

1-Allyl-4-hydroxy-2-oxo-1,2,5,6,7,8-hexahydroquinoline-3-carboxylic acid

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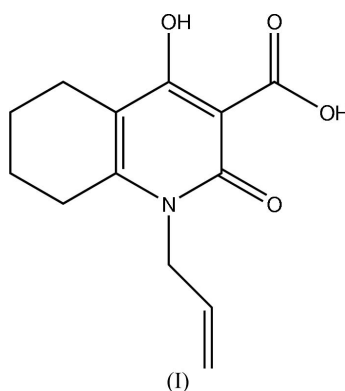
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.047
 wR factor = 0.132
Data-to-parameter ratio = 12.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

An X-ray diffraction study of the title compound, $\text{C}_{13}\text{H}_{15}\text{NO}_4$, has demonstrated that the formation of intramolecular $\text{O}-\text{H}\cdots\text{O}=\text{C}$ hydrogen bonds leads to the lengthening of the carbonyl $\text{C}=\text{O}$ bonds, shortening of the $(\text{C}=\text{C})-\text{C}-\text{O}(-\text{H})$ bond and delocalization of the electron density within the dihydropyridine ring.

Comment

N-substituted 4-hydroxy-2-oxo-1,2-dihydroquinoline-3-carboxylic acids possess interesting pharmacological properties (Rowley *et al.*, 1993; Ukrainets *et al.*, 1994). These compounds are also used for the synthesis of different drugs, for example, the synthetic immunomodulator Roquinimex (trademark name for 4-hydroxy-1-methyl-2-oxo-1,2-dihydroquinoline-3-carboxylic acid methyl-phenyl amide; Jönsson *et al.*, 2004) and the injectable anesthetic Chinoxicaine (trademark name for 4-hydroxy-2-oxo-1-propyl-1,2-dihydroquinoline-3-carboxylic acid diethylaminoethyl amide hydrochloride; Ukrayinecz & Bezuhliy, 2002). Structures of *N*-substituted 4-hydroxy-2-oxo-1,2,5,6,7,8-hexahydroquinoline-3-carboxylic acids should be similar to their 1,2-dihydro analogues but remain almost unexplored. In the present paper, we report the crystal structure of the title *N*-allyl derivative, (I) (Table 1).



The cyclohexene ring of (I) has a half-chair conformation, with atoms C6 and C7 displaced from the plane of the remaining ring atoms by -0.382 (6) and 0.360 (6) Å, respectively. All non-H atoms of the pyridine ring, the carboxy, hydroxy and carbonyl groups, and atom C10 are almost coplanar (Fig. 1). The formation of the $\text{O4}-\text{H4O}\cdots\text{O2}$ and $\text{O3}-\text{H3O}\cdots\text{O1}$ intramolecular hydrogen bonds (Table 2) leads to a significant change of bond lengths in the hydroxy, carboxy and carbonyl groups; the $\text{O1}-\text{C1}$ and $\text{O2}-\text{C13}$ bonds are long compared with standard value for a $\text{C}=\text{O}$ bond (1.210 Å; Bürgi & Dunitz, 1994), and the $\text{O4}-\text{C3}$ bond

Received 16 March 2005

Accepted 12 May 2005

Online 21 May 2005

is shorter than the standard value for the Csp^2-O bond length (1.362 Å). The presence of the hydrogen bonds results in the delocalization of the electron density within the dihydroquinoline ring; the $C2=C3$ and $C4=C9$ bonds are longer and the $C1-C2$ and $C3-C4$ bonds are shorter than standard values for $C=C$ (1.334 Å) and Csp^2-Csp^2 (1.455 Å) bond lengths, respectively (Bürgi & Dunitz, 1994). The structure of this fragment is very similar to that observed for 1-ethyl-4-hydroxy-2-oxo-1,2-dihydroquinoline-3-carboxylic acid (Shishkina *et al.*, 2002).

The repulsion between the substituent at atom N1 and the methylene and carbonyl groups is reflected in short intramolecular contacts $O1 \cdots H10A = 2.39$ Å [van der Waals radii sum is 2.72 Å (Bondi, 1964)], $H8A \cdots C10 = 2.66$ Å (van der Waals sum 2.90 Å), $H8A \cdots H10B = 2.22$ Å (van der Waals sum 2.40 Å) and $H8B \cdots H10B = 2.27$ Å (van der Waals sum 2.40 Å). This hindered arrangement causes a lengthening of the $N1-C1$, $N1-C9$ and $N1-C10$ bonds [standard values for $N-Csp^2$ and $N-Csp^3$ bonds lengths are 1.355 and 1.464 Å, respectively (Bürgi & Dunitz, 1994)]. The double bond of the allyl group is situated almost orthogonal to the plane of the pyridine ring; the $C1-N1-C10-C11$ torsion angle is $81.7(3)^\circ$ and has *+ac* orientation with respect to the $N1-C10$ bond, while the $N1-C10-C11-C12$ torsion angle is $120.3(3)^\circ$.

Experimental

The title compound was prepared by hydrolysis of 1-allyl-4-hydroxy-2-oxo-1,2,5,6,7,8-hexahydroquinoline-3-carboxylic acid ethyl ester, following a published procedure (Ukrainets *et al.*, 2004; Jönsson *et al.*, 2004).

Crystal data

$C_{13}H_{15}NO_4$	Mo $K\alpha$ radiation
$M_r = 249.26$	Cell parameters from 24 reflections
Orthorhombic, <i>Pbca</i>	$\theta = 10-11^\circ$
$a = 16.823(4)$ Å	$\mu = 0.10$ mm $^{-1}$
$b = 7.179(2)$ Å	$T = 293(2)$ K
$c = 19.847(5)$ Å	Needle, colourless
$V = 2397.0(11)$ Å 3	$0.30 \times 0.10 \times 0.10$ mm
$Z = 8$	
$D_x = 1.381$ Mg m $^{-3}$	

Data collection

Siemens P3/PC diffractometer	$h = 0 \rightarrow 20$
$\omega-2\theta$ scans	$k = 0 \rightarrow 8$
2087 measured reflections	$l = -23 \rightarrow 0$
2086 independent reflections	2 standard reflections
935 reflections with $I > 2\sigma(I)$	every 98 reflections
$R_{int} = 0.030$	intensity decay: 5%
$\theta_{max} = 25.1^\circ$	

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.047$	$w = 1/[\sigma^2(F_o^2) + (0.0714P)^2]$
$wR(F^2) = 0.132$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.85$	$(\Delta/\sigma)_{max} < 0.001$
2086 reflections	$\Delta\rho_{max} = 0.18$ e Å $^{-3}$
165 parameters	$\Delta\rho_{min} = -0.17$ e Å $^{-3}$

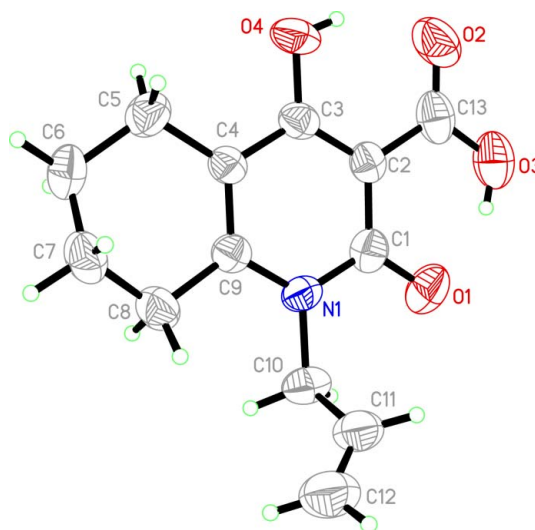


Figure 1

View of the title compound. Non-H atoms are shown with displacement ellipsoids drawn at the 50% probability level.

Table 1

Selected interatomic distances (Å).

$N1-C1$	1.381 (3)	$O4-C3$	1.339 (3)
$N1-C9$	1.397 (3)	$C1-C2$	1.436 (4)
$N1-C10$	1.485 (3)	$C2-C3$	1.395 (4)
$O1-C1$	1.268 (3)	$C2-C13$	1.468 (4)
$O2-C13$	1.236 (4)	$C3-C4$	1.421 (4)
$O3-C13$	1.320 (4)	$C4-C9$	1.384 (4)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O3-H3O \cdots O1$	0.82	1.73	2.485 (3)	153
$O4-H4O \cdots O2$	0.82	1.85	2.576 (3)	147

H atoms bonded to C atoms were placed in calculated positions, while H atoms of the hydroxy groups were located in difference maps. All H atoms were refined in the riding model approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(O)$; constrained distances: 0.97 Å for methylene CH_2 groups, 0.93 Å for $C(allyl)H$ groups and 0.82 Å for hydroxy OH groups.

Data collection: *P3* (Siemens, 1989); cell refinement: *P3*; data reduction: *XDISK* (Siemens, 1991); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1991); software used to prepare material for publication: *SHELXL97*.

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