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3-Cyclohexyl-1-(3,5-dinitrobenzoyl)thiourea

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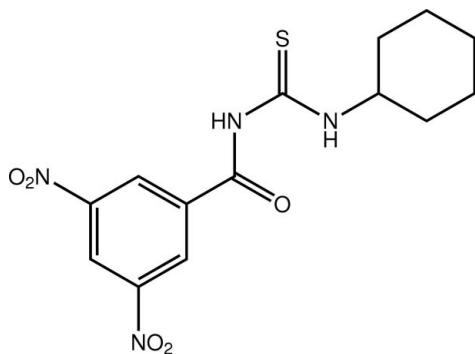
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in main residue; R factor = 0.061; wR factor = 0.211; data-to-parameter ratio = 13.5.

The structure of the title thiourea derivative, $\text{C}_{14}\text{H}_{16}\text{N}_4\text{O}_5\text{S}$, features an almost planar central $\text{C}_2\text{N}_2\text{OS}$ fragment (r.m.s. deviation = 0.005 Å), an arrangement stabilized by an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond. The terminal rings are twisted out of this plane, the dihedral angle formed with the benzene ring being $33.22(10)^\circ$. The cyclohexyl ring is disordered, with two orientations (50:50) being resolved. The mean plane passing through the atoms of each disordered component forms dihedral angles of $65.7(2)$ and $82.4(3)^\circ$ with the central plane. Centrosymmetric dimers mediated by an eight-membered $\{\cdots\text{HNC}=\text{S}\}_2$ synthon occur in the crystal.

Related literature

For the biological activity of thiourea derivatives, see: Venkatachalam *et al.* (2004); Saeed *et al.* (2011). For related thiourea structures, see: Gunasekaran *et al.* (2010); Saeed *et al.* (2010); Dzulkifli *et al.* (2011).



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Experimental

Crystal data

$\text{C}_{14}\text{H}_{16}\text{N}_4\text{O}_5\text{S}$
 $M_r = 352.37$
 Monoclinic, $P2_1/c$
 $a = 12.3404(7)$ Å
 $b = 9.0506(5)$ Å
 $c = 14.6534(6)$ Å
 $\beta = 90.385(5)^\circ$

$V = 1636.57(15)$ Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.23$ mm⁻¹
 $T = 295$ K
 $0.20 \times 0.15 \times 0.10$ mm

Data collection

Agilent Technologies SuperNova
 Dual diffractometer with an Atlas detector
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2010)
 $T_{\min} = 0.955$, $T_{\max} = 0.977$

7954 measured reflections
 3649 independent reflections
 1948 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.061$
 $wR(F^2) = 0.211$
 $S = 1.01$
 3649 reflections
 271 parameters

25 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.22$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O1}$	0.88	1.89	2.639 (4)	142
$\text{N1}-\text{H1}'\cdots\text{O1}$	0.88	1.99	2.639 (4)	130
$\text{N2}-\text{H2}\cdots\text{S1}^i$	0.88	2.65	3.449 (3)	152

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

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: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); 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: HB5839).

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supporting information

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3-Cyclohexyl-1-(3,5-dinitrobenzoyl)thiourea

Sohail Saeed, Naghmana Rashid, Muhammad Sher, Seik Weng Ng and Edward R. T. Tiekink

S1. Comment

Continuing structural studies (Gunasekaran *et al.* 2010; Saeed *et al.* 2010; Dzulkipli *et al.*, 2011) of thiourea derivatives are motivated by their biological potential (Venkatachalam *et al.*, 2004; Saeed *et al.*, 2011) and led to the investigation of the title compound, (I).

The molecular structure of (I), Fig. 1, is highly twisted with dihedral angles formed between the central chromophore (r.m.s. = 0.0054 Å for C7,C8,N1,N2,O1 & S1) and the benzene ring being 33.22 (10) °. Two orientations of equal weight were found for the cyclohexyl ring, each with a chair conformation, and these make angles of 65.74 (24) and 82.42 (30) °, respectively, with the central plane. The N—H atoms are anti as are the S and O atoms. As a consequence, the N1—H atom forms an intramolecular hydrogen bond with the carbonyl-O1 atom to close a pseudo six-membered ring, Table 1; there are two values cited owing to the disorder in the molecule. The nitro groups are effectively co-planar with the benzene ring to which they are bonded as seen in the values of the O2—N3—C11—C10 and O4—N4—C13—C12 torsion angles of 1.2 (5) and -7.0 (5) °, respectively.

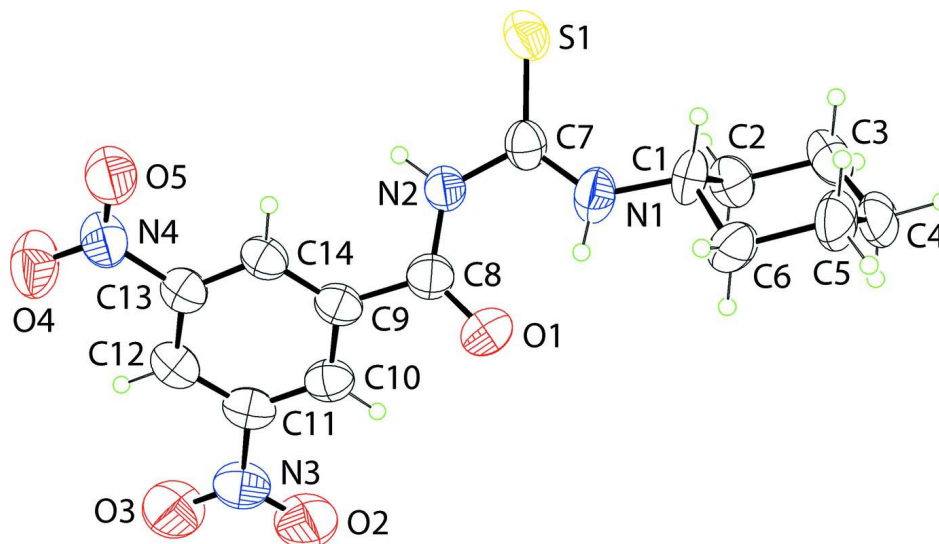
The most prominent feature of the crystal packing is the formation of centrosymmetric eight-membered {···HNC=S}₂ synthon leading to dimeric aggregates, Fig. 2 and Table 1.

S2. Experimental

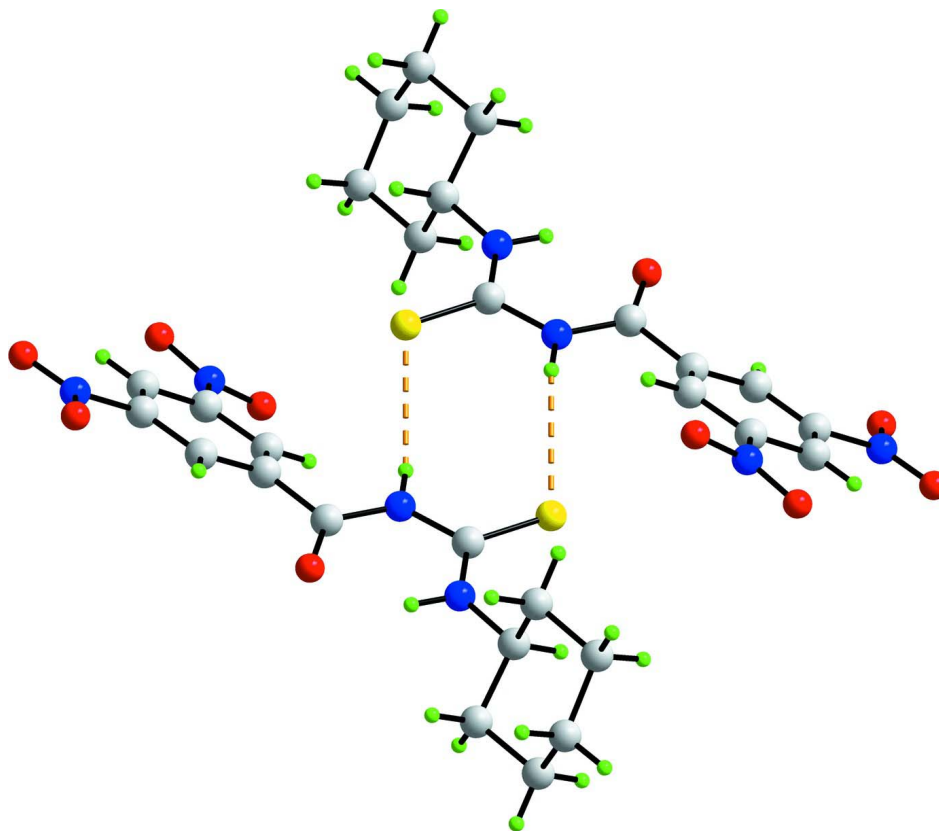
A solution of 3,5-dinitrobenzoyl chloride (0.01 mol) in anhydrous acetone (75 ml) and 3% tetrabutylammonium bromide (TBAB), as a phase-transfer catalyst (PTC), in anhydrous acetone was added drop-wise to a suspension of dry potassium thiocyanate (0.01 mol) in acetone (50 ml). The reaction mixture was refluxed for 50 min. After cooling to room temperature, a solution of cyclohexylamine (0.01 mol) in anhydrous acetone (25 ml) was added drop-wise and the resulting mixture refluxed for 3 h. Hydrochloric acid (0.1 N, 300 ml) was added and the solution was filtered. The solid product was washed with water and purified by re-crystallization from ethanol; Yield: 1.50 g (88%) and *M.pt.* 409 K. IR (KBr, cm⁻¹): 3215 ν(NH), 1673 (C=O), 1527 (benzene ring), 1138 ν(C=S). Anal. Calcd. for C₁₄H₁₆N₄O₅S: C, 47.72; H, 4.58; N, 15.90; S, 9.10%. Found: C, 47.51; H, 4.75; N, 15.88; S, 9.11%.

S3. Refinement

Carbon-bound H-atoms were placed in calculated positions [C—H 0.93 to 0.97 Å, $U_{\text{iso}}(\text{H}) 1.2U_{\text{eq}}(\text{C})$] and were included in the refinement in the riding model approximation. The two amino H-atoms were similarly placed [N—H 0.88 Å, $U_{\text{iso}}(\text{H}) 1.2U_{\text{eq}}(\text{N})$]. The cyclohexyl ring is disordered over two positions; the disorder could not be refined, and was assumed to be a 1:1 type of disorder. The 1,2-related C—C distances were restrained to 1.54±0.01 Å and the 1,3-related ones to 2.51±0.01 Å. The pair of *N*-C_{cyclohexyl} and *N*-C'_{cyclohexyl} distances were restrained to within 0.01 Å of each other.

**Figure 1**

The molecular structure of (I) showing displacement ellipsoids at the 35% probability level. Only one orientation of the disordered cyclohexyl ring is shown.

**Figure 2**

Supramolecular dimer in (I) mediated by N—H...S hydrogen bonding shown as orange dashed lines. Only one orientation of the disordered cyclohexyl ring is shown.

3-Cyclohexyl-1-(3,5-dinitrobenzoyl)thiourea

Crystal data

C₁₄H₁₆N₄O₅S $M_r = 352.37$ Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

 $a = 12.3404$ (7) Å $b = 9.0506$ (5) Å $c = 14.6534$ (6) Å $\beta = 90.385$ (5)° $V = 1636.57$ (15) Å³ $Z = 4$ $F(000) = 736$ $D_x = 1.430$ Mg m⁻³Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2522 reflections

 $\theta = 2.6$ – 29.2 ° $\mu = 0.23$ mm⁻¹ $T = 295$ K

Prism, colorless

 $0.20 \times 0.15 \times 0.10$ mm

Data collection

Agilent Technologies SuperNova Dual
diffractometer with an Atlas detector
Radiation source: SuperNova (Mo) X-ray
Source

Mirror monochromator

Detector resolution: 10.4041 pixels mm⁻¹ ω scansAbsorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2010) $T_{\min} = 0.955$, $T_{\max} = 0.977$

7954 measured reflections

3649 independent reflections

1948 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.024$ $\theta_{\max} = 27.5$ °, $\theta_{\min} = 2.7$ ° $h = -11 \rightarrow 16$ $k = -11 \rightarrow 9$ $l = -19 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.061$ $wR(F^2) = 0.211$ $S = 1.01$

3649 reflections

271 parameters

25 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

 $w = 1/[\sigma^2(F_o^2) + (0.0901P)^2 + 0.5204P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} = 0.001$ $\Delta\rho_{\max} = 0.22$ e Å⁻³ $\Delta\rho_{\min} = -0.28$ e Å⁻³Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.41365 (9)	0.64167 (12)	0.59283 (6)	0.0914 (4)	
O1	0.2058 (2)	0.5972 (4)	0.33847 (18)	0.1069 (9)	
O2	0.1940 (3)	0.2554 (4)	0.0798 (2)	0.1410 (13)	
O3	0.3315 (3)	0.2306 (4)	-0.0052 (3)	0.1351 (13)	
O4	0.6723 (3)	0.4548 (4)	0.0872 (2)	0.1237 (11)	
O5	0.6733 (2)	0.6067 (4)	0.2004 (2)	0.1140 (10)	
N1	0.2226 (2)	0.6699 (3)	0.5124 (2)	0.0864 (9)	
H1	0.1881	0.6629	0.4598	0.104*	0.50
H1'	0.1777	0.6614	0.4657	0.104*	0.50
N2	0.3606 (2)	0.5812 (3)	0.42290 (16)	0.0689 (7)	
H2	0.4294	0.5564	0.4197	0.083*	
N3	0.2883 (4)	0.2769 (4)	0.0621 (3)	0.0964 (10)	
N4	0.6288 (3)	0.5175 (4)	0.1510 (2)	0.0907 (9)	

C1	0.1543 (7)	0.7215 (9)	0.5861 (6)	0.074 (3)	0.50
H1A	0.1990	0.7735	0.6313	0.088*	0.50
C2	0.0991 (8)	0.5900 (9)	0.6317 (6)	0.084 (3)	0.50
H2A	0.1529	0.5203	0.6537	0.101*	0.50
H2B	0.0520	0.5398	0.5885	0.101*	0.50
C3	0.0323 (6)	0.6513 (8)	0.7122 (4)	0.094 (2)	0.50
H3A	-0.0055	0.5709	0.7420	0.113*	0.50
H3B	0.0807	0.6964	0.7567	0.113*	0.50
C4	-0.0488 (6)	0.7647 (9)	0.6791 (5)	0.106 (3)	0.50
H4A	-0.1009	0.7171	0.6389	0.127*	0.50
H4B	-0.0879	0.8040	0.7309	0.127*	0.50
C5	0.0049 (6)	0.8899 (8)	0.6289 (5)	0.098 (3)	0.50
H5A	0.0532	0.9427	0.6700	0.117*	0.50
H5B	-0.0496	0.9586	0.6069	0.117*	0.50
C6	0.0701 (8)	0.8278 (11)	0.5471 (5)	0.091 (3)	0.50
H6A	0.0222	0.7764	0.5051	0.109*	0.50
H6B	0.1054	0.9075	0.5146	0.109*	0.50
C1'	0.1883 (7)	0.7287 (12)	0.6020 (6)	0.121 (6)	0.50
H1B	0.2530	0.7692	0.6321	0.145*	0.50
C2'	0.1393 (7)	0.6133 (12)	0.6672 (7)	0.101 (4)	0.50
H2C	0.1891	0.5312	0.6747	0.121*	0.50
H2D	0.1277	0.6576	0.7267	0.121*	0.50
C3'	0.0326 (8)	0.5580 (10)	0.6290 (9)	0.149 (6)	0.50
H3C	0.0025	0.4846	0.6699	0.179*	0.50
H3D	0.0448	0.5112	0.5704	0.179*	0.50
C4'	-0.0484 (6)	0.6857 (12)	0.6172 (9)	0.146 (5)	0.50
H4C	-0.0635	0.7300	0.6760	0.175*	0.50
H4D	-0.1159	0.6487	0.5917	0.175*	0.50
C5'	0.0006 (8)	0.8018 (12)	0.5529 (10)	0.150 (5)	0.50
H5C	0.0107	0.7586	0.4930	0.180*	0.50
H5D	-0.0492	0.8842	0.5467	0.180*	0.50
C6'	0.1077 (7)	0.8568 (10)	0.5890 (8)	0.106 (3)	0.50
H6C	0.0968	0.9063	0.6469	0.127*	0.50
H6D	0.1376	0.9281	0.5466	0.127*	0.50
C7	0.3244 (3)	0.6318 (3)	0.5079 (2)	0.0695 (8)	
C8	0.3013 (3)	0.5664 (4)	0.3450 (2)	0.0755 (9)	
C9	0.3604 (3)	0.5071 (4)	0.2637 (2)	0.0701 (8)	
C10	0.3015 (3)	0.4227 (4)	0.2026 (2)	0.0757 (9)	
H10	0.2288	0.4025	0.2131	0.091*	
C11	0.3514 (3)	0.3689 (3)	0.1262 (2)	0.0753 (9)	
C12	0.4585 (3)	0.3965 (3)	0.1073 (2)	0.0751 (9)	
H12	0.4916	0.3582	0.0556	0.090*	
C13	0.5142 (3)	0.4833 (3)	0.1684 (2)	0.0700 (8)	
C14	0.4681 (3)	0.5400 (3)	0.2462 (2)	0.0691 (8)	
H14	0.5082	0.5988	0.2860	0.083*	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0966 (7)	0.1080 (8)	0.0695 (5)	0.0228 (6)	0.0037 (5)	-0.0177 (5)
O1	0.0737 (17)	0.145 (3)	0.1016 (18)	0.0229 (17)	-0.0108 (14)	-0.0007 (17)
O2	0.127 (3)	0.161 (3)	0.135 (3)	-0.052 (3)	-0.019 (2)	-0.027 (2)
O3	0.144 (3)	0.126 (3)	0.136 (3)	-0.003 (2)	-0.014 (2)	-0.066 (2)
O4	0.117 (2)	0.121 (2)	0.134 (2)	-0.0209 (19)	0.045 (2)	-0.043 (2)
O5	0.108 (2)	0.131 (2)	0.1037 (19)	-0.0431 (19)	0.0195 (17)	-0.0334 (18)
N1	0.0751 (18)	0.095 (2)	0.0897 (19)	0.0212 (16)	0.0215 (15)	0.0104 (16)
N2	0.0680 (15)	0.0767 (16)	0.0620 (14)	0.0098 (13)	0.0057 (12)	0.0030 (12)
N3	0.111 (3)	0.080 (2)	0.098 (2)	-0.012 (2)	-0.019 (2)	-0.0087 (18)
N4	0.102 (2)	0.086 (2)	0.0843 (19)	-0.0157 (19)	0.0192 (18)	-0.0099 (17)
C1	0.058 (5)	0.071 (6)	0.093 (5)	0.015 (4)	0.026 (4)	0.001 (4)
C2	0.097 (8)	0.077 (5)	0.078 (6)	0.007 (6)	0.011 (6)	-0.005 (5)
C3	0.115 (6)	0.100 (5)	0.069 (4)	-0.024 (5)	0.029 (4)	-0.017 (4)
C4	0.085 (5)	0.131 (8)	0.103 (6)	-0.007 (5)	0.020 (5)	-0.071 (6)
C5	0.085 (5)	0.095 (6)	0.114 (6)	0.018 (4)	0.022 (5)	-0.037 (5)
C6	0.070 (6)	0.099 (7)	0.103 (7)	0.023 (6)	-0.007 (5)	-0.006 (5)
C1'	0.088 (8)	0.119 (11)	0.157 (11)	0.041 (7)	0.051 (7)	0.032 (8)
C2'	0.087 (7)	0.128 (8)	0.087 (7)	-0.004 (6)	0.009 (5)	0.007 (6)
C3'	0.106 (8)	0.160 (12)	0.180 (12)	-0.038 (8)	-0.043 (8)	0.069 (10)
C4'	0.071 (5)	0.162 (10)	0.203 (13)	-0.001 (7)	-0.014 (7)	0.077 (10)
C5'	0.099 (8)	0.124 (9)	0.227 (15)	-0.010 (8)	-0.057 (9)	0.049 (10)
C6'	0.090 (7)	0.092 (7)	0.136 (9)	0.007 (5)	-0.004 (6)	0.000 (6)
C7	0.076 (2)	0.0639 (18)	0.0689 (18)	0.0095 (16)	0.0127 (16)	0.0061 (14)
C8	0.077 (2)	0.076 (2)	0.073 (2)	0.0066 (18)	-0.0027 (17)	0.0090 (16)
C9	0.083 (2)	0.0665 (18)	0.0607 (16)	0.0028 (17)	-0.0076 (15)	0.0113 (15)
C10	0.078 (2)	0.0711 (19)	0.077 (2)	-0.0019 (17)	-0.0112 (17)	0.0110 (17)
C11	0.096 (3)	0.0574 (18)	0.0726 (19)	-0.0027 (18)	-0.0183 (18)	0.0045 (15)
C12	0.102 (3)	0.0593 (18)	0.0641 (18)	0.0011 (18)	-0.0014 (18)	0.0019 (15)
C13	0.083 (2)	0.0607 (17)	0.0665 (18)	-0.0054 (16)	-0.0008 (16)	0.0053 (15)
C14	0.084 (2)	0.0631 (18)	0.0598 (16)	-0.0052 (16)	-0.0035 (16)	0.0051 (14)

Geometric parameters (Å, °)

S1—C7	1.659 (4)	C5—H5B	0.9700
O1—C8	1.215 (4)	C6—H6A	0.9700
O2—N3	1.210 (5)	C6—H6B	0.9700
O3—N3	1.200 (4)	C1'—C6'	1.539 (8)
O4—N4	1.221 (4)	C1'—C2'	1.542 (8)
O5—N4	1.213 (4)	C1'—H1B	0.9800
N1—C7	1.305 (4)	C2'—C3'	1.512 (8)
N1—C1	1.452 (6)	C2'—H2C	0.9700
N1—C1'	1.481 (8)	C2'—H2D	0.9700
N1—H1	0.8800	C3'—C4'	1.537 (9)
N1—H1'	0.8800	C3'—H3C	0.9700
N2—C8	1.358 (4)	C3'—H3D	0.9700

N2—C7	1.402 (4)	C4'—C5'	1.537 (8)
N2—H2	0.8800	C4'—H4C	0.9700
N3—C11	1.474 (5)	C4'—H4D	0.9700
N4—C13	1.471 (5)	C5'—C6'	1.505 (8)
C1—C6	1.524 (8)	C5'—H5C	0.9700
C1—C2	1.527 (8)	C5'—H5D	0.9700
C1—H1A	0.9800	C6'—H6C	0.9700
C2—C3	1.547 (7)	C6'—H6D	0.9700
C2—H2A	0.9700	C8—C9	1.501 (5)
C2—H2B	0.9700	C9—C10	1.380 (5)
C3—C4	1.511 (8)	C9—C14	1.388 (4)
C3—H3A	0.9700	C10—C11	1.370 (5)
C3—H3B	0.9700	C10—H10	0.9300
C4—C5	1.507 (7)	C11—C12	1.375 (5)
C4—H4A	0.9700	C12—C13	1.372 (5)
C4—H4B	0.9700	C12—H12	0.9300
C5—C6	1.552 (8)	C13—C14	1.376 (4)
C5—H5A	0.9700	C14—H14	0.9300
C7—N1—C1	133.3 (5)	C2'—C1'—H1B	107.3
C7—N1—C1'	114.9 (5)	C3'—C2'—C1'	109.8 (7)
C7—N1—H1	113.3	C3'—C2'—H2C	109.7
C1—N1—H1	113.3	C1'—C2'—H2C	109.7
C7—N1—H1'	122.6	C3'—C2'—H2D	109.7
C1'—N1—H1'	122.6	C1'—C2'—H2D	109.7
C8—N2—C7	127.2 (3)	H2C—C2'—H2D	108.2
C8—N2—H2	116.4	C2'—C3'—C4'	110.9 (7)
C7—N2—H2	116.4	C2'—C3'—H3C	109.5
O3—N3—O2	123.5 (4)	C4'—C3'—H3C	109.5
O3—N3—C11	119.1 (4)	C2'—C3'—H3D	109.5
O2—N3—C11	117.4 (4)	C4'—C3'—H3D	109.5
O5—N4—O4	124.5 (3)	H3C—C3'—H3D	108.1
O5—N4—C13	117.9 (3)	C3'—C4'—C5'	109.0 (7)
O4—N4—C13	117.6 (3)	C3'—C4'—H4C	109.9
N1—C1—C6	108.7 (6)	C5'—C4'—H4C	109.9
N1—C1—C2	109.8 (6)	C3'—C4'—H4D	109.9
C6—C1—C2	110.6 (7)	C5'—C4'—H4D	109.9
N1—C1—H1A	109.3	H4C—C4'—H4D	108.3
C6—C1—H1A	109.3	C6'—C5'—C4'	111.1 (7)
C2—C1—H1A	109.3	C6'—C5'—H5C	109.4
C1—C2—C3	107.2 (5)	C4'—C5'—H5C	109.4
C1—C2—H2A	110.3	C6'—C5'—H5D	109.4
C3—C2—H2A	110.3	C4'—C5'—H5D	109.4
C1—C2—H2B	110.3	H5C—C5'—H5D	108.0
C3—C2—H2B	110.3	C5'—C6'—C1'	111.1 (7)
H2A—C2—H2B	108.5	C5'—C6'—H6C	109.4
C4—C3—C2	110.7 (5)	C1'—C6'—H6C	109.4
C4—C3—H3A	109.5	C5'—C6'—H6D	109.4

C2—C3—H3A	109.5	C1'—C6'—H6D	109.4
C4—C3—H3B	109.5	H6C—C6'—H6D	108.0
C2—C3—H3B	109.5	N1—C7—N2	116.4 (3)
H3A—C3—H3B	108.1	N1—C7—S1	125.6 (3)
C5—C4—C3	112.0 (6)	N2—C7—S1	118.0 (2)
C5—C4—H4A	109.2	O1—C8—N2	124.1 (3)
C3—C4—H4A	109.2	O1—C8—C9	119.7 (3)
C5—C4—H4B	109.2	N2—C8—C9	116.2 (3)
C3—C4—H4B	109.2	C10—C9—C14	120.0 (3)
H4A—C4—H4B	107.9	C10—C9—C8	117.2 (3)
C4—C5—C6	109.7 (6)	C14—C9—C8	122.8 (3)
C4—C5—H5A	109.7	C11—C10—C9	119.2 (3)
C6—C5—H5A	109.7	C11—C10—H10	120.4
C4—C5—H5B	109.7	C9—C10—H10	120.4
C6—C5—H5B	109.7	C10—C11—C12	122.6 (3)
H5A—C5—H5B	108.2	C10—C11—N3	118.8 (4)
C1—C6—C5	107.1 (5)	C12—C11—N3	118.6 (3)
C1—C6—H6A	110.3	C13—C12—C11	116.7 (3)
C5—C6—H6A	110.3	C13—C12—H12	121.6
C1—C6—H6B	110.3	C11—C12—H12	121.6
C5—C6—H6B	110.3	C12—C13—C14	123.1 (3)
H6A—C6—H6B	108.5	C12—C13—N4	119.0 (3)
N1—C1'—C6'	110.4 (8)	C14—C13—N4	117.9 (3)
N1—C1'—C2'	115.0 (8)	C13—C14—C9	118.3 (3)
C6'—C1'—C2'	109.4 (6)	C13—C14—H14	120.8
N1—C1'—H1B	107.3	C9—C14—H14	120.8
C6'—C1'—H1B	107.3		
C7—N1—C1—C6	147.6 (6)	C8—N2—C7—N1	-0.6 (5)
C1'—N1—C1—C6	134 (2)	C8—N2—C7—S1	-179.3 (3)
C7—N1—C1—C2	-91.4 (9)	C7—N2—C8—O1	0.6 (6)
C1'—N1—C1—C2	-105 (2)	C7—N2—C8—C9	-179.1 (3)
N1—C1—C2—C3	177.3 (7)	O1—C8—C9—C10	-31.7 (5)
C6—C1—C2—C3	-62.8 (10)	N2—C8—C9—C10	148.0 (3)
C1—C2—C3—C4	57.7 (10)	O1—C8—C9—C14	145.0 (4)
C2—C3—C4—C5	-56.9 (9)	N2—C8—C9—C14	-35.3 (4)
C3—C4—C5—C6	57.5 (9)	C14—C9—C10—C11	1.7 (5)
N1—C1—C6—C5	-175.5 (8)	C8—C9—C10—C11	178.6 (3)
C2—C1—C6—C5	63.9 (10)	C9—C10—C11—C12	-0.5 (5)
C4—C5—C6—C1	-59.6 (10)	C9—C10—C11—N3	179.1 (3)
C7—N1—C1'—C6'	141.7 (5)	O3—N3—C11—C10	179.4 (4)
C1—N1—C1'—C6'	-49.4 (18)	O2—N3—C11—C10	1.2 (5)
C7—N1—C1'—C2'	-94.0 (7)	O3—N3—C11—C12	-1.1 (5)
C1—N1—C1'—C2'	74.9 (19)	O2—N3—C11—C12	-179.3 (4)
N1—C1'—C2'—C3'	-66.7 (10)	C10—C11—C12—C13	-0.8 (5)
C6'—C1'—C2'—C3'	58.1 (11)	N3—C11—C12—C13	179.6 (3)
C1'—C2'—C3'—C4'	-59.8 (12)	C11—C12—C13—C14	0.9 (5)
C2'—C3'—C4'—C5'	58.6 (13)	C11—C12—C13—N4	-179.4 (3)

C3'—C4'—C5'—C6'	-57.3 (14)	O5—N4—C13—C12	172.3 (3)
C4'—C5'—C6'—C1'	57.8 (13)	O4—N4—C13—C12	-7.0 (5)
N1—C1'—C6'—C5'	69.9 (10)	O5—N4—C13—C14	-8.0 (5)
C2'—C1'—C6'—C5'	-57.5 (11)	O4—N4—C13—C14	172.6 (3)
C1—N1—C7—N2	177.5 (5)	C12—C13—C14—C9	0.3 (5)
C1'—N1—C7—N2	-177.5 (5)	N4—C13—C14—C9	-179.4 (3)
C1—N1—C7—S1	-3.8 (7)	C10—C9—C14—C13	-1.6 (5)
C1'—N1—C7—S1	1.1 (6)	C8—C9—C14—C13	-178.3 (3)

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>
N1—H1...O1	0.88	1.89	2.639 (4)	142
N1—H1'...O1	0.88	1.99	2.639 (4)	130
N2—H2...S1 ⁱ	0.88	2.65	3.449 (3)	152

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