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## Structure Reports

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**(Z)-Ethyl 2,4-diphenyl-3-(propylamino)-but-2-enoate**

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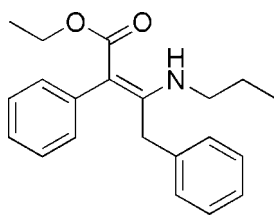
Received 23 December 2008; accepted 25 December 2008

Key indicators: single-crystal X-ray study;  $T = 293$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å; disorder in main residue;  $R$  factor = 0.060;  $wR$  factor = 0.198; data-to-parameter ratio = 14.3.

The title compound,  $\text{C}_{21}\text{H}_{25}\text{NO}_2$ , adopts a *Z* conformation about the  $\text{C}=\text{C}$  double bond. The molecular structure is stabilized by an intramolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bond and the dihedral angle between the aromatic ring planes is  $76.04$  ( $12$ )°. The atoms of the ethyl substituent are disordered over two sets of sites in a  $0.60$  ( $2$ ): $0.40$  ( $2$ ) ratio.

## Related literature

For the synthesis, see: Du *et al.* (2006). For background, see: Xue *et al.* (2007).



## Experimental

## Crystal data

 $\text{C}_{21}\text{H}_{25}\text{NO}_2$  $M_r = 323.42$ 

Monoclinic,  $P2_1/n$   
 $a = 12.186$  (2) Å  
 $b = 8.4771$  (17) Å  
 $c = 19.080$  (4) Å  
 $\beta = 106.33$  (3)°  
 $V = 1891.4$  (7) Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.07$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 $0.28 \times 0.22 \times 0.18$  mm

## Data collection

Rigaku Saturn diffractometer  
 Absorption correction: multi-scan  
 (*CrystalClear*; Rigaku/MS, 2005)  
 $T_{\min} = 0.980$ ,  $T_{\max} = 0.987$

12325 measured reflections  
 3323 independent reflections  
 2290 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.034$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.060$   
 $wR(F^2) = 0.198$   
 $S = 1.06$   
 3323 reflections  
 232 parameters  
 5 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.24$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.19$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O2}$	0.905 (16)	1.925 (17)	2.653 (3)	136.3 (14)

Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

We thank the Technology Fund of Tianjin Bureau of Public Health (grant No. 04KY36) for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2886).

## References

- Du, Y., Liu, R., Linn, G. & Zhao, K. (2006). *Org. Lett.* **8**, 5919–5922.  
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 Xue, J. Y., Xiao, Z. P., Shi, L., Tan, S. H., Li, H. Q. & Zhu, H. L. (2007). *Aust. J. Chem.* **60**, 957–962.

## supporting information

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**(Z)-Ethyl 2,4-diphenyl-3-(propylamino)but-2-enoate****Huimin Jin, Peifan Li, Bin Liu and Xiaoqing Cheng****S1. Comment**

Enamine compounds have been considered to be potential antibacterial agents (Xue *et al.*, 2007) and found important application in the synthesis of N-containing heterocycles (Du *et al.*, 2006). To further study the structure and activity relationship, we determine the crystal structure of the title compound, (I).

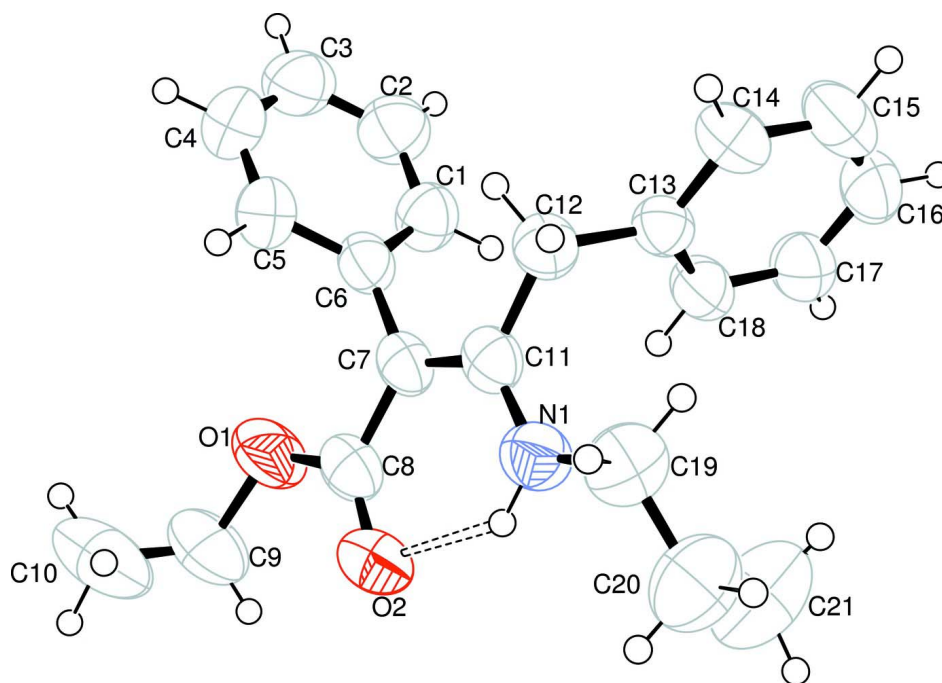
In the molecular structure (Fig. 1), the torsion angles of N1—C11—C7—C6 and C8—C7—C11—C12 are  $-177.32$  (17) and  $-175.14$  (16) $^\circ$ , respectively. Furthermore, the distances C7—C11 and C11—N1 are 1.381 (3), 1.348 (3) Å, respectively. Both of these features confirm the enamine structure formation. The two phenyl rings constructed an angle of  $76.04$  (12) $^\circ$ . The molecule adopts a *Z*-conformation, being stabilised by an intramolecular N—H $\cdots$ O hydrogen bond (Table 1).

**S2. Experimental**

The title compound was prepared according to the method of the literature (Du, *et al.*, 2006). Colourless blocks of (I) were grown from a mixture of ethyl acetate and petroleum ether.

**S3. Refinement**

The non-N H atoms were positioned geometrically (C—H = 0.93–0.98 Å) and refined as riding with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{CH}$  and  $\text{CH}_2)$  or  $1.5U_{\text{eq}}(\text{CH}_3)$ . The N—H distance was refined with the restraint of 0.90 (1) Å, and the C19—C20, C20—C21, C9—C10 and C9—C10' with the restraint of 1.54 (1) Å. The ethyl radical of the ester moiety was found to be disordered, with the site occupancy ratio of 0.40 (2):0.60 (2).

**Figure 1**

The molecular structure of (I) with 50% probability displacement ellipsoids. The dashed line indicates the intramolecular N—H...O hydrogen bond.

### (Z)-Ethyl 2,4-diphenyl-3-(propylamino)but-2-enoate

#### Crystal data

$C_{21}H_{25}NO_2$   
 $M_r = 323.42$   
 Monoclinic,  $P2_1/n$   
 Hall symbol:  $-P\ 2_1n$   
 $a = 12.186\ (2)\ \text{\AA}$   
 $b = 8.4771\ (17)\ \text{\AA}$   
 $c = 19.080\ (4)\ \text{\AA}$   
 $\beta = 106.33\ (3)^\circ$   
 $V = 1891.4\ (7)\ \text{\AA}^3$   
 $Z = 4$

$F(000) = 696$   
 $D_x = 1.136\ \text{Mg m}^{-3}$   
 Mo  $K\alpha$  radiation,  $\lambda = 0.71073\ \text{\AA}$   
 Cell parameters from 4099 reflections  
 $\theta = 2.2\text{--}27.1^\circ$   
 $\mu = 0.07\ \text{mm}^{-1}$   
 $T = 293\ \text{K}$   
 Block, colourless  
 $0.28 \times 0.22 \times 0.18\ \text{mm}$

#### Data collection

Rigaku Saturn  
 diffractometer  
 Radiation source: rotating anode  
 Confocal monochromator  
 Detector resolution:  $7.31\ \text{pixels mm}^{-1}$   
 $\omega$  scans  
 Absorption correction: multi-scan  
 (*CrystalClear*; Rigaku/MSC, 2005)  
 $T_{\min} = 0.980$ ,  $T_{\max} = 0.987$

12325 measured reflections  
 3323 independent reflections  
 2290 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.034$   
 $\theta_{\max} = 25.0^\circ$ ,  $\theta_{\min} = 2.2^\circ$   
 $h = -14 \rightarrow 14$   
 $k = -9 \rightarrow 10$   
 $l = -21 \rightarrow 22$

*Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.060$  $wR(F^2) = 0.198$  $S = 1.06$ 

3323 reflections

232 parameters

5 restraints

Primary atom site location: structure-invariant  
direct methodsSecondary atom site location: difference Fourier  
mapHydrogen site location: inferred from  
neighbouring sitesH atoms treated by a mixture of independent  
and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.1207P)^2 + 0.019P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} = 0.001$  $\Delta\rho_{\max} = 0.24 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick,  
2008),  $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$ 

Extinction coefficient: 0.079 (13)

*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)
O1	0.98880 (16)	0.7415 (2)	0.24987 (10)	0.0936 (6)	
O2	1.07264 (14)	0.97745 (19)	0.24972 (8)	0.0791 (5)	
N1	0.95170 (17)	1.1941 (2)	0.16036 (11)	0.0710 (6)	
C1	0.7770 (2)	0.7362 (3)	0.07667 (13)	0.0723 (7)	
H1	0.8213	0.7655	0.0463	0.087*	
C2	0.6935 (2)	0.6239 (3)	0.05278 (16)	0.0864 (8)	
H2A	0.6812	0.5794	0.0067	0.104*	
C3	0.6286 (2)	0.5778 (3)	0.09719 (18)	0.0906 (8)	
H3A	0.5719	0.5018	0.0814	0.109*	
C4	0.6473 (2)	0.6434 (4)	0.16427 (18)	0.0977 (9)	
H4A	0.6038	0.6115	0.1947	0.117*	
C5	0.7302 (2)	0.7568 (3)	0.18764 (14)	0.0829 (8)	
H5A	0.7415	0.8007	0.2338	0.099*	
C6	0.79728 (17)	0.8072 (2)	0.14437 (11)	0.0591 (6)	
C7	0.88662 (17)	0.9301 (2)	0.16911 (10)	0.0572 (6)	
C8	0.99038 (19)	0.8903 (3)	0.22513 (12)	0.0654 (6)	
C9	1.0852 (3)	0.6899 (4)	0.30878 (18)	0.1159 (11)	0.60 (2)
H9A	1.1075	0.5855	0.2971	0.139*	0.60 (2)
H9B	1.1489	0.7602	0.3112	0.139*	0.60 (2)
C10	1.0665 (11)	0.684 (2)	0.3778 (4)	0.133 (4)	0.60 (2)
H10A	1.1353	0.6512	0.4133	0.199*	0.60 (2)
H10B	1.0063	0.6106	0.3770	0.199*	0.60 (2)
H10C	1.0450	0.7871	0.3904	0.199*	0.60 (2)

C9'	1.0852 (3)	0.6899 (4)	0.30878 (18)	0.1159 (11)	0.40 (2)
H9'A	1.1328	0.7787	0.3306	0.139*	0.40 (2)
H9'B	1.1314	0.6146	0.2913	0.139*	0.40 (2)
C10'	1.0339 (12)	0.616 (2)	0.3618 (8)	0.111 (4)	0.40 (2)
H10D	1.0935	0.5788	0.4030	0.166*	0.40 (2)
H10E	0.9871	0.5285	0.3392	0.166*	0.40 (2)
H10F	0.9879	0.6918	0.3779	0.166*	0.40 (2)
C11	0.87065 (17)	1.0815 (2)	0.14123 (11)	0.0588 (6)	
C12	0.75990 (17)	1.1280 (2)	0.08755 (11)	0.0618 (6)	
H12A	0.7336	1.2250	0.1045	0.074*	
H12B	0.7036	1.0471	0.0876	0.074*	
C13	0.76341 (16)	1.1525 (2)	0.00975 (11)	0.0565 (5)	
C14	0.6757 (2)	1.2334 (3)	-0.03842 (14)	0.0745 (7)	
H14A	0.6150	1.2716	-0.0228	0.089*	
C15	0.6773 (3)	1.2584 (3)	-0.11014 (15)	0.0918 (8)	
H15A	0.6185	1.3148	-0.1419	0.110*	
C16	0.7651 (2)	1.2003 (3)	-0.13433 (14)	0.0849 (8)	
H16A	0.7663	1.2175	-0.1822	0.102*	
C17	0.8495 (2)	1.1182 (3)	-0.08805 (14)	0.0803 (7)	
H17A	0.9085	1.0766	-0.1045	0.096*	
C18	0.84946 (18)	1.0953 (3)	-0.01624 (12)	0.0706 (6)	
H18A	0.9093	1.0397	0.0151	0.085*	
C19	0.9397 (2)	1.3609 (3)	0.14108 (16)	0.0837 (8)	
H19A	0.9085	1.4167	0.1755	0.100*	
H19B	0.8867	1.3728	0.0928	0.100*	
C20	1.0546 (3)	1.4324 (3)	0.14196 (19)	0.1120 (11)	
H20A	1.0446	1.5452	0.1344	0.134*	
H20B	1.1070	1.4167	0.1902	0.134*	
C21	1.1081 (3)	1.3697 (4)	0.0878 (3)	0.1381 (14)	
H21A	1.1817	1.4177	0.0946	0.207*	
H21B	1.0605	1.3929	0.0396	0.207*	
H21C	1.1169	1.2576	0.0937	0.207*	
H1A	1.0180 (11)	1.163 (2)	0.1924 (9)	0.065 (6)*	

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0861 (12)	0.0898 (12)	0.0851 (13)	0.0094 (9)	-0.0083 (10)	0.0275 (9)
O2	0.0732 (11)	0.0914 (11)	0.0625 (10)	0.0011 (8)	0.0026 (8)	-0.0069 (8)
N1	0.0710 (13)	0.0671 (11)	0.0688 (13)	-0.0032 (9)	0.0099 (11)	0.0003 (9)
C1	0.0817 (16)	0.0716 (13)	0.0667 (15)	-0.0068 (11)	0.0257 (13)	-0.0040 (11)
C2	0.0884 (17)	0.0779 (15)	0.0873 (18)	-0.0122 (13)	0.0156 (15)	-0.0051 (13)
C3	0.0755 (16)	0.0790 (16)	0.107 (2)	-0.0092 (12)	0.0090 (16)	0.0195 (15)
C4	0.0823 (18)	0.119 (2)	0.094 (2)	-0.0134 (16)	0.0297 (17)	0.0302 (18)
C5	0.0798 (17)	0.1103 (19)	0.0628 (15)	-0.0056 (14)	0.0270 (14)	0.0115 (13)
C6	0.0591 (12)	0.0645 (12)	0.0522 (12)	0.0079 (9)	0.0132 (10)	0.0094 (10)
C7	0.0629 (12)	0.0659 (12)	0.0428 (11)	0.0084 (9)	0.0148 (9)	0.0027 (9)
C8	0.0712 (14)	0.0745 (13)	0.0487 (12)	0.0090 (11)	0.0138 (11)	-0.0020 (11)

C9	0.096 (2)	0.133 (2)	0.094 (2)	0.0225 (18)	-0.0132 (18)	0.040 (2)
C10	0.102 (6)	0.194 (10)	0.081 (5)	0.043 (6)	-0.010 (4)	-0.022 (5)
C9'	0.096 (2)	0.133 (2)	0.094 (2)	0.0225 (18)	-0.0132 (18)	0.040 (2)
C10'	0.119 (8)	0.129 (9)	0.060 (6)	-0.016 (7)	-0.015 (5)	0.041 (6)
C11	0.0645 (12)	0.0669 (12)	0.0475 (11)	0.0055 (9)	0.0197 (10)	-0.0032 (9)
C12	0.0585 (12)	0.0668 (12)	0.0617 (13)	0.0101 (9)	0.0196 (10)	0.0044 (10)
C13	0.0568 (12)	0.0509 (10)	0.0576 (12)	-0.0005 (8)	0.0092 (10)	0.0023 (9)
C14	0.0717 (15)	0.0751 (14)	0.0690 (16)	0.0136 (11)	0.0071 (12)	0.0039 (12)
C15	0.095 (2)	0.0952 (18)	0.0692 (17)	0.0136 (14)	-0.0033 (15)	0.0173 (14)
C16	0.0953 (19)	0.0984 (18)	0.0556 (14)	-0.0162 (15)	0.0123 (14)	0.0129 (13)
C17	0.0747 (16)	0.1049 (18)	0.0626 (15)	0.0008 (13)	0.0213 (13)	0.0083 (13)
C18	0.0618 (13)	0.0874 (15)	0.0608 (14)	0.0117 (11)	0.0144 (11)	0.0121 (11)
C19	0.0904 (17)	0.0636 (14)	0.099 (2)	-0.0011 (12)	0.0290 (15)	-0.0085 (13)
C20	0.129 (3)	0.0726 (16)	0.138 (3)	-0.0143 (16)	0.044 (2)	-0.0030 (17)
C21	0.136 (3)	0.100 (2)	0.205 (4)	-0.006 (2)	0.092 (3)	-0.004 (2)

*Geometric parameters (Å, °)*

O1—C8	1.349 (3)	C10'—H10E	0.9600
O1—C9	1.446 (3)	C10'—H10F	0.9600
O2—C8	1.227 (3)	C11—C12	1.499 (3)
N1—C11	1.348 (3)	C12—C13	1.511 (3)
N1—C19	1.458 (3)	C12—H12A	0.9700
N1—H1A	0.905 (10)	C12—H12B	0.9700
C1—C2	1.374 (3)	C13—C18	1.369 (3)
C1—C6	1.383 (3)	C13—C14	1.381 (3)
C1—H1	0.9300	C14—C15	1.390 (4)
C2—C3	1.369 (4)	C14—H14A	0.9300
C2—H2A	0.9300	C15—C16	1.370 (4)
C3—C4	1.355 (4)	C15—H15A	0.9300
C3—H3A	0.9300	C16—C17	1.346 (3)
C4—C5	1.375 (4)	C16—H16A	0.9300
C4—H4A	0.9300	C17—C18	1.384 (3)
C5—C6	1.383 (3)	C17—H17A	0.9300
C5—H5A	0.9300	C18—H18A	0.9300
C6—C7	1.485 (3)	C19—C20	1.522 (3)
C7—C11	1.381 (3)	C19—H19A	0.9700
C7—C8	1.448 (3)	C19—H19B	0.9700
C9—C10	1.399 (7)	C20—C21	1.467 (4)
C9—H9A	0.9700	C20—H20A	0.9700
C9—H9B	0.9700	C20—H20B	0.9700
C10—H10A	0.9600	C21—H21A	0.9600
C10—H10B	0.9600	C21—H21B	0.9600
C10—H10C	0.9600	C21—H21C	0.9600
C10'—H10D	0.9600		
C8—O1—C9	117.8 (2)	N1—C11—C12	116.66 (18)
C11—N1—C19	127.2 (2)	C7—C11—C12	120.63 (19)

C11—N1—H1A	114.9 (13)	C11—C12—C13	116.00 (16)
C19—N1—H1A	117.5 (13)	C11—C12—H12A	108.3
C2—C1—C6	122.2 (2)	C13—C12—H12A	108.3
C2—C1—H1	118.9	C11—C12—H12B	108.3
C6—C1—H1	118.9	C13—C12—H12B	108.3
C3—C2—C1	119.7 (3)	H12A—C12—H12B	107.4
C3—C2—H2A	120.2	C18—C13—C14	117.6 (2)
C1—C2—H2A	120.2	C18—C13—C12	122.92 (19)
C4—C3—C2	119.6 (3)	C14—C13—C12	119.44 (18)
C4—C3—H3A	120.2	C13—C14—C15	120.5 (2)
C2—C3—H3A	120.2	C13—C14—H14A	119.7
C3—C4—C5	120.4 (2)	C15—C14—H14A	119.7
C3—C4—H4A	119.8	C16—C15—C14	120.4 (2)
C5—C4—H4A	119.8	C16—C15—H15A	119.8
C4—C5—C6	121.8 (2)	C14—C15—H15A	119.8
C4—C5—H5A	119.1	C17—C16—C15	119.3 (2)
C6—C5—H5A	119.1	C17—C16—H16A	120.3
C1—C6—C5	116.3 (2)	C15—C16—H16A	120.3
C1—C6—C7	121.66 (17)	C16—C17—C18	120.6 (2)
C5—C6—C7	122.1 (2)	C16—C17—H17A	119.7
C11—C7—C8	120.0 (2)	C18—C17—H17A	119.7
C11—C7—C6	121.21 (19)	C13—C18—C17	121.5 (2)
C8—C7—C6	118.74 (18)	C13—C18—H18A	119.2
O2—C8—O1	121.3 (2)	C17—C18—H18A	119.2
O2—C8—C7	126.3 (2)	N1—C19—C20	110.9 (2)
O1—C8—C7	112.4 (2)	N1—C19—H19A	109.5
C10—C9—O1	115.5 (5)	C20—C19—H19A	109.5
C10—C9—H9A	108.4	N1—C19—H19B	109.5
O1—C9—H9A	108.4	C20—C19—H19B	109.5
C10—C9—H9B	108.4	H19A—C19—H19B	108.0
O1—C9—H9B	108.4	C21—C20—C19	115.9 (3)
H9A—C9—H9B	107.5	C21—C20—H20A	108.3
C9—C10—H10A	109.5	C19—C20—H20A	108.3
C9—C10—H10B	109.5	C21—C20—H20B	108.3
H10A—C10—H10B	109.5	C19—C20—H20B	108.3
C9—C10—H10C	109.5	H20A—C20—H20B	107.4
H10A—C10—H10C	109.5	C20—C21—H21A	109.5
H10B—C10—H10C	109.5	C20—C21—H21B	109.5
H10D—C10'—H10E	109.5	H21A—C21—H21B	109.5
H10D—C10'—H10F	109.5	C20—C21—H21C	109.5
H10E—C10'—H10F	109.5	H21A—C21—H21C	109.5
N1—C11—C7	122.7 (2)	H21B—C21—H21C	109.5
C6—C1—C2—C3	-0.9 (4)	C19—N1—C11—C12	7.4 (3)
C1—C2—C3—C4	0.0 (4)	C8—C7—C11—N1	4.6 (3)
C2—C3—C4—C5	0.6 (4)	C6—C7—C11—N1	-177.32 (17)
C3—C4—C5—C6	-0.3 (4)	C8—C7—C11—C12	-175.14 (16)
C2—C1—C6—C5	1.2 (3)	C6—C7—C11—C12	2.9 (3)

C2—C1—C6—C7	-179.1 (2)	N1—C11—C12—C13	71.9 (2)
C4—C5—C6—C1	-0.5 (4)	C7—C11—C12—C13	-108.3 (2)
C4—C5—C6—C7	179.7 (2)	C11—C12—C13—C18	17.9 (3)
C1—C6—C7—C11	74.3 (3)	C11—C12—C13—C14	-163.29 (19)
C5—C6—C7—C11	-105.9 (2)	C18—C13—C14—C15	-1.5 (3)
C1—C6—C7—C8	-107.6 (2)	C12—C13—C14—C15	179.6 (2)
C5—C6—C7—C8	72.1 (3)	C13—C14—C15—C16	1.2 (4)
C9—O1—C8—O2	2.7 (3)	C14—C15—C16—C17	0.3 (4)
C9—O1—C8—C7	-176.8 (2)	C15—C16—C17—C18	-1.4 (4)
C11—C7—C8—O2	-3.9 (3)	C14—C13—C18—C17	0.4 (3)
C6—C7—C8—O2	178.02 (18)	C12—C13—C18—C17	179.3 (2)
C11—C7—C8—O1	175.63 (17)	C16—C17—C18—C13	1.1 (4)
C6—C7—C8—O1	-2.5 (2)	C11—N1—C19—C20	-154.4 (2)
C8—O1—C9—C10	103.2 (10)	N1—C19—C20—C21	64.4 (4)
C19—N1—C11—C7	-172.39 (19)		

*Hydrogen-bond geometry (Å, °)*

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1A...O2	0.91 (2)	1.93 (2)	2.653 (3)	136 (1)