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Specific methionine oxidation of cytochrome c in complexes with zwitterionic lipids by hydrogen peroxide: potential implications for apoptosis†

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Cytochrome c (Cyt-c) has been previously shown to participate in cardiolipin (CL) oxidation and, therefore, in mitochondrial membrane permeabilization during the early events of apoptosis. The gain in this function has been ascribed to specific CL/Cyt-c interactions. Here we report that the cationic protein Cyt-c is also able to interact electrostatically with the main lipid components of the mitochondrial membranes, the zwitterionic lipids phosphatidylcholine (PC) and phosphatidylethanolamine (PE), through the mediation of phosphate anions that bind specifically to amino groups in the surfaces of protein and model membranes. In these complexes, Cyt-c reacts efficiently with H₂O₂ at submillimolar levels, which oxidizes the sulfur atom of the axial ligand Met80. The modified protein is stable and presents significantly enhanced peroxidatic activity. Based on these results, we postulate that the rise of H₂O₂ concentrations to the submillimolar levels registered during initiation of the apoptotic program may represent one signaling event that triggers the gain in peroxidatic function of the Cyt-c molecules bound to the abundant PE and PC membrane components. As the activated protein is a chemically stable species, it can potentially bind and oxidize important targets, such as CL.

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1. Introduction

Cytochrome c (Cyt-c) is a highly conserved monohemic protein that serves as an electron shuttle between the respiratory complexes III and IV in the inner mitochondrial membrane (IMM), where it is also implicated in the generation, oxidation and trapping of reactive oxygen species (ROs).^{1,2} The release of Cyt-c from the mitochondrial intermembrane space to the cytosol, on the other hand, plays a pivotal role in apoptosis, thereby highlighting the importance of this small multifunctional protein in sustaining and terminating cellular life.³ Once in the cytosol, Cyt-c binds Apaf-1, ATP and pro-caspase-9 to form a multimeric complex called apoptosome that triggers a cascade of events leading to cellular suicide.^{2,4-7} The Cyt-c mediated caspase activation apoptotic pathway is essential for a large variety of biological events, including brain development,

immune system homeostasis, and stress-induced and genotoxic-induced cell death and, therefore, is involved in protecting living organisms from diseases such as cancer. It may also amplify other apoptotic pathways.^{2,4-7}

Several mechanisms have been proposed to explain the permeabilization of the outer mitochondrial membrane (OMM) that precedes Cyt-c release and are extensively reviewed elsewhere.^{2,4-9} In this context, different groups have paid particular attention to the interactions between cationic Cyt-c and the anionic phospholipid cardiolipin (CL),^{2,10-24} which represents ca. 10–20% of the total lipid content of the IMM and a much smaller proportion of the OMM.²⁵⁻²⁸ In the IMM, CL plays a crucial role in stabilizing the respiratory complexes and in mediating electron transport by Cyt-c. Indeed, it has been estimated that about 15% of the available Cyt-c is bound to CL.²⁹⁻³² A number of biophysical studies on model systems suggest that, in addition to nonspecific electrostatic interactions, CL forms a specific association with Cyt-c that involves the insertion of one or two hydrocarbon chains into the hydrophobic channels of the protein.^{11,14-20,24} The interplay of electrostatic and hydrophobic interactions triggers conformational changes that include the detachment of the iron axial ligand Met80 and augmented accessibility of potential substrates to the metal center. The result is a gain in peroxidase activity of Cyt-c that has been implicated in the selective oxidation of CL observed during the early events of apoptosis.^{10,11,23} Good arguments in favor of this proposal are the findings that (a) during apoptosis,

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the amount of CL in the OMM increases significantly and (b) the oxidation of CL appears to be a requisite for the release of proapoptotic factors, such as the release of Cyt-c to the cytosol across the OMM.^{10,11} Yet, the mechanistic details and, more specifically, the regulation of these processes remain elusive.

In addition to CL, mitochondrial membranes are rich in zwitterionic phospholipids such as phosphatidylcholine (PC) and phosphatidylethanolamine (PE), which comprise more than 70% of the total lipid content and have a quaternary ammonium and a primary amine bound to the phosphate group, respectively.^{25–27} As previously shown, Cyt-c is not expected to present significant electrostatic affinity for such lipids.^{14,33,34} Very recent studies, however, have demonstrated high affinity binding of Cyt-c to model systems containing amino and ammonium functional groups, that is specifically mediated by inorganic and organic phosphate anions at millimolar levels.^{35,36} In the present work, we have investigated the chemical reactivity of Cyt-c associated to similar model systems and in complexes with PE and PC liposomes. We found that the phosphate mediated binding of Cyt-c to these model lipids results in efficient oxidation of the axial ligand Met80 in the presence of H₂O₂ at submillimolar levels, thus yielding a stable modified protein with high peroxidatic activity. Interestingly, the early stages of the apoptotic program are characterized by a rise in H₂O₂ intracellular levels from submicromolar to submillimolar.^{37,38} Therefore, we propose that such a mechanism might be involved in the induction of peroxidatic activity of Cyt-c that precedes the liberation of proapoptotic molecules from the mitochondria.

2. Experimental

Chemicals

6-mercapto-1-hexanol, 6-mercaptohexanoic acid, 11-mercapto-1-undecanol, 11-mercaptoundecanoic acid, 11-mercaptoundecylphosphoric acid, (11-mercaptoundecyl)-*N,N,N*-trimethylammonium bromide and horse heart cytochrome c (Cyt-c) were purchased from Sigma-Aldrich. 11-amino-1-undecanethiol hydrochloride was purchased from Dojindo. 1,2-dioleoyl-*sn*-glycero-3-phosphocholine (PC) and 1,2-distearoyl-*sn*-glycero-3-phosphoethanolamine (PE) were purchased from Avanti Polar Lipids. Hydrogen peroxide was purchased from Mallinckrodt Baker, Inc. Amplex Red Ultra was purchased from Invitrogen. The M80A mutant of Cyt-c was prepared according to published procedures.³⁹

All chemicals were of the highest available purity and were used without further purification. The water used in all experiments was purified by a Millipore system and its resistance was 18.2 MΩ.

Electrochemistry

Cyclic voltammetry (CV) experiments were performed using a Gamry REF600 potentiostat. The jacketed three-electrode electrochemical cell was placed inside a Faraday cage (Gamry Vista Shield) and was equipped with a polycrystalline Au bead working electrode, a Pt wire counter electrode and a Ag/AgCl (3

M KCl) reference electrode to which all potentials in this work are referenced. Unless stated otherwise, the electrolyte solution was a 10 mM phosphate buffer, pH = 7.0, which was thoroughly deoxygenated by Ar bubbling. All experiments were carried out at room temperature (22–25 °C).

Raman spectroscopy

Resonance Raman (RR) and surface-enhanced resonance Raman (SERR) spectra were acquired with a confocal Raman microscope (Dilor XY) equipped with a liquid-nitrogen cooled CCD detector. The excitation source was the 413 nm line of a krypton ion laser (Spectra-Physics BeamLok 2060). For SERR determinations, the laser beam was focused onto the surface of a home-made rotating Ag electrode by means of a long-working-distance objective (20×, numerical aperture of 0.35). The SERR spectroelectrochemical cell has been described in detail elsewhere.⁴⁰ RR measurements were performed using a cylindrical quartz cell mounted on a home-made rotating device to avoid laser-induced sample degradation. Typically, experiments were performed with laser powers of 3.5 mW measured at the samples.

Electrode treatment and self-assembled monolayer preparation

The Au electrodes used for the CV experiments were first oxidized in 10% HClO₄ by applying a potential of 2.5 V for 2 minutes, sonicated in 10% HCl for 15 minutes and rinsed with water. The electrodes were then cleaned with a 3 : 1 v/v H₂O₂ : H₂SO₄ mixture at 120 °C. Finally, they were cycled between –0.2 and 1.6 V in 10% HClO₄ and thoroughly washed with water and ethanol.

The area of these electrodes was determined by CV from the integration of the reduction peak of the surface oxide. For these determinations, cyclic voltammograms (CVs) were recorded at 0.1 V s^{–1} between –0.25 and 1.6 V in 0.5 M H₂SO₄, and the conversion factor was 0.44 mC cm^{–2} for an oxide monolayer. The average surface area of the electrodes used in this work was calculated to be 0.18 ± 0.08 cm² with a roughness factor of 3.4 ± 1.1.

The Ag ring electrodes employed for the SERR measurements were mechanically polished and then treated with repetitive electrochemical oxidation/reduction cycles in 0.1 M KCl to create a SER-active nanostructured surface.⁴⁰

After the treatment described above, the Au and Ag electrodes were immersed in 2 mM ethanolic solutions of the desired thiols for self-assembly. Unless stated otherwise, the incubation solutions were 2 mM 1 : 1 mixtures of 11-mercapto-1-undecanol, which is added as a diluting component for the self-assembled monolayer (SAM), and a second ω-functionalized undecanethiol with negatively or positively charged ω-groups: carboxylate, phosphate, amino and trimethylammonium. The obtained SAMs are designated as COO-SAM, PO₄-SAM, NH₂-SAM and TA-SAM, respectively.

In the specific case of NH₂-SAM, the incubation solvent was a 4 : 1 v/v mixture of C₂H₅OH–HClO₄, 10^{–4} M, pH = 4, to prevent adsorption *via* the amino group. In all cases, the incubations



were carried out overnight and the SAM-modified electrodes were kept in darkness until use.⁴¹ Prior to their use, the electrodes were rinsed thoroughly with ethanol and water and subjected to electrochemical cycling in the phosphate buffer as a conditioning treatment.³⁶

Preparation of the lipid vesicles

Large unilamellar vesicles were prepared by the extrusion of equimolar PC/PE mixtures. Appropriate amounts of the lipid stock solutions were mixed in chloroform, evaporated to dryness under a gentle nitrogen stream and then left under reduced pressure for 1.5 h to remove any residual solvent. The dry lipids were subsequently hydrated with 20 mM HEPES, pH = 7, at room temperature to yield a lipid concentration of 0.5 mM. Afterwards, the samples were passed 20 times through polycarbonate filters with a pore size of 100 nm (Nuclepore, Whatman), yielding liposomes of the desired composition. The vesicle size, distribution and stability were checked by dynamic light scattering (SLS 90 Plus/BI-MAS equipped with He-Ne laser operating at 632.8 nm and 15 mW).

Synthesis and purification of the SO-Cyt protein variant

Chloramine-T was employed as a selective reagent for oxidizing the Met80 residue of Cyt-c, according to previous reports.^{42,43} The product obtained was designated as *ex situ* generated SO-Cyt. For this purpose, 100 μ L of 1 mM Cyt-c were mixed with 100 μ L of 5 mM chloramine-T and the pH was adjusted to 8.4. After 3 hours at room temperature, the mixture was washed 6 times with 10 mL of 10 mM phosphate buffer, pH = 7, using centrifugal filters (Amicon 10 kDa filters). The progress of the reaction was monitored by the disappearance of the 695 nm band from the UV-visible spectrum of Cyt-c (Thermo Scientific Evolution Array spectrophotometer).

The reaction mixture was passed through a cation exchange sulfopropyl-TSK preparative column (21.5 mm \times 15.0 cm; Tosoh Biosep) at a flow rate of 3 mL min⁻¹. The column was equilibrated with a 5 mM ammonium acetate buffer (pH = 9.0), kept for 5 min in this buffer and then eluted using linear gradients of the same salt from 5 mM to 150 mM (5 to 30 min), from 150 mM to 400 mM (30 to 75 min) and, finally, 500 mM ammonium acetate from 76 to 90 min (see ESI†).

The desired product was characterized by mass spectrometry (MS) employing a 4800 MALDI TOF/TOF instrument (Applied Biosystems) in a positive ion reflector mode. The mass spectra were externally calibrated using a mixture of peptide standards (Applied Biosystems). MS/MS spectra were recorded to confirm the site of oxidation in the analyzed peptides (see ESI†). The mass spectra were analyzed using DATA explorer and GPMW4 software.

Peroxidatic activity determination

The assessment of the peroxidatic activity was performed using an Amplex UltraRed Reagent (Invitrogen). The fluorescence of the oxidation product, resorufin, was monitored at $\lambda_{em} = 585$ nm with $\lambda_{ex} = 570$ nm. Briefly, 0.5 μ M Cyt-c was incubated in HEPES (10 mM plus 100 μ M DTPA), pH = 7. Then, 50 μ M

Amplex Red and 25 μ M H₂O₂ were added to HEPES (100 mM plus 100 μ M DTPA), pH = 7, and the fluorescence was registered using a FLUOstar Optima plate fluorimeter (BMG Labtech) or a Varioskan Flash Multimode Reader (Thermo). The reaction rate was determined by a linear fit of the fluorescence intensity profile.

3. Results and discussion

Reactivity of Cyt-c in biomimetic complexes

Hydrogen peroxide is a well-documented bleaching agent of Cyt-c that produces oxidative damage of the porphyrin ring.⁴⁴ Recent studies, however, suggest that the formation of electrostatic complexes with negatively charged surfaces has a protective effect on Cyt-c, thus preventing its degradation in the presence of H₂O₂ up to concentrations of *ca.* 20 mM.⁴⁵ In this context, we investigated the reactivity of Cyt-c towards H₂O₂ in electrostatic complexes with different model systems that mimic some essential features of the natural protein/membrane interactions. Specifically, we employed liposomes of different compositions and metal electrodes coated with SAMs of ω -functionalized alkanethiols. In the last case, the ω functional groups explored were carboxylate, phosphate, amino and trimethylammonium (COO-SAM, PO₄-SAM, NH₂-SAM and TA-SAM, respectively). The adsorption of Cyt-c on COO- and PO₄-SAMs involves direct electrostatic interactions through the ring of positively charged lysine residues that surround the partially exposed heme edge of the protein.⁴⁶ Roughly the same surface residues are implicated in the adsorption of Cyt-c onto NH₂- and TA-SAMs, although in these cases the interactions are mediated by phosphate ions from the buffer solution.^{35,36} In all cases, the adsorbed protein exhibits quasi-reversible cyclic voltammetric (CV) responses (Fig. S1†) typical of one-electron redox couples. The formal reduction potentials obtained for Cyt-c adsorption on COO-, PO₄- and NH₂-SAMs are very close to the value for the native protein in solution while, in excellent agreement with previous observations, the value for adsorption on TA-SAMs is somewhat more positive.^{35,47} Therefore, the detected electrochemical signals can be ascribed to adsorbed Cyt-c which largely preserves its native structure, particularly at the level of the redox site. It is important to point out that the absence of other electrochemical signals does not rule out the presence of alternative Cyt-c conformations in the adsorbed state that might be kinetically impaired. Indeed, the larger peak separations observed for PO₄-SAM compared to the other SAMs is indicative of a much slower interfacial electron transfer reaction of Cyt-c in this case, which is compatible with a less favorable and more rigid orientation of the protein in such complexes.⁴⁸

Upon addition of H₂O₂ to Cyt-c adsorbed on NH₂-SAMs, we observe the appearance of a second monoelectronic and quasi-reversible voltammetric signal with an apparent reduction potential of -119 mV (Fig. 1). The anodic and cathodic peak currents of the new redox active species rise at the expense of the corresponding signals of native Cyt-c, with a complete transformation at H₂O₂ concentrations slightly above 1 mM. Interestingly, the new redox couple presents enhanced cathodic currents in the presence of H₂O₂, thus indicating the



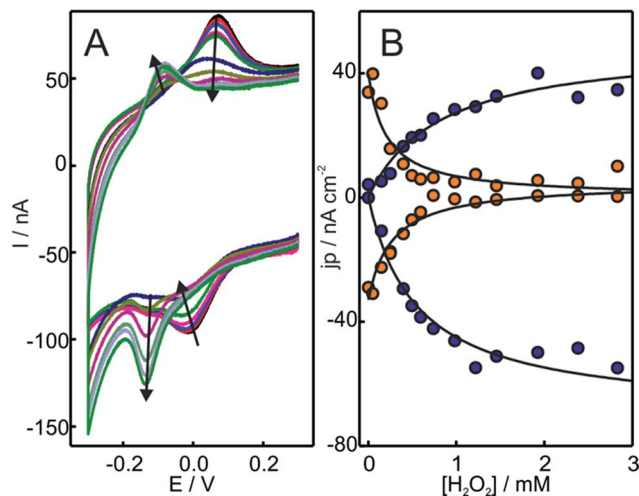


Fig. 1 (A) Voltammetric response of Cyt-c adsorbed on electrodes coated with NH₂-SAMs. The measurements were performed at 0.05 V s⁻¹ in 10 mM phosphate buffer, pH = 7.0, in the presence of increasing concentrations of H₂O₂ from 0 to 500 μM. (B) Background corrected peak current densities for the native Cyt-c redox couple (orange circles) and the newly generated couple (blue circles) as a function of H₂O₂ concentration. The solid lines are only intended as guides.

electrocatalytic reduction of H₂O₂ by the modified protein (pseudoperoxidase activity).

The downshift of the reduction potential of Cyt-c after treatment with H₂O₂ may in principle indicate a more polar heme environment as a consequence of increased solvent accessibility.⁴⁹ However, the addition of several complexing agents such as imidazole, azide, DTPA and methionine has no effect on the electrochemical response (Fig. S2[†]), suggesting that the solvent exposure of the heme crevice is not significantly altered.

The replacement of the electrolyte by fresh phosphate buffer once the H₂O₂-induced transformation is complete results in stable voltammograms that retain the electrochemical features of the newly generated redox couple (Fig. S3[†]). These results suggest a permanent chemical modification of the adsorbed Cyt-c into a stable species that presents pseudoperoxidase activity and, potentially, peroxidase-like characteristics. Remarkably, this behavior is only observed when Cyt-c is adsorbed on the NH₂- and TA-SAMs in the presence of the phosphate buffer (Fig. 1A and S5A[†]). In contrast, when the protein is directly adsorbed on negatively charged COO- and PO₄-SAMs, the addition of H₂O₂ produces a slight decrease of the voltammetric peaks of native Cyt-c, but this partial bleaching is not accompanied by the appearance of new CV signals (Fig. S6[†]). Moreover, pretreatment of the Cyt-c solutions with different concentrations of H₂O₂ (between 1 and 10 mM) prior to incubation with the electrodes coated with NH₂-SAMs leads exclusively to weak voltammetric signals at the potentials expected for the native Cyt-c, but there is no indication of other redox active species (Fig. 2B).

Based on these results, we investigated the effect of unilamellar vesicles of PC and mixtures of PC/PE. Typically, up to

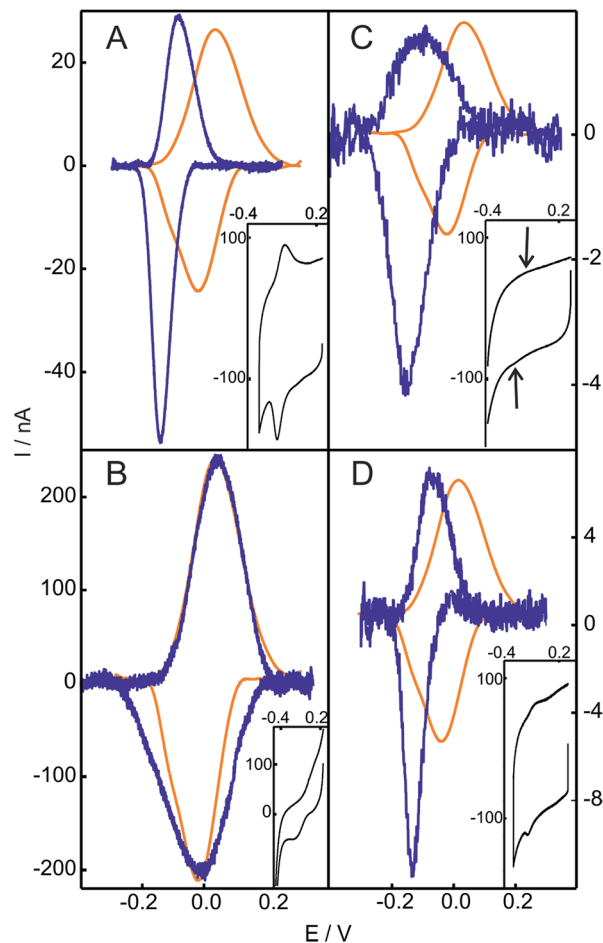


Fig. 2 Baseline subtracted cyclic voltammograms of Cyt-c before (orange) and after (blue) chemical treatment. (A) Cyt-c adsorbed on an NH₂-SAM electrode treated with 2 mM H₂O₂. (B) Cyt-c pretreated in solution with 2 mM H₂O₂ and subsequently adsorbed on an NH₂-SAM electrode. (C) Cyt-c pretreated in solution with 2 mM H₂O₂ in the presence of PE/PC liposomes, and subsequently adsorbed on an NH₂-SAM electrode. (D) *Ex situ* generated and purified SO-Cyt adsorbed on an NH₂-SAM electrode. All measurements were performed at 0.1 V s⁻¹ in 10 mM phosphate buffer, pH = 7.0. The insets are the voltammetric signals before background correction.

1 mM concentrations of H₂O₂ were added to solutions containing 200 μM Cyt-c and 0.5 mM PC (or PC/PE). After this treatment, the Cyt-c solutions were thoroughly washed using centrifugal filters and the supernatant was incubated with electrodes coated with NH₂-SAMs in the phosphate buffer. Similar to the *in situ* treatment of Cyt-c adsorbed on NH₂-SAMs, the modified electrodes exhibit a CV signal with $E_{1/2} = -119$ mV that also presents enhanced cathodic currents in the presence of H₂O₂ (Fig. 2C and S7[†]). Therefore, the H₂O₂-induced chemical modification that leads to a gain in pseudoperoxidatic (and possibly peroxidatic) activity appears to be a general feature of Cyt-c molecules adsorbed on positively charged and zwitterionic membrane models *via* mediating phosphate anions.



Characterization of the catalytically competent species

The structure of Cyt-c in the different biomimetic complexes was investigated by RR and SERR, as these spectroscopic techniques are able to monitor the redox state, spin and coordination pattern of the heme iron, as well as heme-protein interactions.^{40,48,50} In agreement with previous reports, the SERR spectra of Cyt-c adsorbed on COOH-SAMs are identical to the corresponding RR spectra of the native protein in solution (Fig. S8†).⁵⁰ Moreover, in agreement with the electrochemical results, the spectra remain unchanged upon the addition of moderate amounts of H₂O₂ (*ca.* 4 mM). The SERR spectra of Cyt-c adsorbed on PO₄-SAMs are also largely insensitive to the addition of similar amounts of H₂O₂ but in this case they reveal, in addition to the native protein, a *ca.* 30% spectral contribution of a second species consistent with the replacement of the heme axial ligand Met80 by a histidine residue (Fig. S9A†).⁵⁰ A similar contribution of bis-His species is also observed for Cyt-c adsorbed on NH₂-SAMs (Fig. S9B†). The assignment of the bis-His species is based on the spectral parameters of the high frequency region shown in Table S2† and on the appearance of a band at 405 cm⁻¹ assigned to the asymmetric stretching of the heme Fe bound to two imidazolic axial ligands. Upon treatment of the Cyt-c/NH₂-SAM complexes with 1 mM H₂O₂, we detect the appearance of a new spectroscopic component that resembles the RR spectrum of the ferric M80A mutant and, therefore, is compatible with a OH⁻/His coordination pattern (Fig. 3 and S10†).⁵¹ The rise of the OH⁻/His spectral component is concomitant with the appearance of the pseudoperoxidatic redox couple and occurs at the expense of the native spectral component, without significant alteration of the bis-His contribution (Fig. S10†). Nevertheless, within the error of the

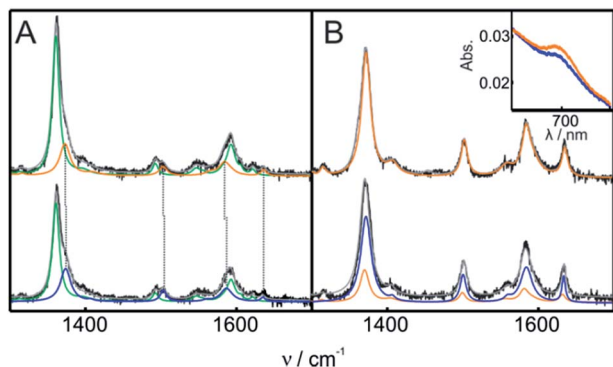


Fig. 3 (A) SERR spectra of Cyt-c adsorbed on an Ag electrode coated with NH₂-SAMs recorded before (top) and after (bottom) the addition of 1 mM H₂O₂ (final concentration). The spectra were acquired while cycling the electrode potential between 0.1 V and -0.3 V at 0.05 V s⁻¹ in 10 mM phosphate buffer, pH = 7. The colors indicate the different spectral components. Green: ferrous native Cyt-c. Orange: ferric native Cyt-c. Blue: OH⁻/His ferric component. (B) RR spectra of 0.2 mM ferric Cyt-c in 10 mM phosphate buffer, pH = 7, containing 0.5 mM PC/PE liposomes. The spectra were recorded before (top) and after (bottom) the addition of 1 mM H₂O₂ (final concentration). The color code of the RR spectral components is the same as that for the SERR spectra in panel (A). The inset shows the UV-vis charge transfer band before (orange) and after (blue) the addition of H₂O₂.

quantification, we cannot discard the possibility that the OH⁻/His species actually originates from the reaction of H₂O₂ with the bis-His species, which is in equilibrium with the main native component.

The fact that the adsorbed Cyt-c is sensitive to H₂O₂ only on amino-terminated SAMs suggests the involvement of hydroxyl radical species ([•]OH) that may be generated by the reaction of the amine groups of the SAMs with H₂O₂.⁵² To test this possibility, we performed a H₂O₂ activation of Cyt-c/NH₂-SAM complexes in the presence of DMSO, a very efficient scavenger of [•]OH.^{53–55} The results were identical to those obtained in the absence of DMSO, thereby indicating that, most likely, [•]OH is not involved in the formation of the pseudoperoxidatic OH⁻/His species.

In contrast to the native Cyt-c, the reduction potential of the OH⁻/His species adsorbed on NH₂- and QA-SAMs exhibits a strong linear pH dependence (Fig. 4A and S5B†), with a slope of -0.057 V per pH unit, which is consistent with the uptake of one proton per incoming electron. For the M80A mutant of Cyt-c cross-linked to electrodes coated with COOH-terminated SAMs, we obtain an almost identical pH-dependence (Fig. 4B). Moreover, the slope is also very similar to the one reported for horseradish peroxidase,⁵⁶ thus further suggesting close similarities of this species to the “professional” peroxidases.

The treatment of Cyt-c/liposome complexes with H₂O₂ yields results very similar to the Cyt-c/NH₂-SAM complexes. On the one

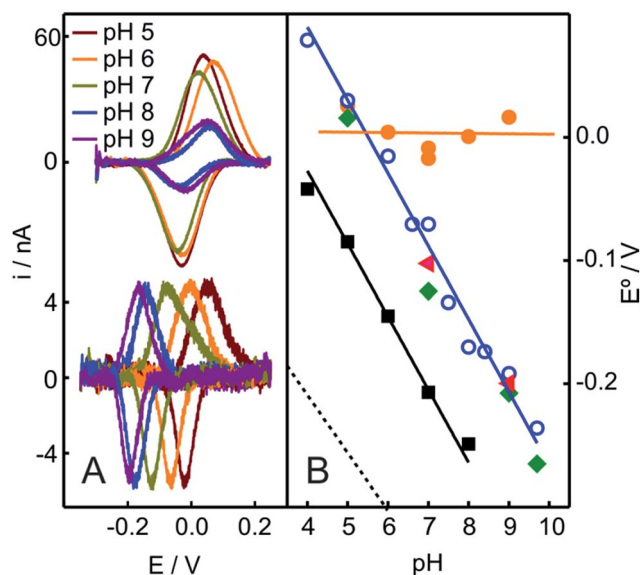


Fig. 4 (A) Voltammetric signals of Cyt-c adsorbed on NH₂-SAMs recorded at different pH values before (top) and after (bottom) the addition of 2 mM H₂O₂ (final concentration). (B) pH-dependencies of the reduction potentials of different protein species adsorbed on NH₂-SAM electrodes. Orange circles: native Cyt-c. Blue circles: Cyt-c adsorbed on the same electrode after *in situ* pretreatment with 1 mM H₂O₂. Green diamonds: Cyt-c pretreated *ex situ* with 1 mM H₂O₂ in the presence of 5 mM PE liposomes. Red triangles: *ex situ* synthesized and purified SO-Cyt. Additionally, the black squares correspond to the M80A Cyt-c mutant cross-linked to a COO-SAM (this work) and the dotted line corresponds to horseradish peroxidase on glassy carbon electrodes, adopted from ref. 56.



hand, the RR spectra of the treated Cyt-c/liposome systems are largely dominated by the same spectral component assigned to a ferric OH^-/His species and, moreover, this spectral change is complemented by a decrease of the characteristic MetS-Fe charge transfer absorption band at 695 nm (Fig. 3B). On the other, the reduction potentials of the Cyt-c products generated by treatment with H_2O_2 in both types of complexes exhibit identical values and pH dependencies (Fig. 4B).

Hence, the spectroscopic and electrochemical results point to the same chemical modification of Cyt-c in the different phosphate-mediated biomimetic complexes that lead to the detachment of the native axial ligand Met80 and the concomitant gain in catalytic activity.

A potential target for the reaction with H_2O_2 is the sulfur atom of the Met80 residue, as shown for methionines in general with a variety of oxidant species.^{42,57–59} Therefore, we treated Cyt-c with chloramine-T, which has been described as a selective reagent to oxidize Met80 and generate the corresponding sulf-oxide species (SO-Cyt).^{42,43} The product was purified by high-performance liquid chromatography and characterized by mass spectrometry-based peptide mapping (see ESI† for further details).

The purified SO-Cyt exhibits an identical RR spectrum to that of the H_2O_2 -generated OH^-/His species obtained in the biomimetic complexes (Fig. S13†). Moreover, as shown in Fig. 2D and 4B, when the *ex situ* generated SO-Cyt is adsorbed on electrodes coated with NH_2 -SAMs, it exhibits nearly identical voltammetric responses and pH dependencies as the *in situ* generated OH^-/His species.

The CV experiments reveal very similar heterogeneous electron transfer rate constants for native Cyt-c and SO-Cyt at all pH values (Table S1†), but a large downshift of the reduction potential of SO-Cyt at physiological pH. Thus, the loss of electron shuttling ability of Cyt-c upon Met80 oxidation is essentially a thermodynamic effect.

In summary, the presented results provide convincing evidence that submillimolar concentrations of H_2O_2 are able to efficiently and selectively oxidize the axial M80 ligand of Cyt-c in phosphate-mediated complexes with the main lipid components of the mitochondrial membrane. The oxidized product SO-Cyt is stable and undergoes a coordination and conformational transition that leads to a gain in pseudoperoxidatic activity. The remaining question of whether this transformation also results in a significant gain in truly peroxidatic activity is addressed next.

Peroxidatic and pseudoperoxidatic activity

The peroxidase-like activity of *in situ* generated SO-Cyt was evaluated following different strategies depending on whether it was obtained on NH_2 -SAMs or on PC/PE liposomes. Both peroxidase and pseudoperoxidase activities refer to the ability to catalyze the reduction of H_2O_2 to water. In the first case, electrons are supplied by a molecule (substrate) in solution, which is then oxidized, and homogeneous reaction rates are measured. In contrast, for pseudoperoxidase activity we refer to the electrocatalytic reduction of H_2O_2 by the immobilized

protein using the heterogeneous electron transfer reaction from the metal electrode to the protein as the electron source. The resulting currents measured by CV in the presence of increasing concentrations of H_2O_2 are a measure of the reaction rate. As shown in Fig. 5, the addition of H_2O_2 results in enhanced cathodic currents, without a significant alteration of other parameters such as peak potentials or FWHMs. This type of electrocatalytic behavior has been previously reported for the M80A mutant of yeast iso-cytochrome c, and was interpreted by a mechanism that involves O_2 generation and its subsequent reduction.⁶⁰ On the other hand, for the same mutant of the mammalian protein, Águila and coworkers showed some evidence of the formation of $\text{Fe}(\text{III})\text{OOH}$ (compound 0) in the reaction with H_2O_2 , which then would evolve to ferryl species (compounds I and II).⁶¹ Assuming that this mechanism is correct, and considering that the reduction potential of the $\text{Fe}(\text{IV})=\text{O}$ intermediate is significantly more positive than that of the other species,⁶² we can quantitatively analyze the cathodic peak currents in Fig. 5 in terms of a Michaelis–Menten formalism (see ESI† for further details).

The catalytic parameters obtained through this analysis are summarized in Table 1. While the k_{cat} values obtained for *in situ* and *ex situ* generated SO-Cyt are similar to native Cyt-c, the oxidation of M80 results in a *ca.* 300–600 times increase in affinity towards H_2O_2 . As a reference, the value of K_{m} reported for horseradish peroxidase on glassy carbon electrodes is *ca.* 53 μM ,⁵⁶ while for the M80A mutant of iso-cytochrome c it is only 3.25 μM , although this unusually high affinity is accompanied by protein bleaching at H_2O_2 concentrations of about 4 μM .⁶⁰

The peroxidatic activity of the different species in solution was assessed fluorometrically using Amplex UltraRed reagent. As summarized in Table 1 and Fig. 6, both the oxidation of Met80 (SO-Cyt) as well as its replacement by non-coordinating alanine (M80A) result in increased peroxidatic activity with respect to native Cyt-c by almost one order of magnitude, in agreement with previous observations.^{39,65} The effect is quantitatively similar for purified SO-Cyt generated *ex situ* and for Cyt-c treated with H_2O_2 in the presence of PE/PC liposomes. In spite

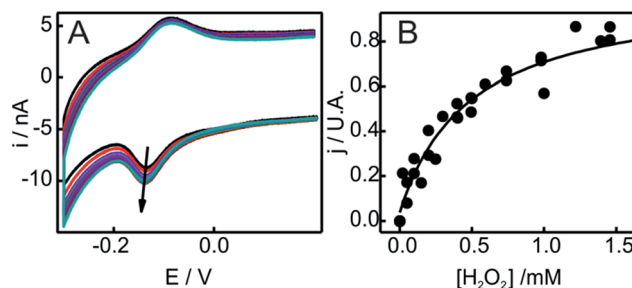


Fig. 5 (A) Voltammetric response as a function of H_2O_2 concentration of SO-Cyt generated *in situ* upon pretreatment with H_2O_2 of Cyt-c adsorbed on an NH_2 -SAM electrode. All measurements were performed in 10 mM phosphate buffer, pH = 7, at 0.05 V s^{-1} . The arrow indicates increasing concentrations of H_2O_2 from 0 to 2 mM. (B) Normalized cathodic peak current densities obtained from panel (A) as a function of H_2O_2 concentration. The line corresponds to the fit to the Michaelis–Menten-like equation (see ESI†).



Table 1 Pseudoperoxidase and peroxidase activities of the different Cyt-c variants^a

	Activity	K_m/mM^{-1}	$k_{\text{cat}}/\text{s}^{-1}$	% of control
WT	Pseudoperoxidase	144.3 ^c	8 ^c	—
	Peroxidase	65 ^d	5.5 ^d	100
SO-Cyt <i>ex situ</i>	Pseudoperoxidase ^b	0.24 (0.08)	2.6 (0.8)	—
	Peroxidase	19 ^e	—	719
SO-Cyt <i>in situ</i>	Pseudoperoxidase ^b	0.44 ^f (0.07)	1.8 ^f (0.6)	—
	Peroxidase	—	—	644 ⁱ
M80A	Pseudoperoxidase	0.00325 ^g	—	—
	Peroxidase	23.2 ^h	72 ^h	780

^a Standard deviations are indicated in brackets. ^b Determined for Cyt-c adsorbed on NH_2 -SAMs, this work. ^c Cyt-c adsorbed on COO -SAMs, from ref. 45. ^d Cyt-c in solution with ABTS, from ref. 63. ^e SO-Cyt in solution with ABTS, from ref. 64. ^f SO-Cyt obtained *in situ* on NH_2 -SAMs, this work. ^g From ref. 60. ^h Iso-cyt M80A mutant, from ref. 65. ⁱ SO-Cyt obtained by pretreatment with H_2O_2 in the presence of PC/PE liposomes.

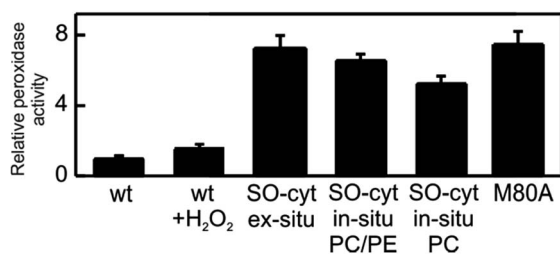


Fig. 6 Peroxidase activities of the different Cyt-c variants relative WT Cyt-c, as determined fluorometrically using Amplex UltraRed reagent.

of the bleaching effect, the treatment of a Cyt-c solution with only H_2O_2 has no significant effect on the measured peroxidatic activity (Fig. 6). Similarly, the mere presence of PE/PC liposomes has no effect unless H_2O_2 is added, thus confirming that the H_2O_2 -induced gain in peroxidatic activity is selective for Cyt-c molecules adsorbed on zwitterionic lipids *via* mediating phosphate anions.

4. Conclusions

Cyt-c is able to interact electrostatically with the main lipid components of the mitochondrial membranes, the zwitterionic lipids PE and PC, in the presence of biologically relevant amounts of mediating phosphate anions. Under these conditions, the adsorbed protein reacts efficiently with H_2O_2 . The reaction does not lead to the bleaching of the heme group, but is restricted to the oxidation of the sulfur in the iron axial ligand Met80, leading to its detachment and to a concomitant increase of the peroxidatic activity by one order of magnitude.

There is no evidence, though, that the PE and PC lipids are peroxidated by Cyt-c to significant amounts.^{10,23} Cardiolipin, in contrast, is effectively peroxidated by Cyt-c during the early events of apoptosis, thus leading to membrane permeabilization. The gain in peroxidatic activity has been ascribed to specific Cyt-c/CL interactions. The present findings, on the other hand, suggest a possible additional pathway for membrane permeabilization. Specifically, the rise of H_2O_2 concentrations to submillimolar levels that characterizes the

initiation of apoptosis^{37,38} appears to be sufficient to chemically modify a fraction of the Cyt-c molecules that interact (*via* phosphate anions) with the PE and PC membrane components. The modified protein is a stable peroxidase that could later bind and catalyze the oxidation of membrane components such as CL, thus facilitating the liberation of pro-apoptotic factors, including unmodified Cyt-c.

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Notes and references

- R. A. Scott and A. G. Mauk, Cytochrome c: a multidisciplinary approach, *Univ Science Books*, Sausalito, California, 1995.
- M. Hüttemann, P. Pecina, M. Rainbolt, T. H. Sanderson, V. E. Kagan, L. Samavati, J. W. Doan and I. Lee, *Mitochondrion*, 2011, **11**, 369.
- X. Liu, C. N. Kim, J. Yang, R. Jemmerson and X. Wang, *Cell*, 1996, **86**, 147.
- Y. Kushnareva and D. Newmeyer, *Ann. N. Y. Acad. Sci.*, 2010, **50**, 1201.
- R. Caroppi, P. Sinibaldi, F. Fiorucci and L. Santucci, *Curr. Med. Chem.*, 2009, **16**, 4058.
- X. Jiang and X. Wang, *Annu. Rev. Biochem.*, 2004, **73**, 87.
- A. V. Kulikov, E. S. Shilov, I. A. Mufazalov, V. Gogvadze, S. A. Nedospasov and B. Zhivotovsky, *Cell. Mol. Life Sci.*, 2012, **69**, 1787.
- C. Garrido, L. Galluzzi, M. Brunet, P. E. Puig, C. Didelot and G. Kroemer, *Cell Death Differ.*, 2006, **13**, 1423.
- V. Gogvadze, S. Orrenius and B. Zhivotovsky, *Biochim. Biophys. Acta*, 2006, **1757**, 639.



- 10 V. E. Kagan, V. A. Tyurin, J. Jiang, Y. Y. Tyurina, V. B. Ritov, A. A. Amoscato, A. N. Osipov, N. A. Belikova, A. A. Kapralov, V. Kini, I. I. Vlasova, Q. Zhao, M. Zou, P. Di, D. A. Svistunenko, I. V. Kurnikov and G. G. Borisenko, *Nat. Chem. Biol.*, 2005, **1**, 223.
- 11 V. E. Kagan, H. A. Bayır, N. A. Belikova, O. Kapralov, Y. Y. Tyurina, V. A. Tyurin, J. Jiang, D. A. Stoyanovsky, P. Wipf, P. M. Kochanek, J. S. Greenberger, B. Pitt, A. A. Shvedova and G. Borisenko, *Free Radicals Biol. Med.*, 2009, **46**, 1439.
- 12 P. Ascenzi, C. Ciaccio, F. Sinibaldi, R. Santucci and M. Coletta, *Biochem. Biophys. Res. Commun.*, 2011, **404**, 190.
- 13 P. Ascenzi, R. Santucci, M. Coletta and F. Polticelli, *Biophys. Chem.*, 2010, **152**, 21.
- 14 V. M. Trusova, G. P. Gorbenko, J. G. Molotkovsky and P. K. J. Kinnunen, *Biophys. J.*, 2010, **99**, 1754.
- 15 L. Macchioni, T. Corazzi, M. Davidescu, E. Francescangeli, R. Roberti and L. Corazzi, *Mol. Cell. Biochem.*, 2010, **341**, 149.
- 16 F. Sinibaldi, B. D. Howes, M. C. Piro, F. Polticelli, C. Bombelli, T. Ferri, M. Coletta, G. Smulevich and R. Santucci, *JBIC, J. Biol. Inorg. Chem.*, 2010, **15**, 689.
- 17 J. Hanske, J. R. Toffey, A. M. Morenz, A. J. Bonilla, K. H. Schiavoni and E. V. Pletneva, *Proc. Natl. Acad. Sci. U. S. A.*, 2012, **109**, 125.
- 18 C. L. Bergstrom, P. A. Beales, Y. Lv, T. K. Vanderlick and J. T. Groves, *Proc. Natl. Acad. Sci. U. S. A.*, 2013, **110**, 6269.
- 19 Y. A. Vladimirov, E. V. Proskurnina and A. V. Alekseev, *Biokhim., Biochem.*, 2013, **78**, 1086.
- 20 J. M. Bradley, G. Silkstone, M. T. Wilson, M. R. Cheesman and J. N. Butt, *J. Am. Chem. Soc.*, 2011, **133**, 19676.
- 21 S. L. Iverson and S. Orrenius, *Arch. Biochem. Biophys.*, 2004, **423**, 37.
- 22 Y.-L. P. Ow, D. R. Green, Z. Hao and T. W. Mak, *Nat. Rev. Mol. Cell Biol.*, 2008, **9**, 532.
- 23 N. Belikova and Y. Vladimirov, *Biochemistry*, 2006, **45**, 4998.
- 24 E. Kalanxhi and C. J. A. Wallace, *Biochem. J.*, 2007, **407**, 179.
- 25 C. Osman, D. R. Voelker and T. Langer, *J. Cell Biol.*, 2011, **192**, 7.
- 26 D. Ardail, J. Privat and M. Egret-Charlier, *J. Biol. Chem.*, 1990, **265**, 18797.
- 27 N. Gebert, A. S. Joshi, S. Kutik, T. Becker, M. McKenzie, X. L. Guan, V. P. Mooga, D. A. Stroud, G. Kulkarni, M. R. Wenk, P. Rehling, C. Meisinger, M. T. Ryan, N. Wiedemann, M. L. Greenberg and N. Pfanner, *Curr. Biol.*, 2009, **19**, 2133.
- 28 M. Garcia Fernandez, L. Troiano, L. Moretti, M. Nasi, M. Pinti, S. Salvioli, J. Dobrucki and A. Cossarizza, *Cell Growth Differ.*, 2002, **13**, 449.
- 29 Z. T. Schug and E. Gottlieb, *Biochim. Biophys. Acta*, 2009, **1788**, 2022.
- 30 J. Montero, M. Mari, A. Colell, A. Morales, G. Basañez, C. Garcia-Ruiz and J. C. Fernández-Checa, *Biochim. Biophys. Acta*, 2010, **1797**, 1217.
- 31 M. Schlame, D. Rua and M. L. Greenberg, *Prog. Lipid Res.*, 2000, **39**, 257.
- 32 V. E. Kagan, G. G. Borisenko, Y. Y. Tyurina, V. A. Tyurin, J. Jiang, A. I. Potapovich, V. Kini, A. A. Amoscato and Y. Fujii, *Free Radicals Biol. Med.*, 2004, **37**, 1963.
- 33 Z. Salamon and G. Tollin, *Biophys. J.*, 1996, **71**, 848.
- 34 T. Ahn, D. B. Oh, B. C. Lee and C. H. Yun, *Biochemistry*, 2000, **39**, 10147.
- 35 D. A. Capdevila, W. A. Marmisollé, F. J. Williams and D. H. Murgida, *Phys. Chem. Chem. Phys.*, 2013, **15**, 5386.
- 36 W. A. Marmisollé, D. A. Capdevila, E. de la Llave, F. J. Williams and D. H. Murgida, *Langmuir*, 2013, **29**, 5351.
- 37 M. Giorgio, M. Trinei, E. Migliaccio and P. G. Pelicci, *Nat. Rev. Mol. Cell Biol.*, 2007, **8**, 722.
- 38 C. Cerella, S. Coppola, V. Maresca, M. De Nicola, F. Radogna and L. Ghibelli, *Ann. N. Y. Acad. Sci.*, 2009, **1171**, 559.
- 39 L. C. Godoy, C. Muñoz-Pinedo, L. Castro, S. Cardaci, C. M. Schonhoff, M. King, V. Tórtora, M. Marin, Q. Miao, J. F. Jiang, A. Kapralov, R. Jemmerson, G. G. Silkstone, J. N. Patel, J. E. Evans, M. T. Wilson, D. R. Green, V. E. Kagan, R. Radi and J. B. Mannick, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**, 2653.
- 40 D. H. Murgida and P. Hildebrandt, *J. Phys. Chem. B*, 2001, **105**, 1578.
- 41 S.-H. Lee, W.-C. Lin, C.-H. Kuo, M. Karakachian, Y.-C. Lin, B.-Y. Yu and J.-J. Shyue, *J. Phys. Chem. C*, 2010, **114**, 10512.
- 42 Y.-R. Chen, L. J. Deterding, B. E. Sturgeon, K. B. Tomer and R. P. Mason, *J. Biol. Chem.*, 2002, **277**, 29781.
- 43 J. Pande, K. Kinnally, K. K. Thallum, B. Verma, Y. Myer, L. Rechsteiner and H. Bosshard, *J. Protein Chem.*, 1987, **6**, 295.
- 44 B. Valderrama, M. Ayala and R. Vazquez-Duhalt, *Chem. Biol.*, 2002, **9**, 555.
- 45 L. Wang and D. H. Waldeck, *J. Phys. Chem. C*, 2008, **112**, 1351.
- 46 D. Alvarez-Paggi, D. F. Martín, P. M. DeBiase, P. Hildebrandt, M. A. Martí and D. H. Murgida, *J. Am. Chem. Soc.*, 2010, **132**, 5769.
- 47 X. Chen, R. Ferrigno, J. Yang and G. M. Whitesides, *Langmuir*, 2002, **18**, 7009.
- 48 D. H. Murgida and P. Hildebrandt, *Acc. Chem. Res.*, 2004, **37**, 854.
- 49 G. Battistuzzi, M. Borsari, J. A. Cowan, A. Ranieri and M. Sola, *J. Am. Chem. Soc.*, 2002, **124**, 5315.
- 50 D. H. Murgida and P. Hildebrandt, *Chem. Soc. Rev.*, 2008, **37**, 937.
- 51 G. Battistuzzi, C. A. Bortolotti, M. Bellei, G. Di Rocco, J. Salewski, P. Hildebrandt and M. Sola, *Biochemistry*, 2012, **51**, 5967.
- 52 G. Cohen and R. Heikkilä, *J. Biol. Chem.*, 1974, **249**, 2447.
- 53 R. C. Scaduto, *Free Radicals Biol. Med.*, 1995, **18**, 271.
- 54 M. G. Steiner and C. F. Babbs, *Arch. Biochem. Biophys.*, 1990, **278**, 478.
- 55 C. Tai, J.-F. Peng, J.-F. Liu, G.-B. Jiang and H. Zou, *Anal. Chim. Acta*, 2004, **527**, 73.
- 56 B. Wang, J.-J. Zhang, Z.-Y. Pan, X.-Q. Tao and H.-S. Wang, *Biosens. Bioelectron.*, 2009, **24**, 1141.
- 57 W. Vogt, *Free Radicals Biol. Med.*, 1995, **18**, 93–105.



- 58 B. Valderrama and R. Vazquez-Duhalt, *J. Mol. Catal. B: Enzym.*, 2005, **35**, 41.
- 59 J. Thariat, F. Collin, C. Marchetti, N. S. Ahmed-Adrar, H. Vitrac, D. Jore and M. Gardes-Albert, *Biochimie*, 2008, **90**, 1442.
- 60 S. Casalini, G. Battistuzzi, M. Borsari, C. A. Bortolotti, G. Di Rocco, A. Ranieri and M. Sola, *J. Phys. Chem. B*, 2010, **114**, 1698.
- 61 S. Águila, A. M. Vidal-Limón, J. B. Alderete, M. Sosa-Torres and R. Vázquez-Duhalt, *J. Mol. Catal. B: Enzym.*, 2013, **85–86**, 187.
- 62 G. Battistuzzi, M. Bellei, C. A. Bortolotti and M. Sola, *Arch. Biochem. Biophys.*, 2010, **500**, 21.
- 63 R. Radi, L. Thomson, H. Rubbo and E. Prodanov, *Arch. Biochem. Biophys.*, 1991, **288**, 112.
- 64 X. Chen, R. Ferrigno, J. Yang and G. Whitesides, *Langmuir*, 2002, 7009.
- 65 A. Ranieri, F. Bernini, C. A. Bortolotti and E. Castellini, *Catal. Sci. Technol.*, 2012, **2**, 2206.

