

Observation of the Fe–O₂ and Fe^{IV}=O stretching Raman bands for dioxygen reduction intermediates of cytochrome *bo* isolated from *Escherichia coli***

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Abstract Reaction intermediates in dioxygen reduction by the *E. coli* cytochrome *bo*-type ubiquinol oxidase were studied by time-resolved resonance Raman spectroscopy using the artificial cardiovascular system. At 0–20 μ s following photolysis of the enzyme–CO adduct in the presence of O₂, we observed the Fe–O₂ stretching Raman band at 568 cm^{–1} which shifted to 535 cm^{–1} with the ¹⁸O₂ derivative. These frequencies are remarkably close to those of other oxyhemoproteins including dioxygen-bound hemoglobin and *aa*₃-type cytochrome *c* oxidase. In the later time range (20–40 μ s), other oxygen-isotope-sensitive Raman bands were observed at 788 and 361 cm^{–1}. Since the 781 cm^{–1} band exhibited a downshift by 37 cm^{–1} upon ¹⁸O₂ substitution, we assigned it to the Fe^{IV}=O stretching mode. This band is considered to arise from the ferryl intermediate, but its appearance was much earlier than the corresponding intermediate of bovine cytochrome *c* oxidase (>100 μ s). The 361 cm^{–1} band showed the ¹⁶O/¹⁸O isotopic frequency shift of 14 cm^{–1} similar to the case of bovine cytochrome *c* oxidase reaction.

Key words: *bo*-Type quinol oxidase; Oxy intermediate; Ferryl intermediate; Iron–dioxygen stretching mode; Ferryl–oxo stretching mode; Time-resolved resonance Raman spectroscopy

1. Introduction

Cytochrome *bo*-type ubiquinol oxidase is a terminal oxidase of the aerobic respiratory chain of *Escherichia coli* (*E. coli*) and is predominantly expressed under highly-aerated growth conditions (see [1] for a review). It catalyzes the four-electron reduction of molecular oxygen at the cytoplasmic side of membranes, which is coupled with the two-electron oxidation of two ubiquinol-8 molecules at the periplasmic side. Redox reactions at different surfaces of the membrane can create a difference in electrochemical potential across the cytoplasmic membrane. In addition to the protolytic scalar reactions, this enzyme is known to function as an electron transfer-linked proton-pump [2] like bovine *aa*₃-type cytochrome *c* oxidase. Recent molecular biological [3–7] and physicochemical [8–10] studies demonstrated that the *bo*-type quinol oxidase belongs to the heme-copper oxidase superfamily and shares a common molecular mechanism for the redox-coupled proton pumping [11].

Subunit I of the *E. coli bo*-type quinol oxidase contains a hexa-coordinated low-spin heme B, a penta-coordinated high-spin heme O and one copper ion (Cu_B) (see [3,12] for recent reviews). The high-spin heme and Cu_B are antiferromagnetically coupled, forming an Fe–Cu_B binuclear center where dioxygen reduction takes place. Subunit II provides the oxidation site for a lipophilic two-electron donor, ubiquinol-8 [13], but does not contain the Cu_A center which accepts electrons from ferrous cytochrome *c* for mammalian cytochrome *c* oxidases. Thus, electrons are transferred from the quinol bound to

subunit II to the binuclear center of subunit I via the low-spin heme.

Resonance Raman (RR) spectroscopy can bring about structural information on hemes and their vicinities [14,15], and is especially powerful for studies of reaction intermediates. In fact, for the reaction of reduced bovine cytochrome *c* oxidase with dioxygen, the oxy [16–18], ferryl [19–21], and hydroxy [20,22] intermediates have been identified by this technique. It is extremely interesting to examine similar intermediates for the reaction of cytochrome *bo* with O₂, although there are large differences between *bo*-type quinol oxidase and *aa*₃-type cytochrome *c* oxidase, regarding electron donors, presence or absence of the Cu_A center, and heme species at the dioxygen reduction site. Therefore, we applied time-resolved RR spectroscopy using the artificial cardiovascular system [23,24] successfully to characterize intermediates involved in dioxygen reduction by *bo*-type quinol oxidase. We detected the Fe–O₂ and Fe^{IV}=O stretching modes at 568 and 788 cm^{–1}, respectively, for the first time, suggesting that the dioxygen reduction mechanism of quinol oxidases is similar to that of *aa*₃-type cytochrome *c* oxidases.

2. Materials and methods

Cytochrome *bo*-type quinol oxidase was purified from the cytochrome *bd*-deficient strain GO103 (Δ *cyd*::Km^r [25]) harboring pHN3795–1 (H. Nakamura, unpublished results), as described previously [8]. pHN3795–1 is a derivative of pBR322 which carries the cytochrome *bo* operon (*cyoABCDE**) and was obtained from pHN3795 [26] as a clone that can support the aerobic growth of the Δ *cyo* Δ *cyd* strain on a nonfermentable carbon source.

The enzyme was dissolved in 100 mM Tris–HCl buffer (pH 7.4) containing 1.0% sucrose monolaurate SM-1200 (Mitsubishi-Kasei Food Co. Ltd., Tokyo), 200 μ M ubiquinone-1 (a kind gift from Dr. S. Ohsono, Eisai Co. Ltd., Tsukuba) and 100 mM sodium ascorbate

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(pH 7.4). About 75 ml of the 50 μ M enzyme solution in the fully reduced CO-bound form was circulated at 5°C at a flow rate of 20 or 40 ml/min and brought under oxygen concentration of 150 μ M using the artificial cardiovascular system. Details of this system are described elsewhere [23,24]. Raman scattering was excited at 406.7 nm by a Kr⁺ ion laser (Spectra Physics, Model 2016), and detected with a cooled (−20 °C) diode array (PAR 1421HQ) attached to a single polychromator (Ritsu Oyo Kogaku, DG-1000), for which the slit width of 200 μ m and slit height of 10 mm were used. A single laser beam with a power of ~5 mW was focused to 40 μ m on a flow cell with a cross section of 0.6 \times 0.6 mm² and was used to photolyze CO for initiation of the reaction and also to excite RR scattering. The transit time of a given molecule across the laser beam was 40 or 20 μ s according to the flow rate used. Raman shifts were calibrated with ethanol as a standard under the same illumination geometry.

3. Results and discussion

Fig. 1 shows time-resolved RR spectra in the 1000 to 300 cm^{-1} region for reaction intermediates of the *bo*-type quinol oxidase with ¹⁶O₂ (A and C) and ¹⁸O₂ (B and D). Spectra A and B represent the raw spectra of intermediates generated in the time range of 0–20 μ s following photolysis, while spectra C and D display those around 0–40 μ s following photolysis. Spectra E and F illustrate the differences between spectra A and B, and between spectra C and D, respectively, in which an intense porphyrin band at 678 cm^{-1} was used as a marker for subtraction.

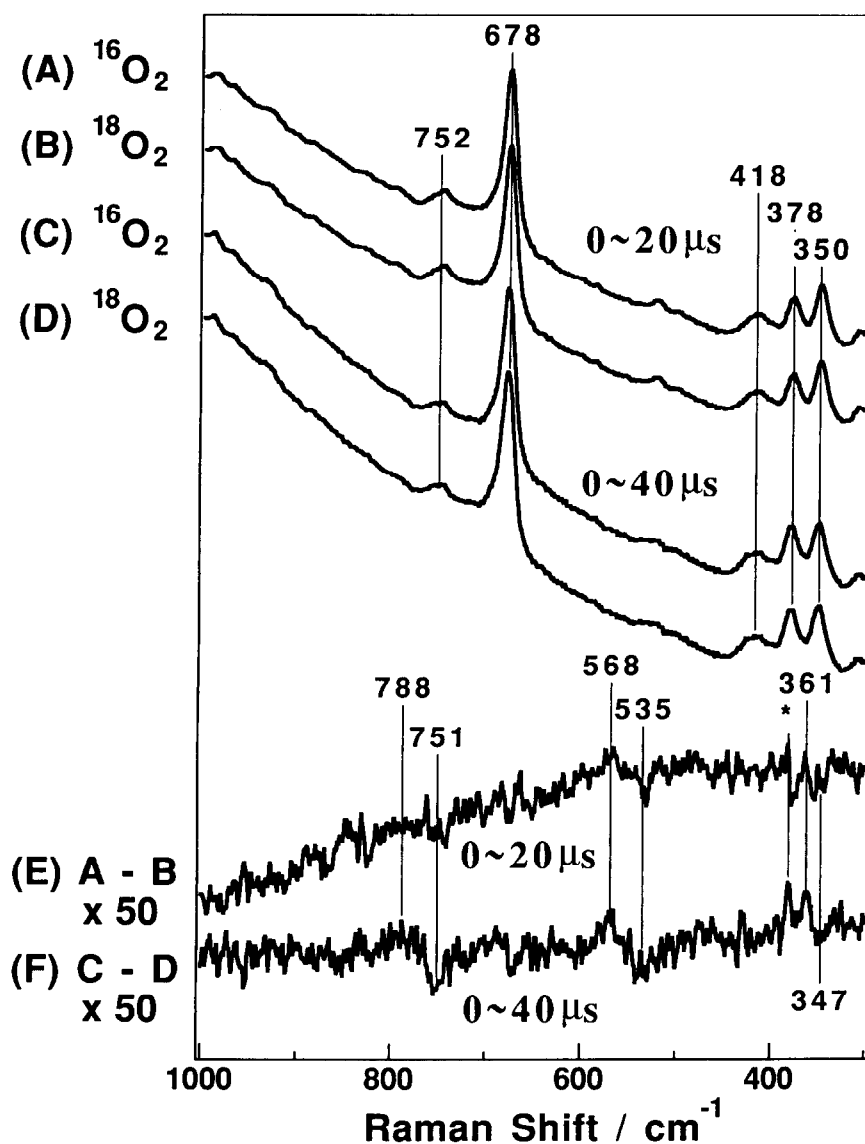


Fig. 1. Time-resolved RR spectra of the *E. coli bo*-type quinol oxidase in the 1000 to 300 cm^{-1} region for ¹⁶O₂ (A,C) and ¹⁸O₂ (B,D) derivatives and their difference spectra (E,F). The ordinate scales of spectra (A), (B), (C) and (D) are normalized by the intensity of porphyrin bands. Traces E and F represent the differences; spectrum E = (spectrum A – spectrum B) \times 50 and spectrum F = (spectrum C – spectrum D) \times 50. The bands marked with an asterisk denote a plasma line from a Kr⁺ ion laser. Transit time of a given molecule across the laser beam is 0–20 μ s for spectra A, B, and E and 0–40 μ s for spectra C, D, and F. Experimental conditions: excitation, 406.7 nm, 5 mW (at the sample) for spectra A and B, and 6 mW (at the sample) for spectra C and D; accumulation time, 3840 s for spectra A and B, and 6080 s for spectra C and D.

A differential pattern exhibiting a peak at 568 cm^{-1} and a trough at 535 cm^{-1} appeared in difference spectra E and F. The frequencies 568 cm^{-1} for $^{16}\text{O}_2$ and 535 cm^{-1} for $^{18}\text{O}_2$ are in reasonable agreement with the $\text{Fe}-\text{O}_2$ stretching frequencies reported for oxyhemoglobin [27,28], oxymyoglobin [29,30], and dioxygen-bound bovine cytochrome *c* oxidase [16–18,24]. Cryogenic flash/trap absorption experiments on the CO adduct of this enzyme in the presence of oxygen suggested the formation of an oxygenated heme at low temperatures [31]. Therefore we assign this band to the $\text{Fe}-\text{O}_2$ stretching mode ($\nu_{\text{Fe}-\text{O}_2}$) of the dioxygen adduct. Note that the $\nu_{\text{Fe}-\text{O}_2}$ frequencies of the *bo*-type oxy-quinol oxidase ($568/535\text{ cm}^{-1}$ for $^{16}\text{O}_2/^{18}\text{O}_2$) and *aa₃*-type oxy-cytochrome *c* oxidase ($571/545\text{ cm}^{-1}$) are alike, suggesting similarity in their $\text{Fe}-\text{O}-\text{O}$ geometry.

Spectrum F shows the presence of other oxygen-isotope-sensitive bands around 785 and 361 cm^{-1} which must be arising from the species generated around $20\text{--}40\text{ }\mu\text{s}$ following the start of the reaction. The 785 cm^{-1} band is shifted to 751 cm^{-1} with the $^{18}\text{O}_2$ derivative. The frequency and its $^{18}\text{O}/^{16}\text{O}$ -isotopic shift (37 cm^{-1}) are close to those observed for the $\text{Fe}^{\text{IV}}=\text{O}$ stretching mode ($\nu_{\text{Fe}^{\text{IV}}=\text{O}}$) of the oxoferryl intermediate for the bovine cytochrome *c* oxidase [19–21], of horseradish peroxidase compound II [32–35], and of other oxoferryl hemeprotein species [36–39]. Therefore we assign this band to the $\nu_{\text{Fe}^{\text{IV}}=\text{O}}$ mode of the oxoferryl intermediate. In the case of bovine cytochrome *c* oxidase, there are two oxygen-isotope-sensitive bands in this frequency region (786 and 804 cm^{-1}) [22,24], but it is not clear from this experiment whether the 785 cm^{-1} feature in Fig. 1 is a single band or not. It should be noted that the rise time (ca. $20\text{--}40\text{ }\mu\text{s}$ at 5°C) of the oxoferryl intermediate of the *bo*-type quinol oxidase is significantly faster than that of bovine cytochrome *c* oxidase, in which the $\nu_{\text{Fe}^{\text{IV}}=\text{O}}$ band was undetectable before 100 ms under similar experimental conditions except for removal of detergents for bovine cytochrome *c* oxidase [24]. Recently, we found that the *E. coli bo*-type quinol oxidase contains a tightly-bound ubiquinone-8 (Q_H) which exists in the proximity of both the quinol oxidation site (Q_L) and the low-spin heme and may serve as a pathway for an electron transfer between these two centers [40]. The faster decay of the oxy intermediate to the oxoferryl intermediate in the *bo*-type quinol oxidase may be due to the unique electron transfer pathway. The other oxygen-isotope-sensitive band at 361 cm^{-1} showed an $^{16}\text{O}/^{18}\text{O}$ isotopic shift of 14 cm^{-1} . A similar band is also reported for an intermediate in dioxygen reduction by bovine cytochrome *c* oxidase [21,24,41], although the nature of the species giving this band has not been characterized yet.

In conclusion, we observed oxygen-isotope-sensitive Raman bands at 568 and 788 cm^{-1} during turnovers of the *E. coli bo*-type quinol oxidase for the first time, and assigned them to the $\nu_{\text{Fe}-\text{O}_2}$ and $\nu_{\text{Fe}^{\text{IV}}=\text{O}}$ of its oxy- and ferryl intermediates, respectively. These frequencies and their $^{16}\text{O}/^{18}\text{O}$ isotopic shifts suggest that structures of these intermediates at the catalytic site are similar to the corresponding intermediates of *aa₃*-type cytochrome *c* oxidases. However, the occurrence of the ferryl intermediate was evidently faster in the *bo*-type quinol oxidase than in the *aa₃*-type cytochrome *c* oxidase.

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References

- [1] Anraku, Y. and Gennis, R.B. (1987) Trends Biochem. Sci. 12, 262–266.
- [2] Puustinen, A., Finel, M., Virkki, M. and Wikström, M. (1989) FEBS Lett. 249, 163–167.
- [3] Mogi, T., Nakamura, H. and Anraku, Y. (1994) J. Biochem. in press.
- [4] Chepur, V., Lemieux, L., Au, D.C.-T. and Gennis, R.B. (1990) J. Biol. Chem. 265, 11185–11192.
- [5] Minagawa, J., Mogi, T., Gennis, R.B. and Anraku, Y. (1992) J. Biol. Chem. 267, 2096–2104.
- [6] Lemieux, L.J., Calhoun, M.W., Thomas, J.W., Ingledew, W.J. and Gennis, R.B. (1992) J. Biol. Chem. 267, 2105–2113.
- [7] Shapleigh, J.P., Hosler, J.P., Tecklenburg, M.M.J., Kim, Y., Babcock, G.T., Gennis, R.B. and Ferguson-Miller, S. (1992) Proc. Natl. Acad. Sci. USA 89, 4786–4790.
- [8] Tsubaki, M., Mogi, T., Anraku, Y. and Hori, H. (1993) Biochemistry 32, 6065–6072.
- [9] Salerno, J.C., Bolgiano, B., Poole, R.K., Gennis, R.B. and Ingledew, W.J. (1990) J. Biol. Chem. 265, 4364–4368.
- [10] Uno, T., Mogi, T., Tsubaki, M., Nishimura, Y. and Anraku, Y. (1994) J. Biol. Chem. 269, 11912–11920.
- [11] Babcock, G.T. and Wikström, M. (1992) Nature 356, 301–309.
- [12] Mogi, T., Saiki, K. and Anraku, Y. (1994) Mol. Microbiol., in press.
- [13] Welter, R., Gu, L.-Q., Yu, C.-A., Rumbley, J. and Gennis, R.B. (1994) Biophys. J. 66, A367.
- [14] Yu, N.-T. (1986) Methods Enzymol. 130, 350–409.
- [15] Yu, N.-T. and Kerr, E.A. (1988) in: Biological Applications of Raman Spectroscopy (Spiro, T.G., Ed.) Vol. 3, pp. 39–95, Wiley-Interscience, New York.
- [16] Ogura, T., Takahashi, S., Shinzawa-Itoh, K., Yoshikawa, S. and Kitagawa, T. (1990) J. Am. Chem. Soc. 112, 5630–5631.
- [17] Han, S., Ching, Y.-C. and Rousseau, D.L. (1990) Proc. Natl. Acad. Sci. USA 87, 2491–2495.
- [18] (a) Varotsis, C., Woodruff, W.H. and Babcock, G.T. (1989) J. Am. Chem. Soc. 111, 6439–6440. (b) Varotsis, C., Woodruff, W.H. and Babcock, G.T. (1989) J. Am. Chem. Soc. 112, 1297.
- [19] Ogura, T., Takahashi, S., Shinzawa-Itoh, K., Yoshikawa, S. and Kitagawa, T. (1990) J. Biol. Chem. 265, 14721–14723.
- [20] Han, S., Ching, Y.-C. and Rousseau, D.L. (1990) Nature 348, 89–90.
- [21] Varotsis, C. and Babcock, G.T. (1990) Biochemistry 29, 7357–7362.
- [22] Ogura, T., Takahashi, S., Shinzawa-Itoh, K., Yoshikawa, S. and Kitagawa, T. (1991) Bull. Chem. Soc. Jpn. 64, 2901–2907.
- [23] Ogura, T., Yoshikawa, S. and Kitagawa, T. (1989) Biochemistry 28, 8022–8027.
- [24] Ogura, T., Takahashi, S., Hirota, S., Shinzawa-Itoh, K., Yoshikawa, S., Appelman, E.H. and Kitagawa, T. (1993) J. Am. Chem. Soc. 115, 8527–8536.
- [25] Oden, K.L., DeVaux, L.C., Vibat, C.R.T., Cronan, J.E., Jr. and Gennis, R.B. (1990) Gene 96, 29–36.
- [26] Nakamura, H., Yamato, I., Anraku, Y., Lemieux, L. and Gennis, R.B. (1990) J. Biol. Chem. 265, 11193–11197.
- [27] Brunner, H. (1974) Naturwissenschaften 61, 129.
- [28] Nagai, K., Kitagawa, T. and Morimoto, H. (1980) J. Mol. Biol. 136, 271–289.
- [29] Kerr, E.A., Yu, N.-T., Bartnicki, D.E. and Mizukami, H. (1985) J. Biol. Chem. 260, 8360–8365.
- [30] Hirota, S., Ogura, T., Appelman, E.H., Shinzawa-Itoh, K., Yoshikawa, S. and Kitagawa, T. (1994) submitted.
- [31] Poole, R.K., Waring, A.J. and Chance, B. (1979) FEBS Lett. 101, 56–58.
- [32] Sitter, A.J., Reczek, C.M. and Terner, J. (1985) J. Biol. Chem. 260, 7515–7522.
- [33] Terner, J., Sitter, A.J. and Reczek, C.M. (1985) Biochim. Biophys. Acta 828, 73–80.
- [34] Hashimoto, S., Tatsuno, Y. and Kitagawa, T. (1986) Proc. Natl. Acad. Sci. USA 83, 2417–2421.
- [35] Hashimoto, S., Tatsuno, Y. and Kitagawa, T. (1984) Proc. Jpn. Acad. Ser. B. Phys. Biol. Sci. 60, 345–348.

- [36] Sitter, A.J., Reczek, C.M. and Terner, J. (1985) *Biochim. Biophys. Acta* 828, 229–235.
- [37] Hashimoto, S., Teraoka, J., Inubushi, T., Yonetani, T. and Kitagawa, T. (1986) *J. Biol. Chem.* 261, 11110–11118.
- [38] Oertling, W.A., Hoogland, H., Babcock, G.T. and Wever, R. (1988) *Biochemistry* 27, 5395–5400.
- [39] Egawa, T., Miki, H., Ogura, T., Makino, R., Ishimura, Y. and Kitagawa, T. (1992) *FEBS Lett.* 305, 206–208.
- [40] Sato-Watanabe, M., Mogi, T., Ogura, T., Kitagawa, T., Miyoshi, H., Iwamura, H. and Anraku, Y. (1994) submitted.
- [41] Varotsis, C., Zhang, Y., Appelman, E.H. and Babcock, G.T. (1993) *Proc. Natl. Acad. Sci. USA* 90, 237–241.