

2-Hydroxy-3,3-dimethyl-7-nitro-3,4-dihydroisoquinolin-1(2H)-one

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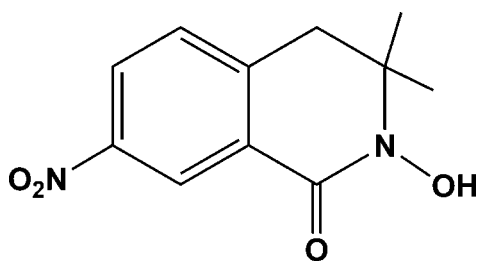
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.151; data-to-parameter ratio = 20.9.

In the title compound, $\text{C}_{11}\text{H}_{12}\text{N}_2\text{O}_4$, a new hydroxamic acid which belonging to the isoquinole family, the heterocyclic ring adopts a half-chair conformation. The nitro group is essentially coplanar with the aromatic ring. Intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds assemble the molecules around inversion centres to form pseudo-dimers.

Related literature

For related literature, see: Bohé & Kammoun (2004); Kurzak *et al.* (1992); Porcheddu & Giacomelli (2006); Weber (1983); Miller (1989); Cremer & Pople (1975).



Experimental

Crystal data

$\text{C}_{11}\text{H}_{12}\text{N}_2\text{O}_4$
 $M_r = 236.23$
Monoclinic, $P2_1/n$

$a = 5.8805$ (9) Å
 $b = 18.605$ (4) Å
 $c = 10.1588$ (17) Å

$\beta = 103.056$ (12)°
 $V = 1082.7$ (3) Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.11$ mm⁻¹
 $T = 296$ K
 $0.60 \times 0.51 \times 0.22$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (Coppens *et al.*, 1965)
 $T_{\min} = 0.938$, $T_{\max} = 0.975$
7659 measured reflections
3286 independent reflections
2051 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.150$
 $S = 1.11$
3286 reflections

157 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.26$ e Å⁻³
 $\Delta\rho_{\min} = -0.27$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O12}-\text{H12}\cdots\text{O11}^i$	0.82	1.99	2.7013 (14)	144
$\text{O12}-\text{H12}\cdots\text{O11}$	0.82	2.20	2.6305 (15)	113

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-III* (Burnett & Johnson, 1996), *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2332).

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supplementary materials

Acta Cryst. (2008). E64, o1050 [doi:10.1107/S1600536808013457]

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Comment

Hydroxamic acids are strong metal ion chelators (Kurzak *et al.*, 1992), they possess a wide spectrum of biological activities, such as antibacterial, antifungal, anti-inflammatory, and anti-asthmatic properties, *etc.* (Weber, 1983; Miller 1989).

The growing number of published synthetic methods further points to the biological significance of hydroxamic acids (Porcheddu & Giacomelli, 2006). Among this family of hydroxamic acids is the title compound (1). We report herein its synthesis and its crystal structure determination. Synthesis of the title compound has been prepared from the corresponding dihydroisoquinoline (2) by metachloroperbenzoic acid oxidation (Fig. 1). Imine (2) was described by Bohé and Kammoun (2004), in three steps from the commercially available tertiary alcohol (3).

In the title compound, the heterocyclic ring adopts a half-chair conformation as indicated by the puckering parameters: $Q = 0.4224(14)\text{Å}$, $\theta = 57.87(18)^\circ$, $\phi = 281.2(2)^\circ$ (Cremer & Pople, 1975). The nitro group attached on C7 is essentially coplanar with the aromatic nucleus (Fig. 2). The methyl substituent in position 3 of the heterocyclic ring is pseudo-axial, the second methyl in position 3 is pseudo-equatorial.

The conformation of (1) is stabilized by an intramolecular hydrogen bond between the hydroxyl O12—H12 group and atom O11 (Table 1). The molecules are assembled by intermolecular O—H...O hydrogen bonds to form pseudo dimer arranged around inversion centre (Table 1, Fig. 3)

Experimental

The title compound was prepared by reaction of imine (2) (100 mg, 0.49 mmol), and methachloroperbenzoic acid 86% (197 mg, 0.98 mmol) in dichloromethane (15 ml). The mixture was stirred at room temperature for 24 h. Then, the mixture was diluted with CH_2Cl_2 and washed with a solution of saturated NaHCO_3 . The organic phase was dried over sodium sulfate, filtered and concentrated under reduced pressure. The concentrate was chromatographed on silica gel, with (ether) as eluent (yield 40%). m.p. 418 K. Spectroscopic analysis, ^1H NMR (300 MHz; DMSO-d_6 , p.p.m): 1.26 (s, 6H, 2Me 3); 3.23 (s, 2H); 7.59 (d, $J = 8.4$, 1H, aromatic H); 8.33 (dd, $J = 8.4$, $J = 2.4$, 1H, aromatic H); 8.56 (d, $J = 2.4$, 1H aromatic H); 9.8 (s wide, 1H, OH). ^{13}C NMR (75 MHz; DMSO-d_6 , p.p.m): 25.24; 41.80; 60.66; 122.02; 126.87; 129.91; 130.41; 144.05; 147.21; 160.08. M.S (EI, 70 ev): m/z : 236 (M^+); 221 [($M-15$) $^+$, base peak]. MS (HR): Found: 236,0844 calcd mass for $\text{C}_{11}\text{H}_{12}\text{N}_2\text{O}_4$: 236,0875. Recrystallizations from dichloromethane afford yellow crystals suitable for diffraction.

Refinement

All H atoms attached to C atoms and O atom were fixed geometrically and treated as riding with C—H = 0.98 Å (methyl), 0.97 Å (methylene), 0.93 Å (aromatic) and O—H = 0.82 Å with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O}, \text{C}_{\text{aromatic}})$.

Figures

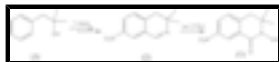


Fig. 1. Chemical pathway of the formation of hydroxamic acid.

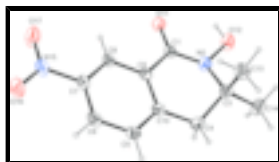


Fig. 2. Molecular view of the title compound with the atom-labelling scheme. Ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii.

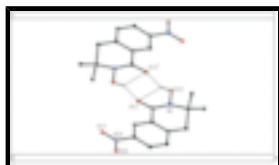


Fig. 3. Partial packing view showing the formation of pseudo dimer through O-H...O hydrogen bonds. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bondings have been omitted for clarity. [Symmetry code: (i) 1-x, 1-y, 1-z]

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Crystal data

$C_{11}H_{12}N_2O_4$

$M_r = 236.23$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 5.8805$ (9) Å

$b = 18.605$ (4) Å

$c = 10.1588$ (17) Å

$\beta = 103.056$ (12)°

$V = 1082.7$ (3) Å³

$Z = 4$

$F_{000} = 496$

$D_x = 1.449$ Mg m⁻³

Melting point: 418 K

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 2421 reflections

$\theta = 2.5$ – 23.2 °

$\mu = 0.11$ mm⁻¹

$T = 296$ K

Prism, colourless

$0.60 \times 0.51 \times 0.22$ mm

Data collection

Bruker SMART CCD area-detector diffractometer

Radiation source: sealed tube

Monochromator: graphite

$T = 296$ K

φ and ω scans

Absorption correction: multi-scan (Coppens *et al.*, 1965)

$T_{\min} = 0.938$, $T_{\max} = 0.975$

7659 measured reflections

3286 independent reflections

2051 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$

$\theta_{\text{max}} = 30.4$ °

$\theta_{\text{min}} = 2.2$ °

$h = -8$ → 8

$k = 0$ → 26

$l = 0$ → 14

Refinement

Refinement on F^2

Secondary atom site location: difference Fourier map

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.044$$

$$wR(F^2) = 0.151$$

$$S = 1.11$$

3286 reflections

157 parameters

Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0543P)^2 + 0.0949P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.26 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$$

Extinction correction: none

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.

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	0.7652 (2)	0.53911 (7)	0.36105 (13)	0.0323 (3)
C3	0.7990 (2)	0.67410 (7)	0.36068 (15)	0.0380 (3)
C4	0.8660 (3)	0.66689 (8)	0.22427 (15)	0.0426 (3)
H4A	0.7256	0.6692	0.1527	0.051*
H4B	0.9645	0.7072	0.2127	0.051*
C5	1.1554 (3)	0.59404 (8)	0.13045 (14)	0.0393 (3)
H5	1.1903	0.6347	0.0857	0.047*
C6	1.2662 (3)	0.52948 (8)	0.11699 (14)	0.0410 (3)
H6	1.3779	0.5268	0.0653	0.049*
C7	1.2075 (2)	0.46956 (7)	0.18163 (13)	0.0339 (3)
C8	1.0435 (2)	0.47108 (7)	0.25982 (13)	0.0330 (3)
H8	1.0052	0.4297	0.3015	0.040*
C9	0.9371 (2)	0.53674 (7)	0.27418 (12)	0.0296 (2)
C10	0.9923 (2)	0.59838 (7)	0.21048 (13)	0.0338 (3)
C13	1.0066 (3)	0.68808 (8)	0.47591 (16)	0.0471 (3)
H13A	0.9566	0.6890	0.5596	0.071*
H13B	1.0752	0.7335	0.4625	0.071*
H13C	1.1200	0.6506	0.4789	0.071*
C14	0.6172 (3)	0.73411 (8)	0.35087 (19)	0.0538 (4)
H14A	0.4823	0.7226	0.2813	0.081*
H14B	0.6829	0.7787	0.3294	0.081*
H14C	0.5727	0.7386	0.4358	0.081*

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N2	0.6880 (2)	0.60494 (6)	0.38233 (13)	0.0402 (3)
N15	1.3238 (2)	0.40117 (7)	0.16643 (13)	0.0438 (3)
O11	0.69294 (19)	0.48475 (5)	0.40857 (11)	0.0437 (3)
O12	0.5448 (2)	0.61049 (6)	0.47313 (13)	0.0542 (3)
H12	0.5129	0.5702	0.4961	0.081*
O16	1.4723 (3)	0.40057 (7)	0.09958 (16)	0.0701 (4)
O17	1.2691 (3)	0.34788 (6)	0.22078 (16)	0.0724 (5)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0312 (5)	0.0344 (6)	0.0338 (6)	-0.0011 (5)	0.0129 (4)	-0.0013 (5)
C3	0.0393 (7)	0.0316 (6)	0.0463 (8)	0.0024 (5)	0.0163 (5)	0.0012 (5)
C4	0.0514 (8)	0.0382 (7)	0.0403 (7)	0.0039 (6)	0.0146 (6)	0.0078 (5)
C5	0.0468 (8)	0.0409 (7)	0.0340 (7)	-0.0071 (5)	0.0172 (5)	0.0027 (5)
C6	0.0422 (7)	0.0506 (8)	0.0357 (7)	-0.0047 (6)	0.0203 (5)	-0.0025 (6)
C7	0.0331 (6)	0.0385 (7)	0.0323 (6)	-0.0022 (5)	0.0118 (4)	-0.0066 (5)
C8	0.0341 (6)	0.0356 (6)	0.0316 (6)	-0.0023 (4)	0.0125 (4)	-0.0028 (5)
C9	0.0251 (5)	0.0356 (6)	0.0300 (5)	-0.0015 (4)	0.0102 (4)	-0.0009 (4)
C10	0.0369 (6)	0.0378 (7)	0.0283 (6)	-0.0015 (5)	0.0104 (4)	0.0001 (5)
C13	0.0491 (8)	0.0449 (8)	0.0478 (8)	-0.0067 (6)	0.0120 (6)	-0.0062 (6)
C14	0.0549 (10)	0.0395 (8)	0.0710 (11)	0.0094 (7)	0.0224 (8)	0.0025 (7)
N2	0.0398 (6)	0.0384 (6)	0.0484 (7)	0.0012 (5)	0.0225 (5)	-0.0004 (5)
N15	0.0435 (7)	0.0449 (7)	0.0484 (7)	-0.0015 (5)	0.0216 (5)	-0.0107 (5)
O11	0.0474 (6)	0.0392 (5)	0.0530 (6)	-0.0016 (4)	0.0290 (5)	0.0036 (4)
O12	0.0577 (7)	0.0439 (6)	0.0759 (8)	0.0026 (5)	0.0466 (6)	-0.0024 (5)
O16	0.0803 (10)	0.0621 (8)	0.0872 (10)	0.0148 (7)	0.0593 (9)	0.0031 (7)
O17	0.0887 (11)	0.0377 (6)	0.1104 (12)	-0.0020 (6)	0.0633 (10)	-0.0055 (7)

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

C1—O11	1.2366 (15)	C7—C8	1.3816 (17)
C1—N2	1.3407 (16)	C7—N15	1.4690 (18)
C1—C9	1.4851 (16)	C8—C9	1.3950 (17)
C3—N2	1.4818 (18)	C8—H8	0.9300
C3—C13	1.511 (2)	C9—C10	1.3905 (17)
C3—C4	1.530 (2)	C13—H13A	0.9600
C3—C14	1.533 (2)	C13—H13B	0.9600
C4—C10	1.497 (2)	C13—H13C	0.9600
C4—H4A	0.9700	C14—H14A	0.9600
C4—H4B	0.9700	C14—H14B	0.9600
C5—C6	1.388 (2)	C14—H14C	0.9600
C5—C10	1.3927 (18)	N2—O12	1.3862 (14)
C5—H5	0.9300	N15—O17	1.2130 (17)
C6—C7	1.3768 (19)	N15—O16	1.2215 (16)
C6—H6	0.9300	O12—H12	0.8200
O11—C1—N2	121.69 (11)	C9—C8—H8	121.1
O11—C1—C9	123.22 (11)	C10—C9—C8	121.09 (11)

N2—C1—C9	115.07 (11)	C10—C9—C1	120.91 (11)
N2—C3—C13	109.91 (12)	C8—C9—C1	118.00 (11)
N2—C3—C4	105.70 (11)	C9—C10—C5	119.18 (12)
C13—C3—C4	112.86 (12)	C9—C10—C4	119.07 (12)
N2—C3—C14	108.53 (12)	C5—C10—C4	121.68 (12)
C13—C3—C14	110.76 (13)	C3—C13—H13A	109.5
C4—C3—C14	108.88 (12)	C3—C13—H13B	109.5
C10—C4—C3	113.20 (11)	H13A—C13—H13B	109.5
C10—C4—H4A	108.9	C3—C13—H13C	109.5
C3—C4—H4A	108.9	H13A—C13—H13C	109.5
C10—C4—H4B	108.9	H13B—C13—H13C	109.5
C3—C4—H4B	108.9	C3—C14—H14A	109.5
H4A—C4—H4B	107.8	C3—C14—H14B	109.5
C6—C5—C10	120.56 (12)	H14A—C14—H14B	109.5
C6—C5—H5	119.7	C3—C14—H14C	109.5
C10—C5—H5	119.7	H14A—C14—H14C	109.5
C7—C6—C5	118.67 (12)	H14B—C14—H14C	109.5
C7—C6—H6	120.7	C1—N2—O12	117.02 (11)
C5—C6—H6	120.7	C1—N2—C3	126.35 (11)
C6—C7—C8	122.71 (12)	O12—N2—C3	112.85 (10)
C6—C7—N15	118.59 (11)	O17—N15—O16	122.81 (13)
C8—C7—N15	118.70 (12)	O17—N15—C7	118.86 (12)
C7—C8—C9	117.77 (11)	O16—N15—C7	118.34 (12)
C7—C8—H8	121.1	N2—O12—H12	109.5

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O12—H12 \cdots O11 ⁱ	0.82	1.99	2.7013 (14)	144
O12—H12 \cdots O11	0.82	2.20	2.6305 (15)	113

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

Fig. 1

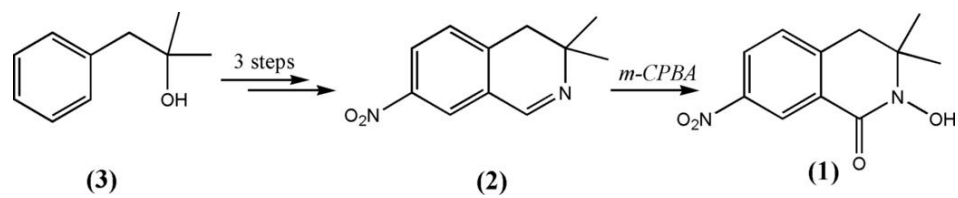


Fig. 2

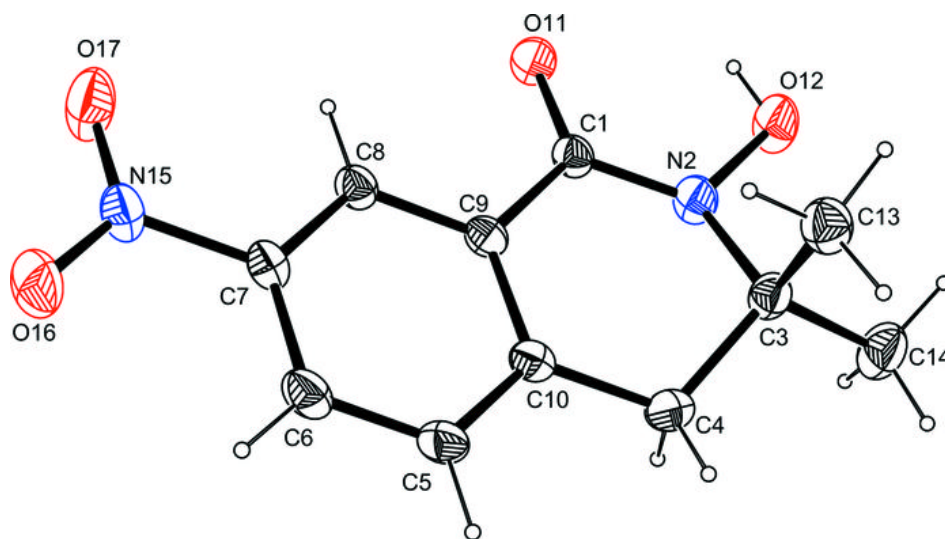


Fig. 3

