

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Methyl 2,4-dihydroxy-5-(4-nitrobenzamido)benzoate

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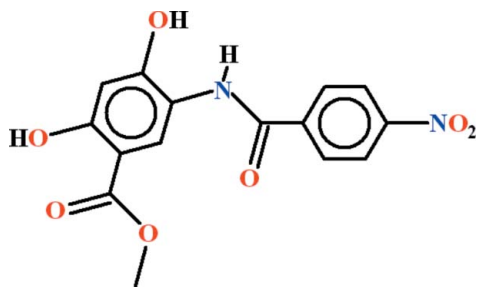
Received 3 January 2013; accepted 3 January 2013

 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.052; wR factor = 0.134; data-to-parameter ratio = 13.1.

In the title compound, $\text{C}_{15}\text{H}_{12}\text{N}_2\text{O}_7$, the dihedral angle between the aromatic rings is 4.58 (13) $^\circ$ and the nitro group is rotated from its attached ring by 18.07 (17) $^\circ$. Intramolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds generate $S(5)$ and $S(6)$ rings, respectively. In the crystal, molecules are linked by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds, generating $[001]$ $C(7)$ chains. The chains are linked by $\text{C}-\text{H}\cdots\text{O}$ interactions, forming a three-dimensional network, which incorporates $R_2^2(7)$ and $R_2^2(10)$ loops.

Related literature

For a related structure, see: Gorelik *et al.* (2010). For graph-set notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

 $\text{C}_{15}\text{H}_{12}\text{N}_2\text{O}_7$
 $M_r = 332.27$

 Monoclinic, $C2/c$
 $a = 30.412$ (6) Å

 $b = 6.9325$ (15) Å
 $c = 14.936$ (3) Å
 $\beta = 111.737$ (8) $^\circ$
 $V = 2925.0$ (11) Å³
 $Z = 8$

 Mo $K\alpha$ radiation
 $\mu = 0.12$ mm⁻¹
 $T = 296$ K
 $0.28 \times 0.18 \times 0.16$ mm

Data collection

 Bruker Kappa APEXII CCD diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.970$, $T_{\max} = 0.980$

 10681 measured reflections
 2885 independent reflections
 1519 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.055$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.134$
 $S = 0.98$
 2885 reflections

 220 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.24$ e Å⁻³

Table 1

 Hydrogen-bond geometry (Å, $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O4}$	0.86	2.16	2.595 (3)	111
$\text{O3}-\text{H3A}\cdots\text{O2}$	0.82	1.91	2.619 (3)	144
$\text{O4}-\text{H4}\cdots\text{O5}^i$	0.82	1.86	2.670 (2)	170
$\text{C8}-\text{H8A}\cdots\text{O3}^{ii}$	0.96	2.51	3.274 (4)	137
$\text{C12}-\text{H12}\cdots\text{O7}^{iii}$	0.93	2.38	3.297 (4)	171
$\text{C15}-\text{H15}\cdots\text{O4}^{ii}$	0.93	2.46	3.370 (3)	165

 Symmetry codes: (i) $x, -y, z + \frac{1}{2}$; (ii) $x, -y, z - \frac{1}{2}$; (iii) $-x + 1, y, -z + \frac{3}{2}$.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 2012) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 2012) and *PLATON*.

The authors acknowledge the provision of funds for the purchase of the diffractometer and encouragement by Dr Muhammad Akram Chaudhary, Vice Chancellor, University of Sargodha, Pakistan.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB7022).

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supporting information

Acta Cryst. (2013). E69, o207 [doi:10.1107/S160053681300024X]

Methyl 2,4-dihydroxy-5-(4-nitrobenzamido)benzoate

Syeda Sohaila Naz, Nazar Ul Islam, M. Nawaz Tahir and Muhammad Raza Shah

S1. Comment

The title compound, (I) (Fig. 1), has been prepared for derivatization and for biological studies. The crystal structure of 4-((4-nitrobenzoyl)amino)benzoic acid (Gorelik *et al.*, 2010) has been published which is related to the title compound.

In (I), the groups A (C1—C8/O1—O4/N1) of methyl 5-amino-2,4-dihydroxybenzoate and B (C9—C15/O5—O7/N2) of 4-nitrobenzoyl are planar with r. m. s. deviation of 0.0252 and 0.0363 Å, respectively. The dihedral angle between A/B is 3.59 (3)°. There exist strong intramolecular H-bondings of N—H···O and O—H···O types (Table 1, Fig. 2) completing S(5) and S(6) ring motifs (Bernstein *et al.*, 1995). There also exist strong intermolecular H-bondings of C—H···O and O—H···O types due to which $R_2^2(7)$ and $R_2^2(10)$ loops are formed (Table 1, Fig. 2) resulting in the formation of three dimensional polymeric network.

S2. Experimental

Equivalent amounts of methyl 5-amino-2,4-dihydroxybenzoate (0.20 g, 1.1 mmol) and 4-nitrobenzoyl chloride (0.20 g, 1.1 mmol) were heated at 333 K for 3 h in dimethylformamide (DMF). The reaction mixture was freeze dried, neutralized with aq. NaHCO₃ (5%) and extracted with dichloromethane (DCM), dried over Na₂SO₄ and evaporated *in vacuo* to give pure product which was recrystallized from methanol and water solution to afford yellow needles.

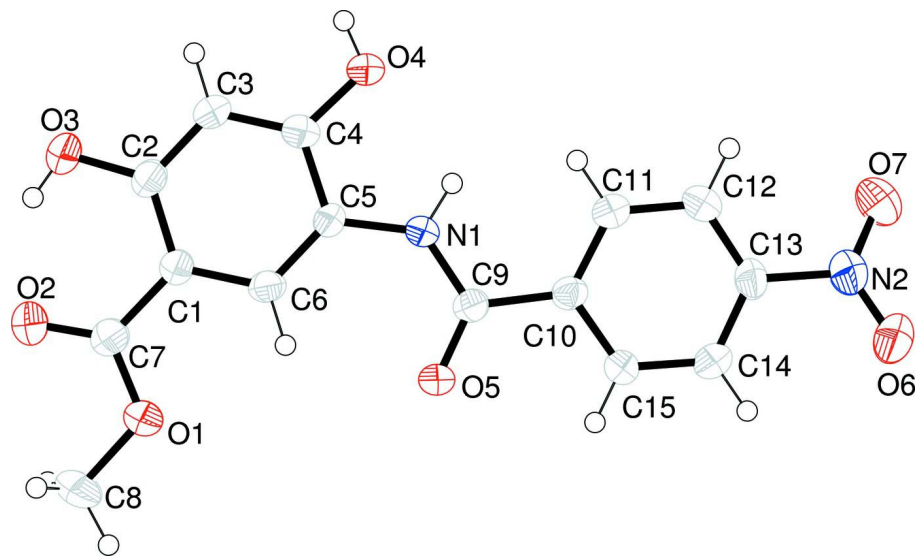
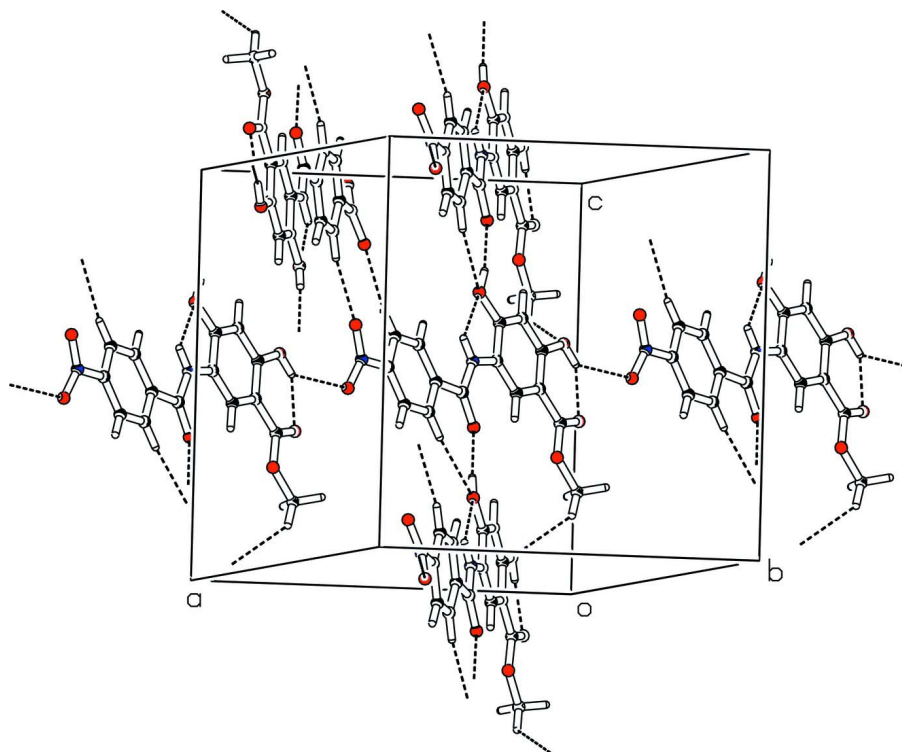


Figure 1

View of the title compound with displacement ellipsoids drawn at the 50% probability level.

**Figure 2**

The partial packing, which shows that molecules form various ring motifs to form three dimensional polymeric network.

Methyl 2,4-dihydroxy-5-(4-nitrobenzamido)benzoate

Crystal data

$C_{15}H_{12}N_2O_7$

$M_r = 332.27$

Monoclinic, $C2/c$

Hall symbol: $-C 2yc$

$a = 30.412 (6) \text{ \AA}$

$b = 6.9325 (15) \text{ \AA}$

$c = 14.936 (3) \text{ \AA}$

$\beta = 111.737 (8)^\circ$

$V = 2925.0 (11) \text{ \AA}^3$

$Z = 8$

$F(000) = 1376$

$D_x = 1.509 \text{ Mg m}^{-3}$

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

Cell parameters from 1519 reflections

$\theta = 1.4\text{--}26.0^\circ$

$\mu = 0.12 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Needle, yellow

$0.28 \times 0.18 \times 0.16 \text{ mm}$

Data collection

Bruker Kappa APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: $8.00 \text{ pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)

$T_{\min} = 0.970$, $T_{\max} = 0.980$

10681 measured reflections

2885 independent reflections

1519 reflections with $I > 2\sigma(I)$

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

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 1.4^\circ$

$h = -35 \rightarrow 37$

$k = -8 \rightarrow 5$

$l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.134$
 $S = 0.98$
 2885 reflections
 220 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
 $w = 1/[\sigma^2(F_o^2) + (0.0575P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.24 \text{ e } \text{Å}^{-3}$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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 (Å^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.10274 (6)	0.1155 (3)	0.32019 (12)	0.0472 (7)
O2	0.05741 (6)	0.1342 (3)	0.40795 (13)	0.0569 (8)
O3	0.09767 (6)	0.1238 (3)	0.59618 (13)	0.0556 (8)
O4	0.26255 (6)	0.0020 (3)	0.72631 (12)	0.0551 (8)
O5	0.25654 (6)	-0.0216 (3)	0.39997 (11)	0.0441 (7)
O6	0.49252 (7)	-0.2190 (4)	0.50312 (16)	0.0817 (10)
O7	0.50869 (7)	-0.1537 (4)	0.65151 (16)	0.0963 (12)
N1	0.26968 (7)	-0.0090 (3)	0.55853 (14)	0.0378 (8)
N2	0.48095 (8)	-0.1732 (4)	0.56944 (19)	0.0571 (10)
C1	0.13985 (9)	0.0859 (4)	0.48802 (17)	0.0319 (9)
C2	0.13784 (9)	0.0904 (4)	0.58047 (19)	0.0364 (9)
C3	0.17868 (9)	0.0616 (4)	0.66047 (18)	0.0415 (10)
C4	0.22102 (9)	0.0301 (4)	0.65078 (17)	0.0371 (9)
C5	0.22394 (8)	0.0249 (3)	0.55857 (17)	0.0303 (9)
C6	0.18331 (8)	0.0531 (3)	0.47889 (16)	0.0314 (9)
C7	0.09636 (10)	0.1146 (4)	0.40347 (18)	0.0390 (10)
C8	0.06012 (10)	0.1408 (5)	0.23493 (19)	0.0624 (13)
C9	0.28430 (8)	-0.0313 (3)	0.48462 (17)	0.0298 (9)
C10	0.33614 (8)	-0.0680 (3)	0.51086 (16)	0.0301 (8)
C11	0.36749 (9)	-0.1024 (4)	0.60418 (17)	0.0425 (10)
C12	0.41477 (9)	-0.1343 (4)	0.62442 (19)	0.0448 (10)
C13	0.43065 (9)	-0.1345 (4)	0.54982 (19)	0.0382 (9)
C14	0.40103 (9)	-0.1016 (4)	0.45639 (19)	0.0482 (10)
C15	0.35354 (9)	-0.0687 (4)	0.43751 (18)	0.0425 (10)
H1	0.29169	-0.01664	0.61475	0.0453*
H3	0.17730	0.06358	0.72164	0.0499*

H3A	0.07616	0.14790	0.54484	0.0833*
H4	0.25843	0.01839	0.77701	0.0826*
H6	0.18481	0.05032	0.41783	0.0377*
H8A	0.06788	0.13162	0.17835	0.0937*
H8B	0.04667	0.26521	0.23692	0.0937*
H8C	0.03765	0.04218	0.23315	0.0937*
H11	0.35623	-0.10400	0.65419	0.0511*
H12	0.43554	-0.15519	0.68741	0.0538*
H14	0.41260	-0.10127	0.40680	0.0578*
H15	0.33301	-0.04676	0.37444	0.0509*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0365 (11)	0.0729 (14)	0.0296 (10)	0.0091 (10)	0.0093 (9)	0.0050 (10)
O2	0.0332 (12)	0.0922 (17)	0.0453 (12)	0.0137 (11)	0.0146 (10)	0.0050 (11)
O3	0.0351 (12)	0.0936 (17)	0.0440 (12)	0.0141 (12)	0.0216 (10)	0.0101 (12)
O4	0.0339 (11)	0.1073 (18)	0.0236 (9)	0.0067 (11)	0.0101 (9)	0.0008 (11)
O5	0.0321 (10)	0.0767 (15)	0.0233 (9)	0.0060 (10)	0.0099 (8)	0.0008 (9)
O6	0.0442 (14)	0.144 (2)	0.0601 (15)	0.0172 (14)	0.0232 (12)	-0.0166 (15)
O7	0.0364 (13)	0.189 (3)	0.0497 (14)	0.0125 (16)	-0.0002 (11)	-0.0097 (16)
N1	0.0263 (12)	0.0631 (17)	0.0222 (10)	0.0026 (11)	0.0069 (9)	0.0004 (11)
N2	0.0335 (15)	0.087 (2)	0.0471 (16)	0.0061 (14)	0.0106 (13)	-0.0062 (15)
C1	0.0295 (15)	0.0362 (17)	0.0312 (14)	0.0024 (12)	0.0128 (12)	0.0022 (12)
C2	0.0346 (16)	0.0428 (18)	0.0368 (15)	0.0043 (14)	0.0189 (13)	0.0039 (13)
C3	0.0367 (17)	0.063 (2)	0.0287 (14)	0.0020 (15)	0.0165 (13)	0.0005 (14)
C4	0.0345 (16)	0.0483 (19)	0.0266 (13)	-0.0013 (14)	0.0090 (12)	0.0004 (13)
C5	0.0266 (15)	0.0385 (17)	0.0284 (13)	-0.0013 (12)	0.0131 (11)	-0.0010 (12)
C6	0.0345 (16)	0.0361 (16)	0.0246 (13)	-0.0033 (12)	0.0122 (12)	-0.0014 (12)
C7	0.0383 (18)	0.0457 (18)	0.0350 (15)	0.0051 (14)	0.0159 (13)	0.0029 (13)
C8	0.0466 (19)	0.098 (3)	0.0312 (16)	0.0178 (19)	0.0013 (14)	0.0066 (17)
C9	0.0317 (15)	0.0314 (16)	0.0276 (14)	-0.0019 (12)	0.0125 (12)	-0.0010 (12)
C10	0.0323 (15)	0.0312 (16)	0.0273 (13)	-0.0008 (12)	0.0116 (11)	-0.0039 (12)
C11	0.0372 (17)	0.063 (2)	0.0296 (14)	0.0045 (15)	0.0149 (13)	0.0008 (14)
C12	0.0348 (17)	0.065 (2)	0.0295 (15)	0.0044 (15)	0.0059 (12)	-0.0001 (14)
C13	0.0244 (15)	0.0484 (18)	0.0405 (16)	0.0013 (14)	0.0104 (12)	-0.0033 (14)
C14	0.0394 (18)	0.077 (2)	0.0328 (15)	0.0059 (16)	0.0189 (13)	0.0019 (15)
C15	0.0301 (16)	0.068 (2)	0.0287 (14)	0.0075 (15)	0.0101 (12)	0.0035 (14)

Geometric parameters (Å, °)

O1—C7	1.328 (3)	C4—C5	1.413 (3)
O1—C8	1.453 (3)	C5—C6	1.376 (3)
O2—C7	1.219 (4)	C9—C10	1.499 (4)
O3—C2	1.347 (4)	C10—C15	1.382 (4)
O4—C4	1.360 (3)	C10—C11	1.387 (3)
O5—C9	1.235 (3)	C11—C12	1.373 (4)
O6—N2	1.210 (4)	C12—C13	1.369 (4)

O7—N2	1.211 (3)	C13—C14	1.371 (4)
O3—H3A	0.8200	C14—C15	1.384 (4)
O4—H4	0.8200	C3—H3	0.9300
N1—C9	1.343 (3)	C6—H6	0.9300
N1—C5	1.411 (3)	C8—H8A	0.9600
N2—C13	1.472 (4)	C8—H8B	0.9600
N1—H1	0.8600	C8—H8C	0.9600
C1—C7	1.465 (4)	C11—H11	0.9300
C1—C6	1.397 (4)	C12—H12	0.9300
C1—C2	1.405 (4)	C14—H14	0.9300
C2—C3	1.382 (4)	C15—H15	0.9300
C3—C4	1.366 (4)		
C7—O1—C8	115.4 (2)	C9—C10—C15	117.9 (2)
C2—O3—H3A	109.00	C11—C10—C15	118.3 (2)
C4—O4—H4	109.00	C9—C10—C11	123.8 (2)
C5—N1—C9	130.3 (2)	C10—C11—C12	121.5 (2)
O6—N2—C13	118.7 (2)	C11—C12—C13	118.5 (2)
O6—N2—O7	123.6 (3)	N2—C13—C12	119.5 (2)
O7—N2—C13	117.7 (2)	C12—C13—C14	122.1 (3)
C9—N1—H1	115.00	N2—C13—C14	118.4 (2)
C5—N1—H1	115.00	C13—C14—C15	118.5 (3)
C6—C1—C7	121.5 (2)	C10—C15—C14	121.1 (2)
C2—C1—C7	119.4 (3)	C2—C3—H3	120.00
C2—C1—C6	119.1 (2)	C4—C3—H3	120.00
O3—C2—C3	117.2 (2)	C1—C6—H6	119.00
C1—C2—C3	119.6 (3)	C5—C6—H6	119.00
O3—C2—C1	123.2 (2)	O1—C8—H8A	109.00
C2—C3—C4	120.8 (2)	O1—C8—H8B	109.00
O4—C4—C3	123.9 (2)	O1—C8—H8C	109.00
O4—C4—C5	115.5 (2)	H8A—C8—H8B	109.00
C3—C4—C5	120.6 (2)	H8A—C8—H8C	109.00
N1—C5—C4	115.0 (2)	H8B—C8—H8C	109.00
C4—C5—C6	118.6 (2)	C10—C11—H11	119.00
N1—C5—C6	126.4 (2)	C12—C11—H11	119.00
C1—C6—C5	121.2 (2)	C11—C12—H12	121.00
O1—C7—O2	122.2 (2)	C13—C12—H12	121.00
O1—C7—C1	114.2 (3)	C13—C14—H14	121.00
O2—C7—C1	123.6 (2)	C15—C14—H14	121.00
N1—C9—C10	116.2 (2)	C10—C15—H15	119.00
O5—C9—N1	121.8 (2)	C14—C15—H15	119.00
O5—C9—C10	122.1 (2)		
C8—O1—C7—O2	-0.2 (4)	C2—C3—C4—O4	179.6 (3)
C8—O1—C7—C1	179.1 (2)	C2—C3—C4—C5	-0.5 (4)
C9—N1—C5—C4	176.6 (2)	O4—C4—C5—N1	0.1 (3)
C9—N1—C5—C6	-3.7 (4)	O4—C4—C5—C6	-179.7 (2)
C5—N1—C9—O5	0.8 (4)	C3—C4—C5—N1	-179.9 (2)

C5—N1—C9—C10	-179.3 (2)	C3—C4—C5—C6	0.3 (4)
O6—N2—C13—C12	162.4 (3)	N1—C5—C6—C1	-179.9 (2)
O6—N2—C13—C14	-17.0 (4)	C4—C5—C6—C1	-0.1 (3)
O7—N2—C13—C12	-19.0 (4)	O5—C9—C10—C11	-172.1 (2)
O7—N2—C13—C14	161.7 (3)	O5—C9—C10—C15	7.5 (3)
C6—C1—C2—O3	179.0 (2)	N1—C9—C10—C11	8.0 (3)
C6—C1—C2—C3	-0.3 (4)	N1—C9—C10—C15	-172.4 (2)
C7—C1—C2—O3	-1.3 (4)	C9—C10—C11—C12	-179.7 (2)
C7—C1—C2—C3	179.4 (3)	C15—C10—C11—C12	0.8 (4)
C2—C1—C6—C5	0.1 (4)	C9—C10—C15—C14	-180.0 (2)
C7—C1—C6—C5	-179.5 (2)	C11—C10—C15—C14	-0.4 (4)
C2—C1—C7—O1	177.3 (2)	C10—C11—C12—C13	-1.0 (4)
C2—C1—C7—O2	-3.4 (4)	C11—C12—C13—N2	-178.5 (3)
C6—C1—C7—O1	-3.0 (4)	C11—C12—C13—C14	0.8 (4)
C6—C1—C7—O2	176.3 (3)	N2—C13—C14—C15	178.8 (3)
O3—C2—C3—C4	-178.9 (3)	C12—C13—C14—C15	-0.5 (4)
C1—C2—C3—C4	0.4 (4)	C13—C14—C15—C10	0.3 (4)

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>
N1—H1...O4	0.86	2.16	2.595 (3)	111
O3—H3 <i>A</i> ...O2	0.82	1.91	2.619 (3)	144
O4—H4...O5 ⁱ	0.82	1.86	2.670 (2)	170
C8—H8 <i>A</i> ...O3 ⁱⁱ	0.96	2.51	3.274 (4)	137
C12—H12...O7 ⁱⁱⁱ	0.93	2.38	3.297 (4)	171
C15—H15...O4 ⁱⁱ	0.93	2.46	3.370 (3)	165

Symmetry codes: (i) *x*, -*y*, *z*+1/2; (ii) *x*, -*y*, *z*-1/2; (iii) -*x*+1, *y*, -*z*+3/2.