

Molecular Cloning of an *aepA* Gene that Activates Production of Extracellular Pectolytic, Cellulolytic, and Proteolytic Enzymes in *Erwinia carotovora* subsp. *carotovora*

H. Murata,¹ J. L. McEvoy,¹ A. Chatterjee,¹ A. Collmer,² and A. K. Chatterjee¹

¹Department of Plant Pathology, University of Missouri, Columbia 65211; and ²Department of Plant Pathology, Cornell University, Ithaca, NY 14853-5908 U.S.A.

Received 14 November 1990. Accepted 3 January 1991.

Strain 71 of *Erwinia carotovora* subsp. *carotovora* produces extracellular enzymes such as pectate lyase (Pel), polygalacturonase (Peh), cellulase (Cel), and protease (Prt). The levels of extracellular Pel, Cel, and Prt were higher in a medium containing crude celery extract than in a medium containing pectate. Using transposons (Tn5, *TnphoA*, and *Tn10-lacZ*), we isolated pleiotropic mutants that were deficient in extracellular levels of these enzymes and attenuated in their ability to macerate plant tissues. The mutants, however, were similar to the parent in their ability to utilize various sugars and to produce periplasmic enzymes. In an *E. c.* subsp. *carotovora* 71 gene library, we detected a cosmid, pAKC264, that restored extracellular enzyme production and tissue maceration in all the mutants. The cosmid appears not to carry *pel*, *peh*, *cel*, or *prt* genes. In *E. c.* subsp. *carotovora* 71, pAKC264 stimulated the production of Pel, Peh, Cel, and

Prt, but it did not affect the levels of the periplasmic enzymes, cyclic phosphodiesterase, or β -lactamase. pAKC602, a subclone of pAKC264, stimulated enzyme production in *E. c.* subsp. *carotovora* 71 and did not complement mutations in *cya*, the gene specifying adenylate cyclase, or *crp*, the gene specifying cyclic AMP receptor protein, in *Escherichia coli*. The *E. c.* subsp. *carotovora* 71 gene that activates extracellular protein production was designated as *aepA*. Sixteen mini-Mu-*lacZ* (*MudI1734*) insertions inactivating *aepA* spanned a DNA region of about 0.8 kilobases and allowed determination of the direction of *aepA* transcription by screening for β -galactosidase production. By further subcloning and localizing the sites of mini-Mu-*lacZ* insertions not inactivating *aepA*, the gene was localized within a 1.1-kilobase DNA segment.

Erwinia carotovora subsp. *carotovora* (Jones) Bergey *et al.* causes tissue-macerating (= soft rot) diseases in a wide variety of plant hosts. Pathogenicity of these bacteria is correlated with their ability to produce extracellular enzymes, specifically pectinases such as pectate lyase (Pel) or endopolygalacturonase (endo-Peh) (Lei *et al.* 1985; Mount *et al.* 1970; Roberts *et al.* 1986; Willis *et al.* 1987). They also produce extracellular protease (Prt), cellulase (Cel), and phospholipase, which are known not to macerate plant tissues. These enzymes may nevertheless contribute to virulence by augmenting the action of pectinases, by inflicting stress onto plant cells previously damaged by pectolytic activities, and by providing the bacterium with readily utilizable sources of carbon and nitrogen through the degradation of polymeric substances.

These ecological considerations and a lack of understanding of the processes affecting extracellular protein production in gram-negative bacteria have generated considerable interest in the *Erwinia* enzymes (for reviews see Chatterjee and Vidaver 1986; Collmer and Keen 1986; Kotoujansky 1987). Aside from structural and functional aspects of the genes specifying extracellular enzymes (see, for example, Hinton *et al.* 1989, 1990; Lei *et al.* 1988; Reverchon *et al.* 1989; Saarilahti *et al.* 1990; Tamaki *et al.* 1988; Van Gijsegem 1989), the issues currently attracting the most attention are 1) biochemical mechanisms by which

proteins are translocated across the tripartite cell envelope (He *et al.*, in press; Ji *et al.* 1989; Murata *et al.* 1990) and 2) molecular aspects of regulation of the expression of genes specifying secreted proteins (Beaulieu and Van Gijsegem 1990; Chatterjee *et al.*, in press; Kelemu and Collmer 1990; Reverchon *et al.* 1989; Yang *et al.* 1989).

Investigations with *Bacillus subtilis* (Ehrenberg) Cohn (Henner *et al.* 1988; Wang and Doi 1990), *Staphylococcus aureus* Rosenbach (Peng *et al.* 1988; Recsei *et al.* 1985), and *Streptomyces griseus* (Krainsky) Waksman and Henrici (Daza *et al.* 1990) have disclosed global regulation of secreted proteins. That Pel, Peh, Cel, Prt, and other extracellular proteins may also be similarly regulated in *E. c.* subsp. *carotovora* was suggested by the phenotypes of mutants isolated by Beraha and Garber (1971). The nitro-soguanidine-induced mutants were pleiotropically defective in the production of these extracellular enzymes as well as phospholipase. They also isolated a one-step revertant that regained the ability to produce all of those enzymes. These findings could have resulted from a defect in a "positive regulator (= activator) gene" required for the production of extracellular proteins. However, since the mutants and the revertant were not subjected to further genetic analyses, the basis for the pleiotropic effect remained unresolved.

In the course of isolating mutants defective in the secretion (Out) pathway (Murata *et al.* 1990), we came across a few mutants wherein the production of Peh, Pel, Cel, and Prt was affected. Such mutants, phenotypically similar to those reported by Beraha and Garber (1971), were different from the bona fide Out⁻ mutants (Murata *et al.*

Address correspondence to A. K. Chatterjee.

1990) in 1) the detrimental effect of mutations on the production of extracellular Prt in addition to Peh, Pel, and Cel, 2) the absence of high levels of cell-bound Peh, Pel, and Cel, and 3) the lack of restoration of the mutational phenotype by the *out*⁺ DNA. We describe here the characteristics of the mutants and a wild-type allele designated as *aepA*⁺ for the activation of extracellular protein production. In addition, our finding that the *aepA*⁺ DNA restores plant tissue macerating ability to the mutants suggests a key role of the *aepA* gene product in pathogenesis of *E. c.* subsp. *carotovora*. (Preliminary accounts of parts of this work will be or have been published [Chatterjee *et al.*, in press; Murata and Chatterjee 1990a, 1990b].)

MATERIALS AND METHODS

Bacterial strains, plasmids, and bacteriophage. Bacterial strains, plasmids, and the bacteriophage that were used in this study are described in Table 1. The strains carrying drug markers were maintained on Luria-Bertani agar sup-

plemented with the appropriate antibiotics. The wild-type strains of *Erwinia* were maintained on yeast extract-glucose-calcium carbonate (YGC) agar. Lambda lysates were kept in SM buffer (Maniatis *et al.* 1982).

Media. Cel detection agar, Luria-Bertani medium, minimal medium, nutrient gelatin agar, polygalacturonate-yeast extract-citrus pectin (PYCP) agar, and YGC agar were described previously (Barras *et al.* 1986; Chatterjee 1980; Chatterjee *et al.* 1985; Murata *et al.* 1990; Thurn and Chatterjee 1985). Salts-yeast extract-glycerol (SYG) medium consisted of 0.4 mM MgSO₄, 7.6 mM (NH₄)₂SO₄, 55 mM potassium phosphate (pH 7.0), 0.025% yeast extract, and 0.2% glycerol. SYG medium was supplemented either with pectate (0.2%) or with plant extracts prepared as follows. Carrot roots, celery petioles, or potato tubers obtained from a local supermarket were cut into pieces; 100 g of the cut material was suspended in 100 ml of distilled water and then autoclaved at 121° C for 20 min. After autoclaving, the liquid material was decanted and centrifuged at 12,000 × *g* at 4° C for 30 min to remove debris. To 95 ml of

Table 1. Bacterial strains, bacteriophage, and plasmids used in this study

Designation	Relevant characteristics ^a	References or sources	Designation	Relevant characteristics ^a	References or sources
<i>Erwinia carotovora</i> subsp. <i>carotovora</i>			Bacteriophage		
Strain 71	Wild type	Zink <i>et al.</i> 1984	λ1045	λ-Tn10- <i>lacZ</i> , Tc ^r	Way <i>et al.</i> 1984
AC5006	Lac ⁻	Derivative of <i>E. c.</i> subsp. <i>carotovora</i> 71; Murata <i>et al.</i> 1990	Plasmids		
AC5012	Tn5 insertion mutant, AepA ⁻ , Km ^r	Derivative of <i>E. c.</i> subsp. <i>carotovora</i> 71; this study	pHCP2	<i>lamB</i> ⁺ , Mob ⁺ , Ap ^r	Clement <i>et al.</i> 1982
AC5026	Tn10- <i>lacZ</i> insertion mutant, AepA ⁻ , Tc ^r	Derivative of AC5006; this study	pJB4JI	pPHIJI::Mu::Tn5, Mob ⁺ , Tra ⁺ , Gm ^r , Km ^r	Hirsch and Beringer 1984
AC5027	TnphoA insertion mutant, AepA ⁻ , Km ^r	Derivative of <i>E. c.</i> subsp. <i>carotovora</i> 71; this study	pRK404	Mob ⁺ , Tc ^r	Ditta <i>et al.</i> 1985
<i>Escherichia coli</i>			pRK609	pRK600::TnphoA, Km ^r , Cm ^r	Long <i>et al.</i> 1988
AC8001	<i>arg</i> , <i>his</i> , Δ <i>lac</i> , Sm ^r , Km ^r , Gm ^r (pJB4JI)	Chatterjee <i>et al.</i> 1983	pRK2013	Mob ⁺ , Tra ⁺ , Km ^r	Figurski and Helsinki 1979
CGSC6027	Δ <i>cya-854</i> , <i>relA1</i> , <i>spoT1</i> , <i>thi-1</i> , λ ⁻	<i>E. coli</i> Genetic Stock Center ^b (B. J. Bachmann); Brickman <i>et al.</i> 1973	pSF6	Mob ⁺ , Sp ^r , Sm ^r	Selvaraj <i>et al.</i> 1984
CGSC7043	Δ <i>crp-45</i> , <i>relA1</i> , <i>rpsL136</i> , <i>spoT1</i> , <i>thi-1</i> , λ ⁻	<i>E. coli</i> Genetic Stock Center (B. J. Bachmann); Brickman <i>et al.</i> 1973	pAKC264	AepA ⁺ , Sp ^r , Sm ^r	pSF6 containing <i>E. c.</i> subsp. <i>carotovora</i> 71 genomic DNA; this study
DH5α	φ80 <i>dlacZ</i> ΔM15, Δ(<i>lacZYA-argF</i>), <i>U169</i> , <i>recA1</i> , <i>thi-1</i>	Bethesda Research Laboratories ^c	pAKC602	AepA ⁺ , Tc ^r	Subcloning of <i>aepA</i> ⁺ DNA into pRK404; this study
DW75	<i>lamB</i> ⁺ , Ap ^r , Cm ^r , Tc ^r	Salmond <i>et al.</i> 1986	pAKC607	AepA ⁺ , Km ^r , Tc ^r	MudI1734 insertion in pAKC602; this study
M8820	Δ(<i>proAB-argF-lacPOZYA</i>), <i>recA</i> ⁺ , Sm ^r	Castilho <i>et al.</i> 1984	pAKC608	AepA ⁺ , Tc ^r	2.2-kilobase (kb) <i>Hind</i> III <i>aepA</i> ⁺ subclone of pAKC607 containing 1.2 kb of <i>E. c.</i> subsp. <i>carotovora</i> 71 DNA and 1.0 kb of Mu DNA in pRK404; this study
MM294A	<i>pro-82</i> , <i>thi-1</i> , <i>endA1</i> , <i>hsdR17</i> , <i>supE44</i>	Long <i>et al.</i> 1988	pAKC609	AepA ⁺ , Tc ^r	Same as pAKC608 except the insert DNA is cloned in the opposite orientation; this study
PO11734	MudI1734 <i>ara</i> ::(Mu cts), Δ(<i>proAB-argF-lacIPOZYA</i>), Km ^r , Sm ^r	Castilho <i>et al.</i> 1984			
HB101	<i>pro</i> , <i>leu</i> , <i>thi</i> , <i>lacY</i> , <i>recA</i> , <i>hsdR</i> , <i>hsdM</i> , Sm ^r	Zink <i>et al.</i> 1984			

^a Km = kanamycin, Tc = tetracycline, Sm = streptomycin, Gm = gentamycin, Ap = ampicillin, Cm = chloramphenicol, Sp = spectinomycin, and ^r = resistant.

^b Yale University, School of Medicine, New Haven, CT.

^c Gaithersburg, MD.

this solution, these stock solutions were added as follows: 1 ml of 40 mM MgSO₄, 1 ml of 760 mM (NH₄)₂SO₄, 1 ml of 20% glycerol, 2 ml of 2.75 M potassium phosphate (pH 7.0), and 0.25 ml of 10% yeast extract. When required, antibiotics were added as follows: ampicillin, 50 µg/ml; chloramphenicol (Cm), 10 µg/ml; gentamycin, 10 µg/ml; kanamycin (Km), 50 µg/ml; spectinomycin, 50 µg/ml; streptomycin, 100 µg/ml; and tetracycline (Tc), 10 µg/ml. The minimal medium was supplemented with 50 µg of amino acid per milliliter as necessary. Media were solidified by the addition of Difco agar (Difco Laboratories, Detroit, MI) (1.5%). Soft (top) agar contained 0.6% Difco agar.

Transposon mutagenesis. Tn5 and Tn10-*lacZ* insertion mutants were obtained by using the plasmid pJB4JI and λ-Tn10-*lacZ*, respectively (Murata *et al.* 1990). Tn*phoA* mutations were generated in *E. c.* subsp. *carotovora* 71 by mating with *Escherichia coli* MM294A carrying the plasmid pRK609 (Long *et al.* 1988). Antibiotic-resistant colonies were scored for defects in pectolytic, proteolytic, and cellulolytic activities on PYCP agar, nutrient gelatin agar, and Cel detection agar, respectively. *aepA-lacZ* transcriptional fusions were derived from pAKC264 and pAKC602 (Table 1 and Fig. 1) with the mini-Mu-*lacZ* element, MudI1734, as described previously (Castilho *et al.* 1984; McEvoy *et al.* 1990). Individual plasmids containing MudI1734 insertions were transferred to the AepA⁻ mutant

AC5012 by triparental matings. The transconjugants that remained deficient in pectolytic, proteolytic, and cellulolytic activities, as determined by agar plate assays, were presumed to result from inactivation of the *aepA*⁺ gene. Plasmids containing *aepA-lacZ* were further characterized by screening the transconjugants for β-galactosidase production on agar media containing X-gal. *Hind*III, which cleaves pAKC602 twice (within the polylinker and the internal fragment; Fig. 1) and MudI1734 twice (0.1 kb from the 5' end of the *lacZ* transcriptional unit and 1.0 kb from the 3' end, McEvoy *et al.* 1990), and *Sst*I, which cleaves once within the insert DNA (Fig. 1) and once within MudI1734 (Metcalf *et al.* 1990), were used in determining the location and orientation of the insertions. *Eco*RI, *Eco*RV, and *Pst*I digests were used to confirm mapping of the insertions.

Preparation and assay of samples for enzymatic activities.

Bacteria were grown in desired media to about 250 Klett units (Klett-Summerson colorimeter, No. 66 filter; Klett Manufacturing Co., Inc., New York), and cells and supernatants were separated by centrifugation (12,100 × *g*) at 4° C. The protocols for preparing periplasmic fractions, cellular extracts, and dialyzed samples; the procedures for the quantitative assays of Cel, cyclic phosphodiesterase, β-lactamase, Pel, Peh, and Prt activities; and the definitions of the enzymatic units have been described previously

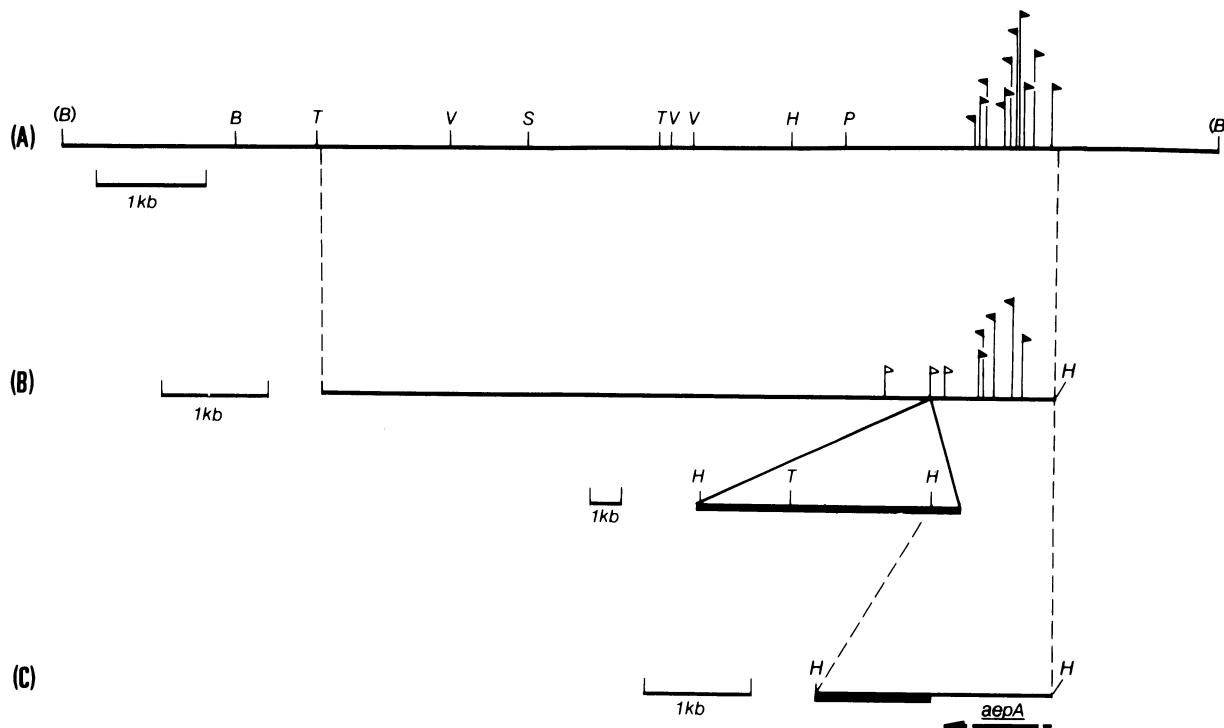


Fig. 1. Cloned portion of the *Erwinia carotovora* subsp. *carotovora* 71 genome harboring the *aepA* gene. **A**, Restriction map of the 10.5-kilobase (kb) region of *E. c.* subsp. *carotovora* 71 DNA contained in the cosmid pAKC264. The solid flags denote the sites and orientations of MudI1734 insertions inactivating *aepA*; flags pointing to the left indicate insertions that produced high levels of β-galactosidase. **B**, A 6.5-kb *Sau*3A subclone of *aepA* DNA contained in pAKC602. The solid flags indicate the sites and orientations of MudI1734 insertions inactivating *aepA*; the open flags indicate insertions not inactivating *aepA*. Solid flags pointing to the left indicate the insertions from which high levels of β-galactosidase were produced. Also shown in detail is the MudI1734 insertion (bold line) found in pAKC607. **C**, A *Hind*III fragment from pAKC607 that contains about 1.0 kb of MudI1734 DNA (bold line) and 1.2 kb of *E. c.* subsp. *carotovora* 71 DNA including the *aepA* gene. This fragment was cloned in both orientations in pRK404 resulting in the *aepA*⁺ plasmids pAKC608 and pAKC609. The direction of transcription and extent of *aepA* are shown at the bottom of the figure. Abbreviations are as follows: B, *Bam*HI; H, *Hind*III; P, *Pst*I; S, *Sal*I; T, *Sst*I; and V, *Eco*RV.

(Barras *et al.* 1986; Murata *et al.* 1990; Thurn and Chatterjee 1985). Protein content was determined by using the BCA reagent (Pierce Chemical Co., Rockford, IL).

Isolation of the *AepA*⁺ cosmid. The procedures used for the screening of an *E. c.* subsp. *carotovora* 71 gene bank have been described in a previous publication (Murata *et al.* 1990). An *AepA*⁻ mutant, AC5012, was used as the recipient. The transconjugants resulting from triparental matings were scored for the production of pectolytic, cellulolytic, and proteolytic activities using agar plate assays.

Recombinant DNA techniques. Standard published procedures were used for the isolation of plasmid and chromosomal DNAs, transformation of *E. coli*, restriction digests, gel electrophoresis, and electroelution (Crouse *et al.* 1983; Maniatis *et al.* 1982). DNA-modifying enzymes were obtained from Promega (Madison, WI).

Plant tissue maceration. The potato tuber assay was described previously (Murata *et al.* 1990). The celery petiole assay was done in the same way as the potato tuber assay except that celery petioles were soaked in 0.5% sodium hypochlorite for 5 instead of 10 min, and inoculation sites were not sealed with Vaseline. Surfaces of celery petioles were punctured, and bacterial suspensions, which were prepared as described previously (Murata *et al.* 1990), were inoculated through the wounds into the vascular tissues.

RESULTS

Effects of crude plant extracts on enzyme production. To develop a medium that could sustain the production

of high levels of Pel, Peh, Cel, and Prt in *E. c.* subsp. *carotovora* 71, we tested the effects of crude extracts from various plant tissues. In preliminary trials, we found higher levels of extracellular Pel in SYG medium containing celery extract than in SYG medium supplemented with pectate, carrot extract, or potato extract. Subsequently, we compared the levels of various enzymes in SYG and SYG supplemented with pectate or celery extract. The data (Table 2) show that *E. c.* subsp. *carotovora* 71 produced higher levels of extracellular Pel as well as Cel and Prt in SYG plus celery extract than in SYG or SYG plus pectate. However, extracellular Peh activity was similar in media supplemented with pectate or celery extract. A particularly striking feature was that 94% of Pel, 92% of Cel, and >99% of Prt activities in SYG plus celery extract were present in the culture supernatant. Because of this stimulatory effect of celery extract on the production of extracellular Pel, Cel, and Prt, we routinely used SYG plus celery extract in our subsequent work.

Isolation of transposon insertion mutants defective in extracellular enzyme production. As was done in our previous studies (Murata *et al.* 1990; Zink *et al.* 1984), we introduced Tn5 into *E. c.* subsp. *carotovora* 71 using pJB4JI and Tn10-*lacZ* into AC5006 using λ ::Tn10. For mutagenesis with TnphoA, the Tra⁺ plasmid pRK609 (Long *et al.* 1988) was used. This plasmid is unstable in *E. c.* subsp. *carotovora* 71 because it contains a ColE1 origin of replication. Therefore, in the absence of Cm selection, a high proportion (>99%) of Km^r Cm^s transconjugants were produced. In gel electrophoretic analysis of plasmid preparations, repre-

Table 2. Effect of pectate and celery extract on extracellular enzyme production in *Erwinia carotovora* subsp. *carotovora* 71

Media ^a	Activity ^b (units/mg protein)									
	Pel		Peh		Cel		Prt		Bla	Cpd
	S	C	S	C	S	C	S	C	C ^c	C
SYG	0.3	2.3	<0.1 ^d	5.6	<0.1	<0.1	<0.1	<0.1	8.5	2.4
SYG + pectate	16.7	13.3	4.1	8.1	5.6	10.4	<0.1	<0.1	5.5	2.4
SYG + celery extract	29.6	1.9	3.5	1.3	34.3	3.0	8.0	<0.1	6.6	1.0

^a Bacteria were grown in minimal salts-yeast extract-glycerol (SYG) medium or in this medium supplemented with pectate (0.2%) or celery extract. For additional details see the text.

^b Supernatants (S) and cell extracts (C) of cultures grown to about 250 Klett units in a shaker at 28°C were assayed. Pel = pectate lyase, Peh = polygalacturonase, Cel = cellulase, Prt = protease, Bla = β -lactamase, and Cpd = cyclic phosphodiesterase.

^c Bla and Cpd activities were only detected in cell extracts.

^d <0.1 units for Peh, Cel, or Prt represent the limits of detection of activities under the assay conditions used.

Table 3. Effect of *aepA*⁺ DNA on the production of extracellular and periplasmic enzymes in *Erwinia carotovora* subsp. *carotovora* 71 and a Tn5 insertion mutant, AC5012^a

Enzyme ^b	Activity ^c (units/mg protein)							
	Strain 71/pSF6 ^d		Strain 71/pAKC264 ^e		AC5012/pSF6		AC5012/pAKC264	
	S	C	S	C	S	C	S	C
Pel	13.7	0.7	72.5	2.0	2.0	2.3	70.5	3.0
Peh	1.9	0.9	13.1	2.1	<0.1 ^f	1.3	10.6	2.0
Cel	15.6	<0.1	83.6	7.3	<0.1	6.5	149.1	15.3
Prt	3.6	<0.1	15.2	<0.1	<0.1	<0.1	12.2	<0.1
Bla	<0.02 ^g	2.4	<0.02	2.5	<0.02	3.3	<0.02	3.0
Cpd	<0.02	1.1	<0.02	0.9	<0.02	2.3	<0.02	2.0

^a Bacteria were grown in minimal salts-yeast extract-glycerol (SYG) medium supplemented with celery extract. For additional details see the text.

^b Pel = pectate lyase, Peh = polygalacturonase, Cel = cellulase, Prt = protease, Bla = β -lactamase, and Cpd = cyclic phosphodiesterase.

^c Supernatants (S) and cell extracts (C) of cultures grown to about 250 Klett units in a shaker at 28°C were assayed for enzymatic activities.

^d pSF6 = the cloning vector (Table 1).

^e pAKC264 = pSF6 + *aepA*⁺ DNA (Table 1 and Fig. 1).

^f <0.1 units of activity for Peh, Cel, or Prt represent the limit of detection under the assay conditions used.

^g <0.02 units of activity for Bla and Cpd represent the limit of detection under the assay conditions used.

sentative Km^r Cm^s derivatives were found not to contain pRK609 (data not shown). These data collectively indicate that the Km^r and Cm^s colonies resulted from the loss of the plasmid and the presence of Tn*phoA* in the *E. c.* subsp. *carotovora* 71 genome.

Agar plate screening and subsequent quantitative assays for the production of extracellular enzymes in strains carrying transposon insertions revealed the occurrence of two classes. Class I was represented by Out⁻ mutants previously described by Murata *et al.* (1990). Class II is represented by three independently isolated AepA⁻ mutants (Table 1); quantitative data for one mutant (AC5012) is given in Table 3. While the levels of cell-bound Pel, Peh, and Cel as well as Prt in this mutant were generally low and comparable to those in the parent, extracellular activities were much reduced. The absence of high levels of Cel, Pel, and Peh in cells and the deficiency in Prt production (Table 3) clearly distinguish the AepA⁻ mutants from the bona fide Out⁻ mutants of *E. c.* subsp. *carotovora* (Hinton and Salmond 1987; Murata *et al.* 1990) or *E. chrysanthemi* Burkholder *et al.* (Andro *et al.* 1984; Thurn and Chatterjee 1985). In addition, the plasmids carrying the *E. c.* subsp. *carotovora* 71 *out* cluster (Murata *et al.* 1990) did not restore enzyme production in any of the AepA⁻ mutants (data not shown). However, these mutants, like the Out⁻ mutants (Murata

et al. 1990), were severely attenuated in their ability to macerate celery petioles or potato tuber tissue (Fig. 2). The AepA⁻ mutants, like their parents, utilized the following compounds as carbon sources: glycerol, arabinose, ribose, xylose, glucose, galactose, galacturonate, mannitol, sucrose, raffinose, melibiose, and polygalacturonate.

Molecular cloning of the *aepA*⁺ DNA. By mobilizing an *E. c.* subsp. *carotovora* 71 library of about 1,000 cosmids into AC5012, we obtained one cosmid, pAKC264, that restored extracellular enzyme production. This cosmid restored extracellular enzyme production in all of the AepA⁻ mutants but not in Out⁻ mutants. Along with extracellular enzyme production, the AepA⁻ mutants carrying pAKC264 regained the ability to macerate plant tissue like the parent strain (Fig. 2). pAKC264 appears not to carry the structural genes for the extracellular enzymes Pel, Cel, Peh, or Prt, because these activities were not detected in HB101 carrying the cosmid (data not shown).

We subcloned the *aepA*⁺ segment of pAKC264 by preparing a *Sau*3A partial digest and ligating the fragments to *Bam*HI-cut pRK404 treated with phosphatase. The Tc^r DH5 α transformants that remained white in the presence of X-gal were picked, and their plasmids were mobilized into AC5012. Of the several plasmids that restored enzyme production in AC5012, we selected pAKC602 for our subsequent analysis, because it carried the smallest insert DNA of about 6.5 kb. pAKC602, like its parent pAKC264, restored enzyme production in AC5012 and AC5027. To localize the *aepA*⁺ DNA, we mutagenized the insert DNAs of pAKC264 and pAKC602 with *Mud*II1734 (mini-*Mu-lacZ*; Castilho *et al.* 1984) and tested Km^r derivatives for their ability to restore enzyme production. The results of such analyses, taken along with the sites of mini-*Mu-lacZ* insertions (indicated in Fig. 1), allowed localization of *aepA* within a 1.1-kb DNA stretch. Indeed, 16 plasmids containing *Mud*II1734 insertions that mapped within about 800 bp of DNA (Fig. 1) failed to restore extracellular enzyme production in the AepA⁻ mutants. To confirm the limits of *aepA*, we subcloned into pRK404 a 2.2-kb fragment spanning from the *Hind*III site of the vector polylinker to the *Hind*III site within the *Mu* DNA of the *Mud*II1734 insertion in pAKC607 (Fig. 1). This fragment, containing about 1.2 kb of *E. c.* subsp. *carotovora* 71 DNA and 1.0 kb of *Mu* DNA, was cloned in both orientations in pRK404 and produced plasmids pAKC608 and pAKC609. Both of these plasmids restored enzyme production in AC5012 and AC5027. Thus, the 1.2-kb *E. c.* subsp. *carotovora* 71 DNA appears to contain sequences encoding a functional *aepA* gene as well as the *aepA* promoter.

Stimulation of extracellular enzyme production by the *aepA*⁺ DNA. The data in Table 3 show that in the SYG medium supplemented with celery extract, AC5012 carrying pAKC264 produced higher levels of enzymes than AC5012 carrying the vector pSF6; the levels of extracellular Pel, Peh, Cel, and Prt were 35-fold, >100-fold, >150-fold, and >10-fold higher, respectively. We should point out that the levels of these enzymes within the cells were not markedly affected by the presence of pAKC264. The data also show that cyclic phosphodiesterase and β -lactamase activities in the mutant were comparable to those produced by wild-type strain 71 of *E. c.* subsp. *carotovora* and that

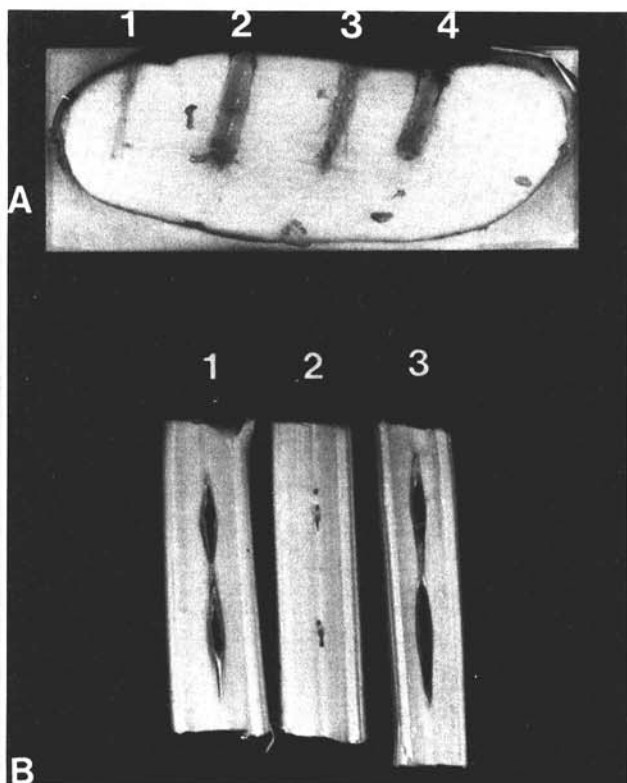


Fig. 2. Tissue maceration by *Erwinia carotovora* subsp. *carotovora* 71, an AepA⁻ strain AC5012, and AC5012 carrying an *aepA*⁺ plasmid. Whole potato tubers and celery petioles were inoculated and incubated as described in the text. After 48 hr of incubation, pathogenic reactions were recorded. **A,** Potato tuber: section 1, buffer control; section 2, AC5012(pAKC602); section 3, AC5012(pRK404); and section 4, *E. c.* subsp. *carotovora* 71(pRK404). **B,** Celery petioles: segment 1, *E. c.* subsp. *carotovora* 71(pRK404); segment 2, AC5012(pRK404); and segment 3, AC5012(pAKC602).

pAKC264 did not affect the levels of these periplasmic enzymes.

pAKC264 also stimulated the production of extracellular enzymes in wild-type strain 71 of *E. c. subsp. carotovora*. In SYG plus celery extract, the levels of Pel, Peh, Cel, and Prt were higher in extracellular fluid of *E. c. subsp. carotovora* 71(pAKC264) than in *E. c. subsp. carotovora* 71(pSF6); Pel was stimulated fivefold; Cel, fivefold; Prt, fourfold; and Peh, sevenfold (Table 3). Although the levels of Pel and Peh in cells were not significantly affected by pAKC264, Cel activity was somewhat higher in cells carrying the AepA⁺ plasmid than in the cells carrying the vector.

Previous studies with *E. c. subsp. carotovora* (Mount *et al.* 1979) have shown a requirement for a functional cAMP-CRP system for Pel production. Consistent with this requirement is the occurrence of cAMP receptor protein (CRP)-binding sites upstream of the coding regions of the various *Erwinia pel* and *peh* genes (Hinton *et al.* 1989; Reverchon *et al.* 1989; Saarihahti *et al.* 1990; Tamaki *et al.* 1988). To rule out the possibility that the stimulatory effect of the cloned DNA was due to the presence of *cya* or *crp* DNAs, pAKC602 was transformed into an *E. coli* Cya⁻ mutant (CGSC6027) and a Crp⁻ mutant (CGSC7043), and the antibiotic-resistant transformants were tested for the utilization of various sugars. The transformants carrying pAKC602, like those carrying the vector pRK404, failed to utilize arabinose, maltose, mannitol, and sorbitol. In light of a previous finding with an *E. chrysanthemi cya*⁺ DNA (Hedegaard and Danchin 1985), complementation of *E. coli* mutations should have occurred if the *E. c. subsp. carotovora* 71 *cya* and *crp* genes were present on pAKC602.

DISCUSSION

The findings of this study show that function of a gene locus, designated as *aepA*, is required for the production of extracellular enzymes in *E. c. subsp. carotovora* 71. Several lines of evidence (see above) established that the pleiotropic mutants were genetically and physiologically distinct from the bona fide Out⁻ mutants of *E. chrysanthemi* and *E. c. subsp. carotovora* (Andro *et al.* 1984; Hinton and Salmond 1987; Murata *et al.* 1990; Thurn and Chatterjee 1985). The cloned DNA restored production of the extracellular enzymes in all of the transposon-generated mutants. Restoration of the AepA phenotype is most likely due to allelic complementation. However, at this juncture we cannot eliminate the possibility of an extragenic suppression of the mutational phenotype by the cloned DNA.

The AepA⁺ plasmid stimulated enzyme production in the wild-type strain of *E. c. subsp. carotovora*. This stimulation, apparently due to a gene dosage effect, appeared to be specific for extracellular proteins, because the levels of periplasmic enzymes were not altered by the presence of the *E. c. subsp. carotovora* 71 *aepA*⁺ DNA (Table 3). Thus, the specificity of the *aepA* function appears to be similar to the *B. subtilis sensS* gene function (Wang and Doi 1990) in regulating extracellular enzyme production.

The pleiotropic phenotype in the AepA⁻ mutants did not result from a defect in the cAMP-CRP system as indicated by the utilization of various sugars whose

catabolism requires *cya* and *crp* functions (Mount *et al.* 1979; Pastan and Adhya 1976) and by the inability of the cloned *aepA*⁺ DNA to complement *cya* and *crp* mutations in *E. coli*. This is in contrast with the findings of De Crecy-Lagard *et al.* (1990) with a *Xanthomonas campestris* pv. *campestris* (Pammel) Dowson gene, *clp*, that apparently regulates various traits, including pathogenicity and extracellular enzyme production. The *clp* gene product (a CAP-like protein; CAP, catabolic activator protein) also partially restored the carbohydrate fermentation pattern of a *cya crp* strain of *E. coli*, and the *clp* gene shared some homology with the *E. coli crp* DNA. Whether the *E. c. subsp. carotovora* 71 *aepA*⁺ DNA, which specifically affects genes for extracellular proteins, has sequence homology with *clp*, *cya*, *crp*, or other genes mediating global regulation is currently under investigation.

We do not yet know how the *aepA* gene product stimulates extracellular enzyme production. However, several possibilities can be considered in explaining the *trans*-dominant effect of *aepA*. It is conceivable that the *aepA* gene product is a component of a cotranslational export machinery required in the production of extracellular proteins, including enzymes such as Pel, Cel, Prt, and Peh. In its normal state, AepA by itself or as a component of secretion machinery may stabilize the mRNA-polysome complex, thereby increasing the half-life of the message and stimulating translation of mRNA. We should note that precedence exists for translational control of exported proteins in *E. coli* (Ferro-Novick *et al.* 1984; Hall *et al.* 1983; Hengge-Aronis and Boos 1986; Watanabe *et al.* 1988). Alternatively, the *aepA* gene product may stimulate gene expression by functioning as a transcriptional activator or as a component of transcriptional machinery such as a sigma factor or a sensor protein. In light of an apparently similar effect of the *Erwinia aepA* gene and the *B. subtilis sensS* gene on the regulation of extracellular enzyme production, we can entertain the possibility that the products of these genes act in a similar fashion. Noteworthy in this context is the finding of Wang and Doi (1990) that the *senS* gene product showed partial homology with sigma factors and possessed a helix-turn-helix motif found in DNA-binding proteins. Much additional work is clearly needed to clarify the role of the *aepA* gene product, to determine the relationship between the expression of *aepA* and activation of extracellular enzyme production, and to identify the components of the signal transduction pathway.

Our demonstration that a gene locus controls the production of multiple extracellular enzymes and pathogenicity of *E. c. subsp. carotovora* 71 has an additional ramification. *E. c. subsp. carotovora* is often considered to be an opportunistic pathogen that causes soft rots in susceptible plants as an inevitable consequence of its secretion of massive quantities of pectic enzymes. The finding of the *aepA* gene, while supporting this concept, also provokes the idea that the bacterium has developed highly sophisticated mechanisms to modulate the production of extracellular enzymes during its interaction with host plants.

ACKNOWLEDGMENTS

This research was supported by grant 87-CRCR-1-2504 from the U.S. Department of Agriculture and a grant from the Food for the 21st Century

program of University of Missouri-Columbia. This article is journal series 11320 of the Missouri Agricultural Experimental Station.

We thank B. J. Bachmann (*E. coli* Genetic Stock Center, Yale University, School of Medicine, New Haven, CT), V. Iyer, N. T. Keen, N. Kleckner, and G. W. Walker for providing bacterial strains, λ phage, and plasmids. We gratefully acknowledge technical contributions of Wesley Chun and the assistance of Eva L. Chatterjee in the preparation of the manuscript.

LITERATURE CITED

- Andro, T., Chambost, J. P., Kotoujansky, A., Cattaneo, J., Bertheau, Y., Barras, F., Van Gijsegem, F., and Coleno, A. 1984. Mutants of *Erwinia chrysanthemi* defective in secretion of pectinase and cellulase. *J. Bacteriol.* 160:1199-1203.
- Barras, F., Thurn, K. K., and Chatterjee, A. K. 1986. Export of *Erwinia chrysanthemi* (EC16) protease by *Escherichia coli*. *FEMS Microbiol. Lett.* 34:343-348.
- Beaulieu, C., and Van Gijsegem, F. 1990. Identification of plant-inducible genes in *Erwinia chrysanthemi* 3937. *J. Bacteriol.* 172:1569-1575.
- Beraha, L., and Garber, E. D. 1971. Avirulence and extracellular enzymes of *Erwinia carotovora*. *Phytopathol. Z.* 70:335-344.
- Brickman, E., Soll, L., and Beckwith, J. 1973. Genetic characterization of mutations which affect catabolite-sensitive operons in *Escherichia coli*, including deletions of the gene for adenyl cyclase. *J. Bacteriol.* 116:582-587.
- Castilho, B. A., Olfson, P., and Casadaban, M. J. 1984. Plasmid insertion mutagenesis and *lac* gene fusion with mini-Mu bacteriophage transposons. *J. Bacteriol.* 158:488-495.
- Chatterjee, A. K. 1980. Acceptance by *Erwinia* spp. of R plasmid R68.45 and its ability to mobilize the chromosome of *Erwinia chrysanthemi*. *J. Bacteriol.* 142:111-119.
- Chatterjee, A. K., and Vidaver, A. K. 1986. Genetics of pathogenicity factors: Application to phytopathogenic bacteria. Pages 1-224 in: *Advances in Plant Pathology*, Vol. 4. D. S. Ingram and P. H. Williams, eds. Academic Press, London.
- Chatterjee, A. K., Thurn, K. K., and Feese, D. A. 1983. Tn5-induced mutations in the enterobacterial phytopathogen *Erwinia chrysanthemi*. *Appl. Environ. Microbiol.* 45:644-650.
- Chatterjee, A. K., Ross, L. M., McEvoy, J. L., and Thurn, K. K. 1985. pULB113, an RP4::mini-Mu plasmid, mediates chromosomal mobilization and R-prime formation in *Erwinia amylovora*, *Erwinia chrysanthemi*, and subspecies of *Erwinia carotovora*. *Appl. Environ. Microbiol.* 50:1-9.
- Chatterjee, A. K., McEvoy, J. L., Murata, H., and Collmer, A. Regulation of the production of pectinases and other extracellular enzymes in the soft-rotting *Erwinia* spp. In: *Molecular Strategies of Pathogens and Host Plants*. S. S. Patil, S. Ouchi, D. Mills, and C. Vance, eds. Springer-Verlag, Berlin. In press.
- Clement, J.-M., Perrin, D., and Hedgpeh, J. 1982. Analysis of λ receptor and β -lactamase synthesis and export using cloned genes in a minicell system. *Mol. Gen. Genet.* 185:302-310.
- Collmer, A., and Keen, N. T. 1986. The role of pectic enzymes in plant pathogenesis. *Annu. Rev. Phytopathol.* 24:383-409.
- Crouse, G. F., Frischauf, A., and Lehrach, H. 1983. An integrated and simplified approach to cloning into plasmids and single-stranded phages. *Methods Enzymol.* 101:78-89.
- Daza, A., Gil, J. A., Vigal, T., and Martin, J. F. 1990. Cloning and characterization of a gene of *Streptomyces griseus* that increases production of extracellular enzymes in several species of *Streptomyces*. *Mol. Gen. Genet.* 222:384-392.
- De Crecy-Lagard, V., Glaser, P., Lejeune, P., Sismeiro, O., Barber, C. E., Daniels, M. J., and Danchin, A. 1990. A *Xanthomonas campestris* pv. *campestris* protein similar to catabolite activation factor is involved in regulation of phytopathogenicity. *J. Bacteriol.* 172:5877-5883.
- Ditta, G., Schmidhauser, T., Yakobson, E., Lu, P., Liang, X.-W., Finlay, D. R., Guiney, D., and Helinski, D. R. 1985. Plasmids related to the broad host range vector, pRK290, useful for gene cloning and for monitoring gene expression. *Plasmid* 13:149-153.
- Ferro-Novick, S., Honma, M., and Beckwith, J. 1984. The product of gene *secC* is involved in the synthesis of exported proteins in *E. coli*. *Cell* 38:211-217.
- Figurski, D. H., and Helinski, D. R. 1979. Replication of an origin-containing derivative of plasmid RK2 dependent on a plasmid function provided in *trans*. *Proc. Natl. Acad. Sci. USA* 76:1648-1652.
- Hall, M. N., Gabay, J., and Schwartz, M. 1983. Evidence for a coupling of synthesis and export of an outer membrane protein in *Escherichia coli*. *EMBO J.* 2:15-19.
- He, S. Y., Lindeberg, M., Chatterjee, A. K., and Collmer, A. Cloned *Erwinia chrysanthemi* *out* genes enable *Escherichia coli* to selectively secrete a diverse family of heterologous proteins to its milieu. *Proc. Natl. Acad. Sci. USA*. In press.
- Hedegaard, L., and Danchin, A. 1985. The *cya* gene region of *Erwinia chrysanthemi* B374: Organisation and gene products. *Mol. Gen. Genet.* 201:38-42.
- Hengge-Aronis, R., and Boos, W. 1986. Translational control of exported proteins in *Escherichia coli*. *J. Bacteriol.* 167:462-466.
- Henner, D. J., Ferrari, E., Perego, M., and Hoch, J. A. 1988. Location of the targets of the *hpr-97*, *sacU32(Hy)*, and *sacQ36(Hy)* mutations in upstream regions of the subtilisin promoter. *J. Bacteriol.* 170:296-300.
- Hinton, J. C. D., and Salmond, G. P. C. 1987. Use of TnphoA to enrich for extracellular enzyme mutants of *Erwinia carotovora* subspecies *carotovora*. *Mol. Microbiol.* 1:381-386.
- Hinton, J. C. D., Sidebotham, J. M., Gill, D. R., and Salmond, G. P. C. 1989. Extracellular and periplasmic isoenzymes of pectate lyase from *Erwinia carotovora* subspecies *carotovora* belong to different gene families. *Mol. Microbiol.* 3:1785-1795.
- Hinton, J. C. D., Gill, D. R., Lalo, D., Plastow, G. S., and Salmond, G. P. C. 1990. Sequence of the *peh* gene of *Erwinia carotovora*: Homology between *Erwinia* and plant enzymes. *Mol. Microbiol.* 4:1029-1036.
- Hirsch, P. R., and Beringer, J. E. 1984. A physical map of pPHIJI and pJB4JI. *Plasmid* 12:139-141.
- Ji, J., Hugouvieux-Cotte-Pattat, N., and Robert-Baudouy, J. 1989. Molecular cloning of the *outJ* gene involved in pectate lyase secretion by *Erwinia chrysanthemi*. *Mol. Microbiol.* 3:285-293.
- Kelemu, S., and Collmer, A. 1990. A new set of *Erwinia chrysanthemi* pectic enzymes produced during growth on plant material. *Phytopathology* 80:983.
- Kotoujansky, A. 1987. Molecular genetics of pathogenesis by soft-rot erwinias. *Annu. Rev. Phytopathol.* 25:405-430.
- Lei, S.-P., Lin, H.-C., Heffernan, L., and Wilcox, G. 1985. Cloning of the pectate lyase genes from *Erwinia carotovora* and their expression in *Escherichia coli*. *Gene* 35:63-70.
- Lei, S.-P., Lin, H.-C., Wang, S.-S., and Wilcox, G. 1988. Characterization of the *Erwinia carotovora pelA* gene and its product pectate lyase A. *Gene* 62:159-164.
- Long, S., McCune, S., and Walker, G. C. 1988. Symbiotic loci of *Rhizobium meliloti* identified by random TnphoA mutagenesis. *J. Bacteriol.* 170:4257-4265.
- Maniatis, T., Fritsch, E. F., and Sambrook, J. 1982. *Molecular Cloning: A Laboratory Manual*. Cold Spring Harbor Laboratory, Cold Spring Harbor, NY. 545 pp.
- McEvoy, J. L., Murata, H., and Chatterjee, A. K. 1990. Molecular cloning and characterization of an *Erwinia carotovora* subsp. *carotovora* pectin lyase gene that responds to DNA-damaging agents. *J. Bacteriol.* 172:3284-3289.
- Metcalfe, W. W., Steed, P. M., and Wanner, B. L. 1990. Identification of phosphate starvation-inducible genes in *Escherichia coli* K-12 by DNA sequence analysis of *psi::lacZ*(Mu dI) transcriptional fusions. *J. Bacteriol.* 172:3191-3200.
- Mount, M. S., Bateman, D. F., and Basham, H. G. 1970. Induction of electrolyte loss, tissue maceration, and cellular death of potato tissue by an endopolygalacturonate *trans*-eliminase. *Phytopathology* 60:924-931.
- Mount, M. S., Berman, P. M., Mortlock, R. P., and Hubbard, J. P. 1979. Regulation of endopolygalacturonate transeliminase in an adenosine 3',5'-cyclic monophosphate-deficient mutant of *Erwinia carotovora*. *Phytopathology* 69:117-120.
- Murata, H., and Chatterjee, A. K. 1990a. Expression of the *Erwinia carotovora* subsp. *carotovora* (Ecc) *aep* gene is required in the production of extracellular proteins. *Phytopathology* 80:984.
- Murata, H., and Chatterjee, A. K. 1990b. Genetic evidence for a trans-acting factor of *Erwinia carotovora* subsp. *carotovora* (Ecc) that stimulates the production of extracellular degradative enzymes. *Phytopathology* 80:1037.
- Murata, H., Fons, M., Chatterjee, A., Collmer, A., and Chatterjee, A. K.

1990. Characterization of transposon insertion *Out*⁻ mutants of *Erwinia carotovora* subsp. *carotovora* defective in enzyme export and of a DNA segment that complements *out* mutations in *E. carotovora* subsp. *carotovora*, *E. carotovora* subsp. *atroseptica*, and *Erwinia chrysanthemi*. *J. Bacteriol.* 172:2970-2978.
- Pastan, I., and Adhya, S. 1976. Cyclic adenosine 5'-monophosphate in *Escherichia coli*. *Bacteriol. Rev.* 40:527-551.
- Peng, H.-L., Novick, R. P., Kreiswirth, B., Kornblum, J., and Schlievert, P. 1988. Cloning, characterization, and sequencing of an accessory gene regulator (*agr*) in *Staphylococcus aureus*. *J. Bacteriol.* 170:4365-4372.
- Recsei, P., Kreiswirth, B., O'Reilly, M., Schlievert, P., Gruss, A., and Novick, R. P. 1985. Regulation of exoprotein gene expression in *Staphylococcus aureus* by *agr*. *Mol. Gen. Genet.* 202:58-61.
- Reverchon, S., Huang, Y., Bourson, C., and Robert-Baudouy, J. 1989. Nucleotide sequences of the *Erwinia chrysanthemi* *ogl* and *pelE* genes negatively regulated by the *kdgR* gene product. *Gene* 85:125-134.
- Roberts, D. P., Berman, P. M., Allen, C., Stromberg, V. K., Lacy, G. H., and Mount, M. S. 1986. Requirement for two or more *Erwinia carotovora* subsp. *carotovora* pectolytic gene products for maceration of potato tuber tissue by *Escherichia coli*. *J. Bacteriol.* 167:279-284.
- Saarilahti, H. T., Heino, P., Pakkanen, R., Kalkkinen, N., Palva, I., and Palva, E. T. 1990. Structural analysis of the *pehA* gene and characterization of its protein product, endopolygalacturonase, of *Erwinia carotovora* subspecies *carotovora*. *Mol. Microbiol.* 4:1037-1044.
- Salmond, G. P. C., Hinton, J. C. D., Gill, D. R., and Perombelon, M. C. M. 1986. Transposon mutagenesis of *Erwinia* using phage λ vectors. *Mol. Gen. Genet.* 203:524-528.
- Selvaraj, G., Fong, Y. C., and Iyer, V. N. 1984. A portable DNA sequence carrying the cohesive site (*cos*) of bacteriophage λ and the *mob* (mobilization) region of the broad-host-range plasmid RK2: A module for the construction of new cosmids. *Gene* 32:235-241.
- Tamaki, S. J., Gold, S., Robeson, M., Manulis, S., and Keen, N. T. 1988. Structure and organization of the *pel* genes from *Erwinia chrysanthemi* EC16. *J. Bacteriol.* 170:3468-3478.
- Thurn, K. K., and Chatterjee, A. K. 1985. Single-site chromosomal Tn5 insertions affect the export of pectolytic and cellulolytic enzymes in *Erwinia chrysanthemi* EC16. *Appl. Environ. Microbiol.* 50:894-898.
- Van Gijsegem, F. 1989. Relationship between the *pel* genes of the *pelADE* cluster in *Erwinia chrysanthemi* strain B374. *Mol. Microbiol.* 3:1415-1424.
- Wang, L.-F., and Doi, R. H. 1990. Complex character of *senS*, a novel gene regulating expression of extracellular-protein genes of *Bacillus subtilis*. *J. Bacteriol.* 172:1939-1947.
- Watanabe, T., Hayashi, S., and Wu, H. C. 1988. Synthesis and export of the outer membrane lipoprotein in *Escherichia coli* mutants defective in generalized protein export. *J. Bacteriol.* 170:4001-4007.
- Way, J. C., Davis, M. A., Morisato, D., Roberts, D. E., and Kleckner, N. 1984. New Tn10 derivatives for transposon mutagenesis and for construction of *lacZ* operon fusions by transposition. *Gene* 32:369-379.
- Willis, J. W., Engwall, J. K., and Chatterjee, A. K. 1987. Cloning of genes for *Erwinia carotovora* subsp. *carotovora* pectolytic enzymes and further characterization of the polygalacturonases. *Phytopathology* 77:1199-1205.
- Yang, Z., Cramer, C. L., and Lacy, G. H. 1989. System for simultaneous study of bacterial and plant gene expression in soft rot of potato. *Mol. Plant-Microbe Interact.* 2:195-201.
- Zink, R. T., Kemble, R. J., and Chatterjee, A. K. 1984. Transposon Tn5 mutagenesis in *Erwinia carotovora* subsp. *carotovora* and *E. carotovora* subsp. *atroseptica*. *J. Bacteriol.* 157:809-814.