

Synthesis and Crystal Structure of *N*-Acryloylcytisine and *N*-(β -Morpholinopropionyl)cytisine

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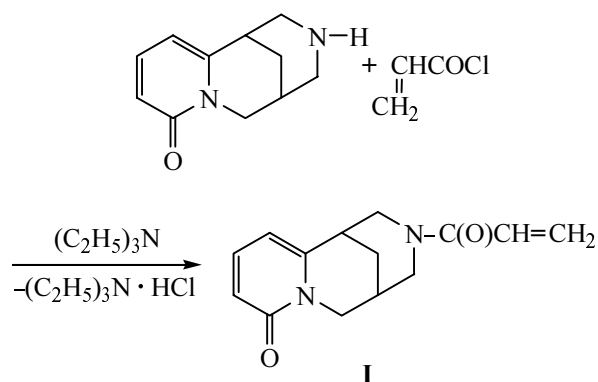
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Abstract—*N*-acryloylcytisine and *N*-(β -morpholinopropionyl)cytisine were synthesized and spatial structures of compounds obtained were established by X-ray diffraction analysis.

Investigations of antimicrobial activity of synthetic derivatives of cytisine revealed that it depended essentially on the chemical structure of the side chains of alkaloid attached to nitrogen [1–3].

Aiming at refining understanding of effect on the physiological activity of cytisine derivatives produced by various substituents with an activated double bond we synthesized *N*-acryloylcytisine (**I**) in 95% yield. The alkaloid was acylated with acryloyl chloride in anhydrous benzene by a standard procedure.



The acryl fragment of compound **I** was modified by adding morpholinodithiocarbamic acid made *in situ* from carbon disulfide and morpholine. The main product isolated from the reaction mixture was *N*-(β -morpholinopropionyl)cytisine (**II**) obtained in 86% yield. Apparently the morpholinodithiocarbamic acid decomposed, and morpholine added to the *N*-acryloylcytisine across the double bond (see the scheme).

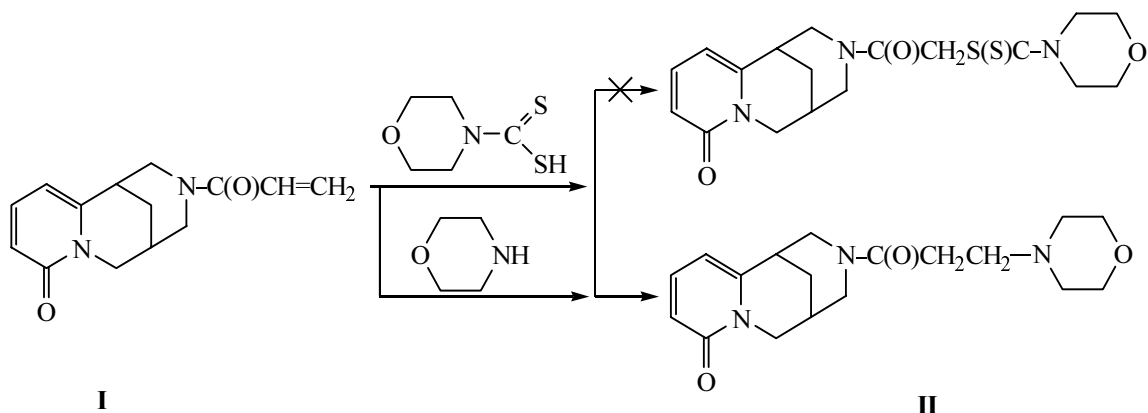
Compound **II** is a colorless crystalline substance soluble in most organic solvents. The structure of com-

pounds **I** and **II** was established by means of IR, ¹H NMR spectroscopy, and X-ray diffraction analysis. General view of molecules **I** and **II** is presented on Figs. 1 and 2.

As demonstrated by the X-ray diffraction study the bond lengths and bond angles in the cytisine skeleton of structures **I** and **II** are close to those in *N*-methylcytisine (**III**) [4], *N*-cyanomethyl-cytisine (**IV**) [5], and *O,O*-dimethyl-*N*-cytisinylamidophosphate (**V**) [6] except for the bond angles at the atom N¹². In molecules **III** and **IV** the atom N¹² is present in a pyramidal coordination (sum of bond angles is 335.7° and 334.0° respectively), whereas in molecules **I** and **II**, like in molecule **V**, the coordination is trigonal-planar (bond angles sum is 360, 359.8, and 354.8° respectively).

The different configuration of nitrogen in molecules **III** and **IV**, on the one hand, and in molecules **I**, **II**, and **V**, on the other hand, is due to mesomeric effect involving a lone electron pair of atom N¹² and conjugation system of the acryloyl moiety or the double bond P¹=O² in compound **V**. Therewith the bond N¹²–C¹⁴ gets shorter, and the bond C¹⁴=O¹⁴ somewhat longer as compared to standard values (Table 1). The dihydropyridine ring in structures **I** and **II** is planar within $\pm(0.015, 0.017)$ Å respectively, the carbonyl oxygen O² is located practically in this plane (its deviation from the plane is 0.06 Å for **I** and 0.05 Å for **II**). The tetrahydropyridine ring N¹C⁶C⁷C⁸C⁹C¹⁰ takes a conformation of *distorted sofa* [ΔC_s^8 6.3 (**I**), ΔC_s^8 7.29 (**II**), with deviation of the bridging atom C⁸ from the mean plane of the other atoms of the ring by 0.74 (**I**), 0.72 (**II**) Å]. Piperidine ring has a *distorted chair* conformation [ΔC_s^7 0.58 Å (min.), ΔC_s^8 3.55 Å (max) (**I**), ΔC_s^7 0.92 Å (min), ΔC_s^8 3.68 Å (max)

Scheme.



(II)]. The morpholine ring N¹⁷C²¹C²²O²⁰C¹⁸C¹⁹ in II molecule is present in the *chair* conformation [ΔC_S^{19} 1.47 Å(min), ΔC_S^{17} 1.84 Å(max)].

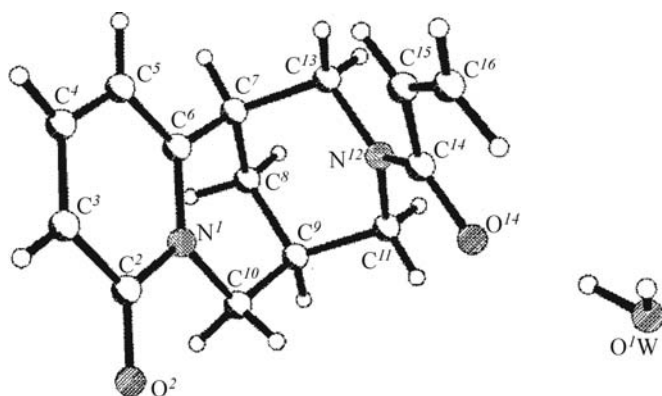


Fig. 1. Structure of N¹²-acryloylcytisine (I) molecule.

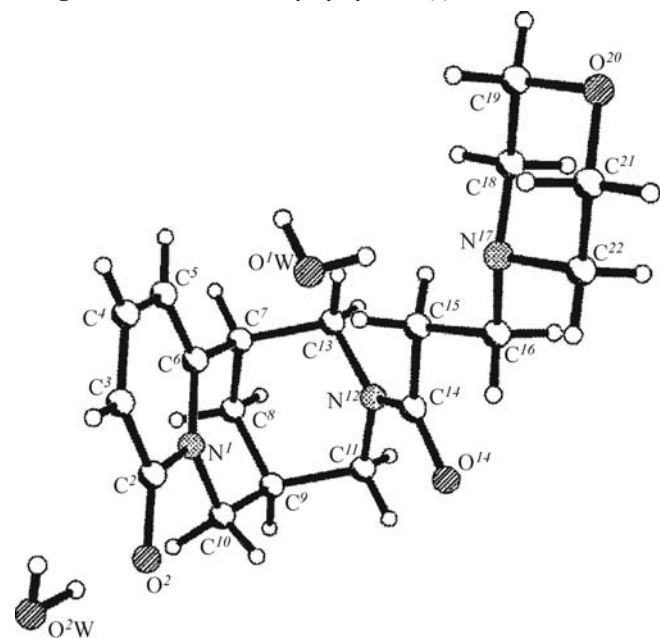


Fig. 2. Structure of N¹²-(b-morpholinopropionyl)cytosine (II).

The comparison of torsion angles in the cytosine fragment of molecules I and II with the respective angles in molecules III and IV shows that the difference does not exceed 8.6° (Table 2).

Thus the conformation of rings in the cytosine molecule crystal is relatively rigid. Therewith the tetrahydropyridine ring is more susceptible to the effect of substituent attached to atom N¹², and the influence of the latter affects the symmetry. The piperidine ring with a good precision conserves its symmetry disregarding the dimensions of the substituent at N¹² that is equatorially directed, and atom N¹² can change its configuration due to the mesomeric effect.

The above reasoning suggests a conclusion that introducing substituents to atom N¹² virtually does not affect the conformation of the cytosine skeleton (Tables 1–3).

In the IR spectra of compounds I and II appears a characteristic absorption band of carbonyl group (C=O)

Table 1. Bond lengths (*d*, Å) in molecules I and II

Bond	<i>d</i> , Å (I)	<i>d</i> , Å (II)	Bond	<i>d</i> , Å (I)	<i>d</i> , Å (II)
N ¹ –C ⁶	1.370(3)	1.368(3)	C ¹¹ –N ¹²	1.467(3)	1.461(3)
N ¹ –C ²	1.403(3)	1.403(3)	N ¹² –C ¹⁴	1.349(3)	1.354(3)
N ¹ –C ¹⁰	1.489(3)	1.484(3)	N ¹² –C ¹³	1.458(3)	1.456(3)
C ² –O ²	1.240(3)	1.237(3)	C ¹⁴ –O ¹⁴	1.235(3)	1.218(3)
C ² –C ³	1.427(3)	1.424(4)	C ¹⁴ –C ¹⁵	1.474(3)	1.520(3)
C ³ –C ⁴	1.346(4)	1.337(4)	C ¹⁵ –C ¹⁶	1.290(4)	1.512(3)
C ⁴ –C ⁵	1.403(4)	1.404(3)	C ¹⁶ –N ¹⁷	–	1.464(3)
C ⁵ –C ⁶	1.361(3)	1.360(3)	N ¹⁷ –C ¹⁸	–	1.450(4)
C ⁶ –C ⁷	1.503(3)	1.507(3)	N ¹⁷ –C ²²	–	1.472(3)
C ⁷ –C ⁸	1.523(3)	1.522(3)	C ¹⁸ –C ¹⁹	–	1.505(4)
C ⁷ –C ¹³	1.535(3)	1.537(3)	C ¹⁹ –O ²⁰	–	1.430(3)
C ⁸ –C ⁹	1.514(3)	1.524(4)	O ²⁰ –C ²¹	–	1.406(4)
C ⁹ –C ¹⁰	1.527(3)	1.515(3)	C ²¹ –C ²²	–	1.502(4)
C ⁹ –C ¹¹	1.532(3)	1.528(3)			

Table 2. Torsion angles (ω , deg) in molecules **I** and **II**

Angle	ω (I)	ω (II)	Angle	ω (I)	ω (II)
C ⁶ N ¹ C ² O ²	176.64(19)	176.5(2)	C ⁸ C ⁹ C ¹¹ N ¹²	58.1(2)	57.8(3)
C ¹⁰ N ¹ C ² O ²	-1.2(3)	-1.1(3)	C ¹⁰ C ⁹ C ¹¹ N ¹²	-65.2(3)	-64.5(3)
C ⁶ N ¹ C ² C ³	-3.2(3)	-3.5(3)	C ⁹ C ¹¹ N ¹² C ¹⁴	122.0(2)	117.6(2)
C ¹⁰ N ¹ C ² C ³	178.99(18)	178.9(2)	C ⁹ C ¹¹ N ¹² C ¹³	-56.5(3)	-56.2(3)
O ² C ² C ³ C ⁴	-177.4(2)	-177.0(3)	C ¹⁴ N ¹² C ¹³ C ⁷	-121.1(2)	-116.8(2)
N ¹ C ² C ³ C ⁴	2.5(3)	3.0(3)	C ¹¹ N ¹² C ¹³ C ⁷	57.3(2)	56.5(2)
C ² C ³ C ⁴ C ⁵	-0.5(3)	-0.5(4)	C ⁶ C ⁷ C ¹³ N ¹²	62.8(2)	63.0(2)
C ³ C ⁴ C ⁵ C ⁶	-0.9(3)	-1.8(4)	C ⁸ C ⁷ C ¹³ N ¹²	-59.8(2)	-59.5(2)
C ⁴ C ⁵ C ⁶ N ¹	0.2(3)	1.4(3)	C ¹³ N ¹² C ¹⁴ O ¹⁴	-176.0(2)	177.8(2)
C ⁴ C ⁵ C ⁶ C ⁷	-176.82(19)	-172.6(2)	C ¹¹ N ¹² C ¹⁴ O ¹⁴	5.7(3)	5.0(4)
C ² N ¹ C ⁶ C ⁵	2.0(3)	1.4(3)	C ¹³ N ¹² C ¹⁴ C ¹⁵	4.8(3)	-2.4(3)
C ¹⁰ N ¹ C ⁶ C ⁵	179.57(18)	178.8(2)	C ¹¹ N ¹² C ¹⁴ C ¹⁵	-173.5(2)	-175.3(2)
C ² N ¹ C ⁶ C ⁷	179.05(16)	175.56(19)	O ¹⁴ C ¹⁴ C ¹⁵ C ¹⁶	4.6(4)	21.7(4)
C ¹⁰ N ¹ C ⁶ C ⁷	-3.3(3)	-7.0(3)	N ¹² C ¹⁴ C ¹⁵ C ¹⁶	-176.2(3)	-158.1(2)
C ⁵ C ⁶ C ⁷ C ⁸	-152.57(19)	-154.5(2)	C ¹⁴ C ¹⁵ C ¹⁶ N ¹⁷	-	-172.3(2)
N ¹ C ⁶ C ⁷ C ⁸	30.4(2)	31.4(2)	C ¹⁵ C ¹⁶ N ¹⁷ C ¹⁸	-	-63.0(3)
C ⁵ C ⁶ C ⁷ C ¹³	85.6(2)	83.9(2)	C ¹⁵ C ¹⁶ N ¹⁷ C ²²	-	175.8(2)
N ¹ C ⁶ C ⁷ C ¹³	-91.4(2)	-90.2(2)	C ¹⁶ N ¹⁷ C ¹⁸ C ¹⁹	-	-177.9(2)
C ⁶ C ⁷ C ⁸ C ⁹	-60.5(2)	-59.5(2)	C ²² N ¹⁷ C ¹⁸ C ¹⁹	-	-56.4(3)
C ¹³ C ⁷ C ⁸ C ⁹	60.9(2)	60.6(2)	N ¹⁷ C ¹⁸ C ¹⁹ O ²⁰	-	58.5(4)
C ⁷ C ⁸ C ⁹ C ¹⁰	64.8(2)	64.5(2)	C ¹⁸ C ¹⁹ O ²⁰ C ²¹	-	-58.0(4)
C ⁷ C ⁸ C ⁹ C ¹¹	-60.0(2)	-59.4(2)	C ¹⁹ O ²⁰ C ²¹ C ²²	-	57.9(3)
C ⁶ N ¹ C ¹⁰ C ⁹	7.6(3)	12.5(3)	C ¹⁸ N ¹⁷ C ²² C ²¹	-	55.9(3)
C ² N ¹ C ¹⁰ C ⁹	-174.56(16)	-169.88(18)	C ¹⁶ N ¹⁷ C ²² C ²¹	-	180.0(2)
C ⁸ C ⁹ C ¹⁰ N ¹	-38.9(2)	-41.6(3)	O ²⁰ C ²¹ C ²² N ¹⁷	-	-57.9(3)
C ¹¹ C ⁹ C ¹⁰ N ¹	84.7(2)	81.1(2)			

Table 3. Bond angles (ω , deg) in structures **I** and **II**

Angle	ω (I)	ω (II)	Angle	ω (I)	ω (II)
C ⁶ N ¹ C ²	122.68(2)	122.60(19)	N ¹ C ¹⁰ C ⁹	114.40(2)	115.0(2)
C ⁶ N ¹ C ¹⁰	123.17(2)	122.91(19)	N ¹² C ¹¹ C ⁹	110.23(2)	111.02(17)
C ² N ¹ C ¹⁰	114.12(2)	114.44(19)	C ¹⁴ N ¹² C ¹³	126.89(2)	126.79(18)
O ² C ² N ¹	119.50(2)	119.2(2)	C ¹⁴ N ¹² C ¹¹	120.26(2)	120.0(2)
O ² C ² C ³	125.3(2)	125.7(2)	C ¹³ N ¹² C ¹¹	112.84(2)	112.84(17)
N ¹ C ² C ³	115.2(2)	115.1(2)	N ¹² C ¹³ C ⁷	109.52(2)	109.73(19)
C ⁴ C ³ C ²	122.0(2)	122.1(2)	O ¹⁴ C ¹⁴ N ¹²	120.87(2)	121.2(2)
C ³ C ⁴ C ⁵	120.3(2)	120.5(2)	O ¹⁴ C ¹⁴ C ¹⁵	120.49(2)	121.6(2)
C ⁶ C ⁵ C ⁴	119.6(2)	119.4(2)	N ¹² C ¹⁴ C ¹⁵	118.63(2)	117.3(2)
C ⁵ C ⁶ N ¹	120.1(2)	120.1(2)	C ¹⁶ C ¹⁵ C ¹⁴	122.6(3)	110.91(19)
C ⁵ C ⁶ C ⁷	120.8(2)	120.9(2)	N ¹⁷ C ¹⁶ C ¹⁵	-	112.68(18)
N ¹ C ⁶ C ⁷	119.04(2)	118.68(19)	C ¹⁸ N ¹⁷ C ¹⁶	-	113.35(19)
C ⁶ C ⁷ C ⁸	111.35(2)	112.0(2)	C ¹⁸ N ¹⁷ C ²²	-	108.6(2)
C ⁶ C ⁷ C ¹³	109.35(2)	107.72(16)	C ¹⁶ N ¹⁷ C ²²	-	109.14(17)
C ⁸ C ⁷ C ¹³	110.03(2)	110.43(18)	N ¹⁷ C ¹⁸ C ¹⁹	-	110.7(2)
C ⁷ C ⁸ C ⁹	106.57(2)	106.37(17)	O ²⁰ C ¹⁹ C ¹⁸	-	111.3(2)
C ¹⁰ C ⁹ C ⁸	109.94(2)	109.51(19)	C ²¹ O ²⁰ C ¹⁹	-	109.8(2)
C ¹⁰ C ⁹ C ¹¹	112.55(2)	112.21(19)	O ²⁰ C ²¹ C ²²	-	111.9(2)
C ⁸ C ⁹ C ¹¹	110.4(2)	110.2(2)	N ¹⁷ C ²² C ²¹	-	110.6(2)

at 1670–1675 cm^{-1} . In the ^1H NMR spectrum of compound **II** the protons attached to the alkaloid skeleton give rise to signals in the characteristic regions of spectra [3], and methylene protons appear as resonances at 2.52–2.75 ppm

The structure of compound **II** was also proved by an independent preparation in reaction of *N*-acryloylcytisine (**I**) with morpholine in alcohol.

EXPERIMENTAL

IR spectra were recorded on spectrometer UR-20 from samples pelletized with KBr, ^1H NMR spectra were registered on VARIAN MERCURY-300 instrument, operating frequency 300 MHz internal reference HMDS.

***N*¹²-Acryloylcytisine (I)** was prepared by a standard procedure from the alkaloid and acryloyl chloride in the presence of triethylamine in anhydrous benzene in 95% yield, mp 130°C.

***N*¹²-(β -Morpholinopropionyl)cytisine (II)**. (a) To a solution of 1.44 g of *N*-acryloylcytisine and 0.38 g of carbon disulfide in ethanol was slowly added 0.435 g of morpholine dissolved in ethanol. The mixture was heated at stirring till the precipitate of morpholine salt with carbon disulfide completely dissolved. On distilling off the solvent 1.57 g (77%) of compound **II** was isolated.

(b) To a solution of 1.44 g of *N*-acryloylcytisine in ethanol was slowly added 0.435 g of morpholine dissolved in ethanol. The mixture was stirred at room temperature for 30–60 min. The solvent was distilled off to obtain 1.96 g (96%) of compound **II**, mp 66–67°C (from benzene). Found, %: C 65.31; H 7.57; N 12.76. $\text{C}_{18}\text{H}_{25}\text{N}_3\text{O}_3$. Calculated, %: C 65.23; H 7.60; N 12.68.

X-ray diffraction study. The unit cell parameters and intensity of 1262 independent reflections of compound **I** were measured on a diffractometer Bruker P4 (MoK_α radiation, graphite monochromator, $\theta/2\theta$ -scanning, $2\theta \leq 48^\circ$). Crystals monoclinic, a 8.3253(7), b 7.2838(5), c 11.3181(8) Å, β 105.512(5)°, V 661.33(9) Å³, d_{calc} 1.317 g/cm³, Z 2. $\text{C}_{16}\text{H}_{23}\text{N}_2\text{O}_3$, space group $P2_1$. In

calculations 1217 reflections were used with $I > 2\sigma(I)$. Extinction correction was done using ψ -curves. The final divergence factors were R 0.030 and wR 0.077.

The unit cell parameters and intensity of 1931 independent reflections of compound **II** were measured on a diffractometer Siemens P2 (CuK_α radiation, graphite monochromator, $\theta/2\theta$ -scanning, $2\theta \leq 70.03^\circ$). Crystals monoclinic, a 11.2792(18), b 7.0995(12), c 11.847(3) Å β 97.370(16)°, V 940.8(3) Å³, d_{calc} 1.297 g/cm³, Z 2. $\text{C}_{18}\text{H}_{29}\text{N}_3\text{O}_5$, space group $P2_1$. In calculations 1217 reflections were used with $I > 2\sigma(I)$. Extinction correction was done using ψ -curves. The final divergence factors were R 0.0394 and wR 0.104.

The structures were solved by the direct method and refined by full-matrix least-squares procedure in anisotropic approximation for nonhydrogen atoms with the use of software SHELXS-97 θ SHELXL-97. Hydrogen atoms were placed geometrically in the *rider* model.

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