

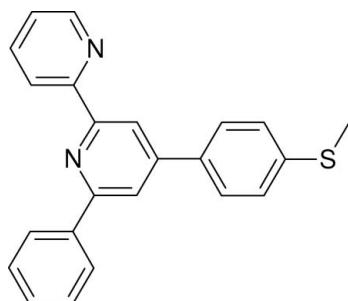
**4-[4-(Methylsulfanyl)phenyl]-6-phenyl-2,2'-bipyridine****Matthew I. J. Polson,<sup>a\*</sup> Franco Scandola<sup>b</sup> and Peter J. Steel<sup>a</sup>**<sup>a</sup>Chemistry Department, University of Canterbury, PO Box 4800, Christchurch, New Zealand, and <sup>b</sup>Dipartimento de Chimica, Università degli Studi di Ferrara, Ferrara 44100, Italy

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Key indicators: single-crystal X-ray study;  $T = 93\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$ ; disorder in main residue;  $R$  factor = 0.063;  $wR$  factor = 0.161; data-to-parameter ratio = 13.2.

The structure of the title compound,  $C_{23}H_{18}N_2S$ , is revealed by X-ray diffraction to be almost planar over all four aromatic rings; the pendant rings are at angles of 10.18, 14.12 and 15.42° relative to the central pyridine ring for the 4-methylsulfanyl, 2-pyridyl and 6-phenyl rings, respectively. The 2,6-aromatic substituents are disordered over two sites in a 0.6:0.4 occupancy ratio.

**Related literature**For related literature, see: Fitchett *et al.* (2005).**Experimental***Crystal data*

$C_{23}H_{18}N_2S$	$V = 1756.3 (5)\text{ \AA}^3$
$M_r = 354.45$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 19.189 (3)\text{ \AA}$	$\mu = 0.19\text{ mm}^{-1}$
$b = 5.3617 (8)\text{ \AA}$	$T = 93 (2)\text{ K}$
$c = 17.084 (3)\text{ \AA}$	$0.45 \times 0.17 \times 0.04\text{ mm}$
$\beta = 92.262 (9)^\circ$	

*Data collection*

Bruker APEXII CCD area-detector diffractometer	19871 measured reflections
Absorption correction: multi-scan ( <i>SADABS</i> ; Bruker, 2007)	3118 independent reflections
$T_{\min} = 0.599$ , $T_{\max} = 0.992$	1442 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.121$

*Refinement*

$R[F^2 > 2\sigma(F^2)] = 0.063$	236 parameters
$wR(F^2) = 0.161$	H-atom parameters constrained
$S = 0.93$	$\Delta\rho_{\max} = 0.50\text{ e \AA}^{-3}$
3118 reflections	$\Delta\rho_{\min} = -0.35\text{ e \AA}^{-3}$

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Version 1.08; Farrugia, 1997); software used to prepare material for publication: *publCIF* (Westrip, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WW2105).

**References**

- Bruker (2007). *APEX2* (Version 2.1-4), *SAINT* (Version 7.34A) and *SADABS* (Version 2004/1). Bruker AXS Inc., Madison, Wisconsin, USA.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Fitchett, C. M., Richardson, C. & Steel, P. J. (2005). *Org. Biomol. Chem.* **3**, 498–502.
- Sheldrick, G. M. (1990). *Acta Cryst. A* **46**, 467–473.
- Sheldrick, G. M. (1997). *SHELXL97*. University of Göttingen, Germany.
- Westrip, S. P. (2008). *publCIF*. In preparation.

# supporting information

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## 4-[4-(Methylsulfanyl)phenyl]-6-phenyl-2,2'-bipyridine

Matthew I. J. Polson, Franco Scandola and Peter J. Steel

### S1. Comment

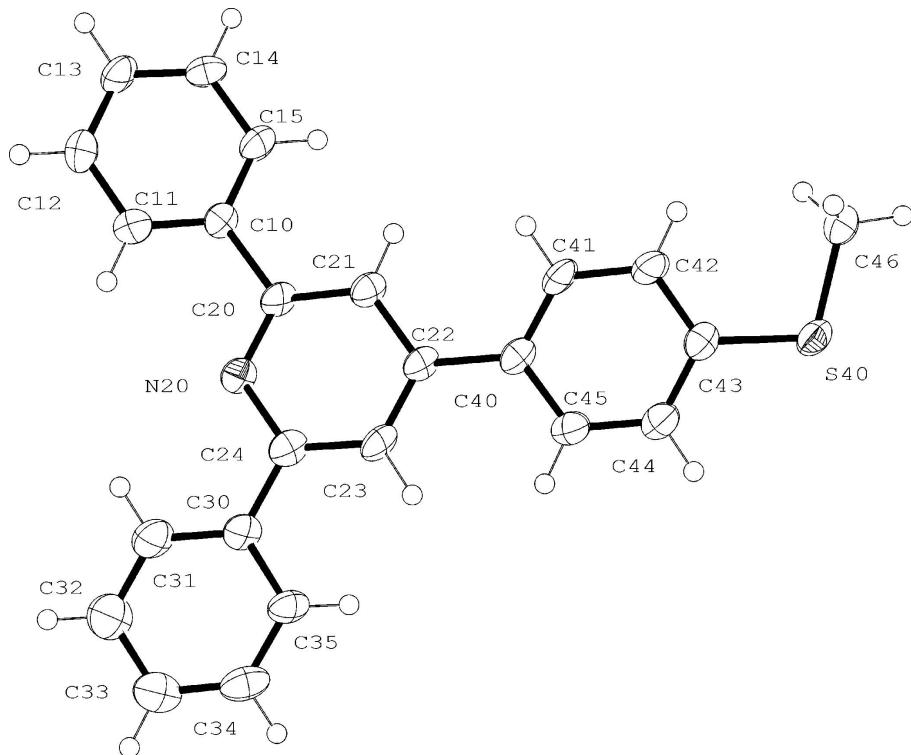
The use of Self Assembled Monolayers (SAMs) in the fabrication of molecular devices is a rapidly expanding field. To incorporate the useful photophysical properties of iridium complexes into a SAM, ligands must be capable of attaching to a surface. The compound (1), a bipyridine based ligand, includes a protected thiol group for attachment to a gold surface and a phenyl group for cyclometallation. Typically bipyridine ligands crystallize with the pyridine N atoms in a *s-trans* arrangement (Fitchett *et al.*, 2005). This is attributed to reduction of C—H/H—C interactions. Here, the pyridine ring and the phenyl ring crystallize in identical conformations, leading to disorder. If C—H/H—C interactions were the dominant force for the arrangement of the ring, one would expect the phenyl ring to adopt a different arrangement due to the additional interaction. This implies that the dominant force for the arrangement of the rings is the attractive C—H/N interaction.

### S2. Experimental

To a solution of 4-(methylsulfanyl)benzaldehyde (5 g), acetophenone (4.5 g), methanol (300 ml) and ammonia (0.81 g/ml, 50 ml) was added a sodium hydroxide solution (1.5 g in 50 ml water) with stirring. Overnight a precipitate of the condensation product formed. This was filtered, air dried and was used in the next step without further purification. This compound (5 g) was ground in a mortar and pestle with 2-acetylpyridine (2.5 g) and sodium hydroxide (0.83 g) until the mixture became a solid again. Excess ammonium hydroxide was added and the mixture dissolved in glacial acetic acid (50 ml) and was refluxed with stirring for 4 h. On cooling, the solution was poured into water (200 ml) and extracted with dichloromethane (3 x 50 ml). Chromatography on silica gel with dichloromethane/methanol (95:5) yielded the pure product (1). Single crystals suitable for X-ray diffraction formed on slow evaporation from dichloromethane solution. Yield = 2.3 g (25%). Spectroscopic data:  $^1\text{H}$  NMR ( $\text{CDCl}_3$ ):  $\delta$  2.54 (3H, s,  $\text{CH}_3\text{S}$ ), 7.35 (1H, ddd, py5'), 7.37 (2H, d, thio3,5), 7.46 (1H, t, ph4), 7.53 (2H, dd, thio-ph2,6), 7.77 (2H, d, ph3,5), 7.87 (1H, td, py4'), 7.95 (1H, d, py5), 8.20 (2H, d, ph2,6), 8.63 (1H, d, py3), 8.68 (1H, d, py3'), 8.72 (1H, dd, py6');  $^{13}\text{C}$  NMR ( $\text{CDCl}_3$ ):  $\delta$  15.5, 117.1, 118.1, 121.6, 123.9, 126.5, 127.1, 127.5, 128.7, 129.1, 135.0, 137.1, 139.4, 140.1, 148.8, 149.5, 156.0, 156.2, 157.2.

### S3. Refinement

The 2-pyridine and 6-phenyl rings are disordered in a 60/40 ratio over the two possible positions. The pyridine ring however always adopts a *s-trans* arrangement to the central pyridine nitrogen, presumably to minimize hydrogen/hydrogen repulsions.

**Figure 1**

The molecular structure of (1), showing displacement ellipsoids at the 50% probability level.

#### 4-[4-(Methylsulfanyl)phenyl]-6-phenyl-2,2'-bipyridine

##### Crystal data

$C_{23}H_{18}N_2S$

$M_r = 354.45$

Monoclinic,  $P2_1/c$

Hall symbol: -P 2ybc

$a = 19.189(3)$  Å

$b = 5.3617(8)$  Å

$c = 17.084(3)$  Å

$\beta = 92.262(9)^\circ$

$V = 1756.3(5)$  Å<sup>3</sup>

$Z = 4$

$F(000) = 744$

$D_x = 1.341$  Mg m<sup>-3</sup>

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

Cell parameters from 1819 reflections

$\theta = 2.7\text{--}25.9^\circ$

$\mu = 0.19$  mm<sup>-1</sup>

$T = 93$  K

Plate, yellow

$0.45 \times 0.17 \times 0.04$  mm

##### Data collection

Bruker APEXII CCD area-detector  
diffractometer

Radiation source: sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Bruker, 2007)

$T_{\min} = 0.599$ ,  $T_{\max} = 0.992$

19871 measured reflections

3118 independent reflections

1442 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.121$

$\theta_{\max} = 25.1^\circ$ ,  $\theta_{\min} = 2.4^\circ$

$h = -22 \rightarrow 22$

$k = -6 \rightarrow 6$

$l = -20 \rightarrow 20$

*Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.063$$

$$wR(F^2) = 0.161$$

$$S = 0.93$$

3118 reflections

236 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

$$w = 1/[\sigma^2(F_o^2) + (0.0562P)^2 + 0.5007P]$$
$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.50 \text{ e \AA}^{-3}$$

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

*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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C10	0.7765 (2)	0.5910 (7)	0.4911 (2)	0.0262 (10)	
C11	0.8384 (2)	0.6255 (8)	0.5330 (3)	0.0467 (13)	
H11	0.8788	0.5333	0.5204	0.056*	
C12	0.8411 (2)	0.7978 (9)	0.5941 (3)	0.0575 (15)	
H12	0.8832	0.8205	0.6245	0.069*	
C13	0.7843 (2)	0.9318 (8)	0.6101 (2)	0.0383 (11)	
H13	0.7862	1.0516	0.6511	0.046*	
C14	0.7242 (2)	0.8947 (9)	0.5672 (3)	0.0440 (12)	
H14	0.6839	0.9883	0.5792	0.053*	
N15	0.71985 (18)	0.7258 (8)	0.5069 (2)	0.0428 (10)	0.60
C15	0.71985 (18)	0.7258 (8)	0.5069 (2)	0.0428 (10)	0.40
H15	0.6775	0.7046	0.4769	0.051*	0.40
N20	0.82507 (15)	0.2544 (7)	0.41731 (17)	0.0296 (8)	
C20	0.7730 (2)	0.4149 (8)	0.4222 (2)	0.0278 (10)	
C21	0.7176 (2)	0.4300 (8)	0.3674 (2)	0.0300 (10)	
H21	0.6811	0.5471	0.3741	0.036*	
C22	0.71595 (19)	0.2711 (8)	0.3022 (2)	0.0269 (9)	
C23	0.7708 (2)	0.1064 (8)	0.2974 (2)	0.0322 (11)	
H23	0.7720	-0.0040	0.2540	0.039*	
C24	0.8246 (2)	0.0983 (8)	0.3549 (2)	0.0314 (10)	
C30	0.8832 (2)	-0.0776 (8)	0.3508 (2)	0.0306 (10)	
C31	0.9434 (2)	-0.0504 (9)	0.3971 (3)	0.0453 (13)	
H31	0.9469	0.0856	0.4327	0.054*	
C32	0.9981 (3)	-0.2125 (10)	0.3934 (3)	0.0545 (14)	
H32	1.0387	-0.1896	0.4262	0.065*	

C33	0.9939 (2)	-0.4123 (9)	0.3407 (3)	0.0454 (13)	
H33	1.0314	-0.5261	0.3361	0.054*	
C34	0.9338 (2)	-0.4371 (9)	0.2964 (3)	0.0441 (12)	
H34	0.9297	-0.5751	0.2617	0.053*	
N35	0.87914 (19)	-0.2752 (8)	0.2989 (2)	0.0354 (10)	0.40
C35	0.87914 (19)	-0.2752 (8)	0.2989 (2)	0.0354 (10)	0.60
H35	0.8389	-0.2979	0.2655	0.042*	0.60
C40	0.65707 (19)	0.2792 (8)	0.2432 (2)	0.0283 (10)	
C41	0.6070 (2)	0.4625 (8)	0.2429 (2)	0.0380 (11)	
H41	0.6118	0.5930	0.2803	0.046*	
C42	0.5494 (2)	0.4669 (8)	0.1906 (2)	0.0393 (12)	
H42	0.5163	0.5981	0.1928	0.047*	
C43	0.54081 (19)	0.2767 (8)	0.1349 (2)	0.0302 (10)	
C44	0.5919 (2)	0.0955 (9)	0.1327 (3)	0.0442 (12)	
H44	0.5880	-0.0321	0.0943	0.053*	
C45	0.6486 (2)	0.0961 (8)	0.1857 (3)	0.0411 (12)	
H45	0.6827	-0.0318	0.1826	0.049*	
S40	0.46881 (5)	0.2541 (2)	0.06834 (6)	0.0347 (3)	
C46	0.4188 (2)	0.5257 (8)	0.0889 (2)	0.0422 (12)	
H46A	0.4083	0.5279	0.1446	0.063*	
H46B	0.3752	0.5232	0.0571	0.063*	
H46C	0.4456	0.6752	0.0763	0.063*	

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C10	0.032 (2)	0.030 (2)	0.016 (2)	-0.001 (2)	0.0034 (18)	0.005 (2)
C11	0.048 (3)	0.050 (3)	0.041 (3)	0.015 (2)	-0.006 (2)	-0.015 (3)
C12	0.056 (3)	0.071 (4)	0.044 (3)	0.019 (3)	-0.024 (2)	-0.021 (3)
C13	0.052 (3)	0.043 (3)	0.019 (3)	0.003 (2)	-0.004 (2)	-0.007 (2)
C14	0.046 (3)	0.050 (3)	0.037 (3)	0.014 (2)	0.004 (2)	-0.019 (3)
N15	0.043 (2)	0.058 (3)	0.028 (2)	0.007 (2)	-0.0014 (17)	-0.019 (2)
C15	0.043 (2)	0.058 (3)	0.028 (2)	0.007 (2)	-0.0014 (17)	-0.019 (2)
N20	0.0384 (19)	0.037 (2)	0.0137 (18)	0.003 (2)	0.0057 (14)	0.0044 (19)
C20	0.037 (2)	0.034 (3)	0.012 (2)	0.001 (2)	0.0076 (19)	0.002 (2)
C21	0.042 (3)	0.029 (3)	0.019 (2)	0.000 (2)	0.002 (2)	0.001 (2)
C22	0.039 (2)	0.029 (2)	0.013 (2)	-0.002 (2)	0.0067 (17)	0.004 (2)
C23	0.048 (3)	0.036 (3)	0.013 (2)	-0.002 (2)	0.003 (2)	-0.001 (2)
C24	0.043 (3)	0.029 (3)	0.022 (3)	-0.002 (2)	0.006 (2)	0.005 (2)
C30	0.039 (3)	0.037 (3)	0.016 (2)	0.003 (2)	0.0086 (19)	0.006 (2)
C31	0.056 (3)	0.056 (3)	0.024 (3)	0.013 (3)	0.001 (2)	-0.003 (2)
C32	0.064 (3)	0.071 (4)	0.029 (3)	0.018 (3)	0.000 (2)	0.007 (3)
C33	0.056 (3)	0.046 (3)	0.035 (3)	0.012 (3)	0.011 (2)	0.013 (3)
C34	0.062 (3)	0.040 (3)	0.032 (3)	0.003 (3)	0.018 (3)	-0.003 (2)
N35	0.048 (2)	0.036 (2)	0.023 (2)	0.006 (2)	0.0093 (18)	0.000 (2)
C35	0.048 (2)	0.036 (2)	0.023 (2)	0.006 (2)	0.0093 (18)	0.000 (2)
C40	0.039 (2)	0.031 (3)	0.016 (2)	-0.005 (2)	0.0059 (17)	0.003 (2)
C41	0.053 (3)	0.045 (3)	0.016 (3)	0.005 (2)	-0.002 (2)	-0.010 (2)

C42	0.055 (3)	0.041 (3)	0.022 (3)	0.012 (2)	0.001 (2)	-0.007 (2)
C43	0.038 (2)	0.029 (3)	0.024 (2)	-0.005 (2)	0.0013 (18)	0.004 (2)
C44	0.056 (3)	0.040 (3)	0.036 (3)	0.006 (3)	-0.008 (2)	-0.011 (2)
C45	0.052 (3)	0.034 (3)	0.037 (3)	0.010 (2)	-0.004 (2)	-0.009 (2)
S40	0.0459 (6)	0.0359 (6)	0.0222 (6)	-0.0007 (6)	-0.0005 (4)	-0.0047 (6)
C46	0.049 (3)	0.053 (3)	0.024 (3)	0.009 (2)	-0.003 (2)	-0.003 (2)

*Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )*

C10—N15	1.342 (5)	C31—C32	1.367 (6)
C10—C11	1.374 (5)	C31—H31	0.9500
C10—C20	1.509 (5)	C32—C33	1.399 (6)
C11—C12	1.394 (6)	C32—H32	0.9500
C11—H11	0.9500	C33—C34	1.360 (6)
C12—C13	1.343 (6)	C33—H33	0.9500
C12—H12	0.9500	C34—N35	1.364 (5)
C13—C14	1.357 (5)	C34—H34	0.9500
C13—H13	0.9500	C40—C41	1.375 (5)
C14—N15	1.371 (5)	C40—C45	1.393 (5)
C14—H14	0.9500	C41—C42	1.392 (5)
N20—C20	1.324 (5)	C41—H41	0.9500
N20—C24	1.355 (5)	C42—C43	1.401 (6)
C20—C21	1.391 (5)	C42—H42	0.9500
C21—C22	1.402 (5)	C43—C44	1.381 (6)
C21—H21	0.9500	C43—S40	1.758 (4)
C22—C23	1.378 (5)	C44—C45	1.388 (6)
C22—C40	1.485 (5)	C44—H44	0.9500
C23—C24	1.398 (5)	C45—H45	0.9500
C23—H23	0.9500	S40—C46	1.787 (4)
C24—C30	1.472 (5)	C46—H46A	0.9800
C30—C31	1.380 (6)	C46—H46B	0.9800
C30—N35	1.382 (5)	C46—H46C	0.9800
N15—C10—C11	120.9 (4)	C30—C31—H31	118.9
N15—C10—C20	118.8 (4)	C31—C32—C33	119.5 (5)
C11—C10—C20	120.1 (4)	C31—C32—H32	120.3
C10—C11—C12	119.0 (4)	C33—C32—H32	120.3
C10—C11—H11	120.5	C34—C33—C32	117.3 (5)
C12—C11—H11	120.5	C34—C33—H33	121.3
C13—C12—C11	120.1 (4)	C32—C33—H33	121.3
C13—C12—H12	120.0	C33—C34—N35	123.8 (5)
C11—C12—H12	120.0	C33—C34—H34	118.1
C12—C13—C14	119.4 (4)	N35—C34—H34	118.1
C12—C13—H13	120.3	C34—N35—C30	118.9 (4)
C14—C13—H13	120.3	C41—C40—C45	116.1 (4)
C13—C14—N15	121.8 (4)	C41—C40—C22	122.4 (4)
C13—C14—H14	119.1	C45—C40—C22	121.5 (4)
N15—C14—H14	119.1	C40—C41—C42	123.4 (4)

C10—N15—C14	118.8 (4)	C40—C41—H41	118.3
C20—N20—C24	117.9 (3)	C42—C41—H41	118.3
N20—C20—C21	123.7 (4)	C41—C42—C43	119.5 (4)
N20—C20—C10	116.4 (3)	C41—C42—H42	120.2
C21—C20—C10	119.9 (4)	C43—C42—H42	120.2
C20—C21—C22	119.4 (4)	C44—C43—C42	117.7 (4)
C20—C21—H21	120.3	C44—C43—S40	118.3 (3)
C22—C21—H21	120.3	C42—C43—S40	123.9 (3)
C23—C22—C21	116.4 (3)	C43—C44—C45	121.4 (4)
C23—C22—C40	122.7 (4)	C43—C44—H44	119.3
C21—C22—C40	120.9 (4)	C45—C44—H44	119.3
C22—C23—C24	121.5 (4)	C44—C45—C40	121.8 (4)
C22—C23—H23	119.3	C44—C45—H45	119.1
C24—C23—H23	119.3	C40—C45—H45	119.1
N20—C24—C23	121.1 (4)	C43—S40—C46	103.4 (2)
N20—C24—C30	116.8 (4)	S40—C46—H46A	109.5
C23—C24—C30	122.1 (4)	S40—C46—H46B	109.5
C31—C30—N35	118.2 (4)	H46A—C46—H46B	109.5
C31—C30—C24	121.9 (4)	S40—C46—H46C	109.5
N35—C30—C24	119.8 (4)	H46A—C46—H46C	109.5
C32—C31—C30	122.2 (5)	H46B—C46—H46C	109.5
C32—C31—H31	118.9		