

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

3-(5-Phenyl-4-phenylsulfonyl-1-*p*-tolyl-1*H*-pyrazol-3-yl)-1,2-dihydroquinoxaline

 Hatem A. Abdel-Aziz,^a Ahmed Bari^a and Seik Weng Ng^{b*}
^aDepartment of Pharmaceutical Chemistry, College of Pharmacy, King Saud University, Riyadh 11451, Saudi Arabia, and ^bDepartment of Chemistry, University of Malaya, 50603 Kuala Lumpur, Malaysia

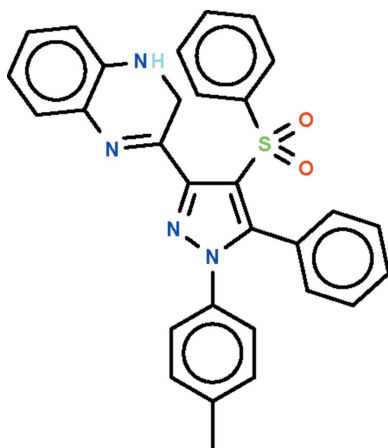
Correspondence e-mail: seikweng@um.edu.my

Received 14 February 2011; accepted 17 February 2011

 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; disorder in main residue; R factor = 0.062; wR factor = 0.158; data-to-parameter ratio = 13.1.

In the crystal structure of the title compound, $\text{C}_{30}\text{H}_{24}\text{N}_4\text{O}_2\text{S}$, the dihydroquinoxaline fused-ring system is disordered over three orientations in a 0.358 (2):0.318 (3):0.324 (3) ratio; the mean planes of the non-H atoms of the disorder components are aligned at 4.0 (3), 11.8 (4) and 41.7 (2)° with respect to the pyrazole ring. The rings of the phenyl and tolyl substituents are aligned at 64.0 (1) and 43.7 (1)° with respect to the pyrazole ring. Weak intermolecular C—H...O hydrogen bonding links the molecules, forming supramolecular chains running along the a axis.

Related literature

 For background to the biological properties of aryl-substituted pyrazoles, see: Abdel-Aziz *et al.* (2010).


Experimental

Crystal data

 $\text{C}_{30}\text{H}_{24}\text{N}_4\text{O}_2\text{S}$
 $M_r = 504.59$
 Monoclinic, $P2_1/c$
 $a = 6.2829$ (2) Å
 $b = 24.5162$ (7) Å
 $c = 15.9180$ (5) Å
 $\beta = 91.925$ (3)°

 $V = 2450.51$ (13) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.17$ mm⁻¹
 $T = 100$ K
 $0.20 \times 0.02 \times 0.02$ mm

Data collection

 Agilent SuperNova Dual diffractometer with an Atlas detector
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2010)
 $T_{\min} = 0.967$, $T_{\max} = 0.997$

 16288 measured reflections
 5524 independent reflections
 3897 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.158$
 $S = 1.02$
 5524 reflections
 422 parameters

 139 restraints
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.38$ e Å⁻³
 $\Delta\rho_{\min} = -0.40$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C19}-\text{H19}\cdots\text{O1}^i$	0.95	2.57	3.430 (3)	150

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

Data collection: *CrysAlis PRO* (Agilent, 2010); 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: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

We thank King Saud University and the University of Malaya for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5163).

References

- Abdel-Aziz, H. A., El-Zahabi, H. S. A. & Dawood, K. M. (2010). *Eur. J. Med. Chem.* **45**, 2427–2432.
 Agilent (2010). *CrysAlis PRO*. Agilent Technologies, Yarnton, England.
 Barbour, L. J. (2001). *J. Supramol. Chem.* **1**, 189–191.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

supporting information

Acta Cryst. (2011). E67, o695 [doi:10.1107/S1600536811005940]

3-(5-Phenyl-4-phenylsulfonyl-1-*p*-tolyl-1*H*-pyrazol-3-yl)-1,2-dihydroquinoxaline

Hatem A. Abdel-Aziz, Ahmed Bari and Seik Weng Ng

S1. Comment

We have reported the antitumor activity of aryl-pyrazoles against CaCo-2 and HEP-2 cell lines (Abdel-Aziz *et al.*, 2010). These compounds were synthesized by a cycloaddition under microwave conditions. The study is now extended to the synthesis of a pyrazole having a dihydroquinoxaline substituent (Scheme I). The dihydroquinoxalinyl substituent of the pyrazolyl ring of C₃₀H₂₄N₄O₂S adopts three orientations. The orientations refined to a 0.358 (2): 0.318 (3): 0.324 (3) ratio. The mean planes of the atoms passing through the non-hydrogen atoms of the disorder components are aligned at 4.0 (3), 11.8 (4) and 41.7 (2)° with respect to the five-membered ring (Fig. 1).

S2. Experimental

The acetyl portion of 1-[5-phenyl-4-(phenylsulfonyl)-1-*p*-tolyl-1*H*-pyrazol-3-yl]ethane was first converted to a bromo-acetyl unit by bromination at 363–373 K to yield 2-bromo-1-[5-phenyl-4-(phenylsulfonyl)-1-*p*-tolyl-1*H*-pyrazol-3-yl]ethanone. The compound (10 mmol) was heated with *o*-phenylenediamine (10 mmol) in ethanol to yield the title compound. Crystals were obtained upon recrystallization from ethanol.

S3. Refinement

Carbon-bound H-atoms were placed in calculated positions [C—H 0.95 to 0.98 Å, $U_{\text{iso}}(\text{H})$ 1.2 to 1.5 $U_{\text{eq}}(\text{C})$] and were included in the refinement in the riding model approximation.

The dihydroquinoxalinyl fused-ring is disordered over three orientations in a 0.358 (2): 0.318 (3): 0.324 (3) ratio. The first two disorder components are close to each other, so that the temperature factors of the singly-primed atoms were set to those of the unprimed atoms. The anisotropic temperature factors of the atoms of the three disorder components were restrained to be nearly isotropic, with the restraint being much tighter for the C1/C1'/C1'' set of atoms. The aromatic rings were refined as rigid hexagons of 1.39 Å sides. The carbon–nitrogen_{tertiary} distances were restrained to 1.35±0.01 Å and the carbon–nitrogen_{secondary} to 1.45±0.01 Å. The carbon–carbon single-bond distances were restrained to 1.50±0.01 Å. The amino H-atoms were treated in the riding mode [N—H 0.88 Å, $U_{\text{iso}}(\text{H})$ 1.2 $U_{\text{eq}}(\text{N})$].

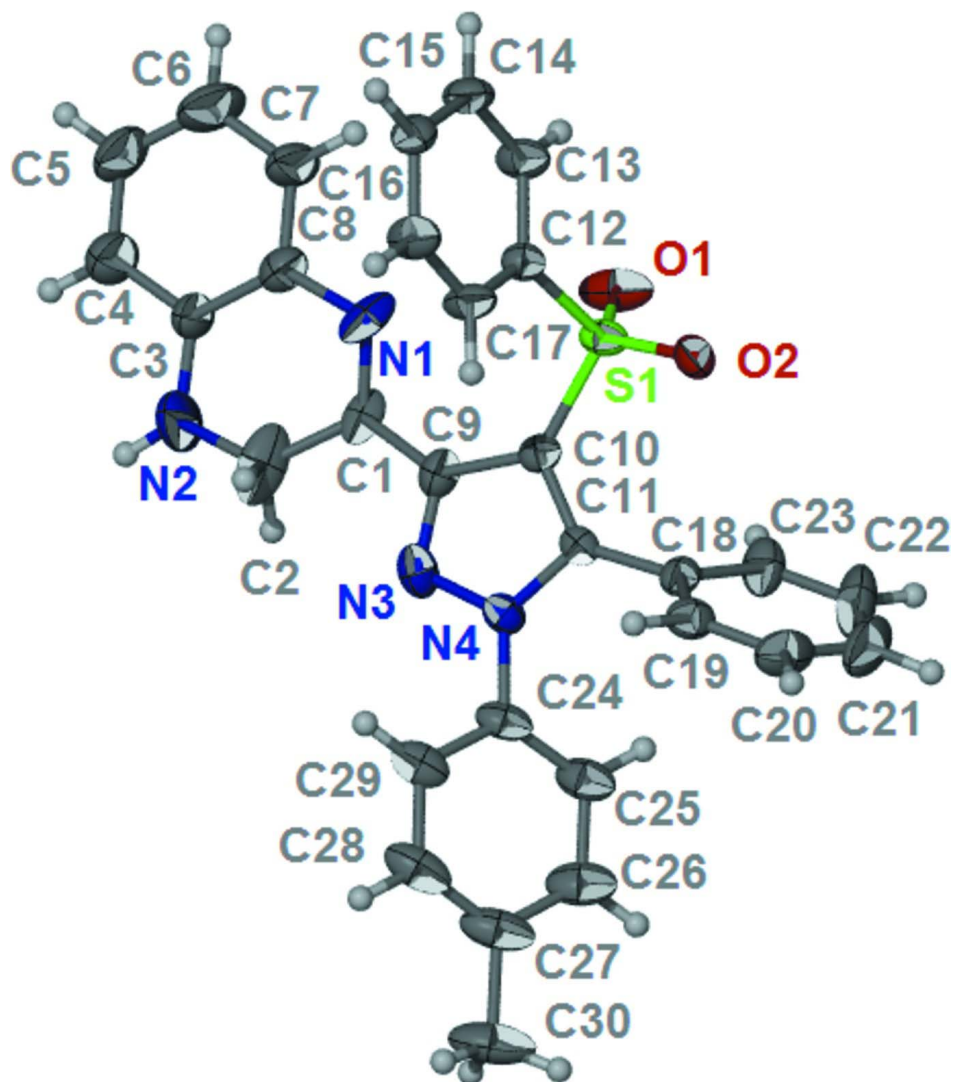


Figure 1

Thermal ellipsoid plot (Barbour, 2001) of $C_{30}H_{24}N_4O_2S$ at the 70% probability level; hydrogen atoms are drawn as spheres of arbitrary radius. The disorder is not shown.

3-(5-Phenyl-4-phenylsulfonyl-1-p-tolyl-1H-pyrazol-3-yl)-1,2-dihydroquinoxaline

Crystal data

$C_{30}H_{24}N_4O_2S$

$M_r = 504.59$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 6.2829 (2) \text{ \AA}$

$b = 24.5162 (7) \text{ \AA}$

$c = 15.9180 (5) \text{ \AA}$

$\beta = 91.925 (3)^\circ$

$V = 2450.51 (13) \text{ \AA}^3$

$Z = 4$

$F(000) = 1056$

$D_x = 1.368 \text{ Mg m}^{-3}$

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

Cell parameters from 4169 reflections

$\theta = 2.5\text{--}29.3^\circ$

$\mu = 0.17 \text{ mm}^{-1}$

$T = 100 \text{ K}$

Prism, yellow

$0.20 \times 0.02 \times 0.02 \text{ mm}$

Data collection

Agilent SuperNova Dual
 diffractometer with an Atlas detector
 Radiation source: SuperNova (Mo) X-ray
 Source
 Mirror monochromator
 Detector resolution: 10.4041 pixels mm⁻¹
 ω scans
 Absorption correction: multi-scan
 (CrysAlis PRO; Agilent, 2010)

$T_{\min} = 0.967$, $T_{\max} = 0.997$
 16288 measured reflections
 5524 independent reflections
 3897 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -8 \rightarrow 8$
 $k = -31 \rightarrow 31$
 $l = -20 \rightarrow 20$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.158$
 $S = 1.02$
 5524 reflections
 422 parameters
 139 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
 $w = 1/[\sigma^2(F_o^2) + (0.0634P)^2 + 1.9372P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.38 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.40 \text{ e } \text{\AA}^{-3}$

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)
S1	0.81495 (10)	0.59167 (2)	0.63024 (4)	0.02549 (18)	
O1	1.0308 (3)	0.59391 (8)	0.66188 (15)	0.0434 (6)	
O2	0.7677 (3)	0.62026 (7)	0.55309 (12)	0.0317 (5)	
N1	0.804 (2)	0.5182 (4)	0.8067 (8)	0.034 (3)	0.358 (2)
N2	0.6304 (13)	0.5044 (4)	0.9751 (5)	0.0518 (19)	0.358 (2)
H2	0.5832	0.5024	1.0264	0.062*	0.358 (2)
N1'	0.854 (3)	0.5300 (7)	0.8110 (12)	0.034 (3)	0.318 (3)
N2'	0.7437 (15)	0.5043 (5)	0.9645 (7)	0.0518 (19)	0.318 (3)
H2'	0.6985	0.4916	1.0125	0.062*	0.318 (3)
N1''	0.8972 (12)	0.5346 (4)	0.7959 (8)	0.018 (2)	0.324 (3)
N2''	0.6519 (13)	0.4503 (3)	0.8722 (6)	0.058 (3)	0.324 (3)
H2''	0.5796	0.4244	0.8967	0.070*	0.324 (3)
N3	0.4322 (4)	0.62374 (9)	0.81706 (14)	0.0300 (5)	
N4	0.3835 (3)	0.66333 (8)	0.76021 (13)	0.0224 (5)	
C1	0.696 (2)	0.5572 (4)	0.8435 (7)	0.027 (2)	0.358 (2)
C2	0.5687 (19)	0.5421 (6)	0.9157 (8)	0.049 (3)	0.358 (2)
H2A	0.5418	0.5764	0.9465	0.059*	0.358 (2)
H2B	0.4290	0.5298	0.8922	0.059*	0.358 (2)
C3	0.7815 (8)	0.4692 (2)	0.9407 (3)	0.0328 (14)	0.358 (2)
C4	0.8492 (9)	0.4250 (2)	0.9894 (3)	0.0443 (16)	0.358 (2)
H4A	0.7923	0.4194	1.0433	0.053*	0.358 (2)
C5	1.0001 (10)	0.3892 (2)	0.9594 (3)	0.0458 (17)	0.358 (2)
H5	1.0464	0.3590	0.9927	0.055*	0.358 (2)
C6	1.0834 (10)	0.3974 (2)	0.8806 (4)	0.054 (2)	0.358 (2)
H6	1.1865	0.3729	0.8600	0.065*	0.358 (2)

C7	1.0157 (10)	0.4416 (2)	0.8318 (3)	0.0432 (18)	0.358 (2)
H7	1.0726	0.4472	0.7780	0.052*	0.358 (2)
C8	0.8647 (9)	0.47745 (19)	0.8619 (3)	0.0314 (16)	0.358 (2)
C1'	0.659 (2)	0.5469 (5)	0.8348 (9)	0.027 (2)	0.318 (3)
C2'	0.634 (2)	0.5474 (6)	0.9271 (9)	0.049 (3)	0.318 (3)
H2'A	0.6885	0.5822	0.9506	0.059*	0.318 (3)
H2'B	0.4809	0.5446	0.9396	0.059*	0.318 (3)
C3'	0.9213 (9)	0.4799 (2)	0.9307 (4)	0.0328 (14)	0.318 (3)
C4'	1.0383 (11)	0.4432 (3)	0.9802 (3)	0.0443 (16)	0.318 (3)
H4'	0.9969	0.4356	1.0358	0.053*	0.318 (3)
C5'	1.2157 (10)	0.4176 (3)	0.9483 (4)	0.0458 (17)	0.318 (3)
H5'	1.2957	0.3925	0.9821	0.055*	0.318 (3)
C6'	1.2763 (11)	0.4288 (3)	0.8669 (4)	0.054 (2)	0.318 (3)
H6'	1.3976	0.4113	0.8451	0.065*	0.318 (3)
C7'	1.1594 (14)	0.4655 (4)	0.8174 (3)	0.0432 (18)	0.318 (3)
H7'	1.2007	0.4731	0.7617	0.052*	0.318 (3)
C8'	0.9819 (12)	0.4911 (3)	0.8493 (4)	0.0314 (16)	0.318 (3)
C1''	0.6918 (14)	0.5421 (4)	0.8112 (7)	0.030 (3)	0.324 (3)
C2''	0.5589 (15)	0.4998 (3)	0.8485 (6)	0.042 (2)	0.324 (3)
H2''A	0.4948	0.5157	0.8989	0.050*	0.324 (3)
H2''B	0.4406	0.4917	0.8078	0.050*	0.324 (3)
C3''	0.8676 (10)	0.4445 (3)	0.8547 (5)	0.053 (3)	0.324 (3)
C4''	0.9628 (12)	0.3945 (3)	0.8730 (6)	0.065 (4)	0.324 (3)
H4''	0.8816	0.3659	0.8962	0.078*	0.324 (3)
C5''	1.1768 (13)	0.3865 (3)	0.8573 (6)	0.075 (4)	0.324 (3)
H5''	1.2419	0.3523	0.8698	0.089*	0.324 (3)
C6''	1.2956 (10)	0.4284 (3)	0.8234 (6)	0.065 (4)	0.324 (3)
H6''	1.4419	0.4229	0.8126	0.078*	0.324 (3)
C7''	1.2003 (12)	0.4783 (3)	0.8051 (5)	0.051 (3)	0.324 (3)
H7''	1.2815	0.5070	0.7819	0.061*	0.324 (3)
C8''	0.9863 (12)	0.4864 (2)	0.8208 (5)	0.044 (3)	0.324 (3)
C9	0.5873 (4)	0.59504 (10)	0.78359 (17)	0.0275 (6)	
C10	0.6366 (4)	0.61620 (9)	0.70379 (16)	0.0213 (5)	
C11	0.5021 (4)	0.66014 (9)	0.69050 (15)	0.0193 (5)	
C12	0.7410 (4)	0.52265 (10)	0.61533 (16)	0.0222 (5)	
C13	0.9002 (4)	0.48606 (10)	0.59728 (18)	0.0306 (6)	
H13	1.0449	0.4974	0.5986	0.037*	
C14	0.8469 (4)	0.43271 (11)	0.57721 (18)	0.0313 (6)	
H14	0.9554	0.4075	0.5637	0.038*	
C15	0.6385 (4)	0.41603 (10)	0.57666 (17)	0.0279 (6)	
H15	0.6029	0.3793	0.5633	0.034*	
C16	0.4805 (4)	0.45288 (10)	0.5956 (2)	0.0336 (7)	
H16	0.3364	0.4412	0.5960	0.040*	
C17	0.5304 (4)	0.50657 (10)	0.61392 (18)	0.0299 (6)	
H17	0.4213	0.5321	0.6254	0.036*	
C18	0.4762 (4)	0.69961 (9)	0.62105 (16)	0.0230 (5)	
C19	0.2844 (4)	0.70325 (11)	0.57537 (16)	0.0279 (6)	
H19	0.1721	0.6785	0.5860	0.033*	

C20	0.2575 (5)	0.74308 (13)	0.51430 (19)	0.0420 (8)
H20	0.1263	0.7456	0.4831	0.050*
C21	0.4199 (7)	0.77902 (13)	0.4985 (2)	0.0522 (9)
H21	0.3998	0.8068	0.4575	0.063*
C22	0.6139 (6)	0.77443 (13)	0.5431 (2)	0.0519 (9)
H22	0.7273	0.7986	0.5312	0.062*
C23	0.6420 (5)	0.73508 (11)	0.60416 (18)	0.0352 (7)
H23	0.7741	0.7322	0.6347	0.042*
C24	0.2196 (4)	0.70160 (11)	0.77998 (16)	0.0273 (6)
C25	0.2472 (4)	0.75683 (11)	0.76539 (19)	0.0338 (6)
H25	0.3723	0.7698	0.7401	0.041*
C26	0.0889 (5)	0.79293 (13)	0.7884 (2)	0.0409 (7)
H26	0.1070	0.8309	0.7790	0.049*
C27	-0.0956 (4)	0.77443 (13)	0.8251 (2)	0.0406 (8)
C28	-0.1194 (4)	0.71885 (13)	0.83758 (19)	0.0383 (7)
H28	-0.2454	0.7057	0.8620	0.046*
C29	0.0362 (4)	0.68207 (12)	0.81537 (17)	0.0310 (6)
H29	0.0174	0.6441	0.8242	0.037*
C30	-0.2606 (5)	0.81477 (15)	0.8538 (2)	0.0539 (10)
H30A	-0.3979	0.7962	0.8585	0.081*
H30B	-0.2750	0.8445	0.8129	0.081*
H30C	-0.2157	0.8297	0.9087	0.081*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0197 (3)	0.0221 (3)	0.0349 (4)	-0.0019 (2)	0.0043 (3)	-0.0087 (3)
O1	0.0207 (9)	0.0356 (11)	0.0736 (16)	-0.0014 (8)	-0.0025 (10)	-0.0245 (11)
O2	0.0408 (11)	0.0249 (9)	0.0305 (10)	-0.0031 (8)	0.0153 (9)	-0.0025 (8)
N1	0.061 (6)	0.015 (5)	0.025 (3)	0.003 (4)	-0.019 (4)	-0.008 (3)
N2	0.062 (5)	0.062 (3)	0.033 (3)	0.014 (5)	0.013 (4)	0.021 (3)
N1'	0.061 (6)	0.015 (5)	0.025 (3)	0.003 (4)	-0.019 (4)	-0.008 (3)
N2'	0.062 (5)	0.062 (3)	0.033 (3)	0.014 (5)	0.013 (4)	0.021 (3)
N1''	0.017 (3)	0.021 (4)	0.018 (4)	-0.001 (3)	0.011 (3)	0.007 (3)
N2''	0.081 (6)	0.034 (4)	0.062 (6)	0.000 (4)	0.027 (5)	0.017 (4)
N3	0.0298 (12)	0.0392 (13)	0.0211 (11)	-0.0052 (10)	0.0004 (9)	0.0040 (10)
N4	0.0209 (10)	0.0268 (10)	0.0193 (10)	-0.0008 (8)	0.0009 (8)	-0.0030 (9)
C1	0.044 (4)	0.017 (4)	0.021 (3)	-0.011 (3)	-0.014 (3)	-0.003 (3)
C2	0.068 (7)	0.044 (4)	0.033 (4)	-0.022 (4)	-0.033 (4)	0.010 (3)
C3	0.041 (3)	0.031 (3)	0.026 (3)	0.008 (3)	0.004 (3)	0.007 (2)
C4	0.052 (4)	0.041 (3)	0.040 (3)	0.010 (3)	0.003 (3)	0.007 (3)
C5	0.054 (4)	0.037 (3)	0.045 (4)	0.013 (3)	-0.006 (3)	0.013 (3)
C6	0.055 (5)	0.048 (4)	0.058 (5)	0.028 (4)	-0.005 (4)	0.006 (4)
C7	0.049 (4)	0.042 (4)	0.039 (4)	0.022 (3)	0.006 (4)	0.009 (3)
C8	0.031 (4)	0.026 (3)	0.038 (4)	0.002 (3)	-0.002 (3)	0.002 (3)
C1'	0.044 (4)	0.017 (4)	0.021 (3)	-0.011 (3)	-0.014 (3)	-0.003 (3)
C2'	0.068 (7)	0.044 (4)	0.033 (4)	-0.022 (4)	-0.033 (4)	0.010 (3)
C3'	0.041 (3)	0.031 (3)	0.026 (3)	0.008 (3)	0.004 (3)	0.007 (2)

C4'	0.052 (4)	0.041 (3)	0.040 (3)	0.010 (3)	0.003 (3)	0.007 (3)
C5'	0.054 (4)	0.037 (3)	0.045 (4)	0.013 (3)	-0.006 (3)	0.013 (3)
C6'	0.055 (5)	0.048 (4)	0.058 (5)	0.028 (4)	-0.005 (4)	0.006 (4)
C7'	0.049 (4)	0.042 (4)	0.039 (4)	0.022 (3)	0.006 (4)	0.009 (3)
C8'	0.031 (4)	0.026 (3)	0.038 (4)	0.002 (3)	-0.002 (3)	0.002 (3)
C1''	0.040 (4)	0.033 (5)	0.017 (4)	0.002 (4)	0.006 (4)	0.009 (4)
C2''	0.055 (5)	0.029 (4)	0.042 (5)	0.003 (4)	0.012 (4)	0.000 (4)
C3''	0.064 (6)	0.056 (6)	0.041 (5)	0.014 (5)	0.018 (5)	0.011 (5)
C4''	0.087 (8)	0.053 (6)	0.057 (6)	0.026 (6)	0.008 (6)	0.014 (5)
C5''	0.088 (8)	0.062 (7)	0.076 (7)	0.030 (6)	0.022 (6)	0.012 (6)
C6''	0.071 (7)	0.067 (7)	0.056 (7)	0.024 (6)	-0.001 (6)	0.012 (6)
C7''	0.049 (6)	0.050 (6)	0.054 (6)	0.023 (5)	0.001 (5)	0.000 (5)
C8''	0.058 (6)	0.042 (6)	0.033 (5)	0.017 (5)	0.003 (5)	0.009 (5)
C9	0.0296 (13)	0.0253 (13)	0.0274 (14)	-0.0026 (11)	-0.0031 (11)	0.0049 (11)
C10	0.0210 (12)	0.0188 (11)	0.0240 (13)	-0.0009 (9)	-0.0013 (10)	-0.0032 (10)
C11	0.0178 (11)	0.0213 (11)	0.0186 (12)	-0.0043 (9)	0.0001 (9)	-0.0045 (10)
C12	0.0223 (12)	0.0224 (12)	0.0220 (13)	0.0006 (10)	-0.0002 (10)	-0.0041 (10)
C13	0.0193 (12)	0.0294 (13)	0.0431 (17)	0.0014 (10)	0.0017 (12)	-0.0059 (12)
C14	0.0283 (14)	0.0254 (13)	0.0406 (16)	0.0070 (11)	0.0065 (12)	-0.0037 (12)
C15	0.0324 (14)	0.0199 (12)	0.0317 (15)	-0.0008 (10)	0.0049 (12)	-0.0048 (11)
C16	0.0223 (13)	0.0262 (13)	0.0528 (19)	-0.0042 (11)	0.0066 (13)	-0.0061 (13)
C17	0.0197 (12)	0.0225 (12)	0.0477 (17)	0.0029 (10)	0.0034 (12)	-0.0058 (12)
C18	0.0301 (13)	0.0190 (11)	0.0200 (13)	0.0008 (10)	0.0009 (10)	-0.0017 (10)
C19	0.0293 (13)	0.0302 (13)	0.0241 (14)	0.0045 (11)	-0.0007 (11)	-0.0040 (11)
C20	0.0542 (19)	0.0457 (18)	0.0258 (15)	0.0225 (16)	-0.0046 (14)	0.0014 (14)
C21	0.094 (3)	0.0345 (17)	0.0286 (17)	0.0108 (18)	0.0036 (18)	0.0107 (14)
C22	0.083 (3)	0.0366 (17)	0.0364 (18)	-0.0200 (17)	0.0049 (18)	0.0102 (15)
C23	0.0440 (17)	0.0307 (14)	0.0307 (15)	-0.0110 (13)	-0.0007 (13)	0.0031 (12)
C24	0.0213 (12)	0.0349 (14)	0.0257 (14)	0.0010 (11)	-0.0010 (10)	-0.0155 (12)
C25	0.0258 (13)	0.0356 (15)	0.0403 (16)	-0.0025 (12)	0.0038 (12)	-0.0164 (13)
C26	0.0339 (15)	0.0380 (16)	0.0503 (19)	0.0039 (13)	-0.0052 (14)	-0.0230 (15)
C27	0.0241 (14)	0.0551 (19)	0.0420 (18)	0.0051 (13)	-0.0048 (12)	-0.0297 (15)
C28	0.0223 (13)	0.060 (2)	0.0330 (16)	-0.0050 (13)	0.0014 (12)	-0.0234 (15)
C29	0.0223 (13)	0.0443 (16)	0.0261 (14)	-0.0013 (12)	-0.0025 (11)	-0.0129 (13)
C30	0.0335 (16)	0.064 (2)	0.064 (2)	0.0130 (15)	-0.0042 (16)	-0.0388 (19)

Geometric parameters (Å, °)

S1—O1	1.432 (2)	C1''—C9	1.512 (8)
S1—O2	1.436 (2)	C2''—H2''A	0.9900
S1—C10	1.754 (2)	C2''—H2''B	0.9900
S1—C12	1.769 (2)	C3''—C4''	1.3900
N1—C1	1.322 (9)	C3''—C8''	1.3900
N1—C8	1.375 (9)	C4''—C5''	1.3900
N2—C2	1.370 (9)	C4''—H4''	0.9500
N2—C3	1.408 (7)	C5''—C6''	1.3900
N2—N2 ⁱ	1.856 (16)	C5''—H5''	0.9500
N2—H2	0.8800	C6''—C7''	1.3900

N1'—C1'	1.358 (10)	C6"—H6"	0.9500
N1'—C8'	1.379 (9)	C7"—C8"	1.3900
N2'—C2'	1.385 (9)	C7"—H7"	0.9500
N2'—C3'	1.390 (8)	C9—C10	1.416 (4)
N2'—H2'	0.8800	C10—C11	1.381 (3)
N1"—C1"	1.335 (8)	C11—C18	1.474 (3)
N1"—C8"	1.361 (8)	C12—C13	1.381 (3)
N2"—C2"	1.392 (8)	C12—C17	1.380 (3)
N2"—C3"	1.400 (8)	C13—C14	1.385 (4)
N2"—H2"	0.8800	C13—H13	0.9500
N3—C9	1.328 (3)	C14—C15	1.371 (4)
N3—N4	1.355 (3)	C14—H14	0.9500
N4—C11	1.359 (3)	C15—C16	1.383 (4)
N4—C24	1.436 (3)	C15—H15	0.9500
C1—C2	1.469 (9)	C16—C17	1.382 (4)
C1—C9	1.480 (8)	C16—H16	0.9500
C2—H2A	0.9900	C17—H17	0.9500
C2—H2B	0.9900	C18—C19	1.389 (4)
C3—C4	1.3900	C18—C23	1.390 (4)
C3—C8	1.3900	C19—C20	1.384 (4)
C4—C5	1.3900	C19—H19	0.9500
C4—H4A	0.9500	C20—C21	1.378 (5)
C5—C6	1.3900	C20—H20	0.9500
C5—H5	0.9500	C21—C22	1.394 (5)
C6—C7	1.3900	C21—H21	0.9500
C6—H6	0.9500	C22—C23	1.377 (4)
C7—C8	1.3900	C22—H22	0.9500
C7—H7	0.9500	C23—H23	0.9500
C1'—C2'	1.483 (10)	C24—C29	1.385 (4)
C1'—C9	1.494 (9)	C24—C25	1.386 (4)
C2'—H2'A	0.9900	C25—C26	1.390 (4)
C2'—H2'B	0.9900	C25—H25	0.9500
C3'—C4'	1.3900	C26—C27	1.391 (4)
C3'—C8'	1.3900	C26—H26	0.9500
C4'—C5'	1.3900	C27—C28	1.386 (5)
C4'—H4'	0.9500	C27—C30	1.515 (4)
C5'—C6'	1.3900	C28—C29	1.384 (4)
C5'—H5'	0.9500	C28—H28	0.9500
C6'—C7'	1.3900	C29—H29	0.9500
C6'—H6'	0.9500	C30—H30A	0.9800
C7'—C8'	1.3900	C30—H30B	0.9800
C7'—H7'	0.9500	C30—H30C	0.9800
C1"—C2"	1.470 (9)		
O1—S1—O2	116.63 (13)	C8"—C3"—N2"	122.5 (6)
O1—S1—C10	111.78 (12)	C5"—C4"—C3"	120.0
O2—S1—C10	106.57 (11)	C5"—C4"—H4"	120.0
O1—S1—C12	108.98 (12)	C3"—C4"—H4"	120.0

O2—S1—C12	107.84 (11)	C4"—C5"—C6"	120.0
C10—S1—C12	104.29 (12)	C4"—C5"—H5"	120.0
C1—N1—C8	112.2 (10)	C6"—C5"—H5"	120.0
C2—N2—C3	108.9 (9)	C7"—C6"—C5"	120.0
C2—N2—N2 ⁱ	98.0 (8)	C7"—C6"—H6"	120.0
C3—N2—N2 ⁱ	135.4 (9)	C5"—C6"—H6"	120.0
C2—N2—H2	125.5	C8"—C7"—C6"	120.0
C3—N2—H2	125.5	C8"—C7"—H7"	120.0
C1'—N1'—C8'	127.4 (13)	C6"—C7"—H7"	120.0
C2'—N2'—C3'	123.8 (10)	N1"—C8"—C7"	117.5 (6)
C2'—N2'—H2'	118.1	N1"—C8"—C3"	122.3 (6)
C3'—N2'—H2'	118.1	C7"—C8"—C3"	120.0
C1"—N1"—C8"	117.2 (7)	N3—C9—C10	110.6 (2)
C2"—N2"—C3"	115.8 (7)	N3—C9—C1	113.7 (6)
C2"—N2"—H2"	122.1	C10—C9—C1	134.2 (6)
C3"—N2"—H2"	122.1	N3—C9—C1'	114.3 (7)
C9—N3—N4	105.3 (2)	C10—C9—C1'	135.1 (7)
N3—N4—C11	112.7 (2)	N3—C9—C1"	130.9 (4)
N3—N4—C24	118.0 (2)	C10—C9—C1"	118.0 (4)
C11—N4—C24	129.3 (2)	C11—C10—C9	105.8 (2)
N1—C1—C2	117.8 (10)	C11—C10—S1	124.51 (19)
N1—C1—C9	113.7 (8)	C9—C10—S1	129.57 (19)
C2—C1—C9	114.3 (8)	N4—C11—C10	105.6 (2)
N2—C2—C1	124.2 (11)	N4—C11—C18	121.6 (2)
N2—C2—H2A	106.3	C10—C11—C18	132.7 (2)
C1—C2—H2A	106.3	C13—C12—C17	120.8 (2)
N2—C2—H2B	106.3	C13—C12—S1	117.43 (19)
C1—C2—H2B	106.3	C17—C12—S1	121.52 (19)
H2A—C2—H2B	106.4	C12—C13—C14	119.3 (2)
C4—C3—C8	120.0	C12—C13—H13	120.3
C4—C3—N2	117.2 (5)	C14—C13—H13	120.3
C8—C3—N2	122.8 (5)	C15—C14—C13	120.5 (2)
C5—C4—C3	120.0	C15—C14—H14	119.8
C5—C4—H4A	120.0	C13—C14—H14	119.8
C3—C4—H4A	120.0	C14—C15—C16	119.7 (2)
C4—C5—C6	120.0	C14—C15—H15	120.1
C4—C5—H5	120.0	C16—C15—H15	120.1
C6—C5—H5	120.0	C15—C16—C17	120.6 (2)
C7—C6—C5	120.0	C15—C16—H16	119.7
C7—C6—H6	120.0	C17—C16—H16	119.7
C5—C6—H6	120.0	C12—C17—C16	119.1 (2)
C6—C7—C8	120.0	C12—C17—H17	120.5
C6—C7—H7	120.0	C16—C17—H17	120.5
C8—C7—H7	120.0	C19—C18—C23	120.1 (2)
N1—C8—C7	114.6 (7)	C19—C18—C11	120.5 (2)
N1—C8—C3	125.3 (7)	C23—C18—C11	119.4 (2)
C7—C8—C3	120.0	C20—C19—C18	119.8 (3)
N1'—C1'—C2'	113.9 (12)	C20—C19—H19	120.1

N1'—C1'—C9	110.2 (9)	C18—C19—H19	120.1
C2'—C1'—C9	119.7 (10)	C21—C20—C19	120.4 (3)
N2'—C2'—C1'	110.7 (11)	C21—C20—H20	119.8
N2'—C2'—H2'A	109.5	C19—C20—H20	119.8
C1'—C2'—H2'A	109.5	C20—C21—C22	119.8 (3)
N2'—C2'—H2'B	109.5	C20—C21—H21	120.1
C1'—C2'—H2'B	109.5	C22—C21—H21	120.1
H2'A—C2'—H2'B	108.1	C23—C22—C21	120.3 (3)
C4'—C3'—C8'	120.0	C23—C22—H22	119.8
C4'—C3'—N2'	118.3 (6)	C21—C22—H22	119.8
C8'—C3'—N2'	121.7 (6)	C22—C23—C18	119.7 (3)
C3'—C4'—C5'	120.0	C22—C23—H23	120.1
C3'—C4'—H4'	120.0	C18—C23—H23	120.1
C5'—C4'—H4'	120.0	C29—C24—C25	121.1 (2)
C4'—C5'—C6'	120.0	C29—C24—N4	118.4 (2)
C4'—C5'—H5'	120.0	C25—C24—N4	120.4 (2)
C6'—C5'—H5'	120.0	C24—C25—C26	118.9 (3)
C7'—C6'—C5'	120.0	C24—C25—H25	120.6
C7'—C6'—H6'	120.0	C26—C25—H25	120.6
C5'—C6'—H6'	120.0	C25—C26—C27	121.1 (3)
C6'—C7'—C8'	120.0	C25—C26—H26	119.4
C6'—C7'—H7'	120.0	C27—C26—H26	119.4
C8'—C7'—H7'	120.0	C28—C27—C26	118.4 (3)
N1'—C8'—C7'	128.0 (9)	C28—C27—C30	121.4 (3)
N1'—C8'—C3'	112.0 (9)	C26—C27—C30	120.1 (3)
C7'—C8'—C3'	120.0	C29—C28—C27	121.6 (3)
N1"—C1"—C2"	122.8 (7)	C29—C28—H28	119.2
N1"—C1"—C9	118.6 (7)	C27—C28—H28	119.2
C2"—C1"—C9	118.5 (7)	C24—C29—C28	118.9 (3)
N2"—C2"—C1"	119.1 (8)	C24—C29—H29	120.6
N2"—C2"—H2"A	107.5	C28—C29—H29	120.6
C1"—C2"—H2"A	107.5	C27—C30—H30A	109.5
N2"—C2"—H2"B	107.5	C27—C30—H30B	109.5
C1"—C2"—H2"B	107.5	H30A—C30—H30B	109.5
H2"A—C2"—H2"B	107.0	C27—C30—H30C	109.5
C4"—C3"—C8"	120.0	H30A—C30—H30C	109.5
C4"—C3"—N2"	117.5 (6)	H30B—C30—H30C	109.5
C9—N3—N4—C11	0.9 (3)	C2'—C1'—C9—N3	-26.6 (14)
C9—N3—N4—C24	-179.4 (2)	N1'—C1'—C9—C10	23.1 (17)
C8—N1—C1—C2	-29.4 (15)	C2'—C1'—C9—C10	158.1 (8)
C8—N1—C1—C9	-167.0 (9)	N1'—C1'—C9—C1	-71 (4)
C3—N2—C2—C1	-22.3 (15)	C2'—C1'—C9—C1	64 (4)
N2 ⁱ —N2—C2—C1	-166.2 (11)	N1'—C1'—C9—C1"	32 (2)
N1—C1—C2—N2	37.2 (17)	C2'—C1'—C9—C1"	167 (4)
C9—C1—C2—N2	174.5 (10)	N1"—C1"—C9—N3	-147.6 (10)
C2—N2—C3—C4	-174.5 (7)	C2"—C1"—C9—N3	36.5 (13)
N2 ⁱ —N2—C3—C4	-50.6 (14)	N1"—C1"—C9—C10	41.6 (14)

C2—N2—C3—C8	6.3 (11)	C2"—C1"—C9—C10	-134.3 (8)
N2 ⁱ —N2—C3—C8	130.2 (11)	N1"—C1"—C9—C1	-96 (2)
C8—C3—C4—C5	0.0	C2"—C1"—C9—C1	87.9 (17)
N2—C3—C4—C5	-179.2 (7)	N1"—C1"—C9—C1'	-132 (3)
C3—C4—C5—C6	0.0	C2"—C1"—C9—C1'	53 (2)
C4—C5—C6—C7	0.0	N3—C9—C10—C11	0.3 (3)
C5—C6—C7—C8	0.0	C1—C9—C10—C11	-164.4 (7)
C1—N1—C8—C7	-167.1 (8)	C1'—C9—C10—C11	175.7 (9)
C1—N1—C8—C3	15.5 (14)	C1"—C9—C10—C11	172.9 (5)
C6—C7—C8—N1	-177.5 (8)	N3—C9—C10—S1	-175.66 (19)
C6—C7—C8—C3	0.0	C1—C9—C10—S1	19.6 (7)
C4—C3—C8—N1	177.3 (8)	C1'—C9—C10—S1	-0.3 (9)
N2—C3—C8—N1	-3.6 (10)	C1"—C9—C10—S1	-3.1 (6)
C4—C3—C8—C7	0.0	O1—S1—C10—C11	119.9 (2)
N2—C3—C8—C7	179.2 (7)	O2—S1—C10—C11	-8.6 (2)
C8'—N1'—C1'—C2'	34 (3)	C12—S1—C10—C11	-122.5 (2)
C8'—N1'—C1'—C9	172.2 (17)	O1—S1—C10—C9	-64.8 (3)
C3'—N2'—C2'—C1'	26.5 (18)	O2—S1—C10—C9	166.7 (2)
N1'—C1'—C2'—N2'	-35.5 (18)	C12—S1—C10—C9	52.8 (3)
C9—C1'—C2'—N2'	-169.0 (11)	N3—N4—C11—C10	-0.7 (3)
C2'—N2'—C3'—C4'	169.7 (10)	C24—N4—C11—C10	179.7 (2)
C2'—N2'—C3'—C8'	-10.8 (16)	N3—N4—C11—C18	-179.4 (2)
C8'—C3'—C4'—C5'	0.0	C24—N4—C11—C18	0.9 (4)
N2'—C3'—C4'—C5'	179.4 (8)	C9—C10—C11—N4	0.2 (3)
C3'—C4'—C5'—C6'	0.0	S1—C10—C11—N4	176.44 (17)
C4'—C5'—C6'—C7'	0.0	C9—C10—C11—C18	178.8 (2)
C5'—C6'—C7'—C8'	0.0	S1—C10—C11—C18	-5.0 (4)
C1'—N1'—C8'—C7'	165.9 (14)	O1—S1—C12—C13	-28.8 (3)
C1'—N1'—C8'—C3'	-17 (2)	O2—S1—C12—C13	98.7 (2)
C6'—C7'—C8'—N1'	176.9 (14)	C10—S1—C12—C13	-148.3 (2)
C6'—C7'—C8'—C3'	0.0	O1—S1—C12—C17	157.2 (2)
C4'—C3'—C8'—N1'	-177.4 (12)	O2—S1—C12—C17	-75.4 (2)
N2'—C3'—C8'—N1'	3.2 (13)	C10—S1—C12—C17	37.7 (3)
C4'—C3'—C8'—C7'	0.0	C17—C12—C13—C14	0.3 (4)
N2'—C3'—C8'—C7'	-179.4 (9)	S1—C12—C13—C14	-173.9 (2)
C8"—N1"—C1"—C2"	-4 (2)	C12—C13—C14—C15	-1.1 (4)
C8"—N1"—C1"—C9	-179.4 (10)	C13—C14—C15—C16	0.5 (4)
C3"—N2"—C2"—C1"	-2.6 (15)	C14—C15—C16—C17	0.9 (4)
N1"—C1"—C2"—N2"	2.7 (18)	C13—C12—C17—C16	1.2 (4)
C9—C1"—C2"—N2"	178.4 (9)	S1—C12—C17—C16	175.0 (2)
C2"—N2"—C3"—C4"	-176.3 (7)	C15—C16—C17—C12	-1.8 (5)
C2"—N2"—C3"—C8"	3.9 (12)	N4—C11—C18—C19	-63.1 (3)
C8"—C3"—C4"—C5"	0.0	C10—C11—C18—C19	118.5 (3)
N2"—C3"—C4"—C5"	-179.7 (9)	N4—C11—C18—C23	113.8 (3)
C3"—C4"—C5"—C6"	0.0	C10—C11—C18—C23	-64.6 (4)
C4"—C5"—C6"—C7"	0.0	C23—C18—C19—C20	-1.4 (4)
C5"—C6"—C7"—C8"	0.0	C11—C18—C19—C20	175.5 (2)
C1"—N1"—C8"—C7"	-179.9 (10)	C18—C19—C20—C21	0.1 (4)

C1"—N1"—C8"—C3"	5.0 (17)	C19—C20—C21—C22	1.4 (5)
C6"—C7"—C8"—N1"	-175.1 (10)	C20—C21—C22—C23	-1.7 (5)
C6"—C7"—C8"—C3"	0.0	C21—C22—C23—C18	0.4 (5)
C4"—C3"—C8"—N1"	174.9 (11)	C19—C18—C23—C22	1.2 (4)
N2"—C3"—C8"—N1"	-5.4 (12)	C11—C18—C23—C22	-175.7 (3)
C4"—C3"—C8"—C7"	0.0	N3—N4—C24—C29	-42.7 (3)
N2"—C3"—C8"—C7"	179.7 (9)	C11—N4—C24—C29	136.9 (3)
N4—N3—C9—C10	-0.7 (3)	N3—N4—C24—C25	136.0 (3)
N4—N3—C9—C1	167.4 (5)	C11—N4—C24—C25	-44.4 (4)
N4—N3—C9—C1'	-177.1 (7)	C29—C24—C25—C26	1.3 (4)
N4—N3—C9—C1"	-172.0 (6)	N4—C24—C25—C26	-177.4 (2)
N1—C1—C9—N3	161.6 (8)	C24—C25—C26—C27	-0.4 (4)
C2—C1—C9—N3	22.5 (11)	C25—C26—C27—C28	-0.5 (5)
N1—C1—C9—C10	-34.1 (13)	C25—C26—C27—C30	176.8 (3)
C2—C1—C9—C10	-173.2 (7)	C26—C27—C28—C29	0.7 (4)
N1—C1—C9—C1'	66 (4)	C30—C27—C28—C29	-176.6 (3)
C2—C1—C9—C1'	-73 (4)	C25—C24—C29—C28	-1.1 (4)
N1—C1—C9—C1"	21.7 (12)	N4—C24—C29—C28	177.6 (2)
C2—C1—C9—C1"	-117 (2)	C27—C28—C29—C24	0.1 (4)
N1'—C1'—C9—N3	-161.6 (12)		

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

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C19—H19 \cdots O1 ⁱⁱ	0.95	2.57	3.430 (3)	150

Symmetry code: (ii) $x-1, y, z$.