

Crystal structure of (pyridin-2-ylmethylidene)(triphenylmethyl)amine

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The title Schiff base compound, C₂₅H₂₀N₂, crystallizes with two independent molecules (*A* and *B*) in the asymmetric unit. In both molecules, the imine group is approximately coplanar with the pyridine ring, with N—C—C—N torsion angles of 170.1 (3) and −172.0 (3) Å. In the crystal, *A* and *B* dimers are linked by pairs of C—H⋯π interactions and further C—H⋯π bonds link the dimers into a three-dimensional network.

Keywords: crystal structure; amine; pyridin-2-ylmethylidene; trityl; Schiff base ligands; magnetism.

CCDC reference: 1021733

1. Related literature

For the use of the pyridin-2-ylmethanimine Schiff base ligands for the development of a new generation of memory devices, multifunctional materials, and other magnetic applications, see: Capes *et al.* (2000); Guionneau *et al.* (2001); Létard *et al.* (1997, 1998); Liu *et al.* (2010); Murray (2008); Goodwin (2004); Gupta & Sutar (2008). For van der Waals radii, see: Bondi (1964)

2. Experimental

2.1. Crystal data

C ₂₅ H ₂₀ N ₂	<i>V</i> = 3696.7 (4) Å ³
<i>M_r</i> = 348.43	<i>Z</i> = 8
Orthorhombic, <i>P</i> 2 ₁ 2 ₁ 2 ₁	Cu <i>K</i> α radiation
<i>a</i> = 7.0719 (5) Å	<i>μ</i> = 0.56 mm ^{−1}
<i>b</i> = 16.3972 (8) Å	<i>T</i> = 173 K
<i>c</i> = 31.880 (2) Å	0.30 × 0.26 × 0.22 mm

2.2. Data collection

Agilent Xcalibur (Sapphire3, Gemini ultra) diffractometer	21735 measured reflections
Absorption correction: multi-scan (<i>CrysAlis PRO</i> and <i>CrysAlis RED</i> ; Agilent, 2012)	6965 independent reflections
<i>T</i> _{min} = 0.849, <i>T</i> _{max} = 0.886	5059 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R</i> _{int} = 0.058

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.057	487 parameters
<i>wR</i> (<i>F</i> ²) = 0.153	H-atom parameters constrained
<i>S</i> = 1.00	Δ <i>ρ</i> _{max} = 0.36 e Å ^{−3}
6965 reflections	Δ <i>ρ</i> _{min} = −0.18 e Å ^{−3}

Table 1

Hydrogen-bond geometry (Å, °).

*Cg*1 is the centroid of the N1, C1–C5 ring, *Cg*2 of ring C8–C13, *Cg*3 of ring C14–C19, *Cg*4 of ring C20–C25, *Cg*5 of ring N1', C1'–C5', *Cg*6 of ring C8'–C13', *Cg*7 of ring C14'–C19' and *Cg*8 of ring C20'–C25'.

D—H⋯A	D—H	H⋯A	H⋯Plane	D⋯A	D—H⋯A
C1—H1⋯Cg8	0.95	2.819	2.786 (3)	3.573	137
C1'—H1'⋯Cg4	0.95	2.789	2.749 (3)	3.545	137
C2—H2⋯Cg5 ⁱ	0.95	2.769	2.751 (3)	3.591	145
C2'—H2'⋯Cg1 ⁱⁱ	0.95	2.789	2.717 (3)	3.555	138
C4—H4⋯Cg2 ⁱⁱⁱ	0.95	3.150	2.982 (3)	3.692	118
C4'—H4'⋯Cg6 ^{iv}	0.95	3.169	2.942 (3)	3.687	116
C9—H9⋯Cg4	0.95	3.104	2.533 (4)	3.670	120
C9'—H9'⋯Cg8	0.95	3.061	2.526 (4)	3.642	121
C10—H10⋯Cg3 ^{iv}	0.95	3.077	3.038 (4)	3.823	136
C10'—H10'⋯Cg7 ⁱⁱⁱ	0.95	3.133	3.067 (4)	3.871	136
C15—H15⋯Cg2	0.95	3.030	2.386 (4)	3.717	130
C15'—H15'⋯Cg6	0.95	3.027	2.387 (4)	3.715	131
C21—H21⋯Cg3	0.95	2.897	2.424 (4)	3.614	133
C21'—H21'⋯Cg7	0.95	2.886	2.435 (4)	3.604	133
C25—H25⋯Cg1 ^{iv}	0.95	2.859	2.824 (3)	3.493	125
C25'—H25'⋯Cg5 ⁱⁱⁱ	0.95	2.865	2.824 (3)	3.496	125

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

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: TK5243).

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S1. Comment

Schiff bases and their complexes have received intense attention owing to their structural and functional diversities (Gupta & Sutar, 2008; Liu *et al.*, 2010). Much effort has been devoted in recent years to the design and synthesis of Schiff bases and their iron(II/III) complexes displaying spin–crossover (SCO) properties with the aim of developing a new generation of memory devices, multifunctional materials, and other magnetic applications (Murray, 2008; Goodwin, 2004 and references therein). The 2–pyridylmethanimine Schiff base derivative has been used as a chelating ligand in mononuclear iron(II) SCO complexes as they often possess field strengths that lie within the right region to facilitate temperature–mediated switching between the high spin (HS) and low spin (LS) states of the iron(II) centers (Guionneau *et al.*, 2001; Capes *et al.*, 2000; Létard *et al.*, 1998; Létard *et al.*, 1997). These ligands are also able to form non–covalent interactions such as hydrogen bonding and weaker electrostatic contacts, which will help to provide diversity to the architectures of structures, and can be associated with subtle effects on the molecular magnetic properties of compounds. For example, the *N*–2–pyridylmethylene–4–(phenylethynyl)aniline (PM–PEA) was employed with [Fe(NCS)₂] precursor complex to form the mononuclear SCO derivative, [Fe(PM–PEA)₂(NCS)₂], exhibiting a thermal hysteresis loop of 37 K (Létard *et al.*, 1997). The crystal structure has been studied at room temperature (HS state) and at 140 K (LS state) suggests that the cooperativity may be attributed to intermolecular π – π stacking between phenyl rings and crucially the strength of the C–H \cdots S interactions. Following the above, a family of iron(II) compounds containing the 2–pyridylmethanimine Schiff base derivatives have been prepared and found to show different SCO behaviors that range from very smooth and incomplete for [Fe(PM–TeA)₂(NCS)₂], PM–TeA = *N*–(2′–pyridylmethyl)–4–aminoterphenyl (Guionneau *et al.*, 2001), smooth with almost no hysteresis for [Fe(PM–AzA)₂(NCS)₂] and exceptionally abrupt for [Fe(PM–BiA)₂(NCS)₂], PM–AzA = *N*–(2′–pyridylmethyl)–4–(phenylazo)aniline, PM–BiA = *N*–(2′–pyridylmethyl)–4–amino–biphenyl (Capes *et al.*, 2000). Our interest in the mononuclear iron(II) SCO complexes has led us to prepare a new *N*–bidentate Schiff base derivative (I), which has the triphenylmethane group attached to the 2–pyridylmethanimine moiety. It is anticipated that weak non–covalent forces such as C–H \cdots π and π – π interactions will help to stabilize the assembly as well as increase the dimensionality of the structure and may also serve as weak exchange pathways between the magnetic metal center. Herein we report the supramolecular structure of (I), which is mediated by both intra– and intermolecular *ef* C–H \cdots π interactions.

The *N*–bidentate Schiff base derivative (I) can be obtained by condensation of 2–pyridinecarboxaldehyde with triphenylmethanamine. Single crystal *X*–ray diffraction analysis reveals that (I) crystallizes in the chiral orthorhombic space group *P*2₁2₁2₁, with two molecules (denoted *A* and *B*) representing the asymmetric unit as shown in Fig. 1. The bond distances within the phenyl (rings 2–4 and 6–8) and pyridine (rings 1 and 5) rings are in the range 1.341 (4) – 1.402 (4) Å, mean 1.381 Å, for molecule *A*, and 1.347 (4) – 1.400 (5) Å, mean 1.381 Å, for molecule *B*. The C–N distances in the

imine bonds are 1.268 (4) and 1.273 (4) Å for the C6—N2 and C6'—N2', respectively, which are in agreement with C=N double bond character for the imine group of Schiff bases type (Guionneau *et al.*, 2001). The C7—N2 and the C7'—N2' distances between the imine and the triphenylmethane groups are 1.483 (4) and 1.473 (4) Å, respectively. These are consistent with the values expected for C—N single bonding (Guionneau *et al.*, 2001). The bond angles C5—C6—N2 of 118.5 (2)° and the corresponding C5'—C6'—N2' of 117.9 (2)° are closed to the ideal value of 120°. This fact is further confirmed its sp^2 character.

The molecules *A* and *B* adopt different conformations (the torsion angles C6—N2—C7—C8 = 138.6 (3)° and C6'—N2'—C7'—C8' = -137.1 (2)°) in the crystal of (I) and are linked in inversion-symmetric pairs by *edge-to-face* (*ef*) C—H \cdots π type to form an *A*–*B* dimer. The imine group in both molecules *A* and *B* is approximately coplanar with the pyridine rings, with the N1—C5—C6—N2 and N1'—C5'—C6'—N2' torsion angles of 170.1 (3) and -172.0 (3) Å, respectively. In molecule *A*, the mean plane of the 2-pyridylmethanimine (ring 1—C6=N2) moiety and atom C7 [maximum deviation = 0.101 (2) Å for atom N2] is inclined to the phenyl rings 2, 3, and 4 by 47.2 (1), 84.7 (1), and 59.8 (1)°, respectively. Similarly, the dihedral angles between the mean plane of the corresponding ring 5—C6'=N2'—C7' [maximum deviation = 0.054 (2) Å for atom C6'] and the phenyl rings 6, 7, and 8 in the molecule *B* are 47.5 (1), 83.9 (1), and 59.4 (1)°, respectively. One of the interesting features of the crystal structure of (I) is the angle between the C=N bond and the rings of the triphenylmethane group. Namely, the angles of N2—C7—C20 (116.3 (2)°) and N2'—C7'—C20' (116.5 (2)°) are much larger than the corresponding angles of N2—C7—C8 (106.4 (2)°), N2'—C7'—C8' (106.2 (2)°), N2—C7—C14 (104.3 (2)°), and N2'—C7'—C14' (104.2 (2)°). This is presumably due to the inversion related intermolecular *ef* C—H \cdots π type in the dimer which lock the molecular conformations.

Several intra- and intermolecular *ef* C—H \cdots π interactions are observed in the crystal structure of (I) and have profound effects on both the molecular and the packing conformations. In the *A*–*B* dimer, hydrogen atoms, H1 and H1' of the pyridine rings (rings 1 and 5) are point toward the centroid (*Cg*) of the phenyl rings (4 and 8) at distance of 2.819 Å (C1—H1 \cdots Cg8 = 136.9°) for the H1 \cdots Cg8 and 2.789 Å (C1'—H1' \cdots Cg4 = 137.1°) for the H1' \cdots Cg4. The 2-pyridylmethanimine moiety of molecules *A* and *B* is almost parallel with the dihedral angle of 0.6 (1)°. It should be noted that the value of C \cdots N contact (C6 \cdots N1' = 3.943 (4) Å, C6—H6 \cdots N1' = 144.6°; C6' \cdots N1 = 3.929 (4) Å, C6'—H6' \cdots N1 = 144.3°) is greatly longer than the sum of the van der Waals radii [1.70 (C) + 1.54 (N) = 3.24 Å] (Bondi, 1964). Thus, no classical C—H \cdots N hydrogen bonds involving the imine and the pyridine groups are observed in the dimer. There are, however, additional intramolecular C—H \cdots π hydrogen bonding between the adjacent phenyl rings (rings 2–4 and 6–8) with the H \cdots Cg distances and the C—H \cdots Cg angle in the range 2.897–3.104 Å and 119.8–133.3°, Table 1. As shown in Fig. 2, the dimers are extended into a three-dimensional supramolecular architecture *via* the intermolecular *ef* C—H \cdots π interactions with the H \cdots Cg distance and the C—H \cdots Cg angle in the range 2.769–3.169 Å and 116.0–145.1°, respectively (Table 1). Finally, no classical hydrogen bonding and π – π stacking between adjacent molecules are observed in the crystal structure of (I). Its packing is mainly based on weak *ef* C—H \cdots π interactions.

S2. Experimental

To a solution of 2-pyridinecarboxaldehyde (1.90 ml, 0.02 mol) in benzene (100 ml), and a few drops of acetic acid as catalyst was added drop wise with triphenylmethanimine (5.20 g, 0.02 mol) at room temperature. The reaction mixture was stirred under reflux at 110 °C. After 6 h of refluxing, the yellow solution was neutralized with Na₂CO₃ (2 mmol), filtered, and concentrated to dryness in *vacuo*. The residue was recrystallized from a mixture of CH₂Cl₂ and petroleum ether (2:1, *v:v*) to give white crystalline solid of (I). Anal. Found (calcd) for C₂₅H₂₀N₂ (348.16): C, 86.15 (86.17); H, 5.81 (5.79); N, 8.05 (8.04).

S3. Refinement

The C-bound hydrogen atoms were placed in geometrically idealized positions based on chemical coordinations and constrained to ride on their parent atom positions with a C—H distances of 0.95 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for the aromatic H atoms.

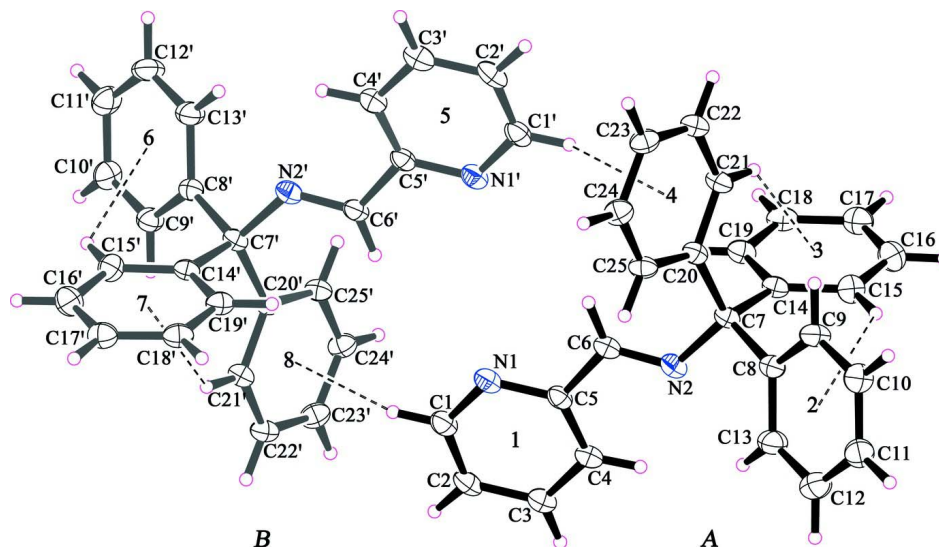


Figure 1

A view of the two symmetry-independent molecules (*A* and *B*) in (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Covalent bonds in *A* and *B* molecules are shaded differently. The labeling scheme, 1–8, applied to the aromatic rings are used to identify the rings in the subsequent discussion.

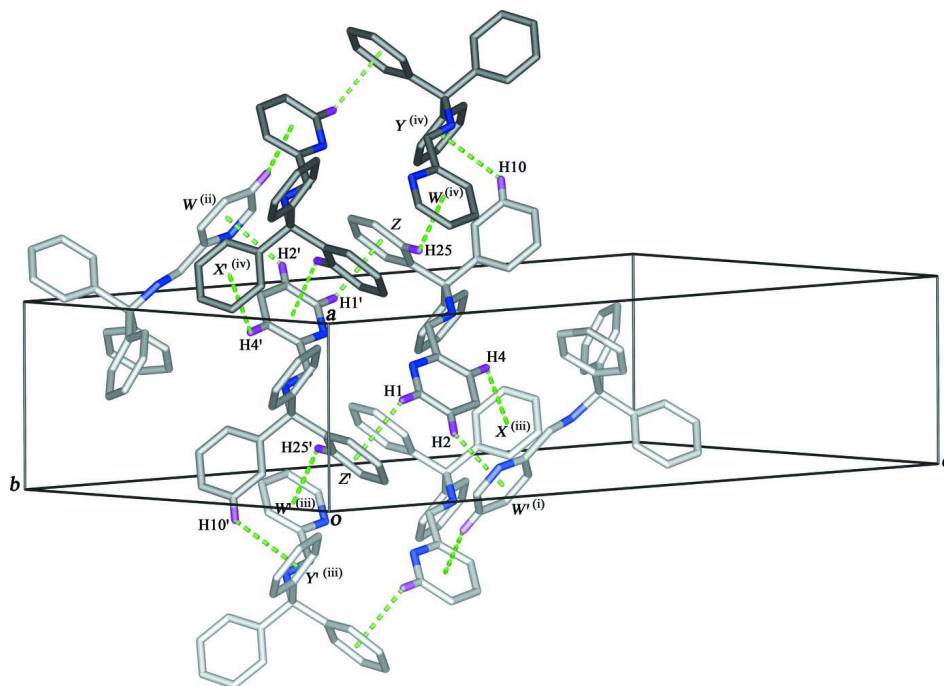


Figure 2

A packing diagram for (I), displaying the intermolecular *ef* C—H... π interactions (dashed lines). For clarity, only H atoms involved in C—H... π hydrogen bonding have been included. [Symmetry codes: (i) $1-x, 1/2+y, 3/2-z$; (ii) $-x, y-1/2, 3/2-z$; (iii) $1+x, y, z$; (iv) $x-1, y, z$].

(Pyridin-2-ylmethylidene)(triphenylmethyl)amine

Crystal data

$C_{25}H_{20}N_2$

$M_r = 348.43$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 7.0719$ (5) Å

$b = 16.3972$ (8) Å

$c = 31.880$ (2) Å

$V = 3696.7$ (4) Å³

$Z = 8$

$F(000) = 1472$

$D_x = 1.252$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å

Cell parameters from 3448 reflections

$\mu = 0.56$ mm⁻¹

$T = 173$ K

Block, yellow

$0.30 \times 0.26 \times 0.22$ mm

Data collection

Agilent Xcalibur (Sapphire3, Gemini ultra)
diffractometer

Radiation source: fine-focus sealed tube

Mirror monochromator

Detector resolution: 8 pixels mm⁻¹

ω scans

Absorption correction: multi-scan

(*CrysAlis PRO* and *CrysAlis RED*; Agilent,
2012)

$T_{\min} = 0.849, T_{\max} = 0.886$

21735 measured reflections

6965 independent reflections

5059 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.058$

$\theta_{\max} = 73.1^\circ, \theta_{\min} = 9.8^\circ$

$h = -8 \rightarrow 8$

$k = -19 \rightarrow 15$

$l = -28 \rightarrow 39$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.057$ $wR(F^2) = 0.153$ $S = 1.00$

6965 reflections

487 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0771P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.36 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), XXX Friedel
pairs

Absolute structure parameter: 0.3 (8)

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.3234 (4)	0.72675 (16)	0.72533 (8)	0.0395 (6)
N2	0.0763 (4)	0.67303 (16)	0.63297 (8)	0.0368 (6)
C1	0.4649 (5)	0.7751 (2)	0.73827 (11)	0.0408 (7)
H1	0.4933	0.7761	0.7674	0.049*
C2	0.5717 (5)	0.8232 (2)	0.71190 (11)	0.0419 (8)
H2	0.6713	0.8561	0.7226	0.050*
C3	0.5300 (5)	0.8225 (2)	0.66932 (10)	0.0419 (8)
H3	0.5988	0.8559	0.6504	0.050*
C4	0.3861 (5)	0.77209 (19)	0.65489 (10)	0.0385 (7)
H4	0.3559	0.7696	0.6259	0.046*
C5	0.2876 (4)	0.72545 (18)	0.68395 (9)	0.0339 (7)
C6	0.1354 (5)	0.66888 (19)	0.67048 (10)	0.0359 (7)
H6	0.0833	0.6304	0.6895	0.043*
C7	-0.0641 (4)	0.61515 (18)	0.61548 (9)	0.0335 (7)
C8	-0.2019 (5)	0.66622 (19)	0.58873 (9)	0.0355 (7)
C13	-0.1588 (5)	0.7448 (2)	0.57610 (11)	0.0465 (8)
H13	-0.0432	0.7689	0.5848	0.056*
C12	-0.2826 (6)	0.7891 (2)	0.55077 (12)	0.0560 (10)
H12	-0.2515	0.8432	0.5425	0.067*
C11	-0.4510 (6)	0.7541 (2)	0.53771 (11)	0.0510 (9)
H11	-0.5344	0.7835	0.5199	0.061*
C10	-0.4964 (5)	0.6768 (2)	0.55064 (11)	0.0470 (9)
H10	-0.6125	0.6529	0.5421	0.056*

C9	-0.3737 (5)	0.6334 (2)	0.57618 (11)	0.0432 (8)
H9	-0.4079	0.5802	0.5852	0.052*
C14	0.0542 (4)	0.55700 (18)	0.58757 (9)	0.0338 (7)
C15	0.0086 (5)	0.5395 (2)	0.54651 (10)	0.0426 (8)
H15	-0.0942	0.5670	0.5335	0.051*
C16	0.1108 (6)	0.4824 (2)	0.52396 (11)	0.0491 (9)
H16	0.0777	0.4712	0.4957	0.059*
C17	0.2588 (5)	0.4419 (2)	0.54197 (11)	0.0475 (9)
H17	0.3262	0.4019	0.5265	0.057*
C18	0.3102 (5)	0.4593 (2)	0.58270 (10)	0.0421 (8)
H18	0.4148	0.4322	0.5952	0.051*
C19	0.2088 (5)	0.51641 (19)	0.60531 (10)	0.0390 (7)
H19	0.2447	0.5283	0.6334	0.047*
C20	-0.1792 (4)	0.56558 (19)	0.64745 (9)	0.0330 (6)
C21	-0.2010 (5)	0.48179 (19)	0.64471 (10)	0.0373 (7)
H21	-0.1317	0.4526	0.6241	0.045*
C22	-0.3221 (5)	0.4391 (2)	0.67150 (10)	0.0403 (7)
H22	-0.3347	0.3817	0.6688	0.048*
C23	-0.4230 (5)	0.4802 (2)	0.70172 (10)	0.0407 (7)
H23	-0.5065	0.4516	0.7198	0.049*
C24	-0.4017 (5)	0.5638 (2)	0.70542 (10)	0.0418 (8)
H24	-0.4700	0.5925	0.7264	0.050*
C25	-0.2821 (5)	0.60567 (19)	0.67889 (10)	0.0374 (7)
H25	-0.2691	0.6630	0.6820	0.045*
N1'	0.1678 (4)	0.52217 (16)	0.76822 (8)	0.0386 (6)
C1'	0.0245 (5)	0.4739 (2)	0.75559 (11)	0.0412 (8)
H1'	-0.0053	0.4722	0.7266	0.049*
N2'	0.4146 (4)	0.57659 (16)	0.86033 (8)	0.0363 (6)
C2'	-0.0814 (5)	0.4267 (2)	0.78293 (11)	0.0447 (8)
H2'	-0.1825	0.3941	0.7728	0.054*
C3'	-0.0381 (5)	0.4277 (2)	0.82482 (11)	0.0453 (8)
H3'	-0.1075	0.3952	0.8441	0.054*
C4'	0.1078 (5)	0.4768 (2)	0.83851 (10)	0.0389 (7)
H4'	0.1404	0.4789	0.8674	0.047*
C5'	0.2058 (4)	0.52291 (18)	0.80935 (9)	0.0335 (6)
C6'	0.3573 (4)	0.58008 (19)	0.82251 (9)	0.0348 (7)
H6'	0.4100	0.6181	0.8033	0.042*
C7'	0.5550 (4)	0.63331 (18)	0.87805 (9)	0.0315 (6)
C8'	0.6910 (5)	0.58111 (19)	0.90443 (9)	0.0333 (6)
C9'	0.8659 (5)	0.6119 (2)	0.91612 (10)	0.0400 (7)
H9'	0.9025	0.6647	0.9069	0.048*
C10'	0.9893 (5)	0.5664 (2)	0.94125 (10)	0.0448 (8)
H10'	1.1085	0.5883	0.9490	0.054*
C11'	0.9380 (6)	0.4904 (2)	0.95454 (11)	0.0506 (9)
H11'	1.0206	0.4597	0.9720	0.061*
C12'	0.7658 (6)	0.4580 (2)	0.94262 (12)	0.0507 (9)
H12'	0.7315	0.4047	0.9515	0.061*
C13'	0.6430 (5)	0.5026 (2)	0.91780 (10)	0.0438 (8)

H13'	0.5251	0.4797	0.9098	0.053*
C14'	0.4373 (4)	0.69107 (18)	0.90604 (9)	0.0330 (7)
C15'	0.4871 (5)	0.7085 (2)	0.94706 (10)	0.0403 (8)
H15'	0.5904	0.6807	0.9596	0.048*
C16'	0.3873 (6)	0.7665 (2)	0.97014 (11)	0.0488 (9)
H16'	0.4250	0.7788	0.9980	0.059*
C17'	0.2338 (5)	0.8060 (2)	0.95266 (10)	0.0444 (8)
H17'	0.1664	0.8458	0.9683	0.053*
C18'	0.1797 (5)	0.7872 (2)	0.91230 (10)	0.0409 (8)
H18'	0.0728	0.8133	0.9003	0.049*
C19'	0.2796 (4)	0.73047 (19)	0.88923 (10)	0.0375 (7)
H19'	0.2404	0.7181	0.8615	0.045*
C20'	0.6711 (4)	0.68358 (18)	0.84642 (9)	0.0308 (6)
C21'	0.6930 (4)	0.76739 (19)	0.84992 (9)	0.0348 (7)
H21'	0.6231	0.7962	0.8706	0.042*
C22'	0.8149 (5)	0.8098 (2)	0.82383 (10)	0.0392 (7)
H22'	0.8284	0.8672	0.8268	0.047*
C23'	0.9176 (5)	0.7686 (2)	0.79330 (10)	0.0423 (8)
H23'	1.0034	0.7973	0.7758	0.051*
C24'	0.8937 (5)	0.68586 (19)	0.78877 (10)	0.0388 (7)
H24'	0.9604	0.6575	0.7674	0.047*
C25'	0.7734 (5)	0.64361 (19)	0.81508 (9)	0.0361 (7)
H25'	0.7599	0.5863	0.8118	0.043*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0380 (14)	0.0299 (15)	0.0505 (15)	-0.0005 (12)	-0.0043 (12)	-0.0053 (12)
N2	0.0331 (14)	0.0311 (14)	0.0461 (14)	-0.0026 (11)	-0.0023 (11)	-0.0050 (11)
C1	0.0409 (17)	0.0314 (19)	0.0502 (18)	0.0022 (15)	-0.0073 (15)	-0.0054 (15)
C2	0.0355 (17)	0.0300 (18)	0.060 (2)	-0.0027 (13)	-0.0094 (15)	-0.0092 (15)
C3	0.0403 (18)	0.0332 (18)	0.0521 (18)	-0.0052 (14)	-0.0007 (15)	-0.0021 (14)
C4	0.0396 (17)	0.0265 (17)	0.0495 (17)	0.0003 (14)	0.0015 (14)	-0.0030 (14)
C5	0.0333 (15)	0.0263 (16)	0.0420 (16)	0.0035 (12)	-0.0037 (13)	-0.0055 (13)
C6	0.0359 (16)	0.0253 (17)	0.0465 (17)	-0.0004 (13)	0.0014 (13)	-0.0039 (13)
C7	0.0322 (14)	0.0267 (16)	0.0415 (16)	-0.0058 (12)	-0.0007 (13)	-0.0021 (12)
C8	0.0372 (16)	0.0290 (17)	0.0401 (15)	-0.0018 (13)	0.0027 (13)	-0.0010 (13)
C13	0.049 (2)	0.037 (2)	0.0540 (19)	-0.0083 (16)	-0.0019 (16)	0.0041 (15)
C12	0.065 (3)	0.045 (2)	0.059 (2)	0.002 (2)	-0.006 (2)	0.0096 (17)
C11	0.056 (2)	0.043 (2)	0.053 (2)	0.0137 (17)	-0.0087 (17)	-0.0002 (16)
C10	0.0377 (18)	0.050 (2)	0.0531 (19)	0.0015 (16)	-0.0034 (15)	0.0010 (16)
C9	0.0402 (18)	0.0307 (18)	0.059 (2)	-0.0017 (14)	-0.0037 (15)	0.0065 (15)
C14	0.0317 (15)	0.0266 (16)	0.0431 (16)	-0.0042 (12)	0.0025 (13)	0.0002 (13)
C15	0.0359 (17)	0.047 (2)	0.0454 (18)	0.0007 (16)	-0.0005 (14)	-0.0005 (15)
C16	0.050 (2)	0.049 (2)	0.0481 (18)	0.0021 (18)	0.0021 (16)	-0.0076 (16)
C17	0.047 (2)	0.043 (2)	0.053 (2)	0.0017 (17)	0.0104 (16)	-0.0014 (16)
C18	0.0352 (17)	0.0341 (19)	0.0571 (19)	-0.0004 (14)	0.0041 (15)	0.0052 (14)
C19	0.0350 (16)	0.0318 (18)	0.0502 (17)	-0.0028 (13)	-0.0005 (14)	0.0003 (14)

C20	0.0318 (15)	0.0264 (16)	0.0408 (15)	0.0008 (12)	-0.0018 (12)	-0.0020 (12)
C21	0.0386 (17)	0.0269 (17)	0.0464 (17)	-0.0060 (14)	0.0037 (14)	-0.0038 (13)
C22	0.0430 (18)	0.0280 (17)	0.0499 (17)	-0.0053 (14)	0.0003 (15)	0.0005 (13)
C23	0.0346 (16)	0.040 (2)	0.0474 (17)	-0.0012 (14)	0.0032 (14)	0.0097 (14)
C24	0.0387 (17)	0.040 (2)	0.0473 (18)	0.0079 (15)	0.0038 (14)	0.0039 (14)
C25	0.0379 (16)	0.0262 (17)	0.0481 (17)	0.0054 (13)	0.0046 (14)	0.0021 (13)
N1'	0.0375 (14)	0.0302 (15)	0.0480 (14)	0.0020 (11)	-0.0039 (11)	-0.0060 (12)
C1'	0.0427 (17)	0.0289 (19)	0.0519 (18)	0.0064 (14)	-0.0076 (14)	-0.0104 (14)
N2'	0.0331 (13)	0.0301 (14)	0.0456 (14)	-0.0055 (11)	-0.0010 (11)	-0.0025 (11)
C2'	0.0357 (17)	0.0340 (18)	0.064 (2)	-0.0061 (14)	-0.0058 (16)	-0.0139 (16)
C3'	0.0401 (18)	0.0351 (19)	0.061 (2)	-0.0049 (14)	0.0048 (16)	-0.0064 (16)
C4'	0.0385 (17)	0.0319 (18)	0.0462 (17)	-0.0006 (14)	0.0031 (14)	-0.0048 (14)
C5'	0.0335 (15)	0.0243 (16)	0.0428 (15)	0.0037 (12)	-0.0029 (13)	-0.0049 (12)
C6'	0.0370 (16)	0.0239 (17)	0.0435 (17)	-0.0012 (12)	-0.0038 (13)	-0.0021 (13)
C7'	0.0290 (14)	0.0265 (16)	0.0388 (15)	-0.0028 (12)	-0.0010 (12)	-0.0025 (12)
C8'	0.0365 (15)	0.0267 (16)	0.0368 (15)	0.0007 (13)	0.0003 (12)	-0.0002 (12)
C9'	0.0385 (17)	0.0355 (19)	0.0459 (18)	-0.0047 (14)	-0.0047 (14)	0.0032 (14)
C10'	0.0358 (17)	0.050 (2)	0.0484 (18)	0.0008 (16)	-0.0064 (14)	0.0027 (16)
C11'	0.051 (2)	0.050 (2)	0.0513 (19)	0.0112 (18)	-0.0074 (17)	0.0075 (17)
C12'	0.060 (2)	0.030 (2)	0.063 (2)	0.0001 (17)	-0.0034 (19)	0.0102 (15)
C13'	0.046 (2)	0.0341 (19)	0.0510 (18)	-0.0077 (15)	-0.0027 (15)	0.0047 (14)
C14'	0.0325 (15)	0.0267 (16)	0.0398 (15)	-0.0073 (12)	0.0017 (12)	0.0016 (12)
C15'	0.0358 (17)	0.045 (2)	0.0399 (16)	0.0008 (15)	-0.0025 (13)	0.0014 (14)
C16'	0.050 (2)	0.047 (2)	0.0494 (19)	0.0037 (18)	0.0028 (16)	-0.0078 (16)
C17'	0.0445 (19)	0.038 (2)	0.0506 (19)	0.0042 (16)	0.0092 (15)	0.0007 (15)
C18'	0.0340 (16)	0.035 (2)	0.0533 (19)	0.0007 (14)	0.0028 (14)	0.0058 (14)
C19'	0.0332 (15)	0.0323 (18)	0.0469 (17)	-0.0017 (13)	-0.0008 (13)	0.0043 (14)
C20'	0.0259 (13)	0.0249 (16)	0.0416 (15)	-0.0033 (11)	-0.0026 (12)	-0.0005 (12)
C21'	0.0344 (16)	0.0277 (17)	0.0422 (16)	-0.0030 (13)	0.0005 (13)	-0.0034 (13)
C22'	0.0396 (17)	0.0289 (18)	0.0492 (17)	-0.0049 (14)	-0.0004 (15)	0.0030 (13)
C23'	0.0366 (17)	0.043 (2)	0.0472 (18)	0.0011 (15)	0.0050 (14)	0.0106 (15)
C24'	0.0386 (17)	0.0349 (19)	0.0430 (17)	0.0078 (14)	0.0067 (14)	0.0047 (14)
C25'	0.0387 (16)	0.0256 (16)	0.0440 (16)	0.0019 (13)	0.0019 (13)	-0.0004 (13)

Geometric parameters (Å, °)

N1—C1	1.341 (4)	N1'—C5'	1.339 (4)
N1—C5	1.344 (4)	N1'—C1'	1.347 (4)
N2—C6	1.268 (4)	C1'—C2'	1.385 (5)
N2—C7	1.483 (4)	C1'—H1'	0.9500
C1—C2	1.379 (5)	N2'—C6'	1.273 (4)
C1—H1	0.9500	N2'—C7'	1.473 (4)
C2—C3	1.389 (5)	C2'—C3'	1.370 (5)
C2—H2	0.9500	C2'—H2'	0.9500
C3—C4	1.389 (5)	C3'—C4'	1.379 (5)
C3—H3	0.9500	C3'—H3'	0.9500
C4—C5	1.388 (4)	C4'—C5'	1.384 (4)
C4—H4	0.9500	C4'—H4'	0.9500

C5—C6	1.485 (4)	C5'—C6'	1.484 (4)
C6—H6	0.9500	C6'—H6'	0.9500
C7—C20	1.537 (4)	C7'—C8'	1.538 (4)
C7—C8	1.542 (4)	C7'—C20'	1.539 (4)
C7—C14	1.550 (4)	C7'—C14'	1.545 (4)
C8—C13	1.384 (4)	C8'—C9'	1.387 (4)
C8—C9	1.388 (5)	C8'—C13'	1.397 (4)
C13—C12	1.395 (5)	C9'—C10'	1.400 (5)
C13—H13	0.9500	C9'—H9'	0.9500
C12—C11	1.386 (6)	C10'—C11'	1.366 (5)
C12—H12	0.9500	C10'—H10'	0.9500
C11—C10	1.371 (5)	C11'—C12'	1.381 (6)
C11—H11	0.9500	C11'—H11'	0.9500
C10—C9	1.387 (5)	C12'—C13'	1.384 (5)
C10—H10	0.9500	C12'—H12'	0.9500
C9—H9	0.9500	C13'—H13'	0.9500
C14—C15	1.378 (4)	C14'—C15'	1.384 (4)
C14—C19	1.399 (4)	C14'—C19'	1.396 (4)
C15—C16	1.385 (5)	C15'—C16'	1.393 (5)
C15—H15	0.9500	C15'—H15'	0.9500
C16—C17	1.366 (5)	C16'—C17'	1.382 (5)
C16—H16	0.9500	C16'—H16'	0.9500
C17—C18	1.378 (5)	C17'—C18'	1.378 (5)
C17—H17	0.9500	C17'—H17'	0.9500
C18—C19	1.383 (4)	C18'—C19'	1.380 (4)
C18—H18	0.9500	C18'—H18'	0.9500
C19—H19	0.9500	C19'—H19'	0.9500
C20—C21	1.385 (4)	C20'—C21'	1.388 (4)
C20—C25	1.402 (4)	C20'—C25'	1.397 (4)
C21—C22	1.397 (4)	C21'—C22'	1.385 (4)
C21—H21	0.9500	C21'—H21'	0.9500
C22—C23	1.375 (5)	C22'—C23'	1.390 (5)
C22—H22	0.9500	C22'—H22'	0.9500
C23—C24	1.383 (5)	C23'—C24'	1.375 (5)
C23—H23	0.9500	C23'—H23'	0.9500
C24—C25	1.379 (5)	C24'—C25'	1.381 (4)
C24—H24	0.9500	C24'—H24'	0.9500
C25—H25	0.9500	C25'—H25'	0.9500
C1—N1—C5	116.9 (3)	C5'—N1'—C1'	116.7 (3)
C6—N2—C7	122.8 (3)	N1'—C1'—C2'	123.2 (3)
N1—C1—C2	124.0 (3)	N1'—C1'—H1'	118.4
N1—C1—H1	118.0	C2'—C1'—H1'	118.4
C2—C1—H1	118.0	C6'—N2'—C7'	123.3 (3)
C1—C2—C3	118.4 (3)	C3'—C2'—C1'	119.1 (3)
C1—C2—H2	120.8	C3'—C2'—H2'	120.5
C3—C2—H2	120.8	C1'—C2'—H2'	120.5
C2—C3—C4	119.0 (3)	C2'—C3'—C4'	118.8 (3)

C2—C3—H3	120.5	C2'—C3'—H3'	120.6
C4—C3—H3	120.5	C4'—C3'—H3'	120.6
C5—C4—C3	118.3 (3)	C3'—C4'—C5'	118.7 (3)
C5—C4—H4	120.8	C3'—C4'—H4'	120.6
C3—C4—H4	120.8	C5'—C4'—H4'	120.6
N1—C5—C4	123.5 (3)	N1'—C5'—C4'	123.5 (3)
N1—C5—C6	115.5 (3)	N1'—C5'—C6'	115.3 (3)
C4—C5—C6	121.0 (3)	C4'—C5'—C6'	121.1 (3)
N2—C6—C5	118.6 (3)	N2'—C6'—C5'	118.0 (3)
N2—C6—H6	120.7	N2'—C6'—H6'	121.0
C5—C6—H6	120.7	C5'—C6'—H6'	121.0
N2—C7—C20	116.3 (2)	N2'—C7'—C8'	106.2 (2)
N2—C7—C8	106.5 (2)	N2'—C7'—C20'	116.5 (2)
C20—C7—C8	108.6 (3)	C8'—C7'—C20'	108.8 (2)
N2—C7—C14	104.4 (2)	N2'—C7'—C14'	104.2 (2)
C20—C7—C14	110.0 (3)	C8'—C7'—C14'	111.2 (2)
C8—C7—C14	111.0 (2)	C20'—C7'—C14'	109.7 (2)
C13—C8—C9	118.0 (3)	C9'—C8'—C13'	118.0 (3)
C13—C8—C7	121.9 (3)	C9'—C8'—C7'	120.1 (3)
C9—C8—C7	120.1 (3)	C13'—C8'—C7'	121.9 (3)
C8—C13—C12	121.0 (4)	C8'—C9'—C10'	121.0 (3)
C8—C13—H13	119.5	C8'—C9'—H9'	119.5
C12—C13—H13	119.5	C10'—C9'—H9'	119.5
C11—C12—C13	119.8 (4)	C11'—C10'—C9'	119.9 (3)
C11—C12—H12	120.1	C11'—C10'—H10'	120.1
C13—C12—H12	120.1	C9'—C10'—H10'	120.1
C10—C11—C12	119.6 (4)	C10'—C11'—C12'	120.0 (3)
C10—C11—H11	120.2	C10'—C11'—H11'	120.0
C12—C11—H11	120.2	C12'—C11'—H11'	120.0
C11—C10—C9	120.3 (4)	C11'—C12'—C13'	120.5 (4)
C11—C10—H10	119.8	C11'—C12'—H12'	119.8
C9—C10—H10	119.8	C13'—C12'—H12'	119.8
C10—C9—C8	121.2 (3)	C12'—C13'—C8'	120.6 (3)
C10—C9—H9	119.4	C12'—C13'—H13'	119.7
C8—C9—H9	119.4	C8'—C13'—H13'	119.7
C15—C14—C19	117.9 (3)	C15'—C14'—C19'	118.0 (3)
C15—C14—C7	123.1 (3)	C15'—C14'—C7'	122.4 (3)
C19—C14—C7	118.9 (3)	C19'—C14'—C7'	119.5 (3)
C14—C15—C16	120.8 (3)	C14'—C15'—C16'	120.7 (3)
C14—C15—H15	119.6	C14'—C15'—H15'	119.6
C16—C15—H15	119.6	C16'—C15'—H15'	119.6
C17—C16—C15	120.7 (3)	C17'—C16'—C15'	120.3 (3)
C17—C16—H16	119.7	C17'—C16'—H16'	119.8
C15—C16—H16	119.7	C15'—C16'—H16'	119.8
C16—C17—C18	119.9 (3)	C18'—C17'—C16'	119.3 (3)
C16—C17—H17	120.1	C18'—C17'—H17'	120.3
C18—C17—H17	120.1	C16'—C17'—H17'	120.3
C17—C18—C19	119.6 (3)	C17'—C18'—C19'	120.4 (3)

C17—C18—H18	120.2	C17'—C18'—H18'	119.8
C19—C18—H18	120.2	C19'—C18'—H18'	119.8
C18—C19—C14	121.1 (3)	C18'—C19'—C14'	121.1 (3)
C18—C19—H19	119.4	C18'—C19'—H19'	119.5
C14—C19—H19	119.4	C14'—C19'—H19'	119.5
C21—C20—C25	116.9 (3)	C21'—C20'—C25'	117.7 (3)
C21—C20—C7	122.8 (3)	C21'—C20'—C7'	122.5 (3)
C25—C20—C7	120.1 (3)	C25'—C20'—C7'	119.6 (3)
C20—C21—C22	121.7 (3)	C22'—C21'—C20'	121.3 (3)
C20—C21—H21	119.1	C22'—C21'—H21'	119.4
C22—C21—H21	119.1	C20'—C21'—H21'	119.4
C23—C22—C21	120.1 (3)	C21'—C22'—C23'	120.1 (3)
C23—C22—H22	120.0	C21'—C22'—H22'	120.0
C21—C22—H22	120.0	C23'—C22'—H22'	120.0
C22—C23—C24	119.2 (3)	C24'—C23'—C22'	119.3 (3)
C22—C23—H23	120.4	C24'—C23'—H23'	120.4
C24—C23—H23	120.4	C22'—C23'—H23'	120.4
C25—C24—C23	120.5 (3)	C23'—C24'—C25'	120.4 (3)
C25—C24—H24	119.7	C23'—C24'—H24'	119.8
C23—C24—H24	119.7	C25'—C24'—H24'	119.8
C24—C25—C20	121.5 (3)	C24'—C25'—C20'	121.2 (3)
C24—C25—H25	119.2	C24'—C25'—H25'	119.4
C20—C25—H25	119.2	C20'—C25'—H25'	119.4
C5—N1—C1—C2	0.9 (5)	C5'—N1'—C1'—C2'	0.1 (5)
N1—C1—C2—C3	0.5 (5)	N1'—C1'—C2'—C3'	-0.9 (5)
C1—C2—C3—C4	-1.5 (5)	C1'—C2'—C3'—C4'	1.0 (5)
C2—C3—C4—C5	1.1 (5)	C2'—C3'—C4'—C5'	-0.3 (5)
C1—N1—C5—C4	-1.3 (5)	C1'—N1'—C5'—C4'	0.7 (4)
C1—N1—C5—C6	177.6 (3)	C1'—N1'—C5'—C6'	-176.8 (3)
C3—C4—C5—N1	0.3 (5)	C3'—C4'—C5'—N1'	-0.6 (5)
C3—C4—C5—C6	-178.5 (3)	C3'—C4'—C5'—C6'	176.7 (3)
C7—N2—C6—C5	175.0 (3)	C7'—N2'—C6'—C5'	-175.9 (3)
N1—C5—C6—N2	170.1 (3)	N1'—C5'—C6'—N2'	-172.0 (3)
C4—C5—C6—N2	-10.9 (5)	C4'—C5'—C6'—N2'	10.5 (4)
C6—N2—C7—C20	17.4 (4)	C6'—N2'—C7'—C8'	-137.2 (3)
C6—N2—C7—C8	138.6 (3)	C6'—N2'—C7'—C20'	-15.8 (4)
C6—N2—C7—C14	-103.9 (3)	C6'—N2'—C7'—C14'	105.2 (3)
N2—C7—C8—C13	15.9 (4)	N2'—C7'—C8'—C9'	162.9 (3)
C20—C7—C8—C13	141.9 (3)	C20'—C7'—C8'—C9'	36.7 (4)
C14—C7—C8—C13	-97.1 (3)	C14'—C7'—C8'—C9'	-84.3 (3)
N2—C7—C8—C9	-164.9 (3)	N2'—C7'—C8'—C13'	-18.0 (4)
C20—C7—C8—C9	-38.9 (4)	C20'—C7'—C8'—C13'	-144.2 (3)
C14—C7—C8—C9	82.1 (4)	C14'—C7'—C8'—C13'	94.8 (3)
C9—C8—C13—C12	-1.3 (5)	C13'—C8'—C9'—C10'	-1.2 (5)
C7—C8—C13—C12	177.9 (3)	C7'—C8'—C9'—C10'	178.0 (3)
C8—C13—C12—C11	-0.4 (6)	C8'—C9'—C10'—C11'	0.1 (5)
C13—C12—C11—C10	1.5 (6)	C9'—C10'—C11'—C12'	1.1 (6)

C12—C11—C10—C9	-0.9 (6)	C10'—C11'—C12'—C13'	-1.1 (6)
C11—C10—C9—C8	-0.9 (5)	C11'—C12'—C13'—C8'	-0.1 (6)
C13—C8—C9—C10	2.0 (5)	C9'—C8'—C13'—C12'	1.2 (5)
C7—C8—C9—C10	-177.2 (3)	C7'—C8'—C13'—C12'	-178.0 (3)
N2—C7—C14—C15	-129.1 (3)	N2'—C7'—C14'—C15'	130.6 (3)
C20—C7—C14—C15	105.4 (3)	C8'—C7'—C14'—C15'	16.5 (4)
C8—C7—C14—C15	-14.8 (4)	C20'—C7'—C14'—C15'	-103.9 (3)
N2—C7—C14—C19	54.7 (4)	N2'—C7'—C14'—C19'	-52.4 (3)
C20—C7—C14—C19	-70.8 (3)	C8'—C7'—C14'—C19'	-166.5 (3)
C8—C7—C14—C19	169.0 (3)	C20'—C7'—C14'—C19'	73.0 (3)
C19—C14—C15—C16	1.2 (5)	C19'—C14'—C15'—C16'	-2.8 (5)
C7—C14—C15—C16	-175.0 (3)	C7'—C14'—C15'—C16'	174.2 (3)
C14—C15—C16—C17	0.2 (6)	C14'—C15'—C16'—C17'	1.6 (6)
C15—C16—C17—C18	-1.5 (6)	C15'—C16'—C17'—C18'	0.5 (6)
C16—C17—C18—C19	1.4 (5)	C16'—C17'—C18'—C19'	-1.2 (5)
C17—C18—C19—C14	0.1 (5)	C17'—C18'—C19'—C14'	-0.1 (5)
C15—C14—C19—C18	-1.4 (5)	C15'—C14'—C19'—C18'	2.1 (5)
C7—C14—C19—C18	175.1 (3)	C7'—C14'—C19'—C18'	-175.0 (3)
N2—C7—C20—C21	-130.8 (3)	N2'—C7'—C20'—C21'	130.7 (3)
C8—C7—C20—C21	109.1 (3)	C8'—C7'—C20'—C21'	-109.3 (3)
C14—C7—C20—C21	-12.5 (4)	C14'—C7'—C20'—C21'	12.6 (4)
N2—C7—C20—C25	55.1 (4)	N2'—C7'—C20'—C25'	-55.4 (4)
C8—C7—C20—C25	-64.9 (4)	C8'—C7'—C20'—C25'	64.6 (3)
C14—C7—C20—C25	173.4 (3)	C14'—C7'—C20'—C25'	-173.5 (3)
C25—C20—C21—C22	1.1 (5)	C25'—C20'—C21'—C22'	-1.3 (5)
C7—C20—C21—C22	-173.2 (3)	C7'—C20'—C21'—C22'	172.7 (3)
C20—C21—C22—C23	-0.3 (5)	C20'—C21'—C22'—C23'	0.3 (5)
C21—C22—C23—C24	-0.6 (5)	C21'—C22'—C23'—C24'	1.4 (5)
C22—C23—C24—C25	0.6 (5)	C22'—C23'—C24'—C25'	-2.0 (5)
C23—C24—C25—C20	0.2 (5)	C23'—C24'—C25'—C20'	0.9 (5)
C21—C20—C25—C24	-1.0 (5)	C21'—C20'—C25'—C24'	0.7 (5)
C7—C20—C25—C24	173.4 (3)	C7'—C20'—C25'—C24'	-173.5 (3)