## Simultaneous reduction of iron–sulfur protein and cytochrome $b_{L}$ during ubiquinol oxidation in cytochrome $bc_{1}$ complex

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Edited by Helmut Beinert, University of Wisconsin, Madison, WI, and approved January 30, 2007 (received for review September 6, 2006)

The key step of the protonmotive Q-cycle mechanism of the cytochrome bc1 complex is the bifurcated oxidation of ubiquinol at the Qp site. It was postulated that the iron-sulfur protein (ISP) accepts the first electron from ubiquinol to generate ubisemiquinone anion to reduce b<sub>L</sub>. Because of the difficulty of following the reduction of ISP optically, direct evidence for the early involvement of ISP in ubiquinol oxidation is not available. Using the ultra-fast microfluidic mixer and the freeze-quenching device, coupled with EPR, we have been able to determine the presteady-state kinetics of ISP and cytochrome b<sub>L</sub> reduction by ubiquinol. The first-phase reduction of ISP starts as early as 100  $\mu$ s with a  $t_{1/2}$  of 250  $\mu$ s. A similar reduction kinetic is also observed for cytochrome b<sub>L</sub>, indicating a simultaneous reduction of both ISP and  $b_{\rm L}$ . These results are consistent with the fact that no ubisemiquinone was detected at the Qp site during oxidation of ubiquinol. Under the same conditions, by using stopped flow, the reduction rates of cytochromes  $b_{\rm H}$  and  $c_1$  were 403 s<sup>-1</sup> ( $t_{1/2}$  1.7 ms) and 164 s<sup>-1</sup> (t<sub>1/2</sub> 4.2 ms), respectively.

EPR | rapid freeze-quenching | stopped-flow | ubiquinol-cytochrome c reductase | electron transfer

he cytochrome  $bc_1$  complex (also known as ubiquinolcytochrome c reductase or complex III) is an essential segment of the electron transfer chain of mitochondria and many respiratory and photosynthetic bacteria (1). It catalyzes electron transfer from ubiquinol to cytochrome c with concomitant translocation of protons across the membrane to generate a proton gradient and membrane potential for ATP synthesis. The complex from bovine heart mitochondria is composed of 11 protein subunits: three core subunits, subunits III, IV, and V, which house *b*-type cytochromes, *c*-type cytochrome, and ironsulfur center, respectively, and eight supernumerary subunits that contain no redox prosthetic groups. Cytochrome  $bc_1$  complexes from other sources contain the three major subunits with one to seven supernumerary subunits (2). In either case, the complex contains four redox prosthetic groups: cytochrome  $b_{\rm L}$  $(b_{566})$  and  $b_{\rm H}$   $(b_{562})$ , cytochrome  $c_1$ , and a high potential ironsulfur cluster (ISC; 2Fe-2S Rieske center).

The 3D structure of the cytochrome  $bc_1$  complexes from beef (3, 4), chicken (5), yeast (6), and Rhodobacter capusulata (7) were recently determined. The complex is crystallized in an intertwined dimer form. The iron-sulfur proteins (ISPs) in the two  $bc_1$  monomers are intertwined with the head domain in one monomer interacting with cytochrome b and cytochrome  $c_1$  of the other monomer. The intertwined dimer observed in the crystalline complex also exists in solution, which was confirmed (8) in the R. sphaeroides  $bc_1$  complex through the formation of a four-subunit (two ISPs and two cytochrome bs) adduct by two intersubunit disulfide bonds between two engineered cysteine pairs: one pair links the ISP head domain to cytochrome b, and the other pair links the ISP tail domain to cytochrome b of another monomer. Two apparently noncommunicating cavities in the dimeric complex are presented: each connecting the Qp pocket of one monomer to the Qn pocket of the other. The distance between the Fe atoms of the two hemes  $b_{\rm L}$  is only 21 Å, which is approximately the same as the distance between heme  $b_{\rm L}$  and  $b_{\rm H}$  in one monomer (Fig. 1). The short distance between the two hemes  $b_{\rm L}$  and the presence of several aromatic amino acid residues at the interface of the two cytochrome *b* proteins have caused investigators (9–12) to speculate about the existence of electron transfer and equilibration between the two hemes  $b_{\rm L}$ . The involvement of an aromatic residue in such an electron transfer has recently been confirmed in the bacterial complex (12).

On the basis of functional data (13, 14) and structural information (3-7), the "protonmotive Q cycle" is the most favored mechanism for electron and proton transfer in the cytochrome  $bc_1$ complex (15, 16). The key step of the Q-cycle mechanism (Fig. 2) is the bifurcation of electrons from ubiquinol at the Qp site. It was postulated that the first electron of ubiquinol is transferred to the "high-potential chain," consisting of the ISP and cytochrome  $c_1$ . Then the second electron of ubiquinol is passed through the "low-potential chain" consisting of cytochromes  $b_{\rm L}$  and  $b_{\rm H}$  to reduce ubiquinone or ubisemiquinone bound at the Qn site. Although crystallographic data clearly indicates the presence of two separated quinone binding domains: one for ubiquinol oxidation (Qp) and the other for quinone reduction (Qn), only the binding of ubiquinone (17–19) and the presence of ubisemiquinone radical (20-22) at the Qn site were demonstrated. The binding of ubiquinol or ubiquinone at the Qp site has not been detected even though the binding of various Qp site inhibitors (18, 23, 24) is well known.

The mechanism of the bifurcated oxidation at the Qp site is still a matter of controversy. Evidences supporting "sequential oxidation" (25–28) and the "concerted oxidation" (29–33) of the cytochrome *b*-bound ubiquinol are both available. In the sequential oxidation, the first electron of the ubiquinol is transferred to the ISC, and the resulting ubisemiquinone is used to reduce cytochrome  $b_L$ . The absence of detectable ubisemiquinone radical at the Qp site (33) questions the validity of the sequential oxidation mechanism, even though several plausible explanations for the absence of Q-radical have been offered (27, 34). Even though the rate constant for the presteady-state reduction of the cytochrome *b* by ubiquinol is much larger than that of cytochrome  $c_1$ , many investigators (35–38) assume that the first electron goes to the ISC, the thermodynamically favorable acceptor. The requirement for ISP (39) during the reduction of cytochrome  $c_1$  by ubiquinol in the

Author contributions: J.Z., T.E., S.-R.Y., L.Y., and C.-A.Y. designed research; J.Z. and T.E. performed research; J.Z., T.E., S.-R.Y., L.Y., and C.-A.Y. analyzed data; and C.-A.Y. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS direct submission.

Abbreviations: ISP, iron–sulfur protein; ISC, iron–sulfur cluster; DM, dodecyl maltoside; OG, octyl glucoside.

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**Fig. 1.** The relative location of the essential redox groups of the bovine heart mitochondrial cytochrome  $bc_1$  complex (59). The  $b_L$ ,  $b_H$ , and  $c_1$  hemes are shown as ball-and-stick models, whereas the *2Fe2S* clusters are shown as ball models. The quinone oxidation pockets are near the intermembrane space (IMS) side of the membrane, and the quinol reduction pockets are near the matrix side of the membrane. Cytochrome *c* is shown as a shaded oval as labeled. Distances between redox centers are given on the left, and the redox potential for each center is given on the right. The high potential ET path is depicted by arrowed lines pointing downward. Circles in the upper and lower parts within the Qp pockets are distal and proximal quinone bindings, respectively.

cytochrome  $bc_1$  complex supports this assumption. The observed smaller rate constant of cytochrome  $c_1$  reduction, compared with the cytochrome *b* reduction rate constant, was attributed to the slow electron transfer between the ISC and heme  $c_1$ . In fact, there is no evidence, so far, showing that the ISC receives an electron from ubiquinol before other redox components, because of the difficulty in detecting the redox change of the ISC in a fast time scale.

Taking advantage of the unique EPR signatures of the ISC (40), cytochromes  $b_{\rm L}$ ,  $b_{\rm H}$ , and  $c_1$  (41–43), and ubisemiquinone radical (21, 22), we recently investigated the presteady-state kinetics of ISP and cytochrome  $b_{\rm L}$  reduction by ubiquinol at the Qp site by using an ultra-fast freeze-quenching technique (44) coupled with EPR spectroscopy. Our results are consistent with the concerted oxidation mechanism (29–32) of the ubiquinol at the Qp site, although we could not fully exclude the sequential mechanism because of our limited time resolution of  $\approx 50 \ \mu$ s. To further confirm this mechanism, measurements with a better time resolution are required.

## **Results and Discussion**

**EPR Characteristics of the Redox Components of Cytochrome**  $bc_1$ **Complex.** The EPR spectra of cytochromes b,  $c_1$ , and the ISC in the cytochrome  $bc_1$  complex have been well characterized. In the oxidized complex, cytochromes  $b_L$  ( $b_{566}$ ),  $b_H$  ( $b_{562}$ ), and  $c_1$  have peaks at g = 3.78, g = 3.45, and g = 3.35, respectively (41–43). The g = 3.78 peak of cytochrome  $b_L$  is sharp and asymmetric and difficult to power saturate even at 7 K. Peaks of cytochromes  $b_H$  and  $c_1$  overlap to show a peak at g = 3.4 (Fig. 3*A*). The roughly symmetric peak (g = 3.45) for cytochrome  $b_H$  can be detected by using ascorbate-reduced complex. The EPR spectrum (g = 3.35) of cytochrome  $c_1$  can be obtained by subtracting the ascorbate-reduced spectrum from the fully oxidized one. Because EPR



**Fig. 2.** The proton motive Q-cycle mechanism. (A) Sequential bifurcated oxidation of quinol at the Qp site. Ubiquinol bound in cytochrome *b* is first oxidized by ISP to generate an ubisemiquinone radical to reduce cytochrome  $b_{L}$ . (B) Concerted bifurcated oxidation of ubiquinol at the Qp site. Bifurcated oxidation of ubiquinol at the Qp site by ISP and cytochrome  $b_{L}$  takes place simultaneously with no generation of ubisemiquinone radical at the Qp site.

spectra of cytochromes  $b_{\rm H}$  and  $c_1$  have similar power saturation behavior, they cannot be resolved by power manipulation (41–43).

The EPR spectrum of the reduced ISC of the Rieske's protein (Fig. 3*B*) shows three peaks,  $g_z = 2.02$ ,  $g_y = 1.89$ , and  $g_x = 1.80$  (40). While the shape and g value of  $g_x$  is affected by the redox states of other components (42, 43) and the presence of ubiquinone (45), the intensity of the signal of  $g_y$  (g = 1.89) is directly proportional to the



**Fig. 3.** EPR spectra of oxidized cytochromes *b* and  $c_1$  (*A*) and the reduced Rieske ISP (*B*). The spectra were taken at 6 K, with the following instrument settings: microwave frequency, 9.45 GHz; modulation amplitude, 6.30 G; modulation frequency, 100 KHz; time constant, 0.655 s.

Table 1. Ubiquinol concentration-dependent reduction rate constants of cytochromes  $b_H$  and  $c_1$ 

$QH_2$ concentration, $\mu M$	Reduction rate constants, s <sup>-1</sup>	
	b <sub>H</sub>	с <sub>1</sub>
84	224	87
167	300	100
333	407	167
500	432	187

The cytochrome  $bc_1$  complex was diluted in 50 mM phosphate buffer, pH 7.4, containing 1 mM EDTA and 0.01% DM to a concentration of 100  $\mu$ M based on cytochrome  $c_1$ . Ubiquinol was diluted in 0.2% DM and 0.2% OG to various concentrations to give a final concentration of 1.7-, 3.34-, 6.66-, and 10-fold greater than that of cytochrome  $c_1$ . Reductions of cytochrome b and cytochrome  $c_1$  were monitored by the increase of absorption at 559 nm and 550 nm, respectively, by using an Applied Photophysics stopped-flow reaction analyzer SX.18MV.

degree of reduction of ISC. Thus, the rate of ISC reduction can be followed by the  $g_y = 1.89$  signal.

The EPR signature of ubisemiquinone radical is a single sharp peak at g = 2.004, which is easily power saturated at low temperature. Because ubisemiquinone at the Qn site is well characterized (20–24), it should not be difficult to differentiate it from any ubisemiquinone radical that is generated at the Qp site. Ubisemiquinone at the Qn site is antimycin A-sensitive and has a reduction kinetic similar to that of cytochrome  $b_{\rm H}$  (37). Ubisemiquinone at the Qp site, if any, would be expected to have a kinetic similar to that of ISC or cytochrome  $b_{\rm L}$ .

Presteady Reduction Rates of Cytochrome b<sub>H</sub> and c<sub>1</sub> by Ubiquinol Determined by Conventional Stopped-Flow Method. The electron transfer rates between the ubiquinol and heme  $b_{\rm H}$  or heme  $c_1$  in the cytochrome  $bc_1$  complex can be determined by mixing the fully oxidized cytochrome  $bc_1$  complex in 50 mM Tris·HCl (pH 8.0) buffer, containing 0.33 M sucrose and 0.01% dodecyl maltoside (DM), with an equal volume of various concentrations of ubiquinol at room temperature in a stopped-flow system (SX.18MV; Applied Photophysics, Leatherhead, UK). The final concentration of cytochrome  $bc_1$  complex is 50  $\mu$ M based on cytochrome  $c_1$ , and the concentrations of ubiquinol are 84, 167, 333, and 500  $\mu$ M. The substrate, ubiquinol, is prepared in the same buffer containing 0.2%DM and 0.2% octyl glucoside (OG) at double concentrations. Reductions of cytochromes  $b_{\rm H}$  and  $c_1$  are monitored by the increase of absorbance at 559 and 550 nm, respectively. Table 1 shows the observed rate constants for  $b_{\rm H}$  and  $c_1$  reduction obtained with various concentrations of ubiquinol.

Although the reduction rate constants of both cytochromes increase as the concentration of substrate increases, the degrees of increase level off as the concentration of ubiquinol approaches 500  $\mu$ M. An  $\approx$ 50% increase in substrate, from 333 to 500  $\mu$ M, results in an increase in the rate constants for cytochromes  $b_{\rm H}$  and  $c_1$ reduction by only 6% and 12%, respectively. Because of the high hydrophobicity of ubiquinol, it is difficult to reach a final concentration >500  $\mu$ M in an aqueous solution with a limited amount of detergent present. All of the subsequent rate measurements were carried out with a substrate concentration of 333  $\mu$ M. Under this condition the reduction rate constants of heme  $b_{\rm H}$  and heme  $c_1$  are 403 and 164 s<sup>-1</sup> with  $t_{1/2}$  of 1.7 and 4.2 ms, respectively.

When the reduction of cytochromes b and  $c_1$  in the stopped-flow experiment is followed with photodiode array scanning, the reduction of  $b_{\rm H}$ , instead of  $b_{\rm L}$ , can be clearly identified. The reaction tracings show the early reduction of heme  $b_{\rm H}$ , starting before 1 ms, followed by the reduction of heme  $c_1$  after 3 ms. The reduction of  $b_{\rm L}$  is completed within the dead time of stopped-flow apparatus. Presteady-State Reduction Rates of Cytochrome  $b_L$  and ISC by Ubiquinol Determined by Ultra-Fast Microfluidic Mixing and the Freeze-Quenching Method. The large reduction rate constant of cytochrome  $b_L$  renders the conventional stopped-flow experiment inadequate for presteady-state reduction kinetics. Also, the close absorption peaks of cytochrome  $b_L$  and  $b_H$  complicate spectroscopic rate analysis. Ultra-fast microfluidic mixing and the freezequenching method (44) coupled with EPR detection offers an excellent way to determine the reduction kinetics of cytochrome  $b_L$ and ISC. The ultra-fast microfluidic mixer has a dead time of 50  $\mu$ s as described (44), hence it is valuable for detecting large reduction rate constants. The well resolved EPR signals of  $b_L$  and  $b_H$  make it easy to follow  $b_L$  reduction without complication from  $b_H$ .

The cytochrome  $bc_1$  complex was diluted in the same storage buffer to a cytochrome  $c_1$  concentration of 100  $\mu$ M and mixed with an equal volume of 666  $\mu$ M QH<sub>2</sub> in 0.2% DM and 0.2% OG in the ultra-fast microfluidic mixer (44). The reaction mixture was freeze-quenched at liquid nitrogen temperature at various time points, from 66  $\mu$ s to 5 ms, packed into EPR tubes, and stored at liquid nitrogen temperature before EPR analysis.

The redox state of ISP and cytochromes b and  $c_1$  were determined by EPR (42, 45) in the same sample, using a Bruker (Billeria, MA) EMX spectrometer at a temperature  $\approx 6$  K. The instrument settings were as follows: microwave frequency, 9.45 GHz; time constant, 0.655 s; and modulation frequency, 100 KHz. The modulation amplitude was 19.57 and 6.30 G for cytochromes and ISP, respectively. A microwave power of 2.15 mW was used, except for the power saturation study. During the EPR experiments it was noticed that air trapped inside the sample caused a distortion in the base line, especially in the region of magnetic field (0.15 to 0.21 Tesla) where EPR signals of  $b/c_1$  appear. To achieve high signal sensitivity, air trapped in the sample was removed by evacuation. The freeze-quenched samples in EPR tubes were first dipped into a mixture of isopentane and liquid nitrogen. The temperature of the samples was maintained at approximately -160°C. The EPR tubes were then subjected to vacuum for 30 s before being subjected to EPR measurements. Fig. 4 shows the EPR spectra of the cytochromes b and  $c_1$  (Fig. 4A and B) and the reduced Rieske ISP (Fig. 4 C and D) at various time points after mixing with ubiquinol. The reduction of ISC was calculated from the intensity of peak to valley at g = 1.89 (g<sub>y</sub>). The reduction of  $b_{\rm L}$  was calculated from the integrated peak area at g = 3.78. Because the freeze-quenching reaction mixtures used for EPR measurements were in powder form and packed into EPR tubes in liquid nitrogen, the packing factor varied from sample to sample. To ascertain the concentration of the sample in the EPR tube at each time point, an internal spin label standard, proxyl, was added to the  $bc_1$  sample before the rapid mixing experiment. The amounts of reduction for each redox center at various time point were normalized by using the spin label signal at g = 1.98. The g = 1.98 peak of proxyl was chosen over the two other stronger signals because it is not overlapping with the  $g_z$  signal of ISP.

As shown in Fig. 4D, there is no detectable semiquinone radical in the samples with a reaction time  $<360 \ \mu s$ . Semiquinone that appears after 366  $\mu s$  is the radical at the O<sub>n</sub> site, because it is sensitive to antimycin treatment and its formation rate is comparable to that of cytochrome  $b_{\rm H}$ . Correlation between semiquinone formation and cytochrome  $b_{\rm H}$  reduction has been reported (21).

Fig. 5 shows the reduction of ISC and heme  $b_L$  as a function of time. Note that the zero time point is based on a control sample prepared under the same conditions in the absence of QH<sub>2</sub>. Reduction of ISC appears to be biphasic. The first phase starts at  $\approx 100 \ \mu$ s and is complete within 1 ms. The reduction of heme  $b_L$  proceeds with similar kinetics but reaches a maximum at  $\approx 400 \ \mu$ s. On the basis of these kinetic traces, the rate constants for electron transfer from ubiquinol to ISC and  $b_L$  at the Qp site is 2,770 s<sup>-1</sup>, which is larger than the rate constants obtained by various indirect methods (36, 46, 47).



**Fig. 4.** EPR spectra of cytochromes *b* and  $c_1$  and the reduced Rieske ISP at various time points after mixing with ubiquinol. The EPR spectra of cytochromes  $b_L$  and  $b_H/c_1$  (*A* and *B*) and ISC (*C* and *D*) in the cytochrome  $bc_1$  complex reacting with ubiquinol are shown at different time points as indicated. *B* and *C* are the EPR spectra of cytochrome  $b_L$  and ISC at  $g_y = 1.89$ , respectively. *D* shows the EPR spectra of ISC with g = 2.00 region. Samples were prepared as described in *Experimental Procedures*. The EPR spectra were taken at 6 K, with the following instrument settings: microwave frequency, 9.45 GHz; microwave power, 2.15 mW; and modulation frequency, 100 KHz; time constant, 0.655 s. The modulation amplitude used was 19.57 and 6.30 G for cytochromes and ISP, respectively. All curves are in the same order as the reaction times given.

One possibility for the biphasic reduction of ISP is the dimeric nature of the cytochrome  $bc_1$  complex. When both monomers are in the oxidized form, the rate of reduction of ISP would be higher than that in the partially reduced complex. The slower reduction of ISP in the second phase cannot be resulted from the oxidation of ISP by cytochrome  $c_1$  because little cytochrome  $c_1$  is reduced in a reaction time < 2 ms. The  $t_{1/2}$  for  $c_1$  reduction was determined by conventional stopped flow to be 4.7 ms. Probably cytochrome  $b_{\rm L}$  is also reduced biphasically. However, if the rate of the second (slower) phase of  $b_{\rm L}$  reduction is comparable to that of  $b_{\rm H}$  reduction then no increase in  $b_{\rm L}$  reduction would be observed, and the kinetic would appear monophasic. In other words, the observed rate of  $b_{\rm L}$ reduction is a "net" rate that includes both the reduction of  $b_{\rm L}$  by ubiquinol and reoxidation by  $b_{\rm H}$ . This proposal is consistent with the hypothesis of the half-of-the-sites activity of the dimeric  $bc_1$ complex (49).

Using a light flash to initiate the electron-transfer cycle in the bacterial cytochrome  $bc_1$  complex, a 1,650 s<sup>-1</sup> rate constant was reported for electron transfer from ubiquinol to ISP at the Qp site of the complex, in a chromatophores preparation from *R. capsulatus* (46). A rate constant of 1,200 s<sup>-1</sup> was reported when using a

binuclear ruthenium complex to rapidly photooxidize cytochrome  $c_1$  in the purified *R. sphaeroides*  $bc_1$  complex (47). Using the same technique of rapid photoreduce/photooxidize cytochrome  $c_1$  with ruthenium dimer (Ru<sub>2</sub>D), oxidant-induced reduction of cytochrome  $b_H$  was observed with a rate constant of 250 s<sup>-1</sup> in the presence of antimycin A with bovine  $b_1$  complex and 1,000 s<sup>-1</sup> with *R. sphaeroides*  $bc_1$  complex (48). The rate constant of the reduction of heme  $b_H$  by ubiquinol was found to be 270 s<sup>-1</sup> with a photoreleasable caged ubiquinol substrate with mitochondrial cytochrome  $bc_1$  (36).

The rate constants found in this study are significantly larger than those observed previously. The similar rate constants for electron transfer from ubiquinol to heme  $b_L$  and from ubiquinol to ISC indicate that either when the first electron of ubiquinol is transferred to the thermodynamically favored ISC the second electron is immediately transferred to heme  $b_L$ , or the two electrons of ubiquinol are transferred to ISC and  $b_L$  simultaneously from a transient ternary complex of ISP–QH<sub>2</sub>–cytochrome b. Either case is consistent with the Q-cycle mechanism with no ubisemiquinone radical formation at thenQp site. The simultaneous reduction of ISC and cytochrome  $b_L$  is also consistent with the high-resolution structural data of bovine  $bc_1$  complex.



**Fig. 5.** The degree of reduction of ISC and heme  $b_L$  against time. ISC ( $\bigcirc$ ) and heme  $b_L$  ( $\bullet$ ) reduction percentage by QH<sub>2</sub>, calculated from EPR spectra, at time points between 0 and 1 ms and 0 to 2.25 ms. The percentage reduction of  $b_L$  was calculated based on percentage decrease in the integrated area of EPR peak at 3.78 using the peak area of the fully oxidized complex as 100%. The percentage reduction of ISC was based on the signal intensity of  $g_y$  in the sample that was premixed with QH<sub>2</sub> as 100%.

No Ubisemiquinone Radical Is Detected at the Qp Site. If a ubisemiquinone radical is present during the oxidation of ubiquinol at the Qp site, it should be easily detected by EPR. However, there is no g = 2.00 signal in the EPR spectra of rapid mixed freeze-quenched samples, which shows reduction of ISP (see Fig. 4D) and cytochrome  $b_{\rm L}$  within 500  $\mu$ s of reaction time. To avoid the error of overlooking the signal of ubisemiquinone radical because of its power saturation behavior, the EPR measurements in g = 2.00 region were carried out at various powers. The absence of ubisemiquinone radical is consistent with the observation that the amounts of ISP reduction are equal to those of cytochrome  $b_{\rm L}$  reduction in the early events of ubiquinol oxidation.

Whether or not a ubisemiquinone radical is generated at the Qp site during the oxidation of ubiquinol has been a subject of debate. It was reported that under special experimental conditions a transient antimycin A-insensitive ubisemiquinone radical was observed (50) at the Qp site. These data were later refuted (33) because the radical was found to be insensitive to Qp site inhibitors such as myxothiazol, (E)- $\beta$ -methoxyacrylate-stilbene, or stigmatellin.

To support the idea that the first electron of ubiquinol goes to ISP and thus generates a highly reducing ubisemiquinone to reduce cytochrome  $b_{\rm L}$  during the oxidation of ubiquinol at the Qp site, several plausible explanations have been offered for the failure to detect a semiquinone radical at the Qp site. One suggestion is that the radical is antiferromagnetically coupled with the reduced ISC and the radical is only separated from ISP when the second electron is transferred to  $b_{\rm L}$  (34). Another suggestion is that the ubisemiquinone radical formed at the Qp site is too transient and the concentration is too low to be detected by EPR (27). The transient nature of the ubisemiquinone radical is easily understood if one pictures that the electron donor of ISP is the ubiquinol- $b_{\rm L}$  complex and not the ubiquinol alone. Once the first electron of ubiquinol in the complex is transferred to ISC, the second electron is immediately transferred to  $b_{\rm L}$ , thus no ubisemiquinone radical would accumulate. It is important to note that a ubisemiquinone radical signal, g = 2.00, starts to accumulate when the reaction time is >500 $\mu$ s. This radical is sensitive to antimycin treatment and has a formation kinetic similar to that of the reduction of cytochrome  $b_{\rm H}$ , hence it is the ubisemiquinone radical of the Qn site.

Structural Basis for Concerted Bifurcated Ubiquinol Oxidation at the **Qp Site.** Although the information concerning the precise ubiquinol binding site in the Qp pocket is still lacking, recent structural analysis of several Qp site inhibitor-loaded crystals suggests that when ubiquinol enters the Qp pocket it arrests the head domain of ISP to the *b*-position with the formation of a transient cytochrome b-QH<sub>2</sub>-ISP complex. This ternary complex is formed with the aid of hydrogen bond formation between Ser-151 and Lys-283 of cytochrome b and the backbone carbonyl atoms of ISP, and the formation of a hydrogen bond between the hydroxyl group at C-1 position of the substrate and the  $N_{\epsilon}$  of the His-161 of ISP, a situation similar to the binding of stigmatellin. The C-4 hydroxyl group of the substrate forms a hydrogen bond with either Tyr-131 or Glu-271, a situation similar to the binding of 5-n-undecyl-6-hydroxy-4,7dioxobenzothiazole (UHDBT) (51). If  $QH_2$  binds in the similar position as UHDBT, the distance between the ISC and the substrate would be shorter than the distance between heme  $b_{\rm L}$  and the substrate in the transient cytochrome b-QH2-ISP complex. Nonetheless, in the actual cytochrome b-QH2-ISP complex, the two distances could be very similar because the presence of a long alky side chain in ubiquinol may pull the ISP closer to  $b_{\rm L}$  than what is observed in the *b*-position. This proposal is supported by the tighter binding of hexahydrodibenzothiophene, than UHDBT, in the yeast complex. Moreover, a water molecule might be present between ubiquinol and Glu-271 or Tyr-131, facilitating its electron transfer to  $b_{\rm L}$ . Thus a comparable electron transfer rate from ubiquinol to ISC and to heme  $b_{\rm L}$  could be achieved. The involvement of a water molecule in electron transfer has been observed in other systems (52). The specific role of Glu-271 in substrate binding has been questioned (53). These factors may account for the observed comparable electron transfer rates from ubiquinol to ISC and heme  $b_{\rm L}$ .

## **Experimental Procedures**

**Materials.** DM and OG were purchased from Anatrace (Maumee, OH). 2,3-Dimethoxy-5-methyl-6-decyl-1,4-benzoquinone bromine  $(Q_0C_{10}Br)$  was synthesized in our laboratory at Oklahoma State University (54). Stigmatellin, myxothiazol, and antimycin A were purchased from Sigma (St. Louis, MO). Other chemicals were of the highest purity commercially available.

**Enzyme Preparations and Assays.** Bovine heart mitochondrial cytochrome  $bc_1$  complex was prepared as reported (55). The purified complex was dissolved in 50 mM Tris·HCl buffer, pH 8.0, containing 0.33 M sucrose and 0.01% DM to a cytochrome c1 concentration of 100  $\mu$ M. In some cases, 3-maleimido-proxyl (proxyl), a spin label, was added to a final concentration of 50  $\mu$ M (from a 50 mM stock solution in ethanol) as an internal standard. The mixtures were frozen at  $-80^{\circ}$ C until use. The purified  $bc_1$  complex contained 9 nmol of cytochrome b and 5.5 nmol of cytochromes b and  $c_1$  were determined spectrophotometrically by using millimolar extinction coefficients of  $\Delta E_{562-575 \text{ nm}} = 28.5 \text{ cm}^{-1} \text{ mM}^{-1}$  and  $\Delta E_{552-540 \text{ nm}} = 17.5 \text{ cm}^{-1} \text{ mM}^{-1}$  for cytochromes b and  $c_1$ , respectively.

For activity assay, the cytochrome  $bc_1$  complex was diluted with 50 mM phosphate buffer, pH 7.4, containing 1 mM EDTA and 0.01% DM to a protein concentration of 0.1 mg/ml. Diluted enzyme solution (5  $\mu$ l) was added to 990  $\mu$ l of an assay mixture containing 50 mM phosphate buffer, pH 7.4, containing 1 mM EDTA and 100  $\mu$ M cytochrome *c* in the presence or absence of inhibitor. Activity was determined by measuring the reduction of cytochrome *c* after the addition of 5  $\mu$ l of 5 mM Q<sub>0</sub>C<sub>10</sub>BrH<sub>2</sub> (QH<sub>2</sub>). A millimolar extinction coefficient of 18.5 cm<sup>-1</sup>·mM<sup>-1</sup> was used to calculate the activity. The *bc*<sub>1</sub> complex used in these experiments had a specific

activity of  $\approx 20 \ \mu \text{mol}$  of cytochrome c reduced per minute per nanomole of cytochrome b.

Stopped-Flow Experiments. For the determination of electron transfer rates between the ubiquinol and heme b or heme  $c_1$ , the cytochrome  $bc_1$  complex was mixed with equal volumes of various concentrations of ubiquinol at room temperature in an Applied Photophysics stopped-flow reaction analyzer SX.18MV. The cytochrome  $c_1$  concentration of  $bc_1$  complex was 100  $\mu$ M. Ubiquinol was diluted in 0.2% DM and 0.2% OG to various concentrations. Reductions of cytochromes b and  $c_1$  were monitored by the increase of absorption at 559 and 550 nm, respectively, and a photodiode array scan between 600 and 500 nm. A set of observed rate constants was measured as a function of the concentration of quinol, 1.7-, 3.34-, 6.66-, and 10-fold greater than that of the concentration of cytochrome  $c_1$ . When an inhibitor was used, the cytochrome  $bc_1$  complex was treated with 5-fold molar excess of the inhibitor over heme  $c_1$  for 15 min before the experiment.

**Freeze-Quenching Experiments.** The cytochrome  $bc_1$  complex, with concentration of cytochrome  $c_1$  at 100  $\mu$ M, was mixed at a 1:1 ratio with 667  $\mu$ M QH<sub>2</sub> in 0.2% DM and 0.2% OG inside the ultrafast microfluidic mixer (44) with a modification in the design of the mixing chamber (T.E., Jorge Durand, and S.-R.Y., unpublished work). The reaction mixture was freeze-quenched at various time points from 66  $\mu$ s to  $\approx$ 3 ms at liquid nitrogen temperature and packed into EPR tubes. All EPR tubes were stored at liquid nitrogen temperature before measurements.

**EPR Experiments.** The redox states of the ISP and cytochromes b and  $c_1$  were determined by EPR (43), using a Bruker EMX Spectrom-

- 1. Trumpower BL, Gennis RB (1994) Annu Rev Biochem 63:675-716.
- 2. Berry E, Guergova-Kuras MH, Huang L-S, Crofts AR (2000) Annu Rev Biochem 69.1007-1077
- 3. Xia D, Yu CA, Kim H, Xia JZ, Kachurin AM, Zhang L, Yu L, Deisenhofer J (1997) Science 277:60-66.
- 4. Iwata S, Lee JW, Okada K, Lee JK, Iwata M, Rasmussen B, Link TA, Ramaswamy S, Jap BK (1998) Science 281:64-71.
- 5. Zhang ZL, Huang L-S, Shulmeister VM, Chi Y-I, Kim KK, Huang L-W, Crofts AR, Berry EA, Kim S-H (1998) Nature 392:677-684.
- 6. Hunte C, Koepke J, Lange C, Robmanith T, Michel H (2000) Structure (London) 8:669-684.
- 7. Berry EA, Huang L-S, Saechao LK, Pon NG, Valkova-Valchanova M, Daldal F (2004) Photosynth Res 81:251-275.
- Xiao K, Chandrasekaran A, Yu L, Yu CA (2001) J Biol Chem 276:46125-46131.
- Soriano GM, Ponamarev MV, Carell CJ, Xia D, Smith J, Cramer WA (1999) J Bionenerg Biomembr 31:201-213.
- 10. Covian R, Gutierrez-Cirlos EB, Trumpower BL (2004) J Biol Chem 279:15040-15049.
- 11. Osyczka A, Moser CC, Daldal F, Dutton PL (2004) Nature 427:607-612.
- Gong X, Yu L, Xia D, Yu C-A (2005) J Biol Chem 280:9251-9257.
- 13. Erecinska M, Chance B, Wilson DF, Dutton PL (1972) Proc Natl Acad Sci USA 69:50-56.
- 14. Alexandre A, Lehninger AL (1979) J Biol Chem 254:11555-11560.
- 15. Mitchell P (1976) J Theor Biol 62:327–367.
- 16. Brandt U, Trumpower B (1994) Crit Rev Biochem Mol Biol 29:165-197.
- 17. Gao X, Wen X, Esser L, Quinn B, Yu L, Yu CA, Xia D (2003) Biochemistry 42:9067-9080
- 18. Kolling DRJ, Samoilova RI, Holland JT, Berry EA, Dikanov SA, Crofts AR (2003) J Biol Chem 278:39747-39754
- 19. Cooley JW, Ohnishi T, Daldal F (2005) Biochemistry 44:10520-10532. 20. Yu CA, Nakaoka S, Yu L, King TE (1978) Biochem Biophys Res Commun 82:1070-1078.
- 21. Yu CA, Nagaoka S, Yu L, King TE (1980) Arch Biochem Biophys 204:59-70.
- Ohnishi T, Trumpower BL (1980) J Biol Chem 255:3278-3284
- Esser L, Quinn B, Li F-F, Zhang M, Elberry M, Yu L, Yu CA, Xia D (2004) J Mol 23. Biol 341:281-302.
- 24. Kim H, Xia D, Yu CA, Xia J-Z, Kachurin AM, Zhang L, Yu L, Deisenhofer J (1998) Proc Natl Acad Sci USA 95:8026–8033.
  25. Link TA (1997) FEBS Lett 412:257–264.
- 26. Brandt U, Von Jagow G (1991) Eur J Biochem 196:163-170.
- Crofts AR (2004) Annu Rev Physiol 66:689-733. 27.
- 28. Hong SJ, Uggulava N, Guergova-Kuras M, Crofts AR (1999) J Biol Chem 274:33931-

eter at a temperature of  $\approx 6$  K. The instrument settings were as follows: microwave frequency, 9.45 GHz; microwave power, 2.15 mW; time constant, 0.655 s; and modulation frequency, 100 KHz. The modulation amplitudes used were 19.57 and 6.30 G for cytochrome and ISP, respectively. Previous experiments indicated that air trapped inside the sample causes a distorted base line, especially in the heme  $b/c_1$  region of the EPR spectrum. To achieve high signal sensitivity, the air remaining within the sample was eliminated by first dipping the EPR tubes containing the freezequenched samples into a mixture of isopentane and liquid nitrogen. This solution was at a temperature of approximately  $-160^{\circ}$ C. Then the EPR tubes were subjected to vacuum for 30 s before being put into the EPR spectrometer. The amount of reduction of ISP was calculated from the intensity of peak to valley around  $g = 1.89 (g_v)$ . Reduction of  $b_{\rm L}$  was calculated from the integrated peak area at g =3.78. As described in Enzyme Preparations and Assays, spin label proxyl was added to enzyme solution as an internal standard. The data collected at each reaction time were normalized by comparison to the spin label intensity at g = 1.98. For detection of ubisemiquinone radical, freeze-quenching experiments were carried out with cytochrome  $bc_1$  complex containing no proxyl internal standard.

Other Biochemical and Biophysical Techniques. Protein concentration was determined by the method of Lowry et al. (56). Cytochrome b (57) and cytochrome  $c_1$  (58) contents were determined according to reported methods.

We thank Dr. Roger Koeppe for critical review of this manuscript. This work was supported by National Institutes of Health Grants GM30721 (to C.-A.Y.) and HL065465 (to S.-R.Y.) and Agricultural Experiment Station Projects 1819 and 2372, Oklahoma State University.

- 29. Yu CA, Wen X, Xiao K, Xia D, Yu L (2002) Biochim Biophys Acta 1555:65-70.
- 30. Kim H, Xia D, Yu CH, Xia J-Z, Kachurin AM, Zhang L, Yu L, Deisenhofer J (1998) Proc Natl Acad Sci USA 95:8026-8033.
- 31. Trumpower BL (2002) Biochim Biophys Acta 1555:166-173.
- 32. Hunte C, Palsdottir H, Trumpower BL (2003) FEBS Lett 545:39-46.
- 33. Junemann S, Heathcote P, Rich PR (1998) J Biol Chem 273:21603-21607.
- 34. Link TA (1997) FEBS Lett 412:257-264.
- 35. De Vries S, Albracht SPJ, Berden JA, Slater EC (1982) J Biol Chem 256:11996-11998.
- 36. Hansen KC, Schulz BE, Wang G, Chan SI (2000) Biochim Biophys Acta 1456:121-
- 137. 37. King TE, Yu CA, Yu L, Chiang YL (1975) in Electron Transfer Chains and Oxidative
- Phosphorylation, eds Quayliariello E, Papa S, Palmieri F, Slater EC, Siliprandi N (Elsevier, New York), pp 105-118.
- 38. Tsai A-L, Olson JS, Palmer G (1983) J Biol Chem 258:2122-2125.
- 39. Trumpower BL, Edwards CA (1979) J Biol Chem 254:8697-8706.
- 40. Orme-Johnson NR, Hansen RE, Beinert H (1974) Biochem Biophys Res Commun 45:871-878.
- 41. Salerno JC (1984) J Biol Chem 259:2331-23366.
- 42. De Vries S, Albracht SPJ, Leeuwerk FJ (1979) Biochim Biophys Acta 546:316-333.
- 43. McCurley JP, Miki T, Yu L, Yu CA (1990) Biochim Biophys Acta 1020:176-186.
- 44. Lin Y, Gary J, Gerfen GJ, Rousseau DL, Yeh S-R (2003) Anal Chem 75:5381-5386.
- 45. Ding H, Robertson DE, Daldal F, Dutton PL (1992) Biochemistry 31:3144-3158.
- 46. Crofts AR, Wang Z (1989) Photosynth Res 22:69-87.
- 47. Sadoski RC, Engstrom G, Tian H, Zhang L, Yu L, Yu CA, Durham B, Millett F (2000) Biochemistry 39:4231-4236.
- 48. Engstrom G, Xiao K-H, Yu CA, Yu L, Durham W, Millett F (2002) J Biol Chem 277:31072-31078.
- 49. Covian R, Trumpower BL (2006) J Biol Chem 281:30925-30932.
- 50. De Vries S, Albracht SPJ, Berden JA, Slater EC (1982) J Biol Chem 256:11996-11998.
- 51. Palsdottir H, Lojero GC, Trumpower BL, Hunte C (2003) J Biol Chem 278:31303-31311.
- 52. Lin J, Balabin IA, Beratan DN (2005) Science 310:1311-1313.
- 53. Osyczka A, Zhang H, Mathe C, Rich P, Moser CC, Dutton PL (2006) Biochemistry 45:10492-10503.
- 54. Yu CA, Yu L (1982) Biochemistry 21:4096-4101.
- 55. Yu CA, Yu L (1980) Biochim Biophys Acta 591:409-420.
- 56. Lowry OH, Rosebrough NJ, Farr AL, Randall RJ (1951) J Biol Chem 193:265-
- 57. Berden JA, Slater EC (1970) Biochim Biophys Acta 216:237-249.
- 58. Yu CA, Yu L, King TE (1972) J Biol Chem 247:1012-1019.
- 59. Esser L, Gong X, Yang S-Q, Yu L, Yu CA, Xia D (2006) Proc Natl Aced Sci USA 35:13045-13050.