

## 3-Aminobenzoic acid–4,4'-bipyridine (2/3)

Pornsuda Lhengwan, Supakit Achiwawanich and Tanwawan Duangthongyou\*

Department of Chemistry, Faculty of Science, Kasetsart University, Bangkok 10903, Thailand

Correspondence e-mail: fscitwd@ku.ac.th

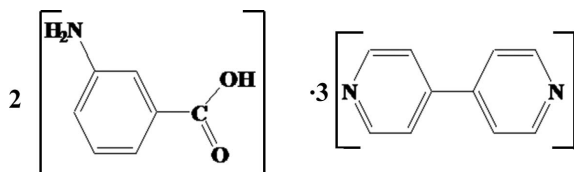
Received 19 July 2012; accepted 23 July 2012

Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å;  $R$  factor = 0.071;  $wR$  factor = 0.252; data-to-parameter ratio = 12.7.

The asymmetric unit of the title compound,  $3\text{C}_{10}\text{H}_8\text{N}_2 \cdot 2\text{C}_7\text{H}_7\text{NO}_2$ , consists of three molecules of 4,4'-bipyridine (bpy) and two molecules of 3-aminobenzoic acid (bza). Two molecules of bza and two molecules of bpy are connected *via*  $\text{O}-\text{H} \cdots \text{N}$ ,  $\text{N}-\text{H} \cdots \text{N}$  and  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds, forming infinite double-stranded zigzag chains along the  $c$  axis. The third molecule of bpy is linked to the chain by weak  $\text{C}-\text{H} \cdots \text{O}$  interactions. Adjacent chains are linked *via*  $\pi-\pi$  interactions [centroid-centroid distances = 3.759 (3)–3.928 (3) Å] involving the pyridine rings of bpy molecules, resulting in a sheet-like structure parallel to (100). These sheets are stacked *via*  $\text{C}-\text{H} \cdots \pi$  interactions, resulting finally in the formation of a three-dimensional supramolecular structure.

### Related literature

For related structures, see: Karpova *et al.* (2004); Koteswara Rao *et al.* (2012); Yao *et al.* (2008); Zhao *et al.* (2007).



### Experimental

#### Crystal data

$3\text{C}_{10}\text{H}_8\text{N}_2 \cdot 2\text{C}_7\text{H}_7\text{NO}_2$

$M_r = 742.82$

Triclinic,  $P\bar{1}$

$a = 9.371$  (3) Å

$b = 11.991$  (4) Å

$c = 17.653$  (6) Å

$\alpha = 94.910$  (11)°

$\beta = 90.224$  (10)°

$\gamma = 102.128$  (11)°

$V = 1931.6$  (11) Å<sup>3</sup>

$Z = 2$

Mo  $K\alpha$  radiation

$\mu = 0.09$  mm<sup>-1</sup>

$T = 296$  K

$0.62 \times 0.34 \times 0.05$  mm

#### Data collection

Bruker APEXII CCD diffractometer  
14412 measured reflections

6518 independent reflections  
3893 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.025$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.071$

$wR(F^2) = 0.252$

$S = 1.07$

6518 reflections

513 parameters

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.31$  e Å<sup>-3</sup>

$\Delta\rho_{\text{min}} = -0.29$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$Cg7$  and  $Cg8$  are the centroids of the  $C31-C36$  and  $C38-C43$  rings, respectively.

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O2-H' \cdots N4$	0.98 (4)	1.67 (4)	2.647 (4)	171 (3)
$O4-H'' \cdots N5$	1.03 (5)	1.60 (5)	2.627 (4)	175 (4)
$N7-H7A \cdots N3^i$	0.86	2.16	3.007 (4)	169 (4)
$N7-H7B \cdots O3$	0.86	2.23	3.038 (4)	156
$N8-H8A \cdots N6^{ii}$	0.86	2.16	3.006 (4)	169 (4)
$N8-H8B \cdots O1$	0.86	2.21	3.019 (4)	158
$C36-H36A \cdots O3$	0.93	2.52	3.300 (4)	141
$C43-H43A \cdots O1$	0.93	2.53	3.304 (4)	141
$C7-H7C \cdots Cg7^{iii}$	0.93	2.87	3.751 (4)	157
$C17-H17A \cdots Cg8^{iv}$	0.93	2.72	3.562 (4)	151
$C19-H19A \cdots Cg8^v$	0.93	2.78	3.643 (4)	154
$C22-H22A \cdots Cg7^{vi}$	0.93	2.72	3.567 (4)	151

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

Data collection: *APEX2* (Bruker, 2011); cell refinement: *SAINT* (Bruker, 2011); data reduction: *SAINT*; 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: *SHELXTL*.

The authors thank Kasetsart University Research and Development Institute and the Department of Chemistry, Faculty of Science, Kasetsart University, for research funds.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: QM2077).

### References

- Bruker (2011). *APEX2*, *SAINT* and *SADABS*. Bruker AXS Inc, Madison, Wisconsin, USA.
- Karpova, E. V., Zakharov, M. A., Gutnikov, S. I. & Alekseyev, R. S. (2004). *Acta Cryst.* **E60**, o2491–o2492.
- Koteswara Rao, V., Zeller, M. & Lovelace-Cameron, S. R. (2012). *Acta Cryst.* **E68**, o1711.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Yao, J. C., Qin, J. H., Sun, Q. B., Qu, L. & Li, Y. G. (2008). *Z. Kristallogr. New Cryst. Struct.* **223**, 11–12.
- Zhao, W.-X., Gao, Y.-X., Dong, S.-F., Li, Y. & Zhang, W.-P. (2007). *Acta Cryst.* **E63**, o2728.

## supplementary materials

*Acta Cryst.* (2012). E68, o2569 [doi:10.1107/S1600536812033181]

**3-Aminobenzoic acid–4,4'-bipyridine (2/3)**

**Pornsuda Lhengwan, Supakit Achiwawanich and Tanwawan Duangthongyou**

**Comment**

The title compound was obtained as a side-product in the synthesis of metal-organic framework materials which similar to previously observed organic compounds (Karpova *et al.* (2004); Koteswara Rao *et al.* (2012); Yao *et al.* (2008); Zhao *et al.* (2007). The asymmetric unit is composed of crystallographically independent of 4,4'-bipyridine (bpy) and 3-aminobenzoic acid (bza) in the stoichiometric ratio of 3:2 (Fig. 1). Three strong intermolecular hydrogen bonds are formed between bpy molecules and bza molecules *via* (1) amino and carboxylic acid groups of bza molecules (N—H $\cdots$ O), (2) amino groups of bza molecules and an N atom of bpy (N—H $\cdots$ N) and (3) a carboxylic group and an N atom of bpy (O—H $\cdots$ N), resulting in infinite double zigzag chain along *c*-axis (Table 1, Fig. 2). The infinite chain is further stabilized by weak C—H $\cdots$ N interactions (distance 2.75 Å) with the third bpy molecule. In the chain the bpy and bza molecules are arranged in a nonplanar structure. Moreover the pyridine rings of bpy molecules exhibit a nonplanar configuration with dihedral angles in the range 28.9–32.66°. In addition, each of the infinite zigzag chains is linked to adjacent chains through  $\pi$ - $\pi$  interactions between the pyridine rings of bpy molecules (distance 3.85 Å) forming a two-dimensional sheet structure (Fig. 2). The two-dimensional sheets are further connected through C—H $\cdots$  $\pi$  interactions (distance 2.74 Å), resulting in a three-dimensional supramolecular structure (Fig. 3)

**Experimental**

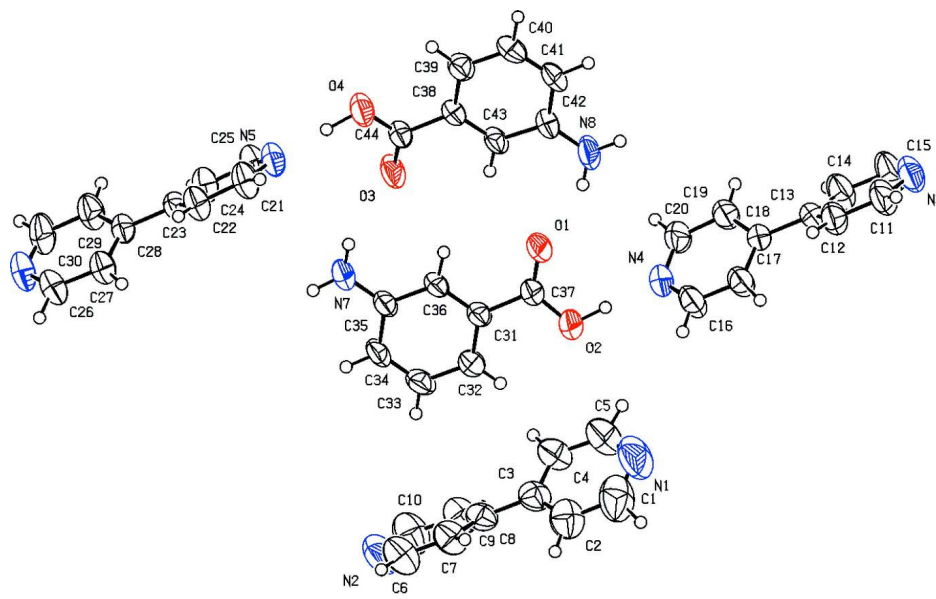
A solution of Mn(OAc)<sub>2</sub>·4H<sub>2</sub>O (1 mmol), 2-aminoterephthalic acid (1 mmol) and 4,4'-bipyridine (1 mmol) in 15 ml of H<sub>2</sub>O:DMF (1:1) was transferred into a Teflon lined autoclave and heat at 170° for 15 h. Then, solution was slowly cooled to room temperature. Yellow crystal was obtained as a minor product from fragmentation of 2-aminoterephthalic acid to 3-aminobenzoic acid during reaction.

**Refinement**

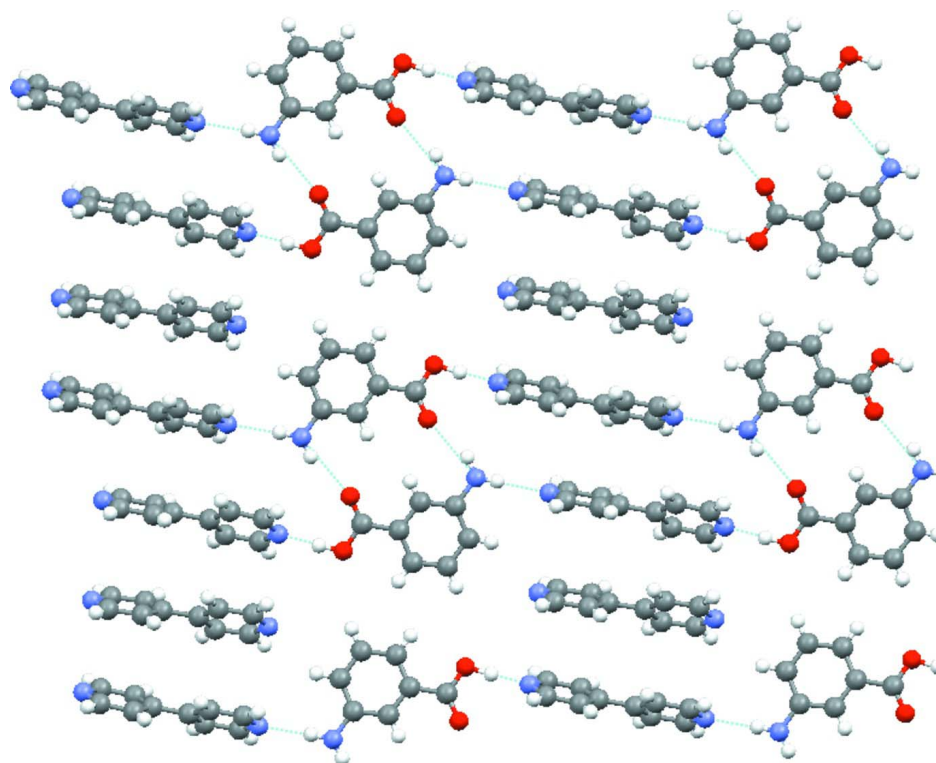
All H atoms of the compound were placed in the calculated positions with C—H = 0.93 Å, N—H = 0.86 Å and included in the cycles of refinement in a rigid model,  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{H})$ . Except carboxylic acid H atom were located in different fourier map and restrained to their hosts.

**Computing details**

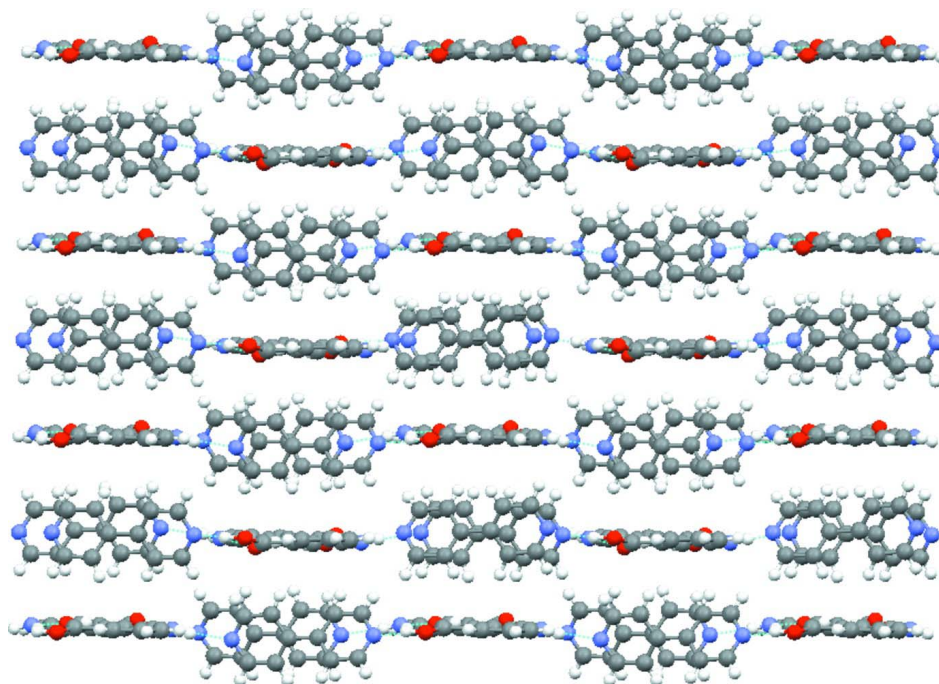
Data collection: *APEX2* (Bruker, 2011); cell refinement: *SAINTE* (Bruker, 2011); data reduction: *SAINTE* (Bruker, 2011); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

**Figure 1**

Molecular structure of the title compound (arbitrary spheres for the H atoms).

**Figure 2**

The two-dimensional-sheet structure showing  $\pi$ - $\pi$  stacking (Hydrogen bonds indicated by dash lines).

**Figure 3**

The three-dimensional-supramolecular structure showing the C-H to  $\pi$  interactions.

### 3-Aminobenzoic acid–4,4'-bipyridine (2/3)

#### Crystal data

$3\text{C}_{10}\text{H}_8\text{N}_2 \cdot 2\text{C}_7\text{H}_7\text{NO}_2$

$M_r = 742.82$

Triclinic,  $P\bar{1}$

$a = 9.371$  (3) Å

$b = 11.991$  (4) Å

$c = 17.653$  (6) Å

$\alpha = 94.910$  (11)°

$\beta = 90.224$  (10)°

$\gamma = 102.128$  (11)°

$V = 1931.6$  (11) Å<sup>3</sup>

$Z = 2$

$F(000) = 780$

$D_x = 1.277$  Mg m<sup>-3</sup>

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

Cell parameters from 3914 reflections

$\theta = 2.2\text{--}23.6^\circ$

$\mu = 0.09$  mm<sup>-1</sup>

$T = 296$  K

Plate, yellow

$0.62 \times 0.34 \times 0.05$  mm

#### Data collection

Bruker APEXII CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

14412 measured reflections

6518 independent reflections

3893 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$

$\theta_{\text{max}} = 25.0^\circ$ ,  $\theta_{\text{min}} = 1.2^\circ$

$h = -11 \rightarrow 11$

$k = -13 \rightarrow 14$

$l = -21 \rightarrow 20$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.071$

$wR(F^2) = 0.252$

$S = 1.07$

6518 reflections

513 parameters

0 restraints

Primary atom site location: structure-invariant direct methods  
 Secondary atom site location: difference Fourier map  
 Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.1384P)^2 + 0.418P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.29 \text{ e } \text{\AA}^{-3}$

*Special details*

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

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O4	0.2374 (3)	0.7665 (2)	0.37029 (13)	0.0653 (7)
O2	0.2525 (3)	0.1595 (2)	0.12426 (13)	0.0686 (7)
C38	0.2173 (3)	0.6986 (3)	0.24016 (15)	0.0419 (7)
C31	0.2317 (3)	0.2219 (3)	0.25340 (16)	0.0453 (7)
N7	0.2381 (3)	0.3911 (3)	0.43764 (14)	0.0716 (9)
H7A	0.2429	0.3827	0.4854	0.086*
H7B	0.2347	0.4570	0.4228	0.086*
O3	0.1895 (3)	0.5792 (2)	0.34188 (12)	0.0728 (8)
C35	0.2356 (3)	0.2994 (3)	0.38509 (16)	0.0480 (8)
C41	0.2283 (3)	0.7337 (3)	0.08730 (17)	0.0527 (8)
H41A	0.2319	0.7465	0.0361	0.063*
C44	0.2138 (3)	0.6754 (3)	0.32162 (16)	0.0449 (7)
O1	0.2077 (4)	0.3316 (2)	0.15097 (12)	0.0843 (9)
C32	0.2392 (4)	0.1155 (3)	0.27634 (17)	0.0541 (8)
H32A	0.2418	0.0545	0.2405	0.065*
C18	0.2790 (3)	0.2763 (3)	-0.16668 (16)	0.0479 (8)
C36	0.2291 (3)	0.3127 (3)	0.30716 (15)	0.0471 (8)
H36A	0.2229	0.3835	0.2912	0.056*
C43	0.2111 (3)	0.6072 (3)	0.18637 (15)	0.0461 (7)
H43A	0.2019	0.5340	0.2021	0.055*
C34	0.2409 (3)	0.1914 (3)	0.40706 (17)	0.0527 (8)
H34A	0.2433	0.1802	0.4585	0.063*
N8	0.2155 (4)	0.5309 (3)	0.05628 (14)	0.0773 (10)
H8A	0.2199	0.5406	0.0086	0.093*
H8B	0.2094	0.4636	0.0711	0.093*
C23	0.2592 (3)	0.6558 (3)	0.66064 (16)	0.0473 (7)
C42	0.2184 (3)	0.6223 (3)	0.10845 (16)	0.0491 (8)
C37	0.2289 (3)	0.2435 (3)	0.17188 (16)	0.0500 (8)
C13	0.2825 (3)	0.3010 (3)	-0.24750 (16)	0.0497 (8)
C39	0.2286 (3)	0.8082 (3)	0.21794 (17)	0.0514 (8)

---

H39A	0.2332	0.8702	0.2541	0.062*
N6	0.2662 (4)	0.5891 (3)	0.89493 (15)	0.0756 (9)
C28	0.2610 (3)	0.6319 (3)	0.74141 (16)	0.0479 (8)
N4	0.2684 (3)	0.2220 (3)	-0.01622 (15)	0.0640 (8)
N3	0.2904 (4)	0.3476 (3)	-0.40037 (16)	0.0818 (10)
N5	0.2546 (3)	0.7082 (3)	0.50962 (14)	0.0612 (8)
C33	0.2426 (4)	0.1014 (3)	0.35363 (18)	0.0579 (9)
H33A	0.2461	0.0300	0.3694	0.070*
C40	0.2329 (3)	0.8241 (3)	0.14098 (18)	0.0554 (8)
H40A	0.2391	0.8972	0.1256	0.067*
C27	0.3880 (4)	0.6226 (3)	0.77845 (17)	0.0606 (9)
H27A	0.4750	0.6300	0.7523	0.073*
C26	0.3852 (4)	0.6022 (4)	0.85425 (19)	0.0717 (11)
H26A	0.4722	0.5975	0.8780	0.086*
C17	0.3960 (4)	0.2479 (3)	-0.13103 (18)	0.0626 (10)
H17A	0.4814	0.2462	-0.1574	0.075*
C19	0.1573 (4)	0.2791 (3)	-0.12358 (18)	0.0623 (9)
H19A	0.0758	0.2989	-0.1445	0.075*
C24	0.1354 (4)	0.6189 (3)	0.61445 (18)	0.0615 (9)
H24A	0.0510	0.5757	0.6334	0.074*
C12	0.4107 (4)	0.3535 (3)	-0.28055 (18)	0.0643 (10)
H12A	0.4973	0.3738	-0.2520	0.077*
C22	0.3807 (4)	0.7180 (3)	0.62811 (18)	0.0619 (9)
H22A	0.4668	0.7438	0.6566	0.074*
C16	0.3865 (4)	0.2222 (4)	-0.05703 (18)	0.0704 (11)
H16A	0.4673	0.2039	-0.0342	0.084*
C25	0.1382 (4)	0.6465 (3)	0.54070 (18)	0.0665 (10)
H25A	0.0541	0.6206	0.5106	0.080*
C11	0.4087 (5)	0.3751 (4)	-0.35570 (19)	0.0762 (11)
H11A	0.4955	0.4112	-0.3764	0.091*
C21	0.3740 (4)	0.7417 (3)	0.55334 (18)	0.0675 (10)
H21A	0.4574	0.7832	0.5325	0.081*
C20	0.1572 (4)	0.2523 (3)	-0.04944 (19)	0.0685 (10)
H20A	0.0746	0.2558	-0.0211	0.082*
C14	0.1599 (4)	0.2723 (4)	-0.29407 (19)	0.0750 (11)
H14A	0.0711	0.2366	-0.2751	0.090*
C29	0.1364 (4)	0.6188 (4)	0.78445 (18)	0.0706 (11)
H29A	0.0476	0.6237	0.7628	0.085*
C30	0.1453 (5)	0.5984 (4)	0.8596 (2)	0.0818 (12)
H30A	0.0603	0.5907	0.8875	0.098*
C15	0.1693 (5)	0.2968 (5)	-0.3688 (2)	0.0928 (15)
H15A	0.0849	0.2761	-0.3991	0.111*
C8	0.7232 (4)	0.0224 (3)	0.2943 (2)	0.0598 (9)
C3	0.7286 (4)	0.0451 (3)	0.2136 (2)	0.0647 (9)
C9	0.5967 (4)	-0.0339 (3)	0.3258 (2)	0.0760 (11)
H9A	0.5130	-0.0604	0.2955	0.091*
N2	0.7093 (6)	-0.0184 (4)	0.4481 (2)	0.1066 (13)
C4	0.6089 (5)	0.0650 (4)	0.1757 (2)	0.0874 (13)
H4B	0.5225	0.0660	0.2012	0.105*

---

C7	0.8441 (4)	0.0586 (4)	0.3433 (2)	0.0790 (11)
H7C	0.9323	0.0976	0.3255	0.095*
N1	0.7348 (8)	0.0833 (4)	0.0594 (2)	0.1276 (18)
C10	0.5937 (6)	-0.0509 (4)	0.4012 (3)	0.0995 (15)
H10A	0.5060	-0.0874	0.4210	0.119*
C2	0.8530 (5)	0.0441 (4)	0.1716 (2)	0.0866 (12)
H2B	0.9366	0.0303	0.1943	0.104*
C6	0.8314 (6)	0.0361 (5)	0.4181 (3)	0.0984 (15)
H6B	0.9135	0.0604	0.4498	0.118*
C5	0.6177 (8)	0.0833 (5)	0.1007 (3)	0.122 (2)
H5B	0.5351	0.0968	0.0768	0.146*
C1	0.8513 (7)	0.0639 (5)	0.0957 (3)	0.1154 (18)
H1B	0.9359	0.0637	0.0684	0.138*
H''	0.239 (4)	0.744 (4)	0.425 (2)	0.101 (13)*
H'	0.248 (4)	0.178 (4)	0.071 (2)	0.099 (13)*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O4	0.1013 (18)	0.0551 (15)	0.0357 (12)	0.0066 (13)	-0.0032 (12)	0.0073 (11)
O2	0.1079 (19)	0.0677 (17)	0.0377 (13)	0.0335 (14)	0.0087 (12)	0.0097 (12)
C38	0.0456 (16)	0.0447 (18)	0.0357 (15)	0.0066 (13)	-0.0013 (12)	0.0125 (14)
C31	0.0505 (17)	0.0510 (19)	0.0385 (16)	0.0160 (14)	0.0045 (13)	0.0141 (14)
N7	0.114 (2)	0.076 (2)	0.0328 (14)	0.0344 (19)	-0.0027 (15)	0.0096 (15)
O3	0.130 (2)	0.0503 (15)	0.0366 (12)	0.0117 (14)	0.0021 (12)	0.0158 (11)
C35	0.0530 (18)	0.061 (2)	0.0345 (15)	0.0193 (15)	0.0021 (13)	0.0147 (15)
C41	0.0566 (19)	0.066 (2)	0.0381 (16)	0.0107 (16)	-0.0010 (13)	0.0227 (16)
C44	0.0517 (17)	0.0478 (19)	0.0350 (15)	0.0080 (14)	0.0001 (12)	0.0087 (15)
O1	0.161 (3)	0.0675 (17)	0.0400 (13)	0.0535 (18)	0.0096 (14)	0.0191 (13)
C32	0.069 (2)	0.051 (2)	0.0467 (18)	0.0176 (16)	0.0069 (15)	0.0143 (16)
C18	0.0554 (18)	0.054 (2)	0.0376 (16)	0.0186 (15)	0.0016 (14)	0.0050 (14)
C36	0.0579 (18)	0.0513 (19)	0.0369 (16)	0.0178 (15)	0.0026 (13)	0.0154 (14)
C43	0.0595 (18)	0.0426 (18)	0.0358 (15)	0.0053 (14)	0.0009 (13)	0.0152 (14)
C34	0.0572 (19)	0.072 (2)	0.0343 (15)	0.0181 (16)	0.0030 (13)	0.0229 (16)
N8	0.139 (3)	0.064 (2)	0.0304 (14)	0.0217 (19)	0.0057 (16)	0.0099 (14)
C23	0.0546 (18)	0.0477 (19)	0.0389 (16)	0.0070 (15)	0.0030 (14)	0.0093 (14)
C42	0.0566 (18)	0.057 (2)	0.0332 (15)	0.0070 (15)	0.0004 (13)	0.0123 (15)
C37	0.066 (2)	0.051 (2)	0.0363 (16)	0.0167 (16)	0.0026 (14)	0.0084 (15)
C13	0.0586 (19)	0.060 (2)	0.0368 (16)	0.0263 (16)	0.0008 (14)	0.0084 (15)
C39	0.063 (2)	0.0462 (19)	0.0448 (17)	0.0091 (15)	-0.0030 (14)	0.0090 (15)
N6	0.091 (2)	0.094 (3)	0.0376 (15)	0.0057 (19)	0.0053 (16)	0.0166 (16)
C28	0.0571 (19)	0.0473 (19)	0.0356 (16)	0.0021 (14)	0.0008 (14)	0.0054 (14)
N4	0.081 (2)	0.078 (2)	0.0371 (14)	0.0252 (16)	0.0047 (14)	0.0070 (14)
N3	0.097 (3)	0.115 (3)	0.0423 (16)	0.036 (2)	0.0021 (17)	0.0206 (18)
N5	0.078 (2)	0.0669 (19)	0.0357 (14)	0.0062 (15)	-0.0010 (14)	0.0074 (14)
C33	0.068 (2)	0.060 (2)	0.054 (2)	0.0214 (17)	0.0041 (16)	0.0254 (18)
C40	0.066 (2)	0.049 (2)	0.054 (2)	0.0092 (16)	-0.0025 (15)	0.0241 (17)
C27	0.066 (2)	0.072 (2)	0.0423 (18)	0.0084 (18)	0.0042 (15)	0.0138 (17)
C26	0.083 (3)	0.083 (3)	0.047 (2)	0.010 (2)	-0.0066 (19)	0.0183 (19)
C17	0.059 (2)	0.092 (3)	0.0444 (18)	0.0281 (19)	0.0010 (15)	0.0155 (18)

C19	0.064 (2)	0.084 (3)	0.0465 (18)	0.0295 (19)	0.0041 (16)	0.0138 (18)
C24	0.061 (2)	0.073 (2)	0.0446 (18)	-0.0025 (17)	0.0021 (15)	0.0102 (17)
C12	0.064 (2)	0.088 (3)	0.0443 (18)	0.0205 (19)	0.0021 (16)	0.0125 (18)
C22	0.058 (2)	0.082 (3)	0.0429 (18)	0.0045 (18)	-0.0003 (15)	0.0156 (18)
C16	0.069 (2)	0.105 (3)	0.0449 (19)	0.030 (2)	-0.0025 (17)	0.023 (2)
C25	0.071 (2)	0.079 (3)	0.0443 (19)	0.003 (2)	-0.0107 (16)	0.0053 (18)
C11	0.086 (3)	0.101 (3)	0.047 (2)	0.025 (2)	0.0108 (19)	0.024 (2)
C21	0.068 (2)	0.086 (3)	0.0453 (19)	0.0020 (19)	0.0060 (17)	0.0181 (19)
C20	0.075 (2)	0.092 (3)	0.0468 (19)	0.033 (2)	0.0143 (17)	0.0145 (19)
C14	0.061 (2)	0.119 (4)	0.050 (2)	0.024 (2)	-0.0042 (17)	0.020 (2)
C29	0.063 (2)	0.104 (3)	0.0454 (19)	0.015 (2)	0.0073 (16)	0.019 (2)
C30	0.081 (3)	0.117 (4)	0.045 (2)	0.011 (2)	0.0157 (19)	0.014 (2)
C15	0.081 (3)	0.150 (5)	0.051 (2)	0.030 (3)	-0.015 (2)	0.021 (3)
C8	0.066 (2)	0.048 (2)	0.067 (2)	0.0104 (17)	-0.0014 (17)	0.0148 (17)
C3	0.077 (2)	0.046 (2)	0.070 (2)	0.0068 (17)	0.000 (2)	0.0151 (18)
C9	0.075 (3)	0.067 (3)	0.079 (3)	-0.006 (2)	-0.006 (2)	0.020 (2)
N2	0.140 (4)	0.104 (3)	0.079 (3)	0.019 (3)	-0.002 (3)	0.041 (2)
C4	0.111 (3)	0.086 (3)	0.074 (3)	0.036 (3)	-0.006 (2)	0.022 (2)
C7	0.069 (2)	0.090 (3)	0.078 (3)	0.012 (2)	-0.005 (2)	0.019 (2)
N1	0.213 (6)	0.098 (3)	0.074 (3)	0.027 (4)	0.007 (3)	0.035 (3)
C10	0.110 (4)	0.095 (4)	0.087 (3)	-0.004 (3)	0.008 (3)	0.036 (3)
C2	0.089 (3)	0.083 (3)	0.080 (3)	-0.003 (2)	0.009 (2)	0.015 (2)
C6	0.100 (4)	0.117 (4)	0.080 (3)	0.023 (3)	-0.020 (3)	0.019 (3)
C5	0.179 (6)	0.121 (5)	0.081 (4)	0.062 (4)	-0.014 (4)	0.029 (3)
C1	0.150 (5)	0.103 (4)	0.082 (3)	-0.003 (4)	0.030 (3)	0.022 (3)

*Geometric parameters (Å, °)*

O4—C44	1.310 (4)	N5—C25	1.332 (5)
O4—H''	1.03 (4)	C33—H33A	0.9300
O2—C37	1.312 (4)	C40—H40A	0.9300
O2—H'	0.98 (4)	C27—C26	1.380 (4)
C38—C43	1.378 (4)	C27—H27A	0.9300
C38—C39	1.387 (4)	C26—H26A	0.9300
C38—C44	1.487 (4)	C17—C16	1.367 (4)
C31—C32	1.386 (4)	C17—H17A	0.9300
C31—C36	1.386 (4)	C19—C20	1.374 (4)
C31—C37	1.485 (4)	C19—H19A	0.9300
N7—C35	1.373 (4)	C24—C25	1.369 (5)
N7—H7A	0.8600	C24—H24A	0.9300
N7—H7B	0.8600	C12—C11	1.374 (5)
O3—C44	1.213 (4)	C12—H12A	0.9300
C35—C34	1.394 (4)	C22—C21	1.378 (4)
C35—C36	1.401 (4)	C22—H22A	0.9300
C41—C40	1.370 (4)	C16—H16A	0.9300
C41—C42	1.402 (5)	C25—H25A	0.9300
C41—H41A	0.9300	C11—H11A	0.9300
O1—C37	1.203 (4)	C21—H21A	0.9300
C32—C33	1.390 (4)	C20—H20A	0.9300
C32—H32A	0.9300	C14—C15	1.375 (5)



C18—C19	1.378 (4)	C14—H14A	0.9300
C18—C17	1.379 (4)	C29—C30	1.374 (5)
C18—C13	1.481 (4)	C29—H29A	0.9300
C36—H36A	0.9300	C30—H30A	0.9300
C43—C42	1.403 (4)	C15—H15A	0.9300
C43—H43A	0.9300	C8—C9	1.380 (5)
C34—C33	1.374 (4)	C8—C7	1.393 (5)
C34—H34A	0.9300	C8—C3	1.472 (5)
N8—C42	1.365 (4)	C3—C4	1.378 (5)
N8—H8A	0.8600	C3—C2	1.385 (6)
N8—H8B	0.8600	C9—C10	1.363 (6)
C23—C22	1.380 (4)	C9—H9A	0.9300
C23—C24	1.386 (4)	N2—C6	1.329 (6)
C23—C28	1.479 (4)	N2—C10	1.331 (6)
C13—C14	1.376 (4)	C4—C5	1.359 (6)
C13—C12	1.389 (5)	C4—H4B	0.9300
C39—C40	1.387 (4)	C7—C6	1.370 (6)
C39—H39A	0.9300	C7—H7C	0.9300
N6—C26	1.317 (5)	N1—C5	1.320 (7)
N6—C30	1.322 (5)	N1—C1	1.336 (7)
C28—C29	1.385 (4)	C10—H10A	0.9300
C28—C27	1.385 (4)	C2—C1	1.381 (6)
N4—C16	1.323 (4)	C2—H2B	0.9300
N4—C20	1.324 (4)	C6—H6B	0.9300
N3—C15	1.320 (5)	C5—H5B	0.9300
N3—C11	1.324 (5)	C1—H1B	0.9300
N5—C21	1.327 (4)		
C44—O4—H''	111 (2)	C16—C17—H17A	120.0
C37—O2—H'	111 (2)	C18—C17—H17A	120.0
C43—C38—C39	120.3 (3)	C20—C19—C18	119.5 (3)
C43—C38—C44	117.9 (3)	C20—C19—H19A	120.2
C39—C38—C44	121.9 (3)	C18—C19—H19A	120.2
C32—C31—C36	120.1 (3)	C25—C24—C23	119.5 (3)
C32—C31—C37	122.1 (3)	C25—C24—H24A	120.2
C36—C31—C37	117.8 (3)	C23—C24—H24A	120.2
C35—N7—H7A	120.0	C11—C12—C13	119.5 (3)
C35—N7—H7B	120.0	C11—C12—H12A	120.2
H7A—N7—H7B	120.0	C13—C12—H12A	120.2
N7—C35—C34	121.6 (3)	C21—C22—C23	119.8 (3)
N7—C35—C36	120.4 (3)	C21—C22—H22A	120.1
C34—C35—C36	118.0 (3)	C23—C22—H22A	120.1
C40—C41—C42	121.0 (3)	N4—C16—C17	123.3 (3)
C40—C41—H41A	119.5	N4—C16—H16A	118.3
C42—C41—H41A	119.5	C17—C16—H16A	118.3
O3—C44—O4	122.1 (3)	N5—C25—C24	123.7 (3)
O3—C44—C38	122.7 (3)	N5—C25—H25A	118.2
O4—C44—C38	115.2 (3)	C24—C25—H25A	118.2
C31—C32—C33	119.1 (3)	N3—C11—C12	123.9 (4)

C31—C32—H32A	120.5	N3—C11—H11A	118.0
C33—C32—H32A	120.5	C12—C11—H11A	118.0
C19—C18—C17	116.6 (3)	N5—C21—C22	123.3 (3)
C19—C18—C13	121.4 (3)	N5—C21—H21A	118.4
C17—C18—C13	122.0 (3)	C22—C21—H21A	118.4
C31—C36—C35	121.0 (3)	N4—C20—C19	123.5 (3)
C31—C36—H36A	119.5	N4—C20—H20A	118.2
C35—C36—H36A	119.5	C19—C20—H20A	118.2
C38—C43—C42	121.4 (3)	C15—C14—C13	119.6 (4)
C38—C43—H43A	119.3	C15—C14—H14A	120.2
C42—C43—H43A	119.3	C13—C14—H14A	120.2
C33—C34—C35	120.8 (3)	C30—C29—C28	119.3 (3)
C33—C34—H34A	119.6	C30—C29—H29A	120.3
C35—C34—H34A	119.6	C28—C29—H29A	120.3
C42—N8—H8A	120.0	N6—C30—C29	124.5 (4)
C42—N8—H8B	120.0	N6—C30—H30A	117.7
H8A—N8—H8B	120.0	C29—C30—H30A	117.7
C22—C23—C24	116.8 (3)	N3—C15—C14	124.3 (4)
C22—C23—C28	121.2 (3)	N3—C15—H15A	117.9
C24—C23—C28	121.9 (3)	C14—C15—H15A	117.9
N8—C42—C41	122.3 (3)	C9—C8—C7	116.4 (4)
N8—C42—C43	120.4 (3)	C9—C8—C3	121.6 (3)
C41—C42—C43	117.3 (3)	C7—C8—C3	122.0 (3)
O1—C37—O2	122.4 (3)	C4—C3—C2	116.8 (4)
O1—C37—C31	122.7 (3)	C4—C3—C8	121.6 (4)
O2—C37—C31	114.9 (3)	C2—C3—C8	121.5 (4)
C14—C13—C12	116.4 (3)	C10—C9—C8	120.3 (4)
C14—C13—C18	122.0 (3)	C10—C9—H9A	119.8
C12—C13—C18	121.6 (3)	C8—C9—H9A	119.8
C38—C39—C40	118.9 (3)	C6—N2—C10	116.2 (4)
C38—C39—H39A	120.5	C5—C4—C3	119.7 (5)
C40—C39—H39A	120.5	C5—C4—H4B	120.2
C26—N6—C30	116.3 (3)	C3—C4—H4B	120.2
C29—C28—C27	116.3 (3)	C6—C7—C8	119.2 (4)
C29—C28—C23	121.9 (3)	C6—C7—H7C	120.4
C27—C28—C23	121.8 (3)	C8—C7—H7C	120.4
C16—N4—C20	116.9 (3)	C5—N1—C1	115.9 (5)
C15—N3—C11	116.2 (3)	N2—C10—C9	123.7 (5)
C21—N5—C25	116.8 (3)	N2—C10—H10A	118.2
C34—C33—C32	121.0 (3)	C9—C10—H10A	118.2
C34—C33—H33A	119.5	C1—C2—C3	119.2 (5)
C32—C33—H33A	119.5	C1—C2—H2B	120.4
C41—C40—C39	121.0 (3)	C3—C2—H2B	120.4
C41—C40—H40A	119.5	N2—C6—C7	124.2 (4)
C39—C40—H40A	119.5	N2—C6—H6B	117.9
C26—C27—C28	119.8 (3)	C7—C6—H6B	117.9
C26—C27—H27A	120.1	N1—C5—C4	124.8 (5)
C28—C27—H27A	120.1	N1—C5—H5B	117.6
N6—C26—C27	123.8 (3)	C4—C5—H5B	117.6

N6—C26—H26A	118.1	N1—C1—C2	123.6 (6)
C27—C26—H26A	118.1	N1—C1—H1B	118.2
C16—C17—C18	120.0 (3)	C2—C1—H1B	118.2
C43—C38—C44—O3	-8.5 (4)	C22—C23—C24—C25	-1.0 (5)
C39—C38—C44—O3	172.5 (3)	C28—C23—C24—C25	178.0 (3)
C43—C38—C44—O4	172.2 (3)	C14—C13—C12—C11	0.9 (5)
C39—C38—C44—O4	-6.8 (4)	C18—C13—C12—C11	-179.8 (3)
C36—C31—C32—C33	0.5 (5)	C24—C23—C22—C21	0.9 (5)
C37—C31—C32—C33	179.5 (3)	C28—C23—C22—C21	-178.1 (3)
C32—C31—C36—C35	0.8 (4)	C20—N4—C16—C17	-2.4 (6)
C37—C31—C36—C35	-178.2 (3)	C18—C17—C16—N4	0.5 (6)
N7—C35—C36—C31	177.7 (3)	C21—N5—C25—C24	1.5 (6)
C34—C35—C36—C31	-1.7 (4)	C23—C24—C25—N5	-0.2 (6)
C39—C38—C43—C42	1.1 (4)	C15—N3—C11—C12	0.1 (6)
C44—C38—C43—C42	-178.0 (3)	C13—C12—C11—N3	-0.8 (6)
N7—C35—C34—C33	-178.1 (3)	C25—N5—C21—C22	-1.6 (6)
C36—C35—C34—C33	1.3 (4)	C23—C22—C21—N5	0.4 (6)
C40—C41—C42—N8	-179.1 (3)	C16—N4—C20—C19	2.6 (6)
C40—C41—C42—C43	1.0 (4)	C18—C19—C20—N4	-0.9 (6)
C38—C43—C42—N8	178.5 (3)	C12—C13—C14—C15	-0.4 (6)
C38—C43—C42—C41	-1.7 (4)	C18—C13—C14—C15	-179.7 (4)
C32—C31—C37—O1	172.3 (3)	C27—C28—C29—C30	0.3 (6)
C36—C31—C37—O1	-8.8 (5)	C23—C28—C29—C30	-179.0 (4)
C32—C31—C37—O2	-8.2 (4)	C26—N6—C30—C29	1.0 (7)
C36—C31—C37—O2	170.7 (3)	C28—C29—C30—N6	-0.6 (7)
C19—C18—C13—C14	-33.3 (5)	C11—N3—C15—C14	0.5 (7)
C17—C18—C13—C14	145.8 (4)	C13—C14—C15—N3	-0.3 (7)
C19—C18—C13—C12	147.5 (4)	C9—C8—C3—C4	32.3 (6)
C17—C18—C13—C12	-33.4 (5)	C7—C8—C3—C4	-146.3 (4)
C43—C38—C39—C40	0.3 (4)	C9—C8—C3—C2	-145.9 (4)
C44—C38—C39—C40	179.3 (3)	C7—C8—C3—C2	35.6 (6)
C22—C23—C28—C29	150.1 (4)	C7—C8—C9—C10	0.5 (6)
C24—C23—C28—C29	-28.8 (5)	C3—C8—C9—C10	-178.2 (4)
C22—C23—C28—C27	-29.2 (5)	C2—C3—C4—C5	-0.5 (7)
C24—C23—C28—C27	151.9 (3)	C8—C3—C4—C5	-178.7 (4)
C35—C34—C33—C32	0.0 (5)	C9—C8—C7—C6	0.5 (6)
C31—C32—C33—C34	-0.9 (5)	C3—C8—C7—C6	179.2 (4)
C42—C41—C40—C39	0.3 (5)	C6—N2—C10—C9	1.7 (8)
C38—C39—C40—C41	-0.9 (5)	C8—C9—C10—N2	-1.7 (8)
C29—C28—C27—C26	-0.5 (5)	C4—C3—C2—C1	0.6 (6)
C23—C28—C27—C26	178.8 (3)	C8—C3—C2—C1	178.9 (4)
C30—N6—C26—C27	-1.2 (6)	C10—N2—C6—C7	-0.6 (8)
C28—C27—C26—N6	1.0 (6)	C8—C7—C6—N2	-0.5 (8)
C19—C18—C17—C16	1.2 (5)	C1—N1—C5—C4	-0.2 (9)
C13—C18—C17—C16	-178.0 (3)	C3—C4—C5—N1	0.3 (9)
C17—C18—C19—C20	-1.0 (5)	C5—N1—C1—C2	0.4 (9)
C13—C18—C19—C20	178.1 (3)	C3—C2—C1—N1	-0.6 (8)

Hydrogen-bond geometry (Å, °)

Cg7 and Cg8 are the centroids of the C31–C36 and C38–C43 rings, respectively.

<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—H' $\cdots$ N4	0.98 (4)	1.67 (4)	2.647 (4)	171 (3)
O4—H'' $\cdots$ N5	1.03 (5)	1.60 (5)	2.627 (4)	175 (4)
N7—H7 <i>A</i> $\cdots$ N3 <sup>i</sup>	0.86	2.16	3.007 (4)	169 (4)
N7—H7 <i>B</i> $\cdots$ O3	0.86	2.23	3.038 (4)	156
N8—H8 <i>A</i> $\cdots$ N6 <sup>ii</sup>	0.86	2.16	3.006 (4)	169 (4)
N8—H8 <i>B</i> $\cdots$ O1	0.86	2.21	3.019 (4)	158
C36—H36 <i>A</i> $\cdots$ O3	0.93	2.52	3.300 (4)	141
C43—H43 <i>A</i> $\cdots$ O1	0.93	2.53	3.304 (4)	141
C7—H7 <i>C</i> $\cdots$ Cg7 <sup>iii</sup>	0.93	2.87	3.751 (4)	157
C17—H17 <i>A</i> $\cdots$ Cg8 <sup>iv</sup>	0.93	2.72	3.562 (4)	151
C19—H19 <i>A</i> $\cdots$ Cg8 <sup>v</sup>	0.93	2.78	3.643 (4)	154
C22—H22 <i>A</i> $\cdots$ Cg7 <sup>vi</sup>	0.93	2.72	3.567 (4)	151

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