

This article was downloaded by:

On: 23 January 2011

Access details: *Access Details: Free Access*

Publisher *Taylor & Francis*

Informa Ltd Registered in England and Wales Registered Number: 1072954 Registered office: Mortimer House, 37-41 Mortimer Street, London W1T 3JH, UK



Journal of Coordination Chemistry

Publication details, including instructions for authors and subscription information:

<http://www.informaworld.com/smpp/title~content=t713455674>

Copper(II) complexes with hydroxyl-containing dipeptides glycyl-*L*-serine and *L*-seryl-*L*-tyrosine

Tsonko Kolev^a; Bojidarka B. Koleva^b; Michael Spiteller^a

^a Institut für Umweltforschung, Universität Dortmund, Otto-Hahn-Strasse 6, 44221 Dortmund,

Germany ^b Lehrstuhl für Analytische Chemie, Ruhr-Universität Bochum, Universitätsstraße 150, 44780 Bochum, Germany

To cite this Article Kolev, Tsonko , Koleva, Bojidarka B. and Spiteller, Michael(2008) 'Copper(II) complexes with hydroxyl-containing dipeptides glycyl-*L*-serine and *L*-seryl-*L*-tyrosine', Journal of Coordination Chemistry, 61: 12, 1897 – 1905

To link to this Article: DOI: 10.1080/00958970701787687

URL: <http://dx.doi.org/10.1080/00958970701787687>

PLEASE SCROLL DOWN FOR ARTICLE

Full terms and conditions of use: <http://www.informaworld.com/terms-and-conditions-of-access.pdf>

This article may be used for research, teaching and private study purposes. Any substantial or systematic reproduction, re-distribution, re-selling, loan or sub-licensing, systematic supply or distribution in any form to anyone is expressly forbidden.

The publisher does not give any warranty express or implied or make any representation that the contents will be complete or accurate or up to date. The accuracy of any instructions, formulae and drug doses should be independently verified with primary sources. The publisher shall not be liable for any loss, actions, claims, proceedings, demand or costs or damages whatsoever or howsoever caused arising directly or indirectly in connection with or arising out of the use of this material.

Copper(II) complexes with hydroxyl-containing dipeptides glycyl-*L*-serine and *L*-seryl-*L*-tyrosine

TSONKO KOLEV†, BOJIDARKA B. KOLEVA*‡
and MICHAEL SPITELLER†

†Institut für Umweltforschung, Universität Dortmund,
Otto-Hahn-Strasse 6, 44221 Dortmund, Germany

‡Lehrstuhl für Analytische Chemie, Ruhr-Universität Bochum,
Universitätsstraße 150, 44780 Bochum, Germany

(Received 11 July 2007; in final form 28 August 2007)

Dipeptides glycyl-*L*-serine and *L*-seryl-*L*-tyrosine are tridentate ligands in coordination with Cu(II) through their NH₂⁻, N⁻ (from deprotonated amide group) and O-atom (by COO⁻ group), forming [Cu^{II}(LH₋₁)H₂O]. The fourth position of square-planar geometry of Cu²⁺ is occupied by H₂O as terminal ligand. Solid-state linear dichroic IR-spectroscopy, UV-Vis, mass spectrometry with ESI and FAB, tandem mass spectrometry (HPLC-MS/MS), TGV and DSC methods, EPR and magnetochemistry data prove the formation of five-membered chelate rings with participation of Cu²⁺ both in solution and in solid state.

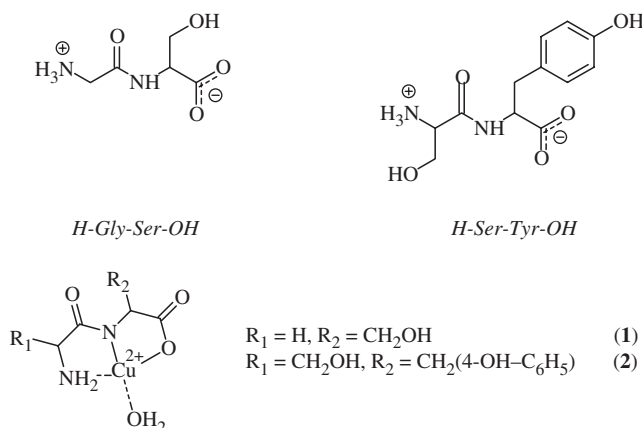
Keywords: Glycyl-*L*-serine; *L*-seryl-*L*-tyrosine; Cu(II) complexes; Solid-state IR-LD; UV-Vis; ESI-MS; FAB-MS; HPLC-MS/MS; TGV; DSC; EPR

1. Introduction

In the course of our systematic spectroscopic and structural elucidation of coordination ability of peptides with essential biometals [1–6] the complexation between Cu²⁺ and *L*-serine containing dipeptides glycyl-*L*-serine (*H-Gly-Ser-OH*) and *L*-seryl-*L*-tyrosine (*H-Ser-Tyr-OH*) (scheme 1) are studied. Investigation in solution and solid state includes the application of linear-polarized IR- (IR-LD) spectroscopy, mass spectrometry with electrospray ionization (ESI) and fast atom bombardment (FAB), hybrid method high performance liquid chromatography (HPLC) with tandem mass spectrometry (MS/MS), UV-Vis spectroscopy, electron magnetic resonance (EPR), thermogravimetry (TGA) and difference scanning calorimetry (DSC). *In vitro* study of Cu(II) complexes with *L*-serine containing peptides is of interest because Cu(II) is an essential biometal and important human proteins contain amino acid residue. A study of Cu-binding proteins in liver and kidney tissue has shown that the proteins are characterized with higher content of glutamic acid, serine, alanine and lysine [7]. The gly-ser fragment has been obtained in chymotrypsin, which is a prominent member

*Corresponding author. Email: BKoleva@chem.uni-sofia.bg

of the family of serine proteases [8]. Thus spectroscopic and structural elucidation of coordination with *L*-seryl containing peptides and proteins are important [9–11].



Scheme 1. Chemical diagram of peptides studied.

2. Experimental

2.1. Materials and methods

Dipeptides were purchased from Bachem and $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ from Acros Organics. The *conventional* (KBr pellets) and *IR-LD spectra* were recorded between 4000 cm^{-1} and 400 cm^{-1} on a Bomem–Michelson 100 FTIR spectrometer equipped with a Perkin Elmer wire-grid polarizer. 150 scans were performed for each spectrum with a resolution of 4 cm^{-1} . A 4-cyano-4'-alkylbicyclohexyl mixture (ZLI-1695 in the Merck notation) was used for orientation of the solid samples as a nematic liquid crystal colloid suspension [12–15].

Elemental analyses were performed according to the classical methods: C and H as CO_2 and H_2O , N through the Duma's method, chlorine by titration with $\text{Hg}(\text{NO}_3)_2$ after a wet digestion of the sample. The Cu^{2+} is obtained after ignition of a sample, dissolving of the residue in dilute HNO_3 and titration with standard solution of EDTA using murexide at pH 8 (ammonia buffer) as an indicator.

The *molecule weight* was determined using FAB mass spectra, measured on a Fusion VG Autospect instrument employing 3-nitrobenzyl alcohol as a matrix.

The *thermogravimetric (TGA) study* was performed using Perkin-Elmer TGS2 apparatus. The *calorimetric (DSC) analyses* are performed on DSC-2C Perkin Elmer equipment in argon.

The *UV-spectra* ($2.5 \cdot 10^{-4}\text{ M}$ aqueous solutions, 1 cm quartz cell) were obtained on a Perkin Elmer Lambda 15 UV-VIS spectrophotometer.

The *EPR* data are obtained in X-band on a Bruker ER 420 spectrometer.

HPLC-MS/MS analysis. The analyses of the samples were performed with a Thermo Finnigan surveyor LC-Pump. Compounds were separated on a Luna C18 column ($150 \times 2\text{ mm}$, $4\text{ }\mu\text{m}$ particle size) from Phenomenex (Torrance, CA, USA). The mobile

Table 1. HPLC-MS/MS conditions.

	<i>t</i> [min]	A [%]	B [%]	Flow rate [$\mu\text{L min}^{-1}$]
0	0.00	100	0	200
1	3.00	100	0	200
2	8.00	65	35	200
3	9.00	0	100	200
4	14.00	0	100	200
5	14.50	100	0	200
6	20.00	100	0	200

phase consisted of water +0.1% HCOOH (A) and acetonitrile +0.1% HCOOH (B) using a gradient program presented in table 1. The compound was detected via UV and a TSQ 7000 (Thermo Electron Corporation, Dreieich, Germany) mass spectrometer. The spectra were obtained using the TSQ 7000 equipped with an ESI ion source and operated with the following conditions: capillary temperature 180 °C; sheath gas 60 psi and spray voltage 4.5 kV. 1 mg mL⁻¹ of the sample was dissolved in acetonitrile and injected into the ion source by an auto sampler (Finnigan Surveyor). Excalibur 1.4 software processed the data obtained.

The *magnetic* measurements were carried out in argon by a Faraday balance.

2.2. Synthesis

The Cu(II) complexes with dipeptides *H-Gly-Ser-OH* (**1**) and *H-Ser-Tyr-OH* (**2**) were synthesized by common procedure: 5 mL 0.1 M solution of CuCl₂ · 2H₂O (170.4 mg) in solvent mixture H₂O:CH₃OH 1 : 1 was added to 5 mL 0.1 M solution of *H-Gly-Ser-OH* (162.2 mg) or *H-Ser-Tyr-OH* (268.2 mg) in the same solvent mixture. The molar ratio of metal to ligand is 1 : 1 in both cases. The blue precipitates formed after about 20 days were filtered off, washed with methanol and dried in air at 298 K. Yields: 72% for **1** and 63% for **2**, respectively. Found%: C, 24.9; H, 4.2; N, 11.7; Cu, 26.4. Calcd for [(C₅H₁₀N₂O₅)Cu] (**1**): C, 24.8; H, 4.1; N, 11.5; Cu, 26.2. Found%: C, 41.4; H, 4.5; N, 8.1; Cu, 18.4. Calcd for [(C₁₂H₁₆N₂O₆)Cu] (**2**): C, 41.4; H, 4.6; N, 8.0; Cu, 18.2.

3. Results and discussion

3.1. Conventional and linear-polarized IR-data

Investigation of Cu²⁺-complexation processes with dipeptides requires detailed IR band assignment of ligand and corresponding changes in the IR-spectra of the complexes isolated. For these reasons the IR-LD analysis started with interpretation of polarized spectra of pure dipeptides *H-Gly-Ser-OH* and *H-Ser-Tyr-OH*. The detail IR-spectroscopic assignment of *H-Gly-Ser-OH* was done [4], where the coordination ability with Au(III) was studied [4]. For *H-Ser-Tyr-OH* the IR-LD characterization is presented in this article with IR-characteristic bands of both ligands and corresponding complexes presented in table 2.

Table 2. Solid-state IR-spectral data of dipeptides *H-Gly-Ser-OH*, *H-Ser-Tyr-OH* and their Cu^{2+} -complexes **1** and **2**.

Assignment*	ν [cm^{-1}]			
	<i>H-Gly-Ser-OH</i> [5]	1	<i>H-Ser-Tyr-OH</i>	2
ν_{OH}	3170	3477	3214	3444
$\nu_{\text{H}_2\text{O}}$	–	3551	–	3473
$\nu_{\text{NH}_3^+}$ -stretches	3100–2300 broad	–	3300–2700	–
ν_{NH}	3315	–	3336	–
$\nu^{\text{as}}_{\text{NH}_2}$, $\nu^{\text{s}}_{\text{NH}_2}$	–	3340, 3206, 3100	–	3355, 3215, 3150
$\nu_{\text{C=O}}$	–	1710	–	1716
$\delta^{\text{as}}_{\text{NH}_3^+}$, $\delta^{\text{as}'}_{\text{NH}_3^+}$	1693, 1623	–	1681, 1633	–
$\delta_{\text{H}_2\text{O}}$	–	1636	–	1639
δ_{NH_2}	–	1681	–	1683
$\nu_{\text{C=O}}$ (Amide I)	1660	1642	1662	1650
8a , 8b	–	–	1610, 1575	1611, 1575
8b	–	–	1588	1590
$\delta^{\text{s}}_{\text{NH}_3^+}$	1590	–	1587	–
δ_{NH} (Amide II)	1555	–	1545	–
$\nu^{\text{as}}_{\text{COO}^-}$	1515	–	1556	–
19b	–	–	1467	1469
$\nu^{\text{s}}_{\text{COO}^-}$	1405	–	1411	–
11- γ_{CH}	–	–	842	848
$\delta_{\text{C=O}}$ (Amide IV)	721	724	742	750
γ_{NH} (Amide V)	668	–	665	–
$\gamma_{\text{C=O}}$ (Amide VI)	624	674	630	644

*The aromatic in-plane and out-of-plane modes are assigned using Wilson's notation [21].

The non-polarized IR-spectrum of *H-Ser-Tyr-OH* contains a broad maximum at $3300\text{--}2700\text{ cm}^{-1}$, assigned to $\nu_{\text{NH}_3^+}$ asymmetric and symmetric stretching modes. The intensive peaks at 3336 and 3214 cm^{-1} (table 2) belong to ν_{NH} and ν_{OH} modes. The low frequency shift of ν_{OH} is explained by participation of OH in the serine side chain in strong intermolecular interactions. The $1700\text{--}1400\text{ cm}^{-1}$ IR-spectral region contains maxima characteristic of --NH_3^+ , COO^- and amide groups and in-plane modes of *p*-disubstituted phenyl ring in the tyrosine side chain (table 2). Confirmation is obtained by the following IR-LD analysis. Simultaneous reduction of 3315 cm^{-1} and 1662 cm^{-1} peaks in the reduced IR-LD spectrum of *H-Ser-Tyr-OH* indicated their character as ν_{NH} and $\nu_{\text{C=O}}$ (Amide I) modes as well as a *transoid* configuration of the amide fragment. A reduction of both Amide V and Amide VI peaks (table 2) in the same dichroic ratio confirms their assignment as γ_{NH} and $\gamma_{\text{C=O}}$ out-of-plane frequencies.

Comparing the IR-spectral data of ligands and their Cu^{2+} -complexes the following main differences are determined (table 2): (i) A disappearance of ν_{NH} bands about 3330 cm^{-1} indicates Cu^{2+} coordination through N-amide supported by a preliminary deprotonation process in the ligands; (ii) Observation of new peaks (table 2) in the complex typical for $\nu^{\text{s}}_{\text{NH}_2}$ and Fermi-resonance split $\nu^{\text{s}}_{\text{NH}_2}$ of primary NH_2 -group, when the asymmetric $\text{HNH} \uparrow \text{X}$ intermolecular interactions are established; (iii) $1700\text{--}1400\text{ cm}^{-1}$ region indicated disappearance of all bending ($\delta_{\text{NH}_3^+}$) maxima of NH_3^+ and observation of δ_{NH_2} about 1680 cm^{-1} , typical for complexes where NH_2^- group is included in coordination. The fact that $\nu_{\text{C=O}}$ (Amide I) is significantly lower frequency and that δ_{NH} (Amide II) disappears in the IR-spectrum of the complex means that the metal ion is coordinated through N-amide (scheme 1); (iv) The observation of a new maximum of $\nu_{\text{C=O}}$ higher than 1700 cm^{-1} with disappearance of $\nu^{\text{as}}_{\text{COO}^-}$ and

$\nu^s_{\text{COO}^-}$ in the IR-spectra of complexes indicates coordination of metal with COO^- in the peptide distorting the C–O bond lengths and reversing C=O double and C–O single bonds in the complex (scheme 1); (v) As expected deprotonation of $-\text{C}(=\text{O})-\text{NH}$ -amide in the complex caused disappearance of the γ_{NH} mode and high frequency shift of $\delta_{\text{C=O}}$ and $\gamma_{\text{C=O}}$ (table 2); (vi) In both complexes new bands belonging to stretching and bending vibrations of H_2O are observed (table 2).

Elimination of δ_{NH_2} in the IR-LD spectra of Cu^{2+} -complexes caused disappearance of corresponding ones assigned to δ_{NH_2} and Fermi-resonance split $\nu^s_{\text{NH}_2}$, where transition moments are co-linear in the frame of one NH_2 -fragment (scheme 1, table 2). These data correlate with previously reported assignment of corresponding isolated amino acids and small peptides [16, 17].

3.2. ESI-MS, FAB-MS and HPLC-MS-MS data

The HPLC ESI-MS/MS spectra of Cu(II) complexes with *H-Gly-Ser-OH* and *H-Ser-Tyr-OH* are shown in figure S1A and S1B. The ESI-MS spectrum of **1** has peaks at m/z 259.02, corresponding to $[[\text{C}_5\text{H}_{11}\text{O}_5\text{N}_2\text{Cu}]^+ + \text{NH}_4^+]$ adduct with a molecule weight of 259.2 (figure S1A). It is well known that the Na^+ and NH_4^+ adducts are typical for ESI-MS in the analysis of peptides and their derivatives [18–20]. The peak at m/z 217.97 corresponds to species with fragmented metal ion. FAB-MS spectrum of **1** show a peak at m/z 241.0 assigned to $[\text{C}_5\text{H}_{11}\text{O}_5\text{N}_2\text{Cu}]^+$ with molecule weight of 241.69. The fragmentation of metal ion leads to an observation of m/z signal at 176.0 of $[\text{C}_5\text{H}_{11}\text{O}_5\text{N}_2]^+$ ion. The ESI-MS data of **2** (figure S1B) show an m/z at 364.5 corresponding again to NH_4^+ adduct of **2** with molecule weight of 365.2 ($[\text{C}_{12}\text{H}_{16}\text{O}_6\text{N}_2\text{Cu}]^+ + \text{NH}_4^+$). The fragmentation of metal ion leads to m/z of 266.9 of $[\text{C}_{12}\text{H}_{16}\text{O}_6\text{N}_2\text{Cu}]^+$. FAB-MS spectrum of **2** show a peak at m/z 351.1 assigned to $[\text{C}_{12}\text{H}_{16}\text{O}_6\text{N}_2\text{Cu}]^+$ ion.

3.3. TGA and DSC data

TGA data of **1** and **2** in the temperature range 300–600 K show a weight loss of 7.4% and 5.1% in **1** and **2**, respectively, corresponding to one H_2O in the Cu^{2+} complexes. The DSC data are shown in figure 1 and an enthalpy effect of $10.3 \text{ kcal mol}^{-1}$ **1** and $8.9 \text{ kcal mol}^{-1}$ **2** is observed, also confirming the inclusion of one solvent molecule in the coordination compounds obtained. The enthalpy effect is observed at the temperature about 125°C in both complexes, supposing that the solvent molecule is included in the coordination sphere.

3.4. UV-Vis spectra

The coordination of dipeptides (scheme 1) with Cu^{2+} causes hypsochromic shifts of the absorption maxima of phenyl chromophor in **2** less than 5 nm in UV-diapason (figure 2). New maxima both in **1** and **2** are observed at 588 nm ($\epsilon = 881 \text{ mol}^{-1} \text{ cm}^{-1}$) and 580 nm ($\epsilon = 901 \text{ mol}^{-1} \text{ cm}^{-1}$), respectively (figure 2). A low intensity shoulder about 600 nm (ϵ within $86\text{--}881 \text{ mol}^{-1} \text{ cm}^{-1}$) is observed in both complexes. These maxima

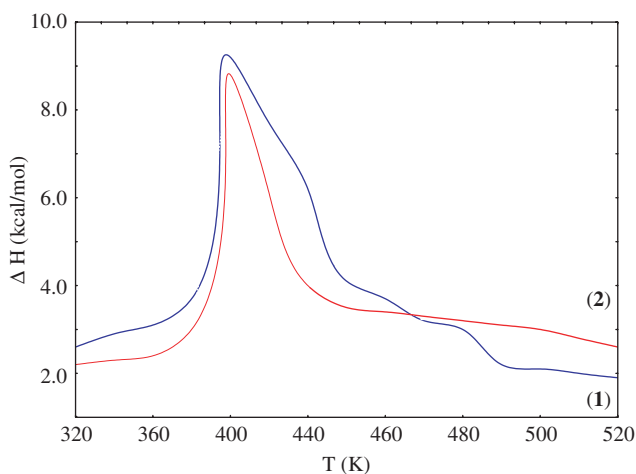


Figure 1. DSC data of **1** and **2** between 300–600 K.

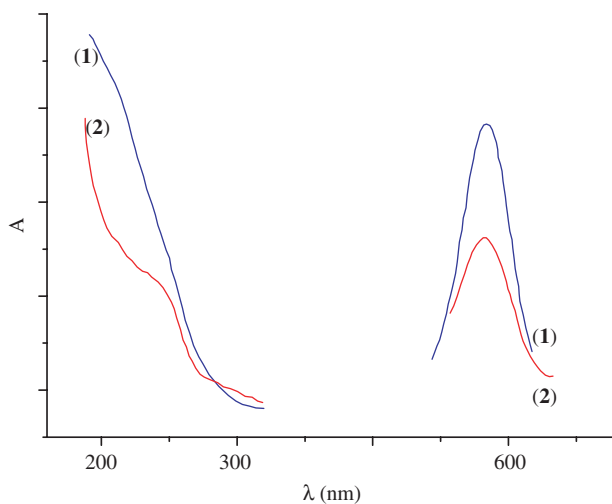


Figure 2. UV-Vis spectrum of Cu^{2+} -complexes of dipeptides studied in water/methanol solvent mixture.

correspond to d–d transitions as a result of the presence of the Cu–O and Cu–N bonds providing direct evidence about the complex formation.

3.5. EPR data

In methanol solution at room temperature isotropic EPR-spectra of the investigated mononuclear complexes with a resolved hyperfine structure for $^{63,65}\text{Cu}$ and parameters $g_{\text{iso}}=2.11$, $A_{\text{iso}}=73.11 \times 10^{-4} \text{ cm}^{-1}$ for **1** and $g_{\text{iso}}=2.09$, $A_{\text{iso}}=69.11 \times 10^{-4}$ for **2** are obtained. Frozen solutions at 77 K give anisotropic signals with additionally resolved SHFS from ^{14}N in the perpendicular region of

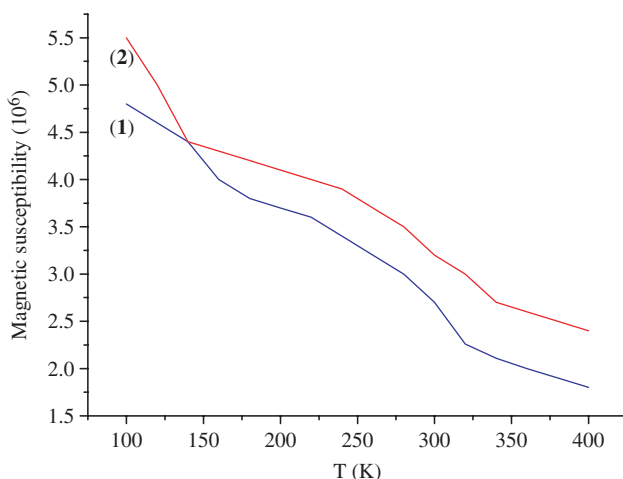


Figure 3. Magnetic susceptibility vs. T for **1** and **2**.

the magnetic field. The obtained parameters are: $g_{\parallel} = 2.19$ ($A_{\parallel} = 189.10 \times 10^{-4} \text{ cm}^{-1}$), $g_{\perp} = 2.02$ ($A_{\perp} = 48.50 \times 10^{-4} \text{ cm}^{-1}$) for **1** and $g_{\parallel} = 2.17$ ($A_{\parallel} = 191.10 \times 10^{-4} \text{ cm}^{-1}$), $g_{\perp} = 2.03$ ($A_{\perp} = 50 \times 10^{-4} \text{ cm}^{-1}$) for **2**. In both cases the A_{N} is 14 and indicates formation of Cu–N bonds. The complexes are also EPR active in the solid state both at 77 K and 298 K, indicating formation of mononuclear Cu^{2+} complexes with dipeptides.

3.6. Magnetochemistry data

The temperature dependence of the magnetic susceptibility for both complexes are typical for a mononuclear paramagnetic complex (figure 3), without any measurable exchange interaction between Cu^{2+} centers. The Curie–Weiss law is not obeyed in the temperature range studied and in such conditions the magnetic moment was calculated using the equation $\mu_{\text{eff}} = 2.828 (\text{k}_{\text{M}}T - \theta)^{1/2}$; $\theta = -80$. At low temperatures, the magnetic moment μ is temperature dependent and corresponds to a tetrahedral coordination of Cu^{2+} (figure 4), but at higher temperatures the μ -value differs from the theoretical one. Most probably the deviation observed is due to a deformation of the tetrahedral structure to a nearly square-planar one, a fact supported by the obtained EPR data.

4. Conclusions

By physical methods such as solid-state IR-LD spectroscopy based on orientation technique of samples as nematic liquid crystal suspensions, mass spectrometry (ESI and FAB), hybrid methods (HPLC MS/MS), UV-Vis, EPR, TGS and DSC analysis, the structures of Cu^{2+} -complexes with *H-Gly-Ser-OH* and *H-Ser-Tyr-OH* have been determined. The metal ion is coordinated through N- and O-atoms from NH_2^- and $-\text{COO}^-$ -groups. The joining of Cu^{2+} ion with N-amide, accompanied with

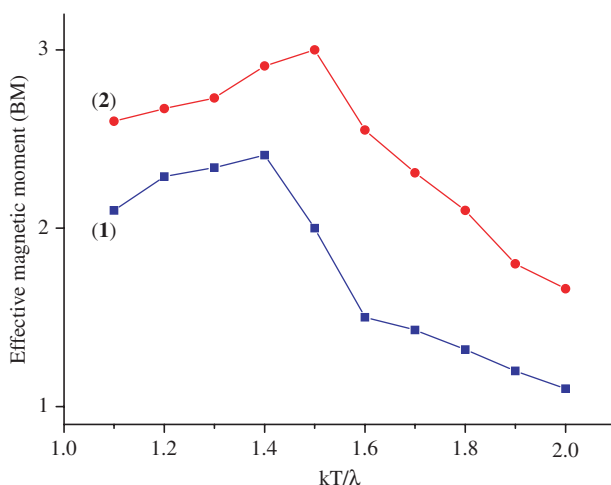


Figure 4. Graphs μ_{eff} vs. T for **1** and **2**.

deprotonation of corresponding $-\text{C}(=\text{O})-\text{NH}$ amide group, is also established. The fourth position of the square-planar geometry is a H_2O molecule as a terminal ligand.

Acknowledgements

B.K. wishes to thank the Alexander von Humboldt Foundation for the grant. T.K. wishes to thank the DAAD for a grant within the priority program “Stability Pact South-Eastern Europe” and the Alexander von Humboldt Foundation.

References

- [1] B.B. Ivanova, M. Mitewa. *J. Coord. Chem.*, **57**, 217 (2004).
- [2] B.B. Ivanova. *J. Coord. Chem.*, **58**, 587 (2005).
- [3] B.B. Ivanova, M.G. Arnaudov, St.T. Todorov. *J. Coord. Chem.*, **59**, 1749 (2006).
- [4] T. Kolev, B.B. Ivanova, S.Y. Zareva. *J. Coord. Chem.*, **60**, 109 (2007).
- [5] Ts. Kolev, S.Y. Zareva, B.B. Koleva, M. Spiteller. *Inorg. Chim. Acta.*, **359**, 4367 (2006).
- [6] B.B. Koleva, Ts. Kolev, M. Spiteller., *Inorg. Chim. Acta* (2006): <http://dx.doi.org/10.1016/j.ica.2006.11.002>.
- [7] A.E. Port, D.M. Hunt. *Biochem. J.*, **183**, 721 (1979).
- [8] N. Singh, T. Jabeen, S. Sharma, I. Roy, M. Gupta, S. Bilgrami, R. Somvanshi, K. Dey, M. Perbandt, C. Betzel, A. Srinivasan, T. Singh. *FEBS Journal*, **272**, 562 (2005).
- [9] G. Zou, G.L. Boyer. *223rd ACS National Meeting*, Orlando, FL, United States, April 7–11, 2002, ANYL-036.
- [10] M.A. Zoroddu, T. Kowalik-Jankowska, H. Kozłowski, K. Salnikow, M. Costa. *J. Inorg. Biochem.*, **84**, 47 (2001).
- [11] R. Liang, S. Senturker, X. Shi, W. Bal, M. Dizdaroglu, K.S. Kasprzak. *Carcinogenesis*, **20**, 893 (1999).
- [12] B.B. Ivanova, M.G. Arnaudov, P.R. Bontchev. *Spectrochim. Acta*, **60(A)**, 855 (2004).
- [13] B.B. Ivanova, D.L. Tsalev, M.G. Arnaudov. *Talanta*, **69**, 822 (2006).
- [14] B.B. Ivanova, V.D. Simeonov, M.G. Arnaudov, D.L. Tsalev. *Spectrochim. Acta*, **67(A)**, 66 (2007).
- [15] B.B. Koleva, T. Kolev, V. Simeonov, T. Spassov, M. Spiteller. *Adv. Colloid. Int. Sci.*, submitted (2007).
- [16] T. Norio, H. Masahiko, T. Kiyoshi, S. Tokura, A. Tsutsumi. *Pept. Chem.*, 245 (1985).

- [17] M. Tsuboi, Y. Ezaki, M. Aida, M. Suzuki, A. Yimit, K. Ushizawa, T. Ueda. *Biospectroscopy*, **4**, 61 (1998).
- [18] D.G. Morgan, M.M. Bursey. *J. Mass Spectrom.*, **30**, 473 (1995).
- [19] Y. Wang, F. Ke, K.W.M. Siu, R.J. Guevremont. *J. Mass Spectrom.*, **31**, 45 (1996).
- [20] D.G. Morgan, M.M. Bursey. *J. Mass Spectrom.*, **30**, 290 (1995).
- [21] G. Varsanyi, *Vibrational Spectra of Benzene Derivatives*, Akademiai kiado, Budapest (1969).