

An efficient bacterial surface display system based on novel outer membrane anchoring elements of
Escherichia coli

한미정*, 김준혁
동양대학교
(mjhan75@dyu.ac.kr*)

In this study, the *Escherichia coli* outer membrane proteomes were analyzed to identify potential candidates for using anchoring motifs. Of many proteins identified by mass spectrometry, the utility of outer membrane protein Y as an anchoring motif was examined. Two enzymes (an α -amylase from *Bacillus subtilis* or a lipase from *Pseudomonas fluorescens* SIK W1) were used for display as a target protein. SDS-PAGE, Western blot, and whole-cell enzyme activity measurement confirmed the successful expression of fusion proteins on the surface of *E. coli*. The fusion protein with Y232 as the anchoring motif had the highest expression level and enzyme activities. These results suggest that protein Y could be used as an anchoring motif of *E. coli* for displaying active enzymes and this system could be employed to various biocatalytic applications. [This work was supported by the Basic Science Research Program (2010-0008826) and Converging Research Center Program (2009-0093652) through the National Research Foundation of Korea funded by the Ministry of Education, Science and Technology]