

Solution Behavior and Zinc Complexation of Dipeptides Made up Solely from Histidine and Cysteine[☆]

Peter Gockel, Raphael Vogler, and Heinrich Vahrenkamp*

Institut für Anorganische und Analytische Chemie der Universität Freiburg,
Albertstraße 21, D-79104 Freiburg, Germany

Received March 21, 1996

Key Words: Dipeptides / Histidine / Cysteine / Zinc complexes, solution studies of

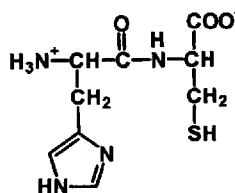
The eight possible types of dipeptides made up from histidine and cysteine, namely the unprotected ones H-His-Cys-OH (**1**) and H-Cys-His-OH (**2**), the *N*-protected ones Ac-His-Cys-OH (**3**) and Ac-Cys-His-OH (**4**), the *C*-protected ones H-His-Cys-OEt (**5**) and H-Cys-His-NH₂ (**6**), and the fully protected ones Ac-His-Cys-OEt (**7**) and Ac-Cys-His-OEt (**8**) were prepared analytically pure. Their acid-base behavior and zinc complexation in solution were studied potentiometrically. In all cases 1:1 (zinc:peptide) complexes are the dominant species near neutral pH. 1:2 complexes could only be detected for **4**, **6**, **7**, and **8**, and precipitations occurred in

basic media. From equimolar mixtures of the peptides and zinc salts (chloride, iodide, perchlorate) solid complexes were obtained upon addition of a base. They always contain coordinating anions (chloride, iodide, or trifluoroacetate which was introduced with the peptide). They have the composition LZnX for L = **5–8** and L₄Zn₅X₂ for L = **1–4**. It is proposed that all complex species observed contain tetrahedral zinc, that they are monomers in dilute solution and thiolate-bridged oligomers in the solid state, and that the peptides as a rule occupy at least three coordination sites.

Cysteine and histidine are the most prominent amino acids for the coordination of zinc in proteins, their thiolate and imidazole donor functions being ideally suited for the moderately soft Lewis acid Zn²⁺ in low coordination numbers^[2–4]. While catalytically active zinc centers in metalloenzymes bear at least one additional oxygen donor (aspartate, glutamate, water, or the substrate), exclusive *N* and *S* coordination is realized in most structural zinc centers in enzymes^[2,3] or the zinc finger proteins^[5,6]. It is for this biological significance that the coordination chemistry of zinc with the amino acids cysteine and histidine and their derivatives draws its justification.

The earliest preparative studies of cysteine- and histidine-containing zinc complexes date back to the fifties^[7,8], and the first review mentioning them appeared in 1973^[9]. The lack of crystallization of the resulting species has made potentiometric investigations of solution equilibria an important method for this subject, having again been applied quite early to the simple systems^[10] and appearing in tabulations^[11] and review articles^[4]. However, while the zinc coordination of the two simple amino acids and their derivatives is well-investigated and while small peptides containing one of them have been used to a reasonable degree for this purpose^[4,9,12–14], very few reports exist on the coordination chemistry of peptides containing cysteine *and* histidine, and to our knowledge only one such dipeptide, benzoyl-Cys-His-OME, has been used for zinc complexation^[15].

In continuation of our systematic studies^[1,10,13,14,16] we therefore chose the series of the eight possible types of Cys/His dipeptides **1–8** for an investigation of their zinc complex chemistry. This series consists of four pairs being either completely unprotected (**1**, **2**), only *N*-protected (**3**, **4**), only *C*-protected (**5**, **6**), or fully protected (**7**, **8**), always having the free thiol and imidazole functions available for coordination. Depending on the degree of protection, these dipeptides are potentially tetradentate (**1**, **2**), tridentate (**3–6**), or bidentate (**7**, **8**).



H-His-Cys-OH

1

Ac-His-Cys-OH

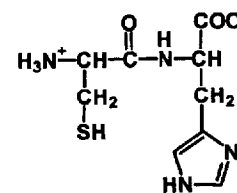
3

H-His-Cys-OEt

5

Ac-His-Cys-OEt

7



H-Cys-His-OH

2

Ac-Cys-His-OH₂

4

H-Cys-His-NH₂

6

Ac-Cys-His-OEt

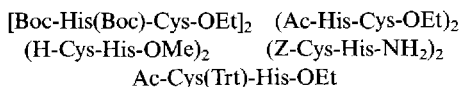
8

Peptide Syntheses

The cysteine/histidine combination is the most difficult one among dipeptides, due to the trifunctional nature of

[⊞] Part 6: Ref.^[1].

both constituents. It therefore took some effort to obtain the eight dipeptides, specifically the unprotected ones **1** and **2**, in analytically pure form. When necessary, we used the Boc (*tert*-butoxycarbonyl) group to protect the histidine imidazole and the Trt (trityl) group to protect the cysteine thiol function, using in addition cystine instead of cysteine in the early stages of the peptide syntheses. Starting from histidine, cystine, histidine ethyl ester, and *N*-acetylcysteine and using the established procedures for protection and coupling^[17], we prepared the following dipeptide derivatives as the immediate precursors of the dipeptides **1–8** (*Z* = benzyloxycarbonyl):



From these the protecting groups were removed according to the sequences described in the experimental part. In general, trifluoroacetic acid was used to remove Boc groups, dithioerythritol was used to cleave the cystine S–S bonds, and alkaline hydrolysis was used to remove ester groups. Of the four *C*-protected dipeptides **5–8**, **6** is the only one bearing an amide instead of an ester function. The reason for this is that H-Cys-His-OEt proved to be somewhat labile towards ester hydrolysis during the potentiometric titrations.

After the deprotection steps the resulting peptides **1–8** were purified by HPLC using the reversed-phase adsorbent Nucleosil 7 C18 and eluting with acetonitrile/water mixtures containing 0.1% trifluoroacetic acid (HTFA). They were isolated as analytically pure colorless powders containing at least 1.5 equivalents of HTFA per equivalent of peptide. The HTFA content was verified by ¹⁹F-NMR spectroscopy ($\delta = -73.5 \pm 2$) and quantified by potentiometric titrations.

Solution Equilibria

The eight dipeptides **1–8** contain variable numbers of acid/base functions and donor sites for metal complexation. The method of potentiometric titration was used to quantify their behavior in solution and to establish their mutual interrelations. The experimental procedures used correspond to those applied by us to the related derivatives of the single amino acids histidine and cysteine^[10] which can also serve as points of reference.

Table 1 lists the pK_a values for the four dissociable functions in the dipeptides. The imidazole function ($pK \approx 6–7$) and the thiol function ($pK \approx 8$) are available in all eight cases, while the carboxylic acid function ($pK \approx 2.5$) and the amino function ($pK \approx 9$) are blocked each in four cases.

The unprotected amino acids cysteine and histidine have carboxylate pK_a values of 1.82 and 1.77, respectively, which are raised to 3.08 and 2.94 in their *N*-acetylated derivatives. The carboxylate pK_a values of **1–4** lie between these extremes. The spatial (**1**, **2**) or chemical (**3**, **4**) removal of the stabilizing NH_3^+ functions raises the pK_a values with respect to those of the free amino acids, but the fact that distant stabilizing functions are still present (NH_3^+ in **1** and **2**,

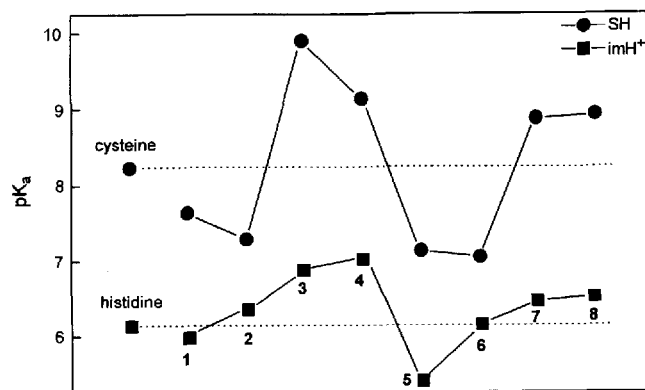
Table 1. pK_a values of the dipeptides

Acid	COOH	imH ⁺	SH	NH ₃ ⁺
HHisCysOH (1)	2.33(10)	5.98(4)	7.64(3)	9.89(2)
HCysHisOH (2)	2.74(3)	6.37(1)	7.30(1)	9.48(1)
AcHisCysOH (3)	2.85(6)	6.82(4)	9.83(3)	--
AcCysHisOH (4)	2.81(10)	7.02(8)	9.13(6)	--
HHisCysOEt (5)	--	5.45(14)	7.17(12)	8.84(8)
HCysHisNH ₂ (6)	--	6.13(1)	7.03(1)	9.26(1)
AcHisCysOEt (7)	--	6.48(2)	8.89(2)	--
AcCysHisOEt (8)	--	6.53(3)	9.00(2)	--

im-H⁺ in **1–4**) limits the rise. Likewise, in comparison with the NH_3^+ pK_a value of cysteine (10.36), those of **2** and **6** with *N*-terminal cysteine are lowered by about 1 unit. In comparison with the NH_3^+ pK_a value of histidine (9.24) those of **1** and **5** with *N*-terminal histidine are not typically changed. This is attributed to the fact that the nearest acid/base function in **1** and **5** (imidazole) is uncharged and that in **2** and **6** (thiolate) is charged when the NH_3^+ function is deprotonated. In the same context it is noticeable that among the eight peptides H-His-Cys-OH (**1**) has the highest acidity of the carboxylic acid function and the highest basicity of the amino function. This indicates that this peptide prefers a conformation which minimizes the distance between these functions in solution.

Figure 1 is a graphical representation of the pK_a variations of the thiol and imidazole functions along the series. The reference points are the corresponding pK_a values of cysteine (8.24) and histidine (6.14). In general, both functions are more acidic for the *N*-unprotected peptides (**1**, **2**, **5**, **6**) and less acidic for the *N*-protected peptides (**3**, **4**, **7**, **8**). Conversely, the effect of changing the C terminus (protected vs. unprotected) is less significant. This indicates that the charge at the N terminus is the dominating mediator for both the SH and the imH⁺ acidities which to our opinion is again a conformational effect.

Figure 1. pK_a values for the imidazole and thiol functions in the peptides **1–8**



For the evaluation of the complexation equilibria potentiometric titrations were performed for 1 mM solutions of the peptides in the presence of ca. 0.3, 0.6, or 1.2 equivalents of $\text{Zn}(\text{NO}_3)_2$. Under these conditions precipitation of

zinc compounds in basic media was minimized (it was, however, observed for 1–3), and the titration curves could be analyzed unambiguously. Figures 2 and 3 show two typical species distributions.

Figure 2. Species distribution in solutions of Ac-Cys-His-OH (4) and $\text{Zn}(\text{NO}_3)_2$ (1.6:1)

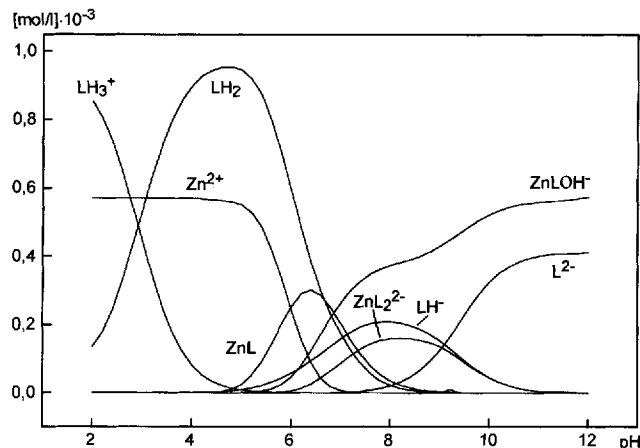
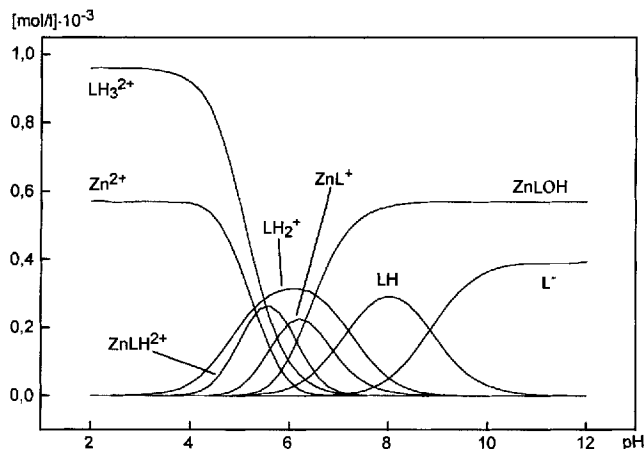


Figure 3. Species distribution in solutions of H-His-Cys-OEt (5) and $\text{Zn}(\text{NO}_3)_2$ (1.7:1)



The species observed in solution are the peptides with varying degrees of protonation, the 1:1 complexes ZnL and ZnLH, the singly deprotonated 1:1 complexes ZnLOH, and the 1:2 complexes ZnL_2 . It must be noted that this notation neglects the charges of the species which are correctly assigned in Figures 2 and 3 and which can be derived by evaluating the charge of the fully protonated ligand LH_n for which $n = 4$ for 1 and 2, $n = 3$ for 3–6, and $n = 2$ for 7 and 8. The notation ZnLOH corresponds to a species with $n = 1$ to which we assign a zinc complex with a deprotonated aqua ligand rather than with a peptide ligand deprotonated at the NH function. Table 2 lists the overall stability constants β for the observed species ZnL ($\beta_{(110)}$), ZnL_2 ($\beta_{(120)}$), ZnLH ($\beta_{(111)}$), and ZnLOH ($\beta_{(11-1)}$), according to the following definitions:

$$\beta_{(110)} = \frac{[\text{ZnL}]}{[\text{Zn}][\text{L}]}$$

$$\beta_{(120)} = \frac{[\text{ZnL}_2]}{[\text{Zn}][\text{L}]^2}$$

$$\beta_{(111)} = \frac{[\text{ZnLH}]}{[\text{Zn}][\text{L}][\text{H}]}$$

$$\beta_{(11-1)} = \frac{[\text{ZnLOH}][\text{H}]}{[\text{Zn}][\text{L}]}$$

$$K_{\text{ZnL}} = \frac{[\text{ZnL}]}{[\text{ZnL}][\text{L}]}$$

$$K_{\text{ZnL}_2} = \frac{[\text{ZnL}_2]}{[\text{ZnL}][\text{L}]}$$

$$K_{\text{ZnLH}} = \frac{[\text{ZnLH}]}{[\text{ZnL}][\text{H}]} = \frac{[\text{ZnLH}]}{[\text{Zn}][\text{LH}]}$$

$$K_{\text{ZnLOH}} = \frac{[\text{ZnLOH}][\text{H}]}{[\text{ZnL}]}$$

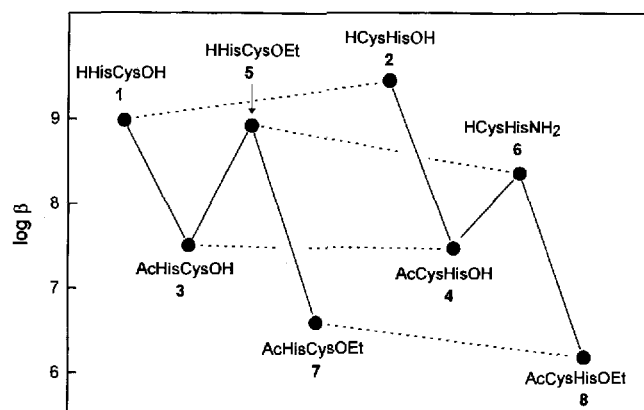
Table 2. Stability constants (log β) for the zinc complex species in solution

Ligand	ZnL	ZnL ₂	ZnLH	ZnLOH
HHisCysOH (1)	9.00 (3)	--	15.35 (3)	--
HCysHisOH (2)	9.41 (6)	--	15.34 (3)	--
AcHisCysOH (3)	7.52 (6)	--	12.80 (8)	--
AcCysHisOH (4)	7.43 (6)	12.97 (8)	--	0.56 (6)
HHisCysOEt (5)	8.36 (10)	--	14.31 (12)	2.00 (20)
HCysHisNH ₂ (6)	8.90 (6)	16.33 (9)	13.75 (6)	2.87 (5)
AcHisCysOEt (7)	6.17 (2)	12.42 (3)	--	-0.98 (12)
AcCysHisOEt (8)	6.57 (6)	11.72 (9)	--	0.20 (7)

In all cases the 1:1 complexes ZnL are observed. Their stabilities are quite high as can be seen by comparison of the log β values for the H-His-Cys ligands 1 (9.00) and 5 (8.36) with that for H-His-OMe (4.45), or of those for the H-Cys-His ligands 2 (9.41) and 6 (8.90) with that for H-Cys-OEt (7.42). The 1:2 complexes ZnL_2 could be identified only for 4, 6, 7, and 8. Their gain in stability with respect to the 1:1 complexes amounts to about 5 log units except for 8 where it is 7.5 log units. Only the latter value corresponds to a significant gain which is comparable with those observed in going from the ZnL to the ZnL_2 complexes of cysteine and cysteine ethyl ester. For all dipeptides 1–8 at least one ZnLH or ZnLOH complex is observed. They are pH-dependent variations of the ZnL species, and their existence underlines the preference for the 1:1 composition. The fact that the ZnLH species is observed for all four *N*-unprotected peptides and only for one *N*-protected peptide seems to indicate that the amino function is the protonated one in ZnLH.

Figure 4 depicts the stability relations between the eight ZnL complexes. The peptides being related only by exchanging His and Cys are connected by dotted lines. It is obvious that the free amino function (*N*-unprotected peptides) is of prime importance for high stabilities. It allows formation of the favorable five- or six-membered chelate rings with *N,S* or *N,N* coordination. In contrast, despite the negative charge provided by the carboxylate in peptides 3 and 4, the resulting six- and seven-membered chelate rings with *S,O* and *N,O* coordination provide significantly less stabilization. And finally the fully protected peptides 7 and 8 offer no favorable chelation and hence provide the lowest stabilities.

It should be noticed, however, that even the relatively low ZnL stabilities for 7 and 8 are still considerable. This becomes obvious by comparison with the log $\beta_{(\text{ZnL})}$ values for

Figure 4. Overall stability constants ($\log \beta$) of the ZnL complexes of peptides 1–8

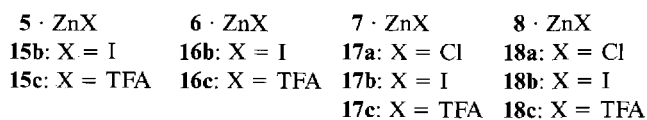
glycine (5.19^[11]) or *N*-acetylcysteine (4.90^[10]) which both can form chelate rings with one or two negative charges. This indicates that the 10-membered chelate rings generated by coordination of 7 or 8 must be relatively strain-free. Along the same line it should be noted that the ZnL complexes of 1 and 2 are the most stable zinc peptide complexes reported until today.

Compared to the dramatic decrease in complex stability upon blocking of the N terminus, the corresponding effect upon blocking of the C terminus is less pronounced. It amounts to $\log \beta$ changes between 0.5 (2/6) and 1.35 (3/7). It is strong enough, though, to point to the fact that in addition to the thiolate and imidazole functions the carboxylate function is also a coordinating unit in the *C*-unprotected peptides.

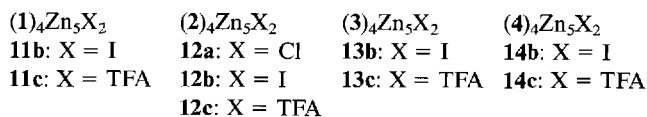
Isolated Complexes

The solution equilibrium studies revealed that the complex species ZnL exist in neutral or weakly basic solutions and that ZnL₂ complexes have a much smaller probability of formation. This was borne out by the preparative studies which failed to produce ZnL₂ compounds but yielded variations of the ZnL composition in all cases. The preparative procedure consisted in the combination of the zinc salt and the peptide, normally in ethanol, and inducing precipitation of the product by adding triethylamine as the reagent for the consumption of the equivalents of acid present. The stoichiometric ratio Zn/L was of little influence on the outcome of the reaction, but purer products were obtained by using a 1:1 ratio. ZnCl₂ and ZnI₂ were used as zinc salts of coordinating anions, Zn(ClO₄)₂ · 6 H₂O was meant to provide zinc ions without coordinating anions. It turned out, however, that the trifluoroacetate (TFA) introduced with the peptides, acted as the coordinating anion in this case.

Using the *C*-protected peptides 5–8, we found a simple verification of the ZnL composition by isolation of the complexes LZnX:



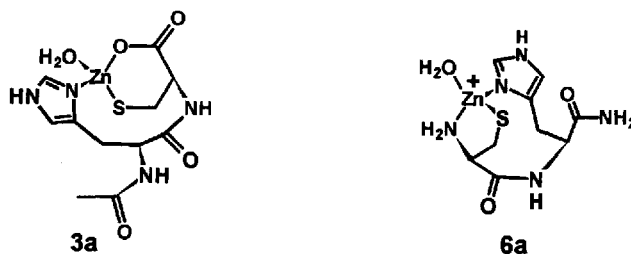
Using the *C*-unprotected peptides 1–4, we could expect products of the composition ZnL. However, in all cases the isolated complexes had the composition L₄Zn₅X₂ which can be written as 4 ZnL · ZnX₂:



All complexes 11–18 have a low solubility in all common solvents, thereby indicating their oligomeric nature. They could, however, be subjected to ¹H-NMR spectroscopy in hot DMSO (see Experimental) which confirmed the presence of the intact ligands. ¹⁹F-NMR spectroscopy confirmed the presence of trifluoroacetate in all complexes c. In the IR spectra (see Experimental) the S–H or O–H bands of the free peptides are missing, as expected. Assuming monodentate coordination by the anions, it can therefore be concluded that the peptides provide at least three donor functions in complexes LZnX and at least four donor functions for the ZnL constituents of complexes L₄Zn₅X₂. Attempts to crystallize any one of the isolated complexes for a structure determination remained unsuccessful.

Conclusions

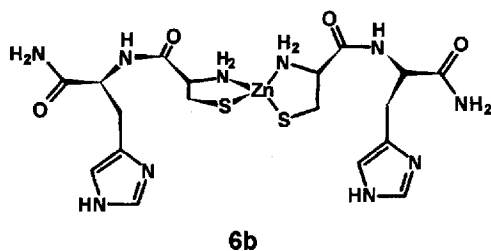
The high stabilities of the ZnL complexes in solution point to favorable chelating properties of the peptide ligands L, the low solubility of the isolated complexes points to their multidentate nature. This allows us to propose constitutional assignments with a reasonable degree of certainty. We propose tridentate coordination in dilute solution as exemplified by 3a and 6a containing 3 and 6 as ligands. Dilute solution conditions (1 mM) prevail during the potentiometric titrations, thereby reducing the probability of intermolecular coordination of the various donor functions and of thiolate bridging.



In concentrated solution or in the solid state oligomerization seems to occur. For this situation it is more difficult to make constitutional proposals due to the multitude of possibilities. We assume, however, that zinc is tetrahedrally coordinated in all cases because this is the standard situation for N/S ligand sets^[2,3]. For ligands 1, 2, 3, and 6 which can form favorable N,S or O,S chelate rings the ligation pattern in the oligomers should be just like it is in 3a and 6a except that the “dangling” donor provides the bridging function. For ligands 4, 5, 7, and 8 which do not favor terminal thiolate coordination due to chelation the possibility of thiolate bridging must be considered. This would in-

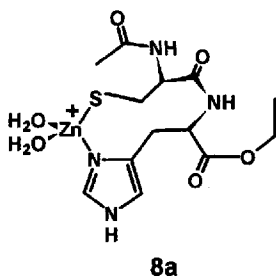
crease the number of donor functions per peptide by one, which could be accounted for by replacement of the ligating water molecules. Alternatively, with coordinating anions, less potent donor functions, e.g. carboxylate, might stay unused.

The preference of the peptides to be tridentate can also explain why ZnL_2 complexes are not favored. In going from a ZnL to a ZnL_2 complex either the first L has to be converted from tridentate to bidentate or the second L has to come in monodentate. There is only one favorable case for this, namely ZnL_2 with $L = H\text{-Cys-His-NH}_2$ (**6**). In this case ZnL_2 is a neutral ZnN_2S_2 complex with two five-ring chelate ligands, one of the most stable situations in zinc complex chemistry. The exceptionally high stability of the resulting complex **6b** confirms this.



We cannot offer a good explanation for the 4:5:2 composition of complexes **11–14** other than that they are 4:1 composites of ZnL and ZnX_2 . A neutral unit ZnL implies two negative charges on L which is easily realized for $L = 1–4$. It also requires the provision of four donor functions by L^{2-} which is no problem for **1** and **2** and which implies bidentate carboxylate or bridging thiolate for **3** and **4**. Why the stoichiometric ratio ZnL/ZnX_2 is persistently 4:1 in these compounds remains an open question.

Compared to the situation in proteins most of the peptides used here are still “unnatural”: **1–6** contain either terminal amino or terminal carboxylate functions which due to their ligating and chelating properties cause too high complex stabilities. Only **7** and **8** which form the least stable ZnL complexes possess the “natural” unsupported thiolate and imidazole donors. However, as pointed out above, the stabilities of their ZnL complexes are still considerable despite the presence of 10-membered chelate rings as in **8a**.



Thus, the dipeptides investigated here are still only milestones on the way to an understanding of zinc complexation by proteins. The favorable interaction between cysteine and histidine in the fully protected peptides **7** and **8** can, however, be taken as a preview of the preorganization of the

polypeptides causing the site specificity of zinc ion incorporation.

This work was supported by the *Deutsche Forschungsgemeinschaft* and the *Fonds der Chemischen Industrie*. We are grateful to Prof. A. Zuberbühler, Basel, for allowing us to use his TITFIT program. Mrs. F. Bitgöl and Mrs. D. Schoch helped with the preparative work.

Experimental

The general working and measuring methods are described in ref.^[16], the HPLC procedures in ref.^[14], the procedures for potentiometric measurements in ref.^[10]. The amino acid starting materials (optically pure L amino acids) and the zinc salts were obtained commercially.

The peptide syntheses started from histidine, cystine, histidine ethyl ester, and *N*-acetylcysteine. Standard protecting and coupling techniques^[17] were used to convert these to the immediate precursors of the peptides **1–8**, viz. the S–S dimers [Boc-His(Boc)-Cys-OEt]₂, (Ac-His-Cys-OEt)₂, (H-Cys-His-OMe)₂ and (Boc-Cys-His-NH₂)₂ and to Ac-Cys(Trt)-His-OEt. For the following procedures the abbreviations HTFA = trifluoroacetic acid and DTE = dithioerythritol are used. The HPLC eluents always contained 0.1% HTFA causing all peptides to be isolated as HTFA adducts. Their precise HTFA content was determined by potentiometric titration and confirmed by elemental analysis. All elemental analyses are summarized in Table 4.

Peptides 1: A solution of 5.67 g (5.84 mmol) of [Boc-His(Boc)-Cys-OEt]₂ in 50 ml of methanol was treated with 1.17 g (7.60 mmol) of DTE. The pH was adjusted to 8 with 0.2 M NaOH and the mixture stirred for 4 h. Then 50 ml of 1 M NaOH was added, and stirring was continued for 2 h. Then the mixture was acidified to pH 2 with 1 M HCl. The solvent was removed in vacuo, the residue taken up in 50 ml of methanol, the solution filtered and the filtrate evaporated to dryness again. The residue was treated five times with 10 ml of ether each in an ultrasound bath. The remaining colorless powder consisted of 4.01 g (96%) of Boc-His-Cys-OH which was spectroscopically pure. – 4.87 g (13.57 mmol) of Boc-His-Cys-OH was treated at 0°C with 16 ml of anhydrous HTFA. The mixture was stirred for 20 h during which time it was allowed to warm up to room temp. All volatile components were removed in vacuo. HPLC (water/acetonitrile, gradient 98/2→90/10) yielded 1.98 g (30%) of **1** · 2 HTFA as a colorless, very hygroscopic powder of m.p. 109°C.

2: A solution of 5.86 g (6.88 mmol) of (H-Cys-His-OMe)₂ · 4 HBr in 16 ml of methanol was treated with a solution of 0.25 g (10.8 mmol) of sodium in 10 ml of methanol. After 5 min the resulting precipitate was filtered and dried in vacuo. The remaining colorless powder consisted of 3.46 g (98%) of (H-Cys-His-OH)₂ which was spectroscopically pure. – A solution of 2.00 g (3.89 mmol) of (H-Cys-His-OH)₂ in 50 ml of water was treated with a solution of 1.37 g (8.90 mmol) of DTE in 20 ml of water. The pH was adjusted to 8.5 with 2 M NaOH, and the resulting solution was stirred under nitrogen for 4 h at room temp. The mixture was acidified to pH 3 with HTFA and freeze-dried. The resulting raw product was washed 3 times each with 10 ml of diethyl ether, then dissolved in 7 ml of water. HPLC (water/acetonitrile, 100:0, for 10 min followed by 80:20) yielded 0.68 g (68%) of **2** as a colorless powder of m.p. 68°C.

7: A solution of 3.35 g (3.80 mmol) of (Ac-His-Cys-OEt · HTFA)₂ in 30 ml of methanol was treated with 0.76 g (4.94 mmol) of DTE. The pH was adjusted to 8 with 0.2 M NaOH and the mixture stirred for 3 h at room temp. Then the mixture was acidi-

fied with HTFA to pH 2 and the solvent removed in vacuo. The remaining oil was dissolved in methanol, the solution filtered and the filtrate evaporated to dryness again. HPLC (water/acetonitrile, 85:15) yielded 2.56 g (65%) of $7 \cdot 1.5$ HTFA \cdot H₂O as a colorless, very hygroscopic powder of m.p. 89°C.

3: 1.39 g (2.68 mmol) of $7 \cdot 1.5$ HTFA \cdot H₂O was dissolved in a 1:1 mixture of 30 ml of methanol and 30 ml of 1 M NaOH and the solution stirred at room temp. for 4 h. Then the mixture was acidified to pH 3 with 1 M HCl and the solvent removed in vacuo. The residue was suspended in methanol, the suspension filtered and the filtrate evaporated to dryness again. HPLC (water/acetonitrile, 92:8) yielded 0.96 g (69%) of $3 \cdot 1.5$ HTFA \cdot EtOH as a colorless powder of m.p. 179°C.

8: 3.89 g (6.82 mmol) of Ac-Cys(Trt)-His-OEt was treated with 20 ml of ethanthiol, followed by 20 ml of HTFA. The mixture was stirred for 2.5 h, and then the solvent was removed in vacuo. The remaining solid was suspended in 50 ml of water and the suspension extracted 3 times each with 20 ml of diethyl ether. The combined organic layers were washed 3 times each with 20 ml of water and then dried in vacuo. HPLC (water/acetonitrile, 90:10) yielded 2.01 g (59%) of $8 \cdot 1.5$ HTFA as a colourless crystalline solid of m.p. 99°C.

4: 0.68 g (1.36 mmol) of Ac-Cys-His-OEt \cdot 1.5 HTFA was dissolved in a 1:1 mixture of 20 ml of methanol and 20 ml of 1 M NaOH and the solution stirred at room temp. for 5 h. Then the mixture was acidified to pH 3 with 1 M HCl and the solvent removed in vacuo. The residue was suspended in methanol, the suspension filtered and the filtrate evaporated to dryness again. HPLC (water/acetonitrile, 92:8) yielded 0.48 g (75%) of $4 \cdot 1.5$ HTFA as a colorless crystalline solid of m.p. 91°C.

5: 8.94 g (9.20 mmol) of [Boc-His(Boc)-Cys-OEt]₂ was treated with 25 ml of HTFA at 0°C. During 15 h the stirred mixture was allowed to warm to room temp. Then the solvent was removed in vacuo. The remaining oil was triturated with dry diethyl ether and then dried in vacuo. The remaining colorless powder consisted of 9.81 g (97%) of (H-His-Cys-OEt \cdot 2 HTFA)₂ which was spectroscopically pure. — A solution of 1.38 g (1.34 mmol) of (H-His-Cys-OEt \cdot 2 HTFA)₂ in 30 ml of methanol was treated with 0.27 g (1.75

mmol) of DTE. The pH was adjusted to 8 with 0.2 M NaOH, and the mixture was stirred for 4 h at room temp. It was then acidified with HTFA to pH 2 and the solvent removed in vacuo. The remaining solid was dissolved in methanol, the solution filtered and the filtrate evaporated to dryness again. HPLC (water/acetonitrile, 90/10) yielded 0.84 g (61%) of $5 \cdot 2.5$ HTFA as a colorless, very hygroscopic powder of m.p. 82–84°C.

6: 3.66 g (4.69 mmol) of (Z-Cys-His-NH₂)₂ (Z = benzyloxycarbonyl) was suspended in 15 ml of glacial acetic acid and the suspension treated with a solution of 15 ml of HBr in glacial acetic acid (33%). After 16 h the resulting precipitate was collected by filtration, washed 3 times each with 40 ml of diethyl ether and dried for 4 d with KOH. The resulting light yellow powder consisted of 3.61 g (92%) of (H-Cys-His-NH₂)₂ \cdot 4 HBr of m.p. 178°C. — A solution of 1.00 g (1.20 mmol) of (H-Cys-His-NH₂)₂ \cdot 4 HBr in 20 ml of water was treated with 0.23 g (1.45 mmol) of DTE. The pH was adjusted to 8.5 with 2 M NaOH, and the mixture was stirred for 4 h at room temp. It was then acidified with HTFA to pH 3, and the solvent was removed in vacuo. The remaining solid was washed 3 times each with 10 ml of diethyl ether. HPLC (water/acetonitrile, 100:0) yielded 0.41 g (70%) of $6 \cdot 2$ HTFA as a colorless powder of m.p. 76°C.

Relevant IR data of the peptides [$\nu(\text{OH})$, $\nu(\text{NH})$, $\nu(\text{SH})$, amide bands, in KBr, $\tilde{\nu}$ in cm⁻¹]: **1:** 3423, 3239, 3147, 2652, 1677, 1635, 1558, 1543. **2:** 3391, 3147, 3043, 2642, 1662, 1561, 1532. **3:** 3300, 3154, 3029, 2700, 1638, 1545. **4:** 3290, 3153, 3042, 2639, 1673, 1542. **5:** 3421, 3390, 3146, 3037, 2669, 1737, 1673, 1558, 1543. **6:** 3394, 3149, 3024, 2632, 1672, 1543. **7:** 3408, 3286, 3166, 3140, 3045, 2648, 1666, 1644, 1531. **8:** 3275, 3153, 3039, 2642, 1671, 1542.

¹H-NMR data of the peptides (calibrated on the residual proton resonance of the deuterated solvent): **1** (DMSO, 300 K): δ = 2.77–2.99 (m, 2H, H_β), 3.18–3.23 (m, 2H, H_β), 4.25–4.31 (m, 1H, H_α); 4.46–4.56 (m, 1H, H_α); 7.44 (s, 1H, 5-H), 8.84 (d, ³J = 7.5 Hz, 1H, NH), 8.97 (s, 1H, 2-H). **2** (D₂O, 300 K): δ = 3.13 (d, ³J = 6.0 Hz, 2H, H_βCys), 3.19 (dd, ³J = 8.7, ²J = 15.0 Hz, 1H, H_βHis), 3.45 (dd, ³J = 5.3, ²J = 15.0 Hz, 1H, H_βHis), 4.30 (t, ³J = 6.0 Hz, 1H, H_αCys), 4.82–4.90 (m, 1H, H_αHis), 7.41 (s, 1H, 5-H), 8.68 (d, ³J = 1.2 Hz, 1H, 2-H).

Table 3. Preparative details for the zinc complexes

Nr.	Peptide		EtOH ml	NEt ₃		X	Zinc salt ZnX ₂		EtOH ml	Nr.	Product	
	mg	mmol		mg	mmol		mg	mmol			mg	%
1	185	0.381	25	154	1.524	I	122	0.381	5	11b \cdot 2EtOH \cdot 2H ₂ O	105	79
1	185	0.381	25	154	1.524	ClO ₄ (\cdot 6H ₂ O)	142	0.381	5	11c \cdot 2EtOH \cdot 2H ₂ O	95	73
2	239	0.440	15	178	1.759	Cl	60	0.440	5	12a \cdot 2EtOH \cdot 3H ₂ O	126	91
2	239	0.440	15	178	1.759	I	140	0.440	5	12b \cdot 2EtOH \cdot 4H ₂ O	137	88
2	239	0.440	15	178	1.759	ClO ₄ (\cdot 6H ₂ O)	164	0.440	5	12c \cdot 2EtOH \cdot 4H ₂ O	137	89
3	122	0.236	15	60	0.589	I	75	0.236	10	13b \cdot 4H ₂ O	74	85
3	122	0.236	15	60	0.589	ClO ₄ (\cdot 6H ₂ O)	88	0.236	10	13c \cdot 4H ₂ O	73	85
4	67	0.142	10	36	0.355	I	45	0.142	5	14b \cdot 4H ₂ O	47	90
4	67	0.142	10	36	0.355	ClO ₄ (\cdot 6H ₂ O)	53	0.142	5	14c \cdot 4H ₂ O	39	76
5	180	0.314	20	111	1.099	I	100	0.314	10	15b	114	76
5	180	0.314	20	111	1.099	ClO ₄ (\cdot 6H ₂ O)	117	0.314	10	15c	117	77
6	62	0.114	15	40	0.400	I	37	0.114	10	16b	29	57
6	62	0.114	15	40	0.400	ClO ₄ (\cdot 6H ₂ O)	43	0.114	10	16c	31	63
7	180	0.348	20 [a]	35 [b]	0.870	Cl	47	0.348	10 [a]	17a	25	17
7	150	0.280	20	71	0.700	I	89	0.280	10	17b \cdot H ₂ O	97	64
7	150	0.280	20	71	0.700	ClO ₄ (\cdot 6H ₂ O)	104	0.280	10	17c \cdot H ₂ O	102	70
8	350	0.701	15 [a]	70 [c]	1.753	Cl	96	0.701	10 [a]	18a \cdot H ₂ O	247	79
8	210	0.421	15	106	1.051	I	134	0.421	10	18b	149	68
8	210	0.421	20	106	1.051	ClO ₄ (\cdot 6H ₂ O)	157	0.421	5	18c	118	56

[a] Water was used as solvent instead of EtOH. — [b] 0.2 M aqueous NaOH was used instead of NEt₃. — [c] 1 M aqueous NaOH was used instead of NEt₃.

3 (DMSO, 300 K): $\delta = 1.16$ (t, $^3J = 6.9$ Hz, 3H, CH₃ ethoxy), 1.82 (s, 3H, CH₃ acetyl), 2.70–3.10 (m, 4H, H _{β}), 3.09 (q, $^3J = 6.9$ Hz, 2H, CH₂ ethoxy), 4.35–4.45 (m, 1H, H _{α}), 4.60–4.71 (m, 1H, H _{α}), 7.33 (s, 1H, 5-H), 8.23 (d, $^3J = 8.3$ Hz, 1H, NH), 8.24 (d, $^3J = 7.7$ Hz, 1H, NH), 8.95 (s, 1H, 2-H).

4 (MeOH, 300 K): $\delta = 2.01$ (s, 3H, CH₃ acetyl), 2.71–2.92 (m, 2H, H _{β}), 3.11–3.41 (m, 2H, H _{β}), 4.36–4.42 (m, 1H, H _{α}), 4.73–4.80 (m, 1H, H _{α}), 7.35 (s, 1H, 5-H), 8.78 (s, 1H, 2-H).

5 (DMSO, 300 K): $\delta = 1.19$ (t, $^3J = 7.1$ Hz, 3H, CH₃ ethoxy), 2.80–2.88 (m, 2H, H _{β}), 3.13–3.21 (m, 2H, H _{β}), 4.16 (q, $^3J = 7.1$ Hz, 2H, CH₂ ethoxy), 4.23–4.29 (m, 1H, H _{α}), 4.51–4.60 (m, 1H, H _{α}), 7.38 (s, 1H, 5-H), 8.22 (d, $^3J = 16.6$ Hz, 1H, NH), 8.85 (s, 1H, 2-H).

6 (D₂O, 300 K): $\delta = 3.12$ –3.15 (m, 2H, C _{β} Cys), 3.28–3.34 (m, 2H, C _{β} His), 4.26–4.32 (m, 1H, C _{α} Cys), 4.74–4.82 (m, 1H, C _{α} His), 7.40 (s, 1H, 5-H), 8.68 (d, $^3J = 1.2$ Hz, 1H, 2-H).

Table 4. Analytical data

Compound	Formula (mol.mass)		Analyses			
			C	H	N	Zn
1•2HTFA	C ₉ H ₁₄ N ₄ O ₃ S•2C ₂ HF ₃ O ₂ (258.3 + 228.1)	calcd.	32.11	3.32	11.52	
		found	31.82	3.43	11.10	
2•2HTFA	C ₉ H ₁₄ N ₄ O ₃ S•2C ₂ HF ₃ O ₂ (258.3 + 228.1)	calcd.	32.11	3.32	11.52	
		found	31.42	3.37	11.21	
3•1.5HTFA•EtOH	C ₁₁ H ₁₆ N ₄ O ₄ S•1.5C ₂ HF ₃ O ₂ •C ₂ H ₆ O (300.3 + 171.0 + 46.1)	calcd.	37.14	4.58	10.83	
		found	37.63	4.49	11.03	
4•1.5HTFA	C ₁₁ H ₁₆ N ₄ O ₄ S•1.5C ₂ HF ₃ O ₂ (300.3 + 171.0)	calcd.	35.67	3.74	11.89	
		found	35.68	3.79	12.05	
5•2.5HTFA	C ₁₁ H ₁₈ N ₄ O ₃ S•2.5C ₂ HF ₃ O ₂ (286.4 + 285.1)	calcd.	33.63	3.62	9.81	
		found	33.16	3.63	9.29	
6•2HTFA	C ₉ H ₁₅ N ₅ O ₂ S•2C ₂ HF ₃ O ₂ (257.3 + 228.1)	calcd.	32.17	3.53	14.43	
		found	32.16	3.73	13.50	
7•1.5HTFA•H₂O	C ₁₃ H ₂₀ N ₄ O ₄ S•1.5C ₂ HF ₃ O ₂ •H ₂ O (328.4 + 171.0 + 18.0)	calcd.	37.14	4.58	10.83	
		found	37.39	4.41	10.00	
8•1.5HTFA	C ₁₃ H ₂₀ N ₄ O ₄ S•1.5C ₂ HF ₃ O ₂ (328.4 + 171.0)	calcd.	38.48	4.34	11.22	
		found	38.25	4.33	11.09	
11b•2EtOH•2H₂O	C ₄₀ H ₆₄ I ₂ N ₁₆ O ₁₆ S ₄ Zn ₅ (1734.0)	calcd.	27.71	3.72	12.92	18.85
		found	27.71	3.59	12.23	18.05
11c•2EtOH•2H₂O	C ₄₄ H ₆₄ F ₆ N ₁₆ O ₂₀ S ₄ Zn ₅ (1706.2)	calcd.	30.97	3.78	13.13	19.16
		found	30.73	4.03	13.16	19.29
12a•2EtOH•3H₂O	C ₄₀ H ₆₆ Cl ₂ N ₁₆ O ₁₇ S ₄ Zn ₅ (1569.1)	calcd.	30.62	4.24	14.28	20.83
		found	30.71	4.09	14.22	20.30
12b•2EtOH•4H₂O	C ₄₀ H ₆₈ I ₂ N ₁₆ O ₁₈ S ₄ Zn ₅ (1770.0)	calcd.	27.14	3.87	12.66	18.47
		found	27.19	3.45	12.11	18.11
12c•2EtOH•4H₂O	C ₄₄ H ₆₈ F ₆ N ₁₆ O ₂₂ S ₄ Zn ₅ (1742.3)	calcd.	30.33	3.93	12.86	18.76
		found	29.75	3.85	12.80	18.98
13b•4H₂O	C ₄₄ H ₆₄ I ₂ N ₁₆ O ₂₀ S ₄ Zn ₅ (1846.0)	calcd.	28.63	3.49	12.14	17.71
		found	28.33	3.24	11.45	17.40
13c•4H₂O	C ₄₈ H ₆₄ F ₆ N ₁₆ O ₂₄ S ₄ Zn ₅ (1818.3)	calcd.	31.71	3.55	12.33	17.98
		found	31.32	3.38	12.49	17.88
14b•4H₂O	C ₄₄ H ₆₄ I ₂ N ₁₆ O ₂₀ S ₄ Zn ₅ (1846.0)	calcd.	28.63	3.49	12.14	17.17
		found	28.94	3.28	11.96	17.33
14c•4H₂O	C ₄₈ H ₆₄ F ₆ N ₁₆ O ₂₄ S ₄ Zn ₅ (1818.3)	calcd.	31.71	3.55	12.33	17.98
		found	31.91	3.56	13.02	17.41
15b	C ₁₁ H ₁₇ IN ₄ O ₃ SZn (477.6)	calcd.	27.66	3.59	11.73	13.69
		found	27.71	3.40	10.84	13.99
15c•H₂O	C ₁₃ H ₁₉ F ₃ N ₄ O ₆ SZn (481.8)	calcd.	32.41	3.98	11.98	13.69
		found	31.79	4.00	11.92	13.88
16b	C ₉ H ₁₄ IN ₅ O ₂ SZn (448.6)	calcd.	24.10	3.15	15.61	14.57
		found	26.08	3.29	15.57	14.90
16c	C ₁₁ H ₁₄ F ₃ N ₅ O ₄ SZn (434.7)	calcd.	30.39	3.25	16.11	15.04
		found	28.57	3.50	16.81	14.90
17a	C ₁₃ H ₁₉ ClN ₄ O ₄ SZn (428.2)	calcd.	36.46	4.47	13.08	15.27
		found	37.02	4.45	12.47	15.66
17b•H₂O	C ₁₃ H ₂₁ IN ₄ O ₅ SZn (537.7)	calcd.	29.04	3.94	10.42	12.16
		found	29.57	3.61	10.08	12.50
17c•H₂O	C ₁₅ H ₂₁ F ₃ N ₄ O ₇ SZn (523.8)	calcd.	34.40	4.04	10.70	12.48
		found	34.87	4.06	11.48	12.78
18a•H₂O	C ₁₃ H ₂₁ ClN ₄ O ₅ SZn (446.2)	calcd.	34.99	4.74	12.56	14.65
		found	35.07	4.18	12.41	15.15
18b	C ₁₃ H ₁₉ N ₄ O ₄ SZn (519.7)	calcd.	30.05	3.69	10.78	12.58
		found	29.82	3.76	10.08	13.18
18c	C ₁₅ H ₁₉ F ₃ N ₄ O ₆ SZn (505.8)	calcd.	35.62	3.79	11.08	12.93
		found	35.54	4.04	11.45	13.24

7 (DMSO, 300 K): $\delta = 1.19$ (t, $^3J = 7.0$ Hz, 3H, CH₃ ethoxy), 1.84 (s, 3H, CH₃ acetyl), 2.73–3.18 (m, 4H, H _{β}), 4.11 (q, $^3J = 7.0$ Hz, 2H, CH₂ ethoxy), 4.40–4.50 (m, 1H, H _{α}), 4.61–4.73 (m, 1H, H _{α}), 7.34 (s, 1H, 5-H), 8.25 (d, $^3J = 7.7$ Hz, 1H, NH), 8.42 (d, $^3J = 7.1$ Hz, 1H, NH), 8.98 (s, 1H, 2-H).

8 (DMSO, 300 K): $\delta = 1.15$ (t, $^3J = 7.1$ Hz, 3H, CH₃ ethoxy), 1.87 (s, 3H, CH₃ acetyl), 2.54–2.81 (m, 2H, H _{β}), 3.01–3.24 (m, 2H, H _{β}), 4.09 (q, $^3J = 7.1$ Hz, 2H, CH₂ ethoxy), 4.28–4.38 (m, 1H, H _{α}), 4.53–4.62 (m, 1H, H _{α}), 7.39 (s, 1H, 5-H), 8.10 (d, $^3J = 7.9$ Hz, 1H, NH), 8.56 (d, $^3J = 7.8$ Hz, 1H, NH), 8.98 (s, 1H, 2-H).

Zinc Complexes. – General Preparative Procedure: Under nitrogen the peptide-HTFA adduct was dissolved in ethanol, and a 3- to 4-fold excess of triethylamine was added to the mixture followed by a solution of the zinc salt in ethanol. Upon stirring for 1–2 h a cloudy precipitate formed which was collected on a low-porosity paper filter and washed five times with cold ethanol. After drying for 3 d in vacuo, the resulting complex remained as a colorless, non-hygroscopic powder which was analytically pure and had a melting point around 200 °C. Table 3 gives the details for the preparations.

Relevant IR data of the complexes [$\nu(\text{NH})$, amide bands, in KBr, cm⁻¹]: **11b**: 3270, 3136, 1615, 1510. **11c**: 3381, 3292, 1650, 1615. **12a**: 3246, 3135, 1673, 1616, 1525. **12b**: 3412, 3252, 3136, 1673, 1620, 1551. **12c**: 3258, 3138, 1674, 1619, 1508. **13b**: 3303, 3140, 1645, 1524. **13c**: 3303, 3149, 1650, 1638, 1531. **14b**: 3300, 3137, 1639, 1528. **14c**: 3386, 3134, 1642, 1531. **15b**: 3273, 3136, 1672, 1580, 1538. **15c**: 3305, 3154, 1674, 1583, 1542. **16b**: 3412, 3266, 1671, 1560, 1541. **16c**: 3418, 3307, 1639, 1602, 1528. **17a**: 3307, 3110, 1656, 1521. **17b**: 3507, 3307, 1656, 1524. **17c**: 3504, 3306, 3139, 1661, 1530. **18a**: 3307, 1655, 1531. **18b**: 3324, 1655, 1639, 1531, 1525. **18c**: 3316, 3149, 1657, 1531.

¹H-NMR data of the complexes (calibrated on the residual proton resonance of the deuterated solvent) (the complexes **11b**, **11c**, **12a**, **12b** and **12c** were not soluble in any deuterated solvent): **13b** (DMSO, 350 K): $\delta = 1.86$ (s, 3H, CH₃ acetyl), 2.75–3.17 (m, 2 · 2H, H _{β}), 4.08–4.24 (m, 1H, H _{α}), 4.42–4.62 (m, 1H, H _{α}), 7.08 (s, 1H, 5-H), 7.70–7.88 (m, 2 · 1H, NH), 8.05 (s, 1H, 2-H).

13c (DMSO, 350 K): $\delta = 1.86$ (s, 3H, CH₃ acetyl), 2.75–3.17 (m, 2 · 2H, H _{β}), 4.03–4.22 (m, 1H, H _{α}), 4.41–4.68 (m, 1H, H _{α}), 7.09 (s, 1H, 5-H), 7.71–7.90 (m, 2 · 1H, NH), 8.07 (s, 1H, 2-H).

14b (DMSO, 350 K): $\delta = 1.86$ (s, 3H, CH₃ acetyl), 2.59–2.73 (m, 2H, H _{β}), 3.10–3.51 (m, 2H, H _{β}), 4.20–4.35 (m, 1H, H _{α}), 4.40–4.61 (m, 1H, H _{α}), 6.96–7.02 (m, 1H, 5-H), 7.53–7.61 (m, 1H, NH), 7.76–7.84 (m, 1H, NH), 7.91–7.98 (m, 1H, 2-H).

14c (DMSO, 350 K): $\delta = 1.86$ (s, 3H, CH₃ acetyl), 2.61–2.73 (m, 2H, H _{β}), 2.92–3.21 (m, 2H, H _{β}), 4.19–4.31 (m, 1H, H _{α}), 4.40–4.60 (m, 1H, H _{α}), 6.96–7.01 (m, 1H, 5-H), 7.58–7.66 (m, 1H, NH), 7.76–7.84 (m, 1H, NH), 7.92–7.96 (m, 1H, 2-H).

15b (DMSO, 350 K): $\delta = 1.18$ (t, $^3J = 7.3$ Hz, 3H, CH₃ ethoxy), 2.58–3.15 (m, 2 · 2H, H _{β}), 3.98–4.12 (m, 3H, CH₂ ethoxy and H _{α}), 4.29–4.49 (m, 1H, H _{α}), 7.08 (s, 1H, 5-H), 7.93 (s, 1H, 2-H).

15c (DMSO, 350 K): $\delta = 1.19$ (t, $^3J = 6.4$ Hz, 3H, CH₃ ethoxy), 2.60–3.12 (m, 2 · 2H, H _{β}), 3.87–4.18 (m, 3H, CH₂ ethoxy and H _{α}), 4.25–4.41 (m, 1H, H _{α}), 7.05 (s, 1H, 5-H), 7.90 (s, 1H, 2-H).

16b (DMSO, 350 K): $\delta = 2.57$ –3.18 (m, 2 · 2H, H _{β}), 4.31–4.60 (m, 2 · 1H, H _{α}), 6.97 (s, 1H, 5-H), 7.02–7.12 (m, 1H, NH), 7.91 (s, 1H, 2-H).

16c (DMSO, 350 K): $\delta = 2.49$ –3.17 (m, 2 · 2H, H _{β}), 4.25–4.66 (m, 2 · 1H, H _{α}), 6.97 (s, 1H, 5-H), 7.02–7.11 (m, 1H, NH), 7.91 (s, 1H, 2-H).

17a (DMSO, 350 K): $\delta = 1.16$ (t, $^3J = 6.6$ Hz, 3H, CH₃ ethoxy), 1.85 (s, 3H, CH₃ acetyl), 2.62–2.97 (m, 2 · 2H, H _{β}), 4.06 (q, $^3J = 6.6$ Hz, 2H, CH₂ ethoxy), 4.12–4.64 (m, 2 · 1H, H _{α}), 6.91 (s, 1H,

5-H), 7.56–7.64 (m, 1H, NH), 7.68 (s, 1H, 2-H), 7.77–7.86 (m, 1H, NH).

17b (DMSO, 350 K): $\delta = 1.17$ (t, $^3J = 7.1$ Hz, 3H, CH₃ ethoxy), 1.85 (s, 3H, CH₃ acetyl), 2.78–3.18 (m, 2 · 2H, H _{β}), 4.07 (q, $^3J = 7.1$ Hz, 2H, CH₂ ethoxy), 4.17–4.28 (m, 1H, H _{α}), 4.37–4.52 (m, 1H, H _{α}), 7.03 (s, 1H, 5-H), 7.65–7.75 (m, 1H, NH), 7.85 (s, 1H, 2-H), 7.85–7.94 (m, 1H, NH).

17c (DMSO, 350 K): $\delta = 1.16$ (t, $^3J = 7.0$ Hz, 3H, CH₃ ethoxy), 1.86 (s, 3H, CH₃ acetyl), 2.78–3.14 (m, 2 · 2H, H _{β}), 4.06 (q, $^3J = 7.0$ Hz, 2H, CH₂ ethoxy), 4.13–4.22 (m, 1H, H _{α}), 4.33–4.48 (m, 1H, H _{α}), 7.02 (s, 1H, 5-H), 7.69–7.78 (m, 1H, NH), 7.83 (s, 1H, 2-H), 7.83–7.90 (m, 1H, NH).

18a (DMSO, 350 K): $\delta = 1.17$ (t, $^3J = 7.1$ Hz, 3H, CH₃ ethoxy), 1.86 (s, 3H, CH₃ acetyl), 2.60–2.71 (m, 2H, H _{β}), 3.01–3.20 (m, 2H, H _{β}), 4.08 (q, $^3J = 7.1$ Hz, 2H, CH₂ ethoxy), 4.25–4.49 (m, 2 · 1H, H _{α}), 7.10–7.30 (m, 2 · 1H, 5-H, NH), 7.64–7.74 (m, 1H, NH), 7.88 (s, 1H, 2-H).

18b (DMSO, 300 K): $\delta = 1.17$ (t, $^3J = 7.2$ Hz, 3H, CH₃ ethoxy), 1.85 (s, 3H, CH₃ acetyl), 2.60–2.80 (m, 2H, H _{β}), 3.04–3.16 (m, 2H, H _{β}), 4.08 (q, $^3J = 7.2$ Hz, 2H, CH₂ ethoxy), 4.14–4.34 (m, 2 · 1H, H _{α}), 7.13 (s, 1H, 5-H), 7.66 (d, $^3J = 6.3$ Hz, 1H, NH), 7.97 (s, 1H, 2-H), 8.05 (d, $^3J = 9.1$ Hz, 1H, NH).

18c (DMSO, 300 K): $\delta = 1.16$ (t, $^3J = 7.1$ Hz, 3H, CH₃ ethoxy), 1.86 (s, 3H, CH₃ acetyl), 2.59–2.78 (m, 2H, H _{β}), 3.00–3.20 (m, 2H, H _{β}), 4.08 (q, $^3J = 7.1$ Hz, 2H, CH₂ ethoxy), 4.20–4.41 (m, 2 · 1H, H _{α}), 7.13 (s, 1H, 5-H), 7.67 (d, $^3J = 6.9$ Hz, 1H, NH), 7.97 (s, 1H, 2-H), 8.02–8.10 (m, 1H, NH).

* Dedicated to Professor Max Herberhold on the occasion of his 60th birthday.

- [1] M. Förster, I. Brasack, A. K. Duhme, H. F. Nolting, H. Vahrenkamp, *Chem. Ber.* **1996**, *129*, 347–353.
- [2] *Zinc Enzymes* (Eds.: I. Bertini, C. Luchinat, W. Maret, M. Zeppezauer), Birkhäuser, Boston, **1986**.
- [3] R. H. Prince in *Comprehensive Coordination Chemistry* (Eds.: G. Wilkinson, R. D. Gillard, J. A. McCleverty), Pergamon Press, Oxford, **1987**, vol. 5, pp. 925–1045.
- [4] *Biocoordination Chemistry: Coordination Equilibria in Biologically Active Systems* (Ed.: K. Burger), Ellis Horwood, New York, **1990**.
- [5] D. Rhodes, A. Klug, *Trends Biol. Sci.* **1987**, *12*, 464–469.
- [6] N. P. Pavletich, C. O. Pabo, *Science* **1991**, *252*, 809–817.
- [7] G. Weitzel, F. Schneider, A. M. Fretzdorff, *Hoppe Seylers Z. Physiol. Chem.* **1957**, *307*, 23–35.
- [8] H. Shindo, T. L. Brown, *J. Am. Chem. Soc.* **1965**, *87*, 1904–1909.
- [9] H. C. Freeman in *Inorganic Biochemistry* (Ed.: G. L. Eichhorn), Elsevier, Amsterdam, **1973**, vol. 1, pp. 121–166.
- [10] P. Gockel, H. Vahrenkamp, A. Zuberbühler, *Helv. Chim. Acta* **1993**, *76*, 511–520, and references cited therein.
- [11] A. E. Martell, R. M. Smith, *Critical Stability Constants*, Plenum Press, New York, 6 volumes since **1974**.
- [12] D. L. Rabenstein, R. Guevremont, C. A. Evans in *Amino Acids and Derivatives as Ambivalent Ligands* (Ed.: H. Sigel), Marcel Dekker, New York, **1979**, pp. 103–141.

- [13] H. Albrich, H. Vahrenkamp, *Chem. Ber.* **1994**, *127*, 1223–1233; A. Meißner, P. Gockel, H. Vahrenkamp, *ibid.* **1994**, *127*, 1235–1241, and references cited therein.
- [14] M. Förster, H. Vahrenkamp, *Chem. Ber.* **1995**, *128*, 541–550, and references cited therein.
- [15] S. Ranganathan, S. Jayaraman, R. Roy, *Tetrahedron* **1992**, *48*, 931–938.
- [16] M. Förster, R. Burth, A. K. Powell, T. Eiche, H. Vahrenkamp, *Chem. Ber.* **1993**, *126*, 2643–2648.
- [17] M. Bodanszky, *Principles of Peptide Synthesis*, Springer, Berlin, **1984**; M. Bodanszky, A. Bodanszky, *The Practice of Peptide Synthesis*, Springer, Berlin, **1984**. [96057]