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Linking Conformation Change to Hemoglobin Activation Via Chain-Selective Time-resolved Resonance Raman Spectroscopy on Protoheme/Mesoheme Hybrids

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Abstract:

Time-resolved Resonance Raman spectra are reported for Hb tetramers, in which the aand β chains are selectively substituted with mesoheme. The Soret absorbtion band shift in meso- relative to proto-heme permits chainselective excitation of heme RR spectra. The evolution of these spectra following HbCO photolysis show that geminate recombination rates and yields are the same for the two chains, consistent with recent results on ¹⁵N-heme isotopomer hybrids. The spectra also reveal systematic shifts in the deoxyheme v_4 and $v_{\text{Fe-His}}$) RR bands, which are anti-correlated. These shifts are resolved for the successive intermediates in the protein structure, which have previously been determined from time-resolved UVRR spectra. Both chains show Fe-His bond compression in the immediate photoproduct, which relaxes during the formation of the first intermediate, R_{deoxy} (0.07 µs), in which the proximal F-helix is proposed to move away from the heme. Subsequently, the Fe-His bond weakens, more so for the a than the β chains. The weakening is gradual for the β chains, but abrupt for the α chains, coinciding with completion of the R-T quaternary transition, at 20µs. Since the transition from fast- to slow-rebinding Hb also occurs at 20 μ s, the drop in the a chain v_{Fe-His} supports the localization of ligation restraint to tension in the Fe-His bond, at least in the a-chains. The mechanism is more complex in the β chains.

Keywords: Protoheme/mesoheme, hybrid hemoglobin, resonance Raman, geminate recombination, allostery.

Introduction

Understanding the connection between heme ligation and protein conformation change has long been a major aim of research on hemoglobin (Hb), the paradigmatic allosteric protein. In Perutz' original stereochemical mechanism (1), this connection is provided by

constraint on the heme Fe, which lies below the heme plane in fivecoordinate high-spin deoxy heme, but must approach the plane in order to bind an exogenous ligand. Constraint on this motion by the protein lowers the ligand affinity, while overcoming the constraint induces changes in the protein structure and alters contacts at the subunit interfaces, thereby destabilizing the T quaternary structure, relative to the R structure that is characteristic of ligated Hb (2–4).

This constraint is transduced by the bond connecting the Fe to the proximal histidine ligand. The stretching vibration of this bond produces a detectable resonance Raman signal in the deoxy form of Hb, whose frequency is distinctively lower than in deoxy-myoglobin or in relaxed forms of Hb, ~216 vs ~222 cm⁻¹ (5–10). This evidence for Fe-His bond weakening is the most direct sign of molecular tension in the T state. The v_{Fe-His} RR band therefore provides a useful monitor for the heme-protein dynamics (11–15).

However, this form of tension is more pronounced in the a than the β chains. A variety of evidence points to selective weakening of the Fe-His bond in the a-chains of dexyHb (6, 9, 16). RR studies on hybrid Hb's have indicated that the band is at ~218 cm⁻¹ in the β chains but at $\sim 207 \text{ cm}^{-1}$ in the a chains. For native Hb, the presence of overlapping contributions from the two chains has complicated the interpretation of time-resolved RR studies, in which thev_{Fe-His} RR band is monitored during evolution of the protein from the R to the T state, following HbCO photolysis (11-15). In the present work, we remove this interference by employing chain-selective substitution of mesoheme, in which the vinyl substituents of protoheme are replaced with ethyl groups. This replacement shifts the Soret absorption band of the heme, thereby allowing RR selectivity via appropriate choice of laser wavelength (16, 17). We show that the a chains are indeed the locus of most of the v_{Fe-His} perturbation, and we link the successive steps in its evolution to tertiary and quaternary changes in the protein that have previously been established by UVRR spectroscopy (13, 18, 19).

Materials and Methods

Protein Preparation

The human hemoglobin (HbA) was isolated from red blood cells according to the previously published procedures(20, 21) and stored as the CO adduct. The apohemoglobin (apoHb) was obtained using acid-acetone method(22, 23). The meso-heme substituted hemoglobin (mesoHb) was obtained by reconstitution of apoHb with Fe(III)

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mesoporphyrin IX chloride (meso-heme, Frontier Scientific, Logan, Utah)(16, 22, 23). The HbA and reconstituted mesoHb were separated into subunits using p-(chloromercuri)benzoate (PMB, Sigma-Aldrich, Milwaukee, WI)(16, 24–26). The isolated subunits were regenerated by 2-mercaptoethanol (BME, Sigma-Aldrich, Milwaukee, WI) and purified according to well established procedures (16, 24–26). The purified ferrous CO subunits were equilibrated with 100 mM phosphate buffer 7.4 before hybrid formation. The $(a_{proto}\beta_{meso})_2$ hybrids were prepared by combining 1.2 molar excess of β_{meso} subunits with a_{proto} subunits, followed by overnight incubation at 4°C to ensure tetramer formation(16, 27, 28). The $(a_{meso}\beta_{proto})_2$ hybrids were prepared is similar way using corresponding appropriate subunits (16, 27, 28). The excess of β subunits was removed using column chromatography(CM52) as described previously (16, 17, 27, 28). In order to further ensure that the samples are in the pure ferrous CO state, sodium dithionite was added under CO atmosphere. Excess dithionite was removed by passing the fully reduced CO samples through Bio-Gel P-6 column equilibrated with a CO-saturated 50 mM phosphate buffer pH 7.4. Purity of the hybrids was confirmed by UVvis spectroscopy (29) (Hewlett-Packard Model 8452 Diode Array Spectrophotometer). The concentrations of all samples used for RR measurements were ~ 0.4 mM in heme in CO-saturated 50 mM phosphate buffer, pH 7.4.

About 350 μ I of sample solution was contained in a CO saturated septum sealed 5 mm glass NMR tube. The solution was stirred via a stationary stirrer bar/magnet combination through spinning of the sample tube. The diameters of the pump and probe laser spots at the samples were estimated to be ~ 80 and 60 μ m respectively and the diameter of the NMR tube was 5 mm. To avoid spatial overlap mismatch between pump and probe laser spots, and to ensure that successive pulses excited a new sample volume, the spinning speed of the NMR tube (monitored on-line by a tachometer (08210, Cole Parmer)) was optimized to 800 rpm for 200 ns to 3 us time delays, 500 rpm for 6 to 50 us delays, and 300 rpm for longer delay times. The shape of the CO recombination profiles were compared with previous studies to verify the reliability of the spinning speed used in the present experiment. Sample integrity was checked by the reproducibility of Raman and absorption spectra.

Solutions of deoxy-HbA (and deoxy-mesoHb, deoxyproto/meso-hybrids) (0.4 mM in heme) were generated in 50 mM phosphate buffer at pH 7.4 by irradiating the deoxygenated protein solutions with visible light at 4 °C under flowing ultra-pure N₂. About

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0.4 ml of solution was transferred to an septum sealed NMR tube under excess N_2 pressure.

Resonance Raman Spectroscopy

The time-resolved resonance Raman (TR³) spectroscopic setup has been described elsewhere(30, 31). Briefly, photolysis pulses at 419 nm (20 ns, 60µJ/pulse, 1 kHz) were generated by frequency doubling the output of a Ti-Sapphire laser (Photonics International Industries), which was pumped (527 nm, \sim 10 mJ/pulse, 70 ns, 1 kHz) by an intracavity frequency-doubled Nd-YLF laser (GM30, Photonics International Inc). Probe pulses (20 ns, 1 kHz) at 440 nm (0.5 µJ/pulse) were obtained by frequency doubling a second Ti-Sapphire laser(Photonics International Industries). Scattered light was collected at 135° with a camera lens (f # 1.2) and focused using a f-matched singlet lens at the entrance slit of the 0.8 m spectrograph (Spex 1401) equipped with a holographic grating (1800 groove mm⁻¹) and intensified CCD (PI/MAX, Princeton Instruments). A time-delay signal generator (DG535, Stanford Research) was used to control the delay time between pump and probe pulses and synchronize the Raman scattered signals from the probe pulse with the gate of the detector. Interfering signals from the pump laser were eliminated by electronically gating the detector. The time-resolution of the current set up is 40 ns. The probe pulse wavelength, 440 nm, was selected to provide selective resonance enhancement of protoheme in deoxy-HbA and deoxy-proto/meso-hybrids. Static RR spectra of deoxy Hb samples were obtained using the same experimental set-up as above.

The RR spectral data were processed with *Grams/AI* (7.0) software (Thermo Galactic). The bound CO fraction was determined at each delay, by deconvoluting the peaks and dividing their heights by those obtained from a probe-only spectrum (in order to compensate for the measurable photolysis induced by the probe pulse). The component spectra of photoproducts were obtained using UVRR derived time constants by means of linear least squares in Matlab(18).

Results

Substitution of the *proto*-heme vinyl substituents by ethyl groups produces a ~10 nm blueshift of the heme Soret absorption band (32, 33), as illustrated in Fig. 1. The broad deoxyHb Soret bands are readily deconvoluted into separate contributions from the *proto*-and *meso*-heme chromophores, and an additional short-wavelength contribution from vibronic side-bands. Interestingly, the heme

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absorptivity in the a chains is higher than that of the β chains as previously noted (32, 33). This difference accounts for the greater RR enhancement seen for a than for β chains in our recent study of ¹⁵N-heme isotopic hybrids of Hb (34).



Figure 1 Soret absorption envelops for the meso-proto hybrid Hb's, showing the

proximity of the 440 nm Raman laser wavelength to the deconvoluted proto-heme band.

Because of the 10 nm separation between the proto- and mesoheme contributions, excitation on the long wavelength side of the absorption band (440 nm) provides selective enhancement of the proto-heme RR spectra (16). Figure 2 shows that the RR spectrum of Hb with meso-heme in both chains can still be recorded at this wavelength (bottom spectrum), but the enhancement is ~ 3 times lower than for native Hb (top spectrum). Saturation of the vinyl double bonds also eliminates the characteristic vinyl C=C stretching mode, at 1622 cm⁻¹ and shifts the porphyrin skeletal modes to varying extents (35). Particularly noticeable is the 17 cm⁻¹ upshift of the v₂ band, to 1585 cm⁻¹. *Proto*-heme has only a weak v_4 band at 1528 cm⁻¹ with equal intensity in the RR spectrum of native Hb and the two hybrids, confirming that contribution of *meso*-heme to the *proto*-hemedominated 440 nm RR spectra is small. Other differences are seen in the low-frequency region, where the 300 cm^{-1} band appears to shift up 27 cm⁻¹. This band has been identified(16, 17, 35) as a porphyrin out-of-plane mode, γ_7 , but it has a substantial contribution from the vinyl substituents (36-38), likely accounting for the up-shift upon the removal of vinyl conjugation. It is noted that the relatively strong y_7 mode of the mesoheme is not seen in the spectra of the hybrid acquired with 440 nm excitation, ensuring that the bands seen in the hybrids are exclusively from the *proto*-heme-bearing chains. Consistent with previous hybrid data (9, 16, 39), the v_{Fe-His} band is found at 218 cm⁻¹ for the β chains, while the a chain envelope has 203 and 214 cm⁻¹ contributions.



Figure 2 The 440 nm - excited RR spectra of native (top) and meso-Hb (bottom), and of the two meso-proto hybrids. Band assignments and frequency shifts are indicated.

Figure 2 also reveals RR differences in the high-frequency region between the a and β chains in deoxyHb. The a chain frequencies are 2.5 and 1.8 cm⁻¹ higher than the β chain frequencies for v₄ and v₃, but 1.2 and 1.5 cm⁻¹ lower for v₂ and v_{C=C}. The v₄ elevation for the a chains was previously noted in ¹⁵N-heme isotopic hybrids (34), and is attributed to the electronic effects associated with strain in the Fe-His bond (see below) since a negative correlation has been established between the deoxy-heme v₄ and v_{Fe-His} frequencies (10). The other heme mode shifts are likely due to the same effects.

Fig. 3 shows the time evolution of v_4 , following HbCO photolysis with a saturating pump pulse. The deoxy-heme v_4 frequencies are the same for the two chains at the earliest time, and evolve to higher values for the a, but not for β chains. The 1376 cm⁻¹ band from COheme reflects geminate recombination at early times, and secondorder recombination of CO from solution at later times. The deoxyand CO-heme contributions are readily deconvoluted, and the peak heights, corrected for the small extent of re-photolysis generated by

the probe pulse (bottom spectra), can be used to construct rebinding yield curves (Fig. 4), as described previously (18). The geminate and second-order rebinding phases are well separated in time, and the plateau between them gives an accurate measure of the geminate yield, which is $42 \pm 2\%$ for native Hb and for both hybrids. Fully *meso*-heme-substituted Hb was included as a control, and gave the same geminate yield, confirming that the substitution has no effect on rebinding. The absence of any detectable difference between the a and β chains confirms our recent result using ¹⁵N heme substitution (34).







Figure 4 The CO recombination progress curves for the indicated Hb forms, calculated via the v_4 intensities.

Fig. 5 shows the time evolution of low-frequency RR difference spectra [(pump+probe) minus (probe-only)] for the two hybrid Hb's. The v_{Fe-His} band is seen only for deoxy-heme. Its observation in the probe-only spectra is due to partial photolysis from the probe pulse. As previously noted (11–15), this vibration has an elevated frequency, ~228 cm⁻¹, *immediately after photolysis*, which then decreases with time. The magnitude of this evolving frequency shift is more pronounced for the a than the β chain (*vide infra*).



Figure 5 Pump/probe minus probe only RR spectra at the indicated time delays. The ~228 cm⁻¹ v_{Fe-His} bands in the probe only spectra (top) are due to partial photolysis. Static deoxy Hb spectra are compared (bottom).

The spectra were analyzed with the aid of a kinetic model derived from time-resolved UVRR studies (18); the exact procedure used to extract the spectra of the various kinetic intermediates from the set of raw spectra obtained here at different times is described in detail in the cited reference. These spectra are displayed in Fig. 6 and and7,7, and are discussed below.



Figure 6 Computed RR spectra in the low frequency region after HbCO photolysis, as determined from time-resolved UVRR analysis(18).



Figure 7 Computed RR spectra in the v_4 region after HbCO photolysis, as determined from time-resolved UVRR analysis(18).

Discussion

As demonstrated previously, the substitution of *meso*-heme for the native *proto*-heme in Hb allows for chain-selective recording of heme RR spectra. In the present study, we used excitation at 440 nm,

where RR scattering is strong from proto-heme, but weak from mesoheme (Figure 1), in order to selectively monitor the protoheme vibrations in the proto/meso hybrid Hb's.

Although the *meso/proto*-heme substitution in Hb does have modest functional effects (40–43) (higher affinity, lower cooperativity) the present results show that geminate recombination is unaffected, and is equally probable in the a and β chains, confirming the result of our recent study using ¹⁵N-heme isotopic hybrids (34). We discussed there the possibility that this equivalence might reflect mechanical coupling within an $\alpha\beta$ dimer, or might be coincidental, the result of convergent evolution toward a cooperative tetramer.

An important additional opportunity is presented by these proto/meso hybrids in that they also allow us to monitor the evolution of the Fe-His stretching vibration. It has long been known that its frequency is depressed in deoxyHb (5-7, 9-14, 39), reflecting the molecular tension associated with the T state. Moreover the frequency lowering is greater for the a than the β chains. This can be seen in the deoxyHb spectra in Fig. 2, showing the $a_{\rm D}$ RR Fe-His band to consist of two components, 203 and 214 cm⁻¹, confirming earlier observations with hybrids Hb's (6, 9, 10, 16). The β_{p} heme has a single band at 218 cm⁻¹, while native Hb has an asymmetric band, reflecting the sum of a and β contributions. The source of the heterogeneity reflected in the two-component a_p RR band has yet to be determined (6, 9, 10, 16), but the greater weakening of the g chain Fe-His bond is clear. Other manifestations of weaker Fe-His bonding in the deoxyHb a chains include a-selective severing of this bond at low pH or in the presence of allosteric effectors when NO is bound (due to the NO trans effect)(44, 45), or when Ni(II) is substituted for Fe(II) (the Ni(II) axial bonding being weak)(46). Also, greater splitting of the Q absorption band is seen in high-resolution emission spectra when Zn(II) is substituted in the a than in the β chains in Hb(47). The time-resolved RR spectra in Fig. 5 show that the Fe-His perturbation is also more extensive in the α chains than in the β chains.

UVRR spectroscopy has previously been used to characterize the protein structure changes following HbCO photolysis (13, 18, 19). The time-course of tryptophan and tyrosine mode intensities defined a set of transitions, associated with well-separated time constants of 0.07, 0.75, 2.9 and 20 μ s, prior to recombination with solution CO to reform HbCO (18). The kinetic model based on these transitions afforded a set of UVRR spectra for the kinetic intermediates, which permitted interpretation of the structural changes. The resulting model of the allosteric reaction coordinate is shown in Figure 8. Photolysis of ligated

heme (A) leaves CO in the distal pocket (B) while the Fe-His bond is compressed, as the Fe atom is driven out of the heme plane by its conversion to a high-spin state. It has long been known that the Fe-His frequency is elevated in the immediate photoproduct of HbCO (11–15). This can be seen in the probe-only RR spectra, shown in Fig. 6. The photoproducts generated by the probe pulse display Fe-His bands at 229 (a_p) and 227 (β_p) cm⁻¹.



Figure 8 Allosteric model for the reaction path following HbCO photolysis showing the v_{Fe-His} values for the intermediates (upper number for a subunit, lower number for β subunit)[see text for descriptions of the intermediates].

The first protein motion, at 0.07 μ s, produces an intermediate called R_{deoxy}, because its difference spectrum can be simulated by Hb constructs in which ligands are absent in one or more subunits but the quaternary structure remains R (48). This spectrum is characterized by large Trp and Tyr intensity losses, which have been shown to result from breaking of tertiary H-bonds between the E and A helices (49, 50), and between the F and H helices (51) (R_{deoxy} in Fig. 8). The E and F helices hold the heme like a clamshell, and the interhelical H-bonds are proposed to break as a result of rotation of the clamshell. This rotation results from the proximal F helix moving away from the heme, impelled by the Fe-His bond compression. The resulting relaxation of the Fe-His bond can be seen in the visible RR difference spectra

extracted for the R_{deoxy} intermediate, showing v_{Fe-His} at 224 and 223 cm⁻¹ (Fig. 6), values similar to those seen in deoxy-myoglobin (Mb), which has no quaternary constraints.

The postulated clamshell rotation can be seen by comparing high-resolution crystal structures of MbCO and deoxyMb, and has been successfully modeled in Mb with DFT/MM calculations(52). Importantly, it has also been observed in crystal structures of Hb from Antarctic fish which have been trapped in an intermediate conformation along the allosteric coordinate because of selective oxidation of β subunits(53, 54). The clamshell rotation brings the distal E helix toward the heme, impelling the photodissociated CO to recombine geminately (40 %) or else to exit the distal pocket. The rate of geminate recombination is essentially the same as that of R_{deoxy} formation. The R_{deoxy} spectrum disappears in 0.75 µs signaling the reformation of the interhelical H-bonds (intermediate R_s)(18). This is suggested to result from subsequent motions of the H and A helices whose C and N termini form salt-bridges in the T state. These motions are viewed as a conformational search phase, resulting in the beginning of inter-dimer rotation and formation of the 'hinge' contact between the FG-corner of the a_P chain and the C-helix of the β_2 chain, at 2.9 μ s (intermediate R_T)(18). The R_T UVRR difference spectra reveal the resulting H-bond between Trp β 35 and Aspa94 at the rotation hinge. Finally, the 'switch' is thrown at 20 μ s (intermediate T'), as shown in the UVRR spectrum by a signal characteristic of the resulting H-bond between Tyra42 on the C-helix of the a_P chain and AspB99 on the FG-corner of the β_2 chain. This completes the transition to the T quaternary structure. However, because of the geminate recombination, the Hb tetramer retains 1–2 CO molecules on average, and the intermediate is labeled T'.

The heme RR spectra (Fig. 6 and and7)7) show divergent behavior for a and β chains following R_{deoxy} v_{Fe-His} between R_{deoxy}, R_S and R_T, but then a sharp drop in T'. In contrast, the β chain v_{Fe-His} decreases gradually at each step, and is higher in T' than is the a chain v_{Fe-His}. These trends in the behavior of the v_{Fe-His} modes are supported by the inverse behavior of the v₄ modes seen in Figure 7(discussed further below). For both chains the T' frequencies of the v_{Fe-His} band remain higher than those seen in deoxyHb (see Fig. 2 and and6).6). We conclude that the presence of the CO ligands in the T' intermediate perturb the Hb structure, even though the UVRR signals indicate the formation of T quaternary contacts. Quantification of the UVRR signals is problematic because of the issue of ligand distribution among the Hb molecules in the sample, and the uncertain relationship of ligand occupancy to quaternary state(55, 56). It is known, however,

that structural perturbations in deoxyHb (induced by site-mutation or chemical modification) which weaken the UVRR difference signals, also shift the v_{Fe-His} band to higher frequencies (13, 51, 57, 58).

Nevertheless, the sharp drop in the a chain v_{Fe-His} when the quaternary 'switch' is thrown at 20 µs supports the idea that the Fe-His perturbation reflects molecular tension. Gibson showed long ago that 20 µs characterizes the transition between fast- and slow-recombining forms of hemoglobin (59). These forms were associated with the R and T states. This was why the penultimate 2.9 µs intermediate was labeled R_T (18), even though the formation of the hinge contact signals a quaternary motion; a recent time-resolved wide angle x-ray scattering study confirms a 3 µs time constant for quaternary motion (60). Perutz initially advanced the notion that protein-induced strain on the Fe-His bond could hinder the approach of the Fe to the heme plane, thereby inhibiting ligation(4). The temporal association of the v_{Fe-His} lowering with the lowering of ligation rate provides direct support for this mechanism.

Perutz also recognized that a different mechanism was required for the β chains, and proposed steric hindrance to ligand entry, since deoxyHb crystal structures showed the distal pocket to be more crowded for the β than the a chains(4). Subsequent measurements, especially those of Unzai et al (61, 62) with Cr/Fe and Zn/Fe hybrid Hb's, in which the R and T states are stabilized by the Cr(III) and Zn(II) ions, have shown little difference between a and β chain binding rates and affinities. Consistent with Perutz' proposal, DFT/MM calculations indicate similar R-T energy differences between the two chains, but with quite different apportioning of the energy to different structural elements in the heme pocket (52). Thus nature appears to have evolved similar energetic responses to allostery in the two chains, a requirement for cooperative binding, by distinctly different mechanisms.

We note that the v₄ frequency shows a slight but distinctive evolution (Figure 7), as was also seen in the ¹⁵N-heme isotopomer hybrid study [24]. Rousseau et al. (10) have noted a negative correlation between v_{Fe-His} and v₄ for a variety of Hb constructs. The present results allow us to test this correlation in the context of native Hb. Figure 9 confirms the correlation for the Hb kinetic intermediates. It also reveals a distinctively greater slope for the a than the β chains. The v₄ mode is responsive to electron density in the porphyrin ring (the upshift seen for HbCO reflects electron withdrawal by the CO ligand), which is modulated by alterations in the donor strength of the

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Fe-His bond. The higher slope is another manifestation of the greater constraint on this bond in the α than the β chains.



Figure 9 The $v_{\text{Fe-His}}/v_4$ correlation for the kinetic intermediates and for the deoxy Hb (T)

Conclusions

Chain-selective substitution of *meso*-heme into Hb coupled with time-resolved pump-probe RR spectroscopy has revealed identical geminate CO recombination yields for a and β chains, confirming the results obtained by selective 15-N *proto*-heme substitution. In addition, the *meso/proto* hybrids have permitted chain selective monitoring of the Fe-His stretching vibration as it evolves through successive intermediates along the R-T allosteric coordinate. Immediately after CO photolysis, the Fe-His frequency is elevated in both chains, reflecting compression of the Fe-His bond as the Fe is driven out of the heme plane. It relaxes with a time constant of 0.07 µs, reflecting relief of the bond compression as the His residue moves away from the heme during the clamshell rotation of the E and F helices. For the a chains, the Fe-His frequency then remains

unperturbed during subsequent tertiary and quaternary intermediates, until the quaternary 'switch' is thrown, at 20 μ s. The drop in a chain Fe-His frequency at this stage, coincident with a diminished CO rebinding rate, demonstrates that the 'switch' inhibits ligand binding by constraining the His ligand in the a chains. For the β chains, the Fe-His frequency is lowered slightly and gradually throughout the allosteric coordinate, ending at a higher value than in the a chains. This behavior indicates a different mechanism of ligand binding inhibition in the β chains, consistent with computational results(52), and with the observation of distal crowding in the β chain binding pocket.

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References

- 1. Perutz MF. Nature. 1970;228:726-734.
- 2. Baldwin J, Chothia C. J Mol Biol. 1979;129:175–220.
- 3. Monod J, Wyman J, Changeux JP. J Mol Biol. 1965;12:88–118.
- 4. Perutz MF. Mechanisms of Cooperativity and Allosteric Regulation in Proteins. Cambridge University Press; Cambridge: 1990.
- 5. Friedman JM, Scott TW, Stepnoski RA, Ikedasaito M, Yonetani T. J Biol Chem. 1983;258:564–572.
- 6. Kaminaka S, Ogura T, Kitagishi K, Yonetani T, Kitagawa T. J Am Chem Soc. 1989;111:3787–3794.
- 7. Kitagawa T, Nagai K, Tsubaki M. FEBS Lett. 1979;104:376–378.
- 8. Nagai K, Kitagawa T. Proc Nat Acad Sci USA Biol Sci. 1980;77:2033– 2037.
- 9. Ondrias MR, Rousseau DL, Kitagawa T, Ikedasaito M, Inubushi T, Yonetani T. J Biol Chem. 1982;257:8766–8770.
- 10. Ondrias MR, Rousseau DL, Shelnutt JA, Simon SR. Biochemistry. 1982;21:3428–3437.
- 11. Friedman JM, Rousseau DL, Ondrias MR. Annu Rev Phys Chem. 1982;33:471–491.

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- 12. Friedman JM, Rousseau DL, Ondrias MR, Stepnoski RA. Science. 1982;218:1244–1246.
- 13. Jayaraman V, Rodgers KR, Mukerji I, Spiro TG. Science. 1995;269:1843– 1848.
- 14. Jayaraman V, Spiro TG. Biospectrosc. 1996;2:311–316.
- 15. Scott TW, Friedman JM. J Am Chem Soc. 1984;106:5677–5687.
- 16. Jeyarajah S, Kincaid JR. Biochemistry. 1990;29:5087–5094.
- 17. Podstawka E, Proniewicz LM. J Inorg Biochem. 2004;98:1502–1512.
- 18. Balakrishnan G, Case MA, Pevsner A, Zhao XJ, Tengroth C, McLendon GL, Spiro TG. J Mol Biol. 2004;340:843–856.
- 19. Rodgers KR, Su C, Subramaniam S, Spiro TG. J Am Chem Soc. 1992;114:3697–3709.
- 20. Antonini E, Brunori M. Hemoglobin and Myoglobin and their Reactions with Ligands. Elsevier; New York: 1971.
- Riggs A, Eraldo Antonini LR-BaEC. Methods Enzymol. Academic Press; 1981. pp. 5–29.
- 22. Ascoli F, Rossi Fanelli MR, Antonini E, Eraldo Antonini LR-BaEC. Methods Enzymol. Academic Press; 1981. pp. 72–87.
- 23. Yip YK, Waks M, Beychok S. J Biol Chem. 1972;247:7237-7244.
- Bucci E, Eraldo Antonini LR-BaEC. Methods Enzymol. Academic Press; 1981. pp. 97–106.
- 25. Geraci G, Parkhurs Lj, Gibson QH. J Biol Chem. 1969;244:4664-4667.
- 26. Yip YK, Waks M, Beychok S. Proc Nat Acad Sci USA. 1977;74:64–68.
- 27. Ikeda-Saito M, Inubushi T, Yonetani T, Eraldo Antonini LR-BaEC. Methods Enzymol. Academic Press; 1981. pp. 113–121.
- 28. Makino N, Sugita Y. J Biol Chem. 1978;253:1174–1178.
- 29. Yamamoto H, Yonetani T. J Biol Chem. 1974;249:7964-7968.
- Zhao XJ, Chen RP, Tengroth C, Spiro TG. Appl Spectrosc. 1999;53:1200– 1205.
- 31. Zhao XJ, Tengroth C, Chen RP, Simpson WR, Spiro TG. Journal of Raman Spectroscopy. 1999;30:773–776.
- 32. Olson JS. Proc Nat Acad Sci USA. 1976;73:1140-1144.
- 33. Sugita Y. J Biol Chem. 1975;250:1251-1256.

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- 34. Balakrishnan G, Zhao XJ, Podstawska E, Proniewicz LM, Kincaid JR, Spiro TG. Biochemistry. 2008. Submitted.
- 35. Spiro TG, Burke JM. J Am Chem Soc. 1976;98:5482–5489.
- 36. Choi S, Spiro TG, Langry KC, Smith KM. J Am Chem Soc. 1982;104:4337–4344.
- Choi S, Spiro TG, Langry KC, Smith KM, Budd DL, Lamar GN. J Am Chem Soc. 1982;104:4345–4351.
- 38. Hu SZ, Smith KM, Spiro TG. J Am Chem Soc. 1996;118:12638-12646.
- 39. Kaminaka S, Zhou YX, Tsuneshige A, Yonetani T, Kitagawa T. J Am Chem Soc. 1994;116:1683–1689.
- 40. Ishimori K, Morishima I. Biochemistry. 1986;25:4892–4898.
- 41. Seybert DW, Moffat K. J Mol Biol. 1977;113:419-430.
- 42. Seybert DW, Moffat K, Gibson QH. Biochem Biophys Res Commun. 1975;63:43–49.
- 43. Seybert DW, Moffat K, Gibson QH. J Biol Chem. 1976;251:45–52.
- 44. Yonetani T, Tsuneshige A, Zhou YX, Chen XS. J Biol Chem. 1998;273:20323–20333.
- 45. Fujii M, Hori H, Miyazaki G, Morimoto H, Yonetani T. J Biol Chem. 1993;268:15386–15393.
- 46. Shelnutt JA, Alston K, Ho JY, Yu NT, Yamamoto T, Rifkind JM. Biochemistry. 1986;25:620–627.
- 47. Sudhakar K, Laberge M, Tsuneshige A, Vanderkooi JM. Biochemistry. 1998;37:7177–7184.
- 48. Mukerji I, Spiro TG. Biochemistry. 1994;33:13132–13139.
- 49. Balakrishnan G, Tsai CH, Wu Q, Case MA, Pevsner A, McLendon GL, Ho C, Spiro TG. J Mol Biol. 2004;340:857–868.
- 50. Wang DJ, Zhao XJ, Shen TJ, Ho C, Spiro TG. J Am Chem Soc. 1999;121:11197–11203.
- 51. Kneipp J, Balakrishnan G, Chen RP, Shen TJ, Sahu SC, Ho NT, Giovannelli JL, Simplaceanu V, Ho C, Spiro TG. J Mol Biol. 2006;356:335–353.
- 52. Guallar V, Jarzecki AA, Friesner RA, Spiro TG. J Am Chem Soc. 2006;128:5427–5435.
- 53. Riccio A, Vitagliano L, di Prisco G, Zagari A, Mazzarella L. Proc Nat Acad Sci USA. 2002;99:9801–9806.

- Vitagliano L, Vergara A, Bonomi G, Merlino A, Verde C, di Prisco G, Howes BD, Smulevich G, Mazzarella L. J Am Chem Soc. 2008;130:10527– 10535.
- 55. Ackers GK, Holt JM, Huang YW, Grinkova Y, Klinger AL, Denisov I. Proteins-Structure Function and Genetics. 2000. pp. 23–43.
- 56. Ciaccio C, Coletta A, De Saneti G, Marini S, Coletta M. Iubmb Life. 2008;60:112–123.
- 57. Ishimori K, Hashimoto M, Imai K, Fushitani K, Miyazaki G, Morimoto H, Wada Y, Morishima I. Biochemistry. 1994;33:2546–2553.
- 58. Juszczak L, Samuni U, Friedman J. Journal of Raman Spectroscopy. 2005;36:350–358.
- 59. Sawicki CA, Gibson QH. J Biol Chem. 1976;251:1533–1542.
- 60. Cammarata M, Levantino M, Schotte F, Anfinrud PA, Ewald F, Choi J, Cupane A, Wulff M, Ihee H. Nat Methods. 2008;5:881–886.
- 61. Unzai S, Eich R, Shibayama N, Olson JS, Morimoto H. J Biol Chem. 1998;273:23150–23159.
- 62. Unzai S, Hori H, Miyazaki G, Shibayama N, Morimoto H. J Biol Chem. 1996;271:12451–12456.

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