interactions in the structures **1-6** are the weak (CH ... O and CH ... Cl) bonds. In addition, in the structure **6** the stacking interactions between the fragments Ph-NO₂ are observed.

The derivatives of glycine and alanine (7 and 8) can be described as "freeze" cation-anion pairs. The Si...Br interatomic distances (2.7095(8) and 2.9704(7) Å, for 7 and 8, respectively) are much larger than those in the compounds of tetracoordinated Si (for instance, in the tris(pentafluorophenylmethyl)silylbromide the corresponding bond length is 2.208 Å). In turn, the Si-O bond in 7 and 8 is considerably shorter that in 1-6. In contrast to 1-6, the coordination polyhedron corresponds to the distorted tetrahedron with weak additional Si-O coordination bond.

To estimate the energy of intermolecular interactions, we have carried out quantum chemical calculations of the isolated molecules **7-8** and their associates. Also, the solid-state MAS ²⁹Si spectra of **7** and **8** were simulated using plane-wave DFT calculations.

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 $1: R=Me, 2: R=Ph, 3: R=p-MeC_{6}H_{4}, 4: R=p-ClC_{6}H_{4}, 5: R=p-BrC_{6}H_{4}, 6: R=p-NO_{2}C_{6}H_{4}, 6: R=p-NO_{2}$

Keywords: N-organosulfonyl-2-aminoacids, hypervalent silicon, crystal packing

MS61.P10

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The influence of crystal packing on the structure of N-organosulfonyl-2-aminoacid complexes containing $OSiC_3F$ coordination unit

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Silylfluorides with pentacoordinated Si atom obtained from N-organosulfonyl-2-aminoacids contain bioactive fragments and have high hydrolytic stability. In this study, we analyzed the influence of crystal packing on the geometry of coordination environment of the silicon atom in six pentacoordinated silylfluorides containing OSiC₃F coordination polyhedron.

Taking into account similar inductive effect of exocyclic substituents in the complexes **a-f**, one would expect the coordination site of the silicon atom to have a similar structure. Indeed, in all cases $OSiC_3F$ coordination polyhedron corresponds to a distorted trigonal pyramid. The deviation of the silicon atom from the plane of the equatorial C atoms varies within a fairly narrow range of 0.05 Å. The Si-O bond length decreases along with an increase in Si-F one. These bonds vary in the ranges of 2.135(2)-2.384(1) and 1.681(2)-1.646(1) Å, respectively. Observed differences in the lengths of the Si-O bond are mainly related to intermolecular interactions rather than to the nature of R and R' substituents. The presence of strong N-H...F bonds causes a strengthening of the coordination bond Si-O. In contrast, the formation of a N-H...O bond with the endocyclic O atom leads to weakening of the Si-O coordination bond.

In the solid state, molecules \mathbf{a} and \mathbf{e} form chains via N-H...F and N-H...O bonds, respectively; in the cases of \mathbf{c} , \mathbf{d} and \mathbf{f} complexes, centrosymmetric dimers are formed via N-H...O bonds with carbonyl (\mathbf{d} and \mathbf{f}) and sulfonyl (\mathbf{c}) groups of neighboring molecules. The

shortest Si-O bond is observed in the structure of \mathbf{e} , where F atom is involved in N-H...F hydrogen bond, which leads to the weakening of Si-F bond. In turn, the longest Si-O bond is found in the structure of \mathbf{b} (2.384(1) Å), which is the maximum value for all the ealier described complexes with OSiC₃F coordination polyhedron; the reason for this being is the intramolecular N-H...O hydrogen bond. Quantum chemical calculations of isolated molecules and their dimers were carried out to evaluate the energy of hydrogen bonds.

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Keywords: N-organosulfonyl-2-aminoacids, hypervalent silicon, crystal packing

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Crystal structure of $[Cu_2(L)_2(H_2O)_2](SO_4)_2$ ·2H₂O (L = pyridoxal thiosemicarbazone)

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Dark red monocrystals of the title complex were prepared by the reaction of methanol solutions of CuSO4.5H2O and piridoxal thiosemicarbazone (L) in mole ratio 1:1. Dinuclear complex cation has centrosymmetric structure. The Cu(II) is situated in a square-pyramidal environment ($\tau = 0.039$). The equatorial plane is formed by chelate ONS coordination of L and one water molecule, while the apical position is occupied by an alcoholic oxygen $O(2)^i$ (i = -x, 1 - y, -z) of the bridging ligand. The same way of coordination was reported for a similar ligand [1]. The Cu atom is shifted torwards the apical oxygen atom by 0.173(12) Å. The Cu-atom ligator bond lengths are in the range 1.917(2)-1.967(2) Å for O(1) N(3) and O(7), while Cu-S(1) and Cu–O(2)ⁱ are longer (2.2998(9) Å and 2.233(2) Å, respectively. Piridoxal thiosemicarbazone is coordinated in neutral, zwitterionic form as tetradentate bridging ligand, *i.e. via* phenol oxygen, hydrazine nitrogen, thioamide sulfur and hydroxymethyl oxygen, which, as the bridging ligator, connects the subunits. Such coordination results in formation of two metallocycles, five-membered (thiosemicarbazide) and six-membered (pyridoxilidene). The coordinated ligand slightly deviates from planarity, and dihedral angles between pyridine ring, sixand five-membered metallocycles are 7.54(10)°, 4.80(7)°, respectively. The six-membered metallocycle is in screw-boat conformation and can be described by the following puckering parameters: Q = 0.200(2) Å, $\theta = 67.9(9)^{\circ}$, $\varphi = 34.3(9)^{\circ}$, while the other rings are planar. Crystal structure of the complex is stabilized by an extended inter- and intramolecular 3D hydrogen-bond network. It can be mentioned that all possible hydrogen donors are involved in hydrogen-bonding, and that sulfate anion acts as multiple hydrogen acceptor.

The data were collected on an Oxford Diffraction Gemini S diffratometer using MoK α radiation (λ =0.71069 Å), and were corrected for Lorentz, polarization and background effects. The structure was solved by direct methods using SIR92 and refined by full matrix least square methods on F^2 using SHELXL-97 to R=0.036. Crystallographic data: C₁₈H₃₂Cu₂N₈O₁₆S₄, Mr = 871.90, monoclinic, space group $P2_1/n$, a = 7.2529(2) Å, b = 11.1766(4) Å, c = 18.8522(6) Å, $\beta = 93.751(3)^\circ$, V = 1524.94(8) Å³, Z = 2, μ (MoK α) = 1.756 mm⁻¹, GooF = 1.059.



The structure of complex cation $[Cu_2(L)_2(H_2O)_2]^{4+}$

 M.B. Ferrari, F. Bisceglie, G. Pelosi, P. Tarasconi, R. Albertini, P.P. Dall'Aglio, S. Pinelli, A. Bergamo, G. Sava, *J. Inorg. Biochem.* 2004, *98* 301–312.

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Incorporation of different lithium aluminates to titanium oxide organometallic ligands

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During the last years our research group has investigated the coordination of the organometallic oxides $[{Ti}(\eta^5-C_5Me_5)(\mu-O)]_3(\mu_3-CR)]$ [R = H(1), Me(2)] to different inorganic fragments of groups 1 and 13 elements. Reactions with group 1 derivatives lead to dehydrogenation processes of the alkylidyne moiety to give $[M(\mu_3-O)_3{Ti}_3(\eta^5-C_5Me_5)_3(\mu_3-C)]_2$ (M = Li, Na, K) complexes [1].

On the other hand, the novel adducts [$\{E_3M\}(\mu_3-O)(\mu-O)_2\{Ti(\eta^5-C_5Me_5)\}_3(\mu_3-CR)$] (R = H, Me; M = B, Al, Ga; E = alkyl, haloalkyl, halide) are obtained when group 13 derivatives react with 1 and 2 [2]. These species exhibit coordination of 1 or 2 through one of the three oxygen atoms of the Ti₃O₃ ring to the metal center. Now, we are studying the chemical reactivity of this family of compounds with different alkyl, amide or alkoxyde lithium derivatives in order to isolate the theoretical possible intermediate species in the reactions between 1 and 2 with the group 1 derivatives. We will show how dimethylamide and 2,6-dimethylphenoxyde lithium derivatives coordinate to the organometallic oxides in a chelate tridentate fashion. Now, this bonding

system is stabilized by the alkyl aluminum fragment, which avoids that the amide or alkoxyde fragments react with the alkylidyne moiety (see figure).

However, when the same reaction is carried out with *p*-tolyl lithium and the triphenylgallium adduct, abstraction and redistribution reactions are observed. The lithium atom is then encapsulated by two units of **1** and the new gallate species stabilizes a phenyllithium unit.



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Keywords: aluminate, gallium, titanium

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Discrete role of chlorine substitutions in the conformation and supramolecular architecture of isostructural arylsulfonamides

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Compounds containing sulfonamide groups SO₂NH occur in many biologically active molecules, including antimicrobial, antithyroid, antitumor and antimalarial drugs [1,2,3,4]. In this study, two arylsulfonamide derivatives differing for two chlorine substitutions at one of their two phenyl rings were synthesised and characterized by X-ray diffraction technique in order to establish structural relationships and the chlorine role in the conformation and crystal assembly. The compounds (4-N(phenyl-sulphonyl-amide-acetophenone)) (I) and 4-N [(2, 5-dichlorophenyl)sulfonyl-amide] acetophenone (II) were obtained by the equimolar coupling between benzene sulfonyl chloride or 2,5-dichlorobenzene sulfonyl chloride and 4-amineacetophenone in dichloromethane or acetone as solvent at 343K for until six hours. The precipitate was re-crystallized in suitable solvents to obtain the single crystals. Crystallographic data for compounds (I) and (II) were collected using a Enraf Nonius difractometer at room temperature. All crystal structures were solved by Direct Methods, and refined by full matrix least square method on F^2 . The non hydrogen atoms were refined anisotropically. All C-H hydrogens were placed geometrically and refined using riding model. The hydrogen H1 bonded to nitrogen was found from the difference Fourier map and its positional parameters were refined freely. The tables, geometrical calculations, molecular packing and drawings were done with the WinGX [5,6]. program package, obtaining the following results: (I) Molecular Formula: $C_{14}H_{13}NO_3S$. Unit Cell Parameters: a = 13.0007(5) Å, b = 8.3615(4) Å, c = 12.5179(6) Å, $\beta = 98.118(3)^{\circ}$, monoclinic, space group P2₁/c, Z = 4, V = 1347.13 (1) Å³. 5393 measured reflections with 3009 unique and 2214 observed. Final indices $R_1 = 0.0483$ for 177 refined parameters. (II) Molecular formula: $C_{14}H_{11}NO_3SCl_2$. Structure: a = 13.3622(2)Å, b = 8.1542(2) Å, c = 15.6845(3) Å, $\beta = 120.180(1)^{\circ}$, monoclinic, space group $P2_1/c$, Z = 4, 1477.24(5) Å³. 5653 measured reflections with 2972 unique and 2463 observed. Final indices $R_1 = 0.049$ for 195 refined parameters. There are several classical and non-classical hydrogen bonds and π - π stacking interactions contributing to stabilize the crystal packing of the compounds.

