

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Methyl 2-{*N*-[2-(2,4-dichlorophenoxy)-acetyl]-4-[(4,6-dimethoxypyrimidin-2-yl)-oxy]anilino}propanoate

Lihong Ning, Hao Peng and Hongwu He*

Key Laboratory of Pesticide and Chemical Biology, College of Chemistry, Central China Normal University, Wuhan 430079, People's Republic of China.

Correspondence e-mail: he1208@mail.ccnu.edu.cn

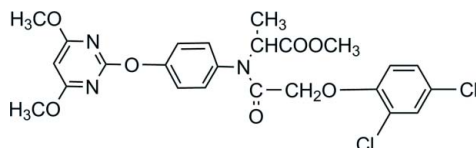
Received 25 May 2012; accepted 5 June 2012

Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.056; wR factor = 0.174; data-to-parameter ratio = 18.9.

In the title molecule, $\text{C}_{24}\text{H}_{23}\text{Cl}_2\text{N}_3\text{O}_7$, the central benzene ring forms dihedral angles of 65.71 (1) and 44.42 (1)° with the pyrimidine ring and the terminal benzene ring, respectively. In the crystal, molecules are linked *via* $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For reference bond-length data, see: Allen *et al.* (1987). For the synthesis of 4-(4,6-dimethoxypyrimidin-2-yloxy)benzenamine, see: Jin *et al.* (2011). For biological properties of fungicides, see: Gozzo & Garlaschelli (1985).



Experimental

Crystal data

 $\text{C}_{24}\text{H}_{23}\text{Cl}_2\text{N}_3\text{O}_7$ $M_r = 536.35$ Triclinic, $P\bar{1}$ $a = 8.2438$ (9) Å $b = 11.2405$ (12) Å $c = 14.2502$ (15) Å $\alpha = 85.178$ (2)° $\beta = 78.702$ (2)° $\gamma = 80.032$ (2)° $V = 1273.6$ (2) Å³ $Z = 2$ Mo $K\alpha$ radiation $\mu = 0.30$ mm⁻¹
 $T = 298$ K $0.16 \times 0.12 \times 0.10$ mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer
15581 measured reflections6204 independent reflections
4390 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.050$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.056$ $wR(F^2) = 0.174$ $S = 1.06$

6204 reflections

329 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.77$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.35$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C9}-\text{H9}\cdots\text{O4}^i$	0.93	2.57	3.402 (3)	150

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

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; 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: *PLATON*.

We gratefully acknowledge financial support of this work by the National Basic Research Program of China (2010CB126100) and the National Natural Science Foundation of China (Nos. 21172090 and 21002037). The research was supported in part by the PCSIRT (No. IRT0953) and the self-determined research funds of CCNU from the college's basic research and operation of MOE (No. CCNU10A01007).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WN2477).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
- Bruker (2001). *SMART* and *SAINT-Plus*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Gozzo, F. & Garlaschelli, L. (1985). *Pestic. Sci.* **16**, 227–286.
- Jin, C. F., Liang, Y. J., He, H. W. & Fu, L. W. (2011). *Eur. J. Med. Chem.* **46**, 429–432.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supporting information

Acta Cryst. (2012). E68, o2046 [https://doi.org/10.1107/S1600536812025494]

Methyl 2-{*N*-[2-(2,4-dichlorophenoxy)acetyl]-4-[(4,6-dimethoxypyrimidin-2-yl)-oxy]anilino}propanoate

Lihong Ning, Hao Peng and Hongwu He

S1. Comment

N-acylalanine fungicides are mainly used in crop protection because of their systemic properties, with both curative and protective activity against fungal pathogens of the Peronosporales (Gozzo & Garlaschelli, 1985). The pyrimidinyl group is widely used in fungicides and drug molecular design (Jin *et al.*, 2011), so we have introduced the pyrimidinyl group into acylalanine derivatives in order to decrease resistance and increase activity.

Here we report the crystal structure of the title compound (Fig. 1). The bond lengths (Allen *et al.*, 1987) and angles show normal values. The central benzene ring forms dihedral angles of 65.71 (1)° and 44.42 (1)° with the pyrimidinyl ring and the terminal benzene ring, respectively. The C9—H9···O4 intermolecular hydrogen bond plays an important role in determining the crystal structure.

S2. Experimental

4-(4,6-Dimethoxypyrimidin-2-yloxy)benzenamine (Jin *et al.*, 2011) (1 mmol) and methyl 2-chloropropanoate (1.2 mmol) were dissolved in 15 ml dimethylformamide, then 1 mmol K₂CO₃ was added with constant stirring. The temperature was maintained at 100 °C for 10 h. The reaction mixture was poured into water and extracted with ethyl acetate, dried with Na₂SO₄, then purified by column chromatography on silica gel with petroleum ether/ethyl acetate (4:1) to give the compound methyl 2-(*N*-(4-(4,6-dimethoxypyrimidin-2-yloxy)phenyl)amino)propanoate.

To a mixture of methyl 2-(*N*-(4-(4,6-dimethoxypyrimidin-2-yloxy) phenyl)amino)propanoate (2 mmol) and triethylamine (2 mmol) in CH₂Cl₂ (20 ml), 2,4-dichlorophenoxyacetyl chloride (2 mmol) was added at 2–5 °C and the mixture was stirred for another 3 h, then washed with saturated sodium hydrogen carbonate solution and dried with Na₂SO₄. The residue was purified by column chromatography on silica gel with petroleum ether/ethyl acetate (3:1) to give the pure title compound as a white solid. Recrystallization from ethanol over a period of one week gave colourless crystals of the title compound.

S3. Refinement

H atoms were geometrically positioned (Csp²—H = 0.93 Å, Cmethine—H = 0.98 Å, Cmethylene—H = 0.97 Å, Cmethyl—H = 0.96 Å) and refined as riding, with U_{iso}(H) = xU_{eq}(C), where x = 1.5 for methyl H and 1.2 for all other H atoms.

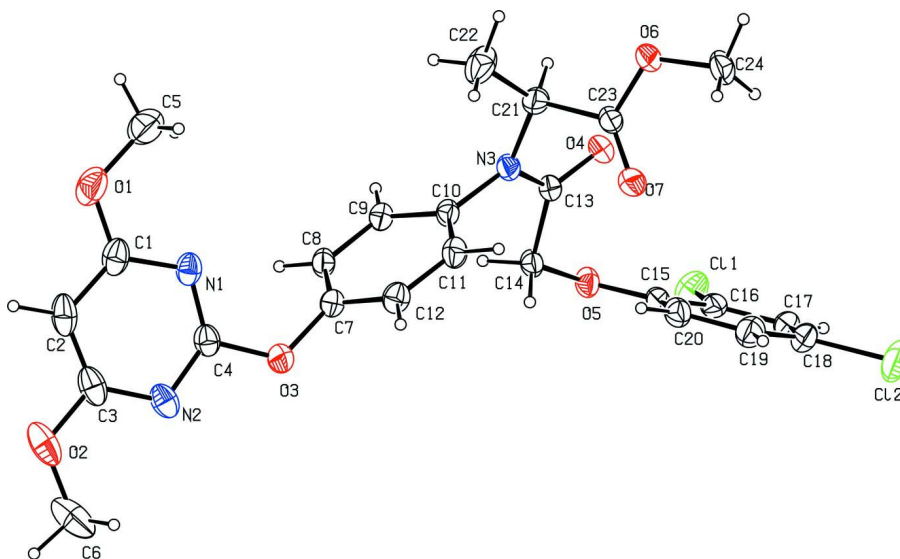


Figure 1

Molecular structure of the title compound, with 50% probability displacement ellipsoids. H atoms are shown as spheres of arbitrary radius.

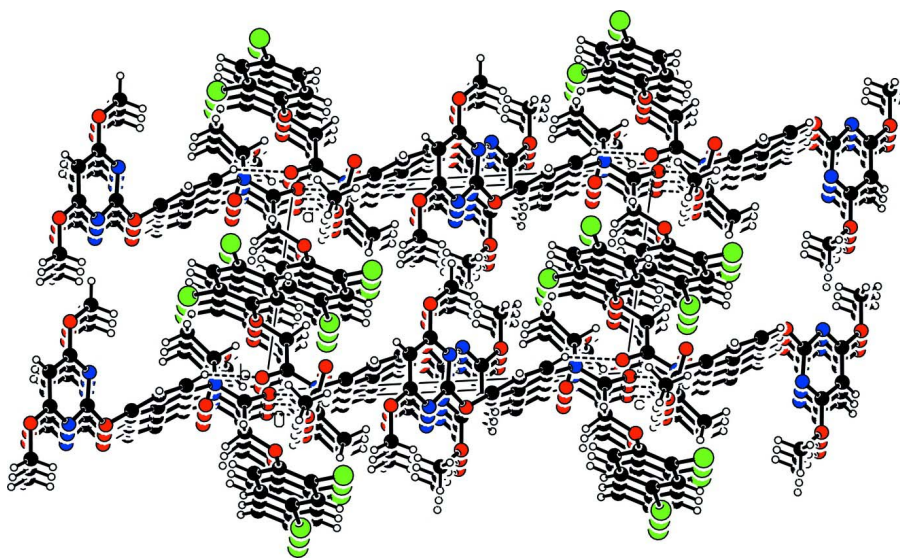


Figure 2

Part of the crystal packing, showing the intermolecular hydrogen bonds as dashed lines.

Methyl 2-[N-[2-(2,4-dichlorophenoxy)acetyl]-4-[(4,6-dimethoxypyrimidin-2-yl)oxy]anilino]propanoate

Crystal data

$C_{24}H_{23}Cl_2N_3O_7$

$M_r = 536.35$

Triclinic, $P\bar{1}$

$a = 8.2438(9) \text{ \AA}$

$b = 11.2405(12) \text{ \AA}$

$c = 14.2502(15) \text{ \AA}$

$\alpha = 85.178(2)^\circ$

$\beta = 78.702(2)^\circ$

$\gamma = 80.032(2)^\circ$

$V = 1273.6(2) \text{ \AA}^3$

$Z = 2$

$F(000) = 556$

$D_x = 1.399 \text{ Mg m}^{-3}$

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

Cell parameters from 5848 reflections
 $\theta = 2.4\text{--}27.7^\circ$
 $\mu = 0.30\text{ mm}^{-1}$

$T = 298\text{ K}$
 Block, colourless
 $0.16 \times 0.12 \times 0.10\text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 phi and ω scans
 15581 measured reflections
 6204 independent reflections

4390 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.050$
 $\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 1.8^\circ$
 $h = -10 \rightarrow 10$
 $k = -14 \rightarrow 14$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.174$
 $S = 1.06$
 6204 reflections
 329 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
 $w = 1/[\sigma^2(F_o^2) + (0.0812P)^2 + 0.2621P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.77\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.35\text{ e \AA}^{-3}$

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	−0.1814 (4)	0.1820 (3)	0.55913 (16)	0.0750 (7)
C2	−0.1019 (5)	0.0866 (3)	0.60971 (18)	0.0888 (9)
H2	−0.1613	0.0343	0.6514	0.107*
C3	0.0693 (5)	0.0735 (2)	0.59483 (17)	0.0809 (8)
C4	0.0665 (3)	0.2312 (2)	0.49075 (15)	0.0619 (5)
C5	−0.4294 (4)	0.3007 (4)	0.5202 (3)	0.1131 (12)
H5A	−0.4154	0.3749	0.5442	0.170*
H5B	−0.5467	0.2964	0.5285	0.170*
H5C	−0.3805	0.2983	0.4533	0.170*
C6	0.3282 (7)	−0.0322 (4)	0.6277 (3)	0.1322 (16)
H6A	0.3745	−0.0543	0.5632	0.198*
H6B	0.3690	−0.0945	0.6716	0.198*
H6C	0.3610	0.0426	0.6385	0.198*
C7	0.0948 (3)	0.3856 (2)	0.36520 (15)	0.0582 (5)

C8	0.0267 (3)	0.34724 (19)	0.29442 (15)	0.0576 (5)
H8	0.0106	0.2673	0.2953	0.069*
C9	-0.0170 (3)	0.42863 (18)	0.22240 (14)	0.0529 (5)
H9	-0.0651	0.4042	0.1749	0.063*
C10	0.0107 (2)	0.54753 (17)	0.22072 (14)	0.0508 (4)
C11	0.0755 (3)	0.5849 (2)	0.29330 (16)	0.0621 (5)
H11	0.0911	0.6649	0.2932	0.075*
C12	0.1170 (3)	0.5032 (2)	0.36626 (16)	0.0646 (6)
H12	0.1599	0.5281	0.4157	0.077*
C13	0.0662 (2)	0.62200 (17)	0.05341 (15)	0.0512 (4)
C14	0.2208 (3)	0.52456 (19)	0.04013 (15)	0.0557 (5)
H14A	0.2863	0.5293	0.0889	0.067*
H14B	0.1868	0.4454	0.0472	0.067*
C15	0.3997 (2)	0.63889 (18)	-0.06993 (14)	0.0522 (5)
C16	0.4732 (2)	0.6621 (2)	-0.16498 (15)	0.0564 (5)
C17	0.5604 (3)	0.7573 (2)	-0.19099 (17)	0.0650 (6)
H17	0.6083	0.7721	-0.2547	0.078*
C18	0.5757 (3)	0.8305 (2)	-0.1213 (2)	0.0724 (7)
C19	0.5045 (3)	0.8096 (2)	-0.0276 (2)	0.0744 (6)
H19	0.5156	0.8594	0.0189	0.089*
C20	0.4161 (3)	0.7144 (2)	-0.00201 (17)	0.0644 (6)
H20	0.3672	0.7011	0.0617	0.077*
C21	-0.1595 (3)	0.7360 (2)	0.1557 (2)	0.0687 (6)
H21	-0.2150	0.7427	0.1001	0.082*
C22	-0.2918 (4)	0.7281 (3)	0.2402 (3)	0.1095 (12)
H22A	-0.3292	0.6513	0.2438	0.164*
H22B	-0.3842	0.7919	0.2355	0.164*
H22C	-0.2486	0.7360	0.2969	0.164*
C23	-0.0768 (3)	0.84909 (18)	0.14879 (15)	0.0550 (5)
C24	-0.1183 (4)	1.0605 (2)	0.1175 (2)	0.0823 (8)
H24A	-0.0790	1.0730	0.1744	0.124*
H24B	-0.2063	1.1253	0.1070	0.124*
H24C	-0.0275	1.0584	0.0636	0.124*
C11	0.45524 (9)	0.57030 (7)	-0.25189 (5)	0.0857 (2)
C12	0.68469 (11)	0.95120 (8)	-0.15360 (8)	0.1130 (3)
N1	-0.0971 (3)	0.25673 (18)	0.49818 (12)	0.0639 (5)
N2	0.1583 (3)	0.14535 (19)	0.53514 (13)	0.0715 (5)
N3	-0.0284 (2)	0.62954 (15)	0.14216 (13)	0.0542 (4)
O1	-0.3482 (3)	0.2002 (2)	0.57170 (15)	0.1022 (7)
O2	0.1513 (4)	-0.0183 (2)	0.64236 (15)	0.1119 (8)
O3	0.1632 (2)	0.30374 (17)	0.43166 (13)	0.0761 (5)
O4	0.03226 (19)	0.69051 (14)	-0.01316 (11)	0.0641 (4)
O5	0.32022 (19)	0.54002 (13)	-0.05232 (10)	0.0600 (4)
O6	-0.1820 (2)	0.94630 (14)	0.12870 (13)	0.0702 (4)
O7	0.0634 (2)	0.84994 (14)	0.16004 (13)	0.0695 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0984 (19)	0.0862 (17)	0.0475 (12)	-0.0416 (15)	-0.0091 (12)	0.0017 (11)
C2	0.137 (3)	0.0818 (19)	0.0513 (13)	-0.0444 (18)	-0.0113 (15)	0.0159 (12)
C3	0.131 (3)	0.0653 (15)	0.0478 (12)	-0.0171 (16)	-0.0222 (14)	0.0056 (11)
C4	0.0814 (16)	0.0601 (13)	0.0479 (11)	-0.0198 (11)	-0.0157 (10)	0.0018 (9)
C5	0.084 (2)	0.147 (3)	0.109 (3)	-0.036 (2)	-0.0128 (18)	0.016 (2)
C6	0.172 (4)	0.110 (3)	0.093 (2)	0.044 (3)	-0.037 (3)	0.011 (2)
C7	0.0544 (11)	0.0615 (13)	0.0562 (11)	-0.0144 (9)	-0.0044 (9)	0.0091 (9)
C8	0.0619 (12)	0.0471 (11)	0.0625 (12)	-0.0166 (9)	-0.0049 (9)	0.0055 (9)
C9	0.0548 (11)	0.0507 (11)	0.0524 (10)	-0.0144 (9)	-0.0041 (8)	-0.0001 (8)
C10	0.0508 (10)	0.0454 (10)	0.0512 (10)	-0.0091 (8)	0.0024 (8)	0.0015 (8)
C11	0.0707 (14)	0.0491 (11)	0.0658 (13)	-0.0169 (10)	-0.0047 (10)	-0.0026 (10)
C12	0.0714 (14)	0.0676 (14)	0.0577 (12)	-0.0214 (11)	-0.0097 (10)	-0.0045 (10)
C13	0.0521 (10)	0.0417 (10)	0.0588 (11)	-0.0099 (8)	-0.0093 (8)	0.0053 (8)
C14	0.0563 (11)	0.0481 (11)	0.0544 (11)	-0.0052 (8)	0.0018 (9)	0.0098 (8)
C15	0.0463 (10)	0.0480 (10)	0.0555 (11)	-0.0004 (8)	-0.0037 (8)	0.0094 (8)
C16	0.0437 (10)	0.0631 (13)	0.0553 (11)	-0.0009 (9)	-0.0026 (8)	0.0052 (9)
C17	0.0448 (11)	0.0729 (15)	0.0679 (13)	-0.0074 (10)	0.0026 (9)	0.0153 (11)
C18	0.0541 (13)	0.0674 (15)	0.0905 (18)	-0.0168 (11)	0.0003 (11)	0.0081 (13)
C19	0.0710 (15)	0.0701 (15)	0.0822 (16)	-0.0164 (12)	-0.0097 (12)	-0.0049 (13)
C20	0.0667 (13)	0.0620 (13)	0.0588 (12)	-0.0073 (10)	-0.0039 (10)	0.0059 (10)
C21	0.0489 (11)	0.0513 (12)	0.0969 (17)	-0.0032 (9)	0.0011 (11)	0.0020 (11)
C22	0.0783 (18)	0.0708 (18)	0.159 (3)	-0.0151 (14)	0.0354 (19)	-0.0167 (19)
C23	0.0559 (12)	0.0469 (11)	0.0537 (11)	-0.0005 (9)	-0.0002 (9)	0.0059 (8)
C24	0.0913 (18)	0.0461 (12)	0.0980 (19)	-0.0044 (12)	-0.0036 (14)	0.0163 (12)
Cl1	0.0820 (4)	0.1117 (6)	0.0616 (4)	-0.0311 (4)	0.0092 (3)	-0.0140 (3)
Cl2	0.1001 (6)	0.0934 (6)	0.1411 (8)	-0.0505 (5)	0.0159 (5)	0.0014 (5)
N1	0.0801 (13)	0.0671 (12)	0.0487 (9)	-0.0268 (10)	-0.0111 (8)	0.0028 (8)
N2	0.1018 (16)	0.0628 (12)	0.0512 (10)	-0.0095 (11)	-0.0233 (10)	0.0038 (9)
N3	0.0520 (9)	0.0426 (9)	0.0618 (10)	-0.0045 (7)	-0.0012 (7)	0.0035 (7)
O1	0.0946 (15)	0.137 (2)	0.0777 (12)	-0.0521 (14)	-0.0042 (11)	0.0191 (12)
O2	0.174 (3)	0.0830 (14)	0.0698 (12)	-0.0018 (15)	-0.0302 (14)	0.0235 (10)
O3	0.0694 (10)	0.0823 (12)	0.0788 (11)	-0.0230 (9)	-0.0233 (8)	0.0288 (9)
O4	0.0676 (9)	0.0551 (8)	0.0673 (9)	-0.0059 (7)	-0.0180 (7)	0.0156 (7)
O5	0.0649 (9)	0.0527 (8)	0.0548 (8)	-0.0097 (7)	0.0049 (6)	0.0042 (6)
O6	0.0625 (9)	0.0498 (9)	0.0895 (11)	0.0002 (7)	-0.0083 (8)	0.0141 (8)
O7	0.0671 (10)	0.0522 (9)	0.0904 (12)	-0.0080 (7)	-0.0223 (8)	0.0041 (8)

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

C1—O1	1.334 (3)	C13—N3	1.350 (3)
C1—N1	1.336 (3)	C13—C14	1.522 (3)
C1—C2	1.382 (4)	C14—O5	1.423 (2)
C2—C3	1.369 (4)	C14—H14A	0.9700
C2—H2	0.9300	C14—H14B	0.9700
C3—N2	1.326 (4)	C15—O5	1.367 (2)

C3—O2	1.342 (3)	C15—C20	1.380 (3)
C4—N1	1.315 (3)	C15—C16	1.394 (3)
C4—N2	1.317 (3)	C16—C17	1.376 (3)
C4—O3	1.363 (3)	C16—C11	1.721 (2)
C5—O1	1.433 (4)	C17—C18	1.379 (4)
C5—H5A	0.9600	C17—H17	0.9300
C5—H5B	0.9600	C18—C19	1.368 (4)
C5—H5C	0.9600	C18—C12	1.737 (2)
C6—O2	1.415 (5)	C19—C20	1.381 (3)
C6—H6A	0.9600	C19—H19	0.9300
C6—H6B	0.9600	C20—H20	0.9300
C6—H6C	0.9600	C21—N3	1.465 (3)
C7—C12	1.368 (3)	C21—C22	1.466 (4)
C7—C8	1.377 (3)	C21—C23	1.531 (3)
C7—O3	1.395 (3)	C21—H21	0.9800
C8—C9	1.376 (3)	C22—H22A	0.9600
C8—H8	0.9300	C22—H22B	0.9600
C9—C10	1.392 (3)	C22—H22C	0.9600
C9—H9	0.9300	C23—O7	1.199 (3)
C10—C11	1.378 (3)	C23—O6	1.321 (2)
C10—N3	1.439 (3)	C24—O6	1.454 (3)
C11—C12	1.383 (3)	C24—H24A	0.9600
C11—H11	0.9300	C24—H24B	0.9600
C12—H12	0.9300	C24—H24C	0.9600
C13—O4	1.216 (2)		
O1—C1—N1	118.8 (2)	H14A—C14—H14B	108.1
O1—C1—C2	118.8 (2)	O5—C15—C20	125.74 (18)
N1—C1—C2	122.4 (3)	O5—C15—C16	116.07 (19)
C3—C2—C1	116.1 (2)	C20—C15—C16	118.2 (2)
C3—C2—H2	121.9	C17—C16—C15	121.3 (2)
C1—C2—H2	121.9	C17—C16—C11	119.07 (17)
N2—C3—O2	118.4 (3)	C15—C16—C11	119.64 (17)
N2—C3—C2	123.6 (2)	C16—C17—C18	119.1 (2)
O2—C3—C2	118.0 (3)	C16—C17—H17	120.5
N1—C4—N2	130.0 (2)	C18—C17—H17	120.5
N1—C4—O3	118.5 (2)	C19—C18—C17	120.7 (2)
N2—C4—O3	111.5 (2)	C19—C18—C12	119.9 (2)
O1—C5—H5A	109.5	C17—C18—C12	119.39 (19)
O1—C5—H5B	109.5	C18—C19—C20	120.0 (3)
H5A—C5—H5B	109.5	C18—C19—H19	120.0
O1—C5—H5C	109.5	C20—C19—H19	120.0
H5A—C5—H5C	109.5	C15—C20—C19	120.8 (2)
H5B—C5—H5C	109.5	C15—C20—H20	119.6
O2—C6—H6A	109.5	C19—C20—H20	119.6
O2—C6—H6B	109.5	N3—C21—C22	115.3 (2)
H6A—C6—H6B	109.5	N3—C21—C23	108.87 (17)
O2—C6—H6C	109.5	C22—C21—C23	114.0 (2)

H6A—C6—H6C	109.5	N3—C21—H21	105.9
H6B—C6—H6C	109.5	C22—C21—H21	105.9
C12—C7—C8	121.4 (2)	C23—C21—H21	105.9
C12—C7—O3	116.8 (2)	C21—C22—H22A	109.5
C8—C7—O3	121.4 (2)	C21—C22—H22B	109.5
C9—C8—C7	119.2 (2)	H22A—C22—H22B	109.5
C9—C8—H8	120.4	C21—C22—H22C	109.5
C7—C8—H8	120.4	H22A—C22—H22C	109.5
C8—C9—C10	119.9 (2)	H22B—C22—H22C	109.5
C8—C9—H9	120.0	O7—C23—O6	124.5 (2)
C10—C9—H9	120.0	O7—C23—C21	125.12 (18)
C11—C10—C9	119.99 (19)	O6—C23—C21	110.36 (19)
C11—C10—N3	121.04 (18)	O6—C24—H24A	109.5
C9—C10—N3	118.97 (18)	O6—C24—H24B	109.5
C10—C11—C12	119.8 (2)	H24A—C24—H24B	109.5
C10—C11—H11	120.1	O6—C24—H24C	109.5
C12—C11—H11	120.1	H24A—C24—H24C	109.5
C7—C12—C11	119.6 (2)	H24B—C24—H24C	109.5
C7—C12—H12	120.2	C4—N1—C1	114.2 (2)
C11—C12—H12	120.2	C4—N2—C3	113.7 (2)
O4—C13—N3	122.16 (19)	C13—N3—C10	122.15 (16)
O4—C13—C14	120.86 (18)	C13—N3—C21	115.18 (17)
N3—C13—C14	116.98 (17)	C10—N3—C21	122.20 (18)
O5—C14—C13	110.13 (16)	C1—O1—C5	118.3 (2)
O5—C14—H14A	109.6	C3—O2—C6	118.4 (3)
C13—C14—H14A	109.6	C4—O3—C7	120.35 (18)
O5—C14—H14B	109.6	C15—O5—C14	117.63 (16)
C13—C14—H14B	109.6	C23—O6—C24	116.33 (18)
O1—C1—C2—C3	-179.1 (2)	N2—C4—N1—C1	-1.4 (4)
N1—C1—C2—C3	0.8 (4)	O3—C4—N1—C1	-178.7 (2)
C1—C2—C3—N2	-0.8 (4)	O1—C1—N1—C4	-179.9 (2)
C1—C2—C3—O2	-179.8 (2)	C2—C1—N1—C4	0.1 (3)
C12—C7—C8—C9	1.1 (3)	N1—C4—N2—C3	1.4 (4)
O3—C7—C8—C9	-170.65 (19)	O3—C4—N2—C3	178.9 (2)
C7—C8—C9—C10	1.3 (3)	O2—C3—N2—C4	178.8 (2)
C8—C9—C10—C11	-2.7 (3)	C2—C3—N2—C4	-0.2 (4)
C8—C9—C10—N3	176.97 (18)	O4—C13—N3—C10	179.28 (19)
C9—C10—C11—C12	1.8 (3)	C14—C13—N3—C10	-1.8 (3)
N3—C10—C11—C12	-177.89 (19)	O4—C13—N3—C21	-8.4 (3)
C8—C7—C12—C11	-2.0 (3)	C14—C13—N3—C21	170.50 (19)
O3—C7—C12—C11	170.1 (2)	C11—C10—N3—C13	107.0 (2)
C10—C11—C12—C7	0.6 (3)	C9—C10—N3—C13	-72.7 (3)
O4—C13—C14—O5	7.9 (3)	C11—C10—N3—C21	-64.8 (3)
N3—C13—C14—O5	-171.06 (18)	C9—C10—N3—C21	115.5 (2)
O5—C15—C16—C17	-178.33 (17)	C22—C21—N3—C13	162.8 (3)
C20—C15—C16—C17	0.1 (3)	C23—C21—N3—C13	-67.6 (3)
O5—C15—C16—C11	1.7 (2)	C22—C21—N3—C10	-24.9 (3)

C20—C15—C16—C11	-179.85 (16)	C23—C21—N3—C10	104.8 (2)
C15—C16—C17—C18	0.4 (3)	N1—C1—O1—C5	-1.0 (4)
C11—C16—C17—C18	-179.63 (18)	C2—C1—O1—C5	178.9 (3)
C16—C17—C18—C19	-0.4 (4)	N2—C3—O2—C6	1.2 (4)
C16—C17—C18—C12	-179.73 (16)	C2—C3—O2—C6	-179.7 (3)
C17—C18—C19—C20	0.0 (4)	N1—C4—O3—C7	-14.1 (3)
C12—C18—C19—C20	179.26 (19)	N2—C4—O3—C7	168.1 (2)
O5—C15—C20—C19	177.7 (2)	C12—C7—O3—C4	127.6 (2)
C16—C15—C20—C19	-0.6 (3)	C8—C7—O3—C4	-60.3 (3)
C18—C19—C20—C15	0.6 (4)	C20—C15—O5—C14	11.7 (3)
N3—C21—C23—O7	-20.9 (3)	C16—C15—O5—C14	-169.97 (17)
C22—C21—C23—O7	109.4 (3)	C13—C14—O5—C15	68.7 (2)
N3—C21—C23—O6	159.17 (19)	O7—C23—O6—C24	1.6 (3)
C22—C21—C23—O6	-70.5 (3)	C21—C23—O6—C24	-178.4 (2)

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>
C9—H9...O4 ⁱ	0.93	2.57	3.402 (3)	150

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