Expression of glutamate decarboxylase from *Lactobacillus* sp. OPK-3 in *Bacillus subtilis* enhances GABA production

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Abstract

To express a foreign glutamate decarboxylase (GAD) in *Bacillus* host system, a recombinant DNA (pLip/GAD) was constructed by ligating the GAD gene from *Lactobacillus* sp. OPK-3 into *Escherichia coli-Bacillus* shuttle vector pLip. The pLip/GAD construct was then transformed into *B. subtilis*. The transformed *Bacillus* strain with the pLip/GAD constructed had higher GAD activity than untransformed *Bacillus* counterpart. When the strains were cultured in LB medium containing 1% monosodium glutamate (MSG), the levels of GABA produced by the transformed and untransformed *B. subtilis* were 220 mg/L and 46 mg/L, respectively. Under 3% MSG conditions, the levels of GABA produced by the transformed and untransformed strains were 401 mg/L and 49 mg/L, respectively. In agreement with these observations, the transformed *Bacillus* strain maintained its normal growth rate despite high MSG concentrations. Thus, by introducing foreign GAD gene, *Bacillus* strains has been genetically engineered to produce high levels of GABA from MSG.

Reference

- 1. Park K.B.,, Ji, G.E., Park, M.S., Oh, S.H. (2005) Expression of rice glutamate decarboxylase in *Bifidobacterium longum* enhances gamma-aminobutyric acid production. *Biotechnology letters*: in press.
- 2. Han S.J., Back J.H., Yoon M.Y., Shin P.K., Cheong P.K., Cheong C.S., Sung M.H., Hong S.P., Chung I.Y., and Han Y.S. (2003) Expression and

- characterization of a novel enantioselective lipase from *Acinetobacter* species SY-01. *Biochimie* 85: 501-510.
- 3. Lee, M.H., Song, J.J., Choi, Y.H., Hong, S.P., Rha, E.H., Kim, K., Lee, S.G., Lee, S.C, Poo, H., Seu Y.B., and Sung, M.H. (2003) High-level expression and secretion of *Bacillus pumilus* lipase B26 in *Bacillus subtilis* chungkookjang. *J. Microbiol. Biotechnol.* 13: 892-896.