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Ethyl 3,6-dihydroxy-6-methyl-4-phenyl-4,5,6,7-tetrahydro-1*H*-indazole-5-carboxylate monohydrate

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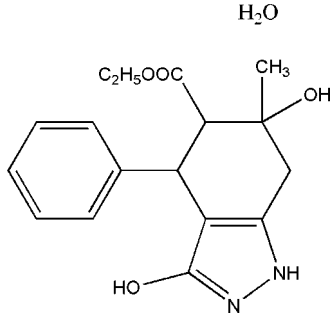
Key indicators: single-crystal X-ray study; T = 296 K; mean σ(C–C) = 0.005 Å; R factor = 0.077; wR factor = 0.198; data-to-parameter ratio = 12.5.

In the title compound, $C_{17}H_{20}N_2O_4\cdot H_2O$, the cyclohexene ring adopts a half-chair conformation while the indazole ring is essentially planar [maximum deviation = 0.0192 (12) Å]. In the crystal, pairs of intermolecular O—H···N hydrogen bonds link the molecules into dimers lying about inversion centers and intramolecular O—H···O hydrogen bonds result in six-membered rings. The dimers are further connected by N—H···O and O—H···O hydrogen bonds.

Related literature

For general background to azoles, see: Genin *et al.* (2000). For a related structure, see: Hema *et al.* (2006).



Experimental

Crystal data

$C_{17}H_{20}N_2O_4\cdot H_2O$
M_r = 334.37
Triclinic, $P\overline{1}$
a = 6.9964 (15) Å
b = 8.8647 (19) Å
c = 15.124 (4) Å
α = 99.363 (6)°
β = 95.281 (6)°
γ = 112.271 (4)°
V = 844.2 (3) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.10 mm^{-1}
T = 296 K
0.20 × 0.20 × 0.20 mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2003)
T_{min} = 0.981, T_{max} = 0.981
6332 measured reflections
2889 independent reflections
2327 reflections with $I > 2\sigma(I)$
R_{int} = 0.065

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.077
$wR(F^2)$ = 0.198
S = 1.00
2889 reflections
231 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.27 e Å^{-3}
$\Delta\rho_{min}$ = −0.37 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1*B*···N2	0.82	1.89	2.705 (2)	171
O4—H4*A*···O3	0.82	2.22	2.897 (2)	141
N1—H1*A*···O4	0.93 (3)	1.94	2.778 (3)	155
O5—H5*B*···O1	0.95 (5)	2.01	2.874 (2)	165
O5—H5*C*···O2	0.84 (4)	1.97	2.844 (3)	179

Data collection: *APEX2* (Bruker, 2005); 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: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

We thank Professor Victor N. Khrustalev for fruitful discussions and help in this work.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2374).

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supplementary materials

Acta Cryst. (2011). E**67**, o480 [**doi:10.1107/S160053681100242X**]

Ethyl 3,6-dihydroxy-6-methyl-4-phenyl-4,5,6,7-tetrahydro-1*H*-indazole-5-carboxylate monohydrate

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Comment

In the field of heterocyclic compounds, azoles are important due to their wide range of applications (Genin *et al.*, 2000). In the microbial evaluation of organic compounds for the development of current research in drug discovery and medicinal chemistry we have prepared the title compound and determined its crystal structure which has been presented in this article.

In the title compound (Fig. 1), the cyclohexene ring adopts a half-chair conformation, C6 lies 0.685 (3) Å out of the plane formed by the rest of the ring atoms. The indazole ring (N1/N2/C3/C3A/C7A) is essentially planar with maximum deviation from the ring plane being 0.0192 (12) Å for C7A. In the crystal structure, intermolecular hydrogen bonds O1—H1B···N2 result in centrosymmetric dimers lying about inversion centers. Intramolecular hydrogen bonds O4—H4A···O3 result in six-membered rings. The dimers are further packed and stabilized by N—H···O and O—H···O hydrogen bonds (Table 1 and Fig. 2).

The molecular dimensions in the title compound are in close agreement with the corresponding molecular dimensions of a closely related compoud (Hema *et al.*, 2006).

Experimental

(rac)-Diethyl-4-hydroxy-4-methyl-6-oxo-2-phenyl-1,3-dicarboxylate (20 mmol) and hydroxylamine hydrochloride (20 mmol) were dissolved in 20 ml e thanol. The mixture was stirred at 345–350 K for 10 h. After cooling to a room temperature colorless crystals were obtained which were filtered and washed with ethanol. The crystals were dissolved in ethanol (50 ml) and recrystallized to yield colourless block-shaped crystals of the title compound suitable for X-ray crystallographic analysis..

Refinement

The hydrogen atoms of the water of hydration and amino group were localized from difference-Fourier maps and included in the refinement with isotropic displacement parameters. The rest of the hydrogen atoms were placed in calculated positions with and refined in the riding model at O—H = 0.82 and C—H = 0.93-0.98 Å with isotropic displacement parameters U_{iso}(H) = 1.2 or 1.5U_{eq}(parent atoms).

Figures

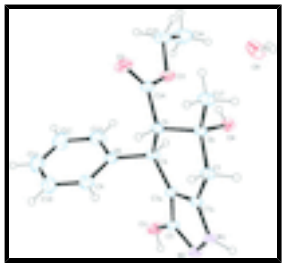

Fig. 1. The molecular structure of the title compound, with the atomic numbering scheme. Displacement ellipsoids were drawn at the 30% probability level.

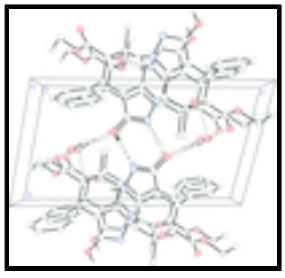

Fig. 2. The hydrogen-bonded (dashed lines) packing in the title compound; H-atoms not involved in hydrogen bonding have been excluded for clarity.

Ethyl 3,6-dihydroxy-6-methyl-4-phenyl-4,5,6,7-tetrahydro-1*H*-indazole-5- carboxylate monohydrate

Crystal data

$C_{17}H_{20}N_2O_4 \cdot H_2O$	$Z = 2$
$M_r = 334.37$	$F(000) = 356$
Triclinic, $P\overline{1}$	$D_x = 1.315$ Mg m^{-3}
Hall symbol: -P 1	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
$a = 6.9964$ (15) Å	Cell parameters from 909 reflections
$b = 8.8647$ (19) Å	$\theta = 2.5$–$30.6°$
$c = 15.124$ (4) Å	$\mu = 0.10$ mm^{-1}
$\alpha = 99.363$ (6)°	$T = 296$ K
$\beta = 95.281$ (6)°	Prism, colourless
$\gamma = 112.271$ (4)°	0.20 × 0.20 × 0.20 mm
$V = 844.2$ (3) Å^3	

Data collection

Bruker APEXII CCD diffractometer	2889 independent reflections
Radiation source: fine-focus sealed tube	2327 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.065$
φ and ω scans	$\theta_{max} = 25.0°$, $\theta_{min} = 2.5°$
Absorption correction: multi-scan (*SADABS*; Bruker, 2003)	$h = -8 \rightarrow 8$
$T_{min} = 0.981$, $T_{max} = 0.981$	$k = -10 \rightarrow 10$
6332 measured reflections	$l = -17 \rightarrow 16$

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.077$	Hydrogen site location: difference Fourier map
$wR(F^2) = 0.198$	H atoms treated by a mixture of independent and constrained refinement
$S = 1.00$	$w = 1/[\sigma^2(F_o^2) + (0.0611P)^2 + 0.4352P]$ where $P = (F_o^2 + 2F_c^2)/3$
2889 reflections	$(\Delta/\sigma)_{max} < 0.001$
231 parameters	$\Delta\rho_{max} = 0.27$ e Å^{-3}
0 restraints	$\Delta\rho_{min} = -0.37$ e Å^{-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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
O1	0.9685 (3)	1.3998 (2)	0.37394 (14)	0.0463 (5)
H1B	0.9998	1.4897	0.4089	0.069*
O2	0.7109 (3)	0.6179 (2)	0.15151 (15)	0.0544 (6)
O3	0.4935 (3)	0.7477 (2)	0.14842 (13)	0.0421 (5)
O4	0.3968 (2)	0.8009 (2)	0.33029 (13)	0.0363 (5)
H4A	0.3737	0.7989	0.2759	0.054*
O5	0.0528 (4)	0.5446 (3)	0.2177 (2)	0.0637 (7)
H5B	0.009 (6)	0.485 (6)	0.264 (3)	0.081 (13)*
H5C	−0.051 (6)	0.563 (5)	0.198 (3)	0.068 (11)*
N1	0.8141 (3)	1.1503 (2)	0.52847 (16)	0.0340 (5)
N2	0.8884 (3)	1.3042 (2)	0.50553 (16)	0.0373 (6)
C3	0.8995 (4)	1.2791 (3)	0.41766 (18)	0.0326 (6)
C3A	0.8266 (3)	1.1038 (3)	0.38069 (17)	0.0302 (6)
C4	0.7975 (3)	1.0105 (3)	0.28485 (17)	0.0306 (6)
H4B	0.6826	1.0230	0.2490	0.037*
C5	0.7311 (3)	0.8213 (3)	0.28392 (18)	0.0312 (6)
H5A	0.8575	0.8062	0.3058	0.037*
C6	0.5710 (3)	0.7581 (3)	0.34881 (18)	0.0316 (6)

C7	0.6747 (4)	0.8481 (3)	0.44704 (18)	0.0325 (6)
H7A	0.5710	0.8228	0.4869	0.039*
H7B	0.7819	0.8107	0.4667	0.039*
C7A	0.7705 (3)	1.0317 (3)	0.45193 (17)	0.0307 (6)
C8	0.9887 (4)	1.0726 (3)	0.23964 (18)	0.0327 (6)
C9	1.1870 (4)	1.1056 (3)	0.2856 (2)	0.0430 (7)
H9A	1.2015	1.0902	0.3449	0.052*
C10	1.3610 (5)	1.1603 (3)	0.2451 (3)	0.0546 (9)
H10A	1.4921	1.1823	0.2771	0.066*
C11	1.3422 (5)	1.1827 (4)	0.1572 (3)	0.0635 (10)
H11A	1.4600	1.2185	0.1294	0.076*
C12	1.1468 (6)	1.1516 (4)	0.1104 (3)	0.0690 (10)
H12A	1.1332	1.1688	0.0515	0.083*
C13	0.9724 (5)	1.0950 (4)	0.1516 (2)	0.0519 (8)
H13A	0.8412	1.0715	0.1193	0.062*
C14	0.6486 (4)	0.7183 (3)	0.18762 (19)	0.0350 (6)
C15	0.3862 (5)	0.6480 (4)	0.0578 (2)	0.0511 (8)
H15A	0.3097	0.5325	0.0606	0.061*
H15B	0.4869	0.6527	0.0172	0.061*
C16	0.2398 (6)	0.7190 (5)	0.0244 (3)	0.0693 (10)
H16A	0.1663	0.6560	−0.0352	0.104*
H16B	0.3173	0.8331	0.0217	0.104*
H16C	0.1409	0.7137	0.0650	0.104*
C17	0.4995 (4)	0.5701 (3)	0.3400 (2)	0.0432 (7)
H17A	0.4114	0.5349	0.3842	0.065*
H17B	0.6197	0.5438	0.3502	0.065*
H17C	0.4224	0.5133	0.2800	0.065*
H1A	0.760 (4)	1.143 (3)	0.582 (2)	0.034 (7)*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0653 (12)	0.0205 (8)	0.0410 (13)	0.0057 (8)	0.0101 (9)	0.0015 (8)
O2	0.0622 (12)	0.0460 (11)	0.0529 (15)	0.0319 (10)	−0.0016 (10)	−0.0141 (10)
O3	0.0449 (10)	0.0372 (9)	0.0360 (12)	0.0172 (8)	−0.0059 (8)	−0.0085 (8)
O4	0.0331 (9)	0.0360 (9)	0.0361 (11)	0.0138 (7)	0.0014 (7)	0.0008 (8)
O5	0.0449 (12)	0.0594 (14)	0.075 (2)	0.0095 (10)	−0.0053 (12)	0.0196 (13)
N1	0.0382 (11)	0.0261 (10)	0.0290 (14)	0.0056 (8)	0.0054 (9)	0.0006 (9)
N2	0.0442 (12)	0.0233 (10)	0.0366 (15)	0.0075 (8)	0.0055 (9)	0.0015 (9)
C3	0.0356 (12)	0.0247 (11)	0.0302 (15)	0.0065 (9)	0.0043 (10)	0.0011 (10)
C3A	0.0316 (11)	0.0210 (11)	0.0312 (15)	0.0065 (8)	0.0028 (9)	−0.0015 (10)
C4	0.0349 (12)	0.0222 (11)	0.0304 (15)	0.0095 (9)	0.0001 (10)	0.0020 (9)
C5	0.0313 (11)	0.0212 (11)	0.0361 (16)	0.0089 (9)	0.0009 (10)	−0.0002 (10)
C6	0.0315 (11)	0.0227 (11)	0.0352 (16)	0.0079 (9)	0.0020 (10)	0.0012 (10)
C7	0.0339 (12)	0.0258 (11)	0.0337 (16)	0.0085 (9)	0.0037 (10)	0.0053 (10)
C7A	0.0291 (11)	0.0249 (11)	0.0316 (15)	0.0082 (9)	−0.0002 (9)	−0.0020 (10)
C8	0.0404 (13)	0.0190 (10)	0.0341 (16)	0.0097 (9)	0.0057 (10)	−0.0003 (10)
C9	0.0459 (15)	0.0370 (13)	0.0455 (19)	0.0168 (11)	0.0067 (12)	0.0072 (12)

C10	0.0438 (15)	0.0390 (15)	0.081 (3)	0.0164 (12)	0.0190 (15)	0.0091 (15)
C11	0.063 (2)	0.0436 (16)	0.080 (3)	0.0124 (14)	0.0382 (19)	0.0118 (17)
C12	0.087 (3)	0.062 (2)	0.048 (2)	0.0133 (18)	0.0266 (18)	0.0183 (17)
C13	0.0565 (17)	0.0467 (16)	0.044 (2)	0.0117 (13)	0.0081 (14)	0.0102 (14)
C14	0.0347 (12)	0.0235 (11)	0.0412 (17)	0.0080 (9)	0.0060 (11)	0.0015 (10)
C15	0.0574 (17)	0.0436 (15)	0.0371 (19)	0.0143 (13)	−0.0075 (13)	−0.0090 (13)
C16	0.074 (2)	0.070 (2)	0.053 (2)	0.0300 (18)	−0.0157 (17)	−0.0018 (17)
C17	0.0453 (14)	0.0245 (12)	0.051 (2)	0.0060 (10)	0.0055 (12)	0.0053 (11)

Geometric parameters (Å, °)

O1—C3	1.308 (3)	C7—C7A	1.493 (3)
O1—H1B	0.8200	C7—H7A	0.9700
O2—C14	1.209 (3)	C7—H7B	0.9700
O3—C14	1.321 (3)	C8—C13	1.379 (4)
O3—C15	1.462 (3)	C8—C9	1.394 (4)
O4—C6	1.425 (3)	C9—C10	1.371 (4)
O4—H4A	0.8200	C9—H9A	0.9300
O5—H5B	0.95 (5)	C10—C11	1.378 (5)
O5—H5C	0.85 (4)	C10—H10A	0.9300
N1—C7A	1.354 (3)	C11—C12	1.385 (6)
N1—N2	1.379 (3)	C11—H11A	0.9300
N1—H1A	0.92 (3)	C12—C13	1.381 (5)
N2—C3	1.326 (4)	C12—H12A	0.9300
C3—C3A	1.432 (3)	C13—H13A	0.9300
C3A—C7A	1.355 (4)	C15—C16	1.482 (5)
C3A—C4	1.502 (3)	C15—H15A	0.9700
C4—C8	1.514 (3)	C15—H15B	0.9700
C4—C5	1.558 (3)	C16—H16A	0.9600
C4—H4B	0.9800	C16—H16B	0.9600
C5—C14	1.517 (3)	C16—H16C	0.9600
C5—C6	1.561 (4)	C17—H17A	0.9600
C5—H5A	0.9800	C17—H17B	0.9600
C6—C17	1.526 (3)	C17—H17C	0.9600
C6—C7	1.532 (3)		
C3—O1—H1B	109.5	C3A—C7A—C7	125.0 (2)
C14—O3—C15	117.9 (2)	C13—C8—C9	117.9 (3)
C6—O4—H4A	109.5	C13—C8—C4	121.4 (2)
H5B—O5—H5C	105 (4)	C9—C8—C4	120.6 (2)
C7A—N1—N2	108.2 (2)	C10—C9—C8	121.2 (3)
C7A—N1—H1A	130.0 (17)	C10—C9—H9A	119.4
N2—N1—H1A	117.5 (17)	C8—C9—H9A	119.4
C3—N2—N1	107.5 (2)	C9—C10—C11	120.2 (3)
O1—C3—N2	123.5 (2)	C9—C10—H10A	119.9
O1—C3—C3A	126.8 (2)	C11—C10—H10A	119.9
N2—C3—C3A	109.6 (2)	C10—C11—C12	119.6 (3)
C7A—C3A—C3	104.3 (2)	C10—C11—H11A	120.2
C7A—C3A—C4	124.9 (2)	C12—C11—H11A	120.2
C3—C3A—C4	130.6 (2)	C13—C12—C11	119.6 (4)

C3A—C4—C8	114.11 (18)	C13—C12—H12A	120.2
C3A—C4—C5	109.1 (2)	C11—C12—H12A	120.2
C8—C4—C5	109.85 (19)	C8—C13—C12	121.4 (3)
C3A—C4—H4B	107.8	C8—C13—H13A	119.3
C8—C4—H4B	107.8	C12—C13—H13A	119.3
C5—C4—H4B	107.8	O2—C14—O3	123.6 (2)
C14—C5—C4	110.5 (2)	O2—C14—C5	125.0 (2)
C14—C5—C6	111.47 (18)	O3—C14—C5	111.4 (2)
C4—C5—C6	112.80 (19)	O3—C15—C16	107.3 (2)
C14—C5—H5A	107.2	O3—C15—H15A	110.3
C4—C5—H5A	107.2	C16—C15—H15A	110.3
C6—C5—H5A	107.2	O3—C15—H15B	110.3
O4—C6—C17	111.01 (18)	C16—C15—H15B	110.3
O4—C6—C7	106.04 (18)	H15A—C15—H15B	108.5
C17—C6—C7	109.6 (2)	C15—C16—H16A	109.5
O4—C6—C5	110.4 (2)	C15—C16—H16B	109.5
C17—C6—C5	110.5 (2)	H16A—C16—H16B	109.5
C7—C6—C5	109.18 (18)	C15—C16—H16C	109.5
C7A—C7—C6	109.0 (2)	H16A—C16—H16C	109.5
C7A—C7—H7A	109.9	H16B—C16—H16C	109.5
C6—C7—H7A	109.9	C6—C17—H17A	109.5
C7A—C7—H7B	109.9	C6—C17—H17B	109.5
C6—C7—H7B	109.9	H17A—C17—H17B	109.5
H7A—C7—H7B	108.3	C6—C17—H17C	109.5
N1—C7A—C3A	110.2 (2)	H17A—C17—H17C	109.5
N1—C7A—C7	124.8 (2)	H17B—C17—H17C	109.5
C7A—N1—N2—C3	−3.1 (3)	C3—C3A—C7A—N1	−3.0 (3)
N1—N2—C3—O1	−178.5 (2)	C4—C3A—C7A—N1	−178.8 (2)
N1—N2—C3—C3A	1.2 (3)	C3—C3A—C7A—C7	177.1 (2)
O1—C3—C3A—C7A	−179.2 (2)	C4—C3A—C7A—C7	1.3 (4)
N2—C3—C3A—C7A	1.1 (3)	C6—C7—C7A—N1	157.6 (2)
O1—C3—C3A—C4	−3.8 (4)	C6—C7—C7A—C3A	−22.5 (3)
N2—C3—C3A—C4	176.5 (2)	C3A—C4—C8—C13	−135.7 (2)
C7A—C3A—C4—C8	−133.3 (2)	C5—C4—C8—C13	101.3 (3)
C3—C3A—C4—C8	52.1 (3)	C3A—C4—C8—C9	45.6 (3)
C7A—C3A—C4—C5	−10.0 (3)	C5—C4—C8—C9	−77.3 (3)
C3—C3A—C4—C5	175.4 (2)	C13—C8—C9—C10	0.7 (4)
C3A—C4—C5—C14	165.98 (18)	C4—C8—C9—C10	179.3 (2)
C8—C4—C5—C14	−68.2 (2)	C8—C9—C10—C11	−0.4 (4)
C3A—C4—C5—C6	40.4 (2)	C9—C10—C11—C12	0.8 (5)
C8—C4—C5—C6	166.2 (2)	C10—C11—C12—C13	−1.4 (5)
C14—C5—C6—O4	−72.4 (2)	C9—C8—C13—C12	−1.3 (4)
C4—C5—C6—O4	52.7 (2)	C4—C8—C13—C12	−180.0 (3)
C14—C5—C6—C17	50.8 (3)	C11—C12—C13—C8	1.7 (5)
C4—C5—C6—C17	175.87 (19)	C15—O3—C14—O2	2.6 (4)
C14—C5—C6—C7	171.44 (19)	C15—O3—C14—C5	−175.3 (2)
C4—C5—C6—C7	−63.5 (2)	C4—C5—C14—O2	126.8 (3)
O4—C6—C7—C7A	−68.0 (2)	C6—C5—C14—O2	−106.9 (3)
C17—C6—C7—C7A	172.1 (2)	C4—C5—C14—O3	−55.4 (3)

C5—C6—C7—C7A	50.9 (2)	C6—C5—C14—O3	70.9 (3)
N2—N1—C7A—C3A	3.9 (3)	C14—O3—C15—C16	−172.8 (3)
N2—N1—C7A—C7	−176.2 (2)		

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O1—H1B···N2^{i}	0.82	1.89	2.705 (2)	171
O4—H4A···O3^{i}	0.82	2.22	2.897 (2)	141
N1—H1A···O4^{ii}	0.93 (3)	1.94	2.778 (3)	155
O5—H5B···O1^{i}	0.95 (5)	2.01	2.874 (2)	165
O5—H5C···O2^{ii}	0.84 (4)	1.97	2.844 (3)	179

Symmetry codes: (i) ; (ii) .

Fig. 1

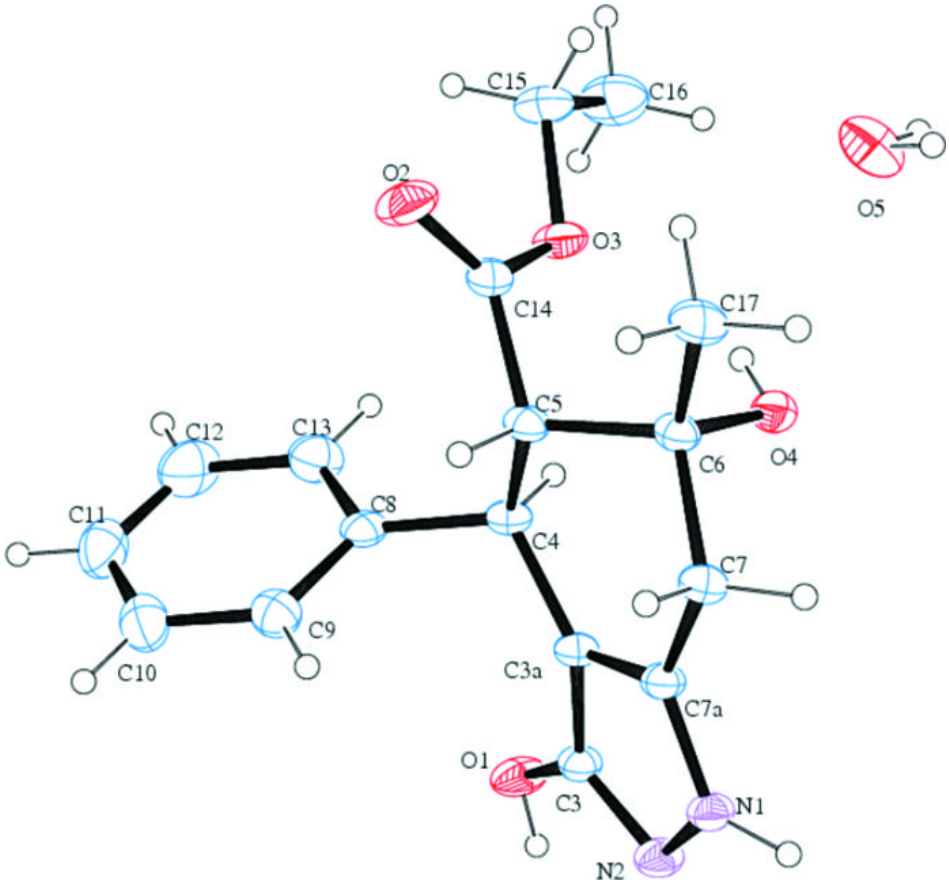
C15
C16
O2
O3
O5
C14
C17
C5
C12
C13
O4
C6
C11
C8
C4
C7
C9
C10
C3a
C7a
O1
C3
N1
N2

Fig. 2