



# Crystal structure of *N*-[(naphthalen-1-yl)-carbamothioyl]cyclohexanecarboxamide

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Received 14 June 2015; accepted 22 June 2015

Edited by W. T. A. Harrison, University of Aberdeen, Scotland

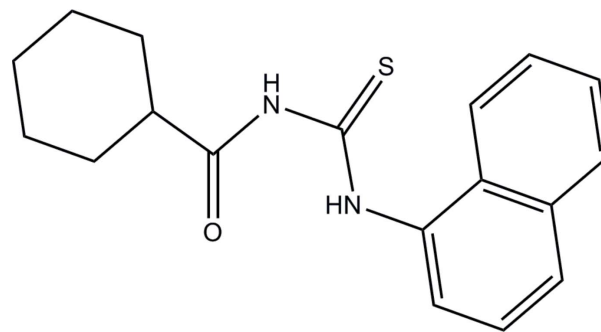
The title compound, C<sub>18</sub>H<sub>20</sub>N<sub>2</sub>OS, displays whole-molecule disorder over two adjacent sets of sites with an occupancy ratio of 0.630 (11):0.370 (11). In each disorder component, the cyclohexyl ring shows a chair conformation with the exocyclic C—C bond in an equatorial orientation. The dihedral angles between the cyclohexyl ring (all atoms) and the naphthyl ring system are 36.9 (6) for the major component and 20.7 (12)° for the minor component. Each component features an intramolecular N—H···O hydrogen bond, which closes an *S*(5) ring. In the crystal, inversion dimers linked by pairs of N—H···S hydrogen bonds generate *R*<sub>2</sub><sup>2</sup>(8) loops for both components. Aromatic  $\pi$ – $\pi$  stacking interactions [shortest centroid–centroid separation = 3.593 (9) Å] and a C—H··· $\pi$  interaction are also observed.

**Keywords:** crystal structure; whole-molecule disorder; thiourea derivatives; intramolecular N—H···O hydrogen bond; N—H···S hydrogen bonds;  $\pi$ – $\pi$  stacking interactions; C—H··· $\pi$  interactions.

**CCDC reference:** 1408027

## 1. Related literature

For background to the varied properties of thiourea derivatives, see: Sun *et al.* (2006); Shen *et al.* (2006). For related structures, see: Hu *et al.* (2011); Gangadharan *et al.* (2015).



## 2. Experimental

### 2.1. Crystal data

|   |   |
|---|---|
| C <sub>18</sub> H <sub>20</sub> N <sub>2</sub> OS | $\gamma = 104.022 (3)^\circ$              |
| $M_r = 312.42$                                    | $V = 835.24 (9) \text{ \AA}^3$            |
| Triclinic, <i>P</i> $\bar{1}$                     | $Z = 2$                                   |
| $a = 7.0464 (5) \text{ \AA}$                      | Mo $K\alpha$ radiation                    |
| $b = 11.0379 (5) \text{ \AA}$                     | $\mu = 0.20 \text{ mm}^{-1}$              |
| $c = 12.4151 (8) \text{ \AA}$                     | $T = 293 \text{ K}$                       |
| $\alpha = 110.873 (3)^\circ$                      | $0.35 \times 0.30 \times 0.25 \text{ mm}$ |
| $\beta = 100.660 (3)^\circ$                       |   |

### 2.2. Data collection

|  |  |
|--|--|
| Bruker APEXII CCD diffractometer                         | 14064 measured reflections             |
| Absorption correction: multi-scan (SADABS; Bruker, 2004) | 2945 independent reflections           |
| $T_{\min} = 0.934$ , $T_{\max} = 0.952$                  | 1930 reflections with $I > 2\sigma(I)$ |
|  | $R_{\text{int}} = 0.031$               |

### 2.3. Refinement

|                                 |  |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.068$ | H atoms treated by a mixture of independent and constrained refinement |
| $wR(F^2) = 0.219$               | $\Delta\rho_{\text{max}} = 0.87 \text{ e \AA}^{-3}$                    |
| $S = 1.05$                      | $\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$                   |
| 2945 reflections                |  |
| 404 parameters                  |  |
| 1056 restraints                 |  |

**Table 1**

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C1–C5/C10 ring.

| <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> |
|------------------------------|-------------|---------------|-----------------------|-------------------------|
| N2—H2A···O1                  | 0.88 (4)    | 1.97 (4)      | 2.667 (15)            | 135 (4)                 |
| N2'—H2A···O1'                | 0.93 (4)    | 2.03 (4)      | 2.62 (3)              | 120 (4)                 |
| N1—H1A···S1 <sup>i</sup>     | 0.87 (3)    | 2.53 (3)      | 3.370 (19)            | 161 (4)                 |
| N1'—H1A···S1 <sup>ii</sup>   | 0.90 (4)    | 2.59 (4)      | 3.44 (3)              | 159 (4)                 |
| C18—H18B···Cg1 <sup>ii</sup> | 0.90        | 2.66          | 3.527 (2)             | 148                     |

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

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 for Windows (Farrugia, 2012); software used to prepare material for publication: PLATON (Spek, 2009).

## Acknowledgements

The authors thank Dr Babu Varghese, SAIF, IIT, Chennai, India, for the data collection. JH thanks UGC for the fellowship.

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7448).

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## supporting information

*Acta Cryst.* (2015). E71, o508–o509 [doi:10.1107/S2056989015011950]

## Crystal structure of *N*-[(naphthalen-1-yl)carbamothioyl]cyclohexane-carboxamide

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### S1. Comment

The design and synthesis of thiourea are of considerable interest because of their use in agriculture, medicine and analytical chemistry (J-H. Hu *et al.*, 2011). Thiourea derivatives are driven by their potential as biological active compounds (Sun *et al.*, 2006) and in the material application such as anti corrosion (Shen *et al.*, 2006). As part of our own studies in this area, the crystal structure of the title compound has been determined and the results are presented herein.

For the major disorder component, the cyclohexane ring (C13—C18) adopts chair conformation [puckering amplitudes and smallest displacement parameters are  $q = 0.568 \text{ \AA}$ ,  $\theta = 177.7 (8)^\circ$ ,  $\varphi = 19 (29)^\circ$  and  $\Delta C_s = 0.9 (14) \text{ \AA}$ ]. Similarly, for the minor disorder component, the cyclohexane ring (C13'—C18') adopts a chair conformation. [puckering amplitudes and smallest displacement parameters are  $q = 0.56 (3) \text{ \AA}$ ,  $\theta = 180 (3)^\circ$ ,  $\varphi = 354 (31)^\circ$  and  $\Delta C_s = 3.0 (4) \text{ \AA}$ ]. The dihedral angles between cyclohexane and benzene rings (C5—C8/C10) and (C5—C10) of naphthalene moiety are  $37.0 (7)$  and  $36.5 (7)^\circ$  (major component). In the case of minor component, the dihedral angles between cyclohexane and benzene rings (C5—C8/C10) and (C5—C10) of naphthalene are  $37.0 (7)$  and  $36.5 (7)^\circ$ , respectively. The molecular conformation is consolidated by an intramolecular N—H $\cdots$ O hydrogen bond, forming S(5) ring motif.

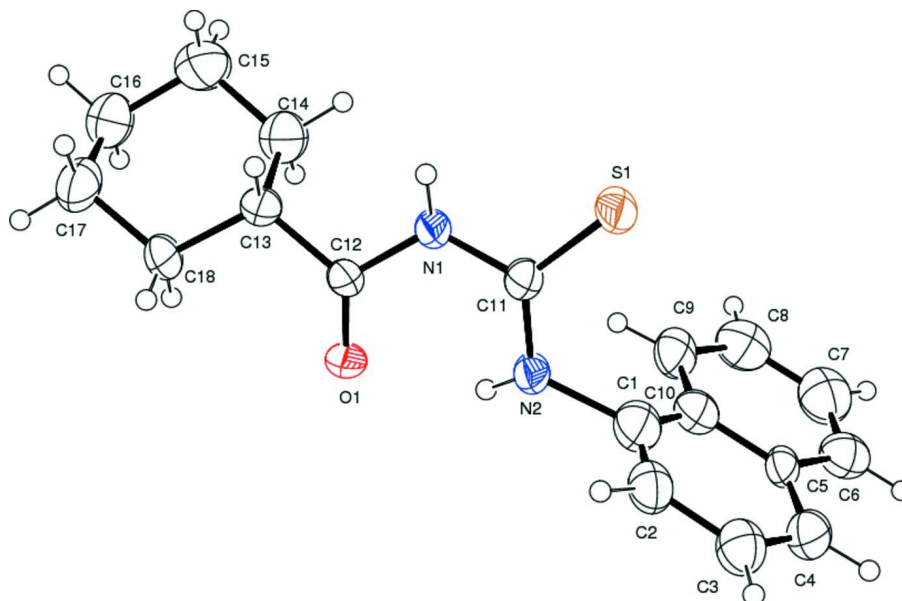
The crystal packing features N—H $\cdots$ S hydrogen bonds with the symmetry code: (i)  $2 - x, -y, 1 - z$ , which links the molecules into centrosymmetric dimers with graph-set descriptor of  $R^2_2(8)$ . The crystal packing also features C—H $\cdots\pi$  (Table 1) and  $\pi$ - $\pi$  interactions ( $Cg2 \cdots Cg2^{ii} = 3.593 (9) \text{ \AA}$ ;  $Cg2$  is the centroid of a ring C5—C10; symmetry code: (ii)  $3 - x, 1 - y, 2 - z$ ). The packing view of the title compound is shown in Fig. 3.

### S2. Experimental

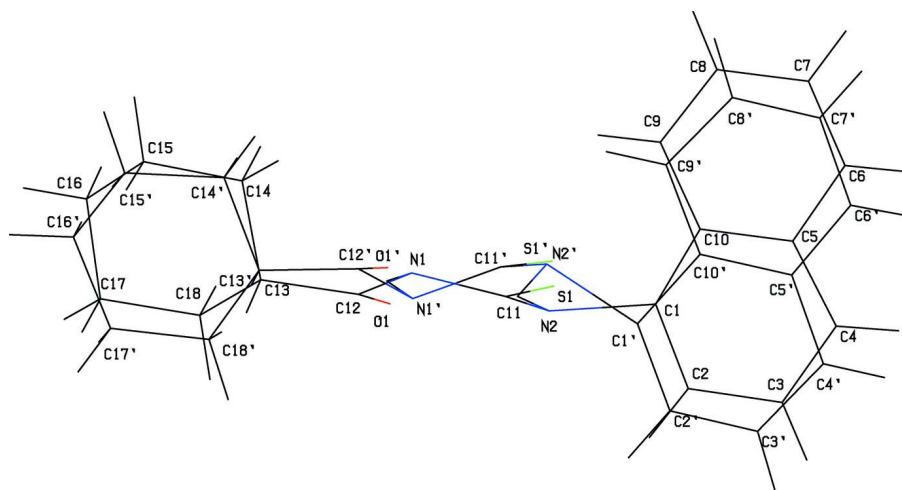
A solution of cyclohexane carbonyl chloride (1.4661 g, 10 mmol) in acetone (60 ml) was added drop wise to a suspension of potassium thiocyanate (0.9718 g, 10 mmol) in anhydrous acetone (60 ml). The reaction mixture was heated under reflux for 45 minutes and then cooled to room temperature. A solution of substituted naphthalen-1-amine (1.43 g, 10 mmol) in acetone (60 ml) was added and the resulting mixture was stirred for 2 h at room temperature. Hydrochloric acid (0.1 N, 500 ml) was added and the resulting white solid was filtered off, washed with water and dried in vacuum. The yield of the isolated product was 89%, giving colourless blocks.

### S3. Refinement

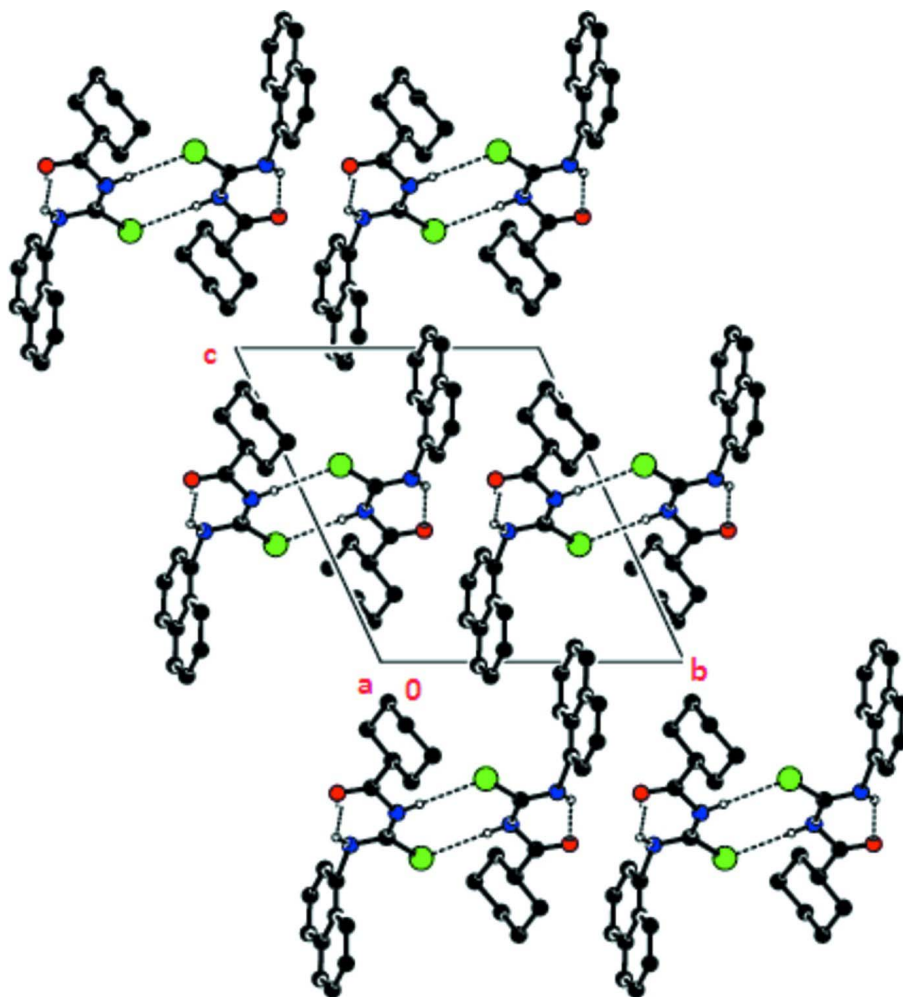
All H atoms were fixed geometrically and allowed to ride on their parent C atoms, with C—H distances fixed in the range  $0.93$ – $0.97 \text{ \AA}$  with  $U_{iso}(H) = 1.5U_{eq}(C)$  for methyl H atoms and  $1.2U_{eq}(C)$  for all other H atoms.

**Figure 1**

The molecular structure of the major component of the title compound, with displacement ellipsoids drawn at 40% probability level.

**Figure 2**

Stick plot of both major and minor components of the title compound, with the atoms label for non-H atoms.



**Figure 3**

The crystal packing of the title compound, viewed along the *b* axis. The hydrogen bonds are shown as dashed lines (see Table 1 for details).

### *N*-[(Naphthalen-1-yl)carbamothioyl]cyclohexanecarboxamide

#### Crystal data

$C_{18}H_{20}N_2OS$

$M_r = 312.42$

Triclinic,  $P\bar{1}$

Hall symbol:  $-P\ 1$

$a = 7.0464$  (5) Å

$b = 11.0379$  (5) Å

$c = 12.4151$  (8) Å

$\alpha = 110.873$  (3)°

$\beta = 100.660$  (3)°

$\gamma = 104.022$  (3)°

$V = 835.24$  (9) Å<sup>3</sup>

$Z = 2$

$F(000) = 332$

$D_x = 1.242$  Mg m<sup>-3</sup>

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

Cell parameters from 1930 reflections

$\theta = 2.1$ – $25.0$ °

$\mu = 0.20$  mm<sup>-1</sup>

$T = 293$  K

Block, white

$0.35 \times 0.30 \times 0.25$  mm

Data collection

|   |  |
|---|--|
| Bruker APEXII CCD diffractometer                            | 14064 measured reflections   |
| Radiation source: fine-focus sealed tube                    | 2945 independent reflections   |
| Graphite monochromator                                      | 1930 reflections with $I > 2\sigma(I)$                                 |
| $\omega$ and $\varphi$ scan                                 | $R_{\text{int}} = 0.031$   |
| Absorption correction: multi-scan<br>(SADABS; Bruker, 2004) | $\theta_{\text{max}} = 25.0^\circ$ , $\theta_{\text{min}} = 2.1^\circ$ |
| $T_{\text{min}} = 0.934$ , $T_{\text{max}} = 0.952$         | $h = -8 \rightarrow 8$   |
|   | $k = -13 \rightarrow 13$   |
|   | $l = -14 \rightarrow 14$   |

Refinement

|  |  |
|--|--|
| Refinement on $F^2$  | Secondary atom site location: difference Fourier map                   |
| Least-squares matrix: full                                     | Hydrogen site location: inferred from neighbouring sites               |
| $R[F^2 > 2\sigma(F^2)] = 0.068$                                | H atoms treated by a mixture of independent and constrained refinement |
| $wR(F^2) = 0.219$  | $w = 1/[\sigma^2(F_o^2) + (0.0972P)^2 + 1.1049P]$                      |
| $S = 1.05$   | where $P = (F_o^2 + 2F_c^2)/3$   |
| 2945 reflections   | $(\Delta/\sigma)_{\text{max}} = 0.053$                                 |
| 404 parameters   | $\Delta\rho_{\text{max}} = 0.87 \text{ e } \text{\AA}^{-3}$            |
| 1056 restraints  | $\Delta\rho_{\text{min}} = -0.25 \text{ e } \text{\AA}^{-3}$           |
| Primary atom site location: structure-invariant direct methods |  |

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 ( $\text{\AA}^2$ )

|    | $x$         | $y$         | $z$         | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1)  |
|----|-------------|-------------|-------------|----------------------------------|------------|
| C1 | 1.4139 (12) | 0.4829 (15) | 0.6995 (10) | 0.072 (3)                        | 0.630 (11) |
| C2 | 1.5803 (14) | 0.5008 (15) | 0.6610 (8)  | 0.058 (2)                        | 0.630 (11) |
| H2 | 1.5690      | 0.4550      | 0.5798      | 0.069*                           | 0.630 (11) |
| C3 | 1.7732 (13) | 0.5902 (11) | 0.7454 (8)  | 0.064 (2)                        | 0.630 (11) |
| H3 | 1.8908      | 0.6065      | 0.7214      | 0.076*                           | 0.630 (11) |
| C4 | 1.7793 (14) | 0.6525 (12) | 0.8658 (8)  | 0.065 (2)                        | 0.630 (11) |
| H4 | 1.9016      | 0.7111      | 0.9257      | 0.078*                           | 0.630 (11) |
| C5 | 1.5918 (13) | 0.6230 (12) | 0.8936 (7)  | 0.0446 (17)                      | 0.630 (11) |
| C6 | 1.5952 (14) | 0.6835 (11) | 1.0116 (8)  | 0.062 (2)                        | 0.630 (11) |
| H6 | 1.7193      | 0.7418      | 1.0700      | 0.075*                           | 0.630 (11) |
| C7 | 1.4217 (16) | 0.6606 (16) | 1.0460 (8)  | 0.082 (3)                        | 0.630 (11) |
| H7 | 1.4294      | 0.7032      | 1.1270      | 0.099*                           | 0.630 (11) |
| C8 | 1.2357 (16) | 0.5753 (16) | 0.9622 (9)  | 0.076 (3)                        | 0.630 (11) |
| H8 | 1.1199      | 0.5574      | 0.9875      | 0.091*                           | 0.630 (11) |
| C9 | 1.2189 (15) | 0.5164 (15) | 0.8419 (8)  | 0.062 (2)                        | 0.630 (11) |

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|      |             |              |             |             |            |
|------|-------------|--------------|-------------|-------------|------------|
| H9   | 1.0919      | 0.4657       | 0.7838      | 0.074*      | 0.630 (11) |
| C10  | 1.4035 (19) | 0.536 (2)    | 0.8090 (10) | 0.063 (3)   | 0.630 (11) |
| C11  | 1.1480 (17) | 0.2490 (13)  | 0.5594 (10) | 0.042 (2)   | 0.630 (11) |
| C12  | 0.839 (2)   | 0.2236 (13)  | 0.4017 (11) | 0.042 (3)   | 0.630 (11) |
| C13  | 0.643 (2)   | 0.113 (2)    | 0.3153 (11) | 0.052 (3)   | 0.630 (11) |
| H13  | 0.6743      | 0.0287       | 0.2776      | 0.062*      | 0.630 (11) |
| C14  | 0.4915 (15) | 0.0832 (11)  | 0.3824 (10) | 0.067 (2)   | 0.630 (11) |
| H14A | 0.4669      | 0.1674       | 0.4258      | 0.080*      | 0.630 (11) |
| H14B | 0.5514      | 0.0538       | 0.4413      | 0.080*      | 0.630 (11) |
| C15  | 0.2882 (18) | −0.0269 (11) | 0.2994 (11) | 0.084 (3)   | 0.630 (11) |
| H15A | 0.1929      | −0.0367      | 0.3455      | 0.101*      | 0.630 (11) |
| H15B | 0.3083      | −0.1146      | 0.2629      | 0.101*      | 0.630 (11) |
| C16  | 0.2014 (19) | 0.0145 (14)  | 0.2016 (11) | 0.088 (3)   | 0.630 (11) |
| H16A | 0.0733      | −0.0568      | 0.1475      | 0.106*      | 0.630 (11) |
| H16B | 0.1720      | 0.0986       | 0.2389      | 0.106*      | 0.630 (11) |
| C17  | 0.3457 (16) | 0.0371 (10)  | 0.1289 (11) | 0.074 (3)   | 0.630 (11) |
| H17A | 0.2864      | 0.0664       | 0.0697      | 0.088*      | 0.630 (11) |
| H17B | 0.3691      | −0.0479      | 0.0865      | 0.088*      | 0.630 (11) |
| C18  | 0.5478 (15) | 0.1477 (9)   | 0.2156 (9)  | 0.059 (3)   | 0.630 (11) |
| H18A | 0.6436      | 0.1599       | 0.1703      | 0.071*      | 0.630 (11) |
| H18B | 0.5240      | 0.2342       | 0.2514      | 0.071*      | 0.630 (11) |
| S1   | 1.2576 (18) | 0.1673 (11)  | 0.6264 (10) | 0.0567 (16) | 0.630 (11) |
| N1   | 0.938 (2)   | 0.1781 (12)  | 0.4817 (12) | 0.044 (3)   | 0.630 (11) |
| N2   | 1.2224 (9)  | 0.3819 (7)   | 0.5845 (5)  | 0.0416 (16) | 0.630 (11) |
| O1   | 0.886 (2)   | 0.3438 (12)  | 0.4191 (12) | 0.055 (3)   | 0.630 (11) |
| C1'  | 1.4044 (16) | 0.4630 (19)  | 0.6642 (11) | 0.032 (3)   | 0.370 (11) |
| C2'  | 1.574 (2)   | 0.480 (3)    | 0.6233 (13) | 0.063 (4)   | 0.370 (11) |
| H2'  | 1.5553      | 0.4330       | 0.5412      | 0.076*      | 0.370 (11) |
| C3'  | 1.7640 (19) | 0.5594 (19)  | 0.6957 (14) | 0.060 (4)   | 0.370 (11) |
| H3'  | 1.8726      | 0.5649       | 0.6630      | 0.072*      | 0.370 (11) |
| C4'  | 1.802 (2)   | 0.635 (2)    | 0.8205 (14) | 0.070 (4)   | 0.370 (11) |
| H4'  | 1.9328      | 0.6931       | 0.8711      | 0.084*      | 0.370 (11) |
| C5'  | 1.632 (3)   | 0.618 (3)    | 0.8638 (14) | 0.067 (4)   | 0.370 (11) |
| C6'  | 1.656 (3)   | 0.682 (3)    | 0.9838 (16) | 0.097 (5)   | 0.370 (11) |
| H6'  | 1.7861      | 0.7371       | 1.0373      | 0.117*      | 0.370 (11) |
| C7'  | 1.489 (3)   | 0.666 (3)    | 1.0264 (14) | 0.077 (5)   | 0.370 (11) |
| H7'  | 1.5100      | 0.7078       | 1.1092      | 0.092*      | 0.370 (11) |
| C8'  | 1.298 (3)   | 0.593 (3)    | 0.9538 (14) | 0.070 (4)   | 0.370 (11) |
| H8'  | 1.1874      | 0.5978       | 0.9844      | 0.084*      | 0.370 (11) |
| C9'  | 1.261 (3)   | 0.507 (3)    | 0.8293 (12) | 0.076 (5)   | 0.370 (11) |
| H9'  | 1.1367      | 0.4381       | 0.7805      | 0.091*      | 0.370 (11) |
| C10' | 1.434 (2)   | 0.536 (3)    | 0.7873 (12) | 0.037 (3)   | 0.370 (11) |
| C11' | 1.100 (3)   | 0.253 (2)    | 0.5789 (19) | 0.049 (4)   | 0.370 (11) |
| C12' | 0.809 (3)   | 0.221 (2)    | 0.424 (2)   | 0.042 (4)   | 0.370 (11) |
| C13' | 0.629 (3)   | 0.110 (4)    | 0.3220 (19) | 0.051 (4)   | 0.370 (11) |
| H13' | 0.6531      | 0.0227       | 0.3069      | 0.061*      | 0.370 (11) |
| C14' | 0.444 (3)   | 0.103 (3)    | 0.364 (2)   | 0.096 (5)   | 0.370 (11) |
| H14C | 0.4248      | 0.1923       | 0.3884      | 0.116*      | 0.370 (11) |

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|      |             |             |             |           |            |
|------|-------------|-------------|-------------|-----------|------------|
| H14D | 0.4647      | 0.0828      | 0.4344      | 0.116*    | 0.370 (11) |
| C15' | 0.256 (4)   | -0.005 (3)  | 0.2673 (18) | 0.102 (6) | 0.370 (11) |
| H15C | 0.2702      | -0.0944     | 0.2488      | 0.123*    | 0.370 (11) |
| H15D | 0.1378      | -0.0039     | 0.2970      | 0.123*    | 0.370 (11) |
| C16' | 0.219 (3)   | 0.017 (2)   | 0.1542 (17) | 0.081 (5) | 0.370 (11) |
| H16C | 0.1000      | -0.0564     | 0.0919      | 0.098*    | 0.370 (11) |
| H16D | 0.1951      | 0.1042      | 0.1694      | 0.098*    | 0.370 (11) |
| C17' | 0.408 (3)   | 0.018 (2)   | 0.116 (2)   | 0.090 (6) | 0.370 (11) |
| H17C | 0.3886      | 0.0311      | 0.0420      | 0.108*    | 0.370 (11) |
| H17D | 0.4230      | -0.0711     | 0.0981      | 0.108*    | 0.370 (11) |
| C18' | 0.602 (3)   | 0.126 (2)   | 0.2064 (17) | 0.083 (5) | 0.370 (11) |
| H18C | 0.7176      | 0.1164      | 0.1762      | 0.099*    | 0.370 (11) |
| H18D | 0.5957      | 0.2172      | 0.2197      | 0.099*    | 0.370 (11) |
| S1'  | 1.222 (3)   | 0.1767 (19) | 0.6457 (17) | 0.065 (4) | 0.370 (11) |
| N2'  | 1.1590 (19) | 0.3881 (14) | 0.6219 (11) | 0.083 (4) | 0.370 (11) |
| N1'  | 0.972 (3)   | 0.176 (2)   | 0.461 (2)   | 0.036 (4) | 0.370 (11) |
| O1'  | 0.838 (4)   | 0.341 (2)   | 0.448 (2)   | 0.057 (4) | 0.370 (11) |
| H1A  | 0.918 (6)   | 0.0899 (17) | 0.454 (4)   | 0.069*    |            |
| H2A  | 1.129 (5)   | 0.415 (4)   | 0.560 (3)   | 0.069*    |            |

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

|     | $U^{11}$  | $U^{22}$  | $U^{33}$  | $U^{12}$    | $U^{13}$     | $U^{23}$  |
|-----|-----------|-----------|-----------|-------------|--------------|-----------|
| C1  | 0.079 (5) | 0.052 (5) | 0.075 (7) | 0.020 (4)   | -0.001 (5)   | 0.030 (6) |
| C2  | 0.071 (4) | 0.050 (5) | 0.046 (5) | 0.024 (3)   | 0.007 (4)    | 0.017 (4) |
| C3  | 0.072 (4) | 0.066 (6) | 0.051 (6) | 0.026 (4)   | 0.016 (4)    | 0.021 (5) |
| C4  | 0.054 (4) | 0.058 (5) | 0.075 (5) | 0.014 (3)   | -0.001 (4)   | 0.034 (5) |
| C5  | 0.052 (4) | 0.031 (3) | 0.039 (4) | 0.011 (3)   | -0.008 (3)   | 0.013 (3) |
| C6  | 0.070 (5) | 0.057 (4) | 0.070 (4) | 0.028 (5)   | 0.022 (4)    | 0.033 (4) |
| C7  | 0.104 (7) | 0.077 (5) | 0.068 (5) | 0.042 (6)   | 0.022 (5)    | 0.027 (4) |
| C8  | 0.084 (6) | 0.081 (6) | 0.080 (5) | 0.037 (5)   | 0.036 (4)    | 0.043 (4) |
| C9  | 0.064 (5) | 0.056 (4) | 0.066 (4) | 0.035 (4)   | 0.004 (3)    | 0.024 (3) |
| C10 | 0.074 (5) | 0.051 (4) | 0.063 (5) | 0.017 (4)   | 0.005 (4)    | 0.032 (4) |
| C11 | 0.047 (5) | 0.037 (3) | 0.035 (4) | 0.011 (3)   | 0.006 (3)    | 0.012 (3) |
| C12 | 0.053 (5) | 0.037 (3) | 0.032 (4) | 0.008 (3)   | 0.007 (3)    | 0.017 (3) |
| C13 | 0.058 (4) | 0.036 (4) | 0.053 (4) | 0.009 (4)   | -0.001 (4)   | 0.022 (3) |
| C14 | 0.058 (5) | 0.069 (4) | 0.070 (5) | 0.021 (3)   | 0.011 (4)    | 0.031 (3) |
| C15 | 0.066 (5) | 0.082 (5) | 0.099 (6) | 0.007 (4)   | 0.013 (5)    | 0.049 (4) |
| C16 | 0.063 (5) | 0.092 (5) | 0.089 (7) | 0.005 (4)   | -0.012 (5)   | 0.046 (5) |
| C17 | 0.072 (6) | 0.056 (4) | 0.070 (5) | 0.018 (4)   | -0.013 (5)   | 0.020 (4) |
| C18 | 0.066 (5) | 0.050 (4) | 0.055 (4) | 0.016 (3)   | -0.009 (3)   | 0.031 (3) |
| S1  | 0.069 (3) | 0.035 (2) | 0.054 (3) | 0.0193 (14) | -0.0060 (17) | 0.017 (2) |
| N1  | 0.056 (5) | 0.031 (3) | 0.036 (5) | 0.009 (3)   | -0.001 (4)   | 0.013 (3) |
| N2  | 0.047 (3) | 0.033 (3) | 0.038 (3) | 0.013 (2)   | 0.001 (2)    | 0.014 (2) |
| O1  | 0.060 (6) | 0.036 (3) | 0.059 (6) | 0.002 (3)   | -0.004 (3)   | 0.028 (3) |
| C1' | 0.049 (5) | 0.025 (5) | 0.015 (4) | 0.012 (4)   | -0.004 (3)   | 0.007 (4) |
| C2' | 0.085 (6) | 0.051 (7) | 0.041 (7) | 0.021 (5)   | 0.001 (5)    | 0.014 (6) |
| C3' | 0.053 (6) | 0.057 (7) | 0.063 (8) | 0.018 (5)   | 0.010 (6)    | 0.020 (7) |



|      |            |           |           |            |            |           |
|------|------------|-----------|-----------|------------|------------|-----------|
| C4'  | 0.080 (7)  | 0.060 (7) | 0.057 (9) | 0.026 (6)  | 0.004 (7)  | 0.018 (8) |
| C5'  | 0.076 (7)  | 0.051 (5) | 0.060 (7) | 0.019 (6)  | 0.001 (6)  | 0.020 (6) |
| C6'  | 0.103 (9)  | 0.075 (7) | 0.090 (8) | 0.025 (8)  | -0.001 (7) | 0.027 (7) |
| C7'  | 0.084 (10) | 0.075 (7) | 0.080 (7) | 0.032 (9)  | 0.027 (7)  | 0.038 (6) |
| C8'  | 0.079 (8)  | 0.074 (7) | 0.063 (6) | 0.028 (7)  | 0.030 (6)  | 0.030 (5) |
| C9'  | 0.058 (7)  | 0.055 (6) | 0.112 (7) | 0.018 (7)  | 0.004 (6)  | 0.044 (6) |
| C10' | 0.049 (5)  | 0.022 (4) | 0.025 (5) | 0.007 (4)  | -0.011 (4) | 0.006 (5) |
| C11' | 0.056 (7)  | 0.032 (5) | 0.050 (7) | 0.023 (5)  | -0.002 (6) | 0.011 (5) |
| C12' | 0.053 (6)  | 0.033 (5) | 0.040 (7) | 0.008 (5)  | 0.007 (5)  | 0.022 (5) |
| C13' | 0.053 (6)  | 0.041 (6) | 0.052 (6) | 0.010 (6)  | -0.003 (6) | 0.024 (5) |
| C14' | 0.073 (8)  | 0.101 (8) | 0.084 (7) | 0.009 (7)  | 0.012 (6)  | 0.024 (7) |
| C15' | 0.079 (8)  | 0.104 (8) | 0.100 (9) | -0.005 (7) | 0.013 (7)  | 0.047 (7) |
| C16' | 0.069 (7)  | 0.072 (6) | 0.082 (9) | 0.011 (6)  | -0.005 (8) | 0.031 (7) |
| C17' | 0.074 (8)  | 0.082 (8) | 0.074 (7) | 0.019 (7)  | -0.011 (7) | 0.008 (6) |
| C18' | 0.064 (8)  | 0.083 (8) | 0.066 (7) | 0.014 (6)  | -0.001 (6) | 0.010 (6) |
| S1'  | 0.074 (7)  | 0.033 (2) | 0.060 (6) | 0.020 (3)  | -0.016 (4) | 0.006 (3) |
| N2'  | 0.097 (7)  | 0.040 (5) | 0.065 (7) | 0.020 (6)  | -0.041 (5) | 0.006 (5) |
| N1'  | 0.045 (6)  | 0.028 (4) | 0.031 (6) | 0.011 (4)  | 0.009 (4)  | 0.009 (4) |
| O1'  | 0.061 (10) | 0.041 (5) | 0.058 (9) | 0.007 (6)  | -0.003 (6) | 0.025 (5) |

*Geometric parameters (Å, °)*

|         |            |           |            |
|---------|------------|-----------|------------|
| C1—C10  | 1.298 (10) | C1'—C2'   | 1.378 (11) |
| C1—C2   | 1.344 (9)  | C1'—C10'  | 1.396 (12) |
| C1—N2   | 1.589 (8)  | C1'—N2'   | 1.616 (10) |
| C2—C3   | 1.416 (9)  | C2'—C3'   | 1.341 (13) |
| C2—H2   | 0.9300     | C2'—H2'   | 0.9300     |
| C3—C4   | 1.392 (8)  | C3'—C4'   | 1.408 (11) |
| C3—H3   | 0.9300     | C3'—H3'   | 0.9300     |
| C4—C5   | 1.418 (9)  | C4'—C5'   | 1.399 (14) |
| C4—H4   | 0.9300     | C4'—H4'   | 0.9300     |
| C5—C6   | 1.370 (9)  | C5'—C6'   | 1.359 (14) |
| C5—C10  | 1.392 (11) | C5'—C10'  | 1.399 (15) |
| C6—C7   | 1.364 (9)  | C6'—C7'   | 1.378 (12) |
| C6—H6   | 0.9300     | C6'—H6'   | 0.9300     |
| C7—C8   | 1.373 (10) | C7'—C8'   | 1.336 (14) |
| C7—H7   | 0.9300     | C7'—H7'   | 0.9300     |
| C8—C9   | 1.369 (9)  | C8'—C9'   | 1.432 (13) |
| C8—H8   | 0.9300     | C8'—H8'   | 0.9300     |
| C9—C10  | 1.424 (10) | C9'—C10'  | 1.422 (14) |
| C9—H9   | 0.9300     | C9'—H9'   | 0.9300     |
| C11—N2  | 1.329 (14) | C11'—N2'  | 1.31 (2)   |
| C11—N1  | 1.451 (19) | C11'—N1'  | 1.40 (3)   |
| C11—S1  | 1.653 (6)  | C11'—S1'  | 1.652 (8)  |
| C12—O1  | 1.214 (6)  | C12'—O1'  | 1.214 (8)  |
| C12—N1  | 1.411 (9)  | C12'—N1'  | 1.413 (11) |
| C12—C13 | 1.502 (7)  | C12'—C13' | 1.500 (9)  |
| C13—C18 | 1.511 (10) | C13'—C14' | 1.490 (14) |

|           |            |              |            |
|-----------|------------|--------------|------------|
| C13—C14   | 1.512 (8)  | C13'—C18'    | 1.491 (13) |
| C13—H13   | 0.9800     | C13'—H13'    | 0.9800     |
| C14—C15   | 1.520 (8)  | C14'—C15'    | 1.502 (13) |
| C14—H14A  | 0.9700     | C14'—H14C    | 0.9700     |
| C14—H14B  | 0.9700     | C14'—H14D    | 0.9700     |
| C15—C16   | 1.521 (8)  | C15'—C16'    | 1.495 (13) |
| C15—H15A  | 0.9700     | C15'—H15C    | 0.9700     |
| C15—H15B  | 0.9700     | C15'—H15D    | 0.9700     |
| C16—C17   | 1.512 (9)  | C16'—C17'    | 1.492 (14) |
| C16—H16A  | 0.9700     | C16'—H16C    | 0.9700     |
| C16—H16B  | 0.9700     | C16'—H16D    | 0.9700     |
| C17—C18   | 1.526 (8)  | C17'—C18'    | 1.498 (13) |
| C17—H17A  | 0.9700     | C17'—H17C    | 0.9700     |
| C17—H17B  | 0.9700     | C17'—H17D    | 0.9700     |
| C18—H18A  | 0.9700     | C18'—H18C    | 0.9700     |
| C18—H18B  | 0.9700     | C18'—H18D    | 0.9700     |
| N1—H1A    | 0.874 (19) | N2'—H2A      | 0.927 (18) |
| N2—H2A    | 0.883 (18) | N1'—H1A      | 0.906 (19) |
|           |            |              |            |
| C10—C1—C2 | 128.3 (8)  | C2'—C1'—C10' | 117.6 (9)  |
| C10—C1—N2 | 124.4 (8)  | C2'—C1'—N2'  | 143.7 (10) |
| C2—C1—N2  | 107.4 (7)  | C10'—C1'—N2' | 98.7 (8)   |
| C1—C2—C3  | 119.2 (6)  | C3'—C2'—C1'  | 123.4 (8)  |
| C1—C2—H2  | 120.4      | C3'—C2'—H2'  | 118.3      |
| C3—C2—H2  | 120.4      | C1'—C2'—H2'  | 118.3      |
| C4—C3—C2  | 117.2 (6)  | C2'—C3'—C4'  | 121.2 (8)  |
| C4—C3—H3  | 121.4      | C2'—C3'—H3'  | 119.4      |
| C2—C3—H3  | 121.4      | C4'—C3'—H3'  | 119.4      |
| C3—C4—C5  | 117.5 (6)  | C5'—C4'—C3'  | 116.1 (10) |
| C3—C4—H4  | 121.3      | C5'—C4'—H4'  | 122.0      |
| C5—C4—H4  | 121.3      | C3'—C4'—H4'  | 121.9      |
| C6—C5—C10 | 117.4 (6)  | C6'—C5'—C10' | 117.6 (11) |
| C6—C5—C4  | 118.2 (6)  | C6'—C5'—C4'  | 120.2 (12) |
| C10—C5—C4 | 124.4 (6)  | C10'—C5'—C4' | 122.3 (10) |
| C7—C6—C5  | 121.9 (7)  | C5'—C6'—C7'  | 120.2 (12) |
| C7—C6—H6  | 119.0      | C5'—C6'—H6'  | 119.9      |
| C5—C6—H6  | 119.0      | C7'—C6'—H6'  | 119.9      |
| C6—C7—C8  | 120.6 (6)  | C8'—C7'—C6'  | 122.9 (9)  |
| C6—C7—H7  | 119.7      | C8'—C7'—H7'  | 118.6      |
| C8—C7—H7  | 119.7      | C6'—C7'—H7'  | 118.5      |
| C9—C8—C7  | 120.7 (6)  | C7'—C8'—C9'  | 120.3 (9)  |
| C9—C8—H8  | 119.6      | C7'—C8'—H8'  | 119.8      |
| C7—C8—H8  | 119.6      | C9'—C8'—H8'  | 119.9      |
| C8—C9—C10 | 117.4 (7)  | C10'—C9'—C8' | 113.6 (10) |
| C8—C9—H9  | 121.3      | C10'—C9'—H9' | 123.2      |
| C10—C9—H9 | 121.3      | C8'—C9'—H9'  | 123.2      |
| C1—C10—C5 | 113.4 (8)  | C5'—C10'—C1' | 119.3 (10) |
| C1—C10—C9 | 124.9 (9)  | C5'—C10'—C9' | 123.6 (10) |

|               |            |                |            |
|---------------|------------|----------------|------------|
| C5—C10—C9     | 121.6 (7)  | C1'—C10'—C9'   | 116.6 (11) |
| N2—C11—N1     | 115.4 (9)  | N2'—C11'—N1'   | 118.4 (15) |
| N2—C11—S1     | 125.6 (9)  | N2'—C11'—S1'   | 120.8 (16) |
| N1—C11—S1     | 118.1 (10) | N1'—C11'—S1'   | 118.6 (18) |
| O1—C12—N1     | 122.6 (12) | O1'—C12'—N1'   | 121 (2)    |
| O1—C12—C13    | 125.0 (14) | O1'—C12'—C13'  | 121 (2)    |
| N1—C12—C13    | 110.6 (11) | N1'—C12'—C13'  | 115 (2)    |
| C12—C13—C18   | 112.6 (11) | C14'—C13'—C18' | 113 (2)    |
| C12—C13—C14   | 110.4 (10) | C14'—C13'—C12' | 108 (2)    |
| C18—C13—C14   | 110.4 (11) | C18'—C13'—C12' | 114 (2)    |
| C12—C13—H13   | 107.7      | C14'—C13'—H13' | 107.2      |
| C18—C13—H13   | 107.8      | C18'—C13'—H13' | 107.2      |
| C14—C13—H13   | 107.7      | C12'—C13'—H13' | 107.2      |
| C13—C14—C15   | 112.9 (10) | C13'—C14'—C15' | 111 (2)    |
| C13—C14—H14A  | 109.0      | C13'—C14'—H14C | 109.3      |
| C15—C14—H14A  | 109.0      | C15'—C14'—H14C | 109.3      |
| C13—C14—H14B  | 109.0      | C13'—C14'—H14D | 109.4      |
| C15—C14—H14B  | 109.0      | C15'—C14'—H14D | 109.3      |
| H14A—C14—H14B | 107.8      | H14C—C14'—H14D | 108.0      |
| C14—C15—C16   | 109.0 (8)  | C16'—C15'—C14' | 112.4 (17) |
| C14—C15—H15A  | 109.9      | C16'—C15'—H15C | 109.1      |
| C16—C15—H15A  | 109.9      | C14'—C15'—H15C | 109.1      |
| C14—C15—H15B  | 109.9      | C16'—C15'—H15D | 109.1      |
| C16—C15—H15B  | 109.9      | C14'—C15'—H15D | 109.1      |
| H15A—C15—H15B | 108.3      | H15C—C15'—H15D | 107.9      |
| C17—C16—C15   | 112.9 (11) | C17'—C16'—C15' | 106 (2)    |
| C17—C16—H16A  | 109.0      | C17'—C16'—H16C | 110.6      |
| C15—C16—H16A  | 109.0      | C15'—C16'—H16C | 110.6      |
| C17—C16—H16B  | 109.0      | C17'—C16'—H16D | 110.5      |
| C15—C16—H16B  | 109.0      | C15'—C16'—H16D | 110.6      |
| H16A—C16—H16B | 107.8      | H16C—C16'—H16D | 108.7      |
| C16—C17—C18   | 108.3 (9)  | C16'—C17'—C18' | 115.5 (19) |
| C16—C17—H17A  | 110.0      | C16'—C17'—H17C | 108.4      |
| C18—C17—H17A  | 110.0      | C18'—C17'—H17C | 108.4      |
| C16—C17—H17B  | 110.0      | C16'—C17'—H17D | 108.4      |
| C18—C17—H17B  | 110.0      | C18'—C17'—H17D | 108.4      |
| H17A—C17—H17B | 108.4      | H17C—C17'—H17D | 107.5      |
| C13—C18—C17   | 112.9 (9)  | C13'—C18'—C17' | 109 (2)    |
| C13—C18—H18A  | 109.0      | C13'—C18'—H18C | 109.9      |
| C17—C18—H18A  | 109.0      | C17'—C18'—H18C | 109.9      |
| C13—C18—H18B  | 109.0      | C13'—C18'—H18D | 109.9      |
| C17—C18—H18B  | 109.0      | C17'—C18'—H18D | 109.9      |
| H18A—C18—H18B | 107.8      | H18C—C18'—H18D | 108.3      |
| C12—N1—C11    | 124.0 (12) | C11'—N2'—C1'   | 114.4 (14) |
| C12—N1—H1A    | 117 (3)    | C11'—N2'—H2A   | 110 (3)    |
| C11—N1—H1A    | 107 (3)    | C1'—N2'—H2A    | 96 (3)     |
| C11—N2—C1     | 120.8 (9)  | C11'—N1'—C12'  | 117 (2)    |
| C11—N2—H2A    | 113 (3)    | C11'—N1'—H1A   | 106 (3)    |

|                 |             |                     |             |
|-----------------|-------------|---------------------|-------------|
| C1—N2—H2A       | 119 (3)     | C12'—N1'—H1A        | 108 (3)     |
| C10—C1—C2—C3    | -2 (3)      | C10'—C1'—C2'—C3'    | 1 (4)       |
| N2—C1—C2—C3     | 178.3 (12)  | N2'—C1'—C2'—C3'     | 179 (3)     |
| C1—C2—C3—C4     | 1.6 (19)    | C1'—C2'—C3'—C4'     | -1 (4)      |
| C2—C3—C4—C5     | -0.9 (17)   | C2'—C3'—C4'—C5'     | 2 (3)       |
| C3—C4—C5—C6     | 180.0 (11)  | C3'—C4'—C5'—C6'     | 177 (3)     |
| C3—C4—C5—C10    | 0 (2)       | C3'—C4'—C5'—C10'    | -3 (4)      |
| C10—C5—C6—C7    | 0 (2)       | C10'—C5'—C6'—C7'    | 0 (4)       |
| C4—C5—C6—C7     | 180.0 (14)  | C4'—C5'—C6'—C7'     | 180 (3)     |
| C5—C6—C7—C8     | 0 (2)       | C5'—C6'—C7'—C8'     | -3 (5)      |
| C6—C7—C8—C9     | -3 (3)      | C6'—C7'—C8'—C9'     | 11 (5)      |
| C7—C8—C9—C10    | 6 (3)       | C7'—C8'—C9'—C10'    | -16 (4)     |
| C2—C1—C10—C5    | 1 (3)       | C6'—C5'—C10'—C1'    | -177 (3)    |
| N2—C1—C10—C5    | -178.9 (14) | C4'—C5'—C10'—C1'    | 3 (4)       |
| C2—C1—C10—C9    | 177.1 (18)  | C6'—C5'—C10'—C9'    | -5 (4)      |
| N2—C1—C10—C9    | -3 (3)      | C4'—C5'—C10'—C9'    | 174 (3)     |
| C6—C5—C10—C1    | 180.0 (16)  | C2'—C1'—C10'—C5'    | -2 (4)      |
| C4—C5—C10—C1    | 0 (3)       | N2'—C1'—C10'—C5'    | 180 (2)     |
| C6—C5—C10—C9    | 4 (2)       | C2'—C1'—C10'—C9'    | -174 (3)    |
| C4—C5—C10—C9    | -176.5 (16) | N2'—C1'—C10'—C9'    | 7 (3)       |
| C8—C9—C10—C1    | 178 (2)     | C8'—C9'—C10'—C5'    | 13 (4)      |
| C8—C9—C10—C5    | -7 (3)      | C8'—C9'—C10'—C1'    | -175 (3)    |
| O1—C12—C13—C18  | 24 (2)      | O1'—C12'—C13'—C14'  | -78 (3)     |
| N1—C12—C13—C18  | -170.9 (14) | N1'—C12'—C13'—C14'  | 123 (3)     |
| O1—C12—C13—C14  | -100.1 (15) | O1'—C12'—C13'—C18'  | 48 (4)      |
| N1—C12—C13—C14  | 65.1 (18)   | N1'—C12'—C13'—C18'  | -111 (3)    |
| C12—C13—C14—C15 | 179.2 (11)  | C18'—C13'—C14'—C15' | 53 (3)      |
| C18—C13—C14—C15 | 54.0 (16)   | C12'—C13'—C14'—C15' | 179.9 (19)  |
| C13—C14—C15—C16 | -54.8 (15)  | C13'—C14'—C15'—C16' | -57 (3)     |
| C14—C15—C16—C17 | 57.3 (14)   | C14'—C15'—C16'—C17' | 57 (3)      |
| C15—C16—C17—C18 | -57.7 (13)  | C15'—C16'—C17'—C18' | -59 (3)     |
| C12—C13—C18—C17 | -178.7 (11) | C14'—C13'—C18'—C17' | -51 (3)     |
| C14—C13—C18—C17 | -54.8 (16)  | C12'—C13'—C18'—C17' | -175 (2)    |
| C16—C17—C18—C13 | 56.1 (14)   | C16'—C17'—C18'—C13' | 56 (3)      |
| O1—C12—N1—C11   | -27 (2)     | N1'—C11'—N2'—C1'    | -120 (2)    |
| C13—C12—N1—C11  | 167.2 (13)  | S1'—C11'—N2'—C1'    | 43 (2)      |
| N2—C11—N1—C12   | 23 (2)      | C2'—C1'—N2'—C11'    | 66 (4)      |
| S1—C11—N1—C12   | -167.5 (12) | C10'—C1'—N2'—C11'   | -116.0 (19) |
| N1—C11—N2—C1    | 159.0 (11)  | N2'—C11'—N1'—C12'   | -43 (3)     |
| S1—C11—N2—C1    | -9.9 (13)   | S1'—C11'—N1'—C12'   | 153.6 (19)  |
| C10—C1—N2—C11   | -80 (2)     | O1'—C12'—N1'—C11'   | 46 (4)      |
| C2—C1—N2—C11    | 99.7 (12)   | C13'—C12'—N1'—C11'  | -156 (2)    |

*Hydrogen-bond geometry (Å, °)*

Cg1 is the centroid of the C1–C5/C10 ring.

| <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> |
|---------------------------------------|-------------|---------------|-----------------------|-------------------------|
| N2—H2 <i>A</i> ···O1                  | 0.88 (4)    | 1.97 (4)      | 2.667 (15)            | 135 (4)                 |
| N2'—H2 <i>A</i> ···O1'                | 0.93 (4)    | 2.03 (4)      | 2.62 (3)              | 120 (4)                 |
| N1—H1 <i>A</i> ···S1 <sup>i</sup>     | 0.87 (3)    | 2.53 (3)      | 3.370 (19)            | 161 (4)                 |
| N1'—H1 <i>A</i> ···S1' <sup>ii</sup>  | 0.90 (4)    | 2.59 (4)      | 3.44 (3)              | 159 (4)                 |
| C18—H18 <i>B</i> ···Cg1 <sup>ii</sup> | 0.90        | 2.66          | 3.527 (2)             | 148                     |

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