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## Structure Reports

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## 6-Butyl-5-(4-methylphenoxy)-3-phenyl-3H-1,2,3-triazolo[4,5-d]pyrimidin-7(6H)-one

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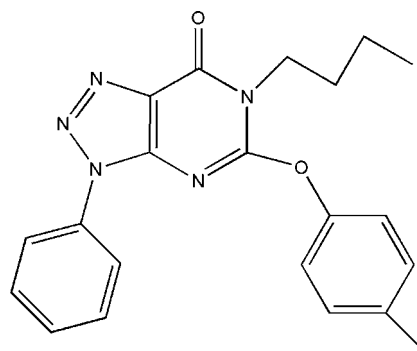
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Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å; disorder in main residue;  $R$  factor = 0.054;  $wR$  factor = 0.182; data-to-parameter ratio = 13.2.

In the title compound,  $\text{C}_{21}\text{H}_{21}\text{N}_5\text{O}_2$ , the triazolopyrimidine ring system is essentially planar [maximum displacement = 0.021 (4) Å] and forms dihedral angles of 41.17 (9) and 67.99 (8)° with the phenyl and benzene rings, respectively. The *n*-butyl side chains is disordered over two positions with an occupancy ratio of 0.77:0.23. An intramolecular C—H···O hydrogen-bonding interaction stabilizes the molecular conformation. In the crystal, molecules are linked by intermolecular C—H···O and C—H···N hydrogen bonds into a three-dimensional network. In addition,  $\pi$ - $\pi$  stacking interactions involving the triazole and pyrimidine rings of adjacent molecules are observed, with centroid-centroid distances of 3.545 (1) Å.

### Related literature

For the synthesis and biological activity of 8-azaguanine derivatives, see: Roblin *et al.* (1945); Ding *et al.* (2004); Mitchell *et al.* (1950); Levine *et al.* (1963); Montgomery *et al.* (1962); Yamamoto *et al.* (1967); Bariana (1971); Holland *et al.* (1975); Zeng *et al.* (2010). For related structures, see: Ferguson *et al.* (1998); Li *et al.* (2004); Zhao, Xie *et al.* (2005); Zhao, Hu *et al.* (2005); Zhao, Wang & Ding (2005); Chen & Shi (2006); Maldonado *et al.* (2006); Xiao *et al.* (2007); Wang *et al.* (2006, 2008); Zeng *et al.* (2006, 2009).



### Experimental

#### Crystal data

$\text{C}_{21}\text{H}_{21}\text{N}_5\text{O}_2$   
 $M_r = 375.43$   
 Monoclinic,  $P2_1/n$   
 $a = 11.0954$  (10) Å  
 $b = 16.4478$  (15) Å  
 $c = 11.3484$  (11) Å  
 $\beta = 107.643$  (1)°  
 $V = 1973.6$  (3) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 298$  K  
 $0.20 \times 0.20 \times 0.20$  mm

#### Data collection

Bruker SMART CCD area-detector diffractometer  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.983$ ,  $T_{\max} = 0.985$   
 20458 measured reflections  
 3876 independent reflections  
 2663 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.055$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.054$   
 $wR(F^2) = 0.182$   
 $S = 1.08$   
 3876 reflections  
 293 parameters  
 11 restraints  
 H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.56$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.24$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C12—H12A···O2	0.97	2.50	3.048 (5)	116
C2—H2···O1 <sup>i</sup>	0.93	2.53	3.230 (3)	133
C3—H3···N2 <sup>ii</sup>	0.93	2.61	3.535 (2)	174

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

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: PLATON (Spek, 2009); software used to prepare material for publication: SHELXTL (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ2502).

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## supporting information

*Acta Cryst.* (2010). E66, o2990–o2991 [https://doi.org/10.1107/S160053681004300X]

## 6-Butyl-5-(4-methylphenoxy)-3-phenyl-3*H*-1,2,3-triazolo[4,5-*d*]pyrimidin-7(6*H*)-one

Hong-Mei Wang, Shou-Heng Deng, Xiao-Hua Zeng, Ping Chen and Li-Li Chen

### S1. Comment

The derivatives of heterocycles containing 8-azaguanine system, which are well known bioisosteres of guanine, are of great importance because of their remarkable biological properties. Some of these activities include antimicrobial or antifungal activities (Roblin *et al.*, 1945; Ding *et al.*, 2004; Zeng *et al.*, 2010), encephaloma cell inhibitor activity (Mitchell *et al.*, 1950; Levine *et al.*, 1963), antileukemic activity (Montgomery *et al.*, 1962), hypersusceptibility inhibitor activity and acesodyne activity (Yamamoto *et al.*, 1967; Bariana, 1971; Holland *et al.*, 1975). In recent years, Ding's group has been engaged in the preparation of derivatives of 8-azaguanine *via* aza-Wittig reaction of beta-ethoxycarbonyl iminophosphoranes with aromatic isocyanates (Zhao, Xie *et al.*, 2005). As a continuation of our research for new biologically active heterocycles, the title compound was obtained from beta-ethoxycarbonyl iminophosphorane with aliphatic isocyanate, and structurally characterized in this context.

In the title compound (Fig. 1), bond lengths and angles within the triazolopyrimidinone moiety are in good agreement with those observed for closely related structures (Zhao, Hu *et al.*, 2005; Zhao, Wang & Ding, 2005). As reported for related compounds (Ferguson *et al.*, 1998; Li *et al.*, 2004; Maldonado *et al.*, 2006; Zeng *et al.*, 2006, 2009; Wang *et al.*, 2006, 2008; Xiao *et al.*, 2007; Chen & Shi, 2006), the triazolopyrimidine ring system is essentially planar, with a maximum displacement of 0.021 (4) Å for atom C8, and forms dihedral angles of 41.17 (9) and 67.99 (8)° with the C1–C6 and C15–C20 rings, respectively. There exists an intramolecular C—H···O hydrogen bonding interaction stabilizing the molecular conformation. In the crystal packing, molecules are linked by intermolecular C—H···O and C—H···N hydrogen bonds (Table 1). In addition,  $\pi$ – $\pi$  stacking interactions involving the triazole and pyrimidine rings of adjacent molecules are observed, with centroid-to-centroid distances of 3.545 (1) Å.

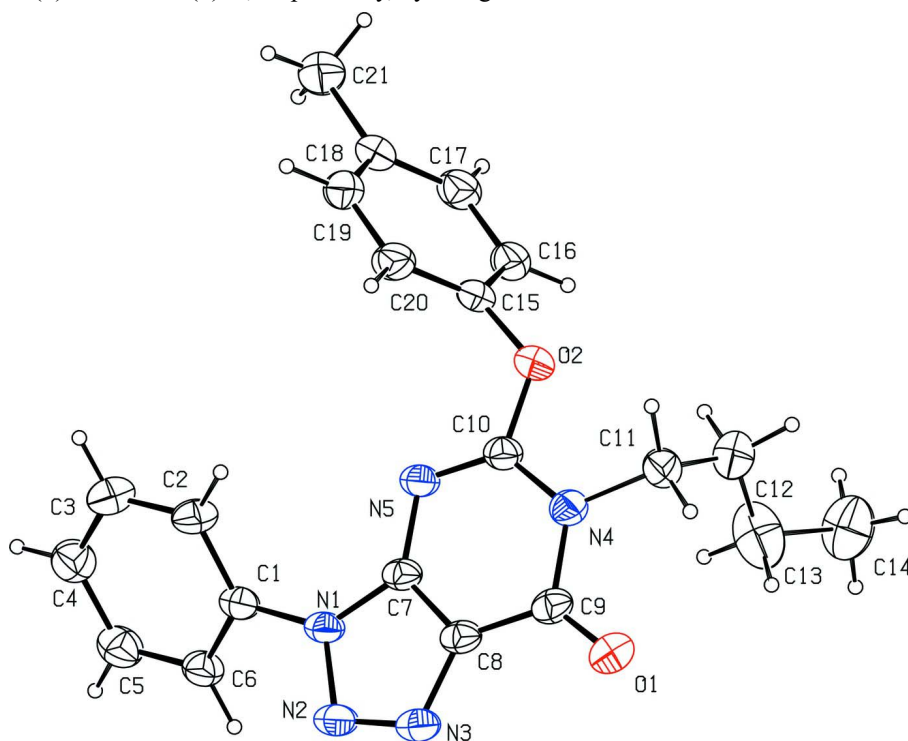
### S2. Experimental

To the solution of carbodiimide in CH<sub>2</sub>Cl<sub>2</sub>/CH<sub>3</sub>CN (1:4 *v/v*, 15 ml) prepared according to the literature method (Zeng *et al.*, 2006), was added 4-methylphenol (3 mmol) and excess K<sub>2</sub>CO<sub>3</sub>, and the reaction mixture was stirred for 12 h. The solvent was removed under reduced pressure and the residue was recrystallized from EtOH to give the title compound (yield 93%; m.p. 406 K). Elemental analysis: calculated for C<sub>21</sub>H<sub>21</sub>N<sub>3</sub>O<sub>2</sub>: C, 67.18; H, 5.64; N, 18.65%. Found: C, 66.62; H, 5.98; N, 18.13%. Crystals suitable for single crystal X-ray diffraction analysis were obtained by slow evaporation of a hexane/dichloromethane (1:3 *v/v*) solution at room temperature.

### S3. Refinement

H atoms were placed at calculated positions and treated as riding atoms, with C—H = 0.93–0.97 Å, and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for CH or  $1.5U_{\text{eq}}(\text{C})$  for CH<sub>3</sub>. The *n*-butyl side chain is disordered over two positions with occupancy factors of 0.77:0.23. During the refinement, the adjacent and interval C—C distances involving the disordered carbon atoms were

restrained to be 1.54 (1) Å and 2.45 (2) Å, respectively, by using the command *DFIX*.



**Figure 1**

The molecular structure of the title compound showing the atom-labeling scheme. Displacement ellipsoids are drawn at 50% probability level. H-atoms are represented by circles of arbitrary size. Only the major component of the disorder is shown.

### 6-Butyl-5-(4-methylphenoxy)-3-phenyl-3H-1,2,3-triazolo[4,5-d]pyrimidin-7(6H)-one

#### Crystal data

$C_{21}H_{21}N_5O_2$

$M_r = 375.43$

Monoclinic,  $P2_1/n$

Hall symbol: -P 2yn

$a = 11.0954$  (10) Å

$b = 16.4478$  (15) Å

$c = 11.3484$  (11) Å

$\beta = 107.643$  (1)°

$V = 1973.6$  (3) Å<sup>3</sup>

$Z = 4$

$F(000) = 792$

$D_x = 1.263$  Mg m<sup>-3</sup>

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

Cell parameters from 6350 reflections

$\theta = 2.3$ – $25.8$ °

$\mu = 0.09$  mm<sup>-1</sup>

$T = 298$  K

Block, colourless

$0.20 \times 0.20 \times 0.20$  mm

#### Data collection

Bruker SMART CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.983$ ,  $T_{\max} = 0.985$

20458 measured reflections

3876 independent reflections

2663 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.055$

$\theta_{\max} = 26.0$ °,  $\theta_{\min} = 2.3$ °

$h = -13 \rightarrow 13$

$k = -20 \rightarrow 20$

$l = -13 \rightarrow 13$

*Refinement*

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.054$	H-atom parameters constrained
$wR(F^2) = 0.182$	$w = 1/[\sigma^2(F_o^2) + (0.1122P)^2 + 0.0419P]$
$S = 1.08$	where $P = (F_o^2 + 2F_c^2)/3$
3876 reflections	$(\Delta/\sigma)_{\max} < 0.001$
293 parameters	$\Delta\rho_{\max} = 0.56 \text{ e } \text{\AA}^{-3}$
11 restraints	$\Delta\rho_{\min} = -0.24 \text{ e } \text{\AA}^{-3}$
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	0.42419 (17)	0.24051 (13)	0.45138 (17)	0.0654 (5)	
C2	0.5223 (2)	0.26001 (14)	0.5541 (2)	0.0797 (6)	
H2	0.5810	0.2208	0.5936	0.096*	
C3	0.5326 (2)	0.33854 (15)	0.5978 (2)	0.0904 (7)	
H3	0.5980	0.3523	0.6682	0.108*	
C4	0.4468 (3)	0.39682 (15)	0.5381 (3)	0.0909 (7)	
H4	0.4543	0.4499	0.5677	0.109*	
C5	0.3502 (2)	0.37608 (16)	0.4348 (3)	0.0923 (7)	
H5	0.2922	0.4154	0.3942	0.111*	
C6	0.3380 (2)	0.29824 (14)	0.3907 (2)	0.0764 (6)	
H6	0.2723	0.2845	0.3205	0.092*	
C7	0.49373 (17)	0.10245 (12)	0.39629 (16)	0.0616 (5)	
C8	0.42311 (18)	0.03487 (12)	0.34748 (17)	0.0659 (5)	
C9	0.4821 (2)	-0.03604 (13)	0.31905 (17)	0.0681 (5)	
C10	0.67331 (18)	0.04538 (13)	0.39924 (17)	0.0670 (5)	
C11	0.6857 (6)	-0.0970 (5)	0.3246 (5)	0.0795 (18)	0.77
H11A	0.6320	-0.1448	0.3109	0.095*	0.77
H11B	0.7589	-0.1073	0.3956	0.095*	0.77
C12	0.7294 (5)	-0.0808 (4)	0.2082 (5)	0.138 (2)	0.77
H12A	0.7829	-0.0328	0.2238	0.166*	0.77
H12B	0.7810	-0.1263	0.1980	0.166*	0.77
C13	0.6379 (6)	-0.0699 (3)	0.1027 (5)	0.149 (2)	0.77
H13A	0.6006	-0.0168	0.1050	0.179*	0.77
H13B	0.5726	-0.1101	0.0979	0.179*	0.77
C14	0.6776 (10)	-0.0757 (5)	-0.0175 (6)	0.142 (4)	0.77

H14A	0.7472	-0.0395	-0.0114	0.213*	0.77
H14B	0.6074	-0.0606	-0.0874	0.213*	0.77
H14C	0.7026	-0.1304	-0.0279	0.213*	0.77
C11'	0.704 (2)	-0.0825 (18)	0.3141 (15)	0.104 (10)	0.23
H11C	0.7847	-0.0555	0.3291	0.125*	0.23
H11D	0.7168	-0.1296	0.3679	0.125*	0.23
C12'	0.6617 (12)	-0.1107 (8)	0.1849 (10)	0.081 (3)	0.23
H12C	0.6172	-0.1610	0.1880	0.097*	0.23
H12D	0.5967	-0.0720	0.1431	0.097*	0.23
C13'	0.7276 (11)	-0.1267 (7)	0.0973 (8)	0.104 (4)	0.23
H13C	0.8104	-0.1024	0.1329	0.125*	0.23
H13D	0.7414	-0.1850	0.1015	0.125*	0.23
C14'	0.695 (2)	-0.1081 (15)	-0.0281 (18)	0.101 (6)	0.23
H14D	0.6086	-0.1234	-0.0676	0.152*	0.23
H14E	0.7492	-0.1374	-0.0650	0.152*	0.23
H14F	0.7046	-0.0508	-0.0381	0.152*	0.23
C15	0.87510 (18)	0.10857 (13)	0.4721 (2)	0.0717 (6)	
C16	0.9372 (2)	0.14531 (16)	0.4005 (2)	0.0856 (7)	
H16	0.9238	0.1289	0.3191	0.103*	
C17	1.0212 (2)	0.20782 (16)	0.4509 (2)	0.0895 (7)	
H17	1.0632	0.2338	0.4020	0.107*	
C18	1.04316 (19)	0.23192 (14)	0.5710 (2)	0.0809 (6)	
C19	0.9782 (2)	0.19289 (17)	0.6399 (2)	0.0904 (7)	
H19	0.9913	0.2088	0.7215	0.108*	
C20	0.8938 (2)	0.13042 (16)	0.5914 (2)	0.0865 (7)	
H20	0.8511	0.1042	0.6396	0.104*	
C21	1.1367 (3)	0.29832 (18)	0.6258 (3)	0.1124 (9)	
H21A	1.2173	0.2847	0.6163	0.169*	
H21B	1.1449	0.3040	0.7121	0.169*	
H21C	1.1074	0.3486	0.5840	0.169*	
N1	0.40888 (13)	0.15838 (10)	0.40749 (14)	0.0651 (4)	
N2	0.28911 (15)	0.12506 (12)	0.36653 (17)	0.0760 (5)	
N3	0.29856 (15)	0.05057 (12)	0.33090 (16)	0.0763 (5)	
N4	0.61476 (15)	-0.02544 (10)	0.34970 (14)	0.0686 (5)	
N5	0.62063 (14)	0.11114 (10)	0.42446 (15)	0.0660 (4)	
O1	0.43198 (15)	-0.09929 (10)	0.27448 (15)	0.0877 (5)	
O2	0.79833 (13)	0.04150 (9)	0.41997 (15)	0.0840 (5)	

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0492 (10)	0.0861 (14)	0.0599 (11)	0.0031 (9)	0.0152 (8)	0.0080 (10)
C2	0.0634 (12)	0.0853 (15)	0.0760 (14)	0.0001 (11)	-0.0002 (10)	0.0097 (11)
C3	0.0771 (15)	0.0954 (17)	0.0843 (15)	-0.0114 (13)	0.0029 (12)	0.0014 (13)
C4	0.0866 (17)	0.0855 (15)	0.1014 (19)	-0.0009 (13)	0.0299 (14)	0.0018 (13)
C5	0.0803 (16)	0.0996 (18)	0.0938 (18)	0.0185 (13)	0.0215 (13)	0.0128 (14)
C6	0.0601 (12)	0.0930 (16)	0.0707 (13)	0.0125 (10)	0.0115 (10)	0.0046 (11)
C7	0.0500 (10)	0.0810 (12)	0.0489 (10)	-0.0003 (9)	0.0079 (7)	0.0085 (8)

C8	0.0558 (11)	0.0851 (13)	0.0507 (10)	-0.0027 (9)	0.0068 (8)	0.0085 (9)
C9	0.0658 (12)	0.0824 (14)	0.0495 (10)	-0.0053 (10)	0.0074 (8)	0.0065 (9)
C10	0.0532 (11)	0.0879 (14)	0.0570 (11)	0.0019 (10)	0.0121 (8)	0.0019 (10)
C11	0.075 (2)	0.086 (3)	0.076 (3)	0.011 (3)	0.020 (2)	-0.004 (2)
C12	0.111 (4)	0.156 (5)	0.147 (5)	-0.007 (3)	0.038 (4)	-0.070 (4)
C13	0.205 (6)	0.147 (4)	0.114 (3)	0.064 (4)	0.077 (4)	0.047 (3)
C14	0.200 (8)	0.147 (7)	0.095 (4)	-0.014 (5)	0.069 (4)	0.014 (4)
C11'	0.129 (17)	0.097 (15)	0.105 (15)	-0.015 (10)	0.064 (12)	-0.019 (10)
C12'	0.077 (7)	0.097 (8)	0.069 (7)	-0.011 (6)	0.022 (6)	-0.018 (6)
C13'	0.115 (10)	0.090 (7)	0.097 (9)	0.002 (7)	0.015 (7)	-0.010 (6)
C14'	0.090 (10)	0.108 (15)	0.106 (12)	-0.014 (9)	0.030 (8)	0.001 (9)
C15	0.0446 (10)	0.0929 (14)	0.0758 (13)	0.0081 (10)	0.0154 (9)	-0.0068 (11)
C16	0.0669 (13)	0.1198 (18)	0.0734 (14)	0.0020 (13)	0.0264 (11)	-0.0123 (13)
C17	0.0689 (14)	0.1146 (18)	0.0916 (17)	-0.0009 (13)	0.0341 (12)	-0.0005 (14)
C18	0.0544 (11)	0.0949 (15)	0.0930 (16)	0.0076 (11)	0.0217 (11)	-0.0096 (12)
C19	0.0702 (14)	0.127 (2)	0.0742 (14)	-0.0021 (13)	0.0221 (11)	-0.0188 (13)
C20	0.0644 (13)	0.1234 (19)	0.0741 (15)	-0.0102 (13)	0.0247 (11)	-0.0068 (13)
C21	0.0855 (18)	0.119 (2)	0.133 (2)	-0.0136 (16)	0.0334 (17)	-0.0260 (18)
N1	0.0459 (8)	0.0832 (11)	0.0604 (9)	0.0038 (8)	0.0072 (7)	0.0055 (8)
N2	0.0471 (9)	0.0998 (13)	0.0749 (11)	-0.0033 (8)	0.0091 (8)	0.0027 (9)
N3	0.0534 (10)	0.0983 (13)	0.0695 (10)	-0.0053 (9)	0.0072 (8)	0.0046 (9)
N4	0.0641 (10)	0.0817 (11)	0.0557 (9)	0.0031 (8)	0.0118 (7)	0.0001 (8)
N5	0.0480 (9)	0.0836 (11)	0.0630 (9)	0.0007 (8)	0.0116 (7)	0.0018 (8)
O1	0.0859 (11)	0.0879 (11)	0.0802 (10)	-0.0134 (8)	0.0117 (8)	-0.0049 (8)
O2	0.0526 (8)	0.1000 (11)	0.0981 (11)	0.0020 (7)	0.0209 (7)	-0.0187 (8)

*Geometric parameters (Å, °)*

C1—C2	1.370 (3)	C14—H14B	0.9600
C1—C6	1.375 (3)	C14—H14C	0.9600
C1—N1	1.432 (3)	C11'—C12'	1.473 (10)
C2—C3	1.376 (3)	C11'—N4	1.502 (10)
C2—H2	0.9300	C11'—H11C	0.9700
C3—C4	1.376 (3)	C11'—H11D	0.9700
C3—H3	0.9300	C12'—C13'	1.426 (9)
C4—C5	1.371 (4)	C12'—H12C	0.9700
C4—H4	0.9300	C12'—H12D	0.9700
C5—C6	1.366 (3)	C13'—C14'	1.39 (2)
C5—H5	0.9300	C13'—H13C	0.9700
C6—H6	0.9300	C13'—H13D	0.9700
C7—N1	1.349 (2)	C14'—H14D	0.9600
C7—N5	1.354 (2)	C14'—H14E	0.9600
C7—C8	1.376 (3)	C14'—H14F	0.9600
C8—N3	1.362 (3)	C15—C20	1.355 (3)
C8—C9	1.422 (3)	C15—C16	1.356 (3)
C9—O1	1.215 (2)	C15—O2	1.410 (3)
C9—N4	1.416 (3)	C16—C17	1.389 (3)
C10—N5	1.302 (3)	C16—H16	0.9300

C10—O2	1.336 (2)	C17—C18	1.369 (3)
C10—N4	1.369 (3)	C17—H17	0.9300
C11—N4	1.491 (4)	C18—C19	1.373 (3)
C11—C12	1.562 (8)	C18—C21	1.505 (3)
C11—H11A	0.9700	C19—C20	1.387 (3)
C11—H11B	0.9700	C19—H19	0.9300
C12—C13	1.326 (6)	C20—H20	0.9300
C12—H12A	0.9700	C21—H21A	0.9600
C12—H12B	0.9700	C21—H21B	0.9600
C13—C14	1.559 (7)	C21—H21C	0.9600
C13—H13A	0.9700	N1—N2	1.381 (2)
C13—H13B	0.9700	N2—N3	1.304 (3)
C14—H14A	0.9600		
C2—C1—C6	121.1 (2)	C13'—C12'—H12C	104.1
C2—C1—N1	119.77 (18)	C11'—C12'—H12C	104.1
C6—C1—N1	119.07 (18)	C13'—C12'—H12D	104.1
C1—C2—C3	118.9 (2)	C11'—C12'—H12D	104.1
C1—C2—H2	120.5	H12C—C12'—H12D	105.5
C3—C2—H2	120.5	C14'—C13'—C12'	130.0 (14)
C4—C3—C2	120.5 (2)	C14'—C13'—H13C	104.8
C4—C3—H3	119.8	C12'—C13'—H13C	104.8
C2—C3—H3	119.8	C14'—C13'—H13D	104.8
C5—C4—C3	119.5 (2)	C12'—C13'—H13D	104.8
C5—C4—H4	120.2	H13C—C13'—H13D	105.8
C3—C4—H4	120.2	C13'—C14'—H14D	109.5
C6—C5—C4	120.8 (2)	C13'—C14'—H14E	109.5
C6—C5—H5	119.6	H14D—C14'—H14E	109.5
C4—C5—H5	119.6	C13'—C14'—H14F	109.5
C5—C6—C1	119.1 (2)	H14D—C14'—H14F	109.5
C5—C6—H6	120.4	H14E—C14'—H14F	109.5
C1—C6—H6	120.4	C20—C15—C16	121.9 (2)
N1—C7—N5	127.68 (18)	C20—C15—O2	121.1 (2)
N1—C7—C8	105.08 (17)	C16—C15—O2	116.70 (19)
N5—C7—C8	127.23 (19)	C15—C16—C17	118.9 (2)
N3—C8—C7	109.43 (18)	C15—C16—H16	120.6
N3—C8—C9	129.96 (19)	C17—C16—H16	120.6
C7—C8—C9	120.58 (19)	C18—C17—C16	121.3 (2)
O1—C9—N4	121.2 (2)	C18—C17—H17	119.4
O1—C9—C8	127.7 (2)	C16—C17—H17	119.4
N4—C9—C8	111.04 (18)	C17—C18—C19	117.8 (2)
N5—C10—O2	120.81 (18)	C17—C18—C21	120.9 (2)
N5—C10—N4	127.48 (18)	C19—C18—C21	121.3 (2)
O2—C10—N4	111.70 (18)	C18—C19—C20	121.9 (2)
N4—C11—C12	110.1 (5)	C18—C19—H19	119.0
N4—C11—H11A	109.6	C20—C19—H19	119.0
C12—C11—H11A	109.6	C15—C20—C19	118.2 (2)
N4—C11—H11B	109.6	C15—C20—H20	120.9



C12—C11—H11B	109.6	C19—C20—H20	120.9
H11A—C11—H11B	108.2	C18—C21—H21A	109.5
C13—C12—C11	115.9 (5)	C18—C21—H21B	109.5
C13—C12—H12A	108.3	H21A—C21—H21B	109.5
C11—C12—H12A	108.3	C18—C21—H21C	109.5
C13—C12—H12B	108.3	H21A—C21—H21C	109.5
C11—C12—H12B	108.3	H21B—C21—H21C	109.5
H12A—C12—H12B	107.4	C7—N1—N2	108.99 (17)
C12—C13—C14	116.1 (6)	C7—N1—C1	131.55 (16)
C12—C13—H13A	108.3	N2—N1—C1	119.47 (15)
C14—C13—H13A	108.3	N3—N2—N1	108.51 (16)
C12—C13—H13B	108.3	N2—N3—C8	107.99 (16)
C14—C13—H13B	108.3	C10—N4—C9	122.28 (17)
H13A—C13—H13B	107.4	C10—N4—C11	122.6 (4)
C12'—C11'—N4	115.4 (11)	C9—N4—C11	115.2 (4)
C12'—C11'—H11C	108.4	C10—N4—C11'	111.9 (13)
N4—C11'—H11C	108.4	C9—N4—C11'	125.1 (12)
C12'—C11'—H11D	108.4	C11—N4—C11'	13.6 (18)
N4—C11'—H11D	108.4	C10—N5—C7	111.38 (17)
H11C—C11'—H11D	107.5	C10—O2—C15	119.97 (16)
C13'—C12'—C11'	132.7 (14)		

*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>
C12—H12 <i>A</i> ...O2	0.97	2.50	3.048 (5)	116
C2—H2...O1 <sup>i</sup>	0.93	2.53	3.230 (3)	133
C3—H3...N2 <sup>ii</sup>	0.93	2.61	3.535 (2)	174

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