Biosynthesis of medium-chain-length poly(3-hydroxyalkanoates) in recombinant Escherichia coli

<u>박종필</u>, 박시재, 이상엽* 한국과학기술원 생명화학공학과 및 생물공정연구센터 전화 (042) 869-3930, Fax (042) 869-3910

Abstract

In this study, we constructed the *fadA* and/or *fadB* mutant *Escherichia coli* strains for the production of medium-chain-length (MCL) polyhydroxyalkanoate (PHA) production. All the *fad* mutant *E. coli* strains were enough to support the MCL-PHA biosynthesis when equipped with MCL-PHA synthase. However, the monomer composition of PHA was highly dependent upon the metabolic capability of host strains and substrate specificity of PHA synthase. The expression of PHA synthase in *fadA* and/or *fadB* mutant *E. coli* strains yielded the accumulation of MCL-PHA, the constituents of which were composed of 3-hydroxybutyrate (3HB), 3-hydroxyhexanoate (3HHx), 3-hydroxyoctanoate (3HO), 3-hydroxydecanoate (3HD), and 3-hydroxydodecanoate (3HDD) depending on the carbon number of used fatty acid.

Acknowledgement

This work was supported by the National Research Laboratory Program (2000-N-NL-01-C-237) of the Korean Ministry of Science and Technology (MOST).

References

- 1. Lee, S. Y., "Bacterial polyhydroxyalkanoates" (1996), Biotechnol. Bioeng, 49, 1-14.
- 2. Langenbach S, Rehm BHA, Steinbüchel A., "Functional expression of the PHA synthase gene *phaC1* from *Pseudomonas aeruginosa* in *Escherichia coli* results in poly(3-hydroxyalkanoate) synthesis" (1997), *FEMS Microbiol Lett*, 150, 303-309.
- 3. Qi Q, Rehm BHA, Steinbüchel A., "Metabolic routing towards polyhydroxyalkanoic acid synthesis in recombinant *Escherichia coli* (fadR): inhibition of fatty acid beta-oxidation by acrylic acid" (1998), FEMS Microbiol. Lett. 167, 89-94.
- 4. Link AJ, Phillips D, Church GM., "Methods for generating precise deletions and insertions in the genome of wild-type *Escherichia coli*: application to open reading frame characterization" (1997), *J. Bacteriol.* 179, 6228-6237.