

3-(4-Chlorophenyl)-1-[(*E*)-1-(4-chlorophenyl)-2-(4-methylphenylsulfanyl)ethenyl]-4-(4-methylphenylsulfanyl)-1*H*-pyrazole

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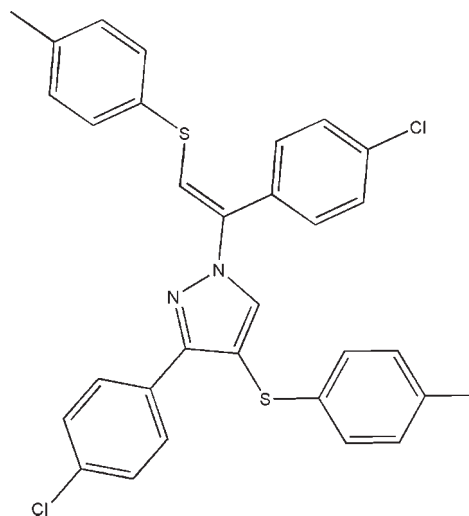
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.052; wR factor = 0.160; data-to-parameter ratio = 29.3.

In the title compound, $\text{C}_{31}\text{H}_{24}\text{Cl}_2\text{N}_2\text{S}_2$, the pyrazole ring adopts planar conformation with a maximum deviation of 0.002 (2) Å. The chlorophenyl rings are twisted out of the plane of the pyrazole ring by 75.1 (1) and 39.5 (1)°. The crystal packing is controlled by weak intermolecular $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For the pharmacological and medicinal properties of pyrazole derivatives, see: Baraldi *et al.* (1998); Bruno *et al.* (1990); Cottineau *et al.* (2002); Londershausen (1996); Chen & Li (1998); Mishra *et al.* (1998); Smith *et al.* (2001). For sp^2 hybridization, see: Beddoes *et al.*, 1986). For bond-length data, see: Jin *et al.* (2004).



Experimental

Crystal data

$\text{C}_{31}\text{H}_{24}\text{Cl}_2\text{N}_2\text{S}_2$
 $M_r = 559.54$
Monoclinic, $P2_1/n$
 $a = 9.7515$ (2) Å
 $b = 10.2097$ (3) Å
 $c = 27.6705$ (6) Å
 $\beta = 96.402$ (1)°

$V = 2737.69$ (11) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.41$ mm⁻¹
 $T = 293$ K
0.30 × 0.25 × 0.20 mm

Data collection

Bruker Kappa APEXII
diffractometer
Absorption correction: multi-scan
(SADABS; Sheldrick, 2001)
 $T_{\min} = 0.883$, $T_{\max} = 0.921$

38656 measured reflections
9852 independent reflections
5773 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.160$
 $S = 1.03$
9852 reflections

336 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.54$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.33$ e Å⁻³

Table 1

C—H $\cdots\pi$ interactions (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C33}-\text{H33C}\cdots\text{Cg3}^i$	0.96	2.90	3.851 (3)	171
$\text{C9}-\text{H9}\cdots\text{Cg5}^{ii}$	0.93	3.03	3.839 (2)	147

Symmetry codes: (i) $-x + 2, -y, -z$; (ii) $-x + 2, -y + 1, -z$. Cg3 and Cg5 are the centroids of the C14–C19 and C26–C31 rings, respectively.

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, (1997)); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5084).

References

- Baraldi, P. G., Manfredini, S., Romagnoli, R., Stevanato, L., Zaid, A. N. & Manservigi, R. (1998). *Nucleosides Nucleotides*, **17**, 2165–2171.
- Beddoes, R. L., Dalton, L., Joule, T. A., Mills, O. S., Street, J. D. & Watt, C. I. F. (1986). *J. Chem. Soc. Perkin Trans. 2*, pp. 787–797.
- Bruker (2004). *APEX2* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruno, O., Bondavalli, F., Ranise, A., Schenone, P., Losasso, C., Cilenti, L., Matera, C. & Marmo, E. (1990). *Farmaco*, **45**, 147–66.
- Chen, H. S. & Li, Z. M. (1998). *Chem. J. Chin. Univ.* **19**, 572–576.
- Cottineau, B., Toto, P., Marot, C., Pipaud, A. & Chenault, J. (2002). *Bioorg. Med. Chem.* **12**, 2105–2108.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Jin, Z.-M., Li, L., Li, M.-C., Hu, M.-L. & Shen, L. (2004). *Acta Cryst.* **C60**, o642–o643.
- Londershausen, M. (1996). *Pestic. Sci.* **48**, 269–274.
- Mishra, P. D., Wahidullah, S. & Kamat, S. Y. (1998). *Indian J. Chem. Sect. B*, **37**, 199–200.
- Sheldrick, G. M. (2001). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Smith, S. R., Denhardt, G. & Terminelli, C. (2001). *Eur. J. Pharmacol.* **432**, 107–119.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supporting information

Acta Cryst. (2009). E65, o2881–o2882 [https://doi.org/10.1107/S1600536809043293]

3-(4-Chlorophenyl)-1-[(*E*)-1-(4-chlorophenyl)-2-(4-methylphenylsulfanyl)ethenyl]-4-(4-methylphenylsulfanyl)-1*H*-pyrazole

P. Ramesh, Ramaiyan Manikannan, S. Muthusubramanian, S. S. Sundaresan and M. N. Ponnuswamy

S1. Comment

Pyrazole derivatives possess significant antiarrhythmic and sedative (Bruno *et al.*, 1990), hypoglycemic (Cottineau *et al.*, 2002), antiviral (Baraldi *et al.*, 1998), and pesticidal (Londershausen *et al.*, 1996) properties. Some pyrazole derivatives are successfully tested for their antifungal (Chen & Li, 1998), antihistaminic (Mishra *et al.*, 1998) and anti-inflammatory (Smith *et al.*, 2001) activities.

The *ORTEP* plot of the molecule is shown in Fig. 1. The pyrazole ring adopts a planar conformation. The sum of the bond angles at N2 of the pyrazole ring (360.0°) is in accordance with *sp*² hybridization (Beddoes *et al.*, 1986). The C—N bond lengths in the pyrazole ring are 1.352 (2) and 1.328 (2) Å, which are shorter than single bond length of 1.443 Å, but longer than a double bond length of 1.269 Å (Jin *et al.*, 2004), indicating electron delocalization. The chlorophenyl rings are twisted from the pyrazole ring at angles of $75.1 (1)^\circ$ and $39.5 (1)^\circ$, respectively. The crystal packing is stabilized by weak C—H $\cdots\pi$ type of intermolecular interactions in addition to van der Waals forces.

S2. Experimental

To a mixture of 1-(4-Chlorophenyl)-2-[(4-methylphenyl)sulfanyl]-1-ethanone *N*-(*Z*)-1-(4-chlorophenyl)-2-[(4-methylphenyl)sulfanyl]ethylidenehydrazone (0.003 mole) and 3 ml of dimethyl formamide kept in ice bath at 0°C , phosphorous oxychloride (0.024 mole) was added dropwise for 5–10 minutes. The reaction mixture was then irradiated under microwaves for 30 sec. The process of the reaction was monitored by TLC. After completion of the reaction, the reaction mixture was poured into crushed ice and extracted with dichloromethane. The organic layer was dried over anhydrous sodium sulfate. The different compounds present in the mixture were separated by column chromatography using petroleum ether and ethyl acetate mixture as eluent. This isolated compound was recrystallized in dichloromethane to obtain 3-(4-Chlorophenyl)-1-(*E*)-1-(4-chlorophenyl)-2-[(4-methylphenyl)sulfanyl]ethenyl-4-[(4-methylphenyl)sulfanyl]-1*H*-pyrazole.

S3. Refinement

All H atoms were positioned geometrically (C—H=0.93–0.96 Å) and allowed to ride on their parent atoms, with $1.5U_{\text{eq}}(\text{C})$ for methyl H and $1.2U_{\text{eq}}(\text{C})$ for other H atoms. The methyl groups were allowed to rotate but not to tip.

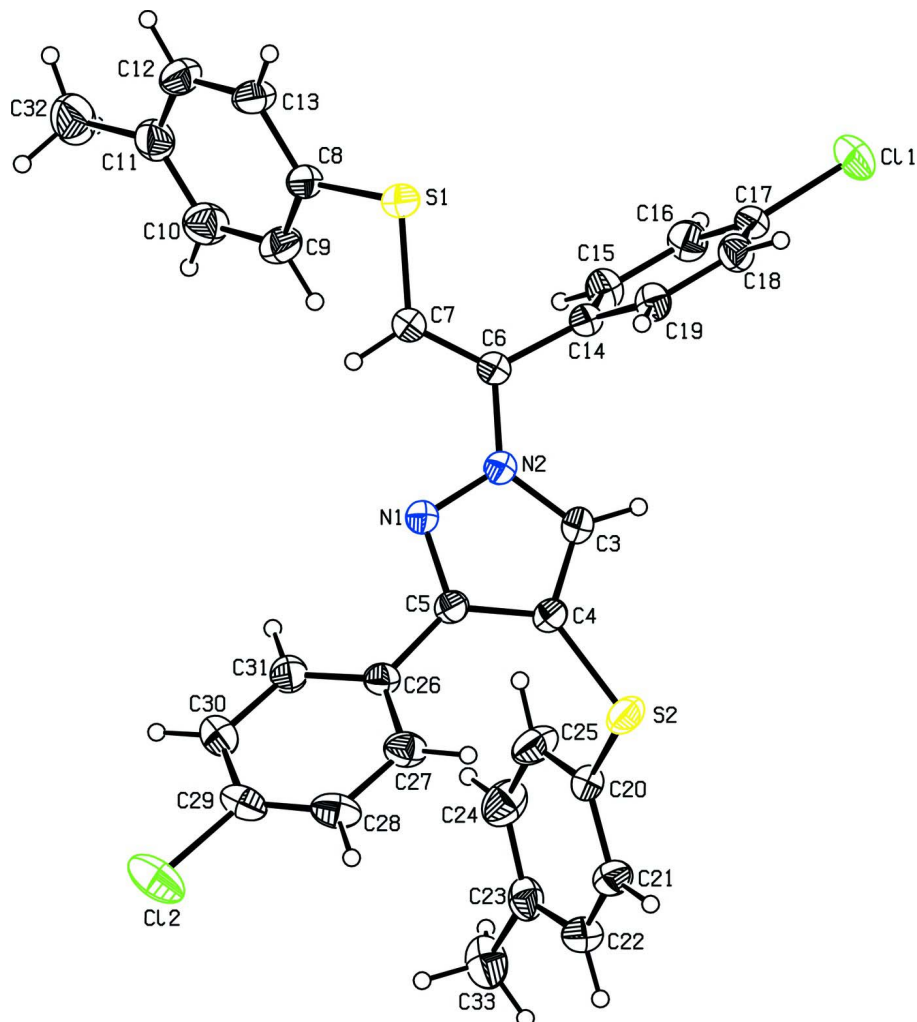


Figure 1

Perspective view of the molecule showing the displacement ellipsoids are drawn at 50% probability level. H atoms have been omitted for clarity.

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Crystal data

$C_{31}H_{24}Cl_2N_2S_2$

$M_r = 559.54$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 9.7515\ (2)\ \text{\AA}$

$b = 10.2097\ (3)\ \text{\AA}$

$c = 27.6705\ (6)\ \text{\AA}$

$\beta = 96.402\ (1)^\circ$

$V = 2737.69\ (11)\ \text{\AA}^3$

$Z = 4$

$F(000) = 1160$

$D_x = 1.358\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 2546 reflections

$\theta = 1.5\text{--}32.5^\circ$

$\mu = 0.41\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Block, colourless

$0.30 \times 0.25 \times 0.20\ \text{mm}$

Data collection

Bruker Kappa APEXII
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω and φ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 2001)
 $T_{\min} = 0.883$, $T_{\max} = 0.921$

38656 measured reflections
9852 independent reflections
5773 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
 $\theta_{\max} = 32.5^\circ$, $\theta_{\min} = 1.5^\circ$
 $h = -14 \rightarrow 14$
 $k = -15 \rightarrow 15$
 $l = -41 \rightarrow 39$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.160$
 $S = 1.03$
9852 reflections
336 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
 $w = 1/[\sigma^2(F_o^2) + (0.0723P)^2 + 0.5516P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.003$
 $\Delta\rho_{\max} = 0.54 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.33 \text{ e } \text{\AA}^{-3}$

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 $F^2 > \sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.08583 (6)	-0.11974 (5)	0.63584 (2)	0.05549 (15)
S2	0.25311 (5)	0.35982 (5)	0.42760 (2)	0.05549 (15)
Cl1	-0.13889 (7)	0.45353 (6)	0.68961 (2)	0.07471 (19)
Cl2	0.48797 (8)	-0.21313 (9)	0.30187 (2)	0.0905 (2)
N1	0.23543 (16)	0.02119 (14)	0.49711 (5)	0.0415 (3)
N2	0.18350 (16)	0.10815 (14)	0.52737 (5)	0.0421 (3)
C3	0.1835 (2)	0.23137 (18)	0.50948 (7)	0.0472 (4)
H3	0.1531	0.3062	0.5242	0.057*
C4	0.23608 (19)	0.22595 (18)	0.46595 (7)	0.0446 (4)
C5	0.26756 (18)	0.09181 (17)	0.45965 (6)	0.0398 (4)
C6	0.13934 (19)	0.06576 (17)	0.57196 (6)	0.0410 (4)
C7	0.1621 (2)	-0.05738 (19)	0.58678 (7)	0.0485 (4)
H7	0.2196	-0.1107	0.5707	0.058*
C8	0.2206 (2)	-0.21076 (18)	0.66790 (6)	0.0461 (4)
C9	0.3584 (2)	-0.1814 (2)	0.66695 (8)	0.0583 (5)
H9	0.3845	-0.1116	0.6484	0.070*

C10	0.4568 (3)	-0.2562 (2)	0.69371 (8)	0.0653 (6)
H10	0.5496	-0.2370	0.6924	0.078*
C11	0.4224 (3)	-0.3585 (2)	0.72236 (7)	0.0606 (5)
C12	0.2849 (3)	-0.3848 (2)	0.72369 (8)	0.0614 (6)
H12	0.2592	-0.4524	0.7433	0.074*
C13	0.1842 (2)	-0.3134 (2)	0.69661 (7)	0.0566 (5)
H13	0.0917	-0.3340	0.6976	0.068*
C14	0.07032 (18)	0.16654 (17)	0.59941 (6)	0.0389 (3)
C15	0.1380 (2)	0.2157 (2)	0.64224 (7)	0.0495 (4)
H15	0.2271	0.1877	0.6527	0.059*
C16	0.0749 (2)	0.3054 (2)	0.66948 (7)	0.0523 (5)
H16	0.1212	0.3388	0.6980	0.063*
C17	-0.0571 (2)	0.34490 (18)	0.65410 (6)	0.0459 (4)
C18	-0.1264 (2)	0.29888 (19)	0.61145 (7)	0.0478 (4)
H18	-0.2155	0.3273	0.6012	0.057*
C19	-0.06171 (19)	0.20999 (18)	0.58410 (6)	0.0443 (4)
H19	-0.1074	0.1789	0.5551	0.053*
C20	0.4342 (2)	0.38614 (17)	0.43419 (7)	0.0441 (4)
C21	0.4884 (2)	0.4601 (2)	0.39927 (7)	0.0521 (5)
H21	0.4301	0.4967	0.3740	0.063*
C22	0.6289 (2)	0.4800 (2)	0.40176 (8)	0.0549 (5)
H22	0.6639	0.5305	0.3780	0.066*
C23	0.7187 (2)	0.42680 (19)	0.43848 (8)	0.0524 (5)
C24	0.6620 (3)	0.3574 (2)	0.47388 (9)	0.0679 (6)
H24	0.7199	0.3227	0.4997	0.081*
C25	0.5222 (2)	0.3377 (2)	0.47219 (8)	0.0649 (6)
H25	0.4869	0.2914	0.4970	0.078*
C26	0.32826 (18)	0.02448 (17)	0.42028 (6)	0.0400 (4)
C27	0.2903 (2)	0.0553 (2)	0.37175 (6)	0.0510 (5)
H27	0.2310	0.1252	0.3636	0.061*
C28	0.3405 (2)	-0.0176 (2)	0.33562 (7)	0.0576 (5)
H28	0.3147	0.0028	0.3031	0.069*
C29	0.4282 (2)	-0.1200 (2)	0.34765 (7)	0.0553 (5)
C30	0.4714 (2)	-0.1501 (2)	0.39539 (8)	0.0567 (5)
H30	0.5336	-0.2178	0.4032	0.068*
C31	0.4201 (2)	-0.07719 (19)	0.43141 (7)	0.0490 (4)
H31	0.4480	-0.0969	0.4638	0.059*
C32	0.5335 (3)	-0.4378 (3)	0.75125 (10)	0.0887 (9)
H32A	0.5850	-0.4854	0.7294	0.133*
H32B	0.4918	-0.4983	0.7718	0.133*
H32C	0.5942	-0.3802	0.7709	0.133*
C33	0.8720 (2)	0.4392 (3)	0.43899 (11)	0.0728 (7)
H33A	0.9079	0.3617	0.4253	0.109*
H33B	0.9136	0.4497	0.4719	0.109*
H33C	0.8929	0.5140	0.4201	0.109*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0571 (3)	0.0534 (3)	0.0597 (3)	0.0077 (2)	0.0230 (2)	0.0147 (2)
S2	0.0492 (3)	0.0490 (3)	0.0690 (3)	0.0058 (2)	0.0096 (2)	0.0254 (2)
C11	0.0911 (5)	0.0722 (4)	0.0659 (3)	0.0193 (3)	0.0308 (3)	-0.0139 (3)
C12	0.0853 (5)	0.1220 (6)	0.0684 (4)	-0.0011 (4)	0.0276 (3)	-0.0391 (4)
N1	0.0492 (8)	0.0357 (7)	0.0409 (7)	0.0026 (6)	0.0111 (6)	0.0010 (6)
N2	0.0519 (9)	0.0344 (7)	0.0416 (7)	0.0020 (6)	0.0125 (6)	0.0017 (6)
C3	0.0532 (11)	0.0349 (9)	0.0553 (10)	0.0047 (8)	0.0140 (8)	0.0039 (7)
C4	0.0460 (10)	0.0392 (9)	0.0498 (9)	0.0025 (8)	0.0112 (8)	0.0094 (7)
C5	0.0398 (9)	0.0391 (9)	0.0411 (8)	0.0002 (7)	0.0066 (7)	0.0051 (7)
C6	0.0451 (9)	0.0391 (9)	0.0402 (8)	0.0021 (7)	0.0107 (7)	0.0008 (7)
C7	0.0563 (11)	0.0428 (10)	0.0494 (9)	0.0078 (8)	0.0193 (8)	0.0051 (8)
C8	0.0559 (11)	0.0412 (9)	0.0433 (9)	-0.0014 (8)	0.0143 (8)	0.0009 (7)
C9	0.0607 (13)	0.0519 (12)	0.0640 (12)	-0.0077 (10)	0.0141 (10)	0.0108 (9)
C10	0.0556 (13)	0.0749 (15)	0.0660 (13)	-0.0008 (11)	0.0096 (10)	0.0038 (11)
C11	0.0736 (15)	0.0634 (14)	0.0451 (10)	0.0151 (11)	0.0085 (9)	-0.0005 (9)
C12	0.0828 (16)	0.0545 (12)	0.0484 (10)	0.0020 (11)	0.0147 (10)	0.0124 (9)
C13	0.0614 (13)	0.0586 (12)	0.0522 (11)	-0.0069 (10)	0.0174 (9)	0.0101 (9)
C14	0.0424 (9)	0.0360 (8)	0.0393 (8)	-0.0002 (7)	0.0090 (7)	0.0014 (6)
C15	0.0450 (10)	0.0531 (11)	0.0491 (10)	0.0040 (8)	-0.0004 (8)	-0.0048 (8)
C16	0.0611 (12)	0.0535 (11)	0.0415 (9)	-0.0004 (9)	0.0014 (8)	-0.0081 (8)
C17	0.0562 (11)	0.0402 (9)	0.0447 (9)	0.0030 (8)	0.0198 (8)	-0.0002 (7)
C18	0.0411 (10)	0.0457 (10)	0.0574 (10)	0.0025 (8)	0.0090 (8)	-0.0002 (8)
C19	0.0433 (10)	0.0427 (9)	0.0465 (9)	-0.0016 (8)	0.0029 (7)	-0.0048 (7)
C20	0.0502 (10)	0.0361 (9)	0.0469 (9)	0.0048 (7)	0.0091 (8)	0.0047 (7)
C21	0.0570 (12)	0.0542 (11)	0.0459 (9)	0.0034 (9)	0.0086 (8)	0.0114 (8)
C22	0.0569 (12)	0.0552 (12)	0.0549 (11)	-0.0034 (10)	0.0162 (9)	0.0050 (9)
C23	0.0513 (11)	0.0397 (10)	0.0659 (12)	-0.0011 (8)	0.0060 (9)	-0.0104 (8)
C24	0.0593 (13)	0.0685 (15)	0.0720 (14)	0.0010 (11)	-0.0097 (11)	0.0181 (12)
C25	0.0635 (14)	0.0664 (14)	0.0637 (13)	-0.0024 (11)	0.0024 (10)	0.0300 (11)
C26	0.0375 (9)	0.0437 (9)	0.0396 (8)	-0.0027 (7)	0.0075 (6)	0.0031 (7)
C27	0.0421 (10)	0.0679 (13)	0.0428 (9)	0.0010 (9)	0.0043 (7)	0.0072 (8)
C28	0.0469 (11)	0.0878 (16)	0.0382 (9)	-0.0089 (11)	0.0059 (8)	0.0003 (9)
C29	0.0462 (11)	0.0725 (14)	0.0498 (10)	-0.0123 (10)	0.0173 (8)	-0.0154 (9)
C30	0.0568 (12)	0.0572 (12)	0.0578 (11)	0.0074 (10)	0.0141 (9)	-0.0059 (9)
C31	0.0563 (11)	0.0492 (10)	0.0416 (9)	0.0053 (9)	0.0060 (8)	0.0022 (7)
C32	0.100 (2)	0.100 (2)	0.0659 (15)	0.0409 (18)	0.0087 (14)	0.0101 (14)
C33	0.0553 (13)	0.0606 (14)	0.1015 (19)	-0.0034 (11)	0.0041 (12)	-0.0168 (13)

Geometric parameters (Å, °)

S1—C7	1.7398 (19)	C16—H16	0.9300
S1—C8	1.766 (2)	C17—C18	1.376 (3)
S2—C4	1.7496 (18)	C18—C19	1.378 (3)
S2—C20	1.776 (2)	C18—H18	0.9300
C11—C17	1.7340 (18)	C19—H19	0.9300

C12—C29	1.736 (2)	C20—C25	1.373 (3)
N1—C5	1.328 (2)	C20—C21	1.377 (3)
N1—N2	1.3565 (19)	C21—C22	1.379 (3)
N2—C3	1.352 (2)	C21—H21	0.9300
N2—C6	1.419 (2)	C22—C23	1.376 (3)
C3—C4	1.362 (3)	C22—H22	0.9300
C3—H3	0.9300	C23—C24	1.374 (3)
C4—C5	1.418 (3)	C23—C33	1.500 (3)
C5—C26	1.467 (2)	C24—C25	1.373 (3)
C6—C7	1.333 (3)	C24—H24	0.9300
C6—C14	1.484 (2)	C25—H25	0.9300
C7—H7	0.9300	C26—C31	1.383 (3)
C8—C9	1.380 (3)	C26—C27	1.389 (2)
C8—C13	1.385 (3)	C27—C28	1.379 (3)
C9—C10	1.377 (3)	C27—H27	0.9300
C9—H9	0.9300	C28—C29	1.369 (3)
C10—C11	1.376 (3)	C28—H28	0.9300
C10—H10	0.9300	C29—C30	1.375 (3)
C11—C12	1.372 (4)	C30—C31	1.381 (3)
C11—C32	1.508 (3)	C30—H30	0.9300
C12—C13	1.376 (3)	C31—H31	0.9300
C12—H12	0.9300	C32—H32A	0.9600
C13—H13	0.9300	C32—H32B	0.9600
C14—C19	1.383 (3)	C32—H32C	0.9600
C14—C15	1.385 (2)	C33—H33A	0.9600
C15—C16	1.374 (3)	C33—H33B	0.9600
C15—H15	0.9300	C33—H33C	0.9600
C16—C17	1.371 (3)		
C7—S1—C8	103.04 (9)	C19—C18—H18	120.5
C4—S2—C20	102.51 (9)	C18—C19—C14	120.68 (17)
C5—N1—N2	105.26 (13)	C18—C19—H19	119.7
C3—N2—N1	111.58 (14)	C14—C19—H19	119.7
C3—N2—C6	127.92 (15)	C25—C20—C21	118.63 (19)
N1—N2—C6	120.49 (14)	C25—C20—S2	123.61 (16)
N2—C3—C4	107.54 (16)	C21—C20—S2	117.76 (14)
N2—C3—H3	126.2	C20—C21—C22	120.14 (18)
C4—C3—H3	126.2	C20—C21—H21	119.9
C3—C4—C5	104.82 (15)	C22—C21—H21	119.9
C3—C4—S2	125.11 (15)	C23—C22—C21	121.67 (19)
C5—C4—S2	130.04 (14)	C23—C22—H22	119.2
N1—C5—C4	110.80 (15)	C21—C22—H22	119.2
N1—C5—C26	118.34 (15)	C24—C23—C22	117.2 (2)
C4—C5—C26	130.85 (15)	C24—C23—C33	121.1 (2)
C7—C6—N2	119.92 (16)	C22—C23—C33	121.7 (2)
C7—C6—C14	124.46 (16)	C25—C24—C23	121.9 (2)
N2—C6—C14	115.62 (14)	C25—C24—H24	119.1
C6—C7—S1	120.93 (15)	C23—C24—H24	119.1

C6—C7—H7	119.5	C20—C25—C24	120.4 (2)
S1—C7—H7	119.5	C20—C25—H25	119.8
C9—C8—C13	119.17 (19)	C24—C25—H25	119.8
C9—C8—S1	123.22 (15)	C31—C26—C27	118.79 (17)
C13—C8—S1	117.56 (16)	C31—C26—C5	119.41 (15)
C10—C9—C8	119.4 (2)	C27—C26—C5	121.71 (17)
C10—C9—H9	120.3	C28—C27—C26	120.04 (19)
C8—C9—H9	120.3	C28—C27—H27	120.0
C11—C10—C9	122.1 (2)	C26—C27—H27	120.0
C11—C10—H10	119.0	C29—C28—C27	119.92 (18)
C9—C10—H10	119.0	C29—C28—H28	120.0
C12—C11—C10	117.8 (2)	C27—C28—H28	120.0
C12—C11—C32	121.8 (2)	C28—C29—C30	121.33 (19)
C10—C11—C32	120.4 (2)	C28—C29—C12	119.50 (16)
C11—C12—C13	121.5 (2)	C30—C29—C12	119.16 (18)
C11—C12—H12	119.3	C29—C30—C31	118.5 (2)
C13—C12—H12	119.3	C29—C30—H30	120.8
C12—C13—C8	120.0 (2)	C31—C30—H30	120.8
C12—C13—H13	120.0	C30—C31—C26	121.40 (18)
C8—C13—H13	120.0	C30—C31—H31	119.3
C19—C14—C15	118.97 (16)	C26—C31—H31	119.3
C19—C14—C6	121.71 (15)	C11—C32—H32A	109.5
C15—C14—C6	119.30 (16)	C11—C32—H32B	109.5
C16—C15—C14	120.76 (18)	H32A—C32—H32B	109.5
C16—C15—H15	119.6	C11—C32—H32C	109.5
C14—C15—H15	119.6	H32A—C32—H32C	109.5
C17—C16—C15	119.21 (17)	H32B—C32—H32C	109.5
C17—C16—H16	120.4	C23—C33—H33A	109.5
C15—C16—H16	120.4	C23—C33—H33B	109.5
C16—C17—C18	121.34 (17)	H33A—C33—H33B	109.5
C16—C17—C11	119.20 (15)	C23—C33—H33C	109.5
C18—C17—C11	119.46 (16)	H33A—C33—H33C	109.5
C17—C18—C19	119.02 (18)	H33B—C33—H33C	109.5
C17—C18—H18	120.5		
C5—N1—N2—C3	-0.3 (2)	C19—C14—C15—C16	0.6 (3)
C5—N1—N2—C6	-179.36 (16)	C6—C14—C15—C16	-177.49 (18)
N1—N2—C3—C4	0.4 (2)	C14—C15—C16—C17	0.8 (3)
C6—N2—C3—C4	179.37 (18)	C15—C16—C17—C18	-1.5 (3)
N2—C3—C4—C5	-0.3 (2)	C15—C16—C17—C11	177.69 (16)
N2—C3—C4—S2	178.16 (14)	C16—C17—C18—C19	0.8 (3)
C20—S2—C4—C3	108.01 (18)	C11—C17—C18—C19	-178.36 (15)
C20—S2—C4—C5	-73.92 (19)	C17—C18—C19—C14	0.6 (3)
N2—N1—C5—C4	0.10 (19)	C15—C14—C19—C18	-1.3 (3)
N2—N1—C5—C26	179.47 (15)	C6—C14—C19—C18	176.76 (17)
C3—C4—C5—N1	0.1 (2)	C4—S2—C20—C25	-18.3 (2)
S2—C4—C5—N1	-178.23 (15)	C4—S2—C20—C21	162.26 (16)
C3—C4—C5—C26	-179.13 (18)	C25—C20—C21—C22	2.6 (3)

S2—C4—C5—C26	2.5 (3)	S2—C20—C21—C22	-177.86 (16)
C3—N2—C6—C7	-171.8 (2)	C20—C21—C22—C23	0.4 (3)
N1—N2—C6—C7	7.1 (3)	C21—C22—C23—C24	-2.7 (3)
C3—N2—C6—C14	7.6 (3)	C21—C22—C23—C33	174.9 (2)
N1—N2—C6—C14	-173.51 (15)	C22—C23—C24—C25	2.0 (4)
N2—C6—C7—S1	-169.62 (14)	C33—C23—C24—C25	-175.6 (2)
C14—C6—C7—S1	11.0 (3)	C21—C20—C25—C24	-3.2 (4)
C8—S1—C7—C6	-138.30 (17)	S2—C20—C25—C24	177.26 (19)
C7—S1—C8—C9	27.9 (2)	C23—C24—C25—C20	0.9 (4)
C7—S1—C8—C13	-154.47 (16)	N1—C5—C26—C31	-37.2 (2)
C13—C8—C9—C10	1.3 (3)	C4—C5—C26—C31	142.1 (2)
S1—C8—C9—C10	178.90 (17)	N1—C5—C26—C27	139.28 (18)
C8—C9—C10—C11	-1.3 (4)	C4—C5—C26—C27	-41.5 (3)
C9—C10—C11—C12	0.0 (3)	C31—C26—C27—C28	2.1 (3)
C9—C10—C11—C32	-179.8 (2)	C5—C26—C27—C28	-174.36 (18)
C10—C11—C12—C13	1.3 (3)	C26—C27—C28—C29	-0.3 (3)
C32—C11—C12—C13	-178.9 (2)	C27—C28—C29—C30	-2.0 (3)
C11—C12—C13—C8	-1.3 (3)	C27—C28—C29—C12	178.72 (16)
C9—C8—C13—C12	-0.1 (3)	C28—C29—C30—C31	2.3 (3)
S1—C8—C13—C12	-177.79 (17)	C12—C29—C30—C31	-178.35 (16)
C7—C6—C14—C19	-108.0 (2)	C29—C30—C31—C26	-0.5 (3)
N2—C6—C14—C19	72.6 (2)	C27—C26—C31—C30	-1.7 (3)
C7—C6—C14—C15	70.0 (3)	C5—C26—C31—C30	174.82 (18)
N2—C6—C14—C15	-109.32 (19)		

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>
C33—H33C...Cg3 ⁱ	0.96	2.90	3.851 (3)	171
C9—H9...Cg5 ⁱⁱ	0.93	3.03	3.839 (2)	147

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