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Pentacoordinate and hexacoordinate ferric hemes in acid medium: EPR, UV–Vis and CD studies of the giant extracellular hemoglobin of *Glossoscolex paulistus*

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Abstract

The equilibrium complexity involving different axially coordinated hemes is peculiar to hemoglobins. The pH dependence of the spontaneous exchange of ligands in the extracellular hemoglobin from *Glossoscolex paulistus* was studied using UV–Vis, EPR, and CD spectroscopies. This protein has a complex oligomeric assembly with molecular weight of 3.1 MDa that presents an important cooperative effect. A complex coexistence of different species was observed in almost all pH values, except pH 7.0, where just aquomet species is present. Four new species were formed and coexist with the aquomethemoglobin upon acidification: (i) a "pure" low-spin hemichrome (Type II), also called hemichrome B, with an usual spin state $(d_{xy})^2(d_{xz},d_{yz})^3$; (iii) a strong g_{max} hemichrome (Type I), also showing an usual spin state $(d_{xy})^2(d_{xz},d_{yz})^3$; (iii) a hemichrome with unusual spin state $(d_{xz},d_{yz})^4(d_{xy})^1$ (Type III); (iv) and a high-spin pentacoordinate species. CD measurements suggest that the mechanism of species formation could be related with an initial process of acid denaturation. However, it is worth mentioning that based on EPR the aquomet species remains even at acidic pH, indicating that the transitions are not complete. The "pure" low-spin hemichrome presents a parallel orientation of the imidazole ring planes but the strong g_{max} hemichrome is a *HALS* (highly anisotropic low-spin) species indicating a reciprocally perpendicular orientation of the imidazole ring planes. The hemichromes and pentacoordinate formation mechanisms are discussed in detail. © 2006 Elsevier B.V. All rights reserved.

Keywords: Hemoglobin; pH; Pentacoordinate; Hexacoordinate; Hemichrome

1. Introduction

Large extracellular hemoglobins have been employed as potential blood substitutes [1]. Strand et al. [2] have proposed that erythrocruorins are useful model systems for developing therapeutic blood substitutes due to their extracellular nature, large size and resistance to oxidation. Hirsch et al. [1] have suggested that the *Lumbricus terrestris* hemoglobin in the ferrous form, at neutral pH, exhibits similar oxygen affinity and cooperativity to that of human hemoglobin. Other advantages to use this hemoglobin class are its natural character and structural stability. It also shows low probability to promote immunogenic

responses since cell membranes are not present and, unlike tetrameric hemoglobins, it does not undergo subunit dissociation upon dilution [3,4]. Recently, Vinogradov [5,6] argued that giant extracellular hemoglobins represent a summit of complexity for oxygen-binding heme protein.

In the present work, the behavior of the extracellular methemoglobin of the annelid *Glossoscolex paulistus* (HbGp)

methemoglobin of the annelid *Glossoscolex paulistus* (HbGp) as a function of pH has been evaluated. This worm is prevalent in sites near the cities of Piracicaba, Araras and Rio Claro in the state of São Paulo, Brazil [7]. Its hemoglobin is a giant biopolymer that shows a molecular mass of 3.1 MDa [8] and a complex assembly with a hexagonal double-layered (HBL) oligomeric structure [9,10] that dissociates in alkaline and acid mediums [11,12]. This hemoprotein shows interesting features regarding natural selection, such as its adaptation in sulfide-rich environments [10,13].

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G. paulistus hemoglobin belongs to the same class of hemoglobin as L. terrestris, which is one of the most studied hemoglobins [14–20], thus showing a highly cooperative oxygen binding process [14,21]. It consists of an arrangement of two heme-containing subunits (monomer and disulfide-bonded trimer T) with molecular mass in the range 15–19 kDa and non-globin linker subunits of 24 to 32 kDa [10,14]. Some studies suggest that the linker chains present not only structural functions but also superoxide dismutase activity [22]. A common model for the quaternary structure, so-called "bracelet model", has been employed to explain its assembly [14].

Many studies about equilibrium between different coordination states of hemoglobins have been developed. The equilibrium of heme proteins involves the coexistence of many species depending on medium conditions [23-25]. In fact, changing pH and ionic strength leads to modifications in the first coordination sphere. These induce alterations of the spin state, ligand orientation, and process of exchange of iron ligands. Furthermore, there are species with different number of coordination, i.e., the coexistence of hexacoordinate, pentacoordinate [24], and tetracoordinate [26,27] forms are possible. It has been described that below pH 4 some heme proteins, such as myoglobin and horseradish peroxidase, lose the sixth ligand, thus converting the heme prosthetic groups to water ligated pentacoordinate [28]. In fact, one of the most significant developments in porphyrin chemistry over the last 20 or so years has been the understanding of the extraordinary flexibility of the porphyrin ring [29], that is associated with heme environment, including axial ligands and spin state of the ferric ion. In heme proteins, asymmetric distortions of the porphyrin macrocycle are induced by axial ligands, asymmetric peripheral substituents, and the anisotropic protein environment [30].

In several recent reports, the hemichrome formation has been correlated to the biological function of tetrameric hemoglobins. On the one hand, Robinson et al. [31], studying the pH-dependent aquomet-to-hemichrome transition of horse methemoglobin in acid medium, have proposed that *in vivo* hemichrome formation is related to a more solvent-exposed heme group. The appearance of bis-histidine in both ferric and ferrous forms of neuroglobin and cytoglobin has also been explained by solvent channel formation [32,33].

The modification mechanism of the first coordination sphere is very relevant in the research of heme proteins. There is a strong correlation between the polypeptide assembly and the ligand properties of the ferric ion. Hence, the conformation changes of the polypeptidic chains can alter the ligands of the ferric center. For example, the pentacoordinate species formation can be associated with the first step of an unfolding process [26,27].

The presence of a pentacoordinate species can have a general significance to all native proteins, not only to myoglobins and hemoglobins [34]. Viola et al. [35], for example, suggest that cytochrome c shows an interplay between redox, structural, ligand binding, and recognition properties. In fact, various metalloporphyrin complexes are pentacoordinate, resembling to the pentacoordinate heme proteins [36,37].

The aim of the present work is to contribute in the understanding of the structure-activity relationship of heme

proteins, specially of the *G. paulistus* giant hemoglobin, characterizing the species in the first coordination sphere of the iron as a function of pH. This work focuses on the transitions of the extracellular hemoglobin in the acidic medium. UV–Vis and electron paramagnetic resonance (EPR) were used to monitor modifications in the spin state as well as in the coordination environment at the metallic center, and circular dichroism (CD) to follow changes in the protein folding upon acidification.

2. Methods and materials

2.1. Protein preparation

The hemoglobin of *G. paulistus* was prepared using freshly drawn blood from worms. The blood sample was purified by ultra centrifugation and gel filtration in Sephadex G-200 column at pH 7.0 [11].

2.2. UV-Vis measurements

Electronic absorption spectra in the wavelength range of 250 to 700 nm were obtained as a function of pH, using a SHIMADZU UV-1601 PC spectrophotometer at room temperature. The absorbance of the samples at 415 nm was kept below 1.0 using a 1 cm path length cell, with concentration approximately 0.2 mg ml⁻¹ based on ε_{415} =3.7 ml mg⁻¹ cm⁻¹ at pH 7.0 [38].

2.3. EPR measurements

X-band (9.5 GHz) EPR spectra were measured on a Brucker Elexsys E580 spectrometer at 4 and 12 K. The temperature was controlled by an Oxford ITC 503 cryogenic system. EPR samples (50 μl) containing approximately 20 mg ml⁻¹ of the protein were frozen by immersion in liquid nitrogen and then placed in the spectrometer rectangular cavity. The microwave power was 4.0 mW and other acquisition conditions, such as modulation amplitude were adjusted to achieve optimal signal-to-noise ratio without signal distortion or saturation. All EPR data were corrected by subtracting a baseline corresponding to the EPR signal of the buffer.

2.4. CD measurements

CD spectra were obtained in a Jasco J715 (Jasco Co., Japan) as a function of pH employing the 0.2 cm path length cylindrical cell. Protein concentration was 0.22 mg ml⁻¹ based on ε_{415} =3.13 ml mg⁻¹ cm⁻¹ [39]. The residual ellipticity, [θ] in mdegree dmol⁻¹ cm², was obtained using the amino acid residue average molecular mass of 117 g mol⁻¹.

3. Results and discussion

3.1. UV-Vis measurements

UV-Vis measurements at pH 7.0 resulted in typical spectrum of a single aquomet species. This species is found in

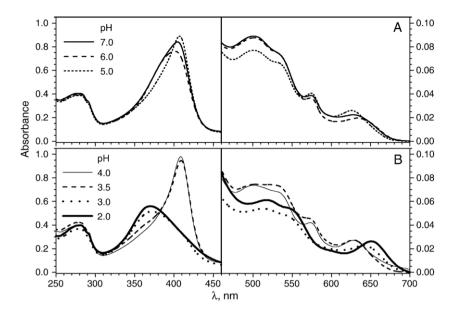


Fig. 1. Absorbance spectra of *Glossoscolex paulistus* hemoglobin. Conditions: 20 mM Tris–HCl buffer; GpHb concentration was 0.2 mg ml⁻¹. (A) pH 7.0, 6.0 and 5.0; (B) pH 4.0, 3.5, 3.0 and 2.0.

a great number of hemoglobins with similar bands [40–44] and is identified by a Soret band at 405 nm, a Q band at 500 nm and an LMCT band at 630 nm (Fig. 1).

The decrease in pH values from pH 6.0 to pH 3.0 causes an absorbance increase and a small red shift of the Soret band simultaneously, indicating the hemichrome formation, i.e., a bis-histidine complex formation, which is characterized by Q bands at 532 and 574 nm [44–47]. Therefore, this spectral change is associated with the distal histidine coordination to the ferric ion. The movement of the distal histidine imidazole towards the metal center is probably the result of a small change in the globin arrangement in the heme pocket.

A drastic spectral transition can be observed around pH 3.0. In fact, a Soret band blue shift relative to the pH 7.0 spectrum can be verified with a simultaneous decrease of absorbance, which indicate the formation of a pentacoordinate species as previously observed for several recent works [26,27,41–43,48–73].

Moreover, the pronounced Soret band shift to 371 nm occurs probably due to an initial unfolding process [42,43,59]. The very broad and asymmetric Soret band may be caused by the coexistence of two pentacoordinated species, i.e., a species with proximal histidine as fifth ligand and a pentacoordinated free hemin with water as fifth ligand [74].

Harris and Loew [75], using a semiempirical quantum chemical INDO/ROHF/CI method, calculated the electronic structure of cytochrome P450 in order to explain the optical spectra of the high- and low-spin states. The comparison of the calculated spectra of the two spin states reveals that, upon change of the Fe(III) from a low- to a high-spin state, a blue shift does indeed occur in the Soret band as observed for HbGp at pH lower than 3.5 (Fig. 1). They interpreted this shift as a direct result of the spin-state change. In the low-spin state, there is an enhanced mixing of the iron d orbital (eg) and porphyrin 4eg ($\pi*$) orbitals, which results in lower energy eg ($\pi*$) states. This

occurs because the higher ligand field of the low-spin state increases the energy of the *eg* orbitals, allowing higher interaction between these metallic orbitals and the porphyrin orbitals. Neya et al. [76] admitted to offer a similar explanation for myoglobins, i.e., the shorter wavelength would be associated with the high-spin state, while longer wavelength would be related to low-spin state.

The pentacoordinated species formation can also be confirmed by the analysis of the LMCT (Ligand-to-Metal Charge Transfer) band (Fig. 1). This band shows a red shift in the same pH range that causes the Soret band blue shift. Recent results showed that an LMCT band between 600 and 650 nm indicates a high-spin species [77,78]. Pentacoordinated high-spin hemes have been assigned to LMCT bands at 645 nm [50,59,61] or, at least, near to 645 nm [26,56, 62,79].

The UV-Vis data can be rationalized in terms of a mechanism for the transition upon acidification as follows. The stability of the aquomet species at pH 7.0 is provided by the presence of a water molecule in the nonpolar pocket, which is stabilized by binding to the heme iron [80]. Thus, the stability of the aquomet species can be explained mainly by two factors: the hydrophobicity of the heme pocket and the strong hydrogen bond between the water ligand and the distal histidine.

The aquomet-hemichrome transition must be due to the protonation of the nitrogen of the distal histidine, which decreases the ability of the histidine imidazole to stabilize the coordinated water molecule. Thus, the water molecule can easily dissociate from the first coordination sphere of the iron. Therefore, the absence of the hydrogen bond between the imidazole and the water molecule could be considered the initial step of the hemichrome formation [46,81].

The hemichrome-pentacoordinated transition can be originated by a competition between the ferric center and the protons for the imine-nitrogen of the imidazole. After imidazole protonation, the iron is shifted from the porphyrin nitrogen

plane towards the fifth ligand. The distal histidine is probably the displaced residue due to the higher tension of the distal coordination compared to the proximal site.

Besides that Boffi et al. reported that a strong σ-donor character of the fifth ligand decreases the coordination of other strong σ -donor in the sixth ligand position [74]. This fact occurs due to the weakening of the Fe-N bond of the sixth ligand by occupation of the d_z^2 antibonding orbital in the highspin configuration, thus increasing electronic repulsion, and precluding a more intense superposition between the orbitals in the σ -bond [40]. Furthermore, the π -donor character of the imidazole to the d^5 metallic center is known in ferric porphyrin complexes and heme proteins [82] as well as in non-porphyrins complexes [83,84], such as tetraammineruthenium(III) complexes [85-87]. This trans-influence must increase the labilization caused by σ -bond. Therefore, the trans-influence caused by σ -bond as well as by π -bond must generate the loss of the sixth coordination, because these two ligations would be weaker, favoring the pentacoordinate species formation, which is a more stable species in these conditions. Corroborating this analysis, other works also confirm this π -donor ability of imidazole ligand [88,89] and other N-heterocyclic ligands [90].

This mechanism is similar to the one occurring *in vivo* during ferrous pentacoordinate formation, indicating a possible correlation between this process and the protein structure. In this way, the movement of the metallic center to the out-of-plane arrangement in the pentacoordinated species yields a *domed* conformation, which is very important to several functions of heme proteins [91].

3.2. EPR measurements

The EPR spectrum at pH 7.0 is characteristic of an axially symmetric high-spin species with g_{\parallel} and g_{\perp} around 2.0 and 6.1, respectively (Fig. 2). In this case, only one component is observed, indicating the existence of one high-spin configuration assigned to the aquomet species. The acidification

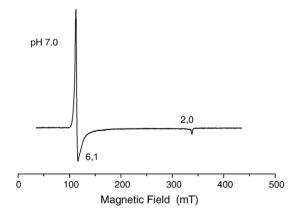


Fig. 2. Electron paramagnetic resonance spectrum of *Glossoscolex paulistus* hemoglobin with the *g*-values indicated. Conditions: 20 mM Tris—HCl buffer at pH 7.0 (T=4 K). EPR conditions: modulation amplitude, 0.1 mT; modulation frequency, 100 kHz; microwave power, 4.0 mW; microwave frequency, 9.4784 GHz.

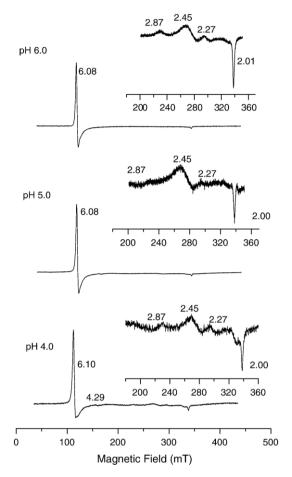


Fig. 3. Electron paramagnetic resonance spectra of *Glossoscolex paulistus* hemoglobin with the *g*-values indicated. Conditions: 20 mM Tris—HCl buffer; GpHb at pH $6.0~(T=4~\rm K$ showing insert with $T=12~\rm K$), GpHb at pH $5.0~(T=4~\rm K$ showing insert with $T=12~\rm K$), GpHb at pH $4.0~(T=4~\rm K$ showing insert with $T=12~\rm K$). EPR conditions: modulation amplitude, $0.1~\rm mT$; modulation frequency, $100~\rm kHz$; microwave power, $4.0~\rm mW$; microwave frequency, $9.4780~\rm GHz$.

causes the appearance of several species, whose spectra are shown in Figs. 3 and 4 as a function of the pH value. Overall, three low-spin species are formed in the spectra (see $g \sim 2$ region of the spectrum—insert of Fig. 3), which are assigned to hemichromes, i.e., bis-histidine complexes. In hemoproteins, it is well established that bis-histidine hemes having two axial histidine ligands display similar EPR spectra with g_z values between 2.9 and 3.6. The g-values of these complexes depend on the relative orientation of the two imidazole rings and on the orientation of these imidazoles with respect to the porphyrin plane [92,93]. These hemichromes show different electronic configurations: two of them are in the more usual spin state $(d_{xy})^2(d_{xz},d_{yz})^3$, and are characterized by g-values at 2.87 and 2.27 (Fig. 3) and by g-values around 3.47 (Fig. 4); the third hemichrome presents an unusual electronic configuration $(d_{xz}, d_{yz})^4 (d_{xy})^1$ with a resonance at 2.45 (Figs. 3 and 4). The g=4.3 line in Fig. 4 has been observed in other ferric proteins and it has been assigned to non-heme iron impurities [94,95]. Consequently, it will not be further considered in our discussion.

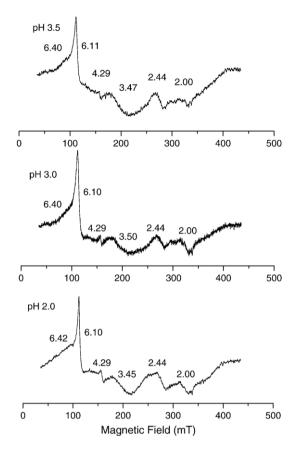


Fig. 4. Electron paramagnetic resonance spectra of *Glossoscolex paulistus* hemoglobin with the g-values indicated. Conditions: 20 mM Tris—HCl buffer. EPR conditions: T=12 K, modulation amplitude, 0.1 mT; modulation frequency, 100 kHz; microwave power, 4.0 mW; microwave frequency, 9.4794 GHz.

Between the hemichromes with more usual electronic configuration, the first species (g-values 2.87 and 2.27) presents mutually parallel imidazole ring orientation (Type II or Bhemichrome) [96], whereas in the second species (g-values around 3.47) those rings are in an orthogonal conformation [Type I hemichrome or *HALS* (highly anisotropic low-spin)] [96–103]. It is worth mentioning that the distinction of these hemichromes was possible only through EPR spectroscopy. The reason is the small difference of energy between these configurations [104,105]. In fact, Zaric et al. [104] investigated the factors determining conformations of imidazole axially coordinated to heme in heme proteins by analyzing 693 hemes in 432 different crystal structures of heme proteins from the Protein Data Bank (PDB). In this expressive collection, 65 were bis-histidine-ligated hemes and the authors explain that the energy balance between the two forms with usual spin state is the result of crystal field stabilization effects favoring the parallel form and steric effects that favor the perpendicular form. In heme model systems, the orientation of axial ligands can depend on crystal field stabilization effects or on steric effects caused by substituents on axial ligands and on the porphyrin. In heme proteins, the heme does not possess bulky substituents, but the protein environment can have a steric influence on the orientation of the axial ligands.

Medakovic and Zaric [105] performed quantum chemical (DFT) calculations on heme model systems with non-substituted Fe-porphyrin core for the different orientations of the axially coordinated imidazoles. Their results indicate that perpendicular orientation (Type I hemichrome) can be explained by steric effects caused by propionic groups of porphyrin ring and by the histidine backbone.

McGarvey [106] has described a more detailed explanation for the HALS (Type I) hemichrome formation considering that imidazole planes, that were initially in a parallel arrangement, adopt a perpendicular configuration, which is similar to a tetragonal distortion. In this way, the heme avoids a classic Jahn-Teller effect, with the axial ligands preferring the perpendicular orientation, which shows smaller orbitals overlap with the iron orbitals than the parallel orientation, thus causing a smaller splitting of iron d_{π} orbitals. Rieger [107] has reported that Type I hemichrome presents an apparent near-degeneracy of the d_{xz} and d_{yz} orbitals that is consistent with the crystal structure, which showed that the two ligands lie in perpendicular planes because of the similar orbitals overlap to d_{xz} and d_{yz} .

The g_{\perp} -value of 2.45 suggests the unusual configuration $(d_{xz},d_{yz})^4(d_{xy})^1$ for the heme-iron [96,108,109], also called Type III hemichrome, that probably occurs because of a great influence of the globin, causing a large distortion of the porphyrin ring. Interactions between the protein backbone and the porphyrin may be responsible for the macrocycle distortions from planarity *in vivo* that could modulate their physical and chemical properties [109]. This porphyrin distortion modifies the superposition of the d_{π} orbitals with the axial ligand orbitals, in such a way that the d_{xy} orbitals become the HOMO. The g_{\parallel} of this spectrum is not observed, but it would be between the g-values of 0.9 and 1.95 [96]. In agreement with Walker [96], this species can be identified as a Type III hemichrome.

After further acidification, an unexpected signal with g greater than 3.2 [96,110] was observed (Fig. 4). Some authors have analyzed this signal in other hemoglobins and ferric complexes [77]. The so-called hemichrome with strong g_{max} (Type I), i.e. g > 3.2, is an approximately low-spin state (S = 1/2), showing a smaller splitting of orbitals than the typical low-spin species as the Type II hemichrome [111,112]. Thus, in bis-coordinated systems with two strong-field ligands, such as imidazoles and most pyridines, the spin state is S = 1/2, showing very different g-values when compared to complexes formed with weak axial ligands, which tends to form an intermediate spin state of S = 3/2 [113,114]. In agreement with many works [100,101,115], the $g \sim 3.5$ corresponds to the g_z -value and the other g-values for the same configuration were not observed.

As for the Type II hemichrome (*g*-values at 2.87 and 2.27), its electronic configuration is $(d_{xy})^2(d_{xz},d_{yz})^3$, and the third *g*-value obey the Griffith condition ($\sum g^2 \sim 16$) [116], yielding a *g*-value of 1.61. The higher proximity of these *g*-values indicates that this EPR spectrum has large rhombicity with axial ligand planes aligned almost parallel to the N_p-Fe-N_p direction, where N_p are the diametrically opposed pyrrole-

nitrogens in the porphyrin ring [96,111]. Unfortunately, it is not possible to perform a similar analysis for the Type I hemichrome (HALS) because only one component of *g*-tensor was experimentally determined.

In complexes with a parallel orientation of two planar axial ligands, such as Type II hemichromes, the porphyrin remains planar [97,117], whereas the porphyrin ring is almost invariably distorted from planarity in porphyrin complexes with two planar axial ligands in a perpendicular orientation [96,104,105]. The *d*-orbital splittings are very important to the reactivity of the ferric center of the hemichromes. The perpendicular alignment of planar axial ligands could lead to a positive shift in reduction potential of up to 50 mV over that observed for parallel alignment, all other structural and environmental factors being equal [96,111,118].

Besides the hemichromes discussed above, pentacoordinated species are also identified by the EPR spectra at pH lower than 3.5 (Fig. 4). The hemichrome-pentacoordinated transition is characterized by an increase of the rhombic features observed by the high-spin signal (g-value around 6.0). In fact, the broadening of the spectral line in $g \approx 6.4$ indicates a second spectral component (Fig. 4). This new spectral component suggests the presence of a pentacoordinate species, corroborating the UV-Vis measurements. Usually, a pentacoordinate system has lower symmetry than a hexacoordinate system. Furthermore, this asymmetry is increased by the *domed* conformation, i.e., the out-of-plane arrangement of the metallic center [91].

A correlation between histidine orientation and degree of the ferric ion displacement from the heme plane with the length of the histidine–iron bond has been reported [31]. The occurrence of an intense resonance at g = 3.5 and an increase of asymmetry in the axial line g = 6.4 at pH 3.5 (Fig. 4) allowed the inference that the formation of the strong g_{max} (Type I) hemichrome is somehow related to the appearance of the pentacoordinate species (Fig. 4A). Hence, the strong g_{max} hemichrome formation could be a prerequisite to formation of the pentacoordinated species. The strong g_{max} (Type I) hemichrome represents a weakening of the axial histidines coordination as compared to the low-spin Type II hemichrome. Nistor et al. [95] have found similar transition in acidic medium to a neuroglobin ferric, suggesting that histidine disruption from hemichrome is a mechanism to generate pentacoordinate species.

Blumberg and Peisach associate the spin transition to yield hemichrome as a consequence of disruption of hydrophobic interactions [47]. In agreement with Tsuka [119], the cooperative breaking of van der Waals contacts between the porphyrin and the globin would be associated with the spin state change. It is possible that the aquomet-hemichrome-pentacoordinate transition could indicate a gradual increase of solvent accessibility allowed by an initial unfolding, corroborating previous studies on whole hemoglobin of G. paulistus [39]. Indeed, Imasato et al. [11] characterized the acid dissociation of the whole hemoglobin of G. paulistus by resonant light scattering (RLS) in pH values below 5.0. This dissociation of oligomeric fractions must enhance the contact of the water solvent molecules with the ferric center, becoming more pronounced the mechanism of transition between species, specially the pentacoordinate species formation, which occurs in pH<5.0.

The EPR results suggest that there is a sequence of species formation as function of pH, which is probably associated to activity–structure relationship of this hemoglobin (Scheme 1).

The EPR spectra indicate that aquometHb remains even in the highly acid pH. The coexistence of various species is a characteristic property of most hemoglobins, which makes the complex equilibrium of this supramolecular system.

3.3. CD measurements

Far-UV circular dichroism spectra for the HbGp as a function of pH are shown in Fig. 5. At pH 7.0, the CD spectrum is characterized by positive bands at 195, and two negative bands at 208 and 222 nm assigned to a considerable α -helix content as expected in hemoglobins. Upon acidification at pH<5, a decrease of both 195 nm and 222 nm bands is observed and it is associated with a decrease of helical content. The stability of the aquomet species at pH 7.0 (observed as a single-component in the EPR spectrum of Fig. 2) is due to the hydrophobicity of the heme pocket. Decreases in pH lead to an increase of the solvent accessibility, destabilizing the aquomet species. This process is probably related mainly to loss of helical content.

The absolute values of residual ellipticity at 195 and 222 nm increase gradually from pH 2.0 up to pH 4.0. This behavior is somehow disturbed at pH greater than 5.0, and the bands get to a

Scheme 1. The broad lines represent the side view of heme surfaces and the fourth model indicates the change of orientation of the distal histidine in relation to the heme, i.e., the side view of the distal histidine coordinated to the ferric center constituting the hemichrome HALS.

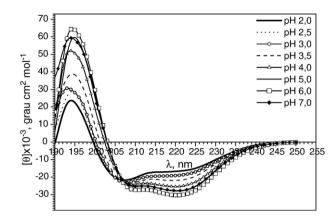


Fig. 5. CD spectra of *Glossoscolex paulistus* hemoglobin as function of pH. Conditions: protein concentration was 0.22 mg ml⁻¹ (ϵ_{415} =3.13 ml mg⁻¹ cm⁻¹).

maximum around pH 6.0. Such an effect can be due to the proximity between pH 6.0 and the protein isoelectric point (pI). The pI for this class of hemoglobins is acid [5] like, for example, the pI for the *L. terrestris* hemoglobin that is 5.5 [120]. This spectral behavior could be associated with the "aggregation" state of the hemoglobin that is maximum at the pI because of minimum electrostatic repulsion [121]. In pH lower than pI the loss of α -helix is drastic, indicating significant influence of the acid medium on the globin.

The relationship between the increase of solvent accessibility in the heme pocket and the loss of secondary structure would indicate that the formation of the pentacoordinate species could be an initial step of the denaturation process. Madura et al. [21] have reported that, in most vertebrate hemoglobins, changes in protein tertiary structure are induced by either ligand binding or changes in oligomeric assembly at the heme neighborhood. Nevertheless no such changes are observed for L. terrestris hemoglobin. Considering the similarity between L. terrestris and G. paulistus hemoglobins, it is possible to infer that the subunit assembly of a giant extracellular hemoglobin is less susceptible to the solvent permeability than vertebrate hemoglobins. Hundahl et al. [122] assert that the small effects of water in the extracellular invertebrate hemoglobins may be correlated with small surfaceto-volume ratios in these high-molecular-weight proteins that pose a limit to water-accessible sites, and suggest smaller quaternary structural changes when compared to dimeric and tetrameric hemoglobins. Other studies are in progress in order to analyze the implication of the pentacoordinate formation in the acid denaturation.

4. Conclusions

The giant extracellular hemoglobin of G. paulistus was studied in the acidic pH range. The coexistence of five species was observed: aquomethemoglobin, strong g_{max} or HALS hemichrome (Type I), low-spin hemichrome (Type II), hemichrome with an unusual spin state $(d_{xz}, d_{yz})^4 (d_{xy})^1$ (Type III) and a high-spin pentacoordinate species with a spin state $(d_{xz}, d_{yz})^2 (d_{xz})^1 (d_{x-y}^2)^1 (d_z^2)^1$.

The hemichrome and pentacoordinate formation are explained by protonation of the histidines associated with the increase of the water accessibility to the heme pocket and the smaller possibility for hydrogen bond formation. The influence of the hydrogen bond in the stability of the sixth ligand is decisive to the consequent heme properties, such as electronic structure, ligand affinities, coordination number, redox potential and macrocycle conformations [73,121,123–127]. Furthermore, this greater accessibility could be related to the subunit dissociation, in a similar way to the alkaline process, where *d* monomer dissociates from the trimer, at pH 9.0. Finally, the presence of the pentacoordinate could be associated to an initial acid denaturation mechanism.

It is likely that the transition from the pure low-spin hemichrome to the *strong* g_{max} hemichrome would be a prerequisite to the formation of the pentacoordinate species. This transition is due to an orientational change of the axial ligands, which represent a weakening of the histidine–metal ligation.

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