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4-Chloro-5-(dimethylamino)-2-[(5-phenyl-1,3,4-oxadiazol-2-yl)methyl]pyridazin-3(2H)-one

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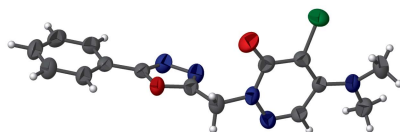
Keywords: crystal structure; pyridazinone; oxadiazole; hydrogen bonding.

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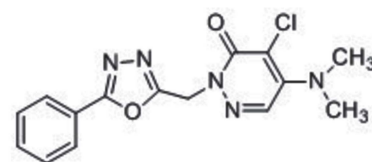
Structural data: full structural data are available from iucrdata.iucr.org

In the structure of the title compound, C₁₅H₁₄ClN₅O₂, the terminal phenyl ring and the adjacent furan ring subtend a dihedral angle of 6.77 (17)°. The 4-chloro-5-(dimethylamino)-pyridazin-3(2H)-one group is linked to the oxadiazole ring by a methylene bridge, and the dihedral angle between the pyridazine and oxadiazole rings is 88.66 (14)°. In the crystal, C—H···O and C—H···N hydrogen bonds extend the structure into a three-dimensional network.

3D view



Chemical scheme



Structure description

Pyridazinones have attracted increasing attention as a scaffold because of their wide spectrum of biological activity (Zou *et al.*, 2002). Along with the development of design and synthetic methodology, pyridazinone derivatives have been widely applied in medicinal and agricultural chemistry (Arora *et al.*, 2022; Vaidergorn *et al.*, 2021; Zhang *et al.*, 2020; Lu *et al.*, 2017; Cao *et al.*, 2005; Xu *et al.*, 2008; Sun *et al.*, 2015). As part of our work in this area, a series of pyridazinone derivatives containing an oxadiazole moiety have been designed and synthesized, and we report here the crystal structure of the title compound.

The molecular structure of the title compound is shown in Fig. 1. The phenyl (C1–C6) and oxadiazole (O1/N1/N2/C7/C8) rings are almost coplanar, subtending a dihedral angle of 6.77 (17)°. The pyridazine ring is almost perpendicular to oxadiazole ring, making a dihedral angle of 88.66 (14)°. The dihedral angle between the phenyl and pyridazine rings is 82.01 (17)°.

The crystal packing is characterized by C—H···N and C—H···O contacts (Fig. 2, Table 1).



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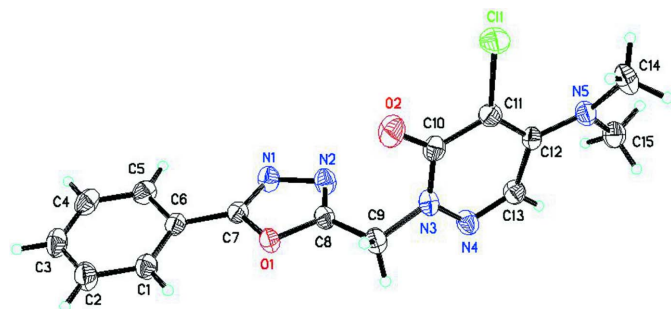


Figure 1
The molecular structure of the title compound, with the atom labelling and displacement ellipsoids drawn at the 50% probability level.

Synthesis and crystallization

To a 100 ml round-bottom flask, 4,5-dichloro-2-[(5-phenyl-1,3,4-oxadiazol-2-yl) methyl]pyridazin-3(2H)-one (1.0 g, 3.1 mmol), dimethylamine (0.6 ml, 6.2 mmol), and potassium carbonate (0.86 g, 6.2 mmol) were added in 30 ml of DMF and stirred for 8 h at 353 K. Afterwards, the reaction mixture was cooled and poured into 60 ml of ice-water. The precipitate formed was collected by filtration and then dried to obtain the pure title compound (yield 0.56 g, 54.6%). It was recrystallized from mixed solvents of ethyl acetate and petroleum (3:5) to give crystals suitable for X-ray diffraction (m.p. 427–429 K).

¹H NMR (400 MHz, CDCl₃): 3.18 (s, 6H), 5.62 (s, 2H), 7.54 (m, 3H), 7.67 (s, 1H), 8.06 (dd, 2H, *J* = 7.6 Hz, *J* = 1.6 Hz). IR (KBr, cm⁻¹): 2973, 2937, 1632, 1600, 1520, 1482, 1449, 1216, 1146, 752, 710.

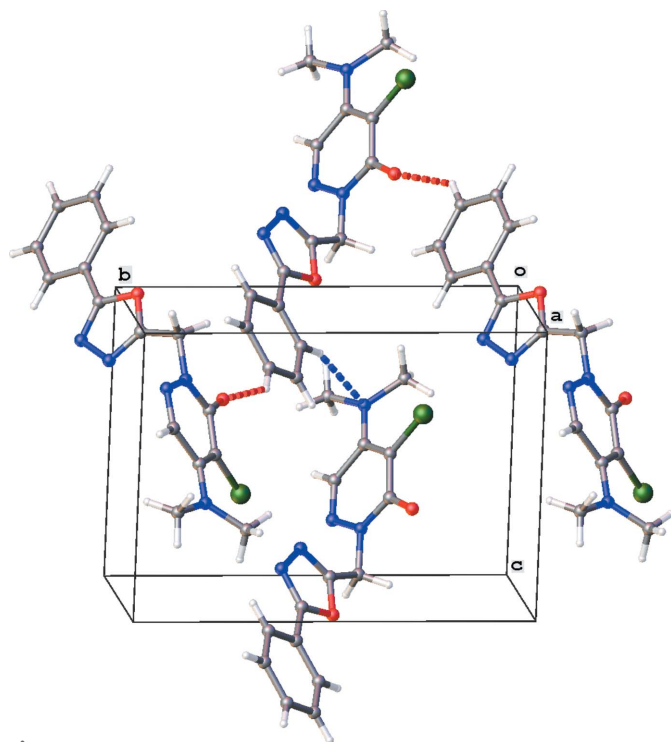


Figure 2
The crystal packing of the title compound. The C–H···N and C–H···O hydrogen bonds are shown as dashed lines (see also Table 1).

Table 1
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>
C3–H3···O2 ⁱ	0.93	2.36	3.125 (5)	140
C1–H1···N5 ⁱⁱ	0.93	2.61	3.477 (5)	156

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

Table 2
Experimental details.

Crystal data	
Chemical formula	C ₁₅ H ₁₄ ClN ₅ O ₂
<i>M</i> _r	331.76
Crystal system, space group	Monoclinic, <i>P</i> 2 ₁
Temperature (K)	293
<i>a</i> , <i>b</i> , <i>c</i> (Å)	7.1533 (16), 11.936 (3), 9.004 (2)
β (°)	98.128 (5)
<i>V</i> (Å ³)	761.1 (3)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	0.27
Crystal size (mm)	0.22 × 0.17 × 0.12
Data collection	
Diffractometer	Bruker SMART CCD area detector
Absorption correction	Multi-scan (SADABS; Bruker, 2002)
<i>T</i> _{min} , <i>T</i> _{max}	0.580, 0.746
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	4542, 2915, 2673
<i>R</i> _{int}	0.030
(sin θ/λ) _{max} (Å ⁻¹)	0.617
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.042, 0.100, 1.04
No. of reflections	2915
No. of parameters	211
No. of restraints	1
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³)	0.28, -0.30
Absolute structure	Flack <i>x</i> determined using 1139 quotients [(<i>I</i> ⁺) – (<i>I</i> [−])] / [(<i>I</i> ⁺) + (<i>I</i> [−])] (Parsons <i>et al.</i> , 2013)
Absolute structure parameter	0.00 (4)

Computer programs: SMART and SAINT (Bruker, 2002), SHELXTL (Sheldrick, 2008) and SHELXL2013 (Sheldrick, 2015).

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2.

Funding information

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References

- Arora, S., Joshi, G., Chaturvedi, A., Heuser, M., Patil, S. & Kumar, R. (2022). *J. Med. Chem.* **65**, 1171–1205.
 Bruker (2002). SMART, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
 Cao, S., Wei, N., Zhao, C. M., Li, L. N., Huang, Q. C. & Qian, X. H. (2005). *J. Agric. Food Chem.* **53**, 3120–3125.

- Lu, D., Liu, F. F., Xing, W. Q., Tong, X. K., Wang, L., Wang, Y. J., Zeng, L. M., Feng, C. L., Yang, L., Zuo, J. P. & Hu, Y. H. (2017). *Infect. Dis.* **3**, 199–205.
- Parsons, S., Flack, H. D. & Wagner, T. (2013). *Acta Cryst.* **B69**, 249–259.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Sheldrick, G. M. (2015). *Acta Cryst.* **C71**, 3–8.
- Sun, R., Liu, C. J., Zhang, H. & Wang, Q. M. (2015). *J. Agric. Food Chem.* **63**, 6847–6865.
- Vaidergorn, M. M., da Silva Emery, F. & Ganesan, A. (2021). *J. Med. Chem.* **64**, 13980–14010.
- Xu, H., Hu, X. H., Zou, X. M., Liu, B., Zhu, Y. Q., Wang, Y., Hu, F. Z. & Yang, H. Z. (2008). *J. Agric. Food Chem.* **56**, 6567–6572.
- Zhang, X. J., Sheng, X. J., Shen, J., Zhang, S. B., Sun, W. J., Shen, C. L., Li, Y., Wang, J., Lv, H., Cui, M., Zhu, Y., Huang, L., Hao, D., Qi, Z., Sun, G., Mao, W., Pan, Y., Shen, L., Li, X., Hu, G., Gong, Z., Han, S., Li, J., Chen, S., Tu, R., Wang, X. & Wu, C. (2020). *ACS Med. Chem. Lett.* **11**, 1863–1868.
- Zou, X. J., Lai, L. H., Jin, G. Y. & Zhang, Z. X. (2002). *J. Agric. Food Chem.* **50**, 3757–3760.

full crystallographic data

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4-Chloro-5-(dimethylamino)-2-[(5-phenyl-1,3,4-oxadiazol-2-yl)methyl]-pyridazin-3(2H)-one

Jingjing Song, Xinyu Jiang, Ziyi Wang, Jingyao Pei and Hongsen Li

4-Chloro-5-(dimethylamino)-2-[(5-phenyl-1,3,4-oxadiazol-2-yl)methyl]pyridazin-3(2H)-one

Crystal data

C₁₅H₁₄ClN₅O₂

M_r = 331.76

Monoclinic, *P*2₁

a = 7.1533 (16) Å

b = 11.936 (3) Å

c = 9.004 (2) Å

β = 98.128 (5)°

V = 761.1 (3) Å³

Z = 2

F(000) = 344

D_x = 1.448 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 2107 reflections

θ = 5.7–49.7°

μ = 0.27 mm⁻¹

T = 293 K

Prismatic, colorless

0.22 × 0.17 × 0.12 mm

Data collection

Bruker SMART CCD area detector
diffractometer

phi and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2002)

T_{min} = 0.580, *T_{max}* = 0.746

4542 measured reflections

2915 independent reflections

2673 reflections with *I* > 2σ(*I*)

R_{int} = 0.030

θ_{\max} = 26.0°, θ_{\min} = 2.3°

h = -8→8

k = -14→14

l = -9→11

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.042

wR (*F*²) = 0.100

S = 1.04

2915 reflections

211 parameters

1 restraint

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.0624*P*)²]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} < 0.001

Δρ_{max} = 0.28 e Å⁻³

Δρ_{min} = -0.30 e Å⁻³

Extinction correction: SHELXL,

*F_c** = *kF_c*[1 + 0.001 × *F_c*²λ³/sin(2θ)]^{-1/4}

Extinction coefficient: 0.029 (8)

Absolute structure: Flack *x* determined using

1139 quotients [(*I*⁺) - (*I*⁻)] / [(*I*⁺) + (*I*⁻)] (Parsons *et al.*, 2013)

Absolute structure parameter: 0.00 (4)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.85200 (15)	0.73449 (8)	0.58094 (11)	0.0698 (3)
N1	0.3672 (4)	1.1087 (2)	0.1248 (3)	0.0522 (7)
N2	0.5386 (4)	1.0502 (2)	0.1672 (3)	0.0525 (7)
N3	0.8278 (4)	0.8804 (2)	0.1888 (3)	0.0450 (6)
N4	0.9890 (4)	0.9391 (2)	0.1928 (3)	0.0503 (7)
N5	1.2442 (4)	0.8512 (2)	0.5493 (3)	0.0501 (7)
O1	0.3908 (3)	0.97490 (18)	-0.0393 (2)	0.0446 (5)
O2	0.6230 (4)	0.7702 (2)	0.2898 (3)	0.0695 (8)
C1	0.0415 (5)	1.0303 (3)	-0.2139 (4)	0.0530 (8)
H1	0.1196	0.9776	-0.2501	0.064*
C2	-0.1339 (5)	1.0519 (4)	-0.2928 (4)	0.0629 (10)
H2	-0.1736	1.0139	-0.3818	0.075*
C3	-0.2509 (5)	1.1296 (3)	-0.2406 (5)	0.0619 (10)
H3	-0.3693	1.1446	-0.2939	0.074*
C4	-0.1906 (6)	1.1849 (3)	-0.1086 (5)	0.0605 (9)
H4	-0.2695	1.2372	-0.0726	0.073*
C5	-0.0155 (5)	1.1641 (3)	-0.0288 (4)	0.0526 (8)
H5	0.0237	1.2023	0.0601	0.063*
C6	0.1024 (4)	1.0857 (3)	-0.0819 (3)	0.0425 (7)
C7	0.2862 (4)	1.0612 (2)	0.0054 (3)	0.0406 (7)
C8	0.5443 (4)	0.9750 (3)	0.0679 (3)	0.0422 (7)
C9	0.6961 (5)	0.8919 (3)	0.0516 (4)	0.0502 (8)
H9A	0.6391	0.8196	0.0245	0.060*
H9B	0.7642	0.9154	-0.0289	0.060*
C10	0.7789 (4)	0.8150 (3)	0.3034 (4)	0.0453 (7)
C11	0.9230 (5)	0.8076 (3)	0.4331 (3)	0.0449 (7)
C12	1.0946 (4)	0.8589 (3)	0.4373 (3)	0.0396 (6)
C13	1.1132 (5)	0.9282 (3)	0.3105 (4)	0.0473 (8)
H13	1.2245	0.9690	0.3132	0.057*
C14	1.2910 (6)	0.7498 (4)	0.6345 (4)	0.0692 (11)
H14A	1.2811	0.7631	0.7382	0.104*
H14B	1.4178	0.7277	0.6249	0.104*
H14C	1.2052	0.6912	0.5968	0.104*
C15	1.4001 (5)	0.9309 (4)	0.5556 (4)	0.0636 (10)
H15A	1.4852	0.9073	0.4882	0.095*
H15B	1.4664	0.9339	0.6559	0.095*
H15C	1.3513	1.0038	0.5266	0.095*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0676 (6)	0.0740 (6)	0.0695 (6)	−0.0110 (5)	0.0159 (4)	0.0263 (5)
N1	0.0468 (16)	0.0478 (15)	0.0600 (17)	0.0006 (13)	0.0010 (13)	−0.0103 (13)
N2	0.0476 (16)	0.0503 (15)	0.0563 (15)	−0.0009 (13)	−0.0047 (12)	−0.0085 (14)
N3	0.0368 (14)	0.0436 (14)	0.0518 (14)	0.0001 (11)	−0.0033 (11)	0.0016 (12)
N4	0.0466 (15)	0.0496 (15)	0.0525 (15)	−0.0048 (12)	−0.0002 (13)	0.0126 (12)
N5	0.0424 (15)	0.0559 (15)	0.0498 (15)	0.0020 (12)	−0.0017 (12)	0.0040 (12)
O1	0.0411 (12)	0.0456 (11)	0.0444 (11)	0.0056 (10)	−0.0034 (9)	−0.0044 (9)
O2	0.0435 (14)	0.0800 (19)	0.0831 (17)	−0.0224 (13)	0.0021 (12)	0.0027 (14)
C1	0.0453 (19)	0.057 (2)	0.0552 (19)	0.0100 (16)	0.0037 (15)	−0.0011 (16)
C2	0.050 (2)	0.070 (2)	0.066 (2)	0.0062 (18)	−0.0032 (17)	0.0008 (19)
C3	0.0411 (19)	0.063 (2)	0.079 (3)	0.0094 (17)	0.0004 (17)	0.0143 (19)
C4	0.051 (2)	0.0516 (19)	0.083 (3)	0.0150 (17)	0.0229 (18)	0.0122 (18)
C5	0.054 (2)	0.0462 (18)	0.0600 (19)	0.0089 (15)	0.0150 (16)	0.0052 (14)
C6	0.0381 (16)	0.0404 (15)	0.0499 (16)	0.0022 (13)	0.0093 (13)	0.0089 (12)
C7	0.0404 (15)	0.0377 (15)	0.0444 (15)	0.0012 (13)	0.0085 (13)	0.0039 (13)
C8	0.0393 (16)	0.0423 (15)	0.0428 (15)	−0.0037 (13)	−0.0015 (12)	−0.0004 (13)
C9	0.0467 (18)	0.0508 (19)	0.0503 (17)	0.0055 (15)	−0.0028 (14)	−0.0043 (15)
C10	0.0389 (17)	0.0411 (15)	0.0557 (18)	−0.0027 (13)	0.0060 (14)	0.0000 (14)
C11	0.0470 (18)	0.0407 (15)	0.0478 (16)	0.0018 (14)	0.0099 (14)	0.0076 (13)
C12	0.0362 (15)	0.0386 (14)	0.0433 (15)	0.0041 (12)	0.0034 (12)	−0.0002 (12)
C13	0.0387 (17)	0.0497 (17)	0.0526 (18)	−0.0100 (14)	0.0033 (15)	0.0062 (15)
C14	0.068 (3)	0.071 (2)	0.064 (2)	0.015 (2)	−0.0101 (18)	0.013 (2)
C15	0.0431 (19)	0.085 (3)	0.060 (2)	−0.0084 (19)	−0.0026 (16)	−0.0018 (19)

Geometric parameters (Å, °)

C11—C11	1.727 (3)	C3—H3	0.9300
N1—C7	1.279 (4)	C4—C5	1.376 (6)
N1—N2	1.415 (4)	C4—H4	0.9300
N2—C8	1.272 (4)	C5—C6	1.389 (5)
N3—N4	1.345 (4)	C5—H5	0.9300
N3—C10	1.378 (4)	C6—C7	1.462 (4)
N3—C9	1.450 (4)	C8—C9	1.494 (5)
N4—C13	1.290 (4)	C9—H9A	0.9700
N5—C12	1.366 (4)	C9—H9B	0.9700
N5—C14	1.447 (5)	C10—C11	1.447 (4)
N5—C15	1.461 (5)	C11—C12	1.367 (5)
O1—C8	1.355 (3)	C12—C13	1.432 (4)
O1—C7	1.367 (4)	C13—H13	0.9300
O2—C10	1.227 (4)	C14—H14A	0.9600
C1—C6	1.376 (5)	C14—H14B	0.9600
C1—C2	1.376 (5)	C14—H14C	0.9600
C1—H1	0.9300	C15—H15A	0.9600
C2—C3	1.376 (6)	C15—H15B	0.9600
C2—H2	0.9300	C15—H15C	0.9600

C3—C4	1.374 (6)		
C7—N1—N2	106.3 (3)	N2—C8—C9	129.4 (3)
C8—N2—N1	105.7 (2)	O1—C8—C9	117.2 (3)
N4—N3—C10	125.6 (3)	N3—C9—C8	111.9 (3)
N4—N3—C9	115.2 (3)	N3—C9—H9A	109.2
C10—N3—C9	119.2 (3)	C8—C9—H9A	109.2
C13—N4—N3	117.3 (3)	N3—C9—H9B	109.2
C12—N5—C14	123.1 (3)	C8—C9—H9B	109.2
C12—N5—C15	119.9 (3)	H9A—C9—H9B	107.9
C14—N5—C15	114.0 (3)	O2—C10—N3	119.7 (3)
C8—O1—C7	102.2 (2)	O2—C10—C11	126.0 (3)
C6—C1—C2	120.7 (3)	N3—C10—C11	114.3 (3)
C6—C1—H1	119.7	C12—C11—C10	122.0 (3)
C2—C1—H1	119.7	C12—C11—Cl1	124.4 (2)
C3—C2—C1	120.3 (4)	C10—C11—Cl1	113.5 (2)
C3—C2—H2	119.9	N5—C12—C11	126.6 (3)
C1—C2—H2	119.9	N5—C12—C13	118.4 (3)
C4—C3—C2	119.2 (4)	C11—C12—C13	114.9 (3)
C4—C3—H3	120.4	N4—C13—C12	125.6 (3)
C2—C3—H3	120.4	N4—C13—H13	117.2
C3—C4—C5	121.1 (3)	C12—C13—H13	117.2
C3—C4—H4	119.5	N5—C14—H14A	109.5
C5—C4—H4	119.5	N5—C14—H14B	109.5
C4—C5—C6	119.6 (4)	H14A—C14—H14B	109.5
C4—C5—H5	120.2	N5—C14—H14C	109.5
C6—C5—H5	120.2	H14A—C14—H14C	109.5
C1—C6—C5	119.2 (3)	H14B—C14—H14C	109.5
C1—C6—C7	121.3 (3)	N5—C15—H15A	109.5
C5—C6—C7	119.5 (3)	N5—C15—H15B	109.5
N1—C7—O1	112.3 (3)	H15A—C15—H15B	109.5
N1—C7—C6	129.0 (3)	N5—C15—H15C	109.5
O1—C7—C6	118.7 (3)	H15A—C15—H15C	109.5
N2—C8—O1	113.4 (3)	H15B—C15—H15C	109.5
C7—N1—N2—C8	1.0 (4)	N4—N3—C9—C8	-98.6 (4)
C10—N3—N4—C13	4.2 (5)	C10—N3—C9—C8	79.8 (4)
C9—N3—N4—C13	-177.5 (3)	N2—C8—C9—N3	16.7 (5)
C6—C1—C2—C3	0.0 (6)	O1—C8—C9—N3	-165.6 (3)
C1—C2—C3—C4	-0.2 (6)	N4—N3—C10—O2	175.9 (3)
C2—C3—C4—C5	0.3 (6)	C9—N3—C10—O2	-2.2 (4)
C3—C4—C5—C6	-0.2 (5)	N4—N3—C10—C11	-3.3 (4)
C2—C1—C6—C5	0.1 (5)	C9—N3—C10—C11	178.6 (3)
C2—C1—C6—C7	178.2 (3)	O2—C10—C11—C12	178.8 (3)
C4—C5—C6—C1	0.0 (5)	N3—C10—C11—C12	-2.0 (4)
C4—C5—C6—C7	-178.1 (3)	O2—C10—C11—Cl1	-4.0 (4)
N2—N1—C7—O1	-0.9 (4)	N3—C10—C11—Cl1	175.2 (2)
N2—N1—C7—C6	177.5 (3)	C14—N5—C12—C11	35.7 (5)

C8—O1—C7—N1	0.5 (3)	C15—N5—C12—C11	-165.2 (3)
C8—O1—C7—C6	-178.1 (3)	C14—N5—C12—C13	-145.3 (3)
C1—C6—C7—N1	176.0 (3)	C15—N5—C12—C13	13.8 (5)
C5—C6—C7—N1	-5.9 (5)	C10—C11—C12—N5	-175.4 (3)
C1—C6—C7—O1	-5.7 (4)	C11—C11—C12—N5	7.8 (5)
C5—C6—C7—O1	172.5 (3)	C10—C11—C12—C13	5.6 (4)
N1—N2—C8—O1	-0.7 (4)	C11—C11—C12—C13	-171.3 (2)
N1—N2—C8—C9	177.1 (3)	N3—N4—C13—C12	0.1 (5)
C7—O1—C8—N2	0.2 (4)	N5—C12—C13—N4	176.0 (3)
C7—O1—C8—C9	-177.9 (3)	C11—C12—C13—N4	-4.9 (5)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C14—H14C \cdots C11	0.96	2.56	3.114 (4)	117
C1—H1 \cdots O1	0.93	2.52	2.836 (4)	100
C3—H3 \cdots O2 ⁱ	0.93	2.36	3.125 (5)	140
C1—H1 \cdots N5 ⁱⁱ	0.93	2.61	3.477 (5)	156

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