

STRUCTURAL STUDIES OF THE BORON COORDINATION COMPOUND WITH THIOUREA

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In the course of systematic structural studies of dicitratoborates the investigation of the crystal structure of $\text{NH}_2\text{CSHNH}_2[(\text{C}_6\text{H}_6\text{O}_7)_2\text{B}]$ has been carried out. The dicitratoborate complex with thiourea (L) was synthesized according to [1]. The structure is formed by cations of thiourea (HL^+) and spiran-type complex anions $[(\text{C}_6\text{H}_6\text{O}_7)_2\text{B}]^-$. In the complex anions two citric acid residues are coordinated bidentately to the BO_4 -tetrahedron. The values of the chemical bond lengths and bond angles in the complex anions are in accordance with those determined earlier for the crystal structures of dicitratoborates. The boron-containing five-membered rings are approximately planar. The cation $(\text{NH}_2\text{CSHNH}_2)^+$ is protonated at the sulphur atom. The protonation gives rise to delocalization of p-electron density at the C-S and N-C bonds. The bond lengths are: C-S 1.738(2) Å; N-C 1.317(3) Å and 1.298(3) Å. The bond $\text{C}(\text{sp}^2)\text{-S}$ is lengthened for 0.10 Å, but the $\text{C}(\text{sp}^2)\text{-N}$ bonds are shortened for 0.04 Å in average in the cation (HL^+) when compared with the neutral molecule (L). In the crystals each organic cation is bonded with three nonidentical complex anions by five hydrogen bonds $\text{N-H}\cdots\text{O}$ (2+2) and $\text{S-H}\cdots\text{O}$. The complex anions are bonded directly by four strong hydrogen bonds $\text{O-H}\cdots\text{O}$ (length from 2.589 to 2.692 Å). Crystal data: $\text{NH}_2\text{CSHNH}_2[(\text{C}_6\text{H}_6\text{O}_7)_2\text{B}]$, triclinic, space group *P*-1, *a* = 9.630(1) Å, *b* = 10.320(2) Å, *c* = 10.914(2) Å; α = 76.86(1)°, β = 88.80(1)°, γ = 64.26(1)°; *V* = 947.7(3) Å³; *Z* = 2; *d*_x = 1.641 g cm⁻³; *R* = 0.0246, *wR*₂ = 0.0661.

[1] G.Sergeeva, N.Burnashova, J.Schwartz. Compounds of borodicitric acid with urea and thiourea, Latvian Journal of Chemistry, (1991), 5: 553-558.

Keywords: CRYSTAL STRUCTURE, BORON COMPLEXES, THIOUREA

BIOLOGICALLY ACTIVE COPPER(II) AND PLATINUM(IV) COMPLEXES WITH CYTOKININ-DERIVED COMPOUNDS

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Biologically active mononuclear Cu(II) and Pt(IV) complexes have been synthesized and structurally characterized in the course of our systematic investigations of cytokinin-derived compounds [1,2]. The composition of the complexes, resulting from a single crystal X-ray analysis, is following: $[\text{Cu}(\text{2OHbapH})_2\text{Cl}_3]\text{Cl}\cdot 2\text{H}_2\text{O}$ (1) (2OHbap = 6-(2-hydroxybenzylamino)purine), $[\text{Cu}(\text{3OHbapH})_2\text{Cl}_3]\text{Cl}\cdot 2\text{H}_2\text{O}$ (2) (3OHbap = 6-(3-hydroxybenzylamino)purine), $[\text{Cu}(\text{4FbapH})_2\text{Cl}_2(\text{H}_2\text{O})]\text{Cl}_2$ (3) (4Fbap = 6-(4-fluorobenzylamino)purine) and $(\text{roscH}_2)_2[\text{PtCl}_4]\text{Cl}_2\cdot 4\text{H}_2\text{O}$ (4) (rosc = 2-(1-ethyl-2-hydroxyethylamino)-6-benzylamino-9-isopropylpurine). A coordination geometry is trigonal-bipyramidal in all the copper(II) complexes. Each of two protonated purine-derived ligands is bonded to the copper(II) ion via the N9 atom. The compound (4) consists of two twice-protonated rosc cations, one $[\text{PtCl}_4]^{2-}$, two Cl⁻ anions and four uncoordinated water molecules. The compounds have been tested for their possible cytotoxic activity against G-361, HOS, K-562 and MCF7 cell lines. A cytokinin activity has been also determined for the copper(II) complexes. X-ray data were collected at 120 K on a four-circle kappa-axis diffractometer KUMA KM-4 equipped with an Oxford Cryostream cooler and a CCD detector. All crystal structures were determined and refined using a SHELXL97 program package. Crystal data: (1) $\text{C}_{24}\text{H}_{26}\text{N}_{10}\text{O}_4\text{Cl}_4\text{Cu}$, monoclinic *P*₂/*n*, *a* = 14.8725(6), *b* = 8.3546(4), *c* = 23.3856(9) Å, β = 95.079(4)°, *Z* = 4, reflections collected / unique 14673 / 5069 [*R*(int) = 0.0501], *R*₁(obs) = 0.0434, *wR*₂(obs) = 0.0937. (2) $\text{C}_{24}\text{H}_{26}\text{N}_{10}\text{O}_4\text{Cl}_4\text{Cu}$, monoclinic *P*₂, *a* = 7.3794(7), *b* = 27.282(3), *c* = 14.369(2) Å, β = 93.193(1)°, *Z* = 4, Reflections collected / unique 10481 / 7780 [*R*(int) = 0.0437], *R*₁(obs) = 0.0569, *wR*₂(obs) = 0.1391. (3) $\text{C}_{24}\text{H}_{26}\text{Cl}_4\text{CuF}_2\text{N}_{10}$, triclinic *P*-1, *a* = 7.2851(5), *b* = 13.4033(7), *c* = 15.3589(9) Å, α = 92.018(4), β = 94.604(5), γ = 90.889(5)°, reflections collected / unique 9769 / 5931 [*R*(int) = 0.0534], *R*₁(obs) = 0.0441, *wR*₂(obs) = 0.1081. (4) $\text{C}_{38}\text{H}_{64}\text{Cl}_8\text{N}_{12}\text{O}_6\text{Pt}$, monoclinic *P*₂/*n*, *a* = 17.8564(14), *b* = 6.9720(4), *c* = 21.353(2) Å, β = 95.784(7)°, Reflections collected / unique 12896 / 4654 [*R*(int) = 0.0992], *R*₁(obs) = 0.0691, *wR*₂(obs) = 0.1714.

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I. Z. Travnicek, M. Malon, Z. Sindelar, K. Dolezal, J. Rolcik, V. Krystof, M. Strnad, and J. Marek, J. Inorg. Biochem., 84(1-2), 23-32 (2001). 2. M. Malon, Z. Travnicek, M. Marysko, R. Zboril, M. Maslan, J. Marek, K. Dolezal, J. Rolcik, V. Krystof, and M. Strnad, Inorg. Chim. Acta, 323,119-129 (2001).

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STRUCTURAL AND SPECTROSCOPIC MOTIFS OF ARTIFICIAL DI-IRON SUB-SITES OF ALL-IRON HYDROGENASE

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The hydrogenases catalyse the reversible interconversion of protons to dihydrogen. Recently, the X-ray crystal structures of all-iron hydrogenase from *Clostridium pasteurianum* and *Desulfovibrio desulfuricans* were reported. The active site, the H-centre, in each is comprised of a 6-Fe cluster, with 2 [4Fe₄S] clusters forming an electron transfer pathway from the site to the surface of the protein. In the active site, a [4Fe₄S] cluster is linked through a cysteinyl sulfur to a novel [2Fe₂S] unit, the sub-site, which is ligated by CO and CN⁻. Model chemistry can provide an understanding of the spectroscopic properties of H-centre redox states and mechanistic insights into how the enzyme works. There has been considerable debate as to whether the epr active redox state of the sub-site comprises an Fe(I)-Fe(II) or Fe(III)-Fe(II) pair. The Fe(I)-Fe(II) state is unprecedented in biology and recent structural and spectroscopic data for synthetic sub-site assemblies, including, the first class of artificial [2Fe₂S] assemblies [1 and references therein], now support its occurrence in the above system. Artificial sub-sites may provide new materials for electrocatalysis of hydrogen evolution/uptake, a key aspect of energy transduction relevant to progress towards an Hydrogen Economy.

1. M. Razavet, S. C. Davies, D. L. Hughes, and C. J. Pickett, Chem. Commun., 2001, 847; M. Razavet, S. J. George, S. A. Fairhurst, S. J. Borg, S. P. Best and C. J. Pickett, Chem. Commun., 2002, in press.

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DERIVATIVES OF SUBSTITUTED 3-TRICHLOROGERMYL PROPIONIC ACID

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Organogermanium chemistry has received a great impetus in the recent years due to the low toxicity of organogermanium compounds, such as germatrans, spirogermanium, germa-gama-lactones, etc. In continuing our work in the studies of biologically active organogermanium compounds, we have synthesized a number of compounds. The crystal structures of three of these have been determined and will be presented in the poster. Crystal data (I): $\text{C}_{31}\text{H}_{32}\text{GeO}_2$, *FW* = 509.15, monoclinic, *P*₂/*c*, *a* = 10.5010(5), *b* = 16.3680(6), *c* = 18.1040(7) Å, β = 123.410(2)°, *V* = 2597.5(2) Å³, *Z* = 4, *D*_x = 1.320 Mg/m³, *T* = 170 K, *F*(000) = 1064, *R* = 0.054, *GoF* = 1.03, for 5870 reflections collected on a KappaCCD diffractometer, and using full-matrix least-squares calculations on *F*² with the aid of SHELXL97. Crystal data (II): $\text{C}_{27}\text{H}_{23}\text{FGeO}_2$, *FW* = 471.04, monoclinic, *P*₂/*c*, *a* = 9.7278(1), *b* = 17.9075(3), *c* = 13.5408(2) Å, β = 110.907(1)°, *V* = 2203.51(5) Å³, *Z* = 4, *D*_x = 1.420 Mg/m³, *T* = 170 K, *F*(000) = 968, *R* = 0.027, *GoF* = 1.03, for 5026 reflections collected on a KappaCCD diffractometer, and using full-matrix least-squares calculations on *F*² with the aid of SHELXL97. Crystal data (III): $\text{C}_{31}\text{H}_{32}\text{GeO}_2$, *FW* = 509.15, monoclinic, *P*₂/*c*, *a* = 10.9842(3), *b* = 13.7103(4), *c* = 20.2450(5) Å, β = 113.857(1)°, *V* = 2788.33(13) Å³, *Z* = 4, *D*_x = 1.213 Mg/m³, *T* = 170 K, *F*(000) = 1064, *R* = 0.043, *GoF* = 1.01, for 6347 reflections collected on a KappaCCD diffractometer, and using full-matrix least-squares calculations on *F*² with the aid of SHELXL97.

References

Sheldrick, G.M. (1997). SHELXL97. Program to Refine Crystal Structures. University of Gottingen, Germany.

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