In vitro directed evolution of TIMP-2 protein for soluble expression from E. Coli

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Abstract

TIMP-2, the second member of the family of tissue inhibitors of matrix metalloproteinases (TMMPs) is 23KDa protein which inhibit matrix metalloproteinase 2 (MMP-2)¹⁾. High-level intracellular expression of many mammalian protein in E.coli, results in the formation of large insoluble aggregates, known as inclusion body²⁾. The TIMP-2 is invariable insoluble when expressed in E.coli¹⁾, because this folding requires the formation of 6 disulfide bonds being incompatible with the reducing environment of the E. coli. but the soluble recombinant protein can often be recovered from E coli by in vitro recombination and mutagenesis³⁾. In this reason, we tried the staggered extension process(StEP) that is one of the in vitro PCR-based recombination methods for mutagenesis⁴⁾. C-terminally located CAT gene expression from the mutant genes made us indentify the soluble from the insoluble through linker sequences⁵⁾. As the results, chloroamphenical resistant varients were selected and some of the mutant showed dramatic soluble expressin of TIMP-2 compared to the wild type. Therefore, StEP has successfully improved folding and solubility of TIMP-2, and the CAT depedent screening may be a simple and effective method in selecting soluble forms from the insoluble. This study was supported by HTPEB (HTPEB 01 - PJ11 - PG9 - 01NT00-0036).

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