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Crystal structure and Hirshfeld analysis of 2-[bis(1-methyl-1*H*-indol-3-yl)methyl]benzoic acid

Suhaila Sapari,^a Sheryn Wong,^a Mohammad Fadzlee Ngatiman,^b Huda Misral^a and
Siti Aishah Hasbullah^{a*}^aCentre of Advanced Materials and Renewable Resources, Faculty of Science and Technology, National University of
Malaysia, 43600 UKM Bangi, Selangor, Malaysia, and ^bCenter for Research and Instrumentation Management, Universiti
Kebangsaan Malaysia, 43600 UKM Bangi, Selangor, Malaysia. *Correspondence e-mail: aishah80@ukm.edu.my

In the title compound, C₂₆H₂₂N₂O₂, the dihedral angles between the 1-methylindole units (*A* and *B*) and the benzoic acid moiety (*C*) are *A/B* = 64.87 (7), *A/C* = 80.92 (8) and *B/C* = 75.05 (8)°. An intramolecular C—H···O interaction arising from the methyne group helps to establish the conformation. In the crystal, *R*₂²(8) carboxylic acid inversion dimers linked by pairs of O—H···O hydrogen bonds are observed. A Hirshfeld surface analysis shows that the greatest contributions are from H···H, C···H/H···C and O···H/H···O contacts (percentage values = 54.6%, 29.6% and 10.1%, respectively).

1. Chemical context

Bisindolyl methane and its derivatives are relatively easy to synthesize and show a broad spectrum of potential biological activities: for example, bis(indolyl)imidazole shows antiplasmodial activity towards *plasmodium falciparum* (Alvarado *et al.*, 2013). Furthermore, they also have good potential as antibacterial (Imran *et al.*, 2014; Challa *et al.*, 2017), antileishmanial (Bharate *et al.*, 2013), antitumor (Carbone *et al.*, 2013), antiplatelet (Grumel *et al.*, 2002) and anticancer (Guo *et al.*, 2010; Jamsheena *et al.*, 2016) agents. Oxidized bis(indolyl)methanes containing an acidic hydrogen-bond-donor group and a basic hydrogen-bond-acceptor group can act as selective colorimetric sensors for either F[−] or HSO₄[−] in an aprotic solvent (He *et al.*, 2006). Arylfuryl-bis(-indolyl)methanes have selective chromogenic and fluorogenic ratiometric receptors for the mercury ion in aqueous solution (Batista *et al.*, 2014). As part of our studies in this area, we now report the acid-catalysed condensation reaction between carboxy benzaldehyde and indole to generate the title compound.

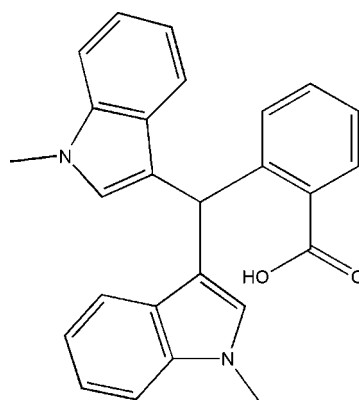
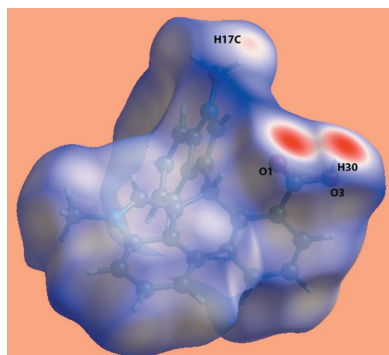


Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O3-H3O\cdots O1^i$	0.82 (4)	1.89 (5)	2.679 (3)	163 (6)
$C8-H8\cdots O1$	0.98	2.20	2.945 (4)	132

Symmetry code: (i) $-x, -y + 2, -z + 2$.

2. Structural commentary

The title compound (Fig. 1) crystallizes in the triclinic system with space group $P\bar{1}$ and $Z = 2$. The molecule consists of two methylated indole ring systems [C8–C17/N1 (*A*) and C18–C26/N2 (*B*)] and a benzoic acid [C1–C7 (*C*)] system linked *via* the tertiary C8 atom, with dihedral angles between them of $A/B = 64.87$ (7), $A/C = 80.92$ (8) and $B/C = 75.05$ (8)°. Significant torsion angles include $C7-C8-C9-C12$ [67.3 (3)] and $C7-C8-C18-C21$ [50.2 (3)°]. An intramolecular $C8-H8\cdots O1$ hydrogen bond (Table 1) may help to establish the conformation.

3. Supramolecular features

In the crystal of the title compound, neighbouring molecules are connected into dimers with an $R_2^2(8)$ graph-set motif *via* pairwise $O3-H3O\cdots O1$ hydrogen bonds (Table 1, Fig. 2).

4. Hirshfeld surface analysis

The Hirshfeld surface and fingerprint (FP) plots for the title compound were generated using *CrystalExplorer17* (McKinnon *et al.*, 2007). A view of the Hirshfeld surface mapped over d_{norm} is shown in Fig. 3. The intense red spots

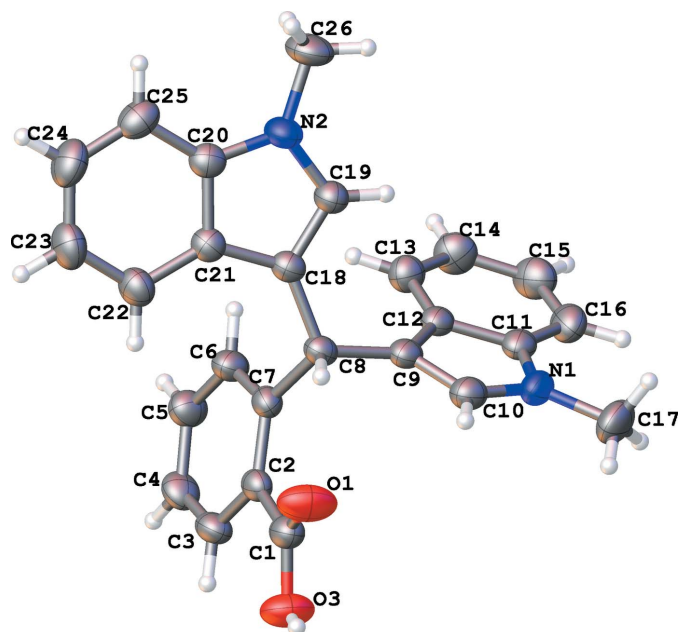


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level.

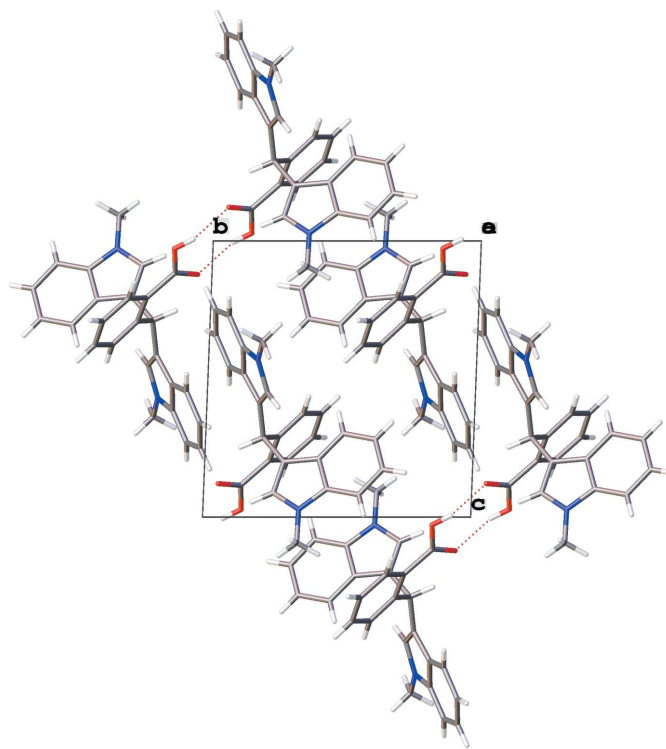


Figure 2

Crystal packing of the title compound viewed down [100] showing inversion dimers linked by pairs of $O-H\cdots O$ hydrogen bonds (dashed lines; Table 1).

near the O1-carbonyl and H30-benzoic acid atoms indicate the short interatomic $O\cdots H/H\cdots O$ contacts relating to the hydrogen bond given in Table 1. The two-dimensional fingerprint plots for the $H\cdots H$, $O\cdots H/H\cdots O$, $C\cdots H/H\cdots C$,

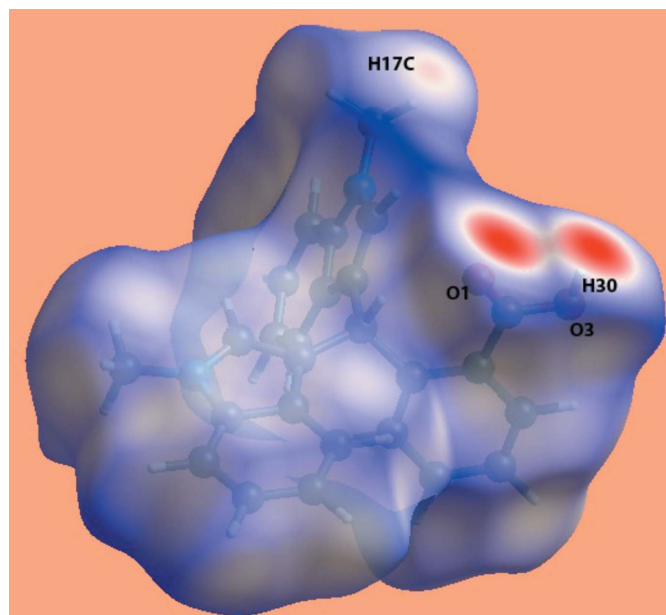


Figure 3

View of the Hirshfeld surface of the title compound mapped over d_{norm} in the range -0.68 to $+1.45$ au.

Table 2

Percentage contributions of interatomic contacts to the Hirshfeld surface of the title compound.

Contact	Percentage contribution
H...H	54.6
O...H/H...O	10.1
C...H/H...C	29.6
N...H/H...N	1.1
C...N/C...N	1.7
C...C	2.8

N...H/H...N, C...C and C...N/C...N contacts are illustrated in Fig. 4. The percentage contributions from the different interatomic contact to the Hirshfeld surface are summarized in Table 2. The fingerprint plot for the H...H contacts, which make the largest contribution to the Hirshfeld surface (54.6%), has a broad appearance with a single tip at $d_e + d_i = 2.2 \text{ \AA}$. The FP plot for the O...H/H...O (10.1%) contacts has prominent ‘forceps-like’ tips at $d_e + d_i = 1.7 \text{ \AA}$, whereas that for C...H/H...C contacts (29.6%) shows two pairs of adjacent peaks with $d_e + d_i = 2.6 \text{ \AA}$. The other remaining interatomic contacts, which make a small percentage contribution, have a negligible effect on the packing.

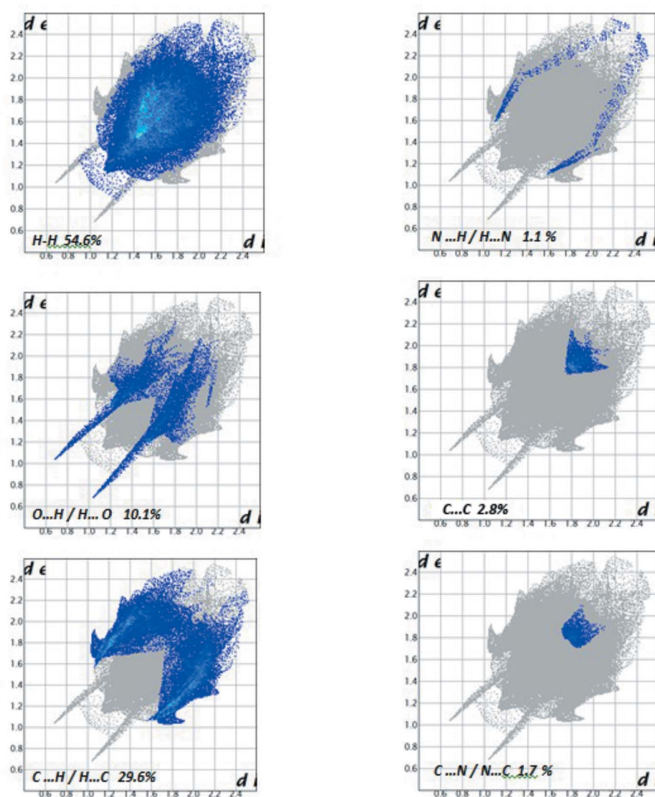


Figure 4
Two-dimensional fingerprint plots of the title compound delineated into H...H, O...H/H...O, C...H/H...C, N...H/H...N, C...C, C...N/N...C contacts.

5. Database survey

A search of the Cambridge Structural Database (Groom *et al.*, 2016) revealed only seven structures of bis(indole-3-yl) derivatives. These include 3,5-bis(indol-3-yl)-1,2,4-triazin-6(1*H*,6*H*)-one methanol solvate (FOLSOP) and 3,6-bis(indol-3-yl)-1,2,4-triazin-4(1*H*,4*H*)-one dimethylformamide solvate (FOLTAC; Garg & Stoltz, 2005), bis(indol-3-yl)(*p*-tolyl)methane (HODROH; Krishna *et al.*, 1999), 1,1-bis(indol-3-yl)-1-phenylethane (MEDJEK; Ganesan *et al.*, 2000), *cyclo-N,N'*-(α,α' -*p*-xylyl)bis(indol-3-yl)-*N*-methylmaleimide (UJALOG), *cyclo-N,N'*-(α,α' -*m*-xylyl)bis(indol-3-yl)-*N*-methylmaleimide (UJALUM) and *cyclo-N,N'*-[1,11-(3,6,9-trioxaundecyl)]bis(indol-3-yl)-*N*-methylmaleimide (UJAMAT; Mandl *et al.*, 2003). Two of these entries (MEDJEK and HODROH) are closely related to the title compound. Two of these entries (MEDJEK and HODROH) are closely related to the title compound with dihedral angles between the 1-methyl indole units of 63.4 (2) and 73.06 (19)° for the two independent molecules in MEDJEK and of 80.8 (1)° in HODROH [64.87 (7)° in the title compound]. In another related compound 4-[bis(1*H*-indol-3-yl)methyl]benzotrile (Deng *et al.*, 2011), the dihedral angle is 72.08 (6)°.

6. Synthesis and crystallization

Equimolar amounts of 2-carboxybenzaldehyde (3.0 mmol) and 1-methylindole (3.0 mmol) was mixed in a reaction vessel. A few drops of anhydrous acetic acid was added and the mixture was then irradiated in a domestic microwave oven at 100 W for 5 min. The crude product obtained was purified by recrystallization from an acetone–EtOH solvent mixture (*v:v* = 1:2) to give the pure product in 13.3% yield. IR (ATR, ν_{max} /cm⁻¹): 3058, 2930 (*broad*, O–H), 1676 (C=O), 1473 (C=C), 1331–1067 (C–O, C–N), 731. ¹H NMR (400 MHz, DMSO-*d*₆) δ (ppm): 3.67 (*s*, 6H, 2 × N-CH₃), 6.70 (*s*, 2H, 2 × H), 6.91 (*t*, 2H, 2 × ArH), 6.99 (*s*, 1H, H), 7.11 (*t*, 2H, 2 × ArH), 7.25–7.30 (*m*, 3H, *J* = 7.6, 6.6, 2.2 Hz, ArH and 2 × ArH), 7.35–7.41 (*m*, 4H, *J* = 8.0, 5.6, 1.2 Hz, ArH and 2 × ArH), 7.77 (*d*, 1H, *J* = 8.0 Hz, ArH) (the OH signal cannot be seen in the ¹H NMR spectrum and hence there are only 21 H atoms in the integration peaks). ¹³C NMR (101 MHz, DMSO-*d*₆) δ (ppm): 32.7, 34.5, 110.1, 117.9, 118.9, 119.5, 121.6, 126.4, 127.4, 128.6, 130.0, 130.1, 131.3, 131.6, 137.4, 145.2, 170.1.

7. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The hydroxy H atom was freely refined. C-bound H atoms were positioned geometrically and refined using a riding model with C–H = 0.93–0.96 and $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{C})$.

Acknowledgements

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Table 3
Experimental details.

Crystal data	
Chemical formula	C ₂₆ H ₂₂ N ₂ O ₂
<i>M_r</i>	394.45
Crystal system, space group	Triclinic, <i>P</i> $\bar{1}$
Temperature (K)	293
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.654 (5), 10.923 (6), 10.964 (5)
α , β , γ (°)	85.85 (2), 82.38 (2), 74.57 (3)
<i>V</i> (Å ³)	989.4 (9)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.55 × 0.39 × 0.30
Data collection	
Diffractometer	Bruker PHOTON 100 CMOS
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2016)
<i>T</i> _{min} , <i>T</i> _{max}	0.548, 0.746
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	37627, 4929, 3077
<i>R</i> _{int}	0.101
(<i>sin</i> θ / λ) _{max} (Å ⁻¹)	0.669
Refinement	
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.069, 0.190, 1.03
No. of reflections	4929
No. of parameters	277
No. of restraints	1
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ (e Å ⁻³)	0.72, -0.35

Computer programs: *APEX3* and *SAINT* (Bruker, 2016), *SHELXT2014/5* (Sheldrick, 2015a), *SHELXL* (Sheldrick, 2015b), *shelXle* (Hübschle *et al.*, 2011), *SHELXTL* (Sheldrick, 2008), *PLATON* (Spek, 2009), *OLEX2* (Dolomanov *et al.*, 2009) and *publCIF* (Westrip, 2010).

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Crystal structure and Hirshfeld analysis of 2-[bis(1-methyl-1*H*-indol-3-yl)methyl]benzoic acid

Suhaila Sapari, Sheryn Wong, Mohammad Fadzlee Ngatiman, Huda Misral and Siti Aishah Hasbullah

Computing details

Data collection: *APEX3* (Bruker, 2016); cell refinement: *SAINT* (Bruker, 2016); data reduction: *SAINT* (Bruker, 2016); program(s) used to solve structure: *SHELXT2014/5* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL* (Sheldrick, 2015b), *shelXle* (Hübschle *et al.*, 2011); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008), *PLATON* (Spek, 2009), *OLEX2* (Dolomanov *et al.*, 2009) and *publCIF* (Westrip, 2010).

2-[Bis(1-methyl-1*H*-indol-3-yl)methyl]benzoic acid

Crystal data

$C_{26}H_{22}N_2O_2$

$M_r = 394.45$

Triclinic, $P\bar{1}$

$a = 8.654$ (5) Å

$b = 10.923$ (6) Å

$c = 10.964$ (5) Å

$\alpha = 85.85$ (2)°

$\beta = 82.38$ (2)°

$\gamma = 74.57$ (3)°

$V = 989.4$ (9) Å³

$Z = 2$

$F(000) = 416$

$D_x = 1.324$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71076$ Å

Cell parameters from 8410 reflections

$\theta = 2.9$ – 27.3 °

$\mu = 0.08$ mm⁻¹

$T = 293$ K

Block, colourless

$0.55 \times 0.39 \times 0.30$ mm

Data collection

Bruker PHOTON 100 CMOS

diffractometer

Radiation source: fine-focus sealed tube

φ and ω scans

Absorption correction: multi-scan

(SADABS; Bruker, 2016)

$T_{\min} = 0.548$, $T_{\max} = 0.746$

37627 measured reflections

4929 independent reflections

3077 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.101$

$\theta_{\max} = 28.4$ °, $\theta_{\min} = 2.9$ °

$h = -11 \rightarrow 11$

$k = -14 \rightarrow 14$

$l = -14 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.069$

$wR(F^2) = 0.190$

$S = 1.03$

4929 reflections

277 parameters

1 restraint

Primary atom site location: structure-invariant

direct methods

Hydrogen site location: mixed

H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0831P)^2 + 0.5551P]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$$\Delta\rho_{\max} = 0.72 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.35 \text{ e } \text{\AA}^{-3}$$

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.

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.1286 (3)	0.9468 (2)	0.8786 (2)	0.0672 (6)
C1	0.0635 (3)	0.8626 (2)	0.8846 (2)	0.0418 (6)
H3O	-0.070 (8)	0.925 (3)	1.005 (6)	0.20 (3)*
C26	0.8219 (3)	0.8016 (3)	0.3643 (3)	0.0575 (7)
H26A	0.840826	0.738582	0.303606	0.086*
H26B	0.901263	0.775487	0.420976	0.086*
H26C	0.829479	0.881343	0.324457	0.086*
C25	0.4893 (3)	0.9294 (2)	0.2705 (2)	0.0465 (6)
H25	0.573439	0.932437	0.209051	0.056*
C24	0.3315 (4)	0.9808 (3)	0.2504 (3)	0.0556 (7)
H24	0.308350	1.019279	0.174135	0.067*
N1	0.6152 (2)	0.6429 (2)	0.97202 (18)	0.0427 (5)
N2	0.6626 (2)	0.81578 (19)	0.43055 (18)	0.0403 (5)
C2	0.1002 (3)	0.7587 (2)	0.7937 (2)	0.0346 (5)
C14	0.7258 (3)	0.3672 (3)	0.7182 (3)	0.0517 (7)
H14	0.748386	0.307509	0.657587	0.062*
C3	-0.0168 (3)	0.6924 (2)	0.7922 (2)	0.0420 (6)
H3	-0.109288	0.711578	0.848612	0.050*
O3	-0.0538 (3)	0.8571 (2)	0.97210 (19)	0.0631 (6)
C4	0.0012 (3)	0.5995 (2)	0.7097 (2)	0.0456 (6)
H4	-0.076746	0.554773	0.711828	0.055*
C5	0.1353 (3)	0.5732 (3)	0.6238 (2)	0.0452 (6)
H5	0.147565	0.512118	0.565909	0.054*
C6	0.2518 (3)	0.6386 (2)	0.6243 (2)	0.0419 (6)
H6	0.341532	0.621037	0.565193	0.050*
C7	0.2402 (3)	0.7292 (2)	0.7094 (2)	0.0341 (5)
C21	0.3939 (3)	0.8673 (2)	0.4803 (2)	0.0339 (5)
C8	0.3842 (3)	0.7849 (2)	0.7119 (2)	0.0332 (5)
H8	0.341561	0.869135	0.746666	0.040*
C22	0.2345 (3)	0.9212 (2)	0.4559 (2)	0.0436 (6)
H22	0.148793	0.919612	0.516422	0.052*

C9	0.4939 (3)	0.7048 (2)	0.7995 (2)	0.0347 (5)
C23	0.2062 (3)	0.9762 (3)	0.3419 (3)	0.0543 (7)
H23	0.100327	1.011219	0.325564	0.065*
C10	0.5135 (3)	0.7397 (2)	0.9123 (2)	0.0414 (6)
H10	0.464125	0.819036	0.944587	0.050*
C11	0.6628 (3)	0.5408 (2)	0.8979 (2)	0.0368 (5)
C13	0.6259 (3)	0.4862 (2)	0.6972 (2)	0.0421 (6)
H13	0.582726	0.506790	0.622801	0.050*
C12	0.5903 (3)	0.5755 (2)	0.7882 (2)	0.0339 (5)
C19	0.6294 (3)	0.7755 (2)	0.5505 (2)	0.0371 (5)
H19	0.707140	0.734204	0.601145	0.044*
C15	0.7928 (3)	0.3347 (3)	0.8272 (3)	0.0542 (7)
H15	0.858248	0.253282	0.839323	0.065*
C17	0.6653 (4)	0.6456 (3)	1.0926 (2)	0.0616 (8)
H17A	0.769095	0.663417	1.083578	0.092*
H17B	0.672004	0.564573	1.134905	0.092*
H17C	0.587916	0.710521	1.139210	0.092*
C16	0.7642 (3)	0.4213 (3)	0.9186 (2)	0.0493 (6)
H16	0.811197	0.400285	0.991351	0.059*
C20	0.5193 (3)	0.8728 (2)	0.3856 (2)	0.0363 (5)
C18	0.4677 (3)	0.8041 (2)	0.5851 (2)	0.0322 (5)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0693 (14)	0.0618 (13)	0.0698 (14)	-0.0252 (11)	0.0249 (11)	-0.0267 (11)
C1	0.0343 (12)	0.0498 (15)	0.0381 (13)	-0.0080 (11)	-0.0023 (10)	0.0048 (11)
C26	0.0394 (14)	0.0683 (19)	0.0607 (18)	-0.0173 (13)	0.0129 (12)	0.0022 (14)
C25	0.0609 (16)	0.0423 (13)	0.0386 (13)	-0.0200 (12)	-0.0030 (12)	0.0028 (11)
C24	0.0698 (19)	0.0517 (16)	0.0477 (16)	-0.0171 (14)	-0.0213 (14)	0.0135 (12)
N1	0.0414 (11)	0.0582 (13)	0.0315 (10)	-0.0179 (10)	-0.0058 (8)	-0.0007 (9)
N2	0.0317 (10)	0.0449 (11)	0.0427 (11)	-0.0119 (8)	0.0025 (8)	0.0025 (9)
C2	0.0281 (10)	0.0387 (12)	0.0344 (11)	-0.0053 (9)	-0.0052 (9)	0.0066 (9)
C14	0.0551 (16)	0.0435 (14)	0.0540 (16)	-0.0087 (12)	-0.0038 (13)	-0.0073 (12)
C3	0.0290 (11)	0.0503 (14)	0.0440 (13)	-0.0097 (10)	0.0006 (10)	0.0045 (11)
O3	0.0614 (13)	0.0700 (14)	0.0543 (12)	-0.0209 (11)	0.0201 (10)	-0.0137 (11)
C4	0.0379 (13)	0.0517 (15)	0.0535 (15)	-0.0237 (11)	-0.0074 (11)	0.0056 (12)
C5	0.0428 (13)	0.0518 (15)	0.0448 (14)	-0.0182 (11)	-0.0032 (11)	-0.0067 (11)
C6	0.0327 (12)	0.0542 (15)	0.0394 (13)	-0.0144 (11)	0.0013 (10)	-0.0040 (11)
C7	0.0287 (10)	0.0411 (12)	0.0329 (11)	-0.0104 (9)	-0.0055 (9)	0.0050 (9)
C21	0.0348 (11)	0.0290 (11)	0.0380 (12)	-0.0089 (9)	-0.0040 (9)	-0.0002 (9)
C8	0.0275 (10)	0.0372 (12)	0.0351 (12)	-0.0096 (9)	-0.0004 (9)	-0.0034 (9)
C22	0.0373 (12)	0.0411 (13)	0.0495 (15)	-0.0047 (10)	-0.0080 (11)	0.0016 (11)
C9	0.0292 (11)	0.0442 (13)	0.0326 (11)	-0.0147 (9)	0.0001 (9)	-0.0016 (9)
C23	0.0486 (15)	0.0500 (16)	0.0623 (18)	-0.0049 (12)	-0.0209 (13)	0.0071 (13)
C10	0.0375 (12)	0.0496 (14)	0.0383 (13)	-0.0154 (11)	0.0003 (10)	-0.0038 (11)
C11	0.0326 (11)	0.0463 (13)	0.0341 (12)	-0.0174 (10)	-0.0008 (9)	0.0023 (10)
C13	0.0420 (13)	0.0439 (14)	0.0416 (13)	-0.0127 (11)	-0.0060 (10)	-0.0024 (11)

C12	0.0290 (10)	0.0422 (12)	0.0323 (11)	-0.0144 (9)	-0.0001 (9)	-0.0003 (9)
C19	0.0323 (11)	0.0415 (13)	0.0369 (12)	-0.0106 (10)	-0.0022 (9)	0.0016 (10)
C15	0.0528 (16)	0.0438 (15)	0.0589 (17)	-0.0042 (12)	-0.0037 (13)	0.0072 (13)
C17	0.0698 (19)	0.087 (2)	0.0350 (14)	-0.0299 (17)	-0.0130 (13)	-0.0032 (14)
C16	0.0462 (14)	0.0571 (16)	0.0433 (14)	-0.0139 (12)	-0.0085 (11)	0.0141 (12)
C20	0.0381 (12)	0.0321 (11)	0.0399 (12)	-0.0125 (9)	-0.0021 (10)	-0.0017 (9)
C18	0.0293 (10)	0.0304 (11)	0.0363 (12)	-0.0082 (9)	-0.0009 (9)	-0.0018 (9)

Geometric parameters (Å, °)

O1—C1	1.193 (3)	C5—H5	0.9300
C1—O3	1.311 (3)	C6—C7	1.384 (3)
C1—C2	1.507 (4)	C6—H6	0.9300
C26—N2	1.445 (3)	C7—C8	1.529 (3)
C26—H26A	0.9600	C21—C22	1.402 (3)
C26—H26B	0.9600	C21—C20	1.410 (3)
C26—H26C	0.9600	C21—C18	1.431 (3)
C25—C24	1.371 (4)	C8—C9	1.507 (3)
C25—C20	1.389 (3)	C8—C18	1.507 (3)
C25—H25	0.9300	C8—H8	0.9800
C24—C23	1.385 (4)	C22—C23	1.370 (4)
C24—H24	0.9300	C22—H22	0.9300
N1—C11	1.368 (3)	C9—C10	1.365 (3)
N1—C10	1.371 (3)	C9—C12	1.441 (3)
N1—C17	1.449 (3)	C23—H23	0.9300
N2—C20	1.370 (3)	C10—H10	0.9300
N2—C19	1.375 (3)	C11—C16	1.388 (4)
C2—C3	1.395 (3)	C11—C12	1.410 (3)
C2—C7	1.402 (3)	C13—C12	1.394 (3)
C14—C13	1.379 (4)	C13—H13	0.9300
C14—C15	1.380 (4)	C19—C18	1.358 (3)
C14—H14	0.9300	C19—H19	0.9300
C3—C4	1.373 (4)	C15—C16	1.380 (4)
C3—H3	0.9300	C15—H15	0.9300
O3—H3O	0.820 (10)	C17—H17A	0.9600
C4—C5	1.374 (3)	C17—H17B	0.9600
C4—H4	0.9300	C17—H17C	0.9600
C5—C6	1.383 (3)	C16—H16	0.9300
O1—C1—O3	120.9 (2)	C9—C8—C7	108.83 (18)
O1—C1—C2	124.9 (2)	C18—C8—C7	112.70 (18)
O3—C1—C2	114.1 (2)	C9—C8—H8	106.9
N2—C26—H26A	109.5	C18—C8—H8	106.9
N2—C26—H26B	109.5	C7—C8—H8	106.9
H26A—C26—H26B	109.5	C23—C22—C21	119.3 (2)
N2—C26—H26C	109.5	C23—C22—H22	120.3
H26A—C26—H26C	109.5	C21—C22—H22	120.3
H26B—C26—H26C	109.5	C10—C9—C12	105.5 (2)

C24—C25—C20	117.9 (2)	C10—C9—C8	125.6 (2)
C24—C25—H25	121.1	C12—C9—C8	128.7 (2)
C20—C25—H25	121.1	C22—C23—C24	121.6 (3)
C25—C24—C23	121.0 (3)	C22—C23—H23	119.2
C25—C24—H24	119.5	C24—C23—H23	119.2
C23—C24—H24	119.5	C9—C10—N1	111.3 (2)
C11—N1—C10	108.1 (2)	C9—C10—H10	124.4
C11—N1—C17	125.0 (2)	N1—C10—H10	124.4
C10—N1—C17	126.9 (2)	N1—C11—C16	129.4 (2)
C20—N2—C19	108.30 (19)	N1—C11—C12	108.1 (2)
C20—N2—C26	126.1 (2)	C16—C11—C12	122.5 (2)
C19—N2—C26	125.6 (2)	C14—C13—C12	119.5 (2)
C3—C2—C7	119.2 (2)	C14—C13—H13	120.2
C3—C2—C1	116.8 (2)	C12—C13—H13	120.2
C7—C2—C1	123.9 (2)	C13—C12—C11	118.0 (2)
C13—C14—C15	121.4 (3)	C13—C12—C9	135.0 (2)
C13—C14—H14	119.3	C11—C12—C9	107.0 (2)
C15—C14—H14	119.3	C18—C19—N2	110.8 (2)
C4—C3—C2	121.6 (2)	C18—C19—H19	124.6
C4—C3—H3	119.2	N2—C19—H19	124.6
C2—C3—H3	119.2	C14—C15—C16	121.0 (2)
C1—O3—H3O	102 (5)	C14—C15—H15	119.5
C3—C4—C5	119.5 (2)	C16—C15—H15	119.5
C3—C4—H4	120.2	N1—C17—H17A	109.5
C5—C4—H4	120.2	N1—C17—H17B	109.5
C4—C5—C6	119.2 (2)	H17A—C17—H17B	109.5
C4—C5—H5	120.4	N1—C17—H17C	109.5
C6—C5—H5	120.4	H17A—C17—H17C	109.5
C5—C6—C7	122.6 (2)	H17B—C17—H17C	109.5
C5—C6—H6	118.7	C15—C16—C11	117.6 (2)
C7—C6—H6	118.7	C15—C16—H16	121.2
C6—C7—C2	117.7 (2)	C11—C16—H16	121.2
C6—C7—C8	118.5 (2)	N2—C20—C25	130.2 (2)
C2—C7—C8	123.7 (2)	N2—C20—C21	107.6 (2)
C22—C21—C20	118.0 (2)	C25—C20—C21	122.2 (2)
C22—C21—C18	134.8 (2)	C19—C18—C21	106.1 (2)
C20—C21—C18	107.2 (2)	C19—C18—C8	126.7 (2)
C9—C8—C18	114.13 (18)	C21—C18—C8	126.96 (19)
C20—C25—C24—C23	-0.1 (4)	C14—C13—C12—C11	1.9 (3)
O1—C1—C2—C3	-161.0 (3)	C14—C13—C12—C9	-179.8 (2)
O3—C1—C2—C3	16.6 (3)	N1—C11—C12—C13	178.30 (19)
O1—C1—C2—C7	16.6 (4)	C16—C11—C12—C13	-1.4 (3)
O3—C1—C2—C7	-165.8 (2)	N1—C11—C12—C9	-0.5 (2)
C7—C2—C3—C4	-0.5 (3)	C16—C11—C12—C9	179.8 (2)
C1—C2—C3—C4	177.2 (2)	C10—C9—C12—C13	-178.4 (2)
C2—C3—C4—C5	-1.9 (4)	C8—C9—C12—C13	5.4 (4)
C3—C4—C5—C6	1.7 (4)	C10—C9—C12—C11	0.1 (2)

C4—C5—C6—C7	0.8 (4)	C8—C9—C12—C11	-176.1 (2)
C5—C6—C7—C2	-3.1 (4)	C20—N2—C19—C18	0.3 (3)
C5—C6—C7—C8	172.9 (2)	C26—N2—C19—C18	179.6 (2)
C3—C2—C7—C6	2.9 (3)	C13—C14—C15—C16	-1.0 (4)
C1—C2—C7—C6	-174.7 (2)	C14—C15—C16—C11	1.5 (4)
C3—C2—C7—C8	-172.9 (2)	N1—C11—C16—C15	-179.9 (2)
C1—C2—C7—C8	9.5 (3)	C12—C11—C16—C15	-0.2 (4)
C6—C7—C8—C9	-90.0 (2)	C19—N2—C20—C25	179.3 (2)
C2—C7—C8—C9	85.8 (3)	C26—N2—C20—C25	0.1 (4)
C6—C7—C8—C18	37.6 (3)	C19—N2—C20—C21	-0.2 (3)
C2—C7—C8—C18	-146.6 (2)	C26—N2—C20—C21	-179.4 (2)
C20—C21—C22—C23	0.4 (3)	C24—C25—C20—N2	-179.5 (2)
C18—C21—C22—C23	179.7 (3)	C24—C25—C20—C21	-0.1 (4)
C18—C8—C9—C10	125.1 (2)	C22—C21—C20—N2	179.5 (2)
C7—C8—C9—C10	-108.1 (2)	C18—C21—C20—N2	0.0 (2)
C18—C8—C9—C12	-59.5 (3)	C22—C21—C20—C25	0.0 (3)
C7—C8—C9—C12	67.3 (3)	C18—C21—C20—C25	-179.5 (2)
C21—C22—C23—C24	-0.6 (4)	N2—C19—C18—C21	-0.3 (3)
C25—C24—C23—C22	0.4 (4)	N2—C19—C18—C8	-174.8 (2)
C12—C9—C10—N1	0.4 (3)	C22—C21—C18—C19	-179.2 (3)
C8—C9—C10—N1	176.70 (19)	C20—C21—C18—C19	0.2 (2)
C11—N1—C10—C9	-0.7 (3)	C22—C21—C18—C8	-4.7 (4)
C17—N1—C10—C9	179.4 (2)	C20—C21—C18—C8	174.7 (2)
C10—N1—C11—C16	-179.6 (2)	C9—C8—C18—C19	-11.6 (3)
C17—N1—C11—C16	0.3 (4)	C7—C8—C18—C19	-136.4 (2)
C10—N1—C11—C12	0.7 (2)	C9—C8—C18—C21	175.0 (2)
C17—N1—C11—C12	-179.4 (2)	C7—C8—C18—C21	50.2 (3)
C15—C14—C13—C12	-0.7 (4)		

Hydrogen-bond geometry (Å, °)

<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>
O3—H3O...O1 ⁱ	0.82 (4)	1.89 (5)	2.679 (3)	163 (6)
C8—H8...O1	0.98	2.20	2.945 (4)	132

Symmetry code: (i) $-x, -y+2, -z+2$.*Percentage contributions of interatomic contacts to the Hirshfeld surface for (I)*

Contact	Percentage contribution
H ... H	54.6
O ... H / H ... O	10.1
C ... H / H ... C	29.6
N ... H / H ... N	1.1
C ... N / N ... C	1.7
C ... C	2.8