

*J. Synchrotron Rad.* (1999). 6, 406–408

## EXAFS studies of the novel iron(III) complexes with an N/S(Se) chromophore simulating ligand environment of the active site of nitrile hydratase

V. G. Vlasenko<sup>a</sup>, A. T. Shuvaev<sup>a</sup>, T. I. Nedoseikina<sup>a</sup>, A. L. Nivorozhkin<sup>b\*</sup>, A. I. Uraev<sup>b</sup>, A. D. Garnovskii<sup>b</sup> and O. Y. Korshunov<sup>b</sup>

<sup>a</sup>*Institute of Physics, Rostov State University, pr. Stachki 194, 344104 Rostov on Don, Russia,*

<sup>b</sup>*Institute of Physical Organic Chemistry, Rostov State University, pr. Stachki 194/2, 344104 Rostov on Don, Russia.*

\*E-mail: vlasenko@iphys.rnd.runnet.ru

\*\*Present address: Department of Chemistry and Chemical Biology, Harvard University, 12 Oxford St, Cambridge, MA 02138, USA.

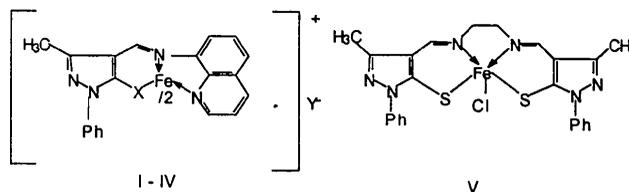
Novel metal chelates with an N/S(Se) chromophore have been synthesized and structurally characterized to model the iron site of the hydrolytic metalloenzyme nitrile hydratase. Iron-sulfur bond lengths Fe-S (2.24 Å) estimated from EXAFS are similar to that reported (Huang et al., 1997) for nitrile hydratase (2.21 - 2.23 Å). Iron-nitrogen bond lengths Fe-N 2.00 Å are also identical to those in the enzyme (1.99 Å). Another investigated metal complex with Fe(III)N<sub>2</sub>S<sub>2</sub>Cl coordination contains a potentially labile terminal chloride ligand which may be substituted by solvent molecules and can be useful as possible analogues of the substrate-bound form of the enzyme.

**Keywords:** EXAFS; metal chelates; nitrile hydratase.

### 1. Introduction

There is an increasing interest in the metal active centers of non-heme metalloproteins in biology. An example of a metalloenzyme with an iron-thiolate active site that does not belong to the heme or Fe-S clusters families is bacterial nitrile hydratase, which catalyzes the conversion of nitriles to corresponding amides (Kobayashi et al., 1992; Yamada H., Kobayashi M., 1996). Combined physicochemical results provided evidence for a presence of the low-spin (S=1/2) Fe<sup>III</sup>N<sub>3</sub>S<sub>2</sub> chromophore with a sixth coordination site occupied by a water molecule or hydroxide ion, which is probably replaced by nitrile in the catalytic cycle (Scarow et al., 1987; Sugiura et al., 1987; Scarow et al., 1996). A recent X-ray structural study (Huang et al., 1997) revealed a FeS<sub>2</sub>N<sub>2</sub>(H<sub>2</sub>O) site containing two deprotonated carboxamido nitrogens. Following the synthesis of several model compounds to imic a N/S ligand environment of the nitrile hydratase active site and some of its spectroscopic properties (Nivorozhkin et al., 1997) we report here an EXAFS study of those and new chelate complexes of Schiff-base ligands

with the supplementary coordinating groups I-IV. A series of complexes I-IV represents the only available set of isoligand Fe(III)N<sub>2</sub>X<sub>2</sub> species with variable donor atoms X=O, S, Se. Complex V contains a potentially labile terminal chloride ligand which may be substituted by solvent molecules and thus has a potential of a suitable starting point to pursue an ultimate goal of nitrile hydrolysis.



I: X=S, Y= FeCl<sub>4</sub><sup>-</sup>; II: X=S, Y= ClO<sub>4</sub><sup>-</sup>; III: X=O; Y= ClO<sub>4</sub><sup>-</sup>; IV: X=Se, Y= ClO<sub>4</sub><sup>-</sup>

### 2. Experimental section

Complexes I-V were synthesized according to ref. (Huang et al., 1997; Nivorozhkin et al., 1997; Uraev et al., 1997) by a reaction of the ligands H<sub>2</sub>L, [N,N'-bis(5-mercapto-3-methyl-1-phenylpyrazol-4-ylmethylene)ethylenediamine] and HL, [5-mercapto-1-phenyl-3-methyl-4-ylmethylene(8-amino)-quinoline] with Fe(ClO<sub>4</sub>)<sub>3</sub>·nH<sub>2</sub>O or FeCl<sub>3</sub>·6H<sub>2</sub>O in ethanol. The samples for XAS were ground to fine powders and mixed with apiezon grease. The mixtures were formed into flat samples of the appropriate thickness ( $\mu \approx 2.7$ ) for EXAFS measurements in transmission mode. X-ray absorption data were obtained using a laboratory XAFS spectrometer with a focussing crystal-monochromator (Shuvaev et al., 1988). A Mo X-ray tube (U=17 kV, I=25 mA) and cylindrically curved quartz (1340) crystal-monochromator of Iohan type in Rowland geometry were used. The energy resolution was determined using the width of the characteristic lines (~ 2 eV in the region of the FeK-edge).

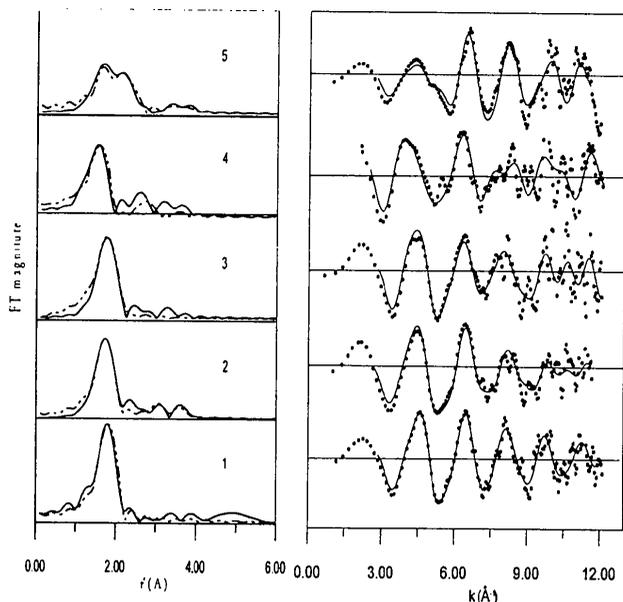
The EXAFS data were weighted by k<sup>2</sup>, and truncated at k=3 and k=12 Å<sup>-1</sup>, following background subtraction. The square filter window between 1.4 and 3.5 Å isolated the significant Fourier transform (FT) peaks, which were then back-transformed into k-dependent data. The best-fit distances (R), coordination numbers (N), and Debye-Waller (DW) factors were obtained from multishell fits to the Fourier-filtered data using empirical phase and amplitude functions. The empirical phase and amplitude functions were derived from FEFF5 (Rehr et al., 1991) of a crystallographically characterized complex I.

### 3. Results

The EXAFS and the k<sup>2</sup>χ Fourier transforms (FT) magnitude for the complexes I-V and the corresponding data fit are shown in Fig. 1. The principal features of FT of the compounds I, II, V include a large peak at r= 1.69-1.73 Å corresponding to the first coordination sphere, and smaller peaks at r=2.5-3.6 Å. The smaller FT peaks in III are similar to those in I, II, V whereas the large peak occurs at r= 1.49 Å due to the absence of the Fe-S/Cl scatterers in the molecule. The splitting of the largest feature in the FT of IV into two peaks is presumably due to the presence of the long Fe-Se distances.

The structure of I was determined by X-ray crystallography (Nivorozhkin et al., 1997) and served as a

reference to obtain EXAFS parameters used in subsequent fits of all the samples. The results of the curve fitting analysis are summarized in Table 1.



**Figure 1**

The left panel shows the magnitude of Fourier transforms of  $k^2\chi$  for compounds: **I** - (1), **II** - (2), **III** - (4), **IV** - (5), **V** - (3). The solid line is the experimental data and the dashed line is the theoretical fit. The right panel shows EXAFS data (dots) and fits (solid line) using parameters presented in Table 1.

The molecular structure of **I** indicates that the iron center is octahedrally coordinated by the two azomethine, and two quinoline nitrogens and two thiolate sulfur atoms in *cis*-positions. Iron-sulfur bond lengths,  $\text{Fe-S}_1=2.227(2)$  Å and  $\text{Fe-S}_2=2.230(2)$  Å, are similar to that reported for nitrile hydratase ( $\text{Fe-S}=2.21$  Å from the EXAFS data (Scarrows et al., 1996)). Iron-nitrogen bond lengths in **I**,  $\text{Fe-N}_{\text{av}}=1.99$  Å, are also identical to those in the enzyme. The EXAFS results on **I** agree with the X-ray diffraction data. Complexes **I-IV** are apparently isostructural. While **I**, **II** and **IV** contain low-spin ( $S=1/2$ )  $\text{Fe(III)N}_3\text{X}_3$  iron centers, **III** is a high-spin species ( $S=5/2$ ).  $S=1/2 \rightarrow S=5/2$  spin transition is usually accompanied by a certain elongation of the metal-ligand bonds (Sim et al., 1981). This trend could be also revealed by a perusal of the Table 1.

#### Acknowledgment

We are grateful to the Russian Fund of Fundamental Research (Grant ' 97-03-33479a) for financial support of this work.

**Table 1**

Results of multishell fits of the EXAFS data for compounds **I-V** and  $\text{Fe}(\text{acac})_3$  (R-average distance, N- number of scattering atoms,  $\sigma^2$  - Debye-Waller factors, Q-parameter of quality of fit)<sup>a</sup>.

Compd/parm	R, Å	N	$\sigma^2$ , Å <sup>2</sup>	atom	Q,%
<b>I</b> <sup>b</sup>	2.00	4	0.0020	N	1.6
	(1.969, 2.005)				
	2.24	2	0.0037	S	
<b>II</b>	(2.227, 2.230)				2.8
	2.18	4	0.0030	Cl <sup>c</sup>	
<b>III</b>	2.03	4	0.0035	N	2.0
	2.24	2	0.0030	S	
	1.97	2	0.0026	O	
	2.05	4	0.0043	N	
<b>IV</b>	3.12	8	0.0031	C	3.0
	3.52	6	0.0040	C	
	2.04	4	0.0020	N	
	2.43	2	0.0040	Se	
<b>V</b>	2.37	1	0.0030	Fe <sup>d</sup>	4.0
	1.95	1	0.0040	C <sup>d</sup>	
	2.00	2	0.0030	N	
<b>Fe(acac)<sub>3</sub></b> <sup>e</sup>	2.24	2	0.0040	S	2.0
	2.28	1	0.0030	Cl	
	2.02	6	0.0027	O	
	(1.998)				

<sup>a</sup>The first-shell Fe-N/O/S distances are typically accurate to  $\pm 0.02$  Å, second and third shell to  $\pm 0.05$  Å. Coordination numbers are generally considered accurate to  $\pm 20$ -25 %. The estimates are determined from comparisons between EXAFS and crystallography of distances and coordination numbers. <sup>b</sup>The coordination numbers for the compound **I** given in the table are for the  $[\text{FeN}_3\text{S}_2]$  centre, as well as  $[\text{FeCl}_4]^-$  anion. For simulations were used average of these coordination numbers. <sup>c</sup> Distance of the Fe-Cl in  $\text{FeCl}_4^-$  anion. The crystallographic distances for **I** are given in parentheses (Nivorozhkin et al., 1997). <sup>d</sup> R and N were obtained from Se K-edge EXAFS spectra. <sup>e</sup> Distance in parentheses is from (Scarrows et al., 1987).

#### References

- Huang, W., Jia J., Cummings, J., Nelson, M.J., Shneider, G., Lindquist, Y. (1997). *Structure*, **5**, 691.  
 Kobayashi, M., Nagasawa, T., Yamada, Y. (1992). *Trends Biotechnol.* **10**, 402-408.  
 Nivorozhkin, A.L., Uraev, A.I., Bondarenko, G.I., Antsyshkina, A.S., Kurbatov, V.P., Garnovskii, A.D., Turta, C.I., Brashoveanu, N.D. (1997). *Chem. Commun.*, 1711-1712.  
 Nivorozhkin, A.L., Toftlund, H., Nielsen, M. (1994). *J. Chem. Soc., Dalton Trans.*, pp. 361-367.  
 Rehr, J.J., Mustre de Leon, J., Zabinsky, S.I. & Albers, R.C. (1991). *J. Am. Chem. Soc.* **113**, 5135.  
 Scarrows, R.C., Maroney, M.J., Palmer, S.M., Que, L.Jr., Roe, A.L., Salowe, S.P., Stubbe, J.-A. (1987). *J. Am. Chem. Soc.*, **109**, 7857-7864.  
 Scarrows, R.C., Brennan, B.A., Cummings, J.G., Jin H., Duong, D.J.,

- Kindt, J.T., Nelson M.J. (1996). *Biochem.* **35**, 10078-10088.
- Shuvaev, A.T., Helmer, B.Y., Lyubeznova, T.A. (1988). *Prib. I Tehn. Exp.* **3**, 234-235.
- Sim, P. G., Sinn, E., Petty, R.H., Merrill, C.L., Wilson, L.J. (1981). *Inorg. Chem.* **20**, 1213.
- Sugiura, Y., Kuwahara, J., Nagasawa, T., Yamada, H. (1987). *J. Am. Chem. Soc.* **109**, 5848-5850.
- Uraev, A.I., Kurbatov, V.P., Garnovskii, A.D. (1997). *Russ. J. Koord. Khim.* **23**, 59.
- Yamada, H., Kobayashi, M. (1996). *Biosci. Biotechnol. Biochem.* **60**, 1391-1400.

(Received 10 August 1998; accepted 18 December 1998)