

Exploring the proton pump mechanism of cytochrome c oxidase in real time

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Cytochrome c oxidase catalyzes most of the biological oxygen consumption on Earth, a process responsible for energy supply in aerobic organisms. This remarkable membrane-bound enzyme also converts free energy from O₂ reduction to an electrochemical proton gradient by functioning as a redox-linked proton pump. Although the structures of several oxidases are known, the molecular mechanism of redox-linked proton translocation has remained elusive. Here, correlated internal electron and proton transfer reactions were tracked in real time by spectroscopic and electrometric techniques after laser-activated electron injection into the oxidized enzyme. The observed kinetics establish the long-sought reaction sequence of the proton pump mechanism and describe some of its thermodynamic properties. The 10- μ s electron transfer to heme *a* raises the pK_a of a "pump site," which is loaded by a proton from the inside of the membrane in 150 μ s. This loading increases the redox potentials of both hemes *a* and *a*₃, which allows electron equilibration between them at the same rate. Then, in 0.8 ms, another proton is transferred from the inside to the heme *a*₃/Cu_B center, and the electron is transferred to Cu_B. Finally, in 2.6 ms, the preloaded proton is released from the pump site to the opposite side of the membrane.

cytochrome oxidase | electron transfer | proton translocation

Electrochemical proton gradients across phospholipid membranes are generated in primary biological energy transduction, whether powered by sunlight (photosynthesis) or by oxidation of hydrogenated foodstuffs by O₂ (respiration). In both cases the proton gradient is then used to synthesize adenosine triphosphate (ATP), the universal energy currency in cellular processes. O₂ consumption is usually catalyzed by cytochrome c oxidase (CcO), the terminal member of the respiratory chain in mitochondria and many bacteria. Electrons from cytochrome *c* on the positively charged P-side of the membrane are transferred, one at a time, to the binuclear heme/copper (*a*₃/Cu_B) oxygen reduction site located $\approx 1/3$ into the membrane domain (Fig. 1). The electron transfer takes a specific route via the bimetallic Cu_A center at the membrane surface, and another heme group (heme *a*) next to the *a*₃/Cu_B site (for reviews, see refs. 1 and 2). Because reduction of O₂ to water requires four electrons, there are four such one-electron transfers in the catalytic cycle. Each of these is associated with uptake of a substrate proton into the *a*₃/Cu_B site from the negatively charged N-side of the membrane to form the equivalent of water, and with translocation (pumping) of another proton across the membrane (3). Proton uptake from the N-side takes place via two pathways (Fig. 1) that are differently engaged in subsequent parts of the catalytic cycle (1). Each of the four electron transfer steps in the catalytic cycle of CcO constitutes one cycle of the proton pump, which is likely to occur by essentially the same mechanism each time. Here, we report on the internal electron transfer and charge translocation kinetics of one such cycle, which is set forth by fast photoinjection of a single electron into the oxidized enzyme.

Results

Photoactivated electron delivery using ruthenium bispyridyl (RubiPy) is a very useful method (4) for studying the reactions

associated with transfer of a single electron through CcO (3, 5–7), which is coupled to one of the four proton-pumping steps of the catalytic cycle. The trajectory of the injected electron may be monitored by time-resolved optical spectroscopy, which reveals the redox states of the metal centers. Vectorial proton movements can be tracked in phospholipid vesicles inlaid with CcO by capacitatively coupled time-resolved electrometry (3, 6–9), which is a sensitive method of detecting charge movements within the dielectric of the enzyme structure as long as they are orientated perpendicular to the membrane. These charge movements are mainly due to proton transfers from the N- toward the P-side (Fig. 1), with only a small contribution from electron transfer between Cu_A and heme *a*. Further electron transfer from heme *a* to the *a*₃/Cu_B site yields no electrometric signal because it occurs parallel to the membrane (10). Consequently, monitoring both electron transfers and proton (charge) translocation in real time can yield valuable mechanistic and thermodynamic insight into the molecular machinery of proton-pumping by CcO.

Electron Transfer. Fig. 2*A* shows the electron transfer kinetics after a laser pulse that converts RubiPy into a strong reductant that injects an electron into CcO (4–7). The quantum efficiency is 10–20% in our conditions, which assures, in practice, that either one or no electron is injected. Because of the short time during which the reaction is monitored (a few milliseconds), we can also exclude electron transfer between CcO molecules, which occurs on a much longer time scale (6). Just before the laser pulse the enzyme was reduced and then reoxidized by O₂ (11, 12), which is essential, because otherwise the oxidized enzyme is in a "resting" state incapable of fast proton-coupled electron transfer (3, 13). Upon excitation, the Cu_A site is reduced first with a time constant (τ) of $<0.5 \mu$ s (the life-time of the excited state of RubiPy), followed by electron equilibration between Cu_A and heme *a* ($\tau \approx 10 \mu$ s), where $\approx 70\%$ of the electron is transferred to the latter (Figs. 2*A* and 3*A*). We emphasize that our methodology yields a full spectrum for each time point, and does not rely on the kinetics at a single wavelength (see *Materials and Methods*), which increases the accuracy of assigning the spectral changes to specific redox centers. The observed noise has a standard deviation of $\approx 2\%$ of the total redox change of heme *a*, and $\approx 5\%$ of Cu_A.

From the Cu_A/heme *a* equilibrium position after the 10- μ s phase

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Abbreviations: CcO, cytochrome c oxidase; DM, dodecyl L-D-maltoside; RubiPy, ruthenium bispyridyl.

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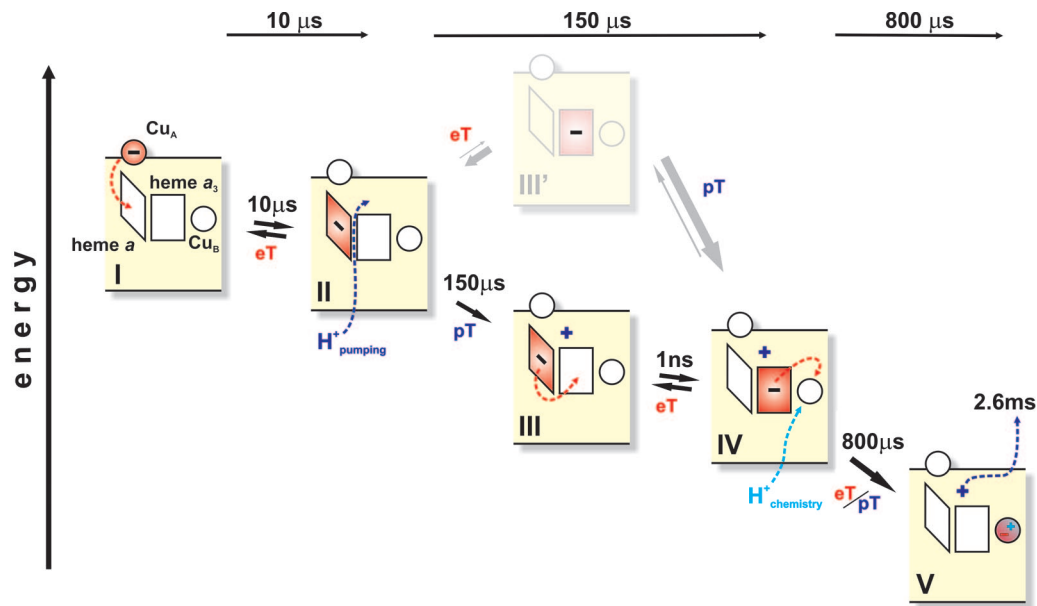


Fig. 4. Reaction scheme. The rhombus and square represent hemes *a* and *a*₃, respectively. The circle above heme *a* represents the Cu_A site, and the circle next to heme *a*₃ represents Cu_B. The minus sign denotes the position of the photoinjected electron. The dark and light blue plus signs denote the pumped and substrate protons, respectively. Dashed arrows indicate electron and proton transfers during the next reaction step. eT, electron transfer; pT, proton transfer.

tribute to most of the membrane potential formation, of which only a small fraction can be due to remaining electron transfer between Cu_A and heme *a*. Most of the electrogenicity must therefore be the result of two processes: uptake of the “substrate proton” from the N-side to the a₃/Cu_B site, and uptake of the pumped proton (Fig. 1). These two reactions are well separated in time, as found earlier (cf. refs. 6 and 7), but which is due to uptake of the substrate proton, and which to the pumped proton? This is a key question, not least because previous work has been contradictory on this point. For example, Ruitenber *et al.* (6) and Faxén *et al.* (29) have suggested that uptake of the substrate proton precedes uptake of the pumped proton, whereas others have presented evidence to the opposite (7, 30). Uptake of the substrate proton directly into the a₃/Cu_B site is expected to raise the *E*_m of that site much more than uptake of the pumped proton to a site located further away. If so, the assignment is straightforward: the 150-μs phase is due to uptake of the pumped proton from the aqueous N-side in a large fraction of the enzyme, and the 800-μs phase is mainly due to uptake of the substrate proton. This distinction leads to a plausible reaction scheme (Fig. 4).

Reaction Scheme. The injected electron initially equilibrates between Cu_A and heme *a*, which are roughly equipotential at this stage. The reduction of heme *a* raises the p*K*_a of a so far unidentified “pump site” above the heme groups (cf. refs. 7, 30, and 31), which takes up a proton in the 150-μs phase. The proton uptake raises the *E*_m of both hemes, and allows electron equilibration between them in the same time window. The rise in the *E*_m of heme a₃ is considerably larger than for heme *a* (see above), which suggests that the pump site is more closely linked to the former. Such a shared proton-binding site has long been predicted (16, 18) and finds clear functional significance here. As described for a photosynthetic reaction center (32), there are four possible states for a two-step mechanism describing such proton-coupled electron transfer, shown here by the upper and lower paths in Fig. 4. The rate may either be limited by proton or by electron transfer. Because of the very fast heme–heme electron tunneling time in CcO (24, 25), we may exclude those two reactions where electron transfer is rate-limiting. From the electron occupancies at the end of the 150-μs phase, we find that

the protonated state with heme *a* reduced (Fig. 4, state III) is highly populated. The data excludes any significant population of the state where heme *a* is reduced without proton uptake (Fig. 4, state II), because of the absence of a measurable population of reduced Cu_A. Combining the optical and electrometric data further strengthens this conclusion. The charge translocation can be described as the sum of electron and proton transfers propagating perpendicular to the membrane plane. The 30% electron transfer from Cu_A to heme *a* across one-third of the dielectric gives 0.1 charge equivalents (30% of 0.33). The location of the pump site is not known, but if we assume that it is at a relative dielectric distance *x* from the P-side of the membrane, proton transfer from the N-side to this site in the 150-μs phase adds 1 – *x* charge equivalents. Because we found the amplitude of this phase (proton + electron transfer) to be ≈0.84 charge equivalents, we may estimate *x* to be ≈0.25.

Our results favor a mechanism in which transfer of the pumped proton occurs before electron transfer to the binuclear center (contrast refs. 7 and 33), even though these two events have the same proton-limited kinetics. This finding is consistent with the conclusion by Brändén *et al.* (34) that protonation of the “pump site” controls this electron transfer, which is of considerable mechanistic interest because much previous work has stressed the importance of heme *a* in the proton pump mechanism (15, 18, 31, 33, 35–37). In the third phase, the substrate proton is transferred from the aqueous N-side, most likely to an OH[–] ligand of Cu_B (38), which raises the *E*_m of Cu_B to a value much higher than that of all other centers. Hence, this step is essentially irreversible (Fig. 4) and provides the main driving force for the entire pump mechanism. Finally, the slowest step (2.6 ms), seen only electrometrically, is proposed to be due to release of the proton from the “pump site” to the P-side by electrostatic repulsion from uptake of the substrate proton (cf. refs. 31 and 36). The observed electrometric amplitude (one charge across ≈32% of the dielectric barrier) agrees reasonably well with the above estimate of the position of the pump site relative to the membrane, and the slow rate is consistent with the suggestion by Salomonsson *et al.* (39) that release of the pumped proton is the rate-limiting step.

Although our work goes some way toward a molecular un-

derstanding of the proton pump mechanism of CcO, some important details remain unsolved, e.g., the identity of the proton-accepting pump site above the hemes. The basis for transferring the first proton to this site, rather than to be consumed at the a_3/Cu_B site, is also not fully understood although our data suggest a key role of heme *a* reduction. The observed thermodynamic effect of heme *a* reduction, i.e., the increase of the pK_a of the pump site, is hardly sufficient to explain the destiny of the first proton because of the proximity of the a_3/Cu_B site, whose pK_a should also increase. The notion that reduction of heme *a* may also kinetically favor proton transfer to the pump site (36) may therefore still be valid.

Materials and Methods

Enzyme Preparation and Reconstitution into Phospholipid Vesicles. CcO from *P. denitrificans* was isolated from bacterial membranes and purified as described (40), with the exception that the second Q-Sepharose column was replaced by a Ni^{2+} -NTA affinity chromatography column (see ref. 41 for details). In addition, the enzyme was washed and concentrated by using pressure dialysis (XM-50 membrane; Millipore, Bedford, MA) with 2 mM Tris-HCl (pH 8)/0.05% (wt/vol) dodecyl L-D-maltoside (DM)/20 mM aniline. The enzyme was reconstituted into vesicles by the Bio-Beads method (SM-2 adsorbent; Bio-Rad, Hercules, CA) as described (42), except that the concentration of CcO during reconstitution was increased to 6 μM .

Samples of detergent-dispersed *Paracoccus* membranes were prepared by solubilization of salt-washed membranes with 1% DM (Anatrace, Maumee, OH) in 20 mM Tris-HCl buffer (pH 7.8)/0.2 mM phenyl-methyl-sulfonyl-fluoride (PMSF), followed by ultracentrifugation (165,000 $\times g$, 30 min), after which the supernatant was washed and concentrated by using pressure dialysis with 2 mM Tris-HCl (pH 8)/0.05% DM/20 mM aniline.

Time-Resolved Measurement of Electric Potential Generation. The development of electric potential across the vesicle membrane was monitored by an electrometric technique (43), as adapted for time-resolved experiments with CcO (44). Details of the sample preparation and the methodology can be found in refs. 8 and 42.

Time-Resolved Spectrophotometric Measurements. Time-resolved multiwavelength absorption changes were followed by using a home-constructed CCD-based instrument. A pulsed 150-W xe-

non arc lamp (Applied Photophysics, Surrey, U.K.) was used as the probe light source. Light from the lamp was passed through a glass filter (OG-550) and fibers to a three-syringes stopped-flow module (SFM-300; Bio-Logic, Grenoble, France) equipped with a fluorescence cuvette (TC-100/10F, optical path 10 mm) with the sample. The light was further directed to a Triax-180 compact imaging spectrograph (HORIBA Jobin Yvon, Edison, NJ), which delivers spectral imaging over a fast kinetic CCD matrix (DV420-UV-FK; Andor Technology, Belfast City, Ireland). The setup was operated by software written by N.B. and allows recording absorption change surfaces with a time resolution of 1–16 μs between the spectra. The reaction was initiated by laser flash-induced electron injection into the enzyme from RubiPy (Tris[2,2'-bipyridyl] ruthenium[II] chloride) (BrilliantB; Quantel, Les Ulis, France; frequency-doubled YAG, 532 nm, pulse energy = 120 mJ).

Electron Injection. To obtain the pulsed oxidized state for absorption measurements, a solution of 80 μM CcO in 2 mM Tris (pH 8)/0.05% DM/20 mM aniline/15 μM *N,N,N',N'*-tetramethyl-*p*-phenylenediamine (TMPD) buffer was first made anaerobic on a vacuum line and then fully reduced by 1 mM potassium ascorbate. Then, anaerobic fully reduced CcO was mixed in the stopped-flow module with oxygen-saturated buffer (2 mM Tris (pH 8)/0.05% DM/20 mM aniline/400 μM RubiPy) resulting in oxidation CcO and formation of the pulsed state. Immediately after the mixing ($\Delta t = 5$ ms), a laser flash initiated the electron injection.

In electrometry, the pulsed state was generated by oxidation of the fully reduced CO-bound CcO in vesicles. Fully reduced enzyme was produced by anaerobic reduction of CcO (4 mM Tris, pH 8/3.5 mg/ml glucose oxidase/50 $\mu\text{g}/\text{ml}$ catalase/50 mM glucose/20 mM aniline/200 μM RubiPy/1 μM ruthenium hexa-amine) in a 1% CO atmosphere. The injection of oxygen-saturated buffer with aniline and RubiPy was performed through a needle directed toward the measuring membrane. The oxygen injection was followed by a series of flashes with 100-ms intervals. The first flash in this series photolyses CO off the enzyme and allows it to react with oxygen. The following flash induces photoinjection of an electron from RubiPy into the enzyme just oxidized (3).

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