

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Ethyl 2-phenyl-5,6-dihydropyrrolo-[2,1-a]isoquinoline-3-carboxylate

 Zhao-Peng Yu,^{a*} Ming Zhao^b and Cheng-Tao Feng^c

^aSchool of Chemistry and Chemical Engineering, Linyi University, Linyi, Shandong 276005, People's Republic of China, ^bDepartment of Chemistry, Huainan Union University, Huainan 232038, People's Republic of China, and ^cDepartment of Chemistry, University of Science and Technology of China, 230026 Hefei, People's Republic of China

Correspondence e-mail: yuzp@mail.ustc.edu.cn

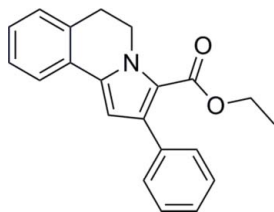
Received 28 May 2012; accepted 31 May 2012

Key indicators: single-crystal X-ray study; $T = 291$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.047; wR factor = 0.117; data-to-parameter ratio = 15.3.

In the title compound, $\text{C}_{21}\text{H}_{19}\text{NO}_2$, the six-membered heterocycle assumes a screw-boat conformation. The phenyl ring is oriented with respect to the pyrrole ring at a dihedral angle of $64.76(10)^\circ$. An intramolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bond helps to stabilize the molecular structure. There are weak $\text{C}-\text{H}\cdots\pi$ interactions between inversion-related molecules in the crystal.

Related literature

For background and applications of lamellarins, see: Bailly (2004); Zou *et al.* (2011). For a related compound, see: Feng *et al.* (2012).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{19}\text{NO}_2$
 $M_r = 317.37$
 Triclinic, $P\bar{1}$

$a = 8.1527(6)$ Å
 $b = 8.4029(6)$ Å
 $c = 12.4220(8)$ Å

$\alpha = 100.117(6)^\circ$
 $\beta = 101.155(5)^\circ$
 $\gamma = 94.312(6)^\circ$
 $V = 816.66(10)$ Å³
 $Z = 2$

Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 291$ K
 $0.42 \times 0.37 \times 0.32$ mm

Data collection

Oxford Diffraction Gemini S Ultra diffractometer
 6799 measured reflections

3340 independent reflections
 2233 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.117$
 $S = 1.02$
 3340 reflections

218 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.14$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the pyrrole ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C8}-\text{H8A}\cdots\text{O1}$	0.97	2.29	2.913 (2)	121
$\text{C8}-\text{H8B}\cdots\text{Cg1}^i$	0.97	2.69	3.6411 (19)	166

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); 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 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

This work was supported by the Natural Science Foundation of the Higher Education Institutions of the Education Bureau of Anhui Province (grant No. KJ2011B154), the Science and Technology Planning Project of Huainan (grant No. 2011 A08101) and Huainan Union University, China (grant No. LYB1005).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5552).

References

- Bailly, C. (2004). *Curr. Med. Chem. Anticancer Agents*, **4**, 363–378.
 Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
 Feng, C.-T., Wang, L.-D., Yan, Y.-G., Liu, J. & Li, S.-H. (2012). *Med. Chem. Res.* **21**, 315–320.
 Oxford Diffraction (2007). *CrysAlis PRO*. Oxford Diffraction Ltd, Abingdon, Oxfordshire, England.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Zou, Y.-Q., Lu, L.-Q., Fu, L., Chang, N.-J., Rong, J., Chen, J.-R. & Xiao, W.-J. (2011). *Angew. Chem. Int. Ed.* **50**, 7171–7175.

supplementary materials

Acta Cryst. (2012). E68, o2021 [doi:10.1107/S1600536812024853]

Ethyl 2-phenyl-5,6-dihydropyrrolo[2,1-*a*]isoquinoline-3-carboxylate**Zhao-Peng Yu, Ming Zhao and Cheng-Tao Feng****Comment**

Lamellarin alkaloids, a new family of marine natural products that contain a pyrrolo[2,1-*a*]isoquinoline core, were found to exhibit a wide spectrum of biological activities (Bailly, 2004). Natural as well as synthetic lamellarins should be excellent candidates for the development of new drugs due to their unique skeletal structure and their important biological activities especially as antitumor agents (Zou *et al.*, 2011). As part of our previous studies concerning anticancer agents, we here report a crystal structure of open chain analogues of lamellarins (Feng *et al.*, 2012).

The conformational analysis show that the conformation of 6-membered hetero-ring is screw boat. The phenyl ring is oriented with respect to the pyrrole ring at 64.76 (10)°. An intramolecular C—H···O hydrogen bond helps to stabilize the molecular structure. There is weak C—H··· π interaction between inversion-related molecules in the crystal.

Experimental

Synthesis or separation ??

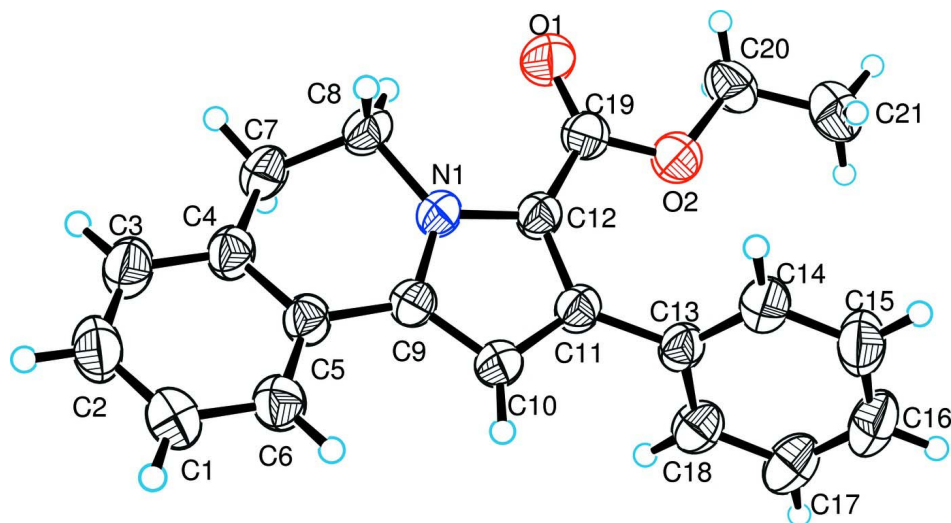
Colourless blocky single crystals of the title compound suitable for X-ray diffraction analysis were obtained by slow evaporation of the mixed solvent ethanol/CH₂Cl₂ (2:1, *v/v*) at room temperature for five days.

Refinement

H atoms were positioned geometrically and allowed to ride on their parent atoms with C—H = 0.93–0.97 Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for the others.

Computing details

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); cell refinement: *CrysAlis PRO* (Oxford Diffraction, 2007); data reduction: *CrysAlis PRO* (Oxford Diffraction, 2007); 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, 1997); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008).

**Figure 1**

Molecular structure of the title compound with displacement ellipsoids drawn at the 45% probability level.

Ethyl 2-phenyl-5,6-dihydropyrrolo[2,1-a]isoquinoline-3-carboxylate

Crystal data

$C_{21}H_{19}NO_2$

$M_r = 317.37$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 8.1527(6)\ \text{\AA}$

$b = 8.4029(6)\ \text{\AA}$

$c = 12.4220(8)\ \text{\AA}$

$\alpha = 100.117(6)^\circ$

$\beta = 101.155(5)^\circ$

$\gamma = 94.312(6)^\circ$

$V = 816.66(10)\ \text{\AA}^3$

$Z = 2$

$F(000) = 336$

$D_x = 1.291\ \text{Mg m}^{-3}$

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

Cell parameters from 1649 reflections

$\theta = 3.3\text{--}29.0^\circ$

$\mu = 0.08\ \text{mm}^{-1}$

$T = 291\ \text{K}$

Block, colourless

$0.42 \times 0.37 \times 0.32\ \text{mm}$

Data collection

Oxford Diffraction Gemini S Ultra diffractometer

Radiation source: Enhance (Mo) X-ray Source

Graphite monochromator

Detector resolution: $15.9149\ \text{pixels mm}^{-1}$

ω scans

6799 measured reflections

3340 independent reflections

2233 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\text{max}} = 26.4^\circ$, $\theta_{\text{min}} = 3.3^\circ$

$h = -9 \rightarrow 10$

$k = -10 \rightarrow 10$

$l = -15 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.117$

$S = 1.02$

3340 reflections

218 parameters

0 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

$w = 1/[\sigma^2(F_o^2) + (0.0468P)^2 + 0.0731P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\text{max}} = 0.14\ \text{e \AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.18\ \text{e \AA}^{-3}$

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.42666 (15)	0.27334 (17)	0.95299 (10)	0.0434 (4)
O1	0.77604 (15)	0.2297 (2)	1.03663 (11)	0.0740 (5)
O2	0.69802 (14)	0.16479 (18)	1.18676 (10)	0.0631 (4)
C5	0.1606 (2)	0.3304 (2)	0.84460 (13)	0.0443 (4)
C18	0.2738 (2)	0.0419 (2)	1.23460 (15)	0.0539 (5)
H18	0.2135	-0.0303	1.1714	0.065*
C8	0.5153 (2)	0.3175 (2)	0.86845 (13)	0.0505 (5)
H8A	0.6167	0.2637	0.8705	0.061*
H8B	0.5475	0.4341	0.8840	0.061*
C12	0.49185 (19)	0.2354 (2)	1.05503 (13)	0.0434 (4)
C13	0.3623 (2)	0.1813 (2)	1.22237 (13)	0.0454 (4)
C10	0.2142 (2)	0.2488 (2)	1.03899 (13)	0.0480 (4)
H10	0.1065	0.2455	1.0537	0.058*
C11	0.3589 (2)	0.2190 (2)	1.10987 (13)	0.0447 (4)
C6	0.0014 (2)	0.3795 (2)	0.84292 (14)	0.0536 (5)
H6	-0.0453	0.3829	0.9058	0.064*
C1	-0.0877 (2)	0.4233 (2)	0.74877 (15)	0.0600 (5)
H1	-0.1942	0.4563	0.7482	0.072*
C9	0.25855 (19)	0.2840 (2)	0.94354 (13)	0.0439 (4)
C17	0.2729 (3)	0.0075 (3)	1.33909 (17)	0.0671 (6)
H17	0.2136	-0.0879	1.3458	0.080*
C4	0.2309 (2)	0.3253 (2)	0.74989 (13)	0.0488 (4)
C20	0.8689 (2)	0.1447 (3)	1.23534 (15)	0.0621 (5)
H20B	0.9460	0.2307	1.2238	0.075*
H20A	0.8983	0.0409	1.2010	0.075*
C16	0.3596 (3)	0.1139 (3)	1.43305 (17)	0.0696 (6)
H16	0.3590	0.0908	1.5035	0.084*
C14	0.4485 (2)	0.2875 (3)	1.31825 (15)	0.0610 (5)
H14	0.5085	0.3829	1.3120	0.073*
C15	0.4469 (3)	0.2539 (3)	1.42279 (15)	0.0708 (6)
H15	0.5052	0.3266	1.4864	0.085*
C7	0.4011 (2)	0.2665 (2)	0.75443 (13)	0.0556 (5)
H7A	0.4538	0.3089	0.7001	0.067*
H7B	0.3873	0.1487	0.7338	0.067*
C19	0.6685 (2)	0.2107 (2)	1.08874 (14)	0.0474 (4)
C3	0.1386 (2)	0.3693 (2)	0.65604 (15)	0.0610 (5)
H3	0.1837	0.3658	0.5925	0.073*

C2	−0.0192 (3)	0.4181 (3)	0.65534 (16)	0.0646 (5)
H2	−0.0796	0.4476	0.5917	0.077*
C21	0.8791 (3)	0.1518 (3)	1.35720 (17)	0.0809 (7)
H21A	0.8039	0.0649	1.3675	0.121*
H21B	0.8479	0.2543	1.3900	0.121*
H21C	0.9921	0.1408	1.3926	0.121*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0427 (7)	0.0525 (9)	0.0368 (8)	0.0057 (6)	0.0130 (6)	0.0080 (6)
O1	0.0462 (7)	0.1197 (14)	0.0634 (9)	0.0135 (8)	0.0206 (6)	0.0257 (8)
O2	0.0443 (7)	0.0917 (11)	0.0587 (8)	0.0135 (7)	0.0087 (6)	0.0292 (7)
C5	0.0485 (9)	0.0444 (11)	0.0373 (9)	0.0012 (8)	0.0068 (7)	0.0047 (7)
C18	0.0594 (11)	0.0530 (12)	0.0541 (11)	0.0124 (9)	0.0195 (9)	0.0124 (9)
C8	0.0518 (10)	0.0596 (12)	0.0454 (10)	0.0072 (9)	0.0218 (8)	0.0115 (9)
C12	0.0434 (9)	0.0488 (11)	0.0396 (9)	0.0073 (8)	0.0117 (7)	0.0090 (8)
C13	0.0432 (9)	0.0564 (12)	0.0411 (10)	0.0138 (8)	0.0133 (7)	0.0130 (8)
C10	0.0416 (9)	0.0614 (12)	0.0429 (10)	0.0067 (8)	0.0130 (7)	0.0107 (8)
C11	0.0452 (9)	0.0502 (11)	0.0405 (9)	0.0077 (8)	0.0122 (7)	0.0094 (8)
C6	0.0510 (10)	0.0624 (13)	0.0461 (10)	0.0072 (9)	0.0094 (8)	0.0080 (9)
C1	0.0551 (11)	0.0656 (14)	0.0553 (11)	0.0085 (9)	0.0009 (9)	0.0125 (10)
C9	0.0432 (9)	0.0487 (11)	0.0395 (9)	0.0042 (8)	0.0108 (7)	0.0057 (8)
C17	0.0864 (14)	0.0631 (14)	0.0653 (14)	0.0175 (11)	0.0318 (11)	0.0274 (11)
C4	0.0575 (10)	0.0492 (11)	0.0375 (9)	0.0010 (8)	0.0090 (8)	0.0057 (8)
C20	0.0442 (10)	0.0721 (15)	0.0652 (13)	0.0141 (9)	0.0003 (9)	0.0098 (10)
C16	0.0869 (15)	0.0860 (17)	0.0510 (12)	0.0336 (13)	0.0261 (11)	0.0304 (12)
C14	0.0657 (12)	0.0681 (14)	0.0482 (11)	−0.0001 (10)	0.0126 (9)	0.0115 (10)
C15	0.0806 (14)	0.0864 (18)	0.0420 (11)	0.0104 (12)	0.0088 (10)	0.0075 (11)
C7	0.0629 (11)	0.0663 (13)	0.0407 (10)	0.0068 (10)	0.0198 (8)	0.0093 (9)
C19	0.0468 (10)	0.0487 (11)	0.0456 (10)	0.0046 (8)	0.0112 (8)	0.0050 (8)
C3	0.0706 (13)	0.0690 (14)	0.0427 (11)	0.0028 (10)	0.0107 (9)	0.0128 (9)
C2	0.0707 (13)	0.0690 (15)	0.0501 (11)	0.0055 (11)	−0.0030 (10)	0.0193 (10)
C21	0.0661 (13)	0.103 (2)	0.0663 (14)	0.0083 (12)	−0.0041 (10)	0.0175 (13)

Geometric parameters (\AA , $^\circ$)

N1—C9	1.3639 (19)	C6—H6	0.9300
N1—C12	1.3775 (19)	C1—C2	1.377 (3)
N1—C8	1.4675 (18)	C1—H1	0.9300
O1—C19	1.2036 (19)	C17—C16	1.371 (3)
O2—C19	1.325 (2)	C17—H17	0.9300
O2—C20	1.442 (2)	C4—C3	1.384 (2)
C5—C6	1.389 (2)	C4—C7	1.502 (2)
C5—C4	1.401 (2)	C20—C21	1.490 (2)
C5—C9	1.461 (2)	C20—H20B	0.9700
C18—C13	1.376 (2)	C20—H20A	0.9700
C18—C17	1.380 (2)	C16—C15	1.367 (3)
C18—H18	0.9300	C16—H16	0.9300
C8—C7	1.508 (2)	C14—C15	1.379 (2)

C8—H8A	0.9700	C14—H14	0.9300
C8—H8B	0.9700	C15—H15	0.9300
C12—C11	1.398 (2)	C7—H7A	0.9700
C12—C19	1.461 (2)	C7—H7B	0.9700
C13—C14	1.385 (2)	C3—C2	1.379 (3)
C13—C11	1.482 (2)	C3—H3	0.9300
C10—C9	1.377 (2)	C2—H2	0.9300
C10—C11	1.396 (2)	C21—H21A	0.9600
C10—H10	0.9300	C21—H21B	0.9600
C6—C1	1.376 (2)	C21—H21C	0.9600
C9—N1—C12	109.29 (12)	C3—C4—C5	118.70 (17)
C9—N1—C8	121.26 (13)	C3—C4—C7	123.36 (15)
C12—N1—C8	129.05 (13)	C5—C4—C7	117.91 (14)
C19—O2—C20	118.09 (13)	O2—C20—C21	107.19 (15)
C6—C5—C4	119.82 (15)	O2—C20—H20B	110.3
C6—C5—C9	121.65 (14)	C21—C20—H20B	110.3
C4—C5—C9	118.53 (15)	O2—C20—H20A	110.3
C13—C18—C17	121.11 (18)	C21—C20—H20A	110.3
C13—C18—H18	119.4	H20B—C20—H20A	108.5
C17—C18—H18	119.4	C15—C16—C17	119.77 (18)
N1—C8—C7	109.17 (13)	C15—C16—H16	120.1
N1—C8—H8A	109.8	C17—C16—H16	120.1
C7—C8—H8A	109.8	C15—C14—C13	120.98 (19)
N1—C8—H8B	109.8	C15—C14—H14	119.5
C7—C8—H8B	109.8	C13—C14—H14	119.5
H8A—C8—H8B	108.3	C16—C15—C14	120.1 (2)
N1—C12—C11	107.38 (13)	C16—C15—H15	120.0
N1—C12—C19	122.42 (13)	C14—C15—H15	120.0
C11—C12—C19	130.12 (15)	C4—C7—C8	112.81 (14)
C18—C13—C14	117.99 (15)	C4—C7—H7A	109.0
C18—C13—C11	120.63 (16)	C8—C7—H7A	109.0
C14—C13—C11	121.33 (16)	C4—C7—H7B	109.0
C9—C10—C11	108.17 (14)	C8—C7—H7B	109.0
C9—C10—H10	125.9	H7A—C7—H7B	107.8
C11—C10—H10	125.9	O1—C19—O2	123.10 (16)
C10—C11—C12	107.05 (14)	O1—C19—C12	125.80 (16)
C10—C11—C13	124.05 (14)	O2—C19—C12	111.10 (14)
C12—C11—C13	128.89 (14)	C2—C3—C4	121.00 (17)
C1—C6—C5	120.41 (16)	C2—C3—H3	119.5
C1—C6—H6	119.8	C4—C3—H3	119.5
C5—C6—H6	119.8	C1—C2—C3	120.08 (17)
C6—C1—C2	119.99 (18)	C1—C2—H2	120.0
C6—C1—H1	120.0	C3—C2—H2	120.0
C2—C1—H1	120.0	C20—C21—H21A	109.5
N1—C9—C10	108.09 (14)	C20—C21—H21B	109.5
N1—C9—C5	120.21 (13)	H21A—C21—H21B	109.5
C10—C9—C5	131.69 (15)	C20—C21—H21C	109.5
C16—C17—C18	120.0 (2)	H21A—C21—H21C	109.5

C16—C17—H17	120.0	H21B—C21—H21C	109.5
C18—C17—H17	120.0		
C9—N1—C8—C7	-36.6 (2)	C4—C5—C9—N1	15.1 (2)
C12—N1—C8—C7	151.50 (17)	C6—C5—C9—C10	13.7 (3)
C9—N1—C12—C11	1.12 (19)	C4—C5—C9—C10	-166.59 (18)
C8—N1—C12—C11	173.80 (16)	C13—C18—C17—C16	0.8 (3)
C9—N1—C12—C19	178.10 (15)	C6—C5—C4—C3	-0.2 (3)
C8—N1—C12—C19	-9.2 (3)	C9—C5—C4—C3	-179.93 (16)
C17—C18—C13—C14	-1.0 (3)	C6—C5—C4—C7	-178.17 (16)
C17—C18—C13—C11	-178.47 (16)	C9—C5—C4—C7	2.1 (2)
C9—C10—C11—C12	-0.5 (2)	C19—O2—C20—C21	160.33 (17)
C9—C10—C11—C13	178.27 (16)	C18—C17—C16—C15	-0.1 (3)
N1—C12—C11—C10	-0.3 (2)	C18—C13—C14—C15	0.5 (3)
C19—C12—C11—C10	-177.02 (18)	C11—C13—C14—C15	177.98 (16)
N1—C12—C11—C13	-179.07 (17)	C17—C16—C15—C14	-0.4 (3)
C19—C12—C11—C13	4.3 (3)	C13—C14—C15—C16	0.2 (3)
C18—C13—C11—C10	63.7 (2)	C3—C4—C7—C8	146.38 (18)
C14—C13—C11—C10	-113.6 (2)	C5—C4—C7—C8	-35.8 (2)
C18—C13—C11—C12	-117.7 (2)	N1—C8—C7—C4	51.0 (2)
C14—C13—C11—C12	64.9 (3)	C20—O2—C19—O1	3.1 (3)
C4—C5—C6—C1	0.0 (3)	C20—O2—C19—C12	-176.31 (15)
C9—C5—C6—C1	179.69 (17)	N1—C12—C19—O1	4.8 (3)
C5—C6—C1—C2	0.1 (3)	C11—C12—C19—O1	-178.94 (18)
C12—N1—C9—C10	-1.46 (19)	N1—C12—C19—O2	-175.75 (15)
C8—N1—C9—C10	-174.81 (15)	C11—C12—C19—O2	0.5 (3)
C12—N1—C9—C5	177.22 (15)	C5—C4—C3—C2	0.3 (3)
C8—N1—C9—C5	3.9 (2)	C7—C4—C3—C2	178.15 (17)
C11—C10—C9—N1	1.2 (2)	C6—C1—C2—C3	0.0 (3)
C11—C10—C9—C5	-177.24 (18)	C4—C3—C2—C1	-0.2 (3)
C6—C5—C9—N1	-164.60 (16)		

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

Cg1 is the centroid of the pyrrole ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C8—H8A \cdots O1	0.97	2.29	2.913 (2)	121
C8—H8B \cdots Cg1 ⁱ	0.97	2.69	3.6411 (19)	166

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