

## X-E Biomolecular Science

Elucidation of a structure-function relationship of metalloproteins and structural chemistry of amyloid are current subjects of this group. The primary technique used for the first project is the stationary and time-resolved resonance Raman spectroscopy excited by visible and UV lasers. Various model compounds of active site of enzymes are also examined with the same technique. IR-microscope dichroism analysis and AFM are the main techniques for the second project. The practical themes that we want to explore for the first project are (1) mechanism of oxygen activation by enzymes, (2) mechanism of active proton translocation and its coupling with electron transfer, (3) structural mechanism of signal sensing and transduction by heme-based sensor proteins, (4) higher order protein structures and their dynamics, and (5) reactions of biological NO. In category (1), we have examined a variety of terminal oxidases, cytochrome P450s, and peroxidases, and also treated their reaction intermediates by using the mixed flow transient Raman apparatus and the Raman/absorption simultaneous measurement device. For (2) the third-generation UV resonance Raman (UVRR) spectrometer was constructed and we are going to apply it to a giant protein like cytochrome *c* oxidase with  $M_r = 210,000$ . Recently, we succeeded in pursuing protein folding of apomyoglobin by combining UV time-resolved Raman and rapid mixing device. We also determined the carboxylic side chains of bovine cytochrome oxidase which undergo protonation/deprotonation changes and hydrogen-bonding status changes in response with electron transfers between metal centers or ligand dissociation from heme *a*<sub>3</sub>. Currently, we focus our attention on detecting tyrosine radical for the P intermediate of terminal oxidases. In (3) we are interested in a mechanism of ligand recognition specific to CO, NO or O<sub>2</sub> and a communication pathway of the ligand binding information to the functional part of the protein. Several gas sensor heme proteins were extensively treated in this year. For (4) we developed a novel technique for UV resonance Raman measurements based on the combination of the first/second order dispersions of gratings and applied it successfully to 235-nm excited RR spectra of several proteins including mutant hemoglobins and myoglobins. Nowadays we can carry out time-resolved UVRR experiments with nanosecond resolution to discuss protein dynamics. With the system, we have succeeded in isolating the spectrum of tyrosinate in ferric Hb M Iwate, which was protonated in the ferrous state, and the deprotonated state of Tyr244 of bovine cytochrome *c* oxidase. The study is extended to a model of Tyr244, an imidazole-bound *para*-cresol coordinated to a metal ion, was synthesized and its UV resonance Raman was investigated. For (5) we purified soluble guanylate cyclase from bovine lung and observed its RR spectra in the presence of allosteric effectors. To further investigate it, we are developing an expression system of this protein. For the amyloid study, we examined FTIR spectra of  $\beta_2$ -microglobulin and its #11-21, K3, and K3-K7 peptides which form a core part of amyloid fibril.

### X-E-1 Construction of a Square-Planar Hydroperoxo-Copper(II) Complex Inducing a Higher Catalytic Reactivity

FUJII, Tatsuya<sup>1</sup>; NAITO, Asako<sup>1</sup>; YAMAGUCHI, Syuhei<sup>1</sup>; WADA, Akira<sup>1</sup>; FUNAHASHI, Yasuhiro<sup>1</sup>; JITSUKAWA, Koichiro<sup>1</sup>; NAGATOMO, Shigenori; KITAGAWA, Teizo; MASUDA, Hideki<sup>1</sup>  
(<sup>1</sup>Nagoya Inst. Tech.)

[*Chem. Commun.* 2700–2701 (2003)]

A novel hydroperoxo-copper(II) complex with a square-planar geometry has been prepared, which has exhibited a higher selectivity and catalytic reactivity for dimethyl sulfide, in contrast to that with a trigonal-bipyramidal one.

### X-E-2 Copper Hydroperoxo Species Activated by Hydrogen-Bonding Interaction with Its Distal Oxygen

YAMAGUCHI, Syuhei<sup>1</sup>; NAGATOMO, Shigenori; KITAGAWA, Teizo; FUNAHASHI, Yasuhiro<sup>1</sup>; OZAWA, Tomohiro<sup>1</sup>; JITSUKAWA, Koichiro<sup>1</sup>; MASUDA, Hideki<sup>1</sup>  
(<sup>1</sup>Nagoya Inst. Tech.)

[*Inorg. Chem.* **42**, 6968–6970 (2003)]

A novel copper(II)-OOH complex with functional ligand that can form a hydrogen bond with the distal oxygen of hydroperoxide has been designed and prepared as a structural/functional model of dopamine-hydroxylases, whose spectroscopic characterization and decomposition rates have indicated that the hydroperoxide is activated through the hydrogen-bonding interaction with the distal oxygen.

### X-E-3 Thermal Stability and Absorption Spectroscopic Behavior of ( $\mu$ -Peroxo)dicopper Complexes Regulated with Intramolecular Hydrogen Bonding Interactions

YAMAGUCHI, Syuhei<sup>1</sup>; WADA, Akira<sup>1</sup>; FUNAHASHI, Yasuhiro<sup>1</sup>; NAGATOMO, Shigenori; KITAGAWA, Teizo; JITSUKAWA, Koichiro<sup>1</sup>; MASUDA, Hideki<sup>1</sup>  
(<sup>1</sup>Nagoya Inst. Tech.)

[*Eur. J. Inorg. Chem.* **2003**, 4378–4386 (2003)]

In order to clarify the effect of hydrogen bonding on the stabilities of ( $\mu$ -peroxo)dicopper complexes with a *trans*-1,2-peroxo form, novel copper complexes with intramolecular hydrogen bonding interaction sites have been synthesized, and their spectroscopic properties and thermal stabilities studied. The selected tripodal tetradentate ligands were tris(2-pyridylmethyl)amine (TPA) derivatives bearing pivalamido and amino groups at the

6-position of the pyridine ring in TPA, {[6-(pivalamido)pyrid-2-yl]methyl}bis(pyrid-2-ylmethyl)amine (MPPA) and [(6-aminopyrid-2-yl)methyl]bis(pyrid-2-ylmethyl)amine (MAPA). The single-crystal X-ray structure of a monomeric Cu<sup>II</sup> complex with N<sub>3</sub><sup>-</sup> namely [Cu(mppa)N<sub>3</sub>]ClO<sub>4</sub> (**1a**), revealed an interligand hydrogen bonding interaction between the substituent NH group of MPPA and the azide nitrogen atom in the axial position. The Cu<sup>I</sup> complexes of MPPA and MAPA were immediately oxygenated with dioxygen in acetone solution at -78 °C to give the μ-peroxo dinuclear copper(II) complexes [{"Cu(mppa)}<sub>2</sub>(O<sub>2</sub>)]<sup>2+</sup> (**1b**) and [{"Cu(mapa)}<sub>2</sub>(O<sub>2</sub>)]<sup>2+</sup> (**2b**). These complexes exhibited two kinds of characteristic absorption bands ( $\pi^*_{\sigma} \rightarrow d_{\sigma}$ ,  $\pi^*_{\nu} \rightarrow d_{\sigma}$ ) originating from the ligand-metal charge transfer (LMCT) of O<sub>2</sub><sup>2-</sup> to Cu. Affected by the hydrogen bonding interaction, the  $\pi^*_{\delta}$  CT band shifted significantly to a higher energy region and the  $\pi^*_{\nu} \rightarrow d_{\sigma}$  CT absorbance decreased due to stabilization of the  $\pi^*$  orbital and restriction of the Cu–O bond rotation. The thermal stabilities of the (μ-peroxo)dinuclear copper(II) complexes were estimated from their decomposition rates which decreased in the order, **2b** [{"Cu(tpa)}<sub>2</sub>(O<sub>2</sub>)]<sup>2+</sup> (**3b**) > **1b** >> [{"Cu(6-metpa)}<sub>2</sub>(O<sub>2</sub>)]<sup>2+</sup> (**4b**) {6-MeTPA = [(6-methylpyrid-2-yl)methyl]bis(pyrid-2-ylmethyl)amine}. The above findings indicate that the interligand hydrogen bonding interaction, although overcome to some extent by the adverse effect of the steric bulk of the NH group, is inclined to stabilize (μ-peroxo)dinuclear copper(II) complexes.

#### X-E-4 CO Binding Study of Mouse Heme-Regulated eIF-2α Kinase: Kinetics and Resonance Raman Spectra

IGARASHI, Jotaro<sup>1</sup>; SATO, Akira<sup>2</sup>; KITAGAWA, Teizo; SAGAMI, Ikuko<sup>3</sup>; SHIMIZU, Toru<sup>1</sup>  
(<sup>1</sup>Tohoku Univ.; <sup>2</sup>GUAS; <sup>3</sup>Tohoku Univ. and Kyoto Prefectural Univ.)

[*Biochim. Biophys. Acta* **1650**, 99–104 (2003)]

Heme-regulated eukaryotic initiation factor (eIF)-2α kinase (HRI) regulates the synthesis of globin chains in reticulocytes with heme availability. In the present study, CO binding kinetics to the 6-coordinated Fe(II) heme of the amino-terminal domain of mouse HRI and resonance Raman spectra of the Fe(II)–CO complex are examined to probe the character of the heme environment. The CO association rate constant,  $k_{\text{on}}$ , and CO dissociation rate constant,  $k_{\text{off}}$ , were 0.0029 μM<sup>-1</sup>s<sup>-1</sup> and 0.003 s<sup>-1</sup>, respectively. These values are very slow compared with those of mouse neuroglobin and sperm whale myoglobin, while the  $k_{\text{off}}$  value of HRI was close to those of the 6-coordinated hemoglobins from *Chlamydomonas* and barley (0.0022 and 0.0011 s<sup>-1</sup>). The dissociation rate constant of an endogenous ligand, which occurs prior to CO association, was 18.3 s<sup>-1</sup>, which was lower than those (197 and 47 s<sup>-1</sup>) of the same 6-coordinate hemoglobins. Resonance Raman spectra suggest that the Fe–C–O adopts an almost linear and upright structure and that the bound CO interacts only weakly with nearby amino acid residues.

#### X-E-5 Resonance Raman Study on Synergistic Activation of Soluble Guanylate Cyclase by Imidazole, YC-1 and GTP

PAL, Biswajit; LI, Zhengqiang<sup>1</sup>; OHTA, Takehiro; TAKENAKA, Shigeo<sup>2</sup>; TSUYAMA, Shingo<sup>2</sup>; KITAGAWA, Teizo  
(<sup>1</sup>IMS and Jilin Univ.; <sup>2</sup>Osaka Prefecture Univ.)

[*J. Inorg. Biochem.* **98**, 824–832 (2004)]

Soluble guanylate cyclase (sGC), a physiological nitric oxide (NO) receptor, is a heme-containing protein and catalyzes the conversion of GTP to cyclic GMP. We found that 200 mM imidazole moderately activated sGC in the coexistence with 3-(5'-hydroxymethyl-2'-furyl)-1-benzylindazole (YC-1), although imidazole or YC-1 alone had little effect for activation. GTP facilitated this process. Resonance Raman spectra of imidazole complex of native sGC and CO-bound sGC (CO-sGC) have demonstrated that a simple heme adduct with imidazole at the sixth coordination position is not present for both sGC and CO-sGC below 200 mM of the imidazole concentration and that the Fe–CO stretching band ( $\nu_{\text{Fe-CO}}$ ) appears at 492 cm<sup>-1</sup> in the presence of imidazole compared with 473 cm<sup>-1</sup> in its absence. Both frequencies fall on the line of His-coordinated heme proteins in the  $\nu_{\text{Fe-CO}}$  vs  $\nu_{\text{C-O}}$  plot. However, it is stressed that the CO-heme of sGC becomes photo-inert in the presence of imidazole, suggesting the formation of five-coordinate CO-heme or of six-coordinate heme with a very weak trans ligand. These observations suggest that imidazole alters not only the polarity of heme pocket but also the coordination structure at the fifth coordination side presumably by perturbing the heme-protein interactions at propionic side chains. Despite the fact that the isolated sGC stays in the reduced state and is not oxidized by O<sub>2</sub>, sGC under the high concentration of imidazole (1.2 M) yielded  $\nu_4$  at 1373 cm<sup>-1</sup> even after its removal by gel-filtration, but addition of dithionite gave the strong  $\nu_4$  band at 1360 cm<sup>-1</sup>. This indicated that imidazole caused autoxidation of sGC.

#### X-E-6 Heme Structures of Five Variants of Hemoglobin M Probed by Resonance Raman Spectroscopy

JIN, Yayoi<sup>1</sup>; NAGAI, Masako<sup>1</sup>; NAGAI, Yukifumi<sup>1</sup>; NAGATOMO, Shigenori; KITAGAWA, Teizo  
(<sup>1</sup>Kanazawa Univ.)

[*Biochemistry* **43**, 8517–8527 (2004)]

The α-abnormal Hb Ms show physiological properties different from the β-abnormal Hb Ms, that is, extremely low oxygen affinity of the normal subunit and extraordinary resistance to both enzymatic and chemical reduction of the abnormal met-subunit. To get insight into the contribution of heme structures to these differences among Hb Ms, we examined the 406.7-nm excited resonance Raman (RR) spectra of five Hb Ms in the frequency region from 1700 to 200 cm<sup>-1</sup>. In the high-frequency region, profound differences between met-α

and met- $\beta$  abnormal subunits were observed for the in-plane skeletal modes (the  $\nu_{C=C}$ ,  $\nu_{37}$ ,  $\nu_2$ ,  $\nu_{11}$  and  $\nu_{38}$  bands), probably reflecting different distortions of heme structure caused by the out-of-plane displacement of the heme iron due to tyrosine coordination. Below 900  $\text{cm}^{-1}$ , Hb M Iwate [ $\alpha(\text{F8})\text{His} \rightarrow \text{Tyr}$ ], exhibited a distinct spectral pattern for the  $\nu_{15}$ ,  $\gamma_{11}$ ,  $\delta(\text{C}_b\text{C}_a\text{C}_b)_{2,4}$  and  $\delta(\text{C}_b\text{C}_c\text{C}_d)_{6,7}$  compared to that of Hb M Boston [ $\alpha(\text{E7})\text{His} \rightarrow \text{Tyr}$ ], although both heme irons are coordinated by Tyr. The  $\beta$ -abnormal Hb Ms, namely, Hb M Hyde Park [ $\beta(\text{F8})\text{His} \rightarrow \text{Tyr}$ ], Hb M Saskatoon [ $\beta(\text{E7})\text{His} \rightarrow \text{Tyr}$ ], and Hb M Milwaukee [ $\beta(\text{E11})\text{Val} \rightarrow \text{Glu}$ ], displayed RR band patterns similar to that of metHb A, but with some minor individual differences. The RR bands characteristic of the met-subunits of Hb Ms totally disappeared by chemical reduction and the ferrous heme of abnormal subunits was no more bonded with Tyr or Glu. They were bonded to the distal (E7) or proximal (F8) His, and this was confirmed by the presence of the  $\nu_{\text{Fe-His}}$  mode at 215  $\text{cm}^{-1}$  in the 441.6-nm excited RR spectra. A possible involvement of heme distortion in differences of reducibility of abnormal subunits and oxygen affinity of normal subunits is discussed.

#### X-E-7 Vibronic Coupling between Soret and Higher Energy Excited States in Iron(II) Porphyrins: Raman Excitation Profiles of $A_{2g}$ Modes in the Soret Region

EGAWA, Tsuyoshi<sup>1</sup>; SUZUKI, Noriyuki<sup>2</sup>; DOKOH, Takashi<sup>2</sup>; HIGUCHI, Tsunehiko<sup>2</sup>; SHIMADA, Hideo<sup>3</sup>; KITAGAWA, Teizo; ISHIMURA, Yuzuru<sup>3</sup>  
(<sup>1</sup>IMS and Keio Univ.; <sup>2</sup>Univ. Tokyo; <sup>3</sup>Keio Univ.)

[*J. Phys. Chem. A* **108**, 568–577 (2004)]

Resonance Raman spectra were observed for heme proteins and iron(II) porphyrins including ferrous-CO and ferrous-isocyanide derivatives of cytochrome P450<sub>cam</sub>, a synthetic iron(II) porphyrin complex having a thiolate axial ligand, ferrous-isocyanide derivative of myoglobin, and synthetic iron(II) porphyrin complexes having either an imidazole or a sulfide axial ligand. Among them, the former three were found to be a hyperporphyrin, giving red and blue Soret absorption bands, whereas others were normal porphyrins giving a single Soret band. When Raman scattering was excited within the Soret region, an anomalously polarized (ap) Raman line, which was assignable to the  $\nu_{19}$  mode belonging to the  $A_{2g}$  species, was observed at 1537–86  $\text{cm}^{-1}$  for all these compounds. Both the synthetic iron (II) porphyrins having the imidazole and sulfide ligands also showed another ap Raman line at 1230  $\text{cm}^{-1}$ , which was assigned to  $\nu_{26}$  of  $A_{2g}$  symmetry. Raman excitation profiles of the  $\nu_{19}$  and  $\nu_{26}$  modes showed a maximum that was displaced from the 0-0 component of the Soret or red Soret band toward higher frequencies by the frequency of the corresponding mode, indicating the 0-1 component. Although Raman lines of these modes were also observed upon excitation at the 0-0 component, they were significantly more intense at the 0-1 component. These results, together with nonadiabatic theories about vibronic contribution to Raman intensity, indicat-

ed the presence of vibronic coupling between the Soret (or red Soret) excited state and some other electronic excited state(s) located in the blue of the Soret band. The present study hence demonstrates that lower occupied orbitals other than those described in the ordinary four-orbital model and its extended form, which is applicable to the hyperporphyrins, contribute to the Soret (or red Soret) excited states.

#### X-E-8 Dioxygen Reactivity of Copper(I) Complexes with Tetradentate Tripodal Ligands Having Aliphatic Nitrogen Donors: Synthesis, Structures, and Properties of Peroxo and Superoxo Complexes

KOMIYAMA, Kazuya<sup>1</sup>; FURUTACHI, Hideki<sup>1</sup>; NAGATOMO, Shigenori; HASHIMOTO, Akifumi<sup>1</sup>; HAYASHI, Hideki<sup>1</sup>; FUJINAMI, Shuhei<sup>1</sup>; SUZUKI, Masatatsu<sup>1</sup>; KITAGAWA, Teizo  
(<sup>1</sup>Kanazawa Univ.)

[*Bull. Chem. Soc. Jpn.* **77**, 59–72 (2004)]

Oxygenation of copper(I) with tetradentate tripodal ligands (L) comprised of a tris(aminoethyl)amine (tren) skeleton having sterically bulky substituent(s) on the terminal nitrogens has been investigated, where L = tris(*N*-benzylaminoethyl)amine ( $\text{L}^{\text{H,Bn}}$ ), tris(*N*-benzyl-*N*-methylaminoethyl)amine ( $\text{L}^{\text{Me,Bn}}$ ), or tris(*N,N*-dimethylaminoethyl)amine ( $\text{L}^{\text{Me,Me}}$ ). All the copper(I) complexes reacted with dioxygen at low temperatures to produce superoxocopper(II) and/or *trans*-( $\mu$ -1,2-peroxo)-dicopper(II) complexes depending on the steric bulkiness of the terminal nitrogens and the reaction conditions. The reaction of a copper(I) complex  $[\text{Cu}(\text{L}^{\text{H,Bn}})]^+$  at  $-90^\circ\text{C}$  in acetone resulted in the formation of a superoxo complex  $[\text{Cu}(\text{L}^{\text{H,Bn}})(\text{O}_2)]^+$  as a less stable species and a peroxo complex  $[\{\text{Cu}(\text{L}^{\text{H,Bn}})\}_2(\text{O}_2)]^{2+}$  as a stable species. The structures of  $[\text{Cu}(\text{L}^{\text{H,Bn}})]\text{ClO}_4$  and  $[\{\text{Cu}(\text{L}^{\text{H,Bn}})\}_2(\text{O}_2)](\text{BPh}_4)_2 \cdot 8(\text{CH}_3)_2\text{CO}$  were determined by X-ray crystallography.  $[\{\text{Cu}(\text{L}^{\text{H,Bn}})\}_2(\text{O}_2)]^{2+}$  has a *trans*-( $\mu$ -1,2-peroxo)-dicopper(II) core with a trigonal bipyramidal structure. The O–O bond distance is 1.450(5) Å with an intermetallic Cu...Cu separation of 4.476(2) Å. The resonance Raman spectrum of  $[\{\text{Cu}(\text{L}^{\text{H,Bn}})\}_2(\text{O}_2)]^{2+}$  measured at  $-90^\circ\text{C}$  in acetone- $d_6$  showed a broad  $\nu(\text{O}-\text{O})$  band at 837–834  $\text{cm}^{-1}$  (788  $\text{cm}^{-1}$  for an  $^{18}\text{O}$  labeled sample) and two  $\nu(\text{Cu}-\text{O})$  bands at 556 and 539  $\text{cm}^{-1}$ , suggesting the presence of two peroxo species in solution.  $[\text{Cu}(\text{L}^{\text{Me,Bn}})]^+$  also produced both superoxo and *trans*- $\mu$ -1,2-peroxo species,  $[\text{Cu}(\text{L}^{\text{Me,Bn}})(\text{O}_2)]^+$  and  $[\{\text{Cu}(\text{L}^{\text{Me,Bn}})\}_2(\text{O}_2)]^{2+}$ . At a lower concentration of  $[\text{Cu}(\text{L}^{\text{Me,Bn}})]^+$  ( $\sim 0.24$  mM) and higher dioxygen concentration ( $P(\text{O}_2) = \sim 1$  atm), the superoxo species is predominantly formed, whereas at a higher concentration of  $[\text{Cu}(\text{L}^{\text{Me,Bn}})]^+$  ( $\sim 1$  mM) and lower dioxygen concentration ( $P(\text{O}_2) = \sim 0.02$  atm) the formation of the peroxo species is observed. The resonance Raman spectrum of  $[\text{Cu}(\text{L}^{\text{Me,Bn}})(\text{O}_2)]^+$  ( $\sim 1$  mM) in acetone- $d_6$  at  $\sim -95^\circ\text{C}$  exhibited a  $\nu(\text{O}-\text{O})$  band at 1120  $\text{cm}^{-1}$  (1059  $\text{cm}^{-1}$  for an  $^{18}\text{O}$  labeled sample) and that of  $[\{\text{Cu}(\text{L}^{\text{Me,Bn}})\}_2(\text{O}_2)]^{2+}$  ( $\sim 3$  mM) in acetone- $d_6$  at  $\sim -90^\circ\text{C}$  showed two  $\nu(\text{O}-\text{O})$  bands at 812 and 797  $\text{cm}^{-1}$  (767 and 753  $\text{cm}^{-1}$  for an  $^{18}\text{O}$  labeled sample),

respectively. A similar observation was also made for  $[\{\text{Cu}(\text{L}^{\text{Me,Me}})\}_2(\text{O}_2)]^{2+}$ . Relationships between the energies of the LMCT and  $d-d$  transitions and those of the  $\nu(\text{O}-\text{O})$  and  $\nu(\text{Cu}-\text{O})$  stretching vibrations and the steric constraints in the  $\text{Cu}(\text{II})-(\text{O}_2^{2-})-\text{Cu}(\text{II})$  core are discussed.

### X-E-9 Dinuclear Copper-Dioxygen Intermediates Supported by Polyamine Ligands

TERAMAE, Shinichi<sup>1</sup>; OSAKO, Takao<sup>2</sup>;  
NAGATOMO, Shigenori; KITAGAWA, Teizo;  
FUKUZUMI, Shunichi<sup>1</sup>; ITOH, Shinobu<sup>2</sup>  
(<sup>1</sup>Osaka Univ.; <sup>2</sup>Osaka City Univ.)

[*J. Inorg. Biochem.* **98**, 746–757 (2004)]

Reactivity of the dicopper(I) and dicopper(II) complexes supported by novel polyamine ligands **L1** (1,1-bis(6-methylpyridin-2-yl)-2,6,10-triaza-2,6,10-tribenzylundecane) and **L2** (5-benzyl-1,9-bis(6-methylpyridin-2-yl)-2,8-bis(6-methylpyridin-2-ylmethyl)-2,5,8-triazanonane) towards  $\text{O}_2$  and  $\text{H}_2\text{O}_2$ , respectively, has been investigated in order to shed light on the ligand effects on  $\text{Cu}_2/\text{O}_2$  chemistry. The dicopper(I) complex of **L1** (**1a**) readily reacted with  $\text{O}_2$  in a 2:1 ratio at a low temperature ( $-94^\circ\text{C}$ ) in acetone to afford a mixture of ( $\mu-\eta^2:\eta^2$ -peroxo)dicopper(II) and bis( $\mu$ -oxo)dicopper(III) complexes. The formation of these species has been confirmed by the electron spin resonance (ESR) silence of the solution as well as their characteristic absorption bands in the UV–visible region [max = 350 and 510 nm due to the peroxo complex and  $\sim 400$  nm due to the bis( $\mu$ -oxo) complex] and the resonance Raman bands at  $729\text{ cm}^{-1}$  [ $\Delta\nu(^{16}\text{O}_2-^{18}\text{O}_2) = 38\text{ cm}^{-1}$ ] due to the peroxo complex and at 611 and  $571\text{ cm}^{-1}$  [ $\Delta\nu(^{16}\text{O}_2-^{18}\text{O}_2) = 22$  and  $7\text{ cm}^{-1}$ , respectively] due to the bis( $\mu$ -oxo) complex. The peroxo and bis( $\mu$ -oxo) complexes were unstable even at the low temperature, leading to oxidative *N*-dealkylation at the ligand framework. The dicopper(I) complex of **L2** (**2a**) also reacted with  $\text{O}_2$  to give ( $\mu$ -hydroxo)dicopper(II) complex (**2b<sup>OH</sup>**) as the product. In this case, however, no active oxygen intermediate was detected even at the low temperature ( $-94^\circ\text{C}$ ). With respect to the copper(II) complexes, treatment of the ( $\mu$ -hydroxo)dicopper(II) complex of **L1** (**1b<sup>OH</sup>**) with an equimolar amount of  $\text{H}_2\text{O}_2$  in acetone at  $-80^\circ\text{C}$  efficiently gave a ( $\mu$ -1,1-hydroperoxo)dicopper(II) complex, the formation of which has been supported by its ESR-silence as well as UV–vis (370 and 650 nm) and resonance Raman spectra [ $881\text{ cm}^{-1}$ ;  $\Delta\nu(^{16}\text{O}_2-^{18}\text{O}_2) = 49\text{ cm}^{-1}$ ]. The ( $\mu$ -1,1-hydroperoxo)dicopper(II) intermediate of **L1** also decomposed slowly at the low temperature to give similar oxidative *N*-dealkylation products. Kinetic studies on the oxidative *N*-dealkylation reactions have been performed to provide insight into the reactivity of the active oxygen intermediates.

### X-E-10 Structural Characterization of a Binuclear Center of a Cu-Containing NO Reductase Homologue from *Roseobacter denitrificans*: EPR and Resonance Raman Studies

MATSUDA, Yuji<sup>1</sup>; UCHIDA, Takeshi; HORI, Hiroshi<sup>2</sup>; KITAGAWA, Teizo; ARATA, Hiroyuki<sup>1</sup>  
(<sup>1</sup>Kyushu Univ.; <sup>2</sup>Osaka Univ.)

[*Biochim. Biophys. Acta* **1656**, 37–45 (2004)]

Aerobic phototrophic bacterium *Roseobacter denitrificans* has a nitric oxide reductase (NOR) homologue with cytochrome *c* oxidase (CcO) activity. It is composed of two subunits that are homologous with NorC and NorB, and contains heme *c*, heme *b*, and copper in a 1:2:1 stoichiometry. This enzyme has virtually no NOR activity. Electron paramagnetic resonance (EPR) spectra of the air-oxidized enzyme showed signals of two low-spin hemes at 15 K. The high-spin heme species having relatively low signal intensity indicated that major part of heme *b*<sub>3</sub> is EPR-silent due to an antiferromagnetic coupling to an adjacent  $\text{Cu}_B$  forming a Fe–Cu binuclear center. Resonance Raman (RR) spectrum of the oxidized enzyme suggested that heme *b*<sub>3</sub> is six-coordinate high-spin species and the other hemes are six-coordinate low-spin species. The RR spectrum of the reduced enzyme showed that all the ferrous hemes are six-coordinate low-spin species.  $\nu(\text{Fe}-\text{CO})$  and  $\nu(\text{C}-\text{O})$  stretching modes were observed at  $523$  and  $1969\text{ cm}^{-1}$ , respectively, for CO-bound enzyme. In spite of the similarity to NOR in the primary structure, the frequency of  $\nu(\text{Fe}-\text{CO})$  mode is close to those of *aa*<sub>3</sub>- and *bo*<sub>3</sub>-type oxidases rather than that of NOR.

### X-E-11 Model Complexes of the Active Site of Galactose Oxidase. Effects of the Metal Ion Binding Sites

TAKI, Masayasu<sup>1</sup>; HATTORI, Haruna<sup>1</sup>; OSAKO, Takao<sup>1</sup>; NAGATOMO, Shigenori; SHIRO, Motoo<sup>2</sup>;  
KITAGAWA, Teizo; ITOH, Shinobu<sup>1</sup>  
(<sup>1</sup>Osaka City Univ.; <sup>2</sup>Rigaku Agency)

[*Inorg. Chim. Acta* **357**, 3369–3381 (2004)]

Model compounds of the active site of galactose oxidase have been developed by using new cofactor model ligands, **L1H** (2-methylthio-4-*tert*-butyl-6-[[bis(pyridin-2-ylmethyl)amino]methyl]phenol) and **L2H** (2-methylthio-4-*tert*-butyl-6-[[bis(6-methylpyridin-2-ylmethyl)amino]methyl]phenol). Treatment of the ligands with copper(II) and zinc(II) perchlorate in the presence of triethylamine followed by anion exchange reaction with  $\text{NaPF}_6$  or  $\text{NaBPh}_4$  provided the corresponding copper(II) and zinc(II) complexes, the crystal structures of which have been determined by X-ray crystallographic analysis. All the copper(II) and zinc(II) complexes have been isolated as a dimeric form in which the phenolate oxygen of each ligand acts as the bridging ligand to form a rhombic  $\text{M}_2(\text{OAr})_2$  core ( $\text{M} = \text{Cu}$  or  $\text{Zn}$ ). The dimeric complexes can be converted into the corresponding monomer complexes by the treatment with exogenous ligand such as acetate ion. The redox potential and the spectroscopic features of the monomer complexes have also been examined. Furthermore, the copper(II)- and zinc(II)-complexes of the phenoxyl radical species of the ligands have been generated in

situ by the oxidation of the phenolate complexes with  $(\text{NH}_4)_2[\text{Ce}^{\text{IV}}(\text{NO}_3)_6]$  (CAN) in  $\text{CH}_3\text{CN}$ , and their spectroscopic features have been explored. The structures and physicochemical properties of the phenolate and phenoxyl radical complexes of **L1** and **L2** have been compared to those of the previously reported copper(II) and zinc(II) complexes of **L3** (2-methylthio-4-*tert*-butyl-6-[[bis(2-pyridin-2-ylethyl)amino]methyl]phenol) in order to get insights into the interaction between the metal ions and the organic cofactor moiety.

#### X-E-12 Refolding Processes of Cytochrome P450<sub>cam</sub> from Ferric and Ferrous Acid Forms to the Native Conformation

EGAWA, Tsuyoshi<sup>1</sup>; HISHIKI, Takako<sup>2</sup>; ICHIKAWA, Yusuke<sup>2</sup>; KANAMORI, Yasukazu<sup>2</sup>; SHIMADA, Hideo<sup>2</sup>; TAKAHASHI, Satoshi<sup>3</sup>; KITAGAWA, Teizo; ISHIMURA, Yuzuru<sup>2</sup>  
(<sup>1</sup>IMS and Keio Univ.; <sup>2</sup>Keio Univ.; <sup>3</sup>IMS and Osaka Univ.)

[*J. Biol. Chem.* **279**, 32008–32017 (2004)]

Changes in heme coordination state and protein conformation of cytochrome P450<sub>cam</sub> (P450<sub>cam</sub>), a *b*-type heme protein, were investigated by employing pH jump experiments coupled with time-resolved optical absorption, fluorescence, circular dichroism, and resonance Raman techniques. We found a partially unfolded form (acid form) of ferric P450<sub>cam</sub> at pH 2.5, in which a Cys–heme coordination bond in the native conformation was ruptured. When the pH was raised to pH 7.5, the acid form refolded to the native conformation through a distinctive intermediate. Formations of similar acid and intermediate forms were also observed for ferrous P450<sub>cam</sub>. Both the ferric and ferrous forms of the intermediate were found to have an unidentified axial ligand of the heme at the 6th coordination sphere, which is vacant in the high spin ferric and ferrous forms at the native conformation. For the ferrous form, it was also indicated that the 5th axial ligand is different from the native cysteinate. The folding intermediates identified in this study demonstrate occurrences of non-native coordination state of heme during the refolding processes of the large *b*-type heme protein, being akin to the well known folding intermediates of cytochromes *c*, in which *c*-type heme is covalently attached to a smaller protein.

#### X-E-13 Simultaneous Resonance Raman Detection of the Heme $a_3$ -Fe-CO and Cu<sub>B</sub>-CO Species in CO-Bound $ba_3$ -Cytochrome *c* Oxidase from *Thermus thermophilus*

PINAKOULAKI, Eftychia<sup>1</sup>; OHTA, Takehiro; SOULIMANE, Tewfik<sup>2</sup>; KITAGAWA, Teizo; VAROTSIS, Constantinos<sup>3</sup>  
(<sup>1</sup>Univ. Crete; <sup>2</sup>Paul Scherrer Inst.; <sup>3</sup>IMS and Univ. Crete)

[*J. Biol. Chem.* **279**, 22791–22794 (2004)]

Understanding of the chemical nature of the dioxygen and nitric oxide moiety of  $ba_3$ -cytochrome *c*

oxidase from *Thermus thermophilus* is crucial for elucidation of its physiological function. In the present work, direct resonance Raman (RR) observation of the Fe–C–O stretching and bending modes and the C–O stretching mode of the Cu<sub>B</sub>–CO complex unambiguously establishes the vibrational characteristics of the heme-copper moiety in  $ba_3$ -oxidase. We assigned the bands at 507 and 568  $\text{cm}^{-1}$  to the Fe–CO stretching and Fe–C–O bending modes, respectively. The frequencies of these modes in conjunction with the C–O mode at 1973  $\text{cm}^{-1}$  showed, despite the extreme values of the Fe–CO and C–O stretching vibrations, the presence of the *a*-conformation in the catalytic center of the enzyme. These data, distinctly different from those observed for the  $caa_3$ -oxidase, are discussed in terms of the proposed coupling of the *a*- and *b*-conformations that occur in the binuclear center of heme-copper oxidases with enzymatic activity. The Cu<sub>B</sub>–CO complex was identified by its  $\nu(\text{CO})$  at 2053  $\text{cm}^{-1}$  and was strongly enhanced with 413.1 nm excitation indicating the presence of a metal-to-ligand charge transfer transition state near 410 nm. These findings provide, for the first time, RR vibrational information on the EPR silent Cu<sub>B</sub>(I) that is located at the O<sub>2</sub> delivery channel and has been proposed to play a crucial role in both the catalytic and proton pumping mechanisms of heme-copper oxidases.

#### X-E-14 Detection of a Photostable Five-Coordinate Heme $a_3$ -Fe-CO Species and Functional Implications of His384/ $\alpha$ 10 in CO-Bound $ba_3$ -Cytochrome *c* Oxidase from *Thermus thermophilus*

OHTA, Takehiro; PINAKOULAKI, Eftychia<sup>1</sup>; SOULIMANE, Tewfik<sup>2</sup>; KITAGAWA, Teizo; VAROTSIS, Constantinos<sup>1</sup>  
(<sup>1</sup>Univ. Crete; <sup>2</sup>Paul Scherrer Inst.; <sup>3</sup>IMS and Univ. Crete)

[*J. Phys. Chem. B* **108**, 5389–5491 (2004)]

Resonance Raman (RR) spectra are reported for the fully reduced carbon monoxy derivative of  $ba_3$ -cytochrome *c* oxidase from *Thermus thermophilus*. The RR spectra show the formation of a photolabile six-coordinate heme-CO and a photostable five-coordinate heme Fe–CO species. The latter species is formed by the cleavage of the proximal heme Fe–His384 bond and is the first five-coordinate Fe–CO species detected in hemecopper oxidases. The frequency of the Fe–CO species observed at 526  $\text{cm}^{-1}$  correlates with either the C–O stretching modes observed at 1967 or 1982  $\text{cm}^{-1}$  and lie on the correlation line of  $\nu(\text{Fe-CO})$  vs  $\nu(\text{C-O})$  for all known five-coordinate heme Fe–CO complexes. The loss of intensity of the heme Fe–His384 mode observed at 193  $\text{cm}^{-1}$  in the photostationary CO-bound spectra is attributed to the loss of the non-hydrogen bonded heme Fe–His384···Gly359 conformer. Taken together, the data indicate that the environment of theruptured His384 that is a part of the *Q*-proton pathway and leads to the highly conserved among all hemecopper oxidases, H<sub>2</sub>O pool, is disrupted upon CO binding to heme  $a_3$ .

### X-E-15 Heme Environment in Aldoxime Dehydratase Involved in Carbon–Nitrogen Triple Bond Synthesis

OINUMA, Ken-Ichi<sup>1</sup>; OHTA, Takehiro; KONISHI, Kazunobu<sup>1</sup>; HASHIMOTO, Yoshiteru<sup>1</sup>; HIGASHIBATA, Hiroki<sup>1</sup>; KITAGAWA, Teizo; KOBAYASHI, Michihiko<sup>1</sup>  
(<sup>1</sup>Univ. Tsukuba)

[*FEBS Lett.* **568**, 44–48 (2004)]

Resonance Raman spectra have been measured to characterize the heme environment in aldoxime dehydratase (OxDa), a novel hemoprotein, which catalyzes the dehydration of aldoxime into nitrile. The spectra showed that the ferric heme in the enzyme is six-coordinate low spin, whereas the ferrous heme is five-coordinate high spin. We assign a prominent vibration that occurs at 226 cm<sup>-1</sup> in the ferrous enzyme to the Fe-proximal histidine stretching vibration. In the CO-bound form of OxDa, the correlation between the Fe–CO stretching (512 cm<sup>-1</sup>) and C–O stretching (1950 cm<sup>-1</sup>) frequencies also supports our assignment of proximal histidine coordination.

### X-E-16 Interactions of Soluble Guanylate Cyclase with Diatomics as Probed by Resonance Raman Spectroscopy

PAL, Biswajit; KITAGAWA, Teizo

[*J. Inorg. Biochem.* in press]

Soluble guanylate cyclase (sGC, EC 4.6.1.2) acts as a sensor for nitric oxide (NO), but is also activated by carbon monoxide in the presence of an allosteric modulator. Resonance Raman studies on the structure-function relations of sGC are reviewed with a focus on the CO-adduct in the presence and absence of allosteric modulator, YC-1, and substrate analogues. It is demonstrated that the sGC isolated from bovine lung contains one species with a five coordinate (5c) ferrous high-spin (HS) heme with the Fe–His stretching mode at 204 cm<sup>-1</sup>, but its CO adduct yields two species with different conformations about the heme pocket with the Fe–CO stretching ( $\nu_{\text{Fe-CO}}$ ) mode at 473 and 489 cm<sup>-1</sup>, both of which are His- and CO-coordinated 6c ferrous adducts. Addition of YC-1 to it changes their population and further addition of GTP yields one kind of 6c ( $\nu_{\text{Fe-CO}} = 489 \text{ cm}^{-1}$ ) in addition to 5c CO-adduct ( $\nu_{\text{Fe-CO}} = 521 \text{ cm}^{-1}$ ). Under this condition the enzymatic activity becomes nearly the same level as that of NO adduct. Addition of  $\gamma$ -S-GTP yields the same effect as GTP does but sGMP and GDP gives much less effects. Unexpectedly, ATP cancels the effects of GTP. The structural meaning of these spectroscopic observation is discussed in detail.

### X-E-17 Resonance Raman Evidence for the Presence of Two Heme Pocket Conformations with Varied Activities in CO-Bound Bovine Soluble Guanylate Cyclase and Their Conversion

LI, Zhengqiang<sup>1</sup>; PAL, Biswajit; TAKENAKA, Shigeo<sup>2</sup>; TSUYAMA, Shingo<sup>2</sup>; KITAGAWA, Teizo  
(<sup>1</sup>IMS and Jilin Univ.; <sup>2</sup>Osaka Prefecture Univ.)

[*Biochemistry* in press]

It was noted previously that resonance Raman (RR) spectra of soluble guanylate cyclase (sGC) observed by five independent research groups were categorized into two types; sGC<sub>1</sub> and sGC<sub>2</sub> (Vogel, K. M., Hu, S. Z., Spiro, T. G., Dierks, E. A., Yu, A. E., and Burstyn, J. N., *J. Biol. Inorg. Chem.* **4**, 804–813 (1999)). We demonstrate here that the RR spectra of sGC isolated from bovine lung contains only sGC<sub>2</sub> but both species for CO-bound form (CO-sGC). The relative population of the two forms was changed from the initial CO-sGC<sub>2</sub> form dominant with the Fe–CO ( $\nu_{\text{Fe-CO}}$ ) and C–O stretching modes ( $\nu_{\text{CO}}$ ) at 472 and 1990 cm<sup>-1</sup>, respectively, to the CO-sGC<sub>1</sub> form with  $\nu_{\text{Fe-CO}}$  and  $\nu_{\text{CO}}$  at 488 and 1973 cm<sup>-1</sup> by adding a xenobiotic, YC-1. Further addition of a substrate, GTP, completed the change. GDP and cGMP had much less effects but a substrate analogue, GTP- $\gamma$ -S was found to have the same effect as GTP. In contrast, ATP has the reverse effect, namely deleted the effect of YC-1 and GTP. In the coexistence of YC-1 and GTP, vinyl vibrations of heme are largely influenced and new CO-isotope sensitive bands were observed at 521, 488, 363, and 227 cm<sup>-1</sup>. The 521 cm<sup>-1</sup> band was assigned to the five-coordinate (5c) species from the model compound studies using ferrous iron-protoporphyrin IX in CTAB micelles. Distinctively from the 472 cm<sup>-1</sup> species, both the 488-cm<sup>-1</sup> and 521-cm<sup>-1</sup> species were apparently unphotodissociable when an ordinary Raman spinning cell was used, meaning rapid recombination of photodissociated CO. On the basis of these observations, binding of YC-1 to the heme pocket is proposed.

### X-E-18 SOUL in Mouse Eyes Is a Novel Hexameric Heme-Binding Protein with Characteristic Optical Absorption, Resonance Raman Spectral and Heme Binding Properties

SATO, Emiko<sup>1</sup>; SAGAMI, Ikuko<sup>2</sup>; UCHIDA, Takeshi; SATO, Akira<sup>3</sup>; KITAGAWA, Teizo; IGARASHI, Jotaro<sup>1</sup>; OLSON, John S.<sup>4</sup>; SHIMIZU, Toru<sup>1</sup>  
(<sup>1</sup>Tohoku Univ.; <sup>2</sup>Tohoku Univ. and Kyoto Prefectural Univ.; <sup>3</sup>GUAS; <sup>4</sup>Rice Univ.)

[*Biochemistry* in press]

SOUL is specifically expressed in the retina and pineal gland, and displays more than 40% sequence homology with p22HBP, a heme protein ubiquitously expressed in numerous tissues. SOUL was purified as a dimer in the absence of heme from the *E. coli* expression system, but displayed a hexameric structure upon heme binding. Heme-bound SOUL displayed optical absorption and resonance Raman spectra typical of 6-coordinate low-spin heme protein, with one heme per monomeric unit for both the Fe(III) and Fe(II) complexes. Spectral data additionally suggest that one of the axial ligands of the Fe(III) heme complex is His. Muta-

tion of His42 (the only His of SOUL) to Ala resulted in loss of heme binding, confirming that this residue is an axial ligand of SOUL. The  $K_d$  value of heme for SOUL was estimated as  $4.8 \times 10^{-9}$  M from the association and dissociation rate constants, suggesting high binding affinity. On the other hand, p22HBP was obtained as a monomer containing one heme per subunit, with a  $K_d$  value of  $2.1 \times 10^{-11}$  M. Spectra of heme-bound p22HBP were different from those of SOUL, but similar to those of heme-bound bovine serum albumin in which heme bound to a hydrophobic cavity with no specific axial ligand coordination. Therefore, the heme-binding properties and coordination structure of SOUL are distinct from those of p22HBP, despite high sequence homology. The physiological role of the new heme binding protein, SOUL, is further discussed in this report.

### X-E-19 Quaternary Structures of Intermediately Ligated Human Hemoglobin A and Influences from Strong Allosteric Effectors; Resonance Raman Investigation

NAGATOMO, Shigenori; NAGAI, Masako<sup>1</sup>;  
MIZUTANI, Yasuhisa<sup>2</sup>; YONETANI, Takashi<sup>3</sup>;  
KITAGAWA, Teizo

(<sup>1</sup>Kanazawa Univ.; <sup>2</sup>IMS and Kobe Univ.; <sup>3</sup>Univ. Pennsylvania)

[*Biophys. J.* submitted]

The Fe-histidine stretching ( $\nu_{\text{Fe-His}}$ ) frequency was determined for deoxy subunits of intermediately ligated human hemoglobin A in equilibrium and CO-photo-dissociated picosecond transient species in the presence and absence of strong allosteric effectors like inositol (hexakis)phosphate (IHP), bezafibrate (BZF) and 2,3-bisphosphoglycerate (BPG). The  $\nu_{\text{Fe-His}}$  frequency of deoxyHb A was unaltered by the effectors. The T to R transition occurred around  $m = 2 \sim 3$  in the absence of effectors but  $m > 3.5$  in their presence, where  $m$  is the average number of ligands bound to Hb and was determined from the intensity of the  $\nu_4$  band measured in the same experiment. The  $\alpha_1$ - $\beta_2$  subunit contacts revealed by UV resonance Raman spectra, which were distinctly different between the T and R states, remained unchanged by the effectors. This observation would solve the recent discrepancy that the strong effectors remove the cooperativity of oxygen binding in the low affinity limit whereas the <sup>1</sup>H NMR spectrum of fully ligated form exhibits the pattern of R state.

### X-E-20 Oxygen Sensing Mechanism of HemAT from *Bacillus subtilis*: A Resonance Raman Spectroscopic Study

OHTA, Takehiro; YOSHIMURA, Hideaki;  
YOSHIOKA, Shiro; AONO, Shigetoshi;  
KITAGAWA, Teizo

[*J. Am. Chem. Soc.* submitted]

HemAT-*Bs* is a heme-based signal transducer protein responsible for aerotaxis of *Bacillus subtilis*, which detects oxygen and transmits the signal to regulatory proteins that control the direction of flagella rotation.

CO and NO are also caught at the same position as O<sub>2</sub> but the signals would be differentiated. Binding of oxygen to the sensor domain of this protein is supposed to alter the protein conformation in the vicinity of heme, which is propagated to the signaling domain through the linker region in a way different from the other case in binding of other gases.

Specific sensing of O<sub>2</sub>, CO, and NO might have been required for the aerophilic bacteria in the early times of the earth, when CO and NO were more abundant than O<sub>2</sub>. In support of this idea our previous resonance Raman (RR) study of the oxygen bound form of HemAT-*Bs* has demonstrated that the Fe–O<sub>2</sub> stretching ( $\nu_{\text{Fe-O}_2}$ ) frequency ( $560 \text{ cm}^{-1}$ ) is noticeably lower than those of general oxygen bound hemoproteins, but similar to the frequencies observed for invertebrate, plant, and bacterial Hbs, suggesting that the bound oxygen is incorporated into a unique hydrogen bonding network in the distal environment. Here we present RR evidences for structural linkage between the distal heme pocket and the signaling domain by using the linker-lacking protein as well as the wild type and the Y70F and T95A mutants of full-length

### X-E-21 The Interaction of Covalently Bound Heme with the Cytochrome *c* Maturation Protein CcmE

UCHIDA, Takeshi; STEVENS, Julie M.<sup>1</sup>;  
DALTRUP, Oliver<sup>1</sup>; HARVAT, Edgar M.<sup>1</sup>; HONG,  
Lin<sup>1</sup>; FERGUSON, Stuart J.<sup>1</sup>; KITAGAWA, Teizo  
(<sup>1</sup>Univ. Oxford)

[*J. Biol. Chem.* submitted]

The heme chaperone CcmE is a novel protein that binds heme covalently *via* a histidine residue as part of its essential function in the process of cytochrome *c* biogenesis in many bacteria as well as plant mitochondria. In the continued absence of a structure of the holo-form of CcmE, identification of the heme ligands is an important step in understanding the molecular function of this protein and the role of covalent heme binding to CcmE during the maturation of *c*-type cytochromes. In this work we present spectroscopic data that provide insight into the ligation of the heme iron in the soluble domain of CcmE from *E. coli*. Resonance Raman spectra demonstrated that one of the heme axial ligands is a histidine residue and the other is likely to be Tyr134. In addition, the properties of the heme resonances of the holo-protein compared with those of a form of CcmE with non-covalently bound heme provide evidence for the modification of one of the heme vinyl side chains by the protein, most likely the 2-vinyl group.

### X-E-22 Role of Tyr288 at the Dioxygen Reduction Site of Cytochrome *bo* Studied by Stable Isotope Labeling and Site-Directed Mutagenesis

UCHIDA, Takeshi; MOGI, Tatsushi<sup>1</sup>;  
NAKAMURA, Hiro<sup>2</sup>; KITAGAWA, Teizo  
(<sup>1</sup>Univ. Tokyo, Tokyo Inst. Tech. and ERATO(JST);  
<sup>2</sup>RIKEN Harima Inst. and Yokohama City Univ.)

[*J. Biol. Chem.* submitted]

To explore the role of a cross-link between side-chains of Tyr288 and His284 at the heme-copper binuclear center, we prepared cytochrome *bo* where D<sub>4</sub>-, 1-<sup>13</sup>C-, or 4-<sup>13</sup>C-Tyr has been biosynthetically incorporated. Unexpectedly, the D<sub>4</sub>-Tyr-labeled enzyme showed the large decrease in the ubiquinol-1 oxidase and CO-binding activities. Optical absorption and resonance Raman spectra identified the defect in the distal side of the heme-copper binuclear center. In the CO-bound D<sub>4</sub>-Tyr-labeled enzyme, a large fraction of the  $\nu_{(\text{Fe}-\text{C})}$  mode was shifted from the normal 520 cm<sup>-1</sup>-band to a broad band centered around 491 cm<sup>-1</sup>, as found for the Y288F mutant. Our results suggest that the substitution of ring hydrogens of Tyr288 with deuteriums slows down the formation of the His-Tyr cross-link essential for the dioxygen reduction at the binuclear center.

### X-E-23 Resonance Raman Characterization of the PAS-A Domain of the Nobel CO-Dependent Gene Regulatory Protein, NPAS2

UCHIDA, Takeshi; SATO, Emiko<sup>1</sup>; SATO, Akira<sup>2</sup>; SAGAMI, Ikuko<sup>3</sup>; SHIMIZU, Toru<sup>1</sup>; KITAGAWA, Teizo

(<sup>1</sup>Tohoku Univ.; <sup>2</sup>GUAS; <sup>3</sup>Tohoku Univ. and Kyoto Prefectural Univ.)

[*J. Biol. Chem.* submitted]

Neuronal PAS domain protein 2, a newly discovered as a heme protein, is expressed in the mammalian forebrain and acts as a CO-dependent transcriptional activator. This protein consists of the N-terminal basic helix-loop-helix domain, and two heme-containing PAS domains (PAS-A and PAS-B). In this study we prepared the isolated PAS-A domain and its mutants, and measured resonance Raman spectra. The CO-bound form gave the  $\nu_{\text{Fe}-\text{CO}}$  and  $\nu_{\text{C}-\text{O}}$  bands at 497 and 1967 cm<sup>-1</sup>, respectively, and the correlation plot between  $\nu_{\text{Fe}-\text{CO}}$  and  $\nu_{\text{C}-\text{O}}$  suggested that a neutral His is a trans ligand of CO. The ferric form is constituted of the dominant 6-coordinate low-spin species and minor 5- and 6-coordinate high-spin species. When its Raman spectrum was excited at 363.8 nm, an intense band assignable to the Fe<sup>3+</sup>-S stretching was observed at 332 cm<sup>-1</sup>, whereas it disappeared in the C170A mutant, suggesting that Cys170 is an axial ligand in the ferric state. The spectrum of the wild-type ferrous PAS-A domain shows a mixture of 5-coordinate high-spin and 6-coordinate low-spin hemes. In the H119A and H171A mutants, the 5-coordinate species increased, while no change was observed for the C170A mutant, which suggest that His119 and His171, not Cys170, are the axial ligands in the ferrous heme, and ligand replacement from Cys to His takes place upon heme reduction. The marker band  $\nu_{11}$  of the reduced form, which is sensitive to the donor strength of the axial ligand, was shifted to a lower frequency than that of cytochrome *c*<sub>3</sub>, suggesting the coordination of a deprotonated histidine. Taken together, the present results support a mechanism that CO binding to heme

causes conformation change in the His171-Cys170 moiety, which leads to physiological signaling.