

The apparent Km values of the purified xylanase I and II were 7.0 and 2.5 mg/ml respectively against oat spelt xylan. The purified enzyme is supposed to endo-type xylanase by the of TLC analysis of reaction products.

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Development of Method of Multiplex PCR for Specific Detection of *L.ivanovii* in Food

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Study of development of multiplex PCR method was conducted to develop a rapid and precise detection method in foods using specific primer of *iap*(invasion-associated protein) gene encoding p60 protein that commonly exist in all of *Listeria spp.* We ensured that *L. ivanovii* and other *Listeria* species could be detected with the multiplex PCR method. We should demonstrate that this method is superior to biochemical detection by re-inspecting biochemically classified species. This multiplex PCR is simple, precise and economic. Also, this detection method have various application potential.

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Purification of the *E.coli* Expressed p60 Protein from *Listeria welshimerii* by Amylose Resin Based Affinity Chromatography

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The *Listeria welshimerii* is an animal and human pathogen and its p60 protein is a major extracellular protein, which is encoded in *iap*(invasion associated protein) gene. These proteins are believed to be involved in the invasion of these bacteria into their host cells. To produce p60 in *E.coli*, the *iap* gene was recombinantly cloned and overexpressed. A purification protocol was developed for MBP(maltose binding protein)-p60 fusion protein by amylose-resin based affinity chromatography. The purified MBP-p60 was detected either as denaturated or neutralized form using a specific p60 monoclonal antibody. The method might be an easy alternative to common purification protocols of p60 from *Listeria spp.* for antibody production.

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Enhanced Production of Avermectin B1a with *Streptomyces avermitilis* by Medium Optimization and Glucose Feeding

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Avermectin is a group of potent anthelmintic and insecticidal antibiotics produced by *Streptomyces avermitilis*. Productivity of avermectin B1a was enhanced by medium optimization and intermittent glucose feeding. Response

surface methodology was used to optimize the concentration of nitrogen sources. A strong stimulation on avermectin B1a production in *S. avermitilis* was observed by 3% glucose feeding at the time of residual glucose being 1%. The avermectin B1a productivity could be further improved by another glucose feeding at 206 hour of cultivation. Avermectin B1a titer was increased by 86.3% and the proportion of avermectin B1a in the total avermectins was increased from 38% to 45% through medium optimization and glucose feeding process. These results would be very useful for enhancing productivity of avermectin B1a in up-scaled processes.

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Development of a Simplified Purification Process of Avermectin B1a from Fermentation Broth of *Streptomyces avermitilis* YA99-40

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Avermectin B1a is the most potent anthelmintic and insecticidal antibiotic among eight avermectin derivatives produced by *Streptomyces avermitilis*. To circumvent difficulty in the purification of avermectin B1a from the other avermectins, especially avermectin B1b, a new *S. avermitilis* mutant YA99-40 was developed by mutagenesis through ultraviolet light irradiation and protoplast fusion. Avermectins were extracted with 5:1 mixture of ethylacetate and acetone, then applied to a column packed with XAD-2000 resin. The avermectins were eluted with 95% methanol. The pooled avermectins B1a was concentrated and further purified by

crystallization. Analysis result of the purified avermectin B1a showed that the proportion of avermectin B1 was more than 95%. This process was simple and applicable to up-scaled purification processes

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Synthesis of 7a-hydroxycephalosporin C by Immobilized Enzyme

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In the chemical synthesis of 7a-methoxylated cephalosporins from cephalosporins substrate, difficulties have been experienced such as low conversion yield and generation of toxic wastes. Enzymatic 7a-methoxylation has been studied to alleviate these problems. The methoxylation of cephalosporin C by enzymes was reported to proceed by two step reactions which involves the formation of 7a-hydroxy cephalosporin C with subsequent methylation to yield 7a-methoxycephalosporin C. We have attempted to synthesize 7a-hydroxycephalosporin C from cephalosporin C as the first step with immobilized cephalosporin 7a-hydroxylase. First of all, the conversion of cephalosporin C to 7a-hydroxycephalosporin C was examined with the cell-free extract of several cephamycin producing strains for the selection of converting strain. *Streptomyces clavuligerus* ATCC 27064 was the most potent strain for the activity of cephalosporin 7a-hydroxylase. The cephalosporin 7a-hydroxylase was purified to near homogeneity through DEAE-sephacel chromatography, ammonium sulfate