	0.38323 (13)	-0.22244 (13)	0.0254 (4)
H5B	0.6563	0.3319	-0.2615	0.030*
C6	0.66242 (19)	0.36513 (13)	-0.12816 (13)	0.0277 (4)
H6B	0.6380	0.3023	-0.1033	0.033*
C7	0.68795 (19)	0.43750 (13)	-0.06922 (13)	0.0264 (4)
H7A	0.6796	0.4253	-0.0043	0.032*
C8	0.7001 (2)	0.82631 (14)	0.13950 (13)	0.0314 (4)
H8A	0.6139	0.8324	0.1092	0.038*
C9	0.7045 (2)	0.91298 (15)	0.18921 (14)	0.0368 (5)
H9A	0.6211	0.9760	0.1947	0.044*
C10	0.8319 (2)	0.90730 (15)	0.23106 (14)	0.0384 (5)
C11	0.9444 (2)	0.81133 (15)	0.22388 (13)	0.0353 (5)
H11A	1.0322	0.8031	0.2532	0.042*

C12	0.9304 (2)	0.72758 (14)	0.17472 (13)	0.0297 (4)
H12A	1.0092	0.6623	0.1712	0.036*
C13	0.8492 (2)	1.00327 (19)	0.28015 (19)	0.0631 (7)
H13A	0.9396	0.9800	0.3122	0.076*
H13B	0.8729	1.0551	0.2326	0.076*
C14	0.7126 (2)	1.05708 (17)	0.34813 (16)	0.0475 (6)
H14A	0.6866	1.0047	0.3945	0.057*
H14B	0.6229	1.0826	0.3157	0.057*
C15	0.7326 (2)	1.15073 (15)	0.39924 (13)	0.0335 (5)
C16	0.6186 (2)	1.24545 (14)	0.40946 (13)	0.0326 (4)
H16A	0.5313	1.2550	0.3795	0.039*
C17	0.6305 (2)	1.32601 (15)	0.46274 (14)	0.0363 (5)
H17A	0.5499	1.3906	0.4681	0.044*
C18	0.8615 (2)	1.22878 (15)	0.49533 (13)	0.0341 (5)
H18A	0.9484	1.2223	0.5251	0.041*
C19	0.8598 (2)	1.14407 (15)	0.44223 (13)	0.0341 (5)
H19A	0.9445	1.0818	0.4350	0.041*
C20	0.25604 (19)	0.19120 (13)	0.40181 (12)	0.0242 (4)
C21	0.27294 (18)	0.10109 (13)	0.33654 (12)	0.0228 (4)
C22	0.26585 (18)	0.12404 (13)	0.24435 (12)	0.0232 (4)
H22A	0.2535	0.1952	0.2242	0.028*
C23	0.27665 (18)	0.04385 (13)	0.18016 (12)	0.0235 (4)
C24	0.28837 (19)	-0.05921 (13)	0.21297 (13)	0.0270 (4)
H24A	0.2917	-0.1145	0.1714	0.032*
C25	0.2952 (2)	-0.08156 (14)	0.30540 (13)	0.0301 (4)
H25A	0.3038	-0.1522	0.3263	0.036*
C26	0.28984 (19)	-0.00229 (13)	0.36817 (13)	0.0268 (4)
H26A	0.2975	-0.0184	0.4312	0.032*
C27	0.8959 (2)	0.14566 (14)	0.03144 (13)	0.0313 (4)
H27A	0.9958	0.1120	-0.0034	0.038*
C28	0.8827 (2)	0.23233 (13)	0.08754 (12)	0.0280 (4)
H28A	0.9720	0.2566	0.0904	0.034*
C29	0.73906 (19)	0.28338 (13)	0.13937 (12)	0.0227 (4)
C30	0.6147 (2)	0.24242 (13)	0.13143 (12)	0.0264 (4)
H30A	0.5137	0.2739	0.1661	0.032*
C31	0.6377 (2)	0.15652 (14)	0.07347 (13)	0.0284 (4)
H31A	0.5501	0.1312	0.0687	0.034*
C32	0.71112 (19)	0.38225 (13)	0.19731 (12)	0.0243 (4)
H32A	0.6204	0.3855	0.2477	0.029*
H32B	0.6829	0.4448	0.1577	0.029*
C33	0.84736 (19)	0.39114 (13)	0.24075 (13)	0.0282 (4)
H33A	0.8779	0.3278	0.2790	0.034*
H33B	0.9373	0.3905	0.1904	0.034*
C34	0.81443 (18)	0.48966 (13)	0.30095 (12)	0.0230 (4)
C35	0.69101 (19)	0.57861 (13)	0.29678 (12)	0.0280 (4)
H35A	0.6204	0.5786	0.2560	0.034*
C36	0.6707 (2)	0.66686 (13)	0.35164 (12)	0.0277 (4)
H36A	0.5854	0.7267	0.3476	0.033*

C37	0.8843 (2)	0.58632 (13)	0.41499 (12)	0.0287 (4)
H37A	0.9527	0.5884	0.4565	0.034*
C38	0.9119 (2)	0.49513 (13)	0.36243 (12)	0.0265 (4)
H38A	0.9975	0.4361	0.3683	0.032*
H1C	0.785 (2)	0.6290 (10)	0.0663 (12)	0.040*
H3A	0.268 (2)	0.2174 (9)	0.5172 (12)	0.040*
H1A	0.731 (2)	0.5562 (16)	-0.3754 (13)	0.034 (5)*
H1B	0.714 (2)	0.4461 (15)	-0.3921 (14)	0.036 (5)*
H4A	0.252 (2)	0.1339 (18)	0.0720 (15)	0.051 (7)*
H4B	0.256 (2)	0.0199 (16)	0.0528 (14)	0.044 (6)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0541 (8)	0.0295 (7)	0.0231 (8)	-0.0198 (6)	-0.0098 (6)	-0.0049 (6)
O2	0.0516 (8)	0.0259 (7)	0.0270 (7)	-0.0185 (6)	-0.0100 (6)	-0.0021 (6)
O3	0.0530 (8)	0.0223 (7)	0.0236 (8)	-0.0093 (6)	-0.0118 (6)	-0.0049 (6)
O4	0.0489 (8)	0.0192 (7)	0.0268 (7)	-0.0078 (6)	-0.0110 (6)	-0.0042 (5)
N1	0.0430 (9)	0.0253 (9)	0.0265 (10)	-0.0107 (8)	-0.0106 (7)	-0.0061 (8)
N2	0.0338 (8)	0.0304 (9)	0.0207 (8)	-0.0117 (7)	-0.0011 (7)	-0.0057 (7)
N3	0.0517 (10)	0.0340 (10)	0.0277 (9)	-0.0169 (8)	-0.0006 (8)	-0.0077 (7)
N4	0.0430 (10)	0.0251 (9)	0.0262 (10)	-0.0120 (8)	-0.0078 (7)	-0.0087 (8)
N5	0.0357 (9)	0.0276 (8)	0.0292 (9)	-0.0113 (7)	-0.0076 (7)	-0.0067 (7)
N6	0.0342 (8)	0.0240 (8)	0.0203 (8)	-0.0078 (6)	-0.0046 (7)	-0.0050 (6)
C1	0.0234 (8)	0.0217 (9)	0.0228 (10)	-0.0052 (7)	-0.0018 (7)	-0.0050 (8)
C2	0.0211 (8)	0.0202 (9)	0.0251 (10)	-0.0032 (7)	-0.0035 (7)	-0.0061 (7)
C3	0.0231 (8)	0.0177 (9)	0.0267 (10)	-0.0047 (7)	-0.0039 (7)	-0.0047 (7)
C4	0.0199 (8)	0.0225 (9)	0.0273 (11)	-0.0005 (7)	-0.0064 (7)	-0.0072 (8)
C5	0.0243 (9)	0.0178 (9)	0.0350 (12)	-0.0030 (7)	-0.0102 (8)	-0.0093 (8)
C6	0.0287 (9)	0.0188 (9)	0.0369 (12)	-0.0075 (7)	-0.0069 (8)	-0.0024 (8)
C7	0.0296 (9)	0.0234 (10)	0.0262 (10)	-0.0067 (7)	-0.0047 (8)	-0.0034 (8)
C8	0.0345 (10)	0.0334 (11)	0.0296 (11)	-0.0109 (8)	-0.0112 (8)	-0.0020 (9)
C9	0.0321 (10)	0.0301 (11)	0.0476 (13)	-0.0047 (8)	-0.0093 (9)	-0.0105 (9)
C10	0.0310 (10)	0.0368 (11)	0.0473 (13)	-0.0073 (8)	-0.0060 (9)	-0.0197 (10)
C11	0.0265 (9)	0.0409 (12)	0.0397 (12)	-0.0073 (8)	-0.0089 (9)	-0.0146 (9)
C12	0.0265 (9)	0.0288 (10)	0.0325 (11)	-0.0063 (8)	-0.0015 (8)	-0.0097 (8)
C13	0.0382 (12)	0.0631 (16)	0.087 (2)	-0.0124 (11)	-0.0032 (12)	-0.0452 (14)
C14	0.0461 (12)	0.0530 (14)	0.0454 (14)	-0.0181 (11)	-0.0027 (11)	-0.0187 (11)
C15	0.0413 (11)	0.0320 (11)	0.0293 (11)	-0.0122 (9)	-0.0067 (9)	-0.0071 (9)
C16	0.0329 (10)	0.0339 (11)	0.0343 (12)	-0.0131 (8)	-0.0084 (8)	0.0014 (9)
C17	0.0359 (10)	0.0294 (11)	0.0380 (12)	-0.0060 (8)	0.0044 (9)	-0.0039 (9)
C18	0.0396 (11)	0.0377 (11)	0.0309 (11)	-0.0179 (9)	-0.0094 (9)	-0.0028 (9)
C19	0.0353 (10)	0.0295 (10)	0.0356 (12)	-0.0058 (8)	-0.0044 (9)	-0.0064 (9)
C20	0.0270 (9)	0.0228 (10)	0.0233 (11)	-0.0065 (7)	-0.0057 (8)	-0.0042 (8)
C21	0.0221 (8)	0.0208 (9)	0.0260 (10)	-0.0063 (7)	-0.0039 (7)	-0.0051 (7)
C22	0.0228 (8)	0.0205 (9)	0.0272 (10)	-0.0070 (7)	-0.0040 (7)	-0.0056 (7)
C23	0.0204 (8)	0.0242 (10)	0.0263 (11)	-0.0069 (7)	-0.0029 (7)	-0.0079 (8)
C24	0.0271 (9)	0.0225 (9)	0.0316 (11)	-0.0065 (7)	-0.0039 (8)	-0.0116 (8)

C25	0.0324 (10)	0.0196 (9)	0.0381 (12)	-0.0068 (8)	-0.0055 (8)	-0.0037 (8)
C26	0.0313 (9)	0.0223 (9)	0.0267 (11)	-0.0060 (7)	-0.0061 (8)	-0.0023 (8)
C27	0.0312 (10)	0.0315 (10)	0.0312 (11)	-0.0091 (8)	-0.0024 (8)	-0.0111 (8)
C28	0.0272 (9)	0.0282 (10)	0.0316 (11)	-0.0114 (7)	-0.0052 (8)	-0.0075 (8)
C29	0.0295 (9)	0.0205 (9)	0.0199 (10)	-0.0071 (7)	-0.0082 (7)	0.0002 (7)
C30	0.0263 (9)	0.0247 (9)	0.0286 (11)	-0.0068 (7)	-0.0056 (8)	-0.0025 (8)
C31	0.0293 (9)	0.0279 (10)	0.0326 (11)	-0.0114 (8)	-0.0115 (8)	-0.0018 (8)
C32	0.0284 (9)	0.0216 (9)	0.0229 (10)	-0.0066 (7)	-0.0034 (7)	-0.0050 (7)
C33	0.0260 (9)	0.0270 (10)	0.0314 (11)	-0.0070 (7)	-0.0027 (8)	-0.0125 (8)
C34	0.0237 (9)	0.0251 (9)	0.0208 (10)	-0.0098 (7)	0.0004 (7)	-0.0050 (7)
C35	0.0257 (9)	0.0300 (10)	0.0296 (11)	-0.0063 (7)	-0.0080 (8)	-0.0088 (8)
C36	0.0298 (9)	0.0257 (10)	0.0256 (10)	-0.0029 (7)	-0.0055 (8)	-0.0061 (8)
C37	0.0374 (10)	0.0281 (10)	0.0218 (10)	-0.0069 (8)	-0.0101 (8)	-0.0050 (8)
C38	0.0302 (9)	0.0242 (9)	0.0242 (10)	-0.0048 (7)	-0.0058 (8)	-0.0028 (8)

Geometric parameters (Å, °)

O1—C1	1.316 (2)	C14—H14A	0.9900
O1—H1C	0.843 (16)	C14—H14B	0.9900
O2—C1	1.2192 (19)	C15—C16	1.374 (3)
O3—C20	1.309 (2)	C15—C19	1.391 (2)
O3—H3A	0.842 (14)	C16—C17	1.367 (2)
O4—C20	1.2245 (19)	C16—H16A	0.9500
N1—C4	1.374 (2)	C17—H17A	0.9500
N1—H1A	0.89 (2)	C18—C19	1.377 (2)
N1—H1B	0.881 (19)	C18—H18A	0.9500
N2—C8	1.330 (2)	C19—H19A	0.9500
N2—C12	1.335 (2)	C20—C21	1.497 (2)
N3—C17	1.332 (2)	C21—C22	1.382 (2)
N3—C18	1.335 (2)	C21—C26	1.391 (2)
N4—C23	1.376 (2)	C22—C23	1.401 (2)
N4—H4A	0.89 (2)	C22—H22A	0.9500
N4—H4B	0.889 (19)	C23—C24	1.398 (2)
N5—C31	1.337 (2)	C24—C25	1.383 (3)
N5—C27	1.338 (2)	C24—H24A	0.9500
N6—C36	1.335 (2)	C25—C26	1.391 (2)
N6—C37	1.339 (2)	C25—H25A	0.9500
C1—C2	1.497 (2)	C26—H26A	0.9500
C2—C3	1.388 (2)	C27—C28	1.386 (2)
C2—C7	1.389 (2)	C27—H27A	0.9500
C3—C4	1.401 (2)	C28—C29	1.383 (2)
C3—H3C	0.9500	C28—H28A	0.9500
C4—C5	1.400 (2)	C29—C30	1.392 (2)
C5—C6	1.383 (3)	C29—C32	1.505 (2)
C5—H5B	0.9500	C30—C31	1.375 (2)
C6—C7	1.390 (2)	C30—H30A	0.9500
C6—H6B	0.9500	C31—H31A	0.9500
C7—H7A	0.9500	C32—C33	1.518 (2)

C8—C9	1.378 (2)	C32—H32A	0.9900
C8—H8A	0.9500	C32—H32B	0.9900
C9—C10	1.385 (2)	C33—C34	1.513 (2)
C9—H9A	0.9500	C33—H33A	0.9900
C10—C11	1.380 (3)	C33—H33B	0.9900
C10—C13	1.521 (3)	C34—C38	1.386 (2)
C11—C12	1.374 (2)	C34—C35	1.388 (2)
C11—H11A	0.9500	C35—C36	1.377 (2)
C12—H12A	0.9500	C35—H35A	0.9500
C13—C14	1.462 (3)	C36—H36A	0.9500
C13—H13A	0.9900	C37—C38	1.378 (2)
C13—H13B	0.9900	C37—H37A	0.9500
C14—C15	1.521 (3)	C38—H38A	0.9500
C1—O1—H1C	109.6 (14)	N3—C18—C19	123.90 (17)
C20—O3—H3A	107.9 (14)	N3—C18—H18A	118.1
C4—N1—H1A	117.4 (12)	C19—C18—H18A	118.1
C4—N1—H1B	117.8 (12)	C18—C19—C15	119.26 (17)
H1A—N1—H1B	117.4 (18)	C18—C19—H19A	120.4
C8—N2—C12	117.35 (14)	C15—C19—H19A	120.4
C17—N3—C18	115.91 (15)	O4—C20—O3	123.30 (15)
C23—N4—H4A	117.2 (14)	O4—C20—C21	122.09 (15)
C23—N4—H4B	117.1 (13)	O3—C20—C21	114.61 (15)
H4A—N4—H4B	116.4 (19)	C22—C21—C26	120.79 (15)
C31—N5—C27	116.04 (14)	C22—C21—C20	118.06 (15)
C36—N6—C37	117.17 (14)	C26—C21—C20	121.11 (15)
O2—C1—O1	123.33 (15)	C21—C22—C23	120.95 (16)
O2—C1—C2	122.23 (16)	C21—C22—H22A	119.5
O1—C1—C2	114.44 (15)	C23—C22—H22A	119.5
C3—C2—C7	120.77 (15)	N4—C23—C24	121.30 (15)
C3—C2—C1	117.48 (15)	N4—C23—C22	120.70 (16)
C7—C2—C1	121.74 (16)	C24—C23—C22	117.97 (16)
C2—C3—C4	121.01 (16)	C25—C24—C23	120.70 (15)
C2—C3—H3C	119.5	C25—C24—H24A	119.7
C4—C3—H3C	119.5	C23—C24—H24A	119.7
N1—C4—C5	121.39 (15)	C24—C25—C26	121.05 (17)
N1—C4—C3	120.93 (16)	C24—C25—H25A	119.5
C5—C4—C3	117.57 (16)	C26—C25—H25A	119.5
C6—C5—C4	121.08 (15)	C25—C26—C21	118.46 (16)
C6—C5—H5B	119.5	C25—C26—H26A	120.8
C4—C5—H5B	119.5	C21—C26—H26A	120.8
C5—C6—C7	120.99 (17)	N5—C27—C28	123.95 (17)
C5—C6—H6B	119.5	N5—C27—H27A	118.0
C7—C6—H6B	119.5	C28—C27—H27A	118.0
C2—C7—C6	118.52 (17)	C29—C28—C27	119.67 (15)
C2—C7—H7A	120.7	C29—C28—H28A	120.2
C6—C7—H7A	120.7	C27—C28—H28A	120.2
N2—C8—C9	123.44 (16)	C28—C29—C30	116.43 (15)

N2—C8—H8A	118.3	C28—C29—C32	123.50 (14)
C9—C8—H8A	118.3	C30—C29—C32	119.95 (15)
C8—C9—C10	119.32 (17)	C31—C30—C29	120.15 (16)
C8—C9—H9A	120.3	C31—C30—H30A	119.9
C10—C9—H9A	120.3	C29—C30—H30A	119.9
C11—C10—C9	116.86 (16)	N5—C31—C30	123.75 (15)
C11—C10—C13	121.56 (17)	N5—C31—H31A	118.1
C9—C10—C13	121.56 (18)	C30—C31—H31A	118.1
C12—C11—C10	120.48 (16)	C29—C32—C33	115.58 (14)
C12—C11—H11A	119.8	C29—C32—H32A	108.4
C10—C11—H11A	119.8	C33—C32—H32A	108.4
N2—C12—C11	122.48 (16)	C29—C32—H32B	108.4
N2—C12—H12A	118.8	C33—C32—H32B	108.4
C11—C12—H12A	118.8	H32A—C32—H32B	107.4
C14—C13—C10	114.94 (18)	C34—C33—C32	114.45 (14)
C14—C13—H13A	108.5	C34—C33—H33A	108.6
C10—C13—H13A	108.5	C32—C33—H33A	108.6
C14—C13—H13B	108.5	C34—C33—H33B	108.6
C10—C13—H13B	108.5	C32—C33—H33B	108.6
H13A—C13—H13B	107.5	H33A—C33—H33B	107.6
C13—C14—C15	114.66 (17)	C38—C34—C35	116.73 (15)
C13—C14—H14A	108.6	C38—C34—C33	120.09 (15)
C15—C14—H14A	108.6	C35—C34—C33	123.16 (14)
C13—C14—H14B	108.6	C36—C35—C34	120.06 (15)
C15—C14—H14B	108.6	C36—C35—H35A	120.0
H14A—C14—H14B	107.6	C34—C35—H35A	120.0
C16—C15—C19	116.69 (16)	N6—C36—C35	123.04 (16)
C16—C15—C14	120.43 (17)	N6—C36—H36A	118.5
C19—C15—C14	122.78 (17)	C35—C36—H36A	118.5
C17—C16—C15	120.10 (17)	N6—C37—C38	123.11 (15)
C17—C16—H16A	119.9	N6—C37—H37A	118.4
C15—C16—H16A	119.9	C38—C37—H37A	118.4
N3—C17—C16	124.04 (18)	C37—C38—C34	119.88 (16)
N3—C17—H17A	118.0	C37—C38—H38A	120.1
C16—C17—H17A	118.0	C34—C38—H38A	120.1

Hydrogen-bond geometry (Å, °)

Cg5 is the centroid of the C2–C7 ring.

<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>
O1—H1C \cdots N2	0.84 (2)	1.79 (2)	2.6294 (19)	176 (2)
O3—H3A \cdots N6 ⁱ	0.84 (1)	1.75 (1)	2.5790 (19)	171 (2)
N1—H1A \cdots O4 ⁱⁱ	0.89 (2)	2.21 (2)	3.061 (2)	158.6 (17)
N1—H1B \cdots N3 ⁱⁱⁱ	0.882 (19)	2.17 (2)	3.048 (2)	177.8 (19)
N4—H4A \cdots O2 ⁱⁱ	0.89 (2)	2.19 (2)	3.035 (2)	157.6 (18)
N4—H4B \cdots N5 ^{iv}	0.89 (2)	2.13 (2)	3.017 (2)	171.9 (17)
C3—H3C \cdots O4 ⁱⁱ	0.95	2.56	3.353 (2)	141
C22—H22A \cdots O2 ⁱⁱ	0.95	2.54	3.327 (2)	141

C28—H28A···O2 ^v	0.95	2.55	3.492 (2)	172
C38—H38A···O4 ^{vi}	0.95	2.50	3.448 (2)	177
C12—H12A···Cg5 ^v	0.95	2.67	3.5510 (18)	154

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