PECTIN TRANS-ELIMINASE

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Ver Since the discovery of pectin by Vanquelin in 1790,1 pectic substances have interested both the 'pure' scientist as well as the technologist. The pectic substances are carbohydrate derivatives of plant origin. They are complex and colloidal in nature and mostly comprise of anhydrogalacturonic acid units linked together in a(1-4) glycosidic linkage to form a chain-like structure of polygalacturonic acid units. While it is true that D-galacturonic acid is the main constituent of pectin, most natural pectins contain about 12% ester methoxyl. A hypothetical pectin with 75% of its carboxyl groups esterified and polygalacturonic acid are depicted in Fig. 1.

PECTIN (WITHOUT END UNITS)

Fig. 1. The Pectic Substances

The complex nature of molecules comprising these substances has not only caused the confusion in their chemistry but has been mainly responsible for the delay in the accumulation of knowledge pertaining to those enzymes which have the power to break these molecules down. However, the immense importance of pectic substances and their enzymes has helped to maintain the tempo of work thereon and it is the purpose of this review to briefly trace the literature on pectic enzymes in general and to discuss in detail what has recently been the subject of a new discovery, viz., pectin transeliminase.

Until 1960, the well described pectinolytic enzymes produced by micro-organisms were all thought to be hydrolytic in nature.² The enzyme polygalacturonase (PG) facilitates the hydrolysis of a(1-4) glycosidic linkages in the polygalacturonic acid skeleton of pectic or pectinic acids with the resulting formation of polygalacturonic acids of small molecular sizes and of monogalacturonic acid. Pectic acid depolymerase (DP) differs from PG in having a sharp pH optimum at 4.5 and to get activated

by sodium chloride. Pectin methyl esterase (PME) catalyses the cleavage of methyl ester groups in pectinic acids and pectins to yield pectic acids and methanol. The mode of action of PG and PME on macromolecular pectin is illustrated in Fig. 2.

ACTION OF METHYLESTERASE

PECTIC ACID

D.GALACTURONIC ACID

PECTINIC ACID

Fig. 2. Action of PG and PME

Investigations on the alkaline degradation of pectin by Neukom and Deuel³ led to the discovery of the new pectinolytic enzyme pectin The mechanism trans-eliminase. degradation was explained on the lines of " β -dealkoxylation" according to which, the removal of activated hydrogen at C-5 resulted in the formation of a double bond between C₄ and C_5 and the cleavage in β -position to the ester carbonyl group. Subsequently, Albersheim, Neukom and Deuel4 observed that such a 'transelimination' also took place on heating pectin in a buffer solution at pH 6.8. The products of degradation showed an absorption maximum at 230-235 mm, reacted with thiobarbituric acid to give a coloured product having an absorption maximum at 547 mm and formed oxalic acid with ozone. These characteristics provided additional evidence for a trans-elimination reaction consequent to the removal of H atom at C-5, formation of a double bond between C-4 and C-5, and elimination of glycosidic residue at C-4. Later, Albersheim, Neukom and Deuel⁵ demonstrated the presence, in commercial

pectinase preparation, "Pectasin-R-10", of an enzyme that brought about such a trans-elimination reaction and designated it as pectin purification of The this trans-eliminase. enzyme was achieved by Albersheim and Killias⁶; they also described its properties. Subsequently, Albersheim was able to obtain it in a crystalline form.7 According to the substrate specificity exhibited, these enzymes have been now designated pectin trans-eliminase (PTE) or polygalacturonic acid trans-eliminase (PATE) (Fig. 3). The product of this reaction has been fully characterised by Hasegawa and Nagel⁸ as 4-0 a-D-(4, 5-dehydrogalacturonosyl) -D-galacturonic acid (Fig. 4).

FIG. 3. Trans-Eliminase Enzymes

ACTION OF POLYGALACTURONIC ACID TRANS-ELIMINASE

FIG. 4. Structure of Dimer

It has now become apparent that a number of microbes may be producing trans-eliminases and it is reasonable to assume that many of the polygalacturonases described prior to 1960 (before the discovery of PTE by the Swiss group of workers) may have to be recharacterized. The polymethyl galacturonase from commercial pectinase—"Hydrolase"—described by Seegmiller and Jansen and pectin depolymerase from Neurospora crassa described by Roboz, Barratt and Tatum are cases in point.

Since the discovery of the enzyme transeliminase, the principal mechanism involved in bacterial degradation has been shown to be due to the trans-eliminative cleavage of glycosidic linkages. While investigating the pectic enzymes of Bacillus polymyra, Nagel¹² observed an altered digalacturonic acid differing from the normal compound. The factors affecting growth and enzyme secretion by this organism

have since been studied elaborately by Nagel and Vaughn^{13–15} and the trans-eliminative mechanism was shown to occur in bacterial degradation of pectic substances. Starr and Moran, ^{16–17} working on the enzymes of the phytopathogenic soft rot bacteria as Erwinia and Bacillus, recorded results substantiating these findings. Starr and Nasuno¹⁸ subsequently indicated the presence of an extracellular enzyme in Xanthomonas sp. which has a specific action on polygalacturonic acid rather than pectin.

Recent work carried out in this laboratory on the culture filtrates of some bacteria characterized as Xanthomonas, the enzyme PTE/ PATE was not detected. 19,20 However, data obtained seem to suggest that these cultures, in addition to PG, possess a hitherto unrecognized mode of action on pectic substances, since all the breakdown products could not quantitatively account for those that might have resulted from the action of PG alone. The enzyme PME was not detected in their On the other hand, the enzyme trans-eliminase has been demonstrated in bacterial cultures of Corynebacterium barkeri,21 Flavobacterium sp.,21 Micrococcus sp.22,23 and Arthrobacter sp.²⁴ Preiss and Ashwell²⁵ reported the presence of this enzyme Pseudomonas sp. and Hsu and Vaughn²⁶ in Aeromonas liquefaciens. Recently, Nagel and Anderson²⁷ studied the action of the enzymes of Bacillus polymyxa on normal and unsaturated oligogalacturonic acids while Dave and Vaughn²⁸ gave evidence to indicate that, in constrast to other polygalacturonic acid trans-eliminases studied, the PATE of Bacillus pumilus produced a large quantity of unsaturated trigalacturonic acid.

MacMillan and Vaughn²⁹ have discovered yet another bacterial polygalacturonic acid transeliminase in a strain of Clostridium multifermentans; the enzyme attacked the terminal groups to produce a preponderance of unsaturated digalacturonic acid. The organism was also reported to produce the demethylating enzyme so that, grown in vitro or in nature, the cultures can bring about the degradation of pectin as far as the dimer stage. However, it appears to be unable to produre the digalacturonase; in other words, it is unlike B. polymyx a^{12} in so far as it does not destroy completely the pectic material, though it may share several other properties with it. Such a terminally acting PATE was also observed in certain Erwinia sp. by Okamoto and Ozawa.30

The presence of the enzyme PTE in different species of the mould Aspergillus was demonstrated by Edstrom and Phaff.³¹ These authors purified the enzyme PTE from the culture fluid of Aspergillus fonsecœus and demonstrated that the enzyme was specific for pectin and differed in several respects from a PATE produced by certain bacteria. The enzyme had an optimum pH at 5·2 and was not absolutely dependent on Ca ions unlike the bacterial transeliminases. These authors further provided the course of action by A. fonsecœus PTE on pectin and on certain oligogalacturonide methyl esters.³²

That the actinomycetes could be considered as a potent group for the production of pectinolytic enzymes have been proved by the successful investigations in this laboratory.^{33,34} Simultaneously Kaiser³⁵ reported the isolation of pectinolytic actinomycetes by exploitation of the method developed by Wieringa.³⁶ It was demonstrated by Agate²¹ and Agate, Bilimoria and Bhat³⁷ that several species of Streptomyces possessed trans-eliminase activity. The streptomycetal trans-eliminase in general resembled the bacterial enzymes, but differed from the fungal trans-eliminase.

A PTE (and PATE) was detected in crude acetone powder prepared from protozoa occurring in the rets of Calotropis and Hibiscus plant straws.³⁸ Subsequently, it was demonstrated conclusively in the protozoan species Plagiopyla.²¹ A PATE, unlike the PTE in S. viridochromogenes was demonstrated in an Epistylis sp. occurring in sewage and sewage products.³⁹ Later on, two other protozoal species from the same ecosystem, viz., Vorticella, Carchesium were found to contain this enzyme²¹ and the inclusion of protozoa in the already existing list of micro-organisms producing PTE was a contribution in the field from this laboratory.

Although these interesting trans-eliminases have been found to be produced by bacteria, actinomycetes, moulds and even protozoa, to date in our laboratory we have not been able to detect this enzyme in the large collection of pectinolytic yeasts that we have, including such pectinolytic species as Cryptococcus laurentii, Saccharomyces kluyveri, Cryptococcus diffluens, Rhodotorula glutinis, R. flava, etc. 19-21-24 This finds confirmation in the fact that Vaughn also could not detect the presence of any trans-eliminase enzyme in the pectinolytic yeasts, e.g., in Saccharomyces fragilis or Rhodotorula glutinis.9

The discovery of this enzyme has broken new grounds in the field of microbial pectinolytic

enzymes and it is hoped that a better understanding of these enzymes would help to bring a little order into the chaos prevailing in the field of pectin chemistry.

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