

10-(Prop-2-yn-1-yl)-2,7-diazaphenothiazine¹

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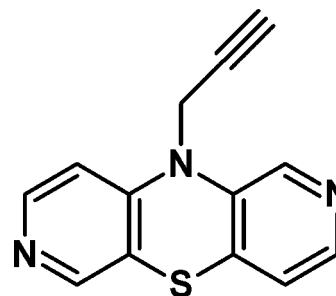
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.049; wR factor = 0.114; data-to-parameter ratio = 15.6.

In the title molecule [systematic name: 10-(prop-2-yn-1-yl)-dipyrido[3,4-*b*:3',4'-*e*][1,4]thiazine], $\text{C}_{13}\text{H}_9\text{N}_3\text{S}$, the dihedral angle between the two pyridine rings is 146.33 (7)° and the angle between two halves of the thiazine ring is 138.84 (8)°, resulting in a butterfly shape for the tricyclic system. The central thiazine ring adopts a boat conformation, with the 2-propynyl substituent at the thiazine N atom located in a pseudo-equatorial position and oriented to the concave side of the diazaphenothiazine system. In the crystal, molecules are arranged *via* $\pi-\pi$ interactions between the pyridine rings [centroid-centroid distances = 3.838 (1) and 3.845 (1) Å] into stacks extending along [001]. There are C—H···C and C—H···N interactions between molecules of neighbouring stacks.

Related literature

For recent literature on the biological activity of phenothiazines, see: Aaron *et al.* (2009); Pluta *et al.* (2011). For the structure of 10-(2-propynyl)phenothiazine and its transformations into anticancer derivatives, see: Bisi *et al.* (2008). For the synthesis and the anticancer and immunosuppressive activity of 2,7-diazaphenothiazines, see: Morak-Młodawska & Pluta (2009); Zimecki *et al.* (2009); Pluta *et al.* (2010). For planar and folded structures of the 2,7-diazaphenothiazine system, see: Morak *et al.* (2002); Morak-Młodawska *et al.* (2010). For alkylation of azaphenothiazines, see: Pluta *et al.* (2009).



Experimental

Crystal data

$\text{C}_{13}\text{H}_9\text{N}_3\text{S}$	$V = 1070.20$ (12) Å ³
$M_r = 239.29$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 14.1150$ (9) Å	$\mu = 0.28$ mm ⁻¹
$b = 10.1909$ (6) Å	$T = 100$ K
$c = 7.6749$ (5) Å	$0.60 \times 0.50 \times 0.35$ mm
$\beta = 104.212$ (3)°	

Data collection

Nonius KappaCCD diffractometer upgraded with APEXII detector	2407 independent reflections
7015 measured reflections	2011 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.049$	154 parameters
$wR(F^2) = 0.114$	H-atom parameters constrained
$S = 1.11$	$\Delta\rho_{\text{max}} = 0.45$ e Å ⁻³
2407 reflections	$\Delta\rho_{\text{min}} = -0.35$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C4}-\text{H4}\cdots\text{N2}^{\text{i}}$	0.95	2.62	3.457 (3)	147
$\text{C13}-\text{H13}\cdots\text{C11}^{\text{ii}}$	0.95	2.78	3.677 (3)	159
$\text{C13}-\text{H13}\cdots\text{C12}^{\text{ii}}$	0.95	2.78	3.686 (3)	161
$\text{C3}-\text{H3}\cdots\text{C13}^{\text{i}}$	0.95	2.78	3.662 (3)	155
$\text{C8}-\text{H8}\cdots\text{C13}^{\text{iii}}$	0.95	2.69	3.407 (3)	133

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

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP III* (Burnett & Johnson, 1996) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2475).

¹ Azinyl sulfides. Part CXXVII.

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supplementary materials

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10-(Prop-2-yn-1-yl)-2,7-diazaphenothiazine

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Comment

Phenothiazines exhibit not only recognized neuroleptic, antihistaminic and antitussive activities but recently also anticancer, antibacterial and reversal multidrug resistance [Aaron *et al.*, (2009); Pluta *et al.*, (2011)]. The modifications of the phenothiazine structures are mainly directed into the introduction of new pharmacophoric substituents at the thiazine nitrogen atom and the substitution of the benzene ring with an azine ring (Pluta *et al.*, 2009, 2011). Synthesis of substituted 10-(2-propynyl)phenothiazines and their transformations into various aminobutynyl derivatives of anticancer and multidrug resistance reverting activities was reported by Bisi *et al.* (2008). We modified the phenothiazine structure *via* the substitution of the benzene ring with the pyridine ring to form 2,7-diazaphenothiazines (Morak-Młodawska & Pluta, 2009) possessing anticancer and immunosuppressive activities (Zimecki *et al.*, 2009; Pluta *et al.* 2010). Alkylation of azaphenothiazines proceeds at the thiazine and/or the azine nitrogen atoms, depending on the reaction conditions (Pluta *et al.*, 2009). *N*-Alkylation of 10*H*-2,7-diazaphenothiazine led to both types of the products showing planar and folded 2,7-diazaphenothiazine ring system (Morak-Młodawska *et al.*, 2010). 10*H*-2,7-Diazaphenothiazine was transformed into the title compound, C₁₃H₉N₃S, a convenient substrate to other 2,7-diazaphenothiazine derivatives using aminomethylation or 1,3-dipolar cycloaddition. The X-ray study showed the propynyl group to be attached to the thiazine nitrogen atom. In the molecule, the butterfly angle between the two pyridine rings is 146.33 (7)° and the angle between two halves of the thiazine ring is 138.84 (8)°. The 2-propynyl substituent is in a pseudo-equatorial position with the angle S5⋯N10–C11 of 163.8 (2)° and directed to the concave side of the diazaphenothiazine system with the angle between the N10/C11/C12/C13 and C4a/C5a/C9a/C10a planes of 86.3 (1)°. The thiazine nitrogen atom shows pyramidal geometry as the sum of the C–N10–C bond angles is 356.1 (1)°. Hydrogen bond C4–H4⋯N2 (Table 1) results in one-dimensional polymeric chain parallel to the *b* axis. Acidic hydrogen atom H13 is in close contact to C11 and C12 atoms of the propynyl substituent (both H⋯C distances equal to 2.78 Å). This suggests, that H13 is involved in C–H⋯C interactions to these two carbon atoms rather than in the C–H⋯ π interaction to the π system of the triple C12=C13 bond (H13⋯center_{C12=C13} distance of 2.96 Å). Additionally, the C12=C13 bond π electrons interact with two aromatic H atoms (H3 and H8) of two other adjacent molecules with short C–H⋯C intermolecular contacts (less than the sum of van der Waals radii) between H3 and H8, and C13 (see Table 1). On the basis of these interactions a three-dimensional network is formed. Molecules π -stack along the *c* axis. Aromatic rings N2/C1/C10a/C4a/C4/C3 π -stack with centroid-to-centroid distance of 3.845 (1) Å, similarly, for rings N7/C6/C5a/C9a/C9/C8 the centroid-to-centroid distance is 3.838 (1) Å (see Figure 2).

Experimental

To a suspension of 10*H*-2,7-diazaphenothiazine (100 mg, 0.5 mmol) in 5 ml DMF potassium *tert*-butoxide (80 mg, 0.72 mmol) was added. The mixture was stirred at room temperature for 1 h. Then a solution of propargyl bromide (80 mg, 0.64 mmol) in toluene was added dropwise. The solution was stirred at room temperature for 24 h and poured into water (15 ml), extracted with methylene chloride (15 ml), dried with Na₂SO₄ and evaporated to the brown oil. The residue was

purified by column chromatography (silica gel, CHCl_3) to yield 10-(2-propynyl)-2,7-diazaphenothiazine (72 mg, 60%), mp. 149–150°C. $^1\text{H NMR}$ in CDCl_3 : δ 2.57 (t, $J = 2.5$ Hz, 1H), 4.54 (d, $J = 2.5$ Hz, 2H), 7.14 (m, 2H, H-9, H-4), 8.12 (s, 1H, H-1), 8.22 (d, $J = 5.5$ Hz, H-3), 8.35 (d, $J = 5.5$ Hz, H-8), 8.40 (s, 1H, H-6). FAB MS: 240 ($M+H$, 100), 201 ($M-\text{CH}_2\text{CCH}+1$, 45).

Refinement

All H atoms in the were treated as riding atoms in geometrically idealized positions, with C–H distances of 0.95 (aromatic and acetylene) or 0.99 Å (methylene), and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Computing details

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP III* (Burnett & Johnson, 1996) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *pubCIF* (Westrip, 2010).

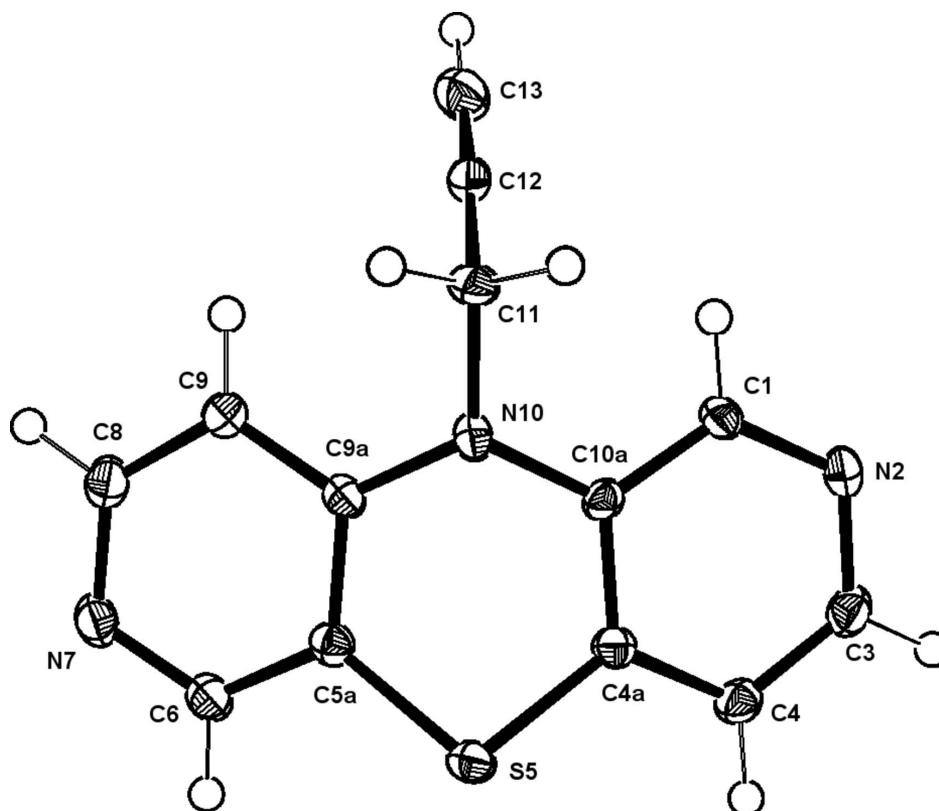
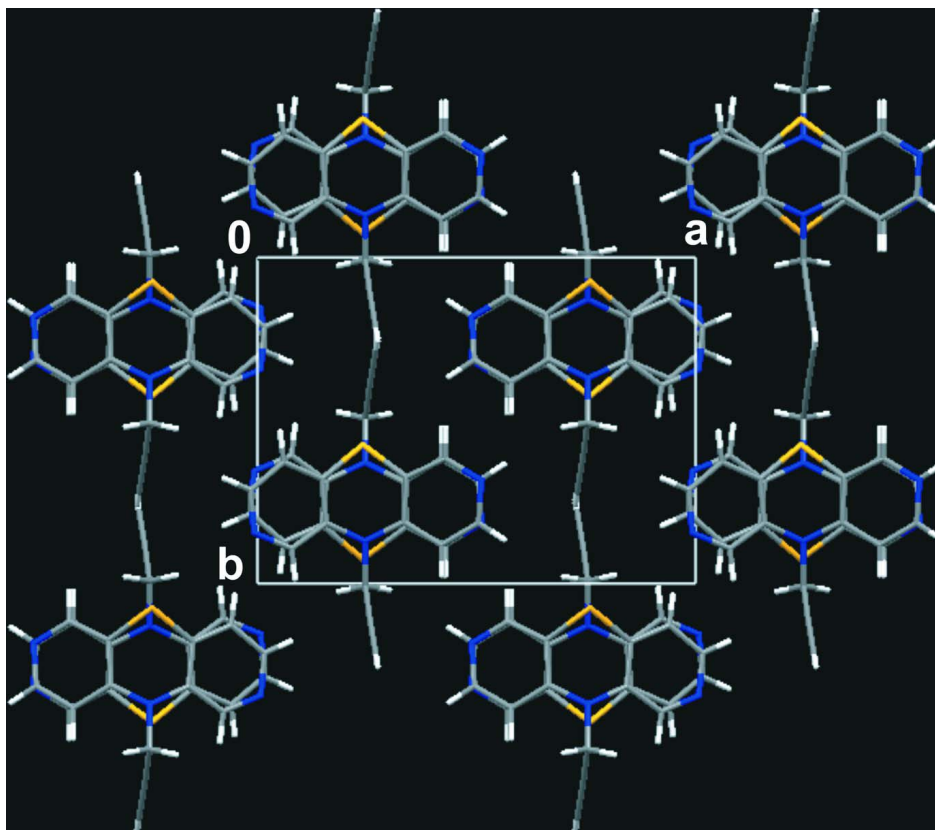


Figure 1

ORTEP drawing with displacement ellipsoids shown at the 50% probability level.

**Figure 2**

Crystal packing shown along the *c* axis.

10-(Prop-2-yn-1-yl)-2,7-diazaphenothiazine

Crystal data

$C_{13}H_9N_3S$

$M_r = 239.29$

Monoclinic, $P2_1/c$

Hall symbol: $-P 2_1/c$

$a = 14.1150$ (9) Å

$b = 10.1909$ (6) Å

$c = 7.6749$ (5) Å

$\beta = 104.212$ (3)°

$V = 1070.20$ (12) Å³

$Z = 4$

$F(000) = 496$

$D_x = 1.485$ Mg m⁻³

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

Cell parameters from 2229 reflections

$\theta = 2.5$ – 27.5 °

$\mu = 0.28$ mm⁻¹

$T = 100$ K

Block, yellow

$0.60 \times 0.50 \times 0.35$ mm

Data collection

Nonius KappaCCD

diffractometer upgraded with APEXII detector

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.3 pixels mm⁻¹

ω scan

7015 measured reflections

2407 independent reflections

2011 reflections with $I > 2\sigma(I)$

$R_{int} = 0.041$

$\theta_{max} = 27.5$ °, $\theta_{min} = 3.4$ °

$h = -18$ → 18

$k = -12$ → 13

$l = -9$ → 9

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.114$
 $S = 1.11$
 2407 reflections
 154 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0285P)^2 + 1.4885P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.45 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.35 \text{ e } \text{\AA}^{-3}$

Special details

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.92188 (15)	0.1074 (2)	0.5605 (3)	0.0172 (4)
H1	0.9145	0.0148	0.5522	0.021*
C3	1.01573 (17)	0.2865 (2)	0.6685 (3)	0.0218 (5)
H3	1.0762	0.3225	0.7335	0.026*
C4	0.94013 (16)	0.3721 (2)	0.5945 (3)	0.0195 (5)
H4	0.9479	0.4640	0.6135	0.023*
C4a	0.85325 (16)	0.3212 (2)	0.4927 (3)	0.0159 (4)
C5a	0.65940 (16)	0.3263 (2)	0.3914 (3)	0.0164 (4)
C6	0.57174 (16)	0.3834 (2)	0.4013 (3)	0.0197 (5)
H6	0.5703	0.4759	0.4161	0.024*
C8	0.49541 (17)	0.1852 (2)	0.3756 (3)	0.0226 (5)
H8	0.4377	0.1351	0.3665	0.027*
C9	0.58017 (16)	0.1180 (2)	0.3718 (3)	0.0192 (5)
H9	0.5801	0.0249	0.3640	0.023*
C9a	0.66564 (15)	0.1889 (2)	0.3797 (3)	0.0154 (4)
C10a	0.84271 (15)	0.1850 (2)	0.4757 (3)	0.0147 (4)
C11	0.75374 (17)	-0.0072 (2)	0.3179 (3)	0.0182 (5)
H11a	0.8152	-0.0251	0.2819	0.022*
H11b	0.6992	-0.0198	0.2102	0.022*
C12	0.74397 (16)	-0.1054 (2)	0.4530 (3)	0.0194 (5)
C13	0.73370 (18)	-0.1884 (3)	0.5542 (4)	0.0284 (6)
H13	0.7254	-0.2552	0.6356	0.034*
S5	0.75938 (4)	0.42507 (5)	0.37659 (8)	0.01880 (16)
N2	1.00796 (13)	0.15579 (19)	0.6532 (3)	0.0204 (4)
N7	0.48877 (14)	0.3163 (2)	0.3912 (3)	0.0226 (4)
N10	0.75434 (13)	0.12936 (18)	0.3721 (2)	0.0158 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0160 (10)	0.0151 (10)	0.0217 (11)	0.0027 (8)	0.0071 (9)	0.0011 (8)
C3	0.0170 (11)	0.0228 (12)	0.0250 (12)	-0.0042 (10)	0.0041 (9)	-0.0051 (10)
C4	0.0194 (11)	0.0165 (10)	0.0244 (12)	-0.0023 (9)	0.0085 (9)	-0.0050 (9)
C4a	0.0165 (10)	0.0147 (10)	0.0178 (11)	0.0005 (8)	0.0070 (8)	0.0003 (8)
C5a	0.0176 (10)	0.0166 (10)	0.0142 (10)	0.0009 (8)	0.0025 (8)	0.0013 (8)
C6	0.0195 (11)	0.0186 (10)	0.0197 (11)	0.0028 (9)	0.0023 (9)	-0.0007 (9)
C8	0.0168 (11)	0.0245 (12)	0.0250 (12)	-0.0011 (9)	0.0024 (9)	-0.0024 (10)
C9	0.0175 (10)	0.0170 (10)	0.0217 (11)	-0.0004 (9)	0.0022 (9)	-0.0013 (9)
C9a	0.0155 (10)	0.0156 (10)	0.0143 (10)	0.0027 (8)	0.0022 (8)	-0.0005 (8)
C10a	0.0141 (10)	0.0144 (10)	0.0169 (11)	-0.0017 (8)	0.0064 (8)	0.0000 (8)
C11	0.0199 (11)	0.0136 (10)	0.0218 (11)	0.0011 (9)	0.0064 (9)	-0.0028 (9)
C12	0.0149 (10)	0.0177 (10)	0.0250 (12)	0.0000 (9)	0.0037 (9)	-0.0034 (9)
C13	0.0256 (13)	0.0242 (12)	0.0360 (15)	0.0034 (10)	0.0086 (11)	0.0067 (11)
S5	0.0187 (3)	0.0143 (3)	0.0241 (3)	0.0015 (2)	0.0066 (2)	0.0035 (2)
N2	0.0131 (9)	0.0226 (10)	0.0248 (10)	0.0021 (8)	0.0033 (7)	-0.0002 (8)
N7	0.0164 (9)	0.0244 (10)	0.0254 (11)	0.0039 (8)	0.0020 (8)	-0.0014 (8)
N10	0.0119 (8)	0.0185 (9)	0.0167 (9)	0.0014 (7)	0.0030 (7)	-0.0005 (7)

Geometric parameters (\AA , $^\circ$)

C1—N2	1.342 (3)	C6—H6	0.9500
C1—C10a	1.393 (3)	C8—N7	1.347 (3)
C1—H1	0.9500	C8—C9	1.385 (3)
C3—N2	1.339 (3)	C8—H8	0.9500
C3—C4	1.387 (3)	C9—C9a	1.395 (3)
C3—H3	0.9500	C9—H9	0.9500
C4—C4a	1.383 (3)	C9a—N10	1.405 (3)
C4—H4	0.9500	C10a—N10	1.422 (3)
C4a—C10a	1.398 (3)	C11—N10	1.452 (3)
C4a—S5	1.758 (2)	C11—C12	1.471 (3)
C5a—C6	1.386 (3)	C11—H11a	0.9900
C5a—C9a	1.407 (3)	C11—H11b	0.9900
C5a—S5	1.760 (2)	C12—C13	1.181 (3)
C6—N7	1.342 (3)	C13—H13	0.9500
N2—C1—C10a	123.9 (2)	C8—C9—H9	120.5
N2—C1—H1	118.1	C9a—C9—H9	120.5
C10a—C1—H1	118.1	C9—C9a—N10	122.98 (19)
N2—C3—C4	123.5 (2)	C9—C9a—C5a	116.82 (19)
N2—C3—H3	118.3	N10—C9a—C5a	120.18 (19)
C4—C3—H3	118.3	C1—C10a—C4a	117.7 (2)
C4a—C4—C3	118.8 (2)	C1—C10a—N10	121.87 (19)
C4a—C4—H4	120.6	C4a—C10a—N10	120.42 (19)
C3—C4—H4	120.6	N10—C11—C12	116.42 (19)
C4—C4a—C10a	119.0 (2)	N10—C11—H11a	108.2
C4—C4a—S5	120.91 (17)	C12—C11—H11a	108.2
C10a—C4a—S5	120.04 (17)	N10—C11—H11b	108.2

C6—C5a—C9a	119.5 (2)	C12—C11—H11b	108.2
C6—C5a—S5	120.26 (17)	H11a—C11—H11b	107.3
C9a—C5a—S5	120.09 (16)	C13—C12—C11	176.5 (3)
N7—C6—C5a	124.1 (2)	C12—C13—H13	180.0
N7—C6—H6	117.9	C4a—S5—C5a	98.00 (10)
C5a—C6—H6	117.9	C3—N2—C1	117.1 (2)
N7—C8—C9	124.9 (2)	C6—N7—C8	115.6 (2)
N7—C8—H8	117.6	C9a—N10—C10a	118.21 (18)
C9—C8—H8	117.6	C9a—N10—C11	118.89 (18)
C8—C9—C9a	119.0 (2)	C10a—N10—C11	119.00 (18)
N2—C3—C4—C4a	-2.9 (4)	C4—C4a—S5—C5a	-148.01 (19)
C3—C4—C4a—C10a	3.3 (3)	C10a—C4a—S5—C5a	35.26 (19)
C3—C4—C4a—S5	-173.43 (17)	C6—C5a—S5—C4a	148.16 (19)
C9a—C5a—C6—N7	-3.6 (4)	C9a—C5a—S5—C4a	-36.6 (2)
S5—C5a—C6—N7	171.68 (18)	C4—C3—N2—C1	0.1 (3)
N7—C8—C9—C9a	-1.8 (4)	C10a—C1—N2—C3	2.2 (3)
C8—C9—C9a—N10	-178.4 (2)	C5a—C6—N7—C8	1.9 (3)
C8—C9—C9a—C5a	0.1 (3)	C9—C8—N7—C6	0.8 (4)
C6—C5a—C9a—C9	2.4 (3)	C9—C9a—N10—C10a	-144.8 (2)
S5—C5a—C9a—C9	-172.88 (17)	C5a—C9a—N10—C10a	36.8 (3)
C6—C5a—C9a—N10	-179.1 (2)	C9—C9a—N10—C11	12.8 (3)
S5—C5a—C9a—N10	5.6 (3)	C5a—C9a—N10—C11	-165.6 (2)
N2—C1—C10a—C4a	-1.7 (3)	C1—C10a—N10—C9a	142.9 (2)
N2—C1—C10a—N10	177.10 (19)	C4a—C10a—N10—C9a	-38.3 (3)
C4—C4a—C10a—C1	-1.2 (3)	C1—C10a—N10—C11	-14.6 (3)
S5—C4a—C10a—C1	175.63 (16)	C4a—C10a—N10—C11	164.16 (19)
C4—C4a—C10a—N10	179.98 (19)	C12—C11—N10—C9a	-76.0 (3)
S5—C4a—C10a—N10	-3.2 (3)	C12—C11—N10—C10a	81.3 (2)

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>
C4—H4...N2 ⁱ	0.95	2.62	3.457 (3)	147
C13—H13...C11 ⁱⁱ	0.95	2.78	3.677 (3)	159
C13—H13...C12 ⁱⁱ	0.95	2.78	3.686 (3)	161
C3—H3...C13 ⁱ	0.95	2.78	3.662 (3)	155
C8—H8...C13 ⁱⁱⁱ	0.95	2.69	3.407 (3)	133

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