**F305** Molecular cloning and characterization of *mucK*, a gene determining growth of *Acinetobatcer calcoaceticus* 93A<sub>2</sub> on exogenous *cis., cis*-muconate

Kye-Heon Oh<sup>1\*</sup> and Peter A. Williams<sup>2</sup>
<sup>1</sup>Department of Life Science, Soonchunhyang University; <sup>2</sup>Department of Biological Sciences, University of Wales, United Kingdom

The DNA sequence of a 1.68 kilobase-pair Xbal/Sall Acinetobacter calcoaceticus 93A2 restriction fragment carrying mucK, encodes a member of a superfamily of membrane transport proteins and within a subgroup involved in the uptake of cis, cis-muconate, was determened. Comparison of the mucK sequences from A. calcoaceticus ADP1 and A. calcoaceticus 93A2 demonstrated 8 base-pair substitution. Mutants from naturally transformed A. calcoaceticus 93A2 were isolated and the lysated DNA was prepared. Using polymerase chain reaction, the DNA fragments containing mucK were amplified and subcloned by pCR-blunt vector. As a result of sequencing DNA, a 29 base-pair deletion of the regulatory gene of mucK was determined. The deletion appeared to be functionally related to the utilization of cis, cis-muconate as an exogenous carbon source.

## F306 Characterization of the *pcbE* Gene Encoding 2-Hydroxypenta -2,4-dienoate Hydratase from *Pseudomonas* sp. DJ-12

Jong-Chul Lim\*, Kyung-Rak Min¹, Youngsoo Kim¹ and Chi-Kyung Kim Department of Microbiology, ¹Department of Pharmacy, Chungbuk National University

Pseudomonas sp. DJ-12 is able to degrade 4-chlorobiphenyl via the meta-cleavage pathway. We have sequenced the downstream region of pcbC of pCU1 which was cloned from the DJ-12 strain. The pcbE gene encoding 2-hydroxypenta-2,4-dienoate hydratase was identified, which is required for the conversion of 2-hydroxypenta-2,4-dienoate to 4-hydroxy-2-oxovalerate in the degradation of biphenyl and 4-chlorobiphenyl. The pcbE gene is apart about 1.5 kb from pcbC gene. A deduced amino acid sequence of the hydratase encoded in pcbE gene exhibited about 45-55% identities with those of 2-hydroxypenta-2,4-dienoate hydratases from Pseudomonas sp. KKS102, Burkholderia cepacia LB400, Pseudomonas putida F1, Pseudomonas sp. DJ77, Pseudomonas pseudoalcaligenes KF707.