

(E)-1-Methyl-4-[2-(2-naphthyl)vinyl]-pyridinium iodide¹

Hoong-Kun Fun,^{a*}§ Kullapa Chanawanno^b and Suchada Chantrapromma^{b¶}

^aX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bCrystal Materials Research Unit, Department of Chemistry, Faculty of Science, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand

Correspondence e-mail: hkfun@usm.my

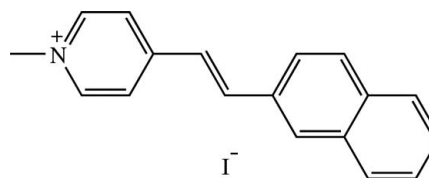
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(I) = 0.000$ Å; disorder in main residue; R factor = 0.056; wR factor = 0.148; data-to-parameter ratio = 20.4.

In the title compound, $C_{18}H_{16}N^+ \cdot I^-$, the cation is disordered over two orientations related by a 180° rotation about its long axis with occupancies of 0.554 (7) and 0.446 (7). Both disorder components exist in an *E* configuration. The dihedral angle between the pyridinium ring and the naphthalene ring system is 4.7 (6°) in the major disorder component and 1.6 (8°) in the minor component. In the crystal structure, centrosymmetrically related cations are stacked along the *a* axis, with significant π - π interactions between the pyridinium ring and the naphthalene ring system [centroid-centroid distance = 3.442 (9) Å]. The iodide ions are located between adjacent columns of cations. The cations are linked to the iodide ions by $C-H \cdots I$ interactions. Weak $C-H \cdots \pi$ interactions involving the methyl group are also observed.

Related literature

For bond-length data, see: Allen *et al.* (1987). For background to non-linear optical materials research, see: Cheng *et al.* (1991a,b); Ogawa *et al.* (2008); Yang *et al.* (2007). For related structures, see: Chanawanno *et al.* (2008); Chantrapromma *et al.* (2006; 2007; 2008; 2009a,b). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

$C_{18}H_{16}N^+ \cdot I^-$	$V = 1568.22$ (5) Å ³
$M_r = 373.29$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 7.2789$ (1) Å	$\mu = 2.03$ mm ⁻¹
$b = 10.9363$ (2) Å	$T = 100$ K
$c = 20.0883$ (4) Å	$0.53 \times 0.30 \times 0.09$ mm
$\beta = 101.280$ (1) $^\circ$	

Data collection

Bruker APEXII CCD area-detector diffractometer	34203 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2005)	6896 independent reflections
$T_{min} = 0.412$, $T_{max} = 0.838$	5373 reflections with $I > 2\sigma(I)$
	$R_{int} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.056$	91 restraints
$wR(F^2) = 0.148$	H-atom parameters constrained
$S = 1.06$	$\Delta\rho_{max} = 3.51$ e Å ⁻³
6896 reflections	$\Delta\rho_{min} = -2.42$ e Å ⁻³
338 parameters	

Table 1

Hydrogen-bond geometry (Å, $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$C18A-H18A \cdots I1^i$	0.96	3.05	3.928 (17)	152
$C18A-H18B \cdots Cg1^i$	0.96	2.63	3.513 (18)	153
$C18A-H18B \cdots Cg2^i$	0.96	2.65	3.517 (18)	150
$C18B-H18E \cdots Cg1^i$	0.96	2.62	3.44 (2)	143
$C18B-H18E \cdots Cg2^i$	0.96	2.66	3.45 (2)	139

Symmetry code: (i) $-x + 1, -y + 1, -z + 2$. $Cg1$ and $Cg2$ are centroids of the $C10A-C15A$ and $C10B-C15B$ rings, respectively.

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

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§ Thomson Reuters ResearcherID: A-3561-2009.

¶ Additional correspondence author, e-mail: suchada.c@psu.ac.th. Thomson Reuters ResearcherID: A-5085-2009.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2807).

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supporting information

Acta Cryst. (2009). E65, o1406–o1407 [doi:10.1107/S1600536809019114]

(E)-1-Methyl-4-[2-(2-naphthyl)vinyl]pyridinium iodide**Hoong-Kun Fun, Kullapa Chanawanno and Suchada Chantrapromma****S1. Comment**

In order to obtain second-order non-linear optical (NLO) single crystals, the main requirements should be the choice of molecules with large hyperpolarizability (β) and these molecules should align into a noncentrosymmetric space group in the crystal. Organic crystals with extensive conjugated π systems with large hyperpolarizability which exhibit NLO properties have been reported (Ogawa *et al.*, 2008; Yang *et al.*, 2007). Styryl pyridinium derivatives are considered to be good conjugated π -systems (Cheng *et al.*, 1991a, 1991b). In our on-going research in searching for NLO materials (Chanawanno *et al.*, 2008; Chantrapromma *et al.*, 2006, 2007, 2008, 2009a, b), we have previously reported the crystal structure of (*E*)-1-methyl-4-[2-(1-naphthyl)vinyl]pyridinium 4-bromobenzenesulfonate (Chantrapromma *et al.*, 2009a). In order to study the effect of different positions of the substituent group and anions, the title compound was synthesized by replacing the 1-naphthyl and 4-bromobenzenesulfonate in the (*E*)-1-methyl-4-[2-(1-naphthyl)vinyl]pyridinium 4-bromobenzenesulfonate with 2-naphthyl and iodide in the title compound. However it crystallized in the monoclinic centrosymmetric space group $P2_1/c$ and would not exhibit second-order nonlinear optical properties.

Fig. 1 shows the asymmetric unit of the title compound which consists of a $C_{18}H_{16}N^+$ cation and a I^- anion. The whole cation is disordered over two sites; the major component *A* and the minor component *B* (Fig. 1), with the refined site-occupancy ratio of 0.554 (7)/0.446 (7). The cation exists in the *E* configuration with respect to the $C6=C7$ double bond. The naphthalenyl moiety is essentially planar in both disorder components as indicated by the interplanar angle between the two aromatic $C8-C10/C15-C17$ and $C10-C15$ rings [1.5 (8) $^\circ$ for the major component *A* and 3.2 (9) $^\circ$ for the minor component *B*]. The major component *A* of cation is slightly twisted with the dihedral angle between the pyridinium and the mean plane through the naphthalenyl moiety ($C8-C17$) being 4.7 (6) $^\circ$ whereas the minor component *B* is almost planar [dihedral angle 1.6 (8) $^\circ$]. The $C4-C5-C6-C7$ and $C6-C7-C8-C17$ torsion angles [0.4 (10) $^\circ$ and 2.1 (10) $^\circ$ in the major component and -179.4 (8) $^\circ$ and 179.9 (8) $^\circ$ in the minor component] in both disorder components indicate that the orientations of the ethynyl moiety in these components are related by 180° rotation about the long axis of the molecule. The bond lengths are in normal ranges (Allen *et al.*, 1987) and are comparable to those observed in related structures (Chanawanno *et al.*, 2008; Chantrapromma *et al.*, 2006, 2007, 2008, 2009a,b).

In the crystal packing (Fig. 2), centrosymmetrically related cations are stacked along the *a* axis, with significant π - π interactions between pyridinium ring and naphthalene ring system [centroid-centroid distance is 3.442 (9) \AA]. The iodide ions are located between adjacent columns of cations. The cations are linked to the iodide ions by $C-H\cdots I$ weak interactions (Table 1). The crystal structure is further stabilized by $C-H\cdots\pi$ interactions involving the methyl group (Table 1; Cg1 and Cg2 are centroids of the $C10A-C15A$ and $C10B-C15B$ rings, respectively).

S2. Experimental

The title compound was prepared by mixing 1:1:1 molar ratio solutions of 1,4-dimethylpyridinium iodide (2 g, 8.5 mmol), 2-naphthaldehyde (1.16 ml, 8.5 mmol) and piperidine (0.84 ml, 8.5 mmol) in methanol (40 ml). The resulting

solution was refluxed for 3 h under a nitrogen atmosphere. The solid compound formed was filtered and washed with diethylether. Yellow plate-shaped single crystals of the title compound suitable for *X*-ray structure determination were recrystallized from methanol by slow evaporation at room temperature over a few weeks (m.p. 557-558 K).

S3. Refinement

The cation is disordered over two orientations with occupancies of 0.554 (7) and 0.446 (7). The same U_{ij} parameters were used for atom pairs N1A/N1B and C15A/C16A and all disordered atoms were subjected to a rigid bond restraint. All H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H = 0.93 Å (aromatic and CH) and 0.96 Å (CH₃). The U_{iso} values were constrained to be $1.5U_{eq}$ of the carrier atom for methyl H atoms and $1.2U_{eq}$ for the remaining H atoms. A rotating group model was used for the methyl groups. The highest residual electron density peak is located at 0.69 Å from I1 and the deepest hole is located at 0.56 Å from I1.

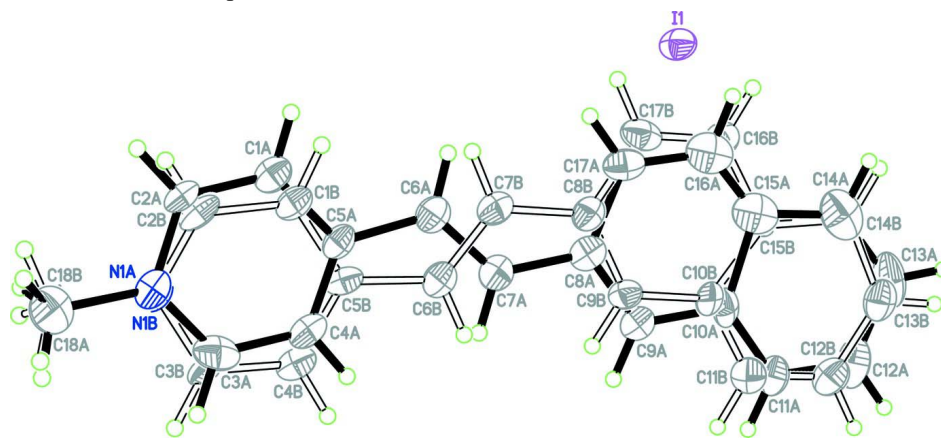


Figure 1

The molecular structure of the title compound, with 50% probability displacement ellipsoids and the atom-numbering scheme. Open bonds show the minor disorder component.

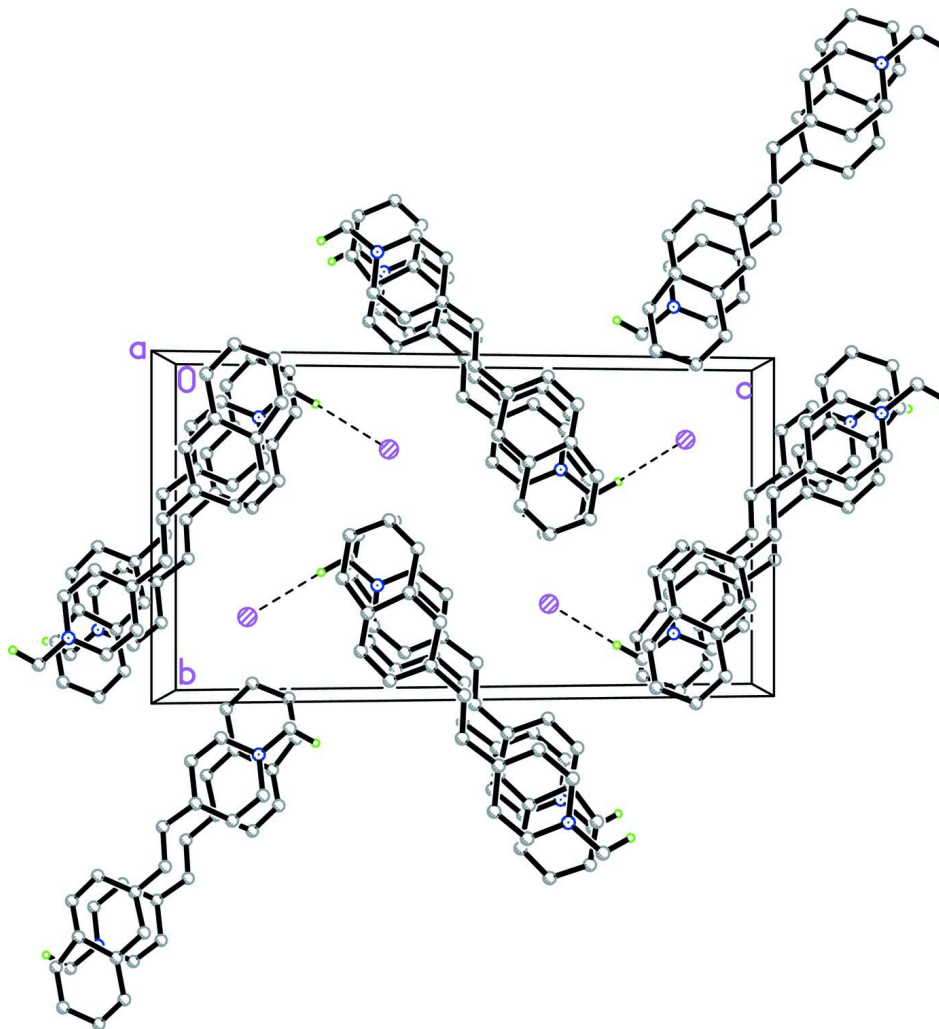


Figure 2

The crystal packing of the major component of the title compound viewed down the *a* axis. H atoms not involved in C—H...I interactions (dashed lines) have been omitted for clarity.

(*E*)-1-Methyl-4-[2-(2-naphthyl)vinyl]pyridinium iodide

Crystal data

$C_{18}H_{16}N^+I^-$

$M_r = 373.29$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 7.2789$ (1) Å

$b = 10.9363$ (2) Å

$c = 20.0883$ (4) Å

$\beta = 101.280$ (1)°

$V = 1568.22$ (5) Å³

$Z = 4$

$F(000) = 736$

$D_x = 1.581$ Mg m⁻³

Melting point = 557–558 K

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

Cell parameters from 6896 reflections

$\theta = 2.1$ – 35.0 °

$\mu = 2.03$ mm⁻¹

$T = 100$ K

Plate, yellow

$0.53 \times 0.30 \times 0.09$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer
Radiation source: sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2005)
 $T_{\min} = 0.412$, $T_{\max} = 0.838$

34203 measured reflections
6896 independent reflections
5373 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 35.0^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -11 \rightarrow 11$
 $k = -17 \rightarrow 16$
 $l = -32 \rightarrow 27$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.148$
 $S = 1.06$
6896 reflections
338 parameters
91 restraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0586P)^2 + 4.6136P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 3.51 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -2.42 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

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

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
I1	0.48129 (4)	0.73286 (2)	0.633723 (12)	0.03874 (9)	
N1A	0.520 (4)	0.174 (3)	1.153 (2)	0.022 (2)	0.555 (7)
C1A	0.3651 (10)	0.1993 (7)	1.0385 (4)	0.0268 (12)	0.555 (7)
H1AA	0.3113	0.1634	0.9973	0.032*	0.555 (7)
C2A	0.452 (2)	0.1269 (8)	1.0898 (7)	0.0230 (16)	0.555 (7)
H2AA	0.4665	0.0440	1.0820	0.028*	0.555 (7)
C3A	0.516 (2)	0.2966 (15)	1.1600 (7)	0.033 (2)	0.555 (7)
H3AA	0.5689	0.3307	1.2018	0.039*	0.555 (7)
C4A	0.4359 (12)	0.3744 (8)	1.1076 (5)	0.0290 (15)	0.555 (7)
H4AA	0.4382	0.4586	1.1145	0.035*	0.555 (7)
C5A	0.3533 (8)	0.3270 (6)	1.0454 (4)	0.0238 (11)	0.555 (7)
C6A	0.2561 (8)	0.4011 (5)	0.9884 (3)	0.0283 (12)	0.555 (7)
H6AA	0.2049	0.3616	0.9480	0.034*	0.555 (7)
C7A	0.2374 (8)	0.5225 (5)	0.9915 (3)	0.0282 (12)	0.555 (7)

H7AA	0.2908	0.5591	1.0325	0.034*	0.555 (7)
C8A	0.1433 (9)	0.6045 (7)	0.9383 (4)	0.0268 (11)	0.555 (7)
C9A	0.1464 (11)	0.7284 (8)	0.9532 (4)	0.0263 (12)	0.555 (7)
H9AA	0.2117	0.7573	0.9947	0.032*	0.555 (7)
C10A	0.047 (3)	0.8108 (13)	0.9036 (9)	0.028 (2)	0.555 (7)
C11A	0.0570 (16)	0.9380 (9)	0.9171 (5)	0.0303 (17)	0.555 (7)
H11A	0.1258	0.9661	0.9582	0.036*	0.555 (7)
C12A	-0.0330 (19)	1.0196 (11)	0.8709 (6)	0.041 (2)	0.555 (7)
H12A	-0.0320	1.1026	0.8811	0.049*	0.555 (7)
C13A	-0.128 (4)	0.976 (2)	0.8070 (8)	0.047 (4)	0.555 (7)
H13A	-0.1882	1.0300	0.7738	0.056*	0.555 (7)
C14A	-0.129 (4)	0.852 (2)	0.7955 (13)	0.047 (3)	0.555 (7)
H14A	-0.1968	0.8284	0.7534	0.057*	0.555 (7)
C15A	-0.0433 (17)	0.7529 (12)	0.8368 (7)	0.0400 (17)	0.555 (7)
C16A	-0.040 (2)	0.6423 (19)	0.8260 (7)	0.0400 (17)	0.555 (7)
H16A	-0.1012	0.6131	0.7840	0.048*	0.555 (7)
C17A	0.0489 (12)	0.5599 (8)	0.8729 (4)	0.0312 (14)	0.555 (7)
H17A	0.0484	0.4769	0.8628	0.037*	0.555 (7)
C18A	0.618 (2)	0.0940 (16)	1.2057 (9)	0.040 (4)	0.555 (7)
H18A	0.6436	0.1367	1.2483	0.061*	0.555 (7)
H18B	0.7343	0.0685	1.1941	0.061*	0.555 (7)
H18C	0.5423	0.0236	1.2095	0.061*	0.555 (7)
N1B	0.544 (5)	0.186 (3)	1.151 (3)	0.022 (2)	0.45
C1B	0.3643 (13)	0.2454 (8)	1.0446 (5)	0.0262 (16)	0.445 (7)
H1BA	0.3072	0.2232	1.0007	0.031*	0.445 (7)
C2B	0.443 (3)	0.1573 (12)	1.0917 (10)	0.029 (2)	0.445 (7)
H2BA	0.4241	0.0752	1.0805	0.034*	0.445 (7)
C3B	0.546 (3)	0.3028 (12)	1.1733 (8)	0.021 (2)	0.445 (7)
H3BA	0.6026	0.3218	1.2177	0.025*	0.445 (7)
C4B	0.4663 (15)	0.3915 (9)	1.1306 (5)	0.0257 (15)	0.445 (7)
H4BA	0.4740	0.4721	1.1457	0.031*	0.445 (7)
C5B	0.3727 (11)	0.3669 (8)	1.0646 (4)	0.0246 (14)	0.445 (7)
C6B	0.2907 (10)	0.4672 (7)	1.0211 (4)	0.0276 (14)	0.445 (7)
H6BA	0.3067	0.5450	1.0401	0.033*	0.445 (7)
C7B	0.1948 (10)	0.4609 (7)	0.9569 (4)	0.0277 (14)	0.445 (7)
H7BA	0.1786	0.3843	0.9366	0.033*	0.445 (7)
C8B	0.1140 (11)	0.5666 (8)	0.9168 (5)	0.0258 (14)	0.445 (7)
C9B	0.1359 (14)	0.6857 (9)	0.9432 (5)	0.0248 (16)	0.445 (7)
H9BA	0.1989	0.6971	0.9877	0.030*	0.445 (7)
C10B	0.066 (3)	0.7876 (14)	0.9045 (12)	0.024 (2)	0.445 (7)
C11B	0.073 (2)	0.9053 (11)	0.9303 (6)	0.031 (2)	0.445 (7)
H11B	0.1359	0.9193	0.9747	0.037*	0.445 (7)
C12B	-0.009 (2)	1.0020 (13)	0.8918 (6)	0.035 (2)	0.445 (7)
H12B	0.0004	1.0804	0.9101	0.042*	0.445 (7)
C13B	-0.110 (4)	0.983 (2)	0.8243 (10)	0.033 (3)	0.445 (7)
H13B	-0.1654	1.0503	0.8000	0.040*	0.445 (7)
C14B	-0.128 (5)	0.8686 (15)	0.7928 (12)	0.024 (3)	0.445 (7)
H14B	-0.1891	0.8547	0.7483	0.029*	0.445 (7)

C15B	−0.0405 (15)	0.7753 (7)	0.8384 (5)	0.0125 (13)*	0.445 (7)
C16B	−0.056 (2)	0.6447 (17)	0.8116 (6)	0.031 (2)	0.445 (7)
H16B	−0.1177	0.6311	0.7671	0.037*	0.445 (7)
C17B	0.0140 (14)	0.5504 (11)	0.8485 (5)	0.0323 (19)	0.445 (7)
H17B	−0.0016	0.4722	0.8301	0.039*	0.445 (7)
C18B	0.624 (3)	0.0915 (12)	1.2029 (8)	0.026 (3)	0.445 (7)
H18G	0.5790	0.0120	1.1874	0.039*	0.445 (7)
H18D	0.5873	0.1095	1.2451	0.039*	0.445 (7)
H18E	0.7586	0.0928	1.2093	0.039*	0.445 (7)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
II	0.05884 (17)	0.02652 (11)	0.03108 (12)	−0.00509 (9)	0.00937 (10)	0.00240 (8)
N1A	0.014 (7)	0.019 (5)	0.037 (3)	−0.005 (4)	0.014 (4)	0.003 (3)
C1A	0.029 (3)	0.021 (3)	0.033 (3)	−0.003 (3)	0.013 (2)	0.000 (3)
C2A	0.024 (3)	0.015 (4)	0.032 (3)	0.000 (3)	0.011 (2)	−0.005 (3)
C3A	0.020 (5)	0.044 (4)	0.034 (5)	−0.010 (3)	0.004 (3)	−0.011 (3)
C4A	0.030 (4)	0.022 (3)	0.037 (4)	0.002 (2)	0.012 (3)	−0.005 (3)
C5A	0.027 (3)	0.017 (3)	0.030 (3)	0.000 (2)	0.012 (2)	0.002 (2)
C6A	0.029 (3)	0.025 (2)	0.033 (3)	0.0001 (19)	0.009 (2)	0.001 (2)
C7A	0.026 (2)	0.026 (2)	0.033 (3)	−0.0007 (19)	0.007 (2)	0.000 (2)
C8A	0.028 (3)	0.025 (3)	0.030 (3)	−0.003 (2)	0.012 (2)	0.000 (2)
C9A	0.026 (3)	0.025 (3)	0.030 (3)	−0.001 (3)	0.011 (2)	0.000 (3)
C10A	0.020 (5)	0.039 (5)	0.026 (3)	0.013 (4)	0.010 (3)	0.008 (4)
C11A	0.029 (3)	0.027 (4)	0.037 (4)	0.002 (3)	0.011 (3)	0.008 (3)
C12A	0.032 (5)	0.035 (4)	0.058 (7)	0.005 (3)	0.017 (5)	0.018 (4)
C13A	0.027 (5)	0.066 (7)	0.049 (8)	0.011 (6)	0.012 (6)	0.025 (6)
C14A	0.027 (7)	0.073 (8)	0.045 (7)	−0.001 (8)	0.013 (5)	0.008 (6)
C15A	0.027 (3)	0.055 (4)	0.042 (3)	−0.009 (3)	0.015 (2)	−0.007 (3)
C16A	0.027 (3)	0.055 (4)	0.042 (3)	−0.009 (3)	0.015 (2)	−0.007 (3)
C17A	0.032 (4)	0.030 (3)	0.033 (4)	−0.008 (2)	0.011 (3)	−0.008 (3)
C18A	0.019 (5)	0.049 (7)	0.050 (7)	0.000 (4)	−0.001 (5)	−0.001 (5)
N1B	0.014 (7)	0.019 (5)	0.037 (3)	−0.005 (4)	0.014 (4)	0.003 (3)
C1B	0.030 (3)	0.017 (4)	0.033 (4)	−0.003 (3)	0.011 (3)	−0.001 (3)
C2B	0.030 (4)	0.018 (5)	0.042 (4)	0.003 (5)	0.019 (3)	−0.007 (4)
C3B	0.016 (5)	0.012 (3)	0.035 (6)	−0.005 (3)	0.004 (4)	−0.007 (3)
C4B	0.031 (4)	0.019 (3)	0.031 (4)	−0.006 (3)	0.017 (3)	−0.003 (3)
C5B	0.030 (3)	0.021 (3)	0.026 (3)	−0.003 (3)	0.012 (3)	−0.007 (2)
C6B	0.029 (3)	0.022 (3)	0.032 (3)	0.001 (2)	0.009 (2)	0.001 (2)
C7B	0.028 (3)	0.025 (3)	0.032 (3)	0.001 (2)	0.009 (2)	−0.004 (2)
C8B	0.024 (3)	0.022 (3)	0.031 (4)	0.000 (3)	0.007 (3)	0.000 (3)
C9B	0.029 (3)	0.024 (4)	0.022 (3)	0.000 (3)	0.008 (2)	−0.003 (3)
C10B	0.009 (4)	0.031 (5)	0.032 (4)	0.007 (4)	0.007 (3)	−0.004 (4)
C11B	0.031 (4)	0.032 (5)	0.030 (5)	0.000 (4)	0.008 (4)	0.002 (4)
C12B	0.036 (5)	0.032 (5)	0.039 (6)	0.003 (4)	0.014 (5)	0.000 (4)
C13B	0.026 (7)	0.037 (5)	0.041 (7)	0.006 (4)	0.015 (6)	0.005 (5)
C14B	0.018 (6)	0.028 (4)	0.026 (4)	0.007 (3)	0.003 (4)	0.015 (3)

C16B	0.030 (5)	0.029 (4)	0.035 (5)	0.002 (3)	0.009 (4)	-0.003 (4)
C17B	0.026 (4)	0.045 (5)	0.026 (4)	-0.003 (3)	0.006 (3)	-0.011 (4)
C18B	0.039 (8)	0.014 (4)	0.032 (5)	0.012 (4)	0.022 (5)	0.012 (4)

Geometric parameters (Å, °)

N1A—C3A	1.34 (3)	N1B—C2B	1.31 (6)
N1A—C2A	1.38 (4)	N1B—C3B	1.36 (4)
N1A—C18A	1.45 (4)	N1B—C18B	1.50 (5)
C1A—C2A	1.354 (13)	C1B—C5B	1.385 (12)
C1A—C5A	1.408 (10)	C1B—C2B	1.393 (18)
C1A—H1AA	0.93	C1B—H1BA	0.93
C2A—H2AA	0.93	C2B—H2BA	0.93
C3A—C4A	1.390 (17)	C3B—C4B	1.350 (17)
C3A—H3AA	0.93	C3B—H3BA	0.93
C4A—C5A	1.378 (10)	C4B—C5B	1.393 (12)
C4A—H4AA	0.93	C4B—H4BA	0.93
C5A—C6A	1.467 (8)	C5B—C6B	1.456 (11)
C6A—C7A	1.337 (8)	C6B—C7B	1.342 (10)
C6A—H6AA	0.93	C6B—H6BA	0.93
C7A—C8A	1.460 (9)	C7B—C8B	1.465 (11)
C7A—H7AA	0.93	C7B—H7BA	0.93
C8A—C9A	1.386 (10)	C8B—C9B	1.404 (12)
C8A—C17A	1.442 (10)	C8B—C17B	1.432 (12)
C9A—C10A	1.429 (15)	C9B—C10B	1.397 (19)
C9A—H9AA	0.93	C9B—H9BA	0.93
C10A—C11A	1.417 (16)	C10B—C11B	1.385 (19)
C10A—C15A	1.51 (2)	C10B—C15B	1.41 (2)
C11A—C12A	1.360 (11)	C11B—C12B	1.378 (15)
C11A—H11A	0.93	C11B—H11B	0.93
C12A—C13A	1.416 (17)	C12B—C13B	1.424 (17)
C12A—H12A	0.93	C12B—H12B	0.93
C13A—C14A	1.38 (3)	C13B—C14B	1.40 (2)
C13A—H13A	0.93	C13B—H13B	0.93
C14A—C15A	1.43 (2)	C14B—C15B	1.435 (15)
C14A—H14A	0.93	C14B—H14B	0.93
C15A—C16A	1.23 (2)	C15B—C16B	1.52 (2)
C16A—C17A	1.372 (18)	C16B—C17B	1.313 (19)
C16A—H16A	0.93	C16B—H16B	0.93
C17A—H17A	0.93	C17B—H17B	0.93
C18A—H18A	0.96	C18B—H18G	0.96
C18A—H18B	0.96	C18B—H18D	0.96
C18A—H18C	0.96	C18B—H18E	0.96
C3A—N1A—C2A	117 (3)	C5B—C1B—C2B	118.5 (10)
C3A—N1A—C18A	123 (3)	C5B—C1B—H1BA	120.7
C2A—N1A—C18A	119 (2)	C2B—C1B—H1BA	120.7
C2A—C1A—C5A	122.2 (7)	N1B—C2B—C1B	123 (2)

C2A—C1A—H1AA	118.9	N1B—C2B—H2BA	118.7
C5A—C1A—H1AA	118.9	C1B—C2B—H2BA	118.7
C1A—C2A—N1A	120.9 (15)	C4B—C3B—N1B	120 (2)
C1A—C2A—H2AA	119.5	C4B—C3B—H3BA	120.2
N1A—C2A—H2AA	119.5	N1B—C3B—H3BA	120.2
N1A—C3A—C4A	123 (2)	C3B—C4B—C5B	122.4 (9)
N1A—C3A—H3AA	118.4	C3B—C4B—H4BA	118.8
C4A—C3A—H3AA	118.4	C5B—C4B—H4BA	118.8
C5A—C4A—C3A	120.0 (9)	C1B—C5B—C4B	116.6 (8)
C5A—C4A—H4AA	120.0	C1B—C5B—C6B	124.0 (8)
C3A—C4A—H4AA	120.0	C4B—C5B—C6B	119.5 (8)
C4A—C5A—C1A	116.0 (6)	C7B—C6B—C5B	127.8 (7)
C4A—C5A—C6A	123.9 (7)	C7B—C6B—H6BA	116.1
C1A—C5A—C6A	120.0 (7)	C5B—C6B—H6BA	116.1
C7A—C6A—C5A	123.4 (6)	C6B—C7B—C8B	124.5 (7)
C7A—C6A—H6AA	118.3	C6B—C7B—H7BA	117.8
C5A—C6A—H6AA	118.3	C8B—C7B—H7BA	117.8
C6A—C7A—C8A	127.9 (6)	C9B—C8B—C17B	118.4 (8)
C6A—C7A—H7AA	116.1	C9B—C8B—C7B	121.3 (8)
C8A—C7A—H7AA	116.1	C17B—C8B—C7B	120.3 (9)
C9A—C8A—C17A	120.8 (6)	C10B—C9B—C8B	121.8 (11)
C9A—C8A—C7A	117.2 (7)	C10B—C9B—H9BA	119.1
C17A—C8A—C7A	122.0 (7)	C8B—C9B—H9BA	119.1
C8A—C9A—C10A	118.7 (10)	C11B—C10B—C9B	123.3 (17)
C8A—C9A—H9AA	120.6	C11B—C10B—C15B	114.6 (13)
C10A—C9A—H9AA	120.6	C9B—C10B—C15B	121.6 (13)
C11A—C10A—C9A	119.1 (14)	C12B—C11B—C10B	121.5 (13)
C11A—C10A—C15A	125.2 (12)	C12B—C11B—H11B	119.3
C9A—C10A—C15A	115.3 (11)	C10B—C11B—H11B	119.3
C12A—C11A—C10A	121.0 (12)	C11B—C12B—C13B	120.7 (13)
C12A—C11A—H11A	119.5	C11B—C12B—H12B	119.7
C10A—C11A—H11A	119.5	C13B—C12B—H12B	119.7
C11A—C12A—C13A	119.0 (13)	C14B—C13B—C12B	123.1 (17)
C11A—C12A—H12A	120.5	C14B—C13B—H13B	118.4
C13A—C12A—H12A	120.5	C12B—C13B—H13B	118.4
C14A—C13A—C12A	117.7 (18)	C13B—C14B—C15B	110.9 (17)
C14A—C13A—H13A	121.1	C13B—C14B—H14B	124.6
C12A—C13A—H13A	121.1	C15B—C14B—H14B	124.6
C13A—C14A—C15A	132 (2)	C10B—C15B—C14B	129.0 (13)
C13A—C14A—H14A	114.2	C10B—C15B—C16B	114.3 (9)
C15A—C14A—H14A	114.2	C14B—C15B—C16B	116.6 (12)
C16A—C15A—C14A	131.7 (17)	C17B—C16B—C15B	122.9 (12)
C16A—C15A—C10A	123.1 (14)	C17B—C16B—H16B	118.5
C14A—C15A—C10A	105.1 (14)	C15B—C16B—H16B	118.5
C15A—C16A—C17A	123.3 (14)	C16B—C17B—C8B	120.8 (11)
C15A—C16A—H16A	118.4	C16B—C17B—H17B	119.6
C17A—C16A—H16A	118.4	C8B—C17B—H17B	119.6
C16A—C17A—C8A	118.6 (10)	N1B—C18B—H18G	109.5

C16A—C17A—H17A	120.7	N1B—C18B—H18D	109.5
C8A—C17A—H17A	120.7	H18G—C18B—H18D	109.5
C2B—N1B—C3B	120 (3)	N1B—C18B—H18E	109.5
C2B—N1B—C18B	123 (3)	H18G—C18B—H18E	109.5
C3B—N1B—C18B	116 (4)	H18D—C18B—H18E	109.5
C5A—C1A—C2A—N1A	6 (2)	C3B—N1B—C2B—C1B	12 (4)
C3A—N1A—C2A—C1A	-8 (3)	C18B—N1B—C2B—C1B	178 (2)
C18A—N1A—C2A—C1A	-177.7 (16)	C5B—C1B—C2B—N1B	-8 (3)
C2A—N1A—C3A—C4A	4 (3)	C2B—N1B—C3B—C4B	-9 (4)
C18A—N1A—C3A—C4A	174.0 (18)	C18B—N1B—C3B—C4B	-176.2 (19)
N1A—C3A—C4A—C5A	1 (3)	N1B—C3B—C4B—C5B	3 (3)
C3A—C4A—C5A—C1A	-3.3 (14)	C2B—C1B—C5B—C4B	2.0 (16)
C3A—C4A—C5A—C6A	176.4 (11)	C2B—C1B—C5B—C6B	-178.1 (12)
C2A—C1A—C5A—C4A	0.1 (13)	C3B—C4B—C5B—C1B	0.3 (17)
C2A—C1A—C5A—C6A	-179.6 (10)	C3B—C4B—C5B—C6B	-179.6 (13)
C4A—C5A—C6A—C7A	-0.4 (10)	C1B—C5B—C6B—C7B	0.7 (13)
C1A—C5A—C6A—C7A	179.2 (6)	C4B—C5B—C6B—C7B	-179.4 (8)
C5A—C6A—C7A—C8A	-179.7 (6)	C5B—C6B—C7B—C8B	179.0 (7)
C6A—C7A—C8A—C9A	-178.5 (6)	C6B—C7B—C8B—C9B	1.5 (12)
C6A—C7A—C8A—C17A	2.1 (10)	C6B—C7B—C8B—C17B	179.9 (8)
C17A—C8A—C9A—C10A	2.7 (14)	C17B—C8B—C9B—C10B	-1.0 (18)
C7A—C8A—C9A—C10A	-176.7 (12)	C7B—C8B—C9B—C10B	177.5 (14)
C8A—C9A—C10A—C11A	-177.1 (13)	C8B—C9B—C10B—C11B	175.4 (16)
C8A—C9A—C10A—C15A	-4 (2)	C8B—C9B—C10B—C15B	4 (3)
C9A—C10A—C11A—C12A	179.0 (13)	C9B—C10B—C11B—C12B	-175.2 (17)
C15A—C10A—C11A—C12A	6 (3)	C15B—C10B—C11B—C12B	-3 (3)
C10A—C11A—C12A—C13A	-4 (2)	C10B—C11B—C12B—C13B	1 (3)
C11A—C12A—C13A—C14A	2 (4)	C11B—C12B—C13B—C14B	-1 (4)
C12A—C13A—C14A—C15A	-2 (5)	C12B—C13B—C14B—C15B	2 (4)
C13A—C14A—C15A—C16A	-178 (3)	C11B—C10B—C15B—C14B	6 (3)
C13A—C14A—C15A—C10A	4 (4)	C9B—C10B—C15B—C14B	178 (2)
C11A—C10A—C15A—C16A	176.1 (17)	C11B—C10B—C15B—C16B	-177.2 (17)
C9A—C10A—C15A—C16A	3 (2)	C9B—C10B—C15B—C16B	-5 (3)
C11A—C10A—C15A—C14A	-6 (3)	C13B—C14B—C15B—C10B	-5 (4)
C9A—C10A—C15A—C14A	-178.7 (19)	C13B—C14B—C15B—C16B	178 (2)
C14A—C15A—C16A—C17A	-179 (2)	C10B—C15B—C16B—C17B	4 (2)
C10A—C15A—C16A—C17A	-1 (2)	C14B—C15B—C16B—C17B	-178 (2)
C15A—C16A—C17A—C8A	0 (2)	C15B—C16B—C17B—C8B	-1 (2)
C9A—C8A—C17A—C16A	-0.9 (13)	C9B—C8B—C17B—C16B	-0.2 (16)
C7A—C8A—C17A—C16A	178.5 (10)	C7B—C8B—C17B—C16B	-178.7 (12)

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C18A—H18A \cdots I1 ⁱ	0.96	3.05	3.928 (17)	152
C18A—H18B \cdots Cg1 ⁱ	0.96	2.63	3.513 (18)	153
C18A—H18B \cdots Cg2 ⁱ	0.96	2.65	3.517 (18)	150

$C18B—H18E\cdots Cg1^i$	0.96	2.62	3.44 (2)	143
$C18B—H18E\cdots Cg2^i$	0.96	2.66	3.45 (2)	139

Symmetry code: (i) $-x+1, -y+1, -z+2$.