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1 **Allylcytisine as a convenient scaffold for the construction of the  $\pi,\sigma$ -coordination compound**  
2 **{Acyt(H<sup>+</sup>)}[Cu<sub>8</sub>{Acyt(H<sup>+</sup>)}Cl<sub>10</sub>] with the unusual anionic 1D-coordination polymer**

3  
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10  
11 **Abstract**

12 The aim of this work is to develop a novel  $\pi$ -coordination compound with unusual architecture  
13 using allylcytisine (Acyt) as a suitable scaffold. The synthesis and structural characterization of  
14 {Acyt(H<sup>+</sup>)}[Cu<sub>8</sub>{Acyt(H<sup>+</sup>)}Cl<sub>10</sub>] (**1**) and Acyt itself have been performed, accompanied by quantum  
15 chemical studies. A distinctive feature of structure **1** is the formation of H-bonded pairs of two  
16 {Acyt(H<sup>+</sup>)} cations, showing the non-equivalent participation of its allyl group regarding to Cu<sup>+</sup>  
17 coordination, thus forcing the organization of the acentric structure **1** with the unusually organized  
18 anionic copper(I) halide 1D-coordination polymer.

19  
20 **Keywords:** Copper(I);  $\eta^2$ -Interaction; Cytisine; Allyl derivative; Crystal structure

21  
22 **1. Introduction**

23 Cytisine is well known alkaloid from plants of the Leguminosae family (especially the seeds of  
24 *Laburnum anagyroides*), which was introduced as a smoking cessation drug due to its high affinity  
25 for nicotinic acetylcholine receptors and acting as their partial agonist [1–3]. Despite significant  
26 advances in the comprehensive studies of the biological activity of cytisine and its derivatives [4–  
27 8], the use of these compounds as co-ligands in coordination chemistry of *d*-metals remains  
28 practically unexplored. According to the Cambridge Structural Database [9], there are only four  
29 coordination compounds of such type: in copper(II) and zinc(II) compounds N-protonated cytisine  
30 is bound to the central ion by the carbonyl O atom only [10], while in an organometallic  
31 coordination compound containing palladium(II) the neutral cytisine molecule, in its bonding with  
32 the Pd(II) cation, uses another center – the most nucleophilic N atom of the amine group [11].

33 To support Cu<sup>+</sup> binding with cytisine, a nitrogen atom N(12)-modification can be used for  
34 incorporating the complexing group. The appearance of the allylic substituent on this amine N atom

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35 may serve as an actual key for the selected coordination of transition metal ions due to the metal-  
36 olefin  $\pi$ -bonding [12], forcing a significant influence on cytosine coordination capabilities. As an  
37 example, several novel allylazole ligands in its  $\pi$ -complexation with copper(I) were studied earlier  
38 [13–15]. The proposed concept is effective in crystal engineering of the unique  $\pi,\sigma$ -coordination  
39 compounds with unknown (or less stable) in the free state  $\text{Cu}^{\text{I}}$  salts [16,17]. Some of these  
40 copper(I)-containing compounds turned out to be very promising in obtaining advanced laser  
41 operated materials [18–20]. N-Allyl derivatives of heterocycles were also found very favourable for  
42 the construction of coordination polymers of different dimensionality as well as the cluster  
43 compounds [21–26].

44 In this report, to insight how the incorporation of allyl substituent to cytosine makes it a  
45 convenient scaffold for the construction of novel  $\pi,\sigma$ -coordination compounds, we described the  
46 synthesis and structural characterization of  $\{\text{Acyt}(\text{H}^+)\}[\text{Cu}_8\{\text{Acyt}(\text{H}^+)\}\text{Cl}_{10}]$  (**1**)  $\pi,\sigma$ -coordination  
47 compound (Acyt – N12-Allylcytosine) with the unusual anionic cupro(I)chloride inorganic chain.  
48 Since the crystal structure of allylcytosine has not been studied earlier by single crystal X-ray  
49 diffraction, here we present a thorough structural analysis of *Acyt* itself.

50

## 51 **2. Experimental section**

### 52 *2.1 Materials and instrumentation*

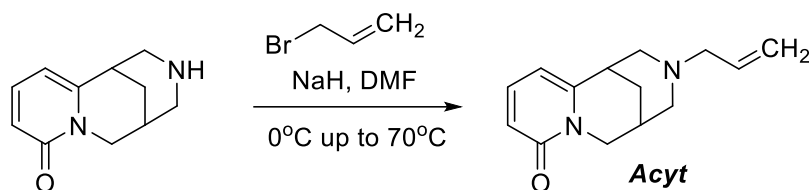
53 Unless mentioned otherwise, all chemicals were obtained from commercial sources and used  
54 without further purification. The NMR experiments:  $^1\text{H}$  NMR (500 MHz),  $^{13}\text{C}\{^1\text{H}\}$  NMR (125  
55 MHz) for allylcytosine *Acyt* were recorded on a Bruker Avance 500 MHz NMR spectrometer. The  
56 chemical shifts are reported in ppm relative to the residual peak of the deuterated DMSO for the  $^1\text{H}$   
57 and  $^{13}\text{C}\{^1\text{H}\}$  NMR spectra. Raman spectra from crystals were recorded with a Horiba Jobin–Yvon  
58 LabRAM HR spectrometer with the use of the 632.81 nm excitation line of a He-Ne laser (17 mW).  
59 Diffraction data for *Acyt* and coordination compound **1** were collected on an Agilent Gemini A  
60 four-circle diffractometer with Atlas CCD detector. Quantum chemical calculations of geometry  
61 optimization and simulation of vibrational (Raman) spectra of the *Acyt* and its coordination  
62 compound **1** have been performed using GAMESS(US) quantum chemistry package [27]. Energy  
63 framework calculations were performed on the DFT/B3LYP/6-31G(d, p) level using the  
64 CrystalExplorer 17.5 software [28].

65

### 66 *2.2 Preparation of the ligand*

67 N(12)-Allylcytosine (3-allyl-1,2,3,4,5,6-hexahydro-8*H*-1,5-methanopyrido[1,2-*a*][1,5]diazocin-  
68 8-one, *Acyt*) was prepared via direct alkylation of cytosine with allyl bromide (3-bromoprop-1-ene)  
69 (Scheme 1). To a solution of cytosine 1.9 g (0.01 mol) in DMF (50 mL) cooled to 0 °C, NaH 0.48 g

70 (0.012 mol) was added in portions and stirred at 0 °C until gas evolution. Allyl bromide 1.3 mL  
 71 (0.015 mol) was added in small portions to the reaction mixture and left at room temperature for 12  
 72 hours. Then the reaction mixture was heated for 1 h at a temperature of 70 °C. The mixture was  
 73 cooled to room temperature. Water (50 mL) was added and the DMF-water azeotrope was removed  
 74 under reduced pressure. The concentrate was diluted with water (10 mL) and extracted with ethyl  
 75 acetate (3 × 10 mL). The organic fraction was dried over Na<sub>2</sub>SO<sub>4</sub>, and the solvent was evaporated  
 76 under reduced pressure. To the resulting residue, hexane (10 mL) was added and boiled for 5 min.  
 77 The mixture was cooled, and allylcytisine was filtered off as white crystals. Yield 1.8 g, 78%; mp =  
 78 113 °C. <sup>1</sup>H NMR (500 MHz, DMSO-*d*<sub>6</sub>) δ 7.29 (dd, *J* = 8.6, 7.2 Hz, 1H), 6.18 (d, *J* = 8.9 Hz, 1H),  
 79 6.04 (d, *J* = 6.6 Hz, 1H), 5.64 – 5.50 (m, 1H), 5.00 (d, *J* = 5.0 Hz, 1H), 4.98 (d, *J* = 15.4 Hz, 1H),  
 80 3.75 (d, *J* = 15.3 Hz, 1H), 3.68 (dd, *J* = 15.3, 6.4 Hz, 1H), 2.98 (s, 1H), 2.89 – 2.77 (m, 4H), 2.36 (s,  
 81 1H), 2.18 (t, *J* = 11.6 Hz, 2H), 1.77 (d, *J* = 12.5 Hz, 1H), 1.65 (d, *J* = 12.6 Hz, 1H); <sup>13</sup>C NMR (126  
 82 MHz, DMSO-*d*<sub>6</sub>) δ 162.19, 152.13, 138.77, 135.04, 117.01, 115.30, 103.72, 60.19, 59.91, 59.51,  
 83 49.59, 34.52, 27.32, 25.19; MS (*m/z*, APCI) = 231 (*M*<sup>+</sup> + 1); Anal. calcd for C<sub>14</sub>H<sub>18</sub>N<sub>2</sub>O: C, 73.01;  
 84 H, 7.88; N, 12.16. Found: C, 73.14; H, 7.79; N, 12.07.



85  
86 Scheme 1. The ligand Acyt synthesis

87  
88 **2.3 Synthesis of the {Acyt(H<sup>+</sup>)}[Cu<sub>8</sub>{Acyt(H<sup>+</sup>)}Cl<sub>10</sub>] (**1**)**

89 Crystals of  $\pi,\sigma$ -coordination compound **1** were obtained under the conditions of the alternating-  
 90 current electrochemical technique starting from the ethanol solution of Acyt and copper(II) chloride.  
 91 A mixture of Acyt (0.66 mmol, 0.152 g) and CuCl<sub>2</sub>·2H<sub>2</sub>O (1.76 mmol, 0.300 g) in 5.0 mL of  
 92 ethanol, acidified by two drops of 37% hydrochloric acid, was prepared and then was placed into a  
 93 small 5.5 mL test-tube. Copper-wire electrodes in cork were inserted to this reactor. The alternating  
 94 current (frequency 50 Hz) of 0.63 V was applied for one month, and after that the reactor was  
 95 moved to the refrigerator, where it was kept at 10-11°C. After two months good quality colourless  
 96 crystals of compound **1** appeared on copper-wire electrodes in a very small amount under a thin  
 97 layer of white amorphous substance. M.p. 155°C.

98  
99 **2.3. Single crystal X-ray diffraction studies**

100 The collected diffraction data for Acyt and coordination compound **1** were processed with the  
 101 CrysAlis PRO program [29]. Both structures were solved by ShelXT and refined by least squares

102 method on  $F^2$  using ShelXL software with the following graphical user interface of OLEX<sup>2</sup> [30–  
 103 32]. The atomic displacement parameters for non-hydrogen atoms were refined using an anisotropic  
 104 model. The amine H atoms of the *Acyt*(H<sup>+</sup>) moieties in coordination compound **1** were derived from  
 105 difference Fourier maps and refined with  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{N})$ . The other H atoms in both *Acyt* and  
 106 **1** structures were refined in ideal positions (riding model), with C—H = 0.99 (methylene) or 0.95 Å  
 107 (otherwise) and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ . Crystal parameters, data collection and refinement details  
 108 are summarized in Table 1. The figures were prepared using the DIAMOND 3.1 software.

109

110 **Table 1.** Selected crystal data and structure refinement parameters of *Acyt* and compound **1**.

Crystal data	<i>Acyt</i>	<b>1</b>
CCDC number	2173890	2173901
Empirical formula	C <sub>14</sub> H <sub>18</sub> N <sub>2</sub> O	C <sub>28</sub> H <sub>38</sub> Cl <sub>10</sub> Cu <sub>8</sub> N <sub>4</sub> O <sub>2</sub>
F. w. (g·mol <sup>-1</sup> )	230.30	1325.44
Crystal system, space group	orthorhombic, <i>P</i> 2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>	triclinic, <i>P</i> 1
<i>a</i> (Å), $\alpha$ (°)	9.4309(4)	7.4509(3), 84.010(4)
<i>b</i> (Å), $\beta$ (°)	10.7697(5)	10.7420(5), 80.982(4)
<i>c</i> (Å), $\gamma$ (°)	11.9283(6)	12.9869(6), 81.270(4)
<i>V</i> (Å <sup>3</sup> )	1211.53(10)	1011.24(8)
<i>Z</i>	4	1
$\mu$ (mm <sup>-1</sup> )	0.634	4.817
<i>F</i> (000)	496	652
Crystal size (mm)	0.06 × 0.21 × 0.33	0.26 × 0.35 × 0.46
Crystal color	colourless	colourless
Calculated density, g/cm <sup>3</sup>	1.263	2.176
<i>Data collection</i>		
Radiation type, wavelength, $\lambda$ (Å)	Cu <i>Ka</i> , 1.54184	Mo <i>Ka</i> , 0.71073
Temperature, <i>K</i>	150	150
Used in refinement reflections	2325	8664
$R[F^2 > 2\sigma(F^2)]$	0.0380	0.0375
$wR(F^2)$	0.0930	0.0852
GooF = <i>S</i>	1.013	1.083
Flack	0.3(5)	-0.030(13)
$\Delta\rho_{\text{max}}/\Delta\rho_{\text{min}}$ (e Å <sup>-3</sup> )	0.160 /-0.154	0.871/-0.792

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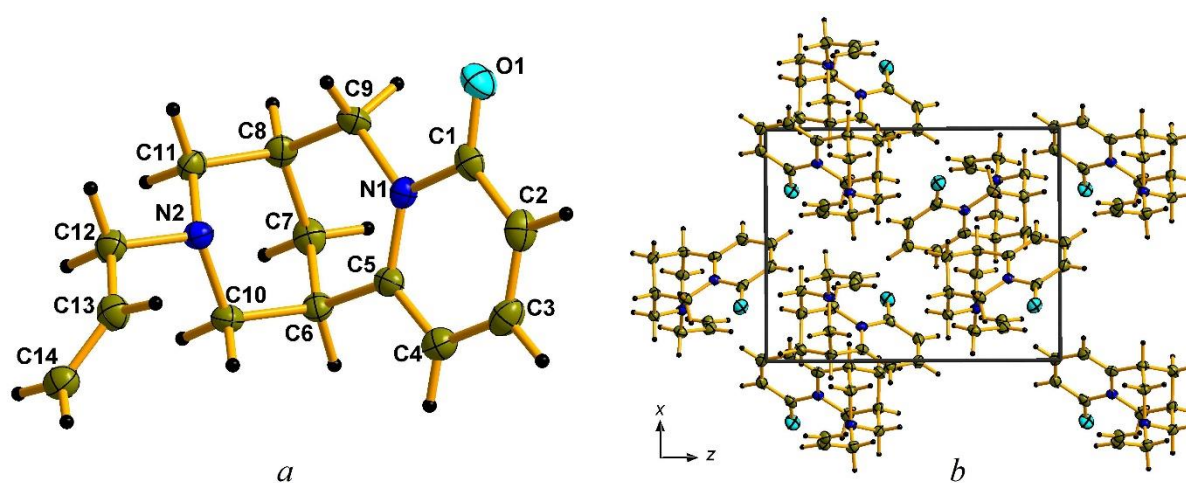
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115 **3. Results and discussion**116 **3.1. Crystal structure**

117 *Acyt* crystallizes in the acentric space group  $P2_12_12_1$ , with one molecule in the asymmetric unit.  
 118 Thus *Acyt* differs from earlier investigated structures of unsubstituted cytosine [33] and its  
 119 benzothienopyrimidinone derivative [34], each of which has two crystallographically independent  
 120 molecules. The geometry of the cytosine fragment in *Acyt* is comparable to that of unsubstituted  
 121 cytosine (Fig. 1, Table 2). The allylamino group has an anticlinal conformation relative to the C12—  
 122 C13 bond (N2—C12—C13—C14,  $-124.9(3)^\circ$ ). Only weak C—H $\cdots$ O bindings are observed among  
 123 O1 center and H-donor C4, C6 & C8 atoms of nearest *Acyt* molecules.



124  
 125 **Fig. 1.** *a*) The molecular structure of *Acyt* with displacement ellipsoids drawn at the 50% probability level. *b*) A view  
 126 along the *b* axis of the crystal packing of the *Acyt*.

127  
 128  
 129 **Table 2.** Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ) of *Acyt* and **1**.

<i>Acyt</i>			
O1—C1	1.239(3)	C10—N2—C12	110.6(2)
C1—N1	1.411(3)	N2—C12—C13	112.2(2)
C13—C14	1.315(4)	C12—C13—C14	124.5(3)
Coordination compound <b>1</b>			
Cu1—Cl1	2.289(2)	Cl1—Cu1— <i>m</i>	123.4(2)
Cu1—Cl2	2.289(2)	Cl2—Cu1— <i>m</i>	121.8(2)
Cu1—Cl3	2.749(2)	Cl3—Cu1— <i>m</i>	101.6(2)
Cu1— <i>m</i>	1.963(7)	Cl1—Cu1—Cl3	102.68(7)
C13A—C14A	1.364(10)	Cl1—Cu2—Cl4	169.95(8)
Cu2—Cl1	2.133(2)	Cl3—Cu3—Cl5	158.91(9)
Cu3—Cl2	2.587(2)	Cl3—Cu3—Cl2	99.13(7)
Cu3—Cl3	2.194(2)	Cl5—Cu4—Cl2	98.18(7)
Cu4—Cl2	2.615(2)	Cl5—Cu4—Cl8 <sup>ii</sup>	119.62(7)
Cu5—Cl2	2.769(2)	Cl6—Cu5—Cl7	162.50(8)
Cu6—Cl7	2.288(2)	Cl2—Cu5—Cl6	95.51(7)
Cu7—Cl4 <sup>i</sup>	2.436(2)	Cl7—Cu6—Cl8	118.64(8)

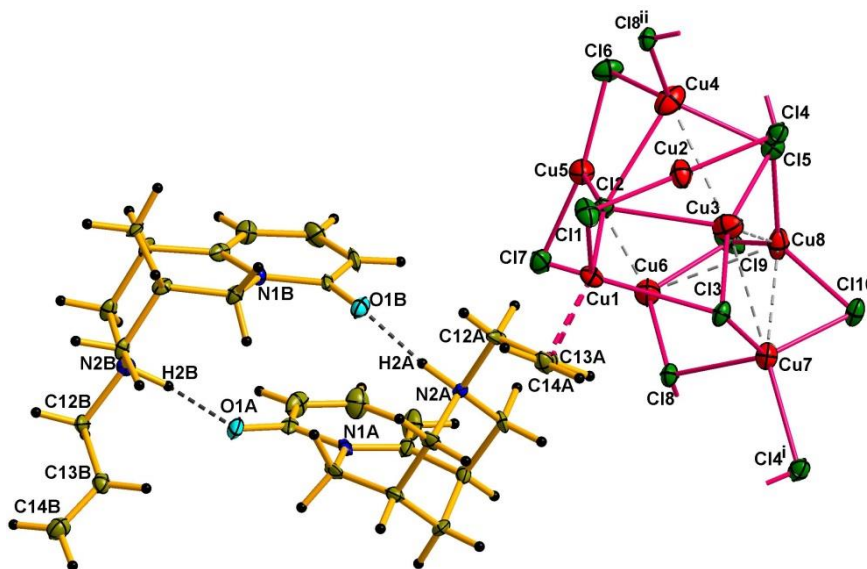
Cu7—C110	2.274(2)	Cl8—Cu6—Cl9	119.87(8)
Cu3—Cu4	2.9806(15)	Cl3—Cu7—Cl10	124.37(8)
Cu5—Cu6	2.8318(14)	Cl8—Cu7—Cl10	117.77(7)
Cu6—Cu8	2.9894(14)	Cl5—Cu8—Cl9	118.55(8)
Cu3—Cu7	2.8512(14)	Cu3—Cu8—Cu6	90.09(4)
Cu3—Cu8	2.8152(14)	Cu6—Cu8—Cu7	82.17(4)
Cu7—Cu8	2.9294(14)	Cu5—Cu6—Cu8	94.80(4)

130 *m* – the mid-point of C13A=C14A bond.

131 Symmetry codes: (i) *x*+1, *y*, *z*; (ii) *x*-1, *y*, *z*.

132

133 Coordination compound **1** crystallizes in the acentric space group *P*1, with two *Acyt*(H<sup>+</sup>)  
 134 cations, eight Cu(I) and ten Cl ions per unit cell. One of the organic cations is  $\pi$ -coordinated to  
 135 Cu(1) through allylic C=C bond, while the carbonyl group and the protonated amino group of this  
 136 *Acyt*(H<sup>+</sup>) are effectively involved in a symmetric N—H $\cdots$ O bonding with the nearest organic cation  
 137 (Fig. 2, Table 3). In addition, pyrimidinone planes of the cations are also involved into  $\pi$ ... $\pi$ -  
 138 stacking (centroid – centroid distance is  $\sim$ 3.3 Å). These cations are two conformers, which arise  
 139 from the rotation of the N-allyl substituent relative to the N2—C12 bond by 102.6(8)°. The  $\pi$ -  
 140 Bonded allylamino group has an anticlinal conformation relative to the C12A—C13A bond (N2A—  
 141 C12A—C13A—C14A,  $-106.7(8)^\circ$ ), while the allylamino group of the nearest cation is  
 142 characterized by an antiperiplanar conformation regarding to the analogous C12B—C13B bond  
 143 (N2B—C12B—C13B—C14B,  $-150.7(7)^\circ$ ). Such extensive cationic pairs {*Acyt*(H<sup>+</sup>)...*Acyt*(H<sup>+</sup>)}  
 144 contribute to the formation of an unusual anionic chain.



145

146 **Fig. 2.** The independent part in crystal structure of **1**. Displacement ellipsoids are drawn at the 50% probability

147

level. Symmetry codes: (i) *x*+1, *y*, *z*; (ii) *x*-1, *y*, *z*.

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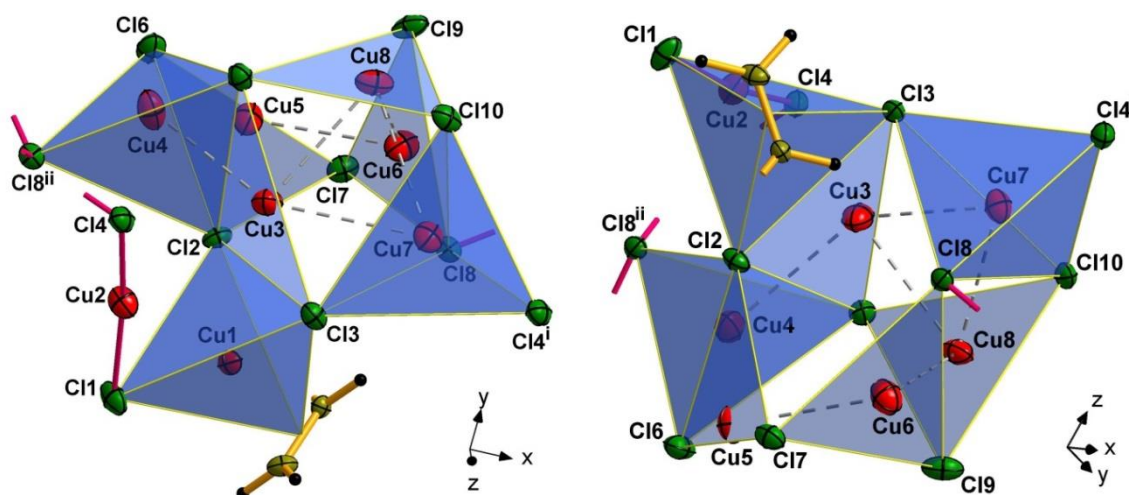


152 **Table 3.** Hydrogen-bond geometry (Å, °) for **1**.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C11B—H11D $\cdots$ Cl4 <sup>i</sup>	0.99	2.83	3.709 (7)	149
N2A—H2A $\cdots$ O1B	0.97 (10)	1.71 (10)	2.667 (7)	170 (8)
N2B—H2B $\cdots$ O1A	0.96 (9)	1.71 (9)	2.652 (8)	168 (8)

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

154 The  $\pi$ -coordinated Cu(1) cation adopts a nearly trigonal pyramidal coordination environment  
 155 (3Cl, (C=C)) (Table 2). The corresponding four-coordinate geometry index  $\tau_4$  [35] for Cu(1) is 0.81.  
 156 The basal plane of the coordination arrangement consists of  $\mu_2$ -Cl(1),  $\mu_4$ -Cl(2) ions and the  $\eta^2$ -allyl  
 157 group. The axial site is occupied by a  $\mu_3$ -Cl(3) ion at 2.749(2) Å. The  $\pi$ -connected to the metal  
 158 centre C13A=C14A bond is elongated (due to back-donation from an occupied 3d metal orbital to a  
 159 low-lying empty  $\pi^*$ -orbital of the olefin) to 1.364(10) Å in comparison with uncoordinated allylic  
 160 C13=C14 bond (Table 2, Fig. 3). Similarly, the Cu(4) ion has a nearly trigonal pyramidal  
 161 coordination environment (4Cl,  $\tau_4 = 0.85$ ) with the axial  $\mu_4$ -Cl(2) ion at 2.615(2) Å.  
 162

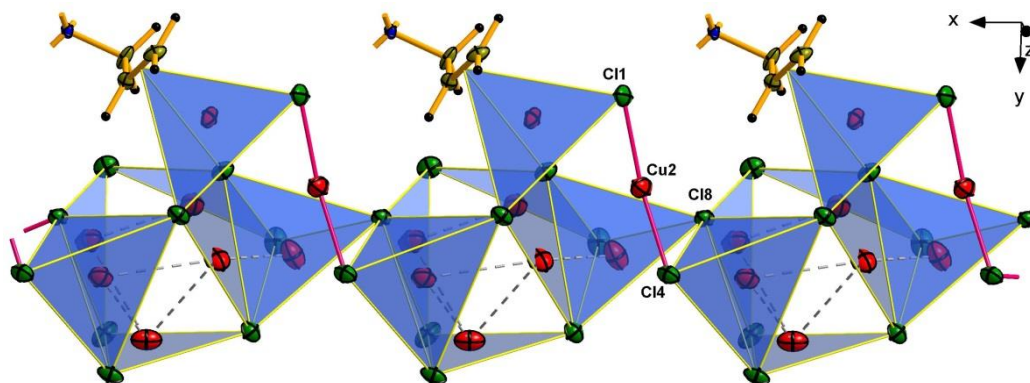


163 **Fig. 3.** A part of inorganic chain in the compound **1** at two different viewing directions and with the depicted Cu  
 164 polyhedra. Symmetry codes: (i)  $x+1, y, z$ ; (ii)  $x-1, y, z$ .  
 165  
 166

167 In the case of Cu(7), its coordination environment ( $\tau_4 = 0.84$ ) is formed by two  $\mu_2$ -Cl and two  
 168  $\mu_3$ -Cl ions, allowing Cu7 to participate in metalophilic interactions with neighboring Cu(3) and Cu(8)  
 169 atoms (Cu(7)...Cu(3) 2.851(1) Å, Cu(7)...Cu(8) 2.929(1) Å). The last values are somewhat higher  
 170 than the sum of the corresponding copper VdW radii (2.80 Å) published by Bondi [36] but still  
 171 much smaller than the corresponding VdW radii sum obtained by Batsanov or Alvarez [37,38]. As a  
 172 result, Cu(3) adopts a close to T-shaped arrangement including two closely attached to it  $\mu_3$ -Cl(3) &  
 173  $\mu_3$ -Cl(5) ions at 2.194(2) Å and 2.211(2) Å, correspondingly. Similar T-shaped arrangement has  
 174 Cu(5), but the closest anions in its environment are  $\mu_2$ -Cl(6) &  $\mu_2$ -Cl(7) ions, while the third one,  $\mu_4$ -  
 175 Cl(2) ion is away from this metal center at 2.769(2) Å. Another closest Cu(6) and Cu(8) possess a

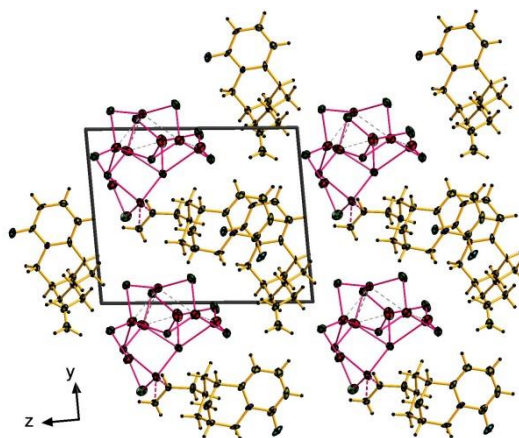


176 trigonal coordination environment. Thus, the formed clusters of Cu(3) – Cu(8) atoms defined within  
 177 the anionic inorganic chain produce the specific  $\{Cu_7\{Acyt(H^+)\}Cl_{10}\}$  subunits (Fig. 4), which are  
 178 linked by linearly-arranged Cu(2) cations and bridging  $\mu_3$ -Cl(8) anions into 1D-polymer, extended  
 179 along the *a* axis.



180  
 181 **Fig. 4.** Anionic inorganic chain in the compound **1** with the depicted Cu(I) polyhedra.  
 182

183 Most probably, the asymmetry of the above subunit and of structure **1** as a whole deals with the  
 184 passivation of the allyl group of the second cation in metal binding, which in turn is enforced by  
 185 H-bond-defined location of both extensive cations (Fig. 5). This gives reason to say that the  
 186 hydrogen bond between these cations competes effectively with Cu–(C=C) interaction specifying  
 187 the unusual architecture of the inorganic fragment, which was impossible in the presence of  
 188 allylazoles with a significant number of donor heteroatoms [12,14].

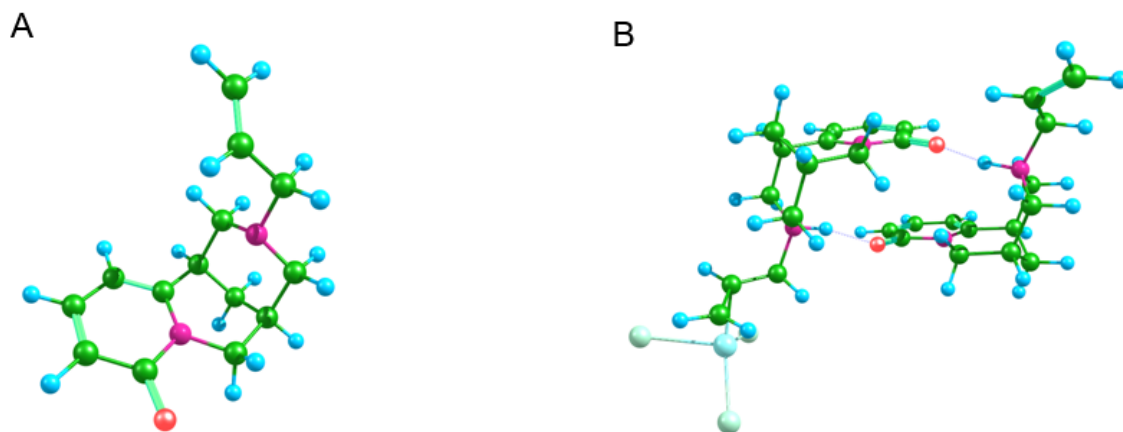


189  
 190 **Fig. 5.** A view along the *a* axis of the crystal packing of the compound **1**.  
 191

### 192 3.2. Raman spectra assignments

193 To simulate vibrational (Raman) spectra of the *Acyt* and its coordination compound  
 194  $\{Acyt(H^+)\}[Cu_8\{Acyt(H^+)\}Cl_{10}]$  we first have performed optimization of the bare *Acyt* (Figure 6A)  
 195 and structural unit, extracted from the crystal, consisting of the two protonated *Acyt* molecules one  
 196 of which forms  $\pi$ -coordination compound with copper(I) and 3 chloride anions  
 197 ( $\{Acyt(H^+)\}[Cu(I)\{Acyt(H^+)\}Cl_3]$ ) shown of Figure 6. Optimizations have been performed at the

198 DFT level of theory using PBE0 [39] functional and aug-cc-pVDZ basis set [40] with effective core  
 199 potential on Cu atom [41]. To preserve coordination from the crystal positions one Cu and three Cl  
 200 atoms have been constrained during the compound optimization. After that, vibrational frequencies  
 201 and corresponding Raman activities have been calculated at the optimized geometries using the  
 202 same level of theory (PBE0/aug-cc-pVDZ).



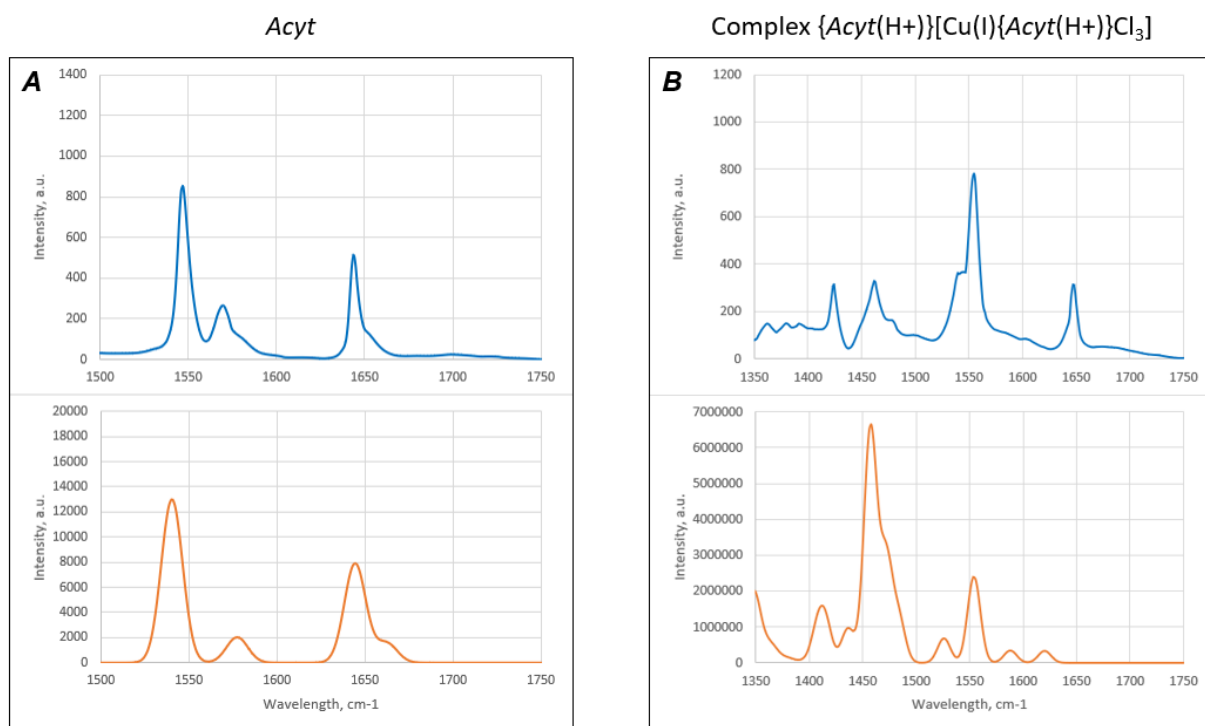
203  
 204 **Fig. 6.** Geometries of *Acyt* (panel A) and coordination compound  $\{Acyt(H^+)\}[Cu(I)\{Acyt(H^+)\}Cl_3]$  (panel B) optimized  
 205 at PBE0/aug-cc-pVDZ level of theory.  
 206

207 To provide band assignments of the experimentally measured Raman spectra of the crystal,  
 208 additional scaling coefficient of 0.95 have been applied to the calculated frequencies, and Raman  
 209 activities have been recalculated into intensities using formalism described in the work by  
 210 Michalska et.al. [42] with experimental conditions of 298K temperature and 632.81 nm (15802.53  
 211  $cm^{-1}$ ) excitation laser wavelength. In addition, spectra have been broadened with gaussian functions  
 212 of 20  $cm^{-1}$  half-width. Results of calculations and comparison between experimental and calculated  
 213 spectra are shown of Figure 7A for *Acyt* molecule and Figure 7B for  $\pi$ -coordination compound with  
 214 copper(I). Assignments of individual bands are summarized in Table 4.

215 As we could conclude on the basis of bare *Acyt* molecule calculations our level of theory is able  
 216 to reproduce Raman spectra of studied system quite well. Symmetric vibration of the allyl C=C  
 217 bond are estimated to be around 1644  $cm^{-1}$ . Shoulder, observed around 1652  $cm^{-1}$  in both measured  
 218 and calculated spectra corresponds to C=O vibration coupled with the symmetric ring mode. Both  
 219 spectral lines at 1570 and 1545  $cm^{-1}$  are also represent vibrational modes of the aromatic ring  
 220 system.

221 In the large complex simulation results we could assign the same vibrational modes of the ring  
 222  $\pi$ -system that are coupled to the C=O vibrations around 1525 and 1590  $cm^{-1}$ . It should be noted that  
 223 C=O vibration frequency is shifted to the lower energies (1525  $cm^{-1}$ ) in comparison to the bare *Acyt*  
 224 molecule (1652  $cm^{-1}$ ), which is due to intermolecular hydrogen bonds, that present between two

225 *Acyt* molecules in the coordination compound. Moreover, we also assign vibrational bands between  
 226 1400 and 1500  $\text{cm}^{-1}$  in the measured spectra to the bending modes of that hydrogen bonds, coupled  
 227 to the ring modes. According to the simulations allyl C=C bond vibration, that forms coordination  
 228 compound with Cu(I) should be also located in that region, giving us a large shift of  $\sim 180 \text{ cm}^{-1}$   
 229 comparison to the free C=C bond ( $1620 \text{ cm}^{-1}$ ) of the second *Acyt* molecule. It appears that due to  
 230 strong overlap with other bands we could observe such  $\pi$ -complexed C=C bond vibration only as a  
 231 shoulder of the peak in experimental spectra at  $\sim 1450 \text{ cm}^{-1}$ .



232  
 233 **Fig. 7.** Comparison of measured (upper panels) and calculated (lower panels) Raman spectra in a selected ranges of for  
 234 bare *Acyt* molecule (panel A) and  $\{Acyt(H^+)\}[Cu(I)\{Acyt(H^+)\}Cl_3]$  (panel B) coordination compound.

237 **Table 4.** Assignments of individual bands in the experimental Raman spectra of *Acyt* molecule and  
 238  $\{Acyt(H^+)\}[Cu_8\{Acyt(H^+)\}Cl_{10}]$  coordination compound.

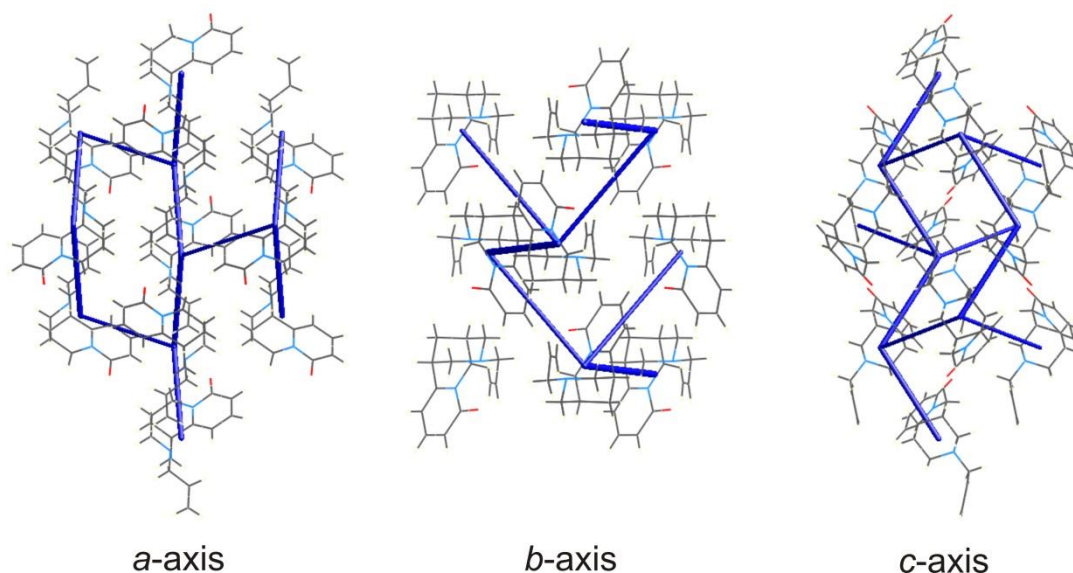
Experimental spectra band, $\text{cm}^{-1}$	Assignment
<i>Acyt</i>	
1546	Ring $\pi$ -system symmetric mode
1571, 1581 (shoulder)	Ring $\pi$ -system mode coupled to C=O vibration
1644	C=C bond mode
1654 (shoulder)	C=O mode
$\{Acyt(H^+)\}[Cu_8\{Acyt(H^+)\}Cl_{10}]$	
1350-1400	$CH_2$ bending modes
1424, 1460, 1478 (shoulder)	Ring modes coupled to C=O and H-bonds bending
1450 (shoulder)	C=C $\pi$ -bond to Cu(I)
1540 (shoulder), 1555	Ring $\pi$ -system mode coupled to C=O vibration
1648	Free C=C bond mode

239

240

## 241 3.3. Energy framework calculations

242 Additionally we have performed energy frameworks computational analysis [14,28] for *Acyt*  
 243 itself. All the calculations were provided for clusters of *Acyt* molecules within a radius of 3.8 Å,  
 244 which were generated around a single fragment. The cylinders in the energy framework represent  
 245 the relative strengths of molecular packing in different directions – interaction energies are  
 246 proportional to the thickness of cylinders joining the centroids of fragments (Fig. 8).



247 **a-axis** **b-axis** **c-axis**  
 248 **Fig. 8.** Comparison energy frameworks of *Acyt* representing the total interaction energy ( $\leq -20$  kJ/mol, blue-colored)  
 249 along different crystallographic axes.

250  
 251 According to the calculation results the main intermolecular interactions which make the most  
 252 contribution to the structure stabilization corresponds to the C—H $\cdots$ O hydrogen bonding between  
 253 cytosine C=O group and hydrogen atoms at C4 & C6 atoms of nearest *Acyt* molecules, covering the  
 254 total energy of  $-23.7$  kJ/mol and  $-33.8$  kJ/mol, correspondingly, with predominance of dispersive  
 255 forces. Both C12—H2B $\cdots$ O1 bond & C11—H11 $\cdots$  $\pi$  interaction with allyl group of the same  
 256 neighboring *Acyt* molecules cover the total energy of  $-19.3$  kJ/mol and also with the predominance  
 257 of dispersive forces. Next step in descending order of energy values occupy weaker C—H $\cdots$ O  
 258 contact of  $-18.8$  kJ/mol with C8 attached hydrogen atom, for which electrostatic and dispersion  
 259 components make approximately the same contribution. The total energy of all interaction in *Acyt*  
 260 structure equals  $-127.4$  kJ/mol.

261  
 262 **4. Conclusions**

263 To sum up, in our work we presented how allylcytisine was used as an instrument for the  
 264 construction of a novel  $\pi,\sigma$ -coordination compound  $\{Acyt(H^+)\}[Cu_8\{Acyt(H^+)\}Cl_{10}]$  (**1**) with an  
 265 acentric structure. We succeeded to obtain crystals of **1** using the alternating-current

266 electrochemical technique and study by single crystal X-ray diffraction method. Within the unusual  
267 anionic chain in **1** there are eight crystallographically independent copper(I) ions, which form the  
268 specific  $\{\text{Cu}_7\{\text{Acyl}(\text{H}^+)\}\text{Cl}_{10}\}$  subunits, linked by linearly-arranged  $\text{Cu}^+$  cations and bridging  $\mu_3\text{-Cl}$   
269 anions into 1D-polymer. The design feature of the subunit is the presence of cuprophilic  
270 interactions defined by geometrically distinct coordination environments of  $\sigma$ -bonded copper(I) ions  
271 that possess distorted tetrahedral, closed to trigonal pyramidal, trigonal, and T-shaped  
272 arrangements. The  $\pi$ -coordination of the only one of two different  $\text{Acyl}(\text{H}^+)$  cation to copper(I) of  
273 the anionic subunit is crucial for its formation since efficient  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds define the  
274 location of both extensive cations and enable second allyl group of second cation to metal binding.  
275 This gives reason to the assumption that the hydrogen bonding between these cations competes  
276 effectively with the  $\text{Cu}-(\text{C}=\text{C})$  interaction, specifying the unusual architecture of the inorganic  
277 fragment discussed.

278

#### 279 *CRedit Authorship Contribution Statement*

280 **Yurii Slyvka:** Conceptualization, Investigation, Methodology, Project administration,  
281 Software, Writing - original draft. **Evgeny Goresnik:** Software, Investigation, Validation, Writing  
282 - original draft. **Nazariy Pokhodylo:** Software, Synthesis, Investigation, Writing - original draft.  
283 **Dmitry Morozov:** Software, Investigation, Writing - original draft. **Mykola Tupychak:** Synthesis,  
284 Investigation. **Marian Mys'kiv:** Software, Investigation.

285

#### 286 **Declaration of Competing Interest**

287 The authors declare that they have no known competing financial interests or personal  
288 relationships that could have appeared to influence the work reported in this paper.

289

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294

#### 295 **Appendix A. Supplementary data**

296 CCDC 2173890 and 2173901 contain the supplementary crystallographic data for this paper.  
297 These data can be obtained free of charge from The Cambridge Crystallographic Data Centre via  
298 [www.ccdc.cam.ac.uk/structures](http://www.ccdc.cam.ac.uk/structures).

299

300

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