

## DEFECTIVE BACTERIOPHAGE LAMBDA CHROMOSOME, POTENTIAL VECTOR FOR DNA FRAGMENTS OBTAINED AFTER CLEAVAGE BY *BACILLUS AMYLOLIQUEFACIENS* ENDONUCLEASE (BAM I)

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### 1. Introduction

There are two essential steps involved in the multiplication of a foreign DNA fragment in bacteria. First, the covalent insertion of this fragment into a vector molecule, capable of self replication in *E. coli*, is accomplished through recombination in vitro. Second, the hybrid DNA molecule is introduced into the bacteria by transformation or transfection. The key enzymatic step in the in vitro recombination procedure is provided by a restriction endonuclease that creates cohesive ends that can be covalently rejoined with DNA ligase. At the present time, two types of vectors are used: plasmids such as psc 101 or colEI [1,2] and derivatives of bacteriophage lambda [3-5]. These phages have lost by mutation the Eco RI cleavage sites located in the essential region of the genome. A main advantage of the phage system, compared to the plasmid system, is that its propagation can be well controlled and therefore accidental dissemination can be more easily prevented. This is important because some genetic engineering experiments could be potentially dangerous, and should therefore be very carefully controlled. Though the derivatives of phage lambda presently used as vectors are safe enough for most of the insertion experiments, it could be important to construct vectors which present higher security criteria.

In this work, we present the cleavage sites localization of *Bacillus amyloliquefaciens* restriction endonuclease (Bam I) [6], in the bacteriophage lambda genome and we propose a derivative of phage lambda which could after having incorporated the foreign

DNA fragment, lose the genes coding for the tail proteins and therefore become defective. The multiplication of such a phage would depend on the presence of a helper phage and could therefore be easily controlled.

### 2. Materials and methods

The localization of the targets for Bam I in the phage lambda genome was determined by the following method. DNA fragments obtained after double hydrolysis with Eco RI and Bam I of DNA from different derivatives of phage lambda having lost some Eco RI cleavage sites by mutation [4] were analysed by polyacrylamide gel electrophoresis. The bacteriophage strains used in this work are listed in fig.1. The mol. wt of the DNA fragments obtained after double hydrolysis were determined according to the reference curve constructed with Eco RI hydrolysates of lambda phage and adenovirus 2 DNA (fig.2).

### 3. Results and discussion

The j fragment is present in  $\lambda$ ,  $\lambda$ plac-I-2 and  $\lambda$ plac-I-2-3 hydrolysed by Eco RI and Bam I, and absent in  $\lambda$  hydrolysed by Eco RI or Bam I (fig.3). The only Eco RI cleavage site present in the three phages is sRI $\lambda$ 2. The j fragment is then located between sRI $\lambda$ 2 and a Bam I cleavage site. Moreover, the Eco RI - C fragment is absent in the double hydrolysates. There-

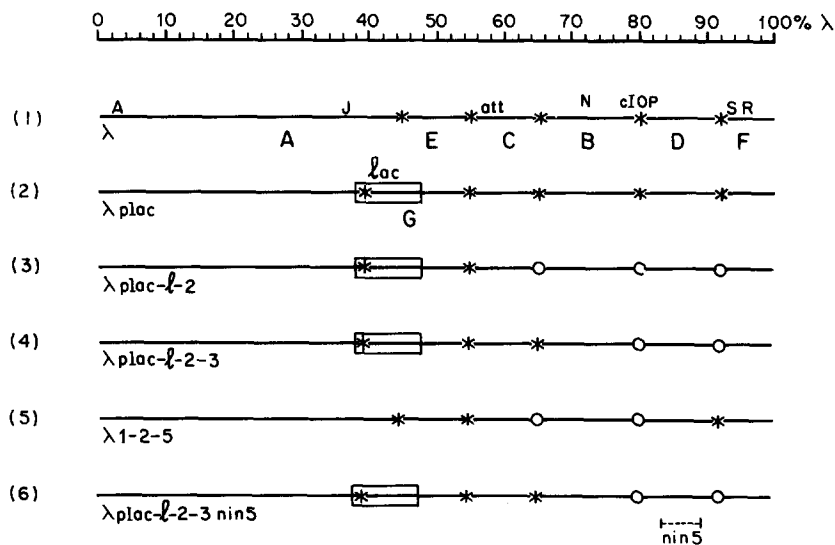
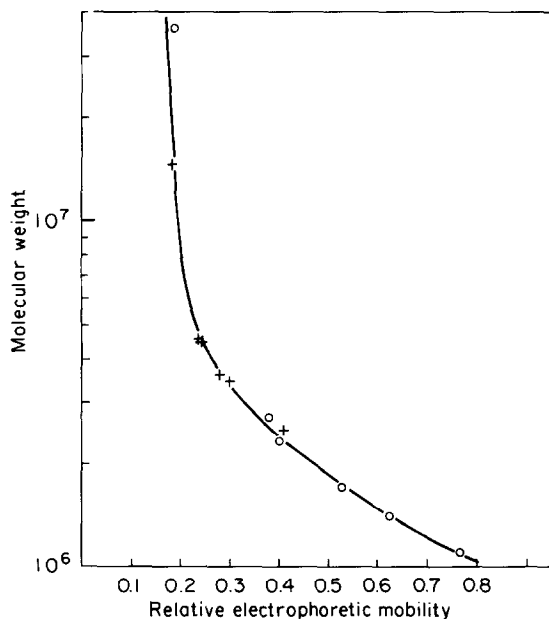


Fig.1.  $\lambda$  derivatives used to locate the BamI endonuclease cleavage sites. The bacteriophage strains are (1)  $\lambda$ CI857S7 referred to as  $\lambda$ . (2)  $\lambda$ plac5CI857S7 referred to as  $\lambda$ plac. (3)  $\lambda$ plac5CI857sRI $\lambda_3^0$  sRI $\lambda_4^0$  sRI $\lambda_5^0$ , referred to as  $\lambda$ plac-l-2 [4]. sRI $\lambda$  refers to the EcoRI cleavage sites in  $\lambda$ , numbered 1 to 5 from the left to the right of the genome. The symbol sRI $\lambda^0$  indicates the loss, by mutation, of an EcoRI site. The letter in  $\lambda$ plac-l-2 corresponds to the EcoRI cleavage site in the lac operon. The number 2 corresponds to the presence of the 2nd EcoRI cleavage site. (4)  $\lambda$ plac5CI857 sRI $\lambda_4^0$  sRI $\lambda_5^0$  referred to as  $\lambda$ plac-l-2-3. (5)  $\lambda$ CI857 sRI $\lambda_3^0$  sRI $\lambda_4^0$ , referred to as  $\lambda$ 1-2-5. This phage is a double recombinant of  $\lambda$ plac-l-2 and  $\lambda$ imm434 obtained after infecting the bacterial strain AR1002 ( $\lambda$  imm434) with  $\lambda$ plac-l-2. (6)  $\lambda$ plac5CI857 sRI $\lambda_4^0$  sRI $\lambda_5^0$  nin5, referred to as  $\lambda$ plac-l-2-3 nin5. This phage was obtained by genetic cross between the  $\lambda$ plac5 and the  $\lambda$ gt $\lambda$ ABC [5] which was a generous gift from R. Davis. The letters A to G correspond to the EcoRI fragments. \* refers to the EcoRI cleavage sites. (o) Refers to the loss by mutation of the EcoRI cleavage sites.



fore the j fragment is located in the C fragment. The mol. wt of the j fragment ( $1.3 \times 10^6$ ) permits the location of this BamI cleavage site at 4.2% on the right of sRI $\lambda_2$ . This site referred to as sBamI $\lambda_3$  is thus located at 58.8% (fig.4).

The g fragment is present in  $\lambda$ , and  $\lambda$ 1-2-5 hydrolysed by EcoRI and BamI, and absent in  $\lambda$ plac-l-2

Fig.2. Reference curve used to determine the mol. wts of  $\lambda$  derivative fragments obtained after double hydrolysis with EcoRI and BamI endonucleases. This curve was constructed according to the EcoRI  $\lambda$  fragments (+) and the EcoRI adenovirus 2 fragments (o). The adenovirus 2 was a gift from P. Boulanger. The reference molecular weight values of the EcoRI  $\lambda$  fragments are given according to B. Allet et al. [9] and the reference molecular weight values of the EcoRI adenovirus 2 fragments are given according to Petterson et al. [10]. The DNA fragments were analysed by electrophoresis in a polyacrylamide gel slab (gradient of 2.5% to 7.5% acrylamide concentration).

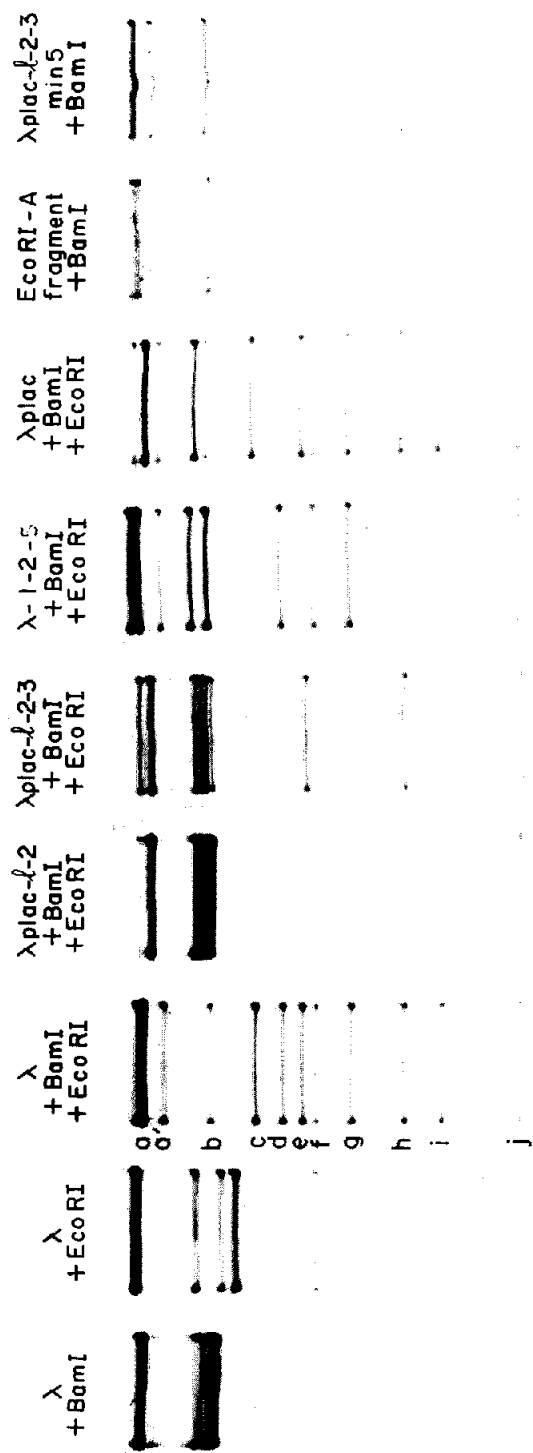


Fig. 3. Electrophoretic separation of DNA fragments from  $\lambda$  and various derivatives obtained after hydrolysis by EcoRI, BamI, and EcoRI + BamI. The location of EcoRI cleavage sites on the chromosome of  $\lambda$  and the various derivatives analysed are shown in fig. 1. The bacteriophage propagation, DNA extraction and hydrolysis by EcoRI were performed as previously described [11]. The EcoRI - A fragment from  $\lambda$  plac5 (fragment 0 to 40%) was prepared by cesium sulfate + silver nitrate isopycnic centrifugation as previously described [12]. (The EcoRI enzyme was a generous gift from R. Roberts. DNA hydrolysis by BamI and the double hydrolysis by EcoRI + BamI were performed for 12 hrs at 37°C in a 6.6 mM Tris-HCl, pH 7.5, 6.6 mM MgCl<sub>2</sub>, 6.6 mM mercaptoethanol, 50 mM NaCl buffer. The analytical polyacrylamide slab gel electrophoresis (2.5% to 7.5% concentration gradient) was performed as previously described [4].

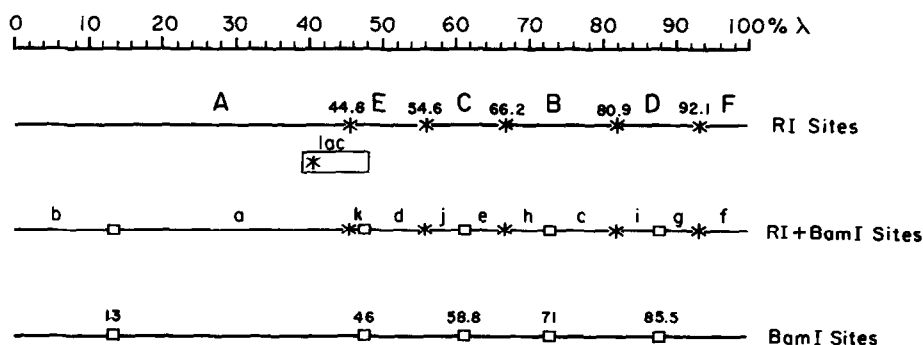


Fig.4. Location of the EcoRI and BamI cleavage sites on the chromosome of  $\lambda$ . The capital letters A to F correspond to the EcoRI fragments of  $\lambda$ . Small letters a to k correspond to the EcoRI + BamI fragments of  $\lambda$ . (\*) EcoRI cleavage sites. ( $\square$ ) BamI cleavage sites. The location of the EcoRI cleavage sites are given according to B. Allet et al. [9].

and  $\lambda$ plac-*l*-2-3 hydrolysed by EcoRI and BamI (fig.3). The only EcoRI cleavage site present in  $\lambda$ -1-2-5 and absent in plac-*l*-2, and plac-*l*-2-3 is sRI $\lambda$ 5. The g fragment is then located between sRI $\lambda$ 5 and a BamI cleavage site. Moreover the EcoRI-D fragment is absent in the double hydrolysis, whereas the EcoRI-F fragment is present. Therefore the g fragment is located in the EcoRI-D fragment. The mol. wt of the g fragment ( $2 \times 10^6$ ) permits the location of this BamI cleavage site at 6.45% on the left of sRI $\lambda$ 5. This site referred to as sBamI $\lambda$ 5 is thus located at 85.5% (fig.4).

The c and i fragments are present in  $\lambda$ plac hydrolysed by EcoRI and BamI and absent in  $\lambda$ plac-*l*-2,  $\lambda$ plac-*l*-2-3 and  $\lambda$ -1-2-5 hydrolysed by EcoRI and BamI (fig.3). The only EcoRI cleavage site present in  $\lambda$ plac and absent in  $\lambda$ plac-*l*-2,  $\lambda$ plac-*l*-2-3 and  $\lambda$ -1-2-5 is sRI $\lambda$ 4. The c and i fragments are therefore located between sRI $\lambda$ 4 and a BamI cleavage site. The comparison of the mol. wt of the c fragment ( $3.15 \times 10^6$ ) and the EcoRI D fragment ( $3.2 \times 10^6$ ) shows that the c fragment cannot be located in the D fragment. Therefore the c fragment is located in the EcoRI-B fragment, and the i fragment (mol wt =  $1.6 \times 10^6$ ) is located in the D fragment. The molecular weight of the c fragment ( $3.15 \times 10^6$ ) permits the location of BamI cleavage site at 10% on the left of sRI $\lambda$ 4. This site referred to as BamI $\lambda$ 4 is thus located at 71% (fig.4).

The d and k fragments are present in  $\lambda$  hydrolysed by EcoRI and BamI and absent in  $\lambda$ plac hydrolysed by EcoRI and BamI (fig.3). Therefore the *lac* insertion suppresses a BamI cleavage site and

this site is located in the EcoRI-E fragment. The *lac* insertion being located between 39.6% and 48.1% [7], and considering the molecular weights of the d ( $2.7 \times 10^6$ ) and the k ( $0.4 \times 10^6$ ) fragments, the d fragment must be located between the BamI cleavage site (of the E fragment) and sRI $\lambda$ 2 and the k fragment must be located between the BamI cleavage site (of the E fragment) and sRI $\lambda$ 1. The molecular weight of the d fragment ( $2.7 \times 10^6$ ) permits the location of the BamI cleavage site at 8.7% on the left of sRI $\lambda$ 2. This site referred to as sBamI $\lambda$ 2 is thus located at 46% (fig.4).

The b fragment is present in  $\lambda$ ,  $\lambda$ plac hydrolysed by EcoRI and BamI and the EcoRI-A fragment purified by cesium sulfate + silver nitrate isopicnic centrifugation [10] and hydrolysed by BamI (fig.3). The b fragment is then located in the A fragment. The sRI $\lambda$ 1 being absent in  $\lambda$ plac, the b fragment is then located between the left end of  $\lambda$  and a BamI cleavage site. The mol. wt of the b fragment ( $4 \times 10^6$ ) permits the location of the BamI cleavage site at 13% in the right of the left end of lambda. This site is referred to as sBamI $\lambda$ 1. (fig.4).

Therefore, there are 5 BamI cleavage sites in the phage lambda genome. Three sites, BamI $\lambda$ 2, BamI $\lambda$ 3 and BamI $\lambda$ 4 are located in the non essential region of the genome (fig.4). One site, BamI $\lambda$ 1 is located in a head protein gene, which is probably the E gene. One site BamI $\lambda$ 5 is located between the replication and the lysis gene (fig.4). In order to use the phage lambda as a vector for BamI DNA fragments the cleavage site BamI $\lambda$ 5 must be deleted. This can be done by



Fig.5. Possible model for a hybrid molecule constructed with the lambda vector BamI and a foreign DNA fragment. (—) lambda DNA, (■) foreign DNA fragment.

introducing the *nin5* deletion into the lambda genome. Effectively the two fragments which are located between *sBamI*λ4 and *sBamI*λ5, and between *sBamI*λ5 and the right end of lambda are absent in *λplac-I-2-3 nin5* (fig.3). Consequently the *λplac-I-2-3 nin5* DNA fragment located between *sBamI*λ4 and the right end of lambda contains all the early genes essential in vegetative growth (N, O and P), the control gene of late functions (Q) and the lysis genes (S and R). BamI restriction enzyme recognizes 6 base pairs and creates cohesive ends (R. Roberts personal communication). Consequently, a hybrid molecule could be constructed in vitro, by introducing a foreign DNA fragment between *sBamI*λ1 and *sBamI*λ4 (fig.5). The structure of such a phage would be comparable to a phage *λgalbio* [8]. This hybrid molecule could be replicated but not be encapsulated in the absence of a helper phage. The propagation of this defective phage could be controlled and its accidental dissemination easily prevented. Moreover such a vector molecule would be able to insert BamI DNA fragments of mol. wts up to  $18 \times 10^6$ . This could not be possible with the lambda vectors constructed for EcoRI DNA fragments.

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