# Synthesis and maturation of $\lambda$ receptor in *Escherichia coli* K-12: *In vivo* and *in vitro* expression of gene *lamB* under *lac* promoter control

(outer membrane protein/phage receptor/protein export/in vitro protein synthesis)

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ABSTRACT The  $\lambda$  receptor is an outer membrane protein from Escherichia coli K-12. lamB, its structural gene, is part of the maltose regulon. We have cloned this gene in a phage so that it is under the control of the *lac* promoter. The phage was devised in such a way that it can infect lamB mutants and that chromosomal lamB mutations can be transferred to it. In vivo, the  $\lambda$  receptor is expressed under *lac* promoter control and is exported normally to the outer membrane, independently of the expression of the other genes of the maltose regulon. In vitro, DNA of the phage allows efficient synthesis of the lamB product. The protein—or pre-λ-receptor—made in vitro contains an NH2-terminal sequence of about 25 amino acids not found in the  $\bar{\lambda}$  receptor. We have detected no inactivation of phage  $\lambda$  by the pre- $\lambda$ -receptor. Conversion of the pre- $\lambda$ -receptor to a form that has the apparent molecular weight of the mature  $\lambda$  receptor was achieved. A lamB mutation that blocks export in vivo also blocks conversion in vitro.

Bacteria export certain classes of proteins through their cytoplasmic membrane (1). To study the molecular mechanisms of this translocation process, which is similar to that of protein secretion in higher cells (2), it is desirable to have a system that allows both genetic and biochemical analyses. The *lamB* gene (3–5) in *Escherichia coli* K-12 is quite suitable for such a combined approach. The *lamB* product is an outer membrane protein (6) involved in the transport of maltose and maltodextrins (7). It serves as the cell surface receptor for phage  $\lambda$  and other phages (8) and can be assayed in cellular extracts by its ability to inactivate phage  $\lambda$  (6). The *lamB* gene is a part of the *malK-lamB* operon of the *malB* region of the *E. coli* chromosome (9). Extensive genetic and restriction site analyses of the region have been performed (10–13).

We wanted to develop a system in which the export of an outer membrane protein could be studied *in vitro*. This could allow us to test the current models of protein export, particularly the role of the signal sequence (14–17). To bypass the positive control factors that are required for *malB* expression (9–18), we cloned *lamB* under the control of the promoter for the *lac* operon on a transducing phage. The DNA of this phage programs the synthesis of the *lamB* product *in vitro* so efficiently that biochemical studies become possible.

## MATERIALS AND METHODS

Strains, Media, Bacteriological Techniques, and Chemicals. The bacterial strains used for S30 preparation were pop945:lacZW4680, rpsL, malQ6, malB $\Delta$ 7 ( $\lambda^+$ ) rpoB and pop997:lacZW4680, rpsL, malA $\Delta$ 108, malB $\Delta$ 7 ( $\lambda^+$ ) rpoB. The relevant markers of the other strains used are given in the text. The phage strains were  $\lambda$ YhÅ34 (Fig. 1),  $\lambda$ CI857h80t68dlacL8UV5 (a gift of Jack Greenblatt),  $\lambda$ vh<sup>+</sup>,  $\lambda$ vho (4), and  $\lambda$ CI57h434. Strain pop3257 or SE2060 [araD139,  $\Delta$ (lac)U169, strA, relA, thi, lamB S60] which is resistant to  $\lambda$  ( $\lambda$ r) and unable to grow on dextrins (Dex<sup>-</sup>) (S. Emr and T. Silhavy, personal communication) was used to generate, by homogenization, phage [ $\lambda$ YhÅ34 (1B8) S60].

Media and bacteriological techniques were as described (10). DNA hydrolysis with restriction enzymes, ligation, and transfection were as described (19).

Cell-Free Protein Synthesis. The system described by Zubay (20) was used without major modifications. The pressure on the French press was 1550 psi ( $10.7 \times 10^6$  pascals). Polyethylene glycol 6000 was added (1.5%, wt/vol) to the final mixture. Isopropyl  $\beta$ -thiogalactoside (IPTG), when present, was at 5 mM. For [<sup>35</sup>S]methionine labeling, the concentration of nonradioactive methionine was decreased to 0.45 mM and [<sup>35</sup>S]methionine was present at 0.04 mCi/ml (1 Ci =  $3.7 \times 10^{10}$  becquerels). Formyl[<sup>35</sup>S]Met-tRNA was synthesized according to Wendell and Stanley (21).

Assay of Pre- $\lambda$ -Receptor Activity. To assay the activity of the pre- $\lambda$ -receptor, the *in vitro* synthesis mixture was dialyzed overnight against the assay buffer. Samples (5  $\mu$ l) were then incubated at 37°C with about 300 plaque-forming units of  $\lambda$ ho in a total volume of 25  $\mu$ l (adjusted with buffer). At 30 min, 100  $\mu$ l of buffer containing strain CR63 (3) (about 2 × 10<sup>8</sup> cells) was added, and plating was done with top agar on tryptone plates. The experiments were performed with in vitro synthesis mixtures programmed with a DNA carrying lamB under lacL8UV5 promotor control, and the appropriate controls were run in parallel. By adding various concentrations of active receptor it was found that the assay was 1/10th as sensitive in presence of synthesis mixture-i.e., in the presence of synthesis mixture the K value was 1/10th that in the assay buffer. No inactivation of  $\lambda$ vho was detected. Because inactivation of 20% of the phage would have been detected, taking into account the decreased sensitivity of the assay, we can calculate that the Kvalue in the *in vitro* synthesis mixture  $(K_{pre})$  is less than 0.5.

Evaluation of the Amount and Relative Specific Activity of the Pre- $\lambda$ -Receptor. The methionine content per unit molecular weight is approximately the same in  $\beta$ -galactosidase [23 methionine residues,  $M_r \approx 116,000$  (22)] and pre- $\lambda$ -receptor [8–9 methionine residues,  $M_r \approx 56,000$  (ref. 23 and unpublished data)]. Syntheses were performed in parallel with DNA carrying lamB or lacZ. By comparing the radioactivity incor-

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Abbreviations:  $K_{\text{pre}}$ , K value in *in vitro* synthesis mixture;  $\lambda r$ ,  $\lambda$ -resistant;  $\lambda s$ ,  $\lambda$ -sensitive; IPTG, isopropyl  $\beta$ -thiogalactoside.

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porated in vitro in the protein bands we could estimate that the amount of pre- $\lambda$ -receptor protein was generally double that of  $\beta$ -galactosidase per unit volume of synthesis mixture. Because the amount of  $\beta$ -galactosidase calculated from the enzymatic activity was about 1.5  $\mu$ g/ml, the amount of pre- $\lambda$ -receptor protein was 3  $\mu$ g/ml. This is approximately the amount of  $\lambda$  receptor protein found in the extract of 2.5–5 × 10<sup>9</sup> cells (6) and corresponds to a K value of 200–400. Because  $K_{\rm pre}$  was found to be less than 0.5, this shows that the specific activity of the pre- $\lambda$ -receptor is no more than 1/800 to 1/400 of that of the  $\lambda$  receptor.

## RESULTS

Cloning and In Vivo Expression of Gene lamB Under lac Promoter Control. Phage  $\lambda$ Yh434 contains a single EcoRI site located early in the lacZ gene (19–25) (Fig. 1). Most of lacZ is deleted in this phage, but lacY is present and is expressed under lac promoter control. We assumed that insertion of the EcoRI fragment [lamB (EcoRI)] containing lamB but not its natural promoter into the EcoRI site of the phage would block lacY expression. We transformed a lacZ<sup>+</sup> lacY<sup>-</sup> strain (pop265) (10) with a ligated mixture of EcoRI-treated  $\lambda$ Yh434 DNA and lamB (EcoRI) fragment and plated the cells on MacConkey's lactose medium at 32°C. Of, 482 plaques, 450 became red within 2 days; 32 remained white, as expected if no lacY complementation occurred. Lysogens were purified from seven white plaques and their progeny phages were characterized.

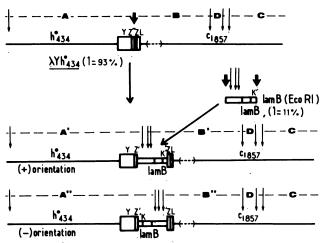


FIG. 1. Cloning of lamB under the control of the lac promoter on a phage vector. The phage vector  $\lambda$ Yh434 was constructed by in vivo recombination. The late functions give the host range of phage 434 and thus allow growth on  $\lambda$ -resistant strains. They originate from phage  $\lambda$ oh434. This hybrid between phage  $\lambda$  and phage 434 was constructed by Irène Mannheimer.  $\lambda oh 434$  is free of any EcoRI cuts.  $\lambda$ Yh434 is a recombinant between  $\lambda$ oh434 and  $\lambda$ Y (19). It carries the promoter mutations lacL8UV5. lamB (EcoRI) is a 5.5-kilobase restriction fragment limited by EcoRI sites. It contains the distal part of malK ( $\approx$ 1 kilobase), gene lamB ( $\approx$ 1.6 kilobases), and additional bacterial DNA ( $\approx$ 3 kilobases). It was purified by EcoRI hydrolysis of the DNA of phage  $\lambda ap malB21$  (25) and separation of the restriction fragments by ultracentrifugation on a sucrose gradient. On the lower part of the figure the two possible orientations for the insertion of the lamB (EcoRI) fragment in the vector have been represented. Expression under lac promoter control occurs only in the + orientation. l, DNA length in % of  $\lambda$  wild-type DNA; ---, HindIII deletion extending from 52.2% to 57.2% of  $\lambda$  (19);  $\downarrow$ , an EcoRI cut;  $\downarrow$ , a Bgl II cut. A, B, A', B', A", and B" correspond to restriction fragments obtained after EcoRI or Bgl II hydrolysis (or both) used to determine the orientation of the inserted lamB (EcoRI) fragment (see Fig. 2). Z is lacZ, Y is lacY, K is malK, and L is the promoter lacL8UV5. Relevant gene fragments have been designated by a letter with the symbol prime (') on the side of the missing piece.

If the lamB (EcoRI) fragment has been inserted, the phages should carry the distal part of malK. Indeed, strain pop1768 having a late mutation in malK (malK4) (9) yielded Mal<sup>+</sup> recombinants when lysogenized with any of the seven phages.

Insertion of the  $lam \vec{B}$  (EcoRI) fragment into phage  $\lambda Y h \vec{4}34$ can occur in either of two possible orientations (Fig. 1). Expression of lamB under lac promoter control should be possible only in the orientation such that the sense strand of *lamB* can be transcribed from the *lac* promoter (+ orientation). We thus expected that only the phages carrying lamB in the + orientation could convert  $\lambda$ -resistant ( $\lambda r$ ) strains to  $\lambda$ -sensitive ( $\lambda s$ ). Furthermore, we expected that expression of lamB would then be independent of malT(9) (the positive regulatory gene of the maltose system) but dependent on derepression of the lac promoter. lamB expression should also be cyclic AMP independent because of the L8UV5 mutations. This was tested by constructing lysogens of a lamB recA strain (pop6510), a recA strain carrying a partial deletion of malK and lamB (pop5208), a strain carrying a deletion in malT (pop101), and a strain carrying a cya deletion (CA8306). Three of the phages  $[\lambda Yh434(1B8), (1B10), and (1B19)]$  gave complementation to  $\lambda$ s for all of these  $\lambda$ r strains and should contain the *lam*B (*Eco*RI) fragment in the + orientation; four  $[\lambda Yh434(1B1)]$ , (1B3), (1B5), and (1B30)] did not, suggesting that they contained the fragment in the opposite orientation (-). This was confirmed by restriction endonuclease mapping of two + phages and one - phage (data not shown). Complementation occurred only in the presence of IPTG as expected with a lacI+ strain; in particular, lysogenic derivatives of the Dex<sup>-</sup> strain (pop6510) became Dex<sup>+</sup> only in presence of IPTG.

Table 1. Assay of  $\lambda$  receptor activity extracted from lysogens for phages carrying the *lamB* (*EcoRI*) fragment

<i>mal</i> T allele*	lamB orientation†	Inducer <sup>‡</sup>	λ rec. act. <sup>§</sup>	β-Gal- ase act.¶
+		None	300	
+		Mal	450	
+		IPTG	320	13,110
-		IPTG	<0.2	7,420
_	-	None	2.6	12
-	-	IPTG	2.8	6,750
-	+	None	2.1	11
_	+	IPTG	1630	5,700
-	+	None	2.7	10
-	+	IPTG	1910	6,500

\* The malT<sup>+</sup> strain is HfrG<sub>6</sub>. The malT strain is HfrG<sub>6</sub> malA $\Delta$ 157; it carries a deletion in malT.

- <sup>†</sup> The control strains (first four lines) contained no phage. The phage carrying the *lamB* (*EcoRI*) fragment in the – orientation was  $\lambda$ Yh434 (1B5); the phages carrying the + orientation were  $\lambda$ Yh434 (1B8) (lines 7 and 8) and  $\lambda$ Yh434 (1B10) (lines 9 and 10).  $\lambda$ Yh434 (1B10) carries mutation *mal*K5 (9). No polar effect of this ochre mutation has been detected on *lamB* expression.
- <sup>‡</sup> Mal, maltose.
- <sup>§</sup> Cell extracts were prepared as described (6) and were assayed for  $\lambda$  receptor activity by measuring the initial kinetics of  $\lambda$ vho inactivation (6). The assay buffer contained 10 mM Tris-HCl at pH 7.5 and 2 mM MgCl<sub>2</sub>. The kinetics of inactivation follow the relationship log  $(\phi/\phi_0) = -K_d \cdot t, \phi_0$  and  $\phi$  are plaque-forming units/ml at times 0 and t (min), respectively.  $K_d$  is the inactivation constant at dilution d of the extract. The activity is measured by the slope of the curve  $K_d = K \times d$ . K is directly proportional to the concentration of active receptor in the extract and hence is a measure of the amount of active receptor (24) which is expressed here as K/mg of protein. The relative error is 20%.
- ¶  $\beta$ -Galactosidase activity, assayed for comparison, is encoded by the chromosomal *lac*Z gene which is under the control of the wild-type *lac* promoter. The activity is expressed in units/mg of protein. The induction ratio for  $\beta$ -galactosidase here is of the order of 550.

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 $\lambda$  receptor activity was assayed in extracts of  $lacI^+$  strains carrying a malT deletion and lysogenic for the phages having inserted the lamB (EcoRI) fragment in each orientation (Table 1). As expected, only in the + orientation was the  $\lambda$  receptor activity inducible by IPTG. The *in vivo* induction ratio was about 700–800. The fully induced level was somewhat higher (about 3 times) than the level in the malT<sup>+</sup> strain (HfrG<sub>6</sub>) induced by maltose. Surprisingly, the level of activity in the absence of IPTG was the same in the + and - orientations.

In Vitro Synthesis of the lamB Product. DNA from a phage carrying lamB inserted in the + orientation was used to program in vitro protein synthesis. The S30 extract was prepared from a lacI<sup>+</sup>  $\Delta$ lacZ  $\Delta$ lamB strain. A major protein band of apparent  $M_r$  56,000 could be detected (Fig. 2a, lanes 5 and 6; Fig. 2b, lanes 1-4; and Fig. 2c, lane 4). This band was not encoded by the vector phage DNA (Fig. 2a, lane 4). The intensity of the band was decreased in the absence of IPTG (Fig. 2a, lanes 7, 8, and 9) so that this protein is synthesized under lac promoter control. Althrough its apparent  $M_r$  is about 2000–3000 greater than that of the  $\lambda$  receptor (Fig. 2b, lanes 5, 6, and 7) we conclude by the following criteria that it is the product of gene lamB. (i) The protein made in vitro was precipitated with anti- $\lambda$ -receptor serum (Fig. 2a, lanes 3, 8, and 9; Fig. 2b, lane 6). (ii) When the DNA carried a small deletion, lamB S60  $[\lambda Yh434 (1B8 S60)]$ , which is internal to lamB and does not change the coding frame (S. Emr and T. Silhavy, personal communication), the protein made in vitro was smaller than that corresponding to lamB<sup>+</sup> (Fig. 2c, lanes 1 and 2; Fig. 3b). (iii) Partial proteolysis of the protein synthesized in vitro and native  $\lambda$  receptor yielded peptides common to both proteins (Fig. 3a). However, three peptides (B', C', and D'), derived from the  $\lambda$  receptor, were absent from the digest of protein made in vitro. They were replaced by peptides B, C, and D, each of which exhibited the same increase in apparent  $M_r$  over its counterpart in the  $\lambda$  receptor. This increase is identical to that found between the  $\lambda$  receptor and the protein made in vitro. All this is consistent with the presence of the same extra sequence of amino acids in the protein made in vitro and in peptides B, C, and D.

Two lines of evidence demonstrate that the extra sequence present in the protein synthesized *in vitro* is located at the NH<sub>2</sub> terminus. (*i*) The difference in apparent  $M_r$  between the *lamB*<sup>+</sup> and *lamB* S60 proteins made *in vitro* is also found between the peptides B<sub>1</sub>, C<sub>1</sub>, and D<sub>1</sub> derived from the *lamB* S60 protein and the peptides B, C and D, respectively (Fig. 3b). Thus, these peptides contain the amino acid sequence deleted in the *lamB* S60 product. Because the S60 deletion maps in the earliest known segment of the *lamB* gene (S. Emr and T. Silhavy,

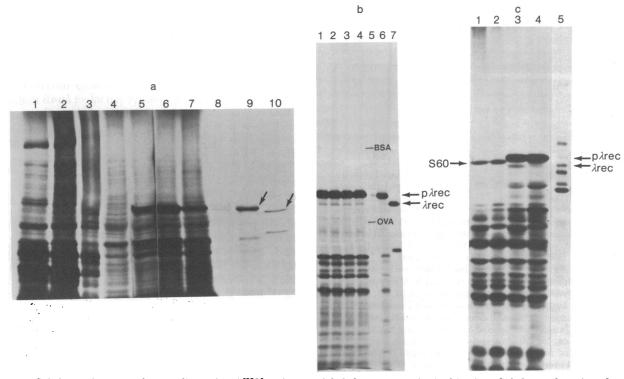


FIG. 2. Gel electrophoresis and autoradiography of [35S] methionine-labeled protein synthesized in vitro. Gel electrophoresis and autoradiography were performed as described (26, 27). Nonradioactive protein markers were located by staining with 1% alkylbenzyldimethylammoniumchloride (Merck). When no immunoprecipitation was performed, each lane contained 10  $\mu$ l of in vitro synthesis mixture. Otherwise, the amount of washed immunoprecipitate per lane corresponded to 50  $\mu$ l of an *in vivo* extract or an *in vitro* synthesis mixture treated with 5  $\mu$ l of anti- $\lambda$ -receptor antiserum and with 10 mg of protein A-Sepharose beads. The *in vivo* extracts (6) were obtained from the  $\lambda$ s strain pop3 labeled with  ${}^{35}SO_4$ . p\rec, lamB product made in vitro; \rec, \receptor made in vivo. (a) A 130  $\times$  75  $\times$  1.5 mm NaDodSO<sub>4</sub>/10% polyacrylamide gel run for 3.5 hr at 30 mA. Lanes: 1, 2, and 3,  $\lambda$ Cl857h80t68d*lac*L8UV5 DNA; 2, with IPTG; 3, with IPTG and  $\lambda$  receptor antiserum; 4,  $\lambda$ Yh434 DNA with IPTG; 5,  $\lambda$ Yh434 (1B19) DNA with IPTG; 6, 7, 8, and 9,  $\lambda$ Yh434 (1B10) DNA; 6, with IPTG; 8, with  $\lambda$  receptor antiserum; 9, with IPTG and  $\lambda$  receptor antiserum; 10, immunoprecipitate of  $\lambda$  receptor. Because the S30 contains some repressor of the *lac* operon, induction of genes under lac control can be seen in vitro. However, the induction ratio is much smaller than in vivo. Compare for example, lanes 1 and 2 for  $\beta$ -galactosidase and lanes 6 and 7 or 8 and 9 for lamB product. (b) a 130 × 300 × 1.5 mm NaDodSO<sub>4</sub>/10% polyacrylamide gel run for 20 hr at 150 V. Lanes: 1–4,  $\lambda$ Yh434 (1B10) DNA with IPTG; 5, position of nonradioactive markers used for  $M_r$  determination (28) (OVA,  $M_r \approx$ 45,000; BSA,  $M_r \approx 69\,000$ ); 6,  $\lambda$ YhÅ34 (1B10) DNA with IPTG and  $\lambda$  receptor antiserum; 7, immune precipitate of  $\lambda$  receptor. (c) A 130  $\times$  200 × 1.5 mm NaDodSO4/10% polyacrylamide gel run for 18 hr at 140 V. Lanes: 1 and 2, λYhÅ34 (1B8 S60) DNA; 3 and 4, Sam 7 hλ derivative of  $\lambda$ Yh434 (1B8); 5, nonprecipitated in vivo extract. Lanes 1 and 3 had inverted membrane vesicles (19  $\mu$ l/100  $\mu$ l of in vitro synthesis). Longer times of exposure of the gel confirmed that no band corresponding to mature  $\lambda$  receptor was produced in lane 1.

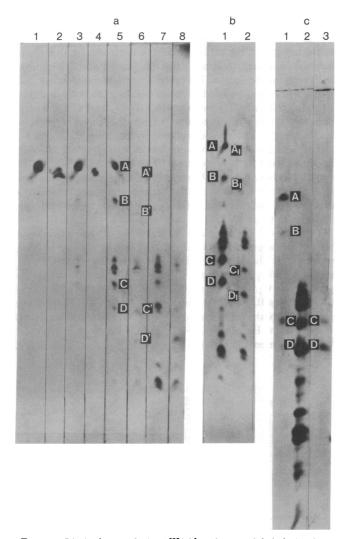


FIG. 3. Limited proteolysis of [35SO] methionine-labeled proteins. After nonlabeled markers were identified, radioactive bands were excised from gels similar to that of Fig. 3a and subjected to limited proteolysis with Staphylococcus aureus V8 protease and electrophoresis on NaDodSO<sub>4</sub>/15% polyacrylamide  $130 \times 300 \times 1.5$  mm gels (29). A constant difference of apparent  $M_r$  can be measured between the following couples of peptides: A/A', B/B', C/C', and D/D',  $\approx$ 2000-3000; A/A<sub>1</sub>, B/B<sub>1</sub>, C/C<sub>1</sub>, and D/D<sub>1</sub>, ≈1200-1500. (a) Lanes: 1, 3, 5, and 7, in vitro made [35S]methionine-labeled lamB product; 2, 4, 6, and 8, immunoprecipitate of in vivo made  $\lambda$  receptor. The amounts of S. aureus V8 protease per lane were 0, 0.01, 0.1, and 1 µg for lanes 1 and 2, lanes 3 and 4, lanes 5 and 6, and lanes 7 and 8. (b) Lanes: 1, in vitro made [35S]methionine-labeled lamB+ product; 2, in vitro made [35S] methionine-labeled lamB S60 product. V8 protease per lane was 0.1 µg. (c) Lanes: 1 and 3, in vitro made formyl[35S]methionine-labeled lamB product; 2, *in vitro* made [<sup>35</sup>S]methion-ine-labeled lamB product. V8 protease per lane was 0.01  $\mu$ g in lane 1 and 0.1  $\mu$ g in lanes 2 and 3.

personal communication), this indicates that peptides B, C, and D must be at or near the NH<sub>2</sub>-terminal fragment of the pre- $\lambda$ -receptor. (ii) In vitro synthesis was performed in the presence of formyl[<sup>35</sup>S]methionine-tRNA which labels only the NH<sub>2</sub> terminus of the protein. Partial proteolysis showed that the label is only in the intact protein and in peptides B, C, and D (Fig. 3c).

We assume that the *in vitro lamB* product is the precursor of the  $\lambda$  receptor predicted by the signal hypothesis (1, 2) and designate it the "pre- $\lambda$ -receptor."

The Pre- $\lambda$ -Receptor: Conversion and Lack of Activity. Addition of inverted membrane vesicles (15) to the synthesis mixture programmed with a  $lamB^+$  template resulted in the appearance of a protein that had the apparent  $M_r$  of the  $\lambda$  receptor (Fig. 2c, lane 3). When the lamB template carried deletion S60, this protein was not produced (Fig. 2c, lanes 1 and 2). This shows that appearance of this protein is due to the conversion of the *in vitro* made  $lamB^+$  product—i.e., the pre- $\lambda$ -receptor. In addition, it suggests strongly that the conversion corresponds to the processing of the signal sequence of the pre- $\lambda$ -receptor (see *Discussion*). Conversion by vesicles was not seen when the vesicles were added after the synthesis is completed (data not shown).

When the ability of the pre- $\lambda$ -receptor to inactivate phage  $\lambda$ vho was assayed, no activity was detected. We can determine that the specific activity of the pre- $\lambda$ -receptor is no more than 1/800th to 1/400th that of the  $\lambda$  receptor. We did not detect any activity of the conversion product under the conditions of the assay.

#### DISCUSSION

In Vivo Expression of lamB Under lac Promoter Control. Insertion of the lamB (EcoRI) fragment in phage  $\lambda$ Yh434 in the + orientation results in the synthesis of the lamB protein under lacL8UV5 promoter control.  $\lambda$  receptor activity is inducible by IPTG in a lacI<sup>+</sup> background. It is expressed independently of the maltose regulon and of the cyclic AMP-dependent control systems. The fact that  $\lambda$  receptor activity can be obtained in the total absence of the malT product proves that, according to the present criteria, neither the malT product nor any other known gene of the maltose regulon is required for translation, positioning, and maturation of the  $\lambda$  receptor.

Because the phage vector has a host range different from  $\lambda$  (h434), the *lam*B-carrying phages can infect *lam*B mutants. It is thus convenient to transfer mutations from a chromosomal *lam*B gene to the *lam*B copy of the phage by looking for homogenotes. This can be used in particular to compare the effects of a mutation *in vivo* and *in vitro*.

In Vitro Expression of lamB Under lac Promoter Control. In vitro synthesis programmed with DNA carrying lamB under lac promoter control yields a major protein. We have shown that this protein is the product of the lamB gene: (i) it is specifically precipitated with anti- $\lambda$  receptor antiserum; (ii) limited proteolysis gives a pattern that is similar to that of the *in vivo* made  $\lambda$  receptor; and (*iii*) it is made with a smaller apparent  $M_r$  when lamB carries the internal deletion S60. The *in vitro* made lamB product has an apparent  $M_r$  that is about 2000– 3000 higher than that of the *in vivo* made  $\lambda$  receptor, and we assume that it is the pre- $\lambda$ -receptor (1-2). The existence of a precursor to the  $\lambda$  receptor had already been found by using various systems (5-30). We show here that the pre- $\lambda$ -receptor carries an extra NH<sub>2</sub>-terminal sequence that suffices to account for its higher apparent  $M_r$ .

This was confirmed by determination of the partial amino acid sequence of the NH<sub>2</sub>-terminal part of the pre- $\lambda$ -receptor: it corresponds to the DNA sequence of the early part of gene *lamB* (unpublished data).

We have observed conversion of the pre- $\lambda$ -receptor by inverted vesicles to a form that has the apparent  $M_r$  of the  $\lambda$  receptor. This conversion was found to be cotranslational (15).

It is not established at present whether the conversion product obtained *in vitro* corresponds to the processing occurring *in vivo*. However, it is interesting that this conversion is not seen when the *lamB* template carries deletion S60 which prevents export *in vivo* (S. Emr and T. Silhavy, personal communication). This mutation as well as two other mutations (5) that are known to prevent *in vivo* processing are located in the signal sequence (unpublished data). One possible explanation is that all these signal sequence mutations block a very early step in export so that processing, which is subsequent, cannot take place. This situation may also be the one prevailing in the *in vitro* system.

No ability of the pre- $\lambda$ -receptor to inactivate phage  $\lambda$ vho was found by us. We found also no phage inactivation after synthesis in the presence of inverted vesicles—i.e., when conversion occurred. However, this does not allow us to conclude that the converted product is not active because (*i*) the converted product might be unable to interact with the phage in the conditions of the assay [for example, it could be sequestered in the vesicles (15)], and (*ii*) the yield of conversion being low, the total activity of the converted product might be near the limit of detection of the assay. At any rate, it is not known at present whether processing of the signal sequence is enough to yield  $\lambda$ receptor activity. Further maturation steps, or cofactors, might be required.

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