Cloning and Amplified Expression in *Streptomyces lividans* of the Gene Encoding the Extracellular β-Lactamase from *Streptomyces cacaoi*

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A 19 kb SphI DNA fragment containing the gene for the extracellular active-site serine β-lactamase of *Streptomyces cacaoi* KCC-SO352 was cloned in *Streptomyces lividans* TK24 using the high-copy-number plasmid pIJ702 as vector. A 30-fold higher yield of β-lactamase was obtained from *S. lividans* strain ML1, carrying the recombinant plasmid pDML51, than from *S. cacaoi* grown under optimal production conditions. In all respects (molecular mass, isoelectric point, kinetics of inhibition by β-lactamase) the overproduced *S. lividans* ML1 β-lactamase was identical to the original *S. cacaoi* enzyme. A considerable reduction of β-lactamase production was caused by elimination of a 12.8 kb portion of the 19 kb DNA fragment by cleavage at an internal SphI site located more than 3 kb upstream of the β-lactamase structural gene. The β-lactamase gene was located within a 1.8 NcoI-BclI fragment but when this fragment was cloned in *S. lividans* pIJ702, the resulting strain produced hardly any more β-lactamase than the original *S. cacaoi*.

INTRODUCTION

Active-site serine β-lactamases (EC 3.5.2.6) and active-site serine DD-peptidases (EC number not attributed) react with β-lactam antibiotics by an acyl enzyme mechanism that involves formation of a serine ester-linked penicilloyl (cephalosporoyl, etc.) enzyme derivative. In most cases, the acyl enzyme formed with the β-lactamases is very short-lived so that the reaction flux rapidly proceeds to product. With the DD-peptidases, the acyl enzyme is very long-lived so that the DD-peptidases become immobilized in an inactive form for a long time; they therefore behave as penicillin-binding proteins (Ghuyser *et al.*, 1984). Questions about the structures and functions of these two classes of β-lactam antibiotic-recognizing enzymes are being investigated by using a battery of techniques such as X-ray crystallography, kinetic and spectroscopic studies, amino acid sequencing, gene analysis and site-directed mutagenesis. The extracellular active-site β-lactamases produced by *Bacillus licheniformis*, *Bacillus cereus* and *Streptomyces albus* G, and the extracellular active-site serine DD-peptidase of *Streptomyces* R61, are unrelated in their primary structures but, nevertheless, show striking similarities in the spatial disposition of their secondary structure elements (α-helices and β-strands), suggesting a close evolutionary relationship (Kelly *et al.*, 1986; Samraoui *et al.*, 1986; O. Dideberg and co-workers, unpublished results). The *Streptomyces* R61 DD-peptidase gene was cloned and amplified in *Streptomyces lividans* on plasmid pIJ702 (Duez *et al.*, 1987). In parallel to this, the same host–vector system was used to study *Streptomyces* β-lactamase genes. Following previous reports on the β-lactamase gene of *Streptomyces albus* G (Dehottay *et al.*, 1986, 1987) this paper...
describes the cloning of the gene that codes for the \( \beta \)-lactamase of *Streptomyces cacaoi*. The biochemical and kinetic properties of this \( \beta \)-lactamase were described by Ogawara (1975), Ogawara *et al.* (1981) and Lenzini & Frère (1985).

**METHODS**

*Streptomyces* strains and plasmids. *Streptomyces lividans* TK24 (str-6) (Hopwood *et al.*, 1983), *Streptomyces coelicolor* A3(2) and *Streptomyces* plasmid vector pIJ702 (Katz *et al.*, 1983) were gifts from D. A. Hopwood and T. Kieser of the John Innes Institute, Norwich, UK. *Streptomyces cacaoi* KCC-SO352 was obtained from A. Seino, Kaken Chemical Co. Ltd, Tokyo, Japan. *Streptomyces albus* G, *Streptomyces* R61 and *Actinomadura* R39 were from the Microbiology Department of the University of Liège. *Escherichia coli* HB101 (Boyer & Roulland-Dussoix, 1969) and plasmid vector pBR322 (Bolivar *et al.*, 1977) were also used.

**Media and growth conditions.** Growth of *Streptomyces* cultures was at 28 °C in YEME medium (Hopwood *et al.*, 1985) with vigorous orbital shaking (New Brunswick G25 shaker, 300 r.p.m.). The following liquid media were also used: (i) P medium (Okanishi *et al.*, 1974) supplemented with 0.5% yeast extract; (ii) Oxoid Mueller–Hinton broth; (iii) Lennox broth (Lennox, 1955); (iv) Merck peptone broth; and (v) E9 broth (Dehottay *et al.*, 1986). R2YE agar (Hopwood *et al.*, 1985) was used for maintenance and plating of *Streptomyces* strains. *E. coli* was grown at 37 °C in Luria–Bertani or M9 medium (Maniatis *et al.*, 1982) with vigorous orbital shaking (see above).

**Recombinant DNA techniques.** Preparation of *Streptomyces* chromosomal and plasmid DNA and transformation of *Streptomyces* protoplasts were done as described by Hopwood *et al.* (1985). Treatment with restriction endonucleases or Klenow fragment, separation of the digested products by agarose gel electrophoresis, treatment with alkaline phosphatase, and ligation experiments were performed as described by Maniatis *et al.* (1982). Elution of the DNA fragments from the gels (Dretzen *et al.*, 1981). \(^{32}\)P-labelling by nick translation (Maniatis *et al.*, 1975), hybridization experiments (Southern, 1975), addition of *BglII* linkers (Maniatis *et al.*, 1982), and isoelectric focusing (Righetti & Drysdale, 1971) were done as described in the relevant references.

**Enzymes and antibiotics.** The enzymes used in recombinant DNA techniques were from Bethesda Research Laboratories, Boehringer Mannheim, Amersham or Sigma. The antibiotics were from Sigma except for nitrocefin (BBL Microbiology Systems), thiostrepton (a gift from J. Kemp, Pfizer Central Research, Sandwich, UK) and \( \beta \)-iodopenicillanate (gift from J. Frère, 1985; and unpublished results). \( \beta \)-Lactamase-producing *Streptomyces* colonies grown on R2YE agar plates were detected on the basis of the red halo that they developed on exposure to nitrocefin (Dehottay *et al.*, 1986).

**RESULTS AND DISCUSSION**

Cloning of the \( \beta \)-lactamase gene from *S. cacaoi*

Fig. 1 shows the restriction maps of the *S. cacaoi* chromosomal DNA fragments used in the experiments described below, and the various derivatives of pIJ702 in which these fragments were inserted. pIJ702 carries a thiostrepton-resistance marker (tsr) and a tyrosinase-encoding gene (mel) which offers three unique restriction sites for insertional inactivation.

Chromosomal DNA from *S. cacaoi* was fully digested with *BglII*, *SphI* or *SstI* and the unfractionated digests were used in shotgun cloning experiments. Shotgun cloning using the *SphI* site of the high-copy-number *Streptomyces* plasmid pIJ702 led to the isolation of one \( \beta \)-lactamase-producing colony, among 3000 melanin-negative, thiostrepton-resistant *S. lividans* TK24 transformants. The resulting plasmid contained a 19 kb insert; it was called pDML51 and the strain bearing it was called *S. lividans* ML1. *SphI* digestion of pDML51 not only regenerated the 5-8 kb pIJ702 and the 19 kb insert but also produced two other fragments of 6-6 and 12-4 kb. Time-course experiments showed that these fragments originated by cleavage of an additional *SphI* site of relatively low susceptibility in the 19 kb insert. Each fragment was recloned into pIJ702 using the *SphI* sites as above. \( \beta \)-Lactamase activity was associated only with the 6-6 kb fragment. The plasmid with this 6-6 kb insert was called pDML52 and the strain bearing it was called *S. lividans* ML2.

The 6-6 kb fragment containing the \( \beta \)-lactamase gene was inserted into the *SphI* site of pBR322 and the recombinant plasmid pDML72 was isolated from ampicillin-resistant,
Cloning of a Streptomyces β-lactamase gene

Fig. 1. Restriction maps and cloning in pIJ702 of chromosomal DNA fragments carrying the S. cacaoi β-lactamase gene. (Plasmids are not drawn to scale.) The restriction map was derived from pMCP38 and pBR322-derived pDML72 (not shown; see text).

tetraphycycle-sensitive E. coli HB101 transformants. pDML72 was 32P-labelled by nick-translation and used as a probe in hybridization experiments against SphI digests of chromosomal DNAs from the β-lactamase-producing Actinomadura R39, S. coelicolor, S. albus G and S. cacaoi and the β-lactamase-non-producing or very poorly producing Streptomyces R61 and S. lividans TK24. Hybridization occurred only with the S. cacaoi 6.6 kb fragment and the pDML51 and pDML52 digests (Fig. 2). With the S. cacaoi DNA SphI digest used for the shotgun cloning, two hybridization bands (6.6 and 19 kb) were observed, indicating incomplete digestion (not shown).

pDML52 was also used to prepare smaller DNA fragments containing the β-lactamase gene (Fig. 1). Three constructions were made. (i) pDML52 was cut with BcII and SphI and the 4.4 kb fragment thus produced was isolated and ligated to pIJ702 restricted with BglII and SphI to produce pMCP38 (in S. lividans strain CMA38). (ii) pDML52 was cut with BcII and the 4.7 kb fragment thus produced was in turn partially digested with PvuII with generation of a 2 kb fragment. After addition of BglII linkers, this fragment was ligated to pIJ702 cut with BglII and treated with bacterial alkaline phosphatase to produce pMCP39 (in S. lividans strain CMA39). (iii) Cleavage of pDML52 with BcII and NeoI generated a 1.8 kb fragment and cleavage of pIJ702 with BglII and SstI generated a 5.5 kb fragment. These fragments were ligated to give rise to a 7.3 kb NeoI–SstI linear fragment. After treatment of the NeoI and SstI ends by the Klenow fragment of E. coli DNA polymerase I, blunt-end ligation yielded the 7.3 kb plasmid pDML53 (in S. lividans strain ML3).

Strains CMA38, CMA39 and ML3 gave, to varying extents, a positive nitrocefin test on R2YE agar. However, as shown below, the abilities of these strains to produce extracellular β-lactamase in liquid media were very different.

Expression and properties of the extracellular β-lactamase of S. lividans ML1

S. lividans ML1 was grown in various liquid media and the β-lactamase activity of the culture filtrates was estimated. The following data are mean values of two independent experiments, the observed variations being less than 10%. Maximal production, 38 mg enzyme l−1, occurred in E9 broth after 3–4 d growth at 28 °C. The yields were 16–17 mg l−1 in P medium and Lennox broth, 12 mg l−1 in YEME, 9 mg l−1 in Merck peptone broth and 6 mg l−1 in Oxoid Mueller–Hinton broth. With S. cacaoi KCC-S0352, production was also maximal in E9 broth (1.4 mg l−1).
Hence, *S. lividans* ML1 was a better β-lactamase producer than *S. cacaoi*, with a 30-fold increase in enzyme excretion observed under optimal conditions (though an effect of the medium on enzyme excretion could not be excluded; Dehottay et al., 1986).

The extracellular β-lactamase of *S. lividans* ML1 grown in E9 medium was purified to protein homogeneity (specific activity 575 μmol min⁻¹ mg protein⁻¹) using the procedure described for the *S. cacaoi* enzyme. Not only the behaviour of the *S. lividans* ML1 β-lactamase at each step of the purification but the molecular mass (34 kDa as revealed by SDS-polyacrylamide gel electrophoresis), isoelectric point (4.95), and kinetics of inhibition by β-iodopenicillanate of the purified enzyme were exactly those expected from the known properties of the *S. cacaoi* β-lactamase. Moreover, the two enzymes co-eluted by MonoQ FPLC chromatography.

Sub-cloning experiments (see Fig. 1) showed that the β-lactamase gene was associated with a 1.8 kb *NcoI–BclI* fragment (with the coding region starting close to the *NcoI* site; unpublished data). However, all these sub-clones resulted in *S. lividans* strains that had a much decreased capacity to produce the extracellular β-lactamase (Table 1, Fig. 3). In particular, cleavage of the original 19 kb DNA fragment at the internal *SphI* site about 3 kb upstream of the β-lactamase structural gene (strain ML2) had a drastic effect, suggesting that the eliminated 12.4 kb segment had important information for β-lactamase gene expression (and/or enzyme secretion). It should be noted that the observed decreased yields in excreted enzyme by the sub-clones did not relate...
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Fig. 3. Growth (○) and β-lactamase excretion (●) of S. lividans ML1 (——) and S. cacaoi (-----) grown in medium E9 at 28°C.

Table 1. Maximal levels of β-lactamase excretion in medium E9 by S. lividans carrying various recombinant plasmids and by S. cacaoi KCC-S0352

For strains and plasmids, see Fig. 1. All the strains except S. lividans TK24 and S. cacaoi KCC-S0352 were grown in medium E9 supplemented with 5 µg thiostrepton ml⁻¹. The yields are those obtained during time-course experiments (Fig. 3). The results represent the mean of two independent experiments; individual variations were less than 10%.

<table>
<thead>
<tr>
<th>Strain/plasmid</th>
<th>Size of S. cacaoi DNA insert (kb)</th>
<th>Mycelium (g dry wt l⁻¹)</th>
<th>Extracellular β-lactamase (mg l⁻¹)</th>
</tr>
</thead>
<tbody>
<tr>
<td>S. lividans ML1/pDML51</td>
<td>19</td>
<td>4.6</td>
<td>4.1</td>
</tr>
<tr>
<td>ML2/pDML52</td>
<td>6.6</td>
<td>4.5</td>
<td>8.4</td>
</tr>
<tr>
<td>CMA38/pMCP38</td>
<td>4.4</td>
<td>6.6</td>
<td>4.8</td>
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<tr>
<td>CMA39/pMCP39</td>
<td>2.0</td>
<td>5.2</td>
<td>1.8</td>
</tr>
<tr>
<td>ML3/pDML53</td>
<td>1.8</td>
<td>4.6</td>
<td>0.14</td>
</tr>
<tr>
<td>TK24/—</td>
<td>—</td>
<td>4.4</td>
<td>ND</td>
</tr>
<tr>
<td>S. cacaoi KCC-S0352/—</td>
<td>—</td>
<td>11.2</td>
<td>1.6</td>
</tr>
</tbody>
</table>

ND, Not detectable.

to variation in the amounts of mycelium produced (Table 1) nor to instability and/or integrity of the plasmids (data not shown). Selection for the plasmids by thiostrepton was applied in all cases (except with S. lividans TK24 and S. cacaoi KCC-S0352) and the plasmids were found to have conserved their insert (at least after 3 d growth). However, the possible effects of plasmid copy number and growth rates (as opposed to growth yield) in strains containing the subclones have not been assessed; these factors cannot be entirely discounted, since they could potentially influence β-lactamase production.

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REFERENCES


