

Immunological Investigation of the Distribution of Cytochromes Related to the Two Terminal Oxidases of *Escherichia coli* in Other Gram-Negative Bacteria

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Monospecific antibodies were raised against the two terminal oxidase complexes of the aerobic respiratory chain of *Escherichia coli*. These are the cytochrome *d* and cytochrome *o* complexes. The antibodies were used to check for the occurrence of cross-reactive antigens in membrane preparations from a variety of gram-negative bacteria by rocket immunoelectrophoresis and immunoblotting techniques. With these criteria, proteins closely related to the cytochrome *d* complex of *E. coli* appeared to be widely distributed. Among the strains containing cytochrome *d*-related material were *Serratia marcescens*, *Photobacterium phosphoreum*, *Salmonella typhimurium*, *Klebsiella pneumoniae*, and *Azotobacter vinelandii*. The data suggest that the *d*-type terminal oxidase in many of these strains is associated in a complex with *b*-type and *a*₁-type cytochromes, as has been found to be the case in *E. coli*. *K. pneumoniae* and *S. typhimurium* were also shown to have material cross-reactive to the *E. coli* cytochrome *o* complex.

The aerobic respiratory chain of *Escherichia coli* is branched and contains two terminal oxidases, the cytochrome *d* complex and the cytochrome *o* complex (10). Both of these oxidases have been purified to homogeneity (25, 26, 31, 33, 37) and have been shown to carry out electrogenic reactions in reconstituted proteoliposomes (24, 28, 33). Electron flow through either oxidase generates a transmembrane voltage difference. Whereas the cytochrome *aa*₃-type terminal oxidases oxidize ferrocycytochrome *c*, the oxidases of *E. coli* appear to directly oxidize ubiquinol in the bacterial membrane (25, 28). The cytochrome *o* complex predominates in the *E. coli* membrane when cells are grown with high aeration (31), whereas the cytochrome *d* complex, which has a higher affinity for oxygen (44), is induced when cells are grown under oxygen-limiting conditions (31; see also references 10, 20, and 41).

The cytochrome *d* complex has been shown to contain three cytochrome components, *b*₅₅₈, *a*₁, and *d*, but only two polypeptides by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) (26, 27, 37). The larger of the two subunits (subunit I; molecular weight, 57,000) has been shown to contain the cytochrome *b*₅₅₈ component of the complex (8). One question of interest which is addressed in this work is whether other gram-negative bacteria, many of which contain *d*-type cytochrome by spectroscopic criteria (20, 41), contain immunocross-reactive material to either or both of these subunits. This was determined by using antibodies raised specifically against either subunit I or subunit II (molecular weight, 43,000) of the cytochrome *d* complex of *E. coli*.

The cytochrome *o* complex of *E. coli* contains heme *b* (25, 31, 33) and copper (25) and is reported to have two (25) or four (31, 33) subunits by SDS-PAGE analysis. Antibodies raised against the native oxidase were used to check for cross-reactivity with preparations from various gram-negative bacteria. Several of these strains contain *o*-type terminal

oxidases by spectroscopic criteria, and it was interesting to investigate their immunological relationships.

MATERIALS AND METHODS

Organisms and growth conditions. The bacterial strains used in this work are listed in Table 1. The local isolates were taken from stocks used for teaching purposes, and most were originally obtained from the American Type Culture Collection. Before use, the strains were checked by metabolic characterization tests. Most bacterial strains were grown in Penassay broth (50 ml or 100 ml; Difco Laboratories, Detroit, Mich.) in 250 ml Klett flasks. The flasks were shaken at 200 rpm at the recommended temperature for each strain, and the cells were harvested at stationary phase. *Photobacterium phosphoreum* required 3% NaCl in the medium. Several strains, *Rhodospseudomonas palustris*, *Rhodospirillum rubrum*, *Rhodospirillum fulvum*, and *Rhodospseudomonas sphaeroides*, were grown in Sistrom basal medium (47). *Rhodospirillum fulvum* was grown microaerophilically in the dark for 2.5 days in medium supplemented with 5 μg of para-aminobenzoic acid per ml. *Pseudomonas putida* (48), *Klebsiella aerogenes* (11), and *Paracoccus denitrificans* (12) were grown in media and conditions previously described. *Azotobacter vinelandii* (19) was grown microaerophilically in Winogradsky medium with sucrose, agar, and molybdenum. *E. coli* SHSP19 (45), which does not synthesize heme, was grown semianaerobically.

Immunological methods. The antibody preparations used in this work have all been previously described. Polyclonal antibodies against *E. coli* cytochrome *o* were raised against immunopurified cytochrome (31). Polyclonal antibodies were raised against subunit II of the cytochrome *d* complex, which had been excised from an SDS-polyacrylamide gel after electrophoresis (29). A monoclonal antibody (A14-5) which has been shown to react with subunit I of the cytochrome *d* complex was also used (32).

Harvested cells were washed and disrupted by sonication. Membranes were prepared and solubilized by using Zwittergent 3-12 as previously described (31). For SDS-PAGE immunoblotting (2), samples containing 60 μg of membrane

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TABLE 1. Distribution of cross-reactive proteins to *E. coli* cytochromes *d* and *o* in various gram-negative bacteria

Bacterial strains	Source	Spectroscopic evidence for cytochrome <i>d</i>		Immunological evidence for cytochrome:	
		Reference ^c	This work	<i>d</i>	<i>o</i>
<i>Escherichia coli</i> K-12 (MR43L)	W. Shipp (46)	+ (9, 37, 42)	+	+	+
<i>Serratia marcescens</i>	LI ^a		ND ^b	+	-
<i>Enterobacter aerogenes</i>	LI	+ (38)	+	+	-
<i>Pseudomonas putida</i>	LI	+ (48, 49)	ND	-	-
<i>Proteus vulgaris</i>	LI	+ (39)	+	+	-
<i>Pseudomonas fluorescens</i>	LI		ND	-	-
<i>Escherichia coli</i> W191-6	(31)		+	+	+
<i>Proteus mirabilis</i>	LI		ND	+	-
<i>Pseudomonas aeruginosa</i>	LI	- (35)	ND	-	-
<i>Rhodopseudomonas palustris</i>	LI	- (20)	ND	-	-
<i>Paracoccus denitrificans</i>	LI	+ (12)	ND	-	-
<i>Photobacterium phosphoreum</i>	T. Baldwin	+ (54)	+	+	-
<i>Escherichia coli</i> SHSP19	J. Cronan		ND	+	-
<i>Rhodospirillum rubrum</i>	LI	- (20)	ND	-	-
<i>Rhodospirillum fulvum</i>	LI		ND	-	-
<i>Rhodopseudomonas sphaeroides</i>	LI	- (20)	ND	-	-
<i>Arthrobacter pyridinolis</i>	T. Krulwich	+ (40)	-	-	-
<i>Acinetobacter HO1N</i>	W. Finnerty	+ (5)	+	+	-
<i>Klebsiella pneumoniae</i>	R. Ugalde	+ (11)	-	+	+
<i>Salmonella typhimurium</i>	LI	+ (4)	+	+	+
<i>Azotobacter vinelandii</i>	R. Ugalde	+ (13, 14, 18, 19, 21, 22, 36)	+	+	-
<i>Vitreoscilla</i> sp.		- (20)	ND	-	-

^a LI, University of Illinois local isolate.

^b ND, Not determined.

^c + and - denote the reported presence or absence, respectively, of cytochrome *d*.

protein each were used. Details are given elsewhere (29). For dot immunoblotting, samples of Zwittergent 3-12 solubilized membranes (50 µg of protein) were first mixed with SDS-PAGE sample buffer and then filtered through nitrocellulose labeled with ¹²⁵I-protein A and autoradiographed as described previously (30). Rocket immunoelectrophoresis was performed as before (31) with the Zwittergent-solubilized membranes (approximately 100 µg of protein).

RESULTS AND DISCUSSION

Cytochrome *d* complex. A number of bacteria other than *E. coli* have been reported to contain cytochrome *d* (previously called cytochrome *a*₂) based on spectroscopic criteria (20, 41). In addition to the strains listed in Table 1, other strains have also been reported to have cytochrome *d*, including *Haemophilus parainfluenzae* (15), *Achromobacter* strain D

(1), and *Pasteurella tularensis* (6). In most cases the presence of cytochrome *a*₁ in the membrane correlates with the presence of cytochrome *d* (20, 41). Very few studies have included biochemical characterization, although a study has been reported on the cytochrome *d* from *Photobacterium phosphoreum* (54). Often the amount of cytochrome *d* can be optimized by using selective growth conditions, usually limited-oxygen conditions. No attempt was made in the current work to optimize for cytochrome *d* production in all the bacterial strains which were examined. Figure 1 shows an SDS-PAGE immunoblot of membrane preparations with monoclonal antibodies directed against subunit I of the cytochrome *d* complex (32). Many of these strains contained a protein closely related to subunit I of the *E. coli* cytochrome *d* complex, which has previously been shown to be cytochrome *b*₅₅₈ (8). Most strains with cytochrome *d* by

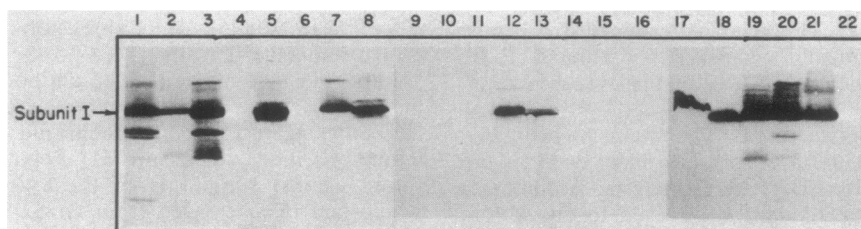


FIG. 1. SDS-PAGE immunoblotting of membrane preparations from various gram-negative bacteria with a monoclonal antibody preparation directed against subunit I of the cytochrome *d* terminal oxidase complex of *E. coli* K-12. Lanes: 1, *E. coli* K-12 (MR43L); 2, *Serratia marcescens*; 3, *Enterobacter aerogenes*; 4, *Pseudomonas putida*; 5, *Proteus vulgaris*; 6, *Pseudomonas fluorescens*; 7, *E. coli* W191-6; 8, *Proteus mirabilis*; 9, *Pseudomonas aeruginosa*; 10, *Rhodopseudomonas palustris*; 11, *Paracoccus denitrificans*; 12, *Photobacterium phosphoreum*; 13, *E. coli* SHSP19; 14, *Rhodospirillum rubrum*; 15, *Rhodospirillum fulvum*; 16, *Rhodopseudomonas sphaeroides*; 17, *Arthrobacter pyridinolis*; 18, *Acinetobacter* HO1N; 19, *Klebsiella pneumoniae*; 20, *Salmonella typhimurium*; 21, *Azotobacter vinelandii*; 22, *Vitreoscilla* sp.

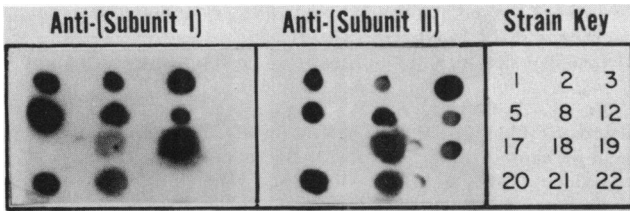


FIG. 2. Dot immunoblot of membrane preparations from various gram-negative bacteria with the monoclonal antibody directed against subunit I and a polyclonal antibody preparation directed against subunit II of the *E. coli* cytochrome *d* complex. The strain numbers given in the key correspond to the lane numbers given in the legend to Fig. 1.

spectroscopic criteria contained this subunit. It is reasonable to conclude that a *b*-type cytochrome is associated with cytochrome *d* in the other bacterial strains as it is in *E. coli*. The presence of the bands in Fig. 1 other than that identified as subunit I does not reflect a lack of specificity of the antibody, which was a monoclonal antibody. Bands below subunit I (e.g., lanes 1 and 3) are due to proteolysis and have been previously noted (32). Occasional smearing above subunit I (e.g., lanes 19 and 20) probably resulted from a tendency of this protein to aggregate, which has also been previously observed (37). Note that the sample from *Arthobacter pyridinolis* (Fig. 1, lane 17) ran anomalously, with most of the protein migrating as a single band. Blotting results were probably due to nonspecific trapping in this case. In all other cases, the membrane proteins were well resolved on the gel, as indicated by protein staining with Coomassie blue, and the blotting results were quite specific. The five strains reported to contain no cytochrome *d* by spectroscopic criteria (Table 1) also contained no cross-reactive material (Fig. 1).

Immunoblotting after SDS-PAGE with anti-subunit II of the cytochrome *d* complex showed clear evidence of subunit II in close relatives of *E. coli*, including *Enterobacter aerogenes*, *Klebsiella pneumoniae*, and *Salmonella typhimurium*. However, subunit II does not transfer efficiently from the polyacrylamide gel to the nitrocellulose, and this could result in false negative results. In order to avoid this step, dot immunoblotting was used. Fig. 2 shows that all

those strains which had a protein which cross-reacted to subunit I also had material cross-reactive to subunit II. *A. pyridinolis* did not appear to contain either subunit, at least not when grown under the conditions we used in these experiments. *Vitreoscilla* sp. was not reported to contain cytochrome *d*, and this is consistent with the data in Fig. 2; thus, *Vitreoscilla* sp. served as a negative control.

Immunoprecipitation studies were also performed with antibodies against the native cytochrome *d* complex from *E. coli*. Only in the case of *E. coli* was 100% of the cytochrome *d* precipitated by this antibody (not shown). In most other cases, very little cytochrome *d* was precipitated from a Zwittergent 3-12 solution after the addition of the antibody preparation. These findings suggest significant divergence between the *E. coli* enzyme and the cytochrome *d* present in the other strains.

Another experiment was performed to examine the membranes of *E. coli* strain SHSP19 (45). This strain does not synthesize heme in the absence of δ -amino levulinic acid. It has been reported that membranes prepared from this strain must contain cytochrome apoproteins because respiratory oxidase activity can be reconstituted by the addition of hematin and ATP (43). The immunoblotting results (Fig. 1) showed that the cytochrome *d* complex was present in the membrane, even though no heme was present.

Cytochrome *o* complex. Immunocross-reactivity data with cytochrome *o* are less complete than those obtained with cytochrome *d* because the antibody preparation against cytochrome *o* did not immunoblot well. Experiments were thus limited to rocket immunoelectrophoresis, which requires immunoprecipitation. As was seen with the anti-cytochrome *d*, it is likely that some strains may contain cross-reactive material which would be manifest in immunoblotting experiments but which will not be apparent by immunoprecipitation. The data (Fig. 3) show that cytochrome *o* similar to that found in *E. coli* is in close relatives, including *E. aerogenes*, *K. pneumoniae*, and *S. typhimurium*. In each of these cases the major immunoprecipitin arc stained for the presence of heme, confirming that the cross-reactive component is a cytochrome. The minor arcs which were observed in some cases did not contain heme. All other strains failed to show any heme-staining rocket immunoprecipitin arc, suggesting that there were no proteins strongly cross-reactive to cytochrome *o*. This would indicate that the

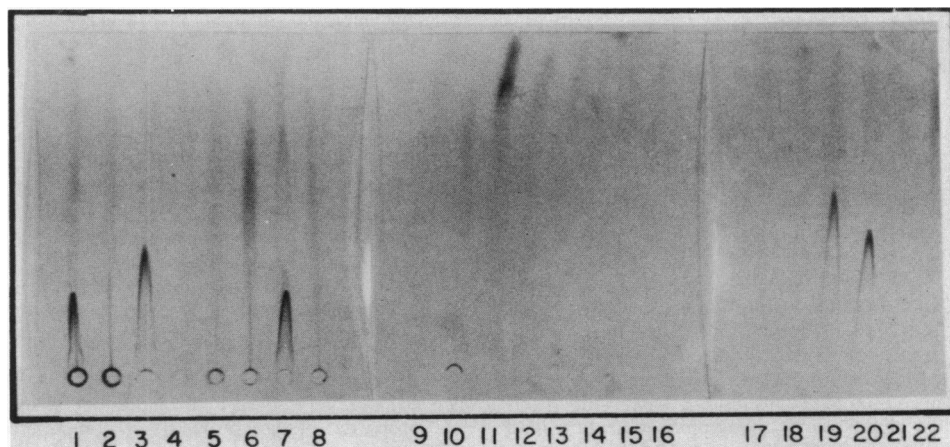


FIG. 3. Rocket immunoelectrophoresis of membrane preparations from various gram-negative bacteria with an antibody preparation which is monospecific against the *E. coli* cytochrome *o*. Lane numbers are as in legend to Fig. 1.

cytochrome *o* species reported in many of these organisms are not very closely related to the *E. coli* enzyme.

Note that most of the cytochrome *o* species which have been purified from organisms other than *E. coli* either function as cytochrome *c* oxidases or contain a *c*-type cytochrome as a tightly bound component. These include cytochrome *o* from *Rhodopseudomonas sphaeroides*, (50, 51), *R. capsulata* (16, 17), *R. palustris* (23), *P. aeruginosa* (29, 55), *A. vinelandii* (56, 57, 58), and *Methylophilus methylotrophus* (3). Hence, these may be quite distinct from the *E. coli* enzyme which neither contains a *c*-type cytochrome nor has cytochrome *c* oxidase activity (12, 29, 33). The well-characterized cytochrome *o* of *Vitreoscilla* sp. is a soluble enzyme of unknown function (52, 53) and is probably not closely related to the *E. coli* enzyme.

E. coli SHSP19 (heme deficient) was also examined by rocket immunoelectrophoresis for the presence of apo-cytochrome *o*. Surprisingly, no rocket was apparent, suggesting the absence of cytochrome *o* when the strain is not synthesizing heme. Others (43) have presented electrochemical evidence that apocytochrome *o* must be present in membranes of cells unable to synthesize heme. Further studies will be required to clarify this situation. Possibly detergent solubilization results in denaturation of the cytochrome in the absence of heme.

In summary, the data presented here clearly show that some of the close relatives of *E. coli* contain both of the terminal oxidases characterized in *E. coli* and that at least the cytochrome *d* complex is widely distributed among gram-negative bacteria. Furthermore, subunits I and II of the complex appear always to be present together, suggesting an association of the *b*-type cytochrome (i.e., subunit I) with cytochrome *d* in all the cases examined. Presumably, cytochrome *a*₁ is associated in a complex with cytochrome *d* in these other bacterial species as well. In many cases the published reduced-minus-oxidized difference spectra of membrane preparations from these bacteria show cytochrome *a*₁ along with cytochrome *d* (20, 41).

Finally, the immunoblotting data show that the apo-cytochrome *d* complex is synthesized and inserted into the membrane in the absence of heme biosynthesis. These findings complement the work of others and show that other apocytochromes are present in the membranes of heme-deficient *E. coli* (43).

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