

Molecular Cloning of Pectinase Genes from *Erwinia carotovora* Subspecies *carotovora* (Strain SCRI193)*

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Abstract

Several members of the genus *Erwinia* are phytopathogenic causing soft rot, black leg or stem rot of various plants. The effects on the plant are probably due to the concerted action of extracellular enzymes, including pectinases, cellulases and proteases. As part of our study of the secretion process in Gram negative bacteria we have cloned five structural genes for various pectinases from *Erwinia carotovora*. These have been preliminarily characterized and include genes encoding the three main pectolytic enzymes produced under laboratory conditions: two pectate lyases and a polygalacturonase. The other two enzymes are also pectate lyases and appear to have minor activities in this strain.

Keywords: *Erwinia carotovora*, molecular cloning, pectate lyase, polygalacturonase

Abbreviations: spp, species; EDTA, ethylenediaminetetraacetic acid; SDS, sodium dodecyl sulphate; Kb, Kilobase pairs

*Reviewed

Introduction

Erwinia carotovora subspecies *carotovora* (*Ecc*) is an aetiological agent of soft rot in stored crops such as potato and carrot. The rotting of potato tubers can be a significant economic problem for some companies who may store potatoes up to 10 months after harvesting prior to processing. It has been known for sometime that factors such as temperature, pO₂, and humidity are important in predisposing the potato tuber to *Ecc*-mediated soft rot. However, it is not always easy, or economical, to control such factors sufficiently on a large scale in order to prevent rot. Consequently attention has turned towards a direct, detailed analysis of the major determinants of soft rot biodeterioration, namely extracellular macerating enzymes. *Erwinia* spp. make a range of pectinolytic, cellulolytic and proteolytic enzymes which are exported into the extracellular medium. This feature of *Erwinia* makes it doubly interesting to study because it is not only a phytopathogen, but also an excellent model system for the investigation of protein secretion. As part of our analysis of the determinants of *Ecc* pathogenicity, and the mechanism of protein export we have attempted to define the spectrum of pectolytic enzyme activities in one strain of *Ecc* (SCRI193). This strain has been chosen because it is amenable to a range of genetic techniques, including transformation (Hinton et al., 1985) and transposon mutagenesis (Salmond et al., 1986), thereby allowing a molecular analysis of all of the features we are investigating. Our initial aim was to identify a variety of enzyme activities in culture supernatants and then to clone (in *Escherichia coli*) the structural genes for such enzymes. The long term aim is to analyze the structure of each enzyme and the mechanisms involved in the export process.

1. Materials and Methods

Bacterial strains

Erwinia carotovora subspecies *carotovora* SCRI193 (Hinton et al., 1985), *Erwinia chrysanthemi* B374 (National Collection of Plant Pathogenic Bacteria 898). *Escherichia coli* NM522 ([lac-pro]⁺ hsd Δ 5:F' lacZ Δ M151acI^q) (Gough and Murray, 1983). *Erwinia* strains were grown at 30°C in Luria broth supplemented with 0.5% sodium polygalacturonate in order to induce the synthesis of pectinases. *E. coli* strains were grown at 37°C in Luria broth with 100 µg/ml ampicillin when appropriate.

Preparation of DNA from *Erwinia carotovora*

A 20 ml culture of *Ecc* SCRI193 was harvested and resuspended in 2 ml of 0.15 M NaCl, 0.1 M EDTA pH 8 containing 2 mg/ml lysozyme, taken

through two cycles of freezing and thawing in a dry ice/methanol bath and incubated at 37°C for 15 min. The cells were lysed by adding 2 ml of 1% SDS, 0.1 M NaCl, 0.1 M Tris HCl pH 9. 10 µl of protease K (10 mg/ml) were added and the incubation continued for 2 hr. The lysate was gently extracted several times with an equal volume of phenol followed by chloroform. DNA was then spooled onto a glass rod after the addition of an equal volume of absolute ethanol, blotted dry on tissues and rehydrated in 0.5 ml 1 mM EDTA, 10 mM Tris HCl pH 7.5

Construction of genomic libraries

Chromosomal DNA was digested to completion with *Bam*HI, *Eco*RI and *Hind*III and ligated to correspondingly digested pUC8 treated with calf intestinal alkaline phosphatase. The same methods were used to construct an *Hind*III library in pBR322. Ligated DNAs were transformed into competent NM522 and plated onto M9 media supplemented with 0.5% glycerol, 0.1% yeast extract and 0.5% sodium polygalacturonate containing 100 µg/ml ampicillin (PGA plates). Transformants were replicated onto Luria agar plates (containing 100 µg/ml ampicillin and 0.5% glucose). Pectinase-positive clones were revealed by the appearance of a white halo after flooding the transformation plates with 7.5% cupric acetate. Potential clones were then rescued from the replicates and plasmid DNA prepared by the rapid isolation procedure of Holmes and Quigley (1981).

Enzyme assays

Pectate lyase (PL) activity was assayed by monitoring the increase in absorbance at 235 nm in reaction mixtures containing sodium polygalacturonate or pectin (Bulmers Type 104, 70% methylated). Assays contained 2.4 mg/ml substrate in 50 mM Tris HCl pH 7.6 and were carried out at room temperature. Polygalacturonase (PG) activity was assayed by measuring the production of reducing groups using dinitrosalicylic acid (Rombouts and Pilnik, 1980). Reaction mixtures contained 2.4 mg/ml substrate in 40 mM sodium phosphate buffer pH 7, 1 mM EDTA and were incubated at 30°C. β -lactamase was assayed by monitoring the increase in absorbance at 490 nm in reaction mixtures containing the chromogenic β -lactam nitrocefin (O'Callaghan et al., 1972) in 10 mM sodium phosphate buffer pH 7 and were carried out at room temperature.

Table 1. Pectinase activities of *Ecc* SCRI193 and strains of *E. coli* carrying various pectinase clones.

Strain	Pectate Lyase		Pectin/50 mM no add.	Tris pH 8.6 1.5 mM CaCl ₂	% Activity in supernatant PL β -lactamase	
	PGA/50 mM no add.	Tris pH 8.6 1.5 mM CaCl ₂			no add.	Tris pH 8.6 1.5 mM CaCl ₂
<i>Ecc</i>	0.96 ¹	-	-	-	94	
B1	0.15	1.85	7.17	-	34	50
B6	0.1				6	25
E4	Nil					
H2	0.69	0.66	1.02	0.9	29	54
H5	1.45	1.34	2.12	nd ²	50	-

¹ ODU₂₃₅/min/ml

² Rapid initial rate which rapidly decayed

Strain	Polygalacturonase PGA/40 mM PO ₄ pH 7 1 mM EDTA	% Activity in supernatant β -lactamase	
		PG	
<i>Ecc</i>	0.026 ¹		
E4	0.166	52	48

¹ ODU₅₄₀/min/ml

H2 (B6) and H5 showed little response. However, all of the enzymes were completely inhibited by the addition of EDTA indicating that they all have a requirement for divalent cations as would be expected for lyases (Rombouts and Pilnik, 1980). We have not investigated this aspect any further nor have we established whether these enzymes are endo- or exo-pectinases.

The enzymes were then characterized further by isoelectric focusing followed by blotting with an agarose-substrate gel in order to detect pectinase activities (not shown). Supernatants from *Ecc* SCRI193 grown in the presence of 0.5% PGA contain two major pectate lyases with approximate pI's of 9.1 and 9.3 (and migrating with a similar pI to PL's *d* and *e* of *Ech* B374 (van Gijsegem et al., 1985) used for comparison) and a single polygalacturonase with an approximate pI of 9.3. Minor activities can sometimes be observed if the blotting time is extended, however there were no major activities corresponding to PL's *b* and *c* of *Ech* B374. The clone E4 was confirmed to encode the polygalacturonase by this method. Interestingly the two major

pectate lyase activities appear to be encoded by a single clone B1. Although we have not yet shown that these enzymes are encoded by two genes we feel that this possibility is the most likely. This would be analogous to the situation in *Ech* where the genes for the two major PL's of this pI, *d* and *e* are linked (van Gijsegem et al., 1985; Reverchon et al., 1985). Furthermore Lei et al. (1985) have found three closely linked pectate lyase genes in a different strain of *Ecc*, (two of the lyases had pI's of 9.4 and the third a pI of 9.1). This suggests that the other clones (H2 and H5) must encode lyases with minor activities in *Ecc* or else further lyases with pI's around 9 as was found by Lei et al. (1985). The former seems to be the case for H2 which encodes a lyase with a pI of 8 (migrating between *b*, pI 7.8 and *c*, pI 8.2 of *Ech* B374). We have not yet identified the enzyme encoded by H5 in this system.

Conclusion

We have succeeded in isolating five pectinase genes from *Ecc* SCRI193: four pectate lyase genes and one polygalacturonase gene. We aim to determine the DNA sequence of these genes in order to search for potential domains of homology (or structural similarities) in the gene and protein sequences. Such homologies between the pectinases and cellulases and proteases might relate to common mechanisms of secretion for these enzymes as has been suggested by genetic studies (Andro et al., 1984; Thurn and Chatterjee, 1985; G.P.C. Salmond et al., pers. comm., 1986). This molecular approach together with the genetic approaches available will enable us to dissect the secretion process in this strain of *Erwinia carotovora*.

Acknowledgement

We thank Alan Boyd for useful advice and discussion.

REFERENCES

- 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.
- Collmer, A., Schoedel, C., Roeder, D.L., Ried, J.L., and Rissler, J.F. 1985. Molecular cloning in *Escherichia coli* of *Erwinia chrysanthemi* genes encoding multiple forms of pectate lyase. *J. Bacteriol.* **161**, 913-920.

- Gough, J.A. and Murray, N.E. 1983. Sequence diversity among related genes for recognition of specific targets in DNA molecules. *J. Mol. Biol.* **166**, 1-19.
- Hinton, J.C.D., Perombelon, M.C.M., and Salmond, G.P.C. 1985. Efficient transformation of *Erwinia carotovora* subsp. *carotovora* and *Erwinia carotovora* subsp. *atroseptica*. *J. Bacteriol.* **161**, 786-788.
- Holmes, D.S. and Quigley, M. 1981. A rapid boiling method for the preparation of bacterial plasmids. *Anal. Biochem.* **114**, 193-197.
- 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.
- O'Callaghan, C.H., Morris, A., Kirby, S., and Shingler, A.H. 1972. Novel method for detection of β -lactamases by using a chromogenic cephalosporin substrate. *Antimicrob Agents Chemother.* **1**, 283-288.
- Ried, J.L. and Collmer, A. 1985. Activity stain for rapid characterization of pectic enzymes in isoelectric focusing and sodium dodecyl sulfate-polyacrylamide gels. *Appl. Environ. Microbiol.* **50**, 615-622.
- Reverchon, S., Hugouvieux-Cotte-Pattat, N., and Robert-Baudouy, J. 1985. Cloning of genes encoding pectolytic enzymes from a genomic library of the phytopathogenic bacterium, *Erwinia chrysanthemi*. *Gene* **35**, 121-130.
- Rombouts, F.M. and Pilnik, W. Pectic enzymes. In: *Microbial Enzymes and Bioconversions*. A.H. Rose, ed. (Economic Microbiology. Vol. 5). Academic Press, London, 1980, pp. 227-282.
- Salmond, G.P.C., Hinton, J.C.D., Gill, D.R., and Perombelon, M.C.M. 1986. Transposon mutagenesis of *Erwinia*: Phage λ as a suicide vector. *Molec. Gen. Genet.* (in press).
- Van Gijsegem, F., Toussaint, A., and Schoonejans, E. 1985. *In vivo* cloning of the pectate lyase and cellulase genes of *Erwinia chrysanthemi*. *EMBO J.* **4**, 787-792.
- Zinc, R.T. and Chatterjee, A.K. 1985. Cloning and expression in *Escherichia coli* of pectinase genes of *Erwinia carotovora* subsp. *carotovora*. *Appl. Environ. Microbiol.* **49**, 714-717.