

# Characterization of cytochrome *b* in the isolated ubiquinol–cytochrome *c*<sub>2</sub> oxidoreductase from *Rhodopseudomonas sphaeroides* GA

Nadia Gabellini and Günter Hauska

*Institut für Botanik, Universität Regensburg, 8400 Regensburg, FRG*

Received 7 January 1983

Extinction coefficients for cytochrome *b* and *c*<sub>1</sub> in the isolated cytochrome *bc*<sub>1</sub> complex from *Rhodopseudomonas sphaeroides* GA have been determined. They are 25 mM<sup>-1</sup>.cm<sup>-1</sup> at 561 nm for cytochrome *b* and 17.4 mM<sup>-1</sup>.cm<sup>-1</sup> at 553 nm for cytochrome *c*<sub>1</sub>, for the difference between the reduced and the oxidized state. Cytochrome *b* is present in two forms in the complex. One form has an *E*<sub>m7</sub> of 50 mV, an  $\alpha$ -peak of 557 nm at liquid N<sub>2</sub> temperature and of 561 nm at RT, which is red-shifted by antimycin A. The other form has an *E*<sub>m7</sub> of –90 mV, a double  $\alpha$ -peak of 555 and 561 nm at liquid N<sub>2</sub> temperature corresponding to 559 and 566 nm at RT. The absorption at 566 nm is red-shifted by myxothiazol. The two shifts are independent of each other. Both midpoint potentials of cytochromes *b* are pH-dependent. The redox center compositions of the cytochrome *bc*<sub>1</sub> complexes from *Rhodopseudomonas sphaeroides* and from mitochondria are identical.

<i>Bacterial bc<sub>1</sub> complex</i>	<i>Rps. sphaeroides</i>	<i>Cytochrome b</i>	<i>Myxothiazol</i>	<i>Antimycin A</i>
	<i>Ubiquinol–cytochrome c oxidoreductase</i>			

## 1. INTRODUCTION

A cytochrome *bc*<sub>1</sub> complex with ubiquinol–cytochrome *c*<sub>2</sub> oxidoreductase activity has been isolated recently from *Rhodopseudomonas sphaeroides* GA [1]. The preparation was sensitive to antimycin A and UHDBT. It contained two hemes *b* and substoichiometric amounts of the Rieske FeS center and of ubiquinone/cytochrome *c*<sub>1</sub>. On SDS–PAGE 3 main polypeptides of *M*<sub>r</sub> 40000, 34000 and 25000 were resolved. Two midpoint potentials of 50 and –60 mV at pH 7 could be ascribed to cytochrome *b* from redox titrations. These are similar to midpoint potentials of cytochromes *b* in the parent chromatophore membrane [2,3], the lower corresponding to a component with an  $\alpha$ -peak at 566 nm plus a shoulder at 559 nm, the higher corresponding to a component with an  $\alpha$ -peak at 561 nm. Such a spectral heterogeneity of cytochrome *b* has not been found at first in the isolated cytochrome *bc*<sub>1</sub> complex [1],

the  $\alpha$ -peak remaining at 561 nm throughout the redox titration.

Here, we provide unequivocal evidence for the two forms of cytochrome *b* in the isolated complex, which in addition to redox titration [1] is based on the independent spectral effects of antimycin A and myxothiazol, and on low temperature spectroscopy.

## 2. MATERIALS AND METHODS

The cytochrome *bc*<sub>1</sub> complex was isolated as described [1], except that Triton X-100 was avoided for resuspending the pellet after ammonium sulfate fractionation. Instead the octylglucoside concentration was raised to 120 mM in the suspending mixture. In this case the complex banded at 30% sucrose in the subsequent density gradient centrifugation [1]. Oxidoreductase activity and oxidant-induced reduction were measured as described [1], with the specifications given in the

legends. With ubiquinol-2 a maximal turnover of  $20 \text{ s}^{-1}$  was found for cytochrome  $c_1$  in the complex. Low temperature spectra and the spectral shifts caused by myxothiazol and antimycin A were measured in an Aminco DW2 UV/Vis spectrophotometer as described in the legends. Pyridine hemochrome was prepared and assayed as in [4]. Myxothiazol was a kind gift from Dr Thierbach (Braunschweig). Other sources for materials were listed in [1,5].

### 3. RESULTS AND DISCUSSION

From the redox difference spectra and the corresponding pyridine hemochrome spectra in fig.1 extinction coefficients of  $17.5 \text{ mM}^{-1} \cdot \text{cm}^{-1}$  at 553 nm for cytochrome  $c_1$ , and of  $25 \text{ mM}^{-1} \cdot \text{cm}^{-1}$  at 561 nm for cytochrome  $b$  in the isolated complex can be calculated, which are similar to the values known for cytochromes  $b$  and  $c_1$  in the mitochondrial cytochrome  $bc_1$  complex [4,6,7]. Two hemes  $b$ /heme  $c$  are present in the bacterial complex (0.6 and  $0.3 \mu\text{M}$  in fig.1), confirming [1]. No reliable measurement is known for cytochrome  $b$  and  $c_1$  in the chromatophore membrane [8].

In [1] redox heterogeneity of cytochrome  $b$  was found, which was not paralleled by spectral heterogeneity, however. The  $\alpha$ -peak remained at

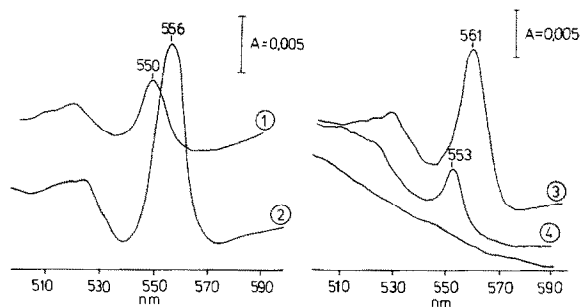


Fig.1. Pyridine hemochrome and cytochrome spectra. The measurements are described in section 2. Pyridine hemochromes were prepared as in [4]: (1) protein bound pyridine hemochrome, dithionite-reduced minus ferricyanide-oxidized; (2) acetone-soluble pyridine hemochrome, dithionite-reduced minus ferricyanide-oxidized; (3) cytochrome  $bc_1$  complex, dithionite minus ascorbate; (4) cytochrome  $bc_1$  complex, ascorbate minus ferricyanide. A baseline is included in the right part of the figure.

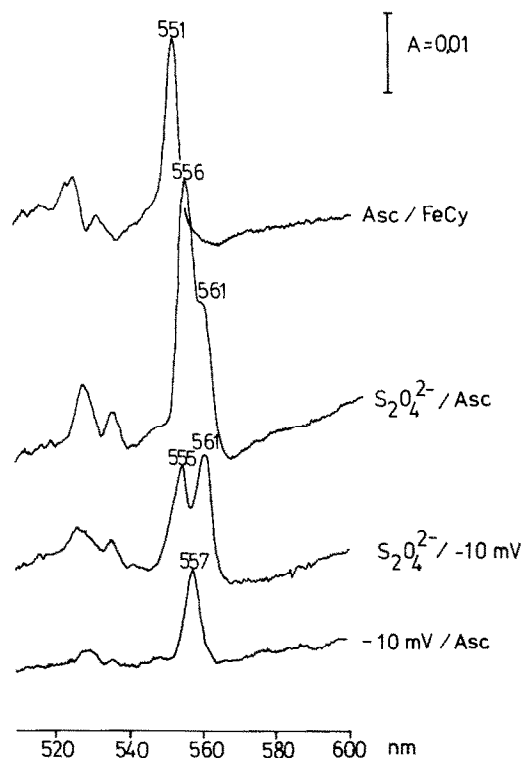


Fig.2. Low temperature redox difference spectra. The spectra were recorded with the low temperature equipment for the Aminco DW2 UV/Vis spectrophotometer over liquid nitrogen. The 2 mm-cuvettes were used. The cytochrome  $bc_1$  complex was diluted to  $0.4 \mu\text{M}$  cytochrome  $c_1$  in 5 mM glycylglycine (pH 7.4), 20% (v/v) glycerol,  $20 \mu\text{M}$  duroquinone. Before freezing the solution was either oxidized by ferricyanide (FeCy), or reduced by ascorbate (Asc) or dithionite, or was poised to an  $E_h$  of  $-10 \text{ mV}$ . Difference spectra between sample and reference cuvettes were taken as indicated.

561 nm during redox titrations. When Triton X-100 is left out during the preparation of the complex a shoulder at 559 nm with the main peak at 566 nm was detectable at low potentials. A clear spectral resolution of the two forms of cytochrome  $b$  in the complex is possible at low temperature (fig.2). In liquid nitrogen the component with the higher midpoint potential shows an  $\alpha$ -peak at 557 nm, the component with the lower midpoint potential shows two peaks at 555 and 561 nm, of almost equal height. The two components probably represent cytochrome  $b$ -561 and cytochrome  $b$ -566 (second peak at 559 nm at RT) in the chromatophore

membrane, recently resolved by redox difference, and flash spectroscopy [3,9]. Spectra of two forms of cytochrome *b* have been reported before for a less purified preparation of the cytochrome *bc*<sub>1</sub> complex [10], but they do not match our results.

Redox potential measurements [1] were repeated with our modified preparation, avoiding Triton X-100. The two  $E_{m7}$ -values match exactly the values reported for the cytochromes *b* in chromatophores [2,3], and both are pH-dependent (table 1). In chromatophores at least one cytochrome *b* has a pH-dependent midpoint potential [11].

The cytochromes *b*-561 and *b*-566 in the bacterial complex correspond to the cytochromes *b*-562 and *b*-565 in the mitochondrial complex, which have  $E_{m7}$ -values of 92 and -34 mV, respectively [7,12]. Also these midpoint potentials are pH-dependent [13].

In fig.3 spectral shifts of the  $\alpha$ -peaks of cytochrome *b* in the isolated complex by antimycin A and myxothiazol are shown. Both cause a red shift, but the effects are observed in the absence as well as in the presence of the other antibiotic. Thus they seem to be independent of each other, antimycin affecting the component resembling *b*-561 and myxothiazol the component resembling *b*-566. These results are identical to a report for the isolated mitochondrial cytochrome *bc*<sub>1</sub> complex [14]. Independent spectral changes of cytochrome *b* by antimycin A and myxothiazol have also been

Table 1  
pH-Dependence of the midpoint potentials of cytochrome *b*

pH	Midpoint potential (mV)	
	<i>b<sub>h</sub></i>	<i>b<sub>l</sub></i>
6.0	+ 100	- 47
7.0	+ 50	- 90
8.0	+ 35	- 110

Redox titration of the isolated cytochrome *bc*<sub>1</sub> complex was carried out as in [1]. The  $E_m$ -values were obtained with the assumption of two  $n = 1$  components present in equal amounts [1]. The pH was adjusted with glycylglycine, Hepes and MES at 50 mM total buffer concentration; *b<sub>h</sub>* and *b<sub>l</sub>* stand for high-potential and low potential cytochrome *b*, respectively

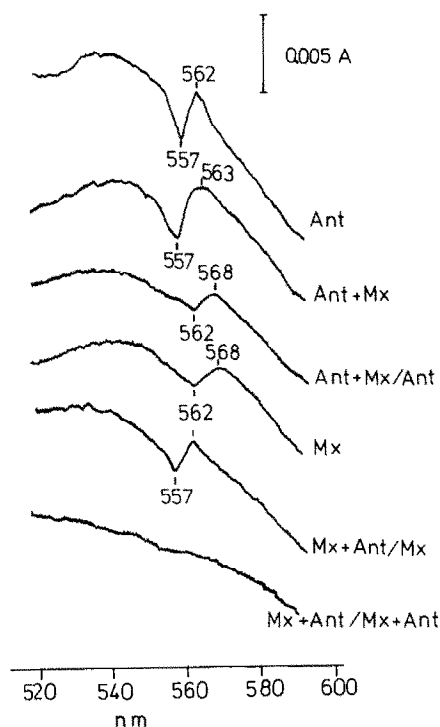


Fig.3. Shifts of the  $\alpha$ -peaks of the cytochromes *b* by antimycin A and myxothiazol. The difference spectra were recorded at RT with the cytochrome *bc*<sub>1</sub> complex in 50 mM Na-acetate (pH 5.8), 0.9  $\mu$ M in cytochrome *c*<sub>1</sub>. The complex was reduced by dithionite, antimycin A was added to 10  $\mu$ M to the sample cuvette and the difference spectrum to the reference cuvette without antimycin A was recorded (Ant). Then myxothiazol was added to 10  $\mu$ M to the sample cuvette (Ant + Mx), followed by addition of antimycin to the reference cuvette (Ant + Mx/Ant). In a second set of recordings (lower 3 spectra) the sequence of additions was changed as indicated.

reported for the chromatophore membrane [15] but a blue shift was observed with myxothiazol in contrast to the red shift with antimycin A, which has been found before [16]. Since, also in the mitochondrial membrane, the spectral change caused by myxothiazol [17] does not resemble the one found for the isolated *bc*<sub>1</sub> complex [14], we must conclude that cytochrome *b*-566 changes during isolation of the complexes from the membrane, even if Triton X-100 is avoided. The spectral shift by myxothiazol is best observable at slightly acidic pH (see [15]).

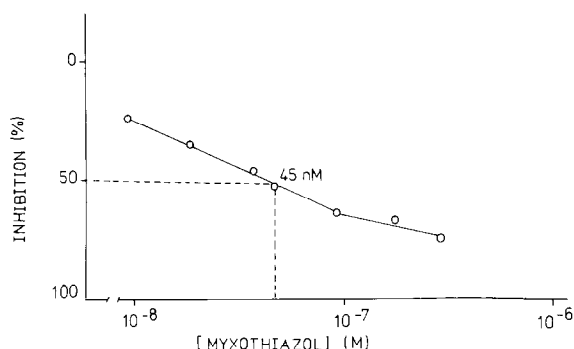


Fig. 4. Inhibition of ubiquinol-cytochrome *c* oxidoreductase activity by myxothiazol. The reaction mixture contained the cytochrome *bc*<sub>1</sub> complex, 37 nM in cytochrome *c*<sub>1</sub>, 50 mM MES-NaOH (pH 6.2), 20  $\mu$ M cytochrome *c* from horse heart, and the indicated concentrations of myxothiazol. The reaction was started by the addition of ubiquinol-3 to 50  $\mu$ M. The uninhibited rate was 15  $\mu$ mol cytochrome *c* reduced  $\cdot$ nmol cytochrome *c*<sub>1</sub><sup>-1</sup>  $\cdot$ h<sup>-1</sup>.

Myxothiazol is an inhibitor of the ubiquinol--cytochrome *c* oxidoreductase activity of the isolated complex, almost as potent as antimycin A [1]; 50% inhibition is reached at a molar ratio of myxothiazol to cytochrome *c*<sub>1</sub> of 1.3 (fig. 4). Fig. 5 shows that myxothiazol inhibits ferricyanide-induced reduction of cytochrome *b*, but does not inhibit the antimycin A-sensitive, direct reduction of cytochrome *b* by ubiquinol. The results in fig. 5 fit a Q-cycle [18] or a b-cycle [19] mechanism for the ubiquinol-cytochrome *c* oxidoreductase, where myxothiazol inhibits the reduction of low-potential cytochrome *b*-566 by ubisemiquinone, formed from ubiquinol by oxidation with the Rieske FeS center, and antimycin A inhibits the reversible reoxidation of the high-potential cytochrome *b*-561 by ubiquinone in a different site or state. Exactly the same conclusion as to the sites of inhibition of the two antibiotics has been reached before in [15] for the chromatophore membrane, and in [14] for the isolated, mitochondrial cytochrome *bc*<sub>1</sub> complex.

#### ACKNOWLEDGEMENTS

The generous gift of myxothiazol by Dr G. Thierbach (Braunschweig) is gratefully acknowledged. We thank Dr A.R. Crofts for pro-

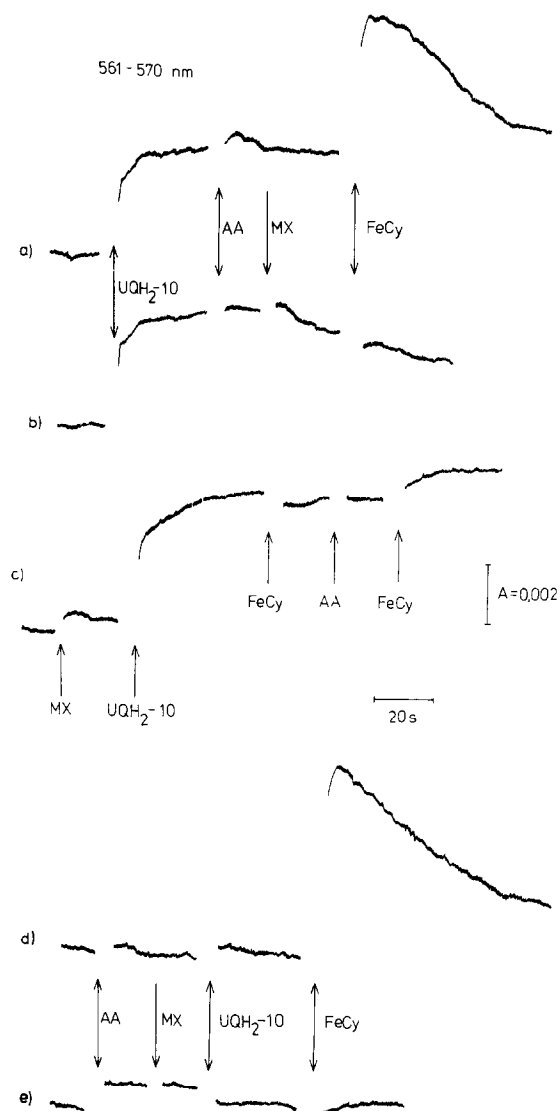


Fig. 5. Inhibition of oxidant-induced reduction of cytochrome *b* by myxothiazol. The traces were recorded in the dual wavelength mode with 561 nm as the measuring and 570 nm as the reference wavelength. The cytochrome *bc*<sub>1</sub> complex was diluted to 0.54  $\mu$ M in 50 mM glycylglycine (pH 7.4). A 1-ml cuvette was used without stirring. The additions during the experiments (a-e) were as indicated by arrows: Ubiquinol-10 (UQH<sub>2</sub>-10), 50 mM in 10% Triton X-100 to 250  $\mu$ M; antimycin A to 10  $\mu$ M (AA); myxothiazol to 45  $\mu$ M (MX); ferricyanide to 100  $\mu$ M (FeCy). The spectral changes after addition of UQH<sub>2</sub>-10 in traces a, b and c correspond to about 25%, the changes after addition of ferricyanide in traces a and d correspond to 30 and 40% of the dithionite-reducible cytochrome *b*.

viding access to unpublished work. Our research was supported by the Deutsche Forschungsgemeinschaft (SFB 43 C2).

## REFERENCES

- [1] Gabellini, N., Bowyer, J.R., Hurt, E., Melandri, B.A. and Hauska, G. (1982) *Eur. J. Biochem.* 126, 105–111.
- [2] Dutton, P.L. and Jackson, J.B. (1972) *Eur. J. Biochem.* 30, 495–510.
- [3] Bowyer, J.R., Meinhardt, S.W., Tierney, G.V. and Crofts, A.R. (1981) *Biochim. Biophys. Acta* 635, 167–186.
- [4] Rieske, J.S. (1967) *Methods Enzymol.* 10, 488–493.
- [5] Hurt, E. and Hauska, G. (1981) *Eur. J. Biochem.* 117, 591–599.
- [6] Hatefi, Y., Haavik, H.G. and Griffith, D.E. (1962) *J. Biol. Chem.* 237, 1681–1685.
- [7] Nelson, B.D. and Gellerfors, P. (1976) *Biochim. Biophys. Acta* 357, 358–364.
- [8] Bowyer, J.R. and Crofts, A.R. (1981) *Biochim. Biophys. Acta* 636, 218–233.
- [9] Meinhardt, S.W. and Crofts, A.R. (1983) *Biochim. Biophys. Acta*, in press.
- [10] Takamiya, K., Doi, M. and Okimatsu, H. (1982) *Plant Cell Physiol.* 23, 987–997.
- [11] Petty, K.M. and Dutton, P.L. (1976) *Arch. Biochem. Biophys.* 172, 346–353.
- [12] Von Jagow, G., Schagger, H., Engel, W.D., Riccio, P., Kolb, H.J. and Klingenberg, M. (1978) *Methods Enzymol.* 53, 92–98.
- [13] Von Jagow, G., Engel, W.D., Schagger, H., Machleidt, W. and Machleidt, I. (1981) in: *Vectorial Reactions in Electron and Ion Transport in Mitochondria and Bacteria* (Palmieri, F. et al. eds) *Dev. Bioenerg. Biomembr.* vol.8, pp.149–161, Elsevier Biomedical, Amsterdam, New York.
- [14] Von Jagow, G. and Engel, W.D. (1981) *FEBS Lett.* 136, 19–24.
- [15] Meinhardt, S.W. and Crofts, A.R. (1982) *FEBS Lett.* 149, 223–227.
- [16] Van den Berg, W.H., Prince, R.C., Bashford, C.L., Takamiya, K.-I., Bonner, W.D. and Dutton, P.L. (1979) *J. Biol. Chem.* 254, 8594–8604.
- [17] Thierbach, G. and Reichenbach, H. (1981) *Biochim. Biophys. Acta* 638, 282–289.
- [18] Mitchell, P. (1976) *J. Theor. Biol.* 62, 327–367.
- [19] Wikström, M., Krab, K. and Saraste, M. (1981) *Annu. Rev. Biochem.* 50, 623–655.