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# Crystal structure and Hirshfeld surface analysis of 4-{[(anthracen-9-yl)methyl]amino}benzoic acid

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In the molecule of the title anthracene derivative,  $C_{22}H_{17}NO_2$ , the benzene ring is inclined to the mean plane of the anthracene ring system (r.m.s. deviation = 0.024 Å) by 75.21 (9)°. In the crystal, molecules are linked by pairs of  $O-H\cdots O$ hydrogen bonds, forming classical carboxylic acid inversion dimers with an  $R_2^2(8)$ ring motif. The dimers are linked by  $C-H\cdots\pi$  interactions, forming a supramolecular framework.

### 1. Chemical context

Anthraldehyde has been used in the synthesis of several Schiff base compounds that exhibit fluorescent properties as a result of strong  $\pi$ - $\pi$  conjugation (Asiri *et al.*, 2011; Pavitha *et al.*, 2017). Many complexes synthesized using anthraldehyde have shown remarkable sensing properties and have been used as chemo sensors (Obali & Ucan, 2012; Zhou *et al.*, 2012). Schiff base compounds are also of interest because of their biological applications, which include antibacterial, anticancer and antiviral (Asiri & Khan, 2010; Cheng *et al.*, 2010) activities. Herein, we report on the crystal and molecular structures of the title Schiff base compound, 4-{[(anthracen-9-yl)methyl]amino}benzoic acid, synthesized *via* reaction of 9-anthraldehyde with 4-aminobenzoic acid (PABA) followed by reduction with sodium borohydride.







### 2. Structural commentary

The molecular structure of the title compound is illustrated in Fig. 1. The molecule is non-planar, with the benzene ring (C2–C7) being inclined to the mean plane of the anthracene ring system (C9–C22; r.m.s. deviation = 0.024 Å) by 75.21 (9)°, and the torsional angle of the bridge, C5–N1–C8–C9, is 142.6 (2)°. The C8–N1 bond length of 1.457 (3) Å, is comparable to the C–N bond-length values obtained for the



Figure 1

The molecular structure of the tittle compound, with atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

similar ligand 5-[(anthracen-9-ylmethyl)amino]isophthalic acid (see §5. *Database survey*).

The C1=O2 and C1-O1 bond lengths of 1.238 (3) and 1.325 (3) Å, respectively, are in the expected ranges (Cambridge Structural Database; Groom *et al.*, 2016).

#### 3. Supramolecular features

In the crystal, a classical carboxylic acid inversion dimer is formed enclosing an  $R_2^2(8)$  ring motif (Table 1 and Fig. 2). The dimers pack along the *a*-axis direction in a herringbone fashion. They are linked by a series of  $C-H\cdots\pi$  interactions



Figure 2

A partial view along the b axis of crystal packing of the title compound. The hydrogen bonds (Table 1) are shown as dashed lines.



Figure 3

A view along the *b* axis of crystal packing of the title compound. The O- $H \cdots O$  hydrogen bonds and the C- $H \cdots \pi$  interactions are indicated by dashed lines (Table 1). For clarity, only the H atoms (grey balls) involved in these interactions have been included.

Table 1Hydrogen-bond geometry (Å, °).

Cg1, Cg2, and Cg4 are the centroids of the C2–C7, C9/C10/C15–C17/C22 and C17–C22 rings, respectively. Approximative geometrical parameters are given for the weak N–H.. $\pi$  interaction.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1-H1\cdots O2^i$	1.05 (4)	1.58 (3)	2.621 (3)	172 (3)
$N1-H1A\cdots Cg1^{ii}$	0.93 (3)	3.49	4.140	129
$C4-H4\cdots Cg4^{iii}$	0.93	2.98(1)	3.752 (3)	141 (1)
$C6-H6\cdots Cg1^{ii}$	0.93	2.69 (1)	3.410 (3)	135 (1)
$C16-H16\cdots Cg4^{iv}$	0.93	2.83 (1)	3.644 (3)	147 (1)
$C18-H18\cdots Cg2^{iv}$	0.93	2.69 (1)	3.452 (3)	140 (1)

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

(Table 1 and Fig. 3), forming a supramolecular three-dimensional structure. The NH hydrogen atom (H1A) is not involved in hydrogen bonding but is directed towards the benzene ring (C2–C7). Approximate geometrical details of this weak N–H··· $\pi$  interaction are given in Table 1.

### 4. Hirshfeld analysis

The Hirshfeld surface analysis (Spackman & Jayatilaka, 2009) and the associated two-dimensional fingerprint plots (McKinnon *et al.*, 2007) were performed with *Crystal-Explorer17* (Turner *et al.*, 2017). The Hirshfeld surfaces are colour-mapped with the normalized contact distance,  $d_{\text{norm}}$ , from red (distances shorter than the sum of the van der Waals radii) through white to blue (distances longer than the sum of the van der Waals radii).

The Hirshfeld surface of the title compound mapped over  $d_{\text{norm}}$ , in the colour range -0.7519 to 1.6997 a.u., is given in Fig. 4. The positions of the strong  $O-H\cdots O$  hydrogen bonds are indicated by the red regions on the Hirshfeld surface.





The Hirshfeld surface of the title compound mapped over  $d_{\text{norm}}$ , in the colour range -0.7519 to 1.6997 a.u..

### research communications



Figure 5

(a) The two-dimensional fingerprint plots of the title compound, and delineated into (b)  $H \cdots H$  (42.7%), (c)  $C \cdots H/H \cdots C$  (40.0%) and (d)  $O \cdots H/H \cdots O$  (12.3%) contacts.

The two-dimensional fingerprint plots are given in Fig. 5. They reveal that the principal contributions to the overall surface involve  $H \cdots H$  contacts at 42.7% (Fig. 5b), followed by  $C \cdots H/H \cdots C$  contacts at 40.0% (Fig. 5c) and  $O \cdots H/H \cdots O$ contacts at 12.3% (Fig. 5d). Apart from the  $C \cdots C$  contacts, contributing 2.1%, all other atom ... atom contact contributions are negligible.

#### 5. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.40, update August 2019; Groom et al., 2016) for the N-(anthracen-9-ylmethyl)aniline skeleton gave six hits (see supporting information file S1), all of which concern polymeric metal complexes of the ligand 5-[(anthracen-9-vlmethyl)amino]isophthalic acid; for example, a series of four gadolinium coordination polymers (CSD refcodes VOLSOG, VOLSUM, VOLTAT, VOLTIB; Singh et al., 2014). The bridging C–N bond length varies from ca.1.389 to 1.494 Å, compared to the C8-N1 bond length of 1.457 (3) Å in the title compound.

A search for the 1-(anthracen-9-yl)-N-phenylmethanimine skeleton gave 21 hits (see supporting information file S2). none of which involve a benzoic acid moiety.

### 6. Synthesis and crystallization

4-Aminobenzoic acid (0.33 g, 2.42 mmol) was added to a solution of 9-anthraldehyde (0.5 g, 2.42 mmol) dissolved in ethanol and the whole mixture was heated at 343 K under reflux for 5-6 h. The mixture was then stirred at room temperature to obtain a yellow prec product, which was monitored through precipitate, which was then air dried, was yield. This was further reduced with sod taken in excess (0.183 g, 4.84 mmol) by temperature at 277-278 K until the colour had changed from bright yellow to dull yello was filtered, washed with water and acidified The product thus obtained was dissolved in kept for crystallization. Block-like pale-yellow crystals of the title compound were obtained after a few days.

### 7. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The OH and NH hydrogen atoms

u at 545 K under	
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lium borohydride	-
maintaining the	Deferences
of the precipitate	Kelerences
w. The precipitate	Asiri, A. M., Al-Youbi, A. O., Khan,
d with acetic acid.	<i>Cryst.</i> E <b>67</b> , o3419.
n hot ethanol and	Asiri, A. M. & Khan, S. A. (2010). <i>M</i>
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Temperature (K)	100
a, b, c (Å)	14.985 (2), 6.0116 (9), 19.106 (3)
$\beta$ (°)	106.796 (5)
$V(\dot{A}^3)$	1647.7 (4)
Z	4
Radiation type	Μο Κα
$\mu (\text{mm}^{-1})$	0.09
Crystal size (mm)	$0.4 \times 0.27 \times 0.18$
Data collection	
Diffractometer	Bruker APEXII CCD
Absorption correction	Multi-scan (SADABS; Krause et al., 2015)
$T_{\min}, T_{\max}$	0.629, 0.746
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	25595, 2913, 1975
R <sub>int</sub>	0.118
$(\sin \theta / \lambda)_{\rm max} ({\rm \AA}^{-1})$	0.596
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.048, 0.141, 1.12
No. of reflections	2913
No. of parameters	235
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta \rho_{\rm max},  \Delta \rho_{\rm min}  ({\rm e}  {\rm \AA}^{-3})$	0.38, -0.32

C22H17NO2 327.39

Monoclinic, P21/c

et al.. 2015), olex2.refine (Bourhis et al., 2015), OLEX2 (Dolomanov et al., 2009), Mercury (Macrae et al., 2008), PLATON (Spek, 2009) and publCIF (Westrip, 2010).

were located in a difference-Fourier map and refined freely. The C-bound H atoms were included in calculated positions and allowed to ride on their parent C atom: C-H = 0.93-0.97Å with  $U_{iso}(H) = 1.2U_{eq}(C)$ .

### Acknowledgements

Table 2

 $M_{r}$ 

Crystal data Chemical formula

Experimental details.

Crystal system, space group

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### Crystal structure and Hirshfeld surface analysis of 4-{[(anthracen-9-yl)methyl]amino}benzoic acid

### Adeeba Ahmed, Md. Serajul Haque Faizi, Aiman Ahmad, Musheer Ahmad and Igor O. Fritsky

### **Computing details**

Data collection: *APEX2* (Bruker, 2016); cell refinement: *SAINT* (Bruker, 2016); data reduction: *SAINT* (Bruker, 2016); program(s) used to solve structure: *olex2.solve* (Bourhis *et al.*, 2015); program(s) used to refine structure: *olex2.refine* (Bourhis *et al.*, 2015); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009), *PLATON* (Spek, 2009) and *publCIF* (Westrip, 2010).

4-{[(Anthracen-9-yl)methyl]amino}benzoic acid

### Crystal data

 $C_{22}H_{17}NO_2$   $M_r = 327.39$ Monoclinic,  $P2_1/c$  a = 14.985 (2) Å b = 6.0116 (9) Å c = 19.106 (3) Å  $\beta = 106.796$  (5)° V = 1647.7 (4) Å<sup>3</sup> Z = 4

### Data collection

Bruker APEXII CCD diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (SADABS; Krause *et al.*, 2015)  $T_{\min} = 0.629$ ,  $T_{\max} = 0.746$ 25595 measured reflections

### Refinement

Refinement on  $F^2$ Least-squares matrix: full  $R[F^2 > 2\sigma(F^2)] = 0.048$  $wR(F^2) = 0.141$ S = 1.122913 reflections 235 parameters 0 restraints 29 constraints F(000) = 688.3239  $D_x = 1.320 \text{ Mg m}^{-3}$ Mo K\alpha radiation,  $\lambda = 0.71073 \text{ Å}$ Cell parameters from 3139 reflections  $\theta = 3.1-28.2^{\circ}$   $\mu = 0.09 \text{ mm}^{-1}$  T = 100 KBlock, pale-yellow  $0.4 \times 0.27 \times 0.18 \text{ mm}$ 

2913 independent reflections 1975 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.118$   $\theta_{max} = 25.1^{\circ}, \ \theta_{min} = 2.8^{\circ}$   $h = -20 \rightarrow 20$   $k = -8 \rightarrow 8$  $l = -25 \rightarrow 25$ 

Primary atom site location: iterative H atoms treated by a mixture of independent and constrained refinement  $w = 1/[\sigma^2(F_o^2) + (0.0452P)^2 + 0.9105P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{max} = 0.0003$  $\Delta\rho_{max} = 0.38 \text{ e} \text{ Å}^{-3}$  $\Delta\rho_{min} = -0.32 \text{ e} \text{ Å}^{-3}$ 

	x	у	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
01	0.47155 (12)	0.7404 (3)	0.95012 (10)	0.0304 (5)	
02	0.57543 (12)	1.0125 (3)	0.95315 (10)	0.0317 (5)	
N1	0.66609 (14)	0.3342 (4)	0.72544 (12)	0.0252 (5)	
C1	0.54000 (17)	0.8318 (4)	0.92904 (14)	0.0238 (6)	
C2	0.57055 (16)	0.7035 (4)	0.87497 (13)	0.0212 (6)	
C3	0.63813 (16)	0.7902 (4)	0.84570 (14)	0.0230 (6)	
H3	0.66266 (16)	0.9303 (4)	0.86054 (14)	0.0276 (7)*	
C4	0.66973 (17)	0.6741 (4)	0.79531 (13)	0.0222 (6)	
H4	0.71436 (17)	0.7367 (4)	0.77612 (13)	0.0266 (7)*	
C5	0.63421 (16)	0.4606 (4)	0.77302 (13)	0.0209 (6)	
C6	0.56515 (17)	0.3731 (4)	0.80193 (13)	0.0224 (6)	
H6	0.53997 (17)	0.2335 (4)	0.78701 (13)	0.0269 (7)*	
C7	0.53463 (16)	0.4920 (4)	0.85202 (13)	0.0215 (6)	
H7	0.48944 (16)	0.4312 (4)	0.87096 (13)	0.0258 (7)*	
C8	0.74688 (17)	0.3903 (4)	0.70105 (14)	0.0243 (6)	
H8a	0.72642 (17)	0.4653 (4)	0.65418 (14)	0.0291 (7)*	
H8b	0.78696 (17)	0.4913 (4)	0.73586 (14)	0.0291 (7)*	
C9	0.80133 (16)	0.1835 (4)	0.69386 (13)	0.0208 (6)	
C10	0.80253 (16)	0.1014 (4)	0.62496 (13)	0.0204 (6)	
C11	0.75646 (17)	0.2083 (4)	0.55691 (14)	0.0257 (6)	
H11	0.72308 (17)	0.3384 (4)	0.55740 (14)	0.0308 (7)*	
C12	0.76034 (18)	0.1242 (5)	0.49175 (14)	0.0294 (7)	
H12	0.72945 (18)	0.1972 (5)	0.44864 (14)	0.0353 (8)*	
C13	0.81084 (18)	-0.0732 (5)	0.48869 (15)	0.0310 (7)	
H13	0.81360 (18)	-0.1278 (5)	0.44382 (15)	0.0372 (8)*	
C14	0.85503 (17)	-0.1822 (4)	0.55102 (14)	0.0269 (6)	
H14	0.88740 (17)	-0.3124 (4)	0.54841 (14)	0.0322 (7)*	
C15	0.85289 (16)	-0.1007 (4)	0.62098 (14)	0.0216 (6)	
C16	0.89943 (16)	-0.2114 (4)	0.68500 (14)	0.0225 (6)	
H16	0.93082 (16)	-0.3429 (4)	0.68203 (14)	0.0270 (7)*	
C17	0.90025 (16)	-0.1302 (4)	0.75358 (13)	0.0202 (6)	
C18	0.94936 (16)	-0.2430 (4)	0.81929 (14)	0.0255 (6)	
H18	0.98095 (16)	-0.3741 (4)	0.81619 (14)	0.0307 (7)*	
C19	0.95099 (18)	-0.1630 (4)	0.88627 (15)	0.0289 (7)	
H19	0.98374 (18)	-0.2383 (4)	0.92840 (15)	0.0347 (8)*	
C20	0.90250 (17)	0.0354 (4)	0.89133 (15)	0.0287 (6)	
H20	0.90346 (17)	0.0897 (4)	0.93712 (15)	0.0344 (8)*	
C21	0.85437 (17)	0.1482 (4)	0.83019 (14)	0.0243 (6)	
H21	0.82321 (17)	0.2784 (4)	0.83510 (14)	0.0291 (7)*	
C22	0.85075 (16)	0.0706 (4)	0.75862 (14)	0.0204 (6)	
H1	0.457 (2)	0.850 (6)	0.9884 (19)	0.074 (11)*	
H1a	0.635 (2)	0.201 (5)	0.7113 (15)	0.042 (9)*	
		(-)			

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters  $(\hat{A}^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
01	0.0321 (10)	0.0298 (10)	0.0357 (12)	-0.0061 (8)	0.0198 (9)	-0.0083 (9)
O2	0.0368 (11)	0.0274 (10)	0.0352 (12)	-0.0070 (9)	0.0172 (9)	-0.0115 (9)
N1	0.0215 (12)	0.0251 (12)	0.0322 (14)	-0.0041 (10)	0.0127 (10)	-0.0071 (10)
C1	0.0234 (14)	0.0238 (14)	0.0239 (15)	0.0007 (11)	0.0065 (11)	0.0001 (11)
C2	0.0200 (13)	0.0223 (13)	0.0209 (14)	0.0021 (11)	0.0053 (11)	-0.0011 (11)
C3	0.0204 (13)	0.0206 (13)	0.0275 (15)	0.0013 (11)	0.0059 (11)	-0.0024 (11)
C4	0.0197 (13)	0.0236 (13)	0.0242 (14)	-0.0003 (11)	0.0078 (11)	0.0018 (11)
C5	0.0195 (13)	0.0207 (13)	0.0219 (14)	0.0026 (10)	0.0051 (11)	0.0004 (11)
C6	0.0215 (13)	0.0194 (13)	0.0251 (15)	0.0012 (11)	0.0050 (11)	-0.0012 (11)
C7	0.0185 (13)	0.0229 (13)	0.0230 (14)	0.0015 (11)	0.0058 (11)	0.0036 (11)
C8	0.0234 (13)	0.0218 (13)	0.0295 (16)	-0.0009 (11)	0.0105 (12)	-0.0011 (11)
С9	0.0193 (13)	0.0202 (13)	0.0241 (14)	-0.0013 (10)	0.0081 (11)	-0.0007 (11)
C10	0.0147 (12)	0.0227 (13)	0.0242 (15)	-0.0019 (10)	0.0063 (11)	-0.0005 (11)
C11	0.0216 (13)	0.0277 (14)	0.0279 (15)	0.0013 (11)	0.0076 (11)	0.0009 (12)
C12	0.0250 (14)	0.0397 (16)	0.0215 (15)	0.0021 (12)	0.0037 (12)	0.0020 (12)
C13	0.0267 (14)	0.0393 (16)	0.0258 (16)	0.0018 (13)	0.0057 (12)	-0.0067 (13)
C14	0.0238 (13)	0.0287 (14)	0.0276 (15)	0.0006 (12)	0.0066 (12)	-0.0089 (12)
C15	0.0170 (12)	0.0235 (13)	0.0239 (15)	-0.0030 (10)	0.0054 (11)	-0.0043 (11)
C16	0.0202 (13)	0.0190 (13)	0.0297 (15)	0.0000 (11)	0.0094 (11)	-0.0029 (11)
C17	0.0171 (12)	0.0212 (13)	0.0235 (14)	-0.0022 (10)	0.0076 (11)	0.0009 (11)
C18	0.0195 (13)	0.0249 (14)	0.0318 (16)	-0.0010 (11)	0.0066 (11)	0.0024 (12)
C19	0.0234 (14)	0.0345 (16)	0.0272 (16)	-0.0043 (12)	0.0047 (11)	0.0064 (12)
C20	0.0285 (15)	0.0338 (15)	0.0245 (16)	-0.0051 (12)	0.0087 (12)	-0.0032 (12)
C21	0.0216 (13)	0.0261 (14)	0.0268 (15)	-0.0023 (11)	0.0096 (11)	-0.0029 (12)
C22	0.0181 (12)	0.0203 (13)	0.0242 (14)	-0.0054 (10)	0.0085 (11)	-0.0030 (11)

Atomic displacement parameters  $(Å^2)$ 

Geometric parameters (Å, °)

01—C1	1.325 (3)	C10—C15	1.443 (3)
01—H1	1.05 (4)	C11—H11	0.9300
O2—C1	1.238 (3)	C11—C12	1.360 (3)
N1—C5	1.372 (3)	C12—H12	0.9300
N1—C8	1.457 (3)	C12—C13	1.418 (4)
N1—H1a	0.92 (3)	C13—H13	0.9300
C1—C2	1.465 (3)	C13—C14	1.353 (4)
С2—С3	1.392 (3)	C14—H14	0.9300
С2—С7	1.401 (3)	C14—C15	1.433 (3)
С3—Н3	0.9300	C15—C16	1.390 (3)
С3—С4	1.379 (3)	C16—H16	0.9300
C4—H4	0.9300	C16—C17	1.395 (3)
C4—C5	1.408 (3)	C17—C18	1.429 (3)
C5—C6	1.408 (3)	C17—C22	1.435 (3)
С6—Н6	0.9300	C18—H18	0.9300
C6—C7	1.375 (3)	C18—C19	1.360 (4)
С7—Н7	0.9300	С19—Н19	0.9300

С8—Н8а 0.9700 С19—С20	1.414 (4)
C8—H8b 0.9700 C20—H20	0.9300
C8—C9 1.515 (3) C20—C21	1.363 (3)
C9—C10 1.411 (3) C21—H21	0.9300
C9—C22 1.418 (3) C21—C22	1.431 (3)
C10—C11 1.436 (3)	
H1—O1—C1 106.6 (18) H11—C11—C10	119.13 (14)
C8—N1—C5 124.4 (2) C12—C11—C10	121.7 (2)
H1a—N1—C5 115.7 (18) C12—C11—H11	119.13 (16)
H1a—N1—C8 119.8 (18) H12—C12—C11	119.57 (16)
O2—C1—O1 122.5 (2) C13—C12—C11	120.9 (3)
C2—C1—O1 115.1 (2) C13—C12—H12	119.57 (16)
C2—C1—O2 122.5 (2) H13—C13—C12	119.99 (16)
C3—C2—C1 119.9 (2) C14—C13—C12	120.0 (3)
C7—C2—C1 121.9 (2) C14—C13—H13	119.99 (16)
C7—C2—C3 118.2 (2) H14—C14—C13	119.39 (16)
H3—C3—C2 119.07 (15) C15—C14—C13	121.2 (2)
C4—C3—C2 121.9 (2) C15—C14—H14	119.39 (15)
C4—C3—H3 119.07 (15) C14—C15—C10	119.3 (2)
H4—C4—C3 120.11 (15) C16—C15—C10	119.6 (2)
C5—C4—C3 119.8 (2) C16—C15—C14	121.1 (2)
C5—C4—H4 120.11 (14) H16—C16—C15	119.16 (14)
C4—C5—N1 122.1 (2) C17—C16—C15	121.7 (2)
C6—C5—N1 119.3 (2) C17—C16—H16	119.16 (14)
C6-C5-C4 118.6 (2) C18-C17-C16	121.5 (2)
H6—C6—C5 119.71 (14) C22—C17—C16	119.5 (2)
C7—C6—C5 120.6 (2) C22—C17—C18	119.0 (2)
С7—С6—Н6 119.71 (15) Н18—С18—С17	119.22 (15)
C6—C7—C2 121.0 (2) C19—C18—C17	121.6 (2)
H7—C7—C2 119.49 (14) C19—C18—H18	119.22 (16)
H7—C7—C6 119.49 (15) H19—C19—C18	120.26 (16)
H8a—C8—N1 109.44 (13) C20—C19—C18	119.5 (3)
H8b—C8—N1 109.44 (13) C20—C19—H19	120.26 (16)
H8b—C8—H8a 108.0 H20—C20—C19	119.47 (16)
C9—C8—N1 111.0 (2) C21—C20—C19	121.1 (3)
C9—C8—H8a 109.44 (14) C21—C20—H20	119.47 (16)
C9—C8—H8b 109.44 (13) H21—C21—C20	119.29 (16)
C10—C9—C8 121.6 (2) C22—C21—C20	121.4 (2)
C22—C9—C8 118.2 (2) C22—C21—H21	119.29 (14)
C22—C9—C10 120.2 (2) C17—C22—C9	119.6 (2)
C11—C10—C9 123.8 (2) C21—C22—C9	123.0 (2)
C15—C10—C9 119.4 (2) C21—C22—C17	117.4 (2)
C15—C10—C11 116.8 (2)	
$C_{2}$ N1 C5 C4 $S_{2}$ $S_{3}$ (4) C0 C10 C11 C12	170.2 (2)
0-11-0-11-01-012	1/9.3 (3)
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	-0.3(4)

O1—C1—C2—C3	176.1 (2)	C9-C10-C15-C16	0.0 (4)
O1—C1—C2—C7	-4.8 (4)	C11—C10—C15—C14	0.4 (3)
O2—C1—C2—C3	-3.8 (4)	C11—C10—C15—C16	179.6 (2)
O2—C1—C2—C7	175.3 (2)	C10-C11-C12-C13	-0.3 (4)
C1—C2—C3—C4	179.2 (2)	C11—C12—C13—C14	0.8 (4)
C7—C2—C3—C4	0.1 (4)	C12—C13—C14—C15	-0.7 (4)
C1—C2—C7—C6	-179.1 (2)	C13-C14-C15-C10	0.1 (4)
C3—C2—C7—C6	0.0 (4)	C13—C14—C15—C16	-179.1 (3)
C2—C3—C4—C5	-0.8 (4)	C10-C15-C16-C17	-1.1 (4)
C3—C4—C5—N1	-177.2 (2)	C14—C15—C16—C17	178.0 (2)
C3—C4—C5—C6	1.4 (4)	C15—C16—C17—C18	-179.1 (2)
N1-C5-C6-C7	177.3 (2)	C15—C16—C17—C22	1.0 (4)
C4—C5—C6—C7	-1.4 (4)	C16—C17—C18—C19	179.4 (2)
C5—C6—C7—C2	0.7 (4)	C22-C17-C18-C19	-0.7 (4)
N1-C8-C9-C10	109.5 (3)	C16—C17—C22—C9	0.3 (4)
N1-C8-C9-C22	-69.6 (3)	C16—C17—C22—C21	-179.5 (2)
C8—C9—C10—C11	2.5 (4)	C18—C17—C22—C9	-179.7 (2)
C8—C9—C10—C15	-177.9 (2)	C18—C17—C22—C21	0.6 (4)
C22—C9—C10—C11	-178.4 (2)	C17-C18-C19-C20	0.5 (4)
C22—C9—C10—C15	1.3 (4)	C18—C19—C20—C21	-0.2 (4)
C8—C9—C22—C17	177.8 (2)	C19—C20—C21—C22	0.1 (4)
C8—C9—C22—C21	-2.5 (4)	C20—C21—C22—C9	179.9 (3)
C10-C9-C22-C17	-1.4 (4)	C20—C21—C22—C17	-0.4 (4)
C10-C9-C22-C21	178.3 (2)		

### Hydrogen-bond geometry (Å, °)

Cg1, Cg2, and Cg4 are the centroids of the C2–C7, C9/C10/C15–C17/C22 and C17–C22 rings, respectively. Approximative geometrical parameters are given for the weak N—H. $\pi$  interaction.

<i>D</i> —Н	H···A	$D \cdots A$	D—H···A
1.05 (4)	1.58 (3)	2.621 (3)	172 (3)
0.93 (3)	3.49	4.140	129
0.93	2.98 (1)	3.752 (3)	141 (1)
0.93	2.69(1)	3.410 (3)	135 (1)
0.93	2.83 (1)	3.644 (3)	147 (1)
0.93	2.69 (1)	3.452 (3)	140 (1)
	<i>D</i> —H 1.05 (4) 0.93 (3) 0.93 0.93 0.93 0.93 0.93	$D$ —H $H \cdots A$ 1.05 (4)1.58 (3)0.93 (3)3.490.932.98 (1)0.932.69 (1)0.932.83 (1)0.932.69 (1)	$D$ —H $H \cdots A$ $D \cdots A$ 1.05 (4)1.58 (3)2.621 (3)0.93 (3)3.494.1400.932.98 (1)3.752 (3)0.932.69 (1)3.410 (3)0.932.83 (1)3.644 (3)0.932.69 (1)3.452 (3)

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