## E121

## Heterologous Expression of Yeast Prepro-a-Factor in Rat Pituitary GH<sub>3</sub> Cells

이명애\*<sup>1</sup>, 한상열<sup>1</sup>, 박상대<sup>1</sup>, 홍승환<sup>1,2</sup> 서울대학교 자연대학 <sup>1</sup>분자생물학과, <sup>2</sup>유전공학 연구소

Yeast pheromone  $\alpha$ -factor is a 13-amino acid peptide hormone that is synthesized as part of a larger precursor, prepro- $\alpha$ -factor, consisting of a signal peptide and a proregion of 69 amino acids, which contains three potential glycosylation sites. The carboxy-terminal half of the precursor contains four tandem copies of mature  $\alpha$ -factor, each preceded by spacer peptides of six or eight amino acids (variations of Lys-Arg-Glu-Ala-Asp-Ala-Glu-Ala), which are excised to produce mature  $\alpha$ -factor. To investigate the molecular basis for intracellular sorting, proteolytic processing, and storage of peptide hormone precursors, yeast pp  $\alpha$  was heterologously expressed in rat pituitary GH<sub>3</sub> cell. In GH<sub>3</sub> cells, as in yeast, the nascent polypeptide is efficiently targeted to the ER, where it undergoes cleavage of its amino-terminal signal peptide and core glycosylation to form glycosylated pro- $\alpha$ -factor. Subsequently, this species rapidly disappears from cells with a half life of  $\alpha$ 0 min, and are secreted to the medium. In these cells pp $\alpha$ 1 was accurately processed to the mature  $\alpha$ -factor with an efficiency of  $\alpha$ 20%. However, only 10% of the newly synthesized mature  $\alpha$ -factor and unprocessed precursor were stored intracellularly, whereas 90% was sorted to the constitutive pathway and secreted rapidly into the medium with kinetics identical to endogenous growth hormone. We demonstrated that expression of yeast pp $\alpha$ 6 in GH<sub>3</sub> rat pituitary cells results in the secretion of mature  $\alpha$ -factor and unprocessed p $\alpha$ 6 into the medium, suggesting that this wild type-prohormone could transit through the mammalian secretory pathway as endogenous protein. Our results show that signal peptide of yeast pp $\alpha$ 6 does direct it into the ER, and proregion does to the distal elements of the Golgi apparatus, respectively, but processing in putative cleavage site are not efficient.

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담배거세미 나방(Spodoptera litura)의 유충으로부터 유도합성된 항균활성 단백질 Spodopsin IA, IB, IC의 정제 및 특성

최충식\*, 여성문¹, 채권석², 김학렬 고려대학교 생물학과, 단국대학교 생물학과¹, 기초과학지원연구소 생명과학부²

담배거세미나방(Spodoptera litura)의 6령 유충의 복강에 micro syringe를 사용하여 약 10<sup>6</sup> 세포의 Salmonella Typhymurium을 주입시켜 항균활성단백질을 유도 합성시킨 후 ion-exchange column, gel filteration column과 마지막으로 reverse phase column을 이용하여 항균단백질을 정제한 다음 그 특성을 조사하였다. 정제된 항균단백질들 중 그람음성세균인 E. coli나 그람양성세균인 M. luteus등에 대해 강한 항균활성을 나타내는 약 4kDa의 분자량을 갖고있는 단백질을 Spodopsin IA, IB, IC 라고 명명하고 이들 각각을 tricine electrophoresis와 mass spectrometry 그리고 amino acid sequencer를 이용하여 정확한 분자량과 아미노산 서열을 조사하였으며 pl를 비롯한 각 단백질의 성질을 비교조사하였다.