

[12:20 – 13:00]

## **Beta-amyloid peptide degradation by aminopeptidase and its functional role in Alzheimer's disease pathogenesis**

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Beta-amyloid peptide ( $A\beta$ ) is a major component of senile plaques and its aggregation is considered to play a critical role in pathogenesis of Alzheimer's disease (AD). Aggregation of  $A\beta$  could result from both increased synthesis and decreased degradation of  $A\beta$ . Our laboratory is interested in understanding the mechanism of  $A\beta$  degradation in brain. Recently our laboratory identified a bacterial gene (SKAP) from *Streptomyces sp* KK565 whose protein product has an activity to cleave  $A\beta$  and thus reduce the  $A\beta$ -induced neurotoxicity. The sequence analysis showed that this gene was closely related to aminopeptidase. Maldi-Tof analysis showed that the recombinant SKAP protein expressed in *E. coli* cleaves both  $A\beta$  40 and  $A\beta$  42 at the N-terminal of  $A\beta$  while an aminopeptidase from *Streptomyces griseus* (SGAP) cleaves at the C-terminal. We also identified a mammalian homolog of SKAP and the recombinant mammalian protein expressed in Sf-9 insect cells showed a similar proteolytic activity to SGAP, cutting  $A\beta$  at the C-terminus. I will discuss the detailed mechanism of the enzyme action and its functional implication in AD.

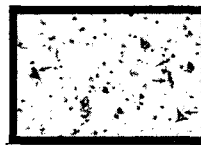
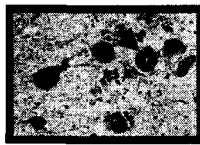
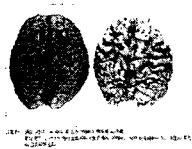
**Beta-amyloid peptide degradation by aminopeptidase  
and its functional role in Alzheimer's disease  
pathogenesis**

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**Alzheimer's Disease (AD)**



- Pathology** : senile plaque ( $\beta$ -amyloid peptide), neurofibrillary tangle (tau), neocortical atrophy, neuron and synapse loss
- Risk factors** : 1) increased age 2) family history  
3) gender 4) head injury
- Genes involved in AD:**
  - 1) **Causative genes:** amyloid precursor protein (APP)  
presenilin 1, 2 (PS1, PS2)
  - 2) **Risk factor genes:** ApoE<sub>4</sub>, alpha<sub>2</sub>-macroglobulin ( $\alpha_2$ M), LRP

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## Genes linked to AD

**Table 1. Overview of Established AD Genes influencing the A $\beta$  Life Cycle**

Gene (Location [Mb]) <sup>a</sup>	Genetic Mechanism	Biochemical Phenotype
<i>APOE</i> (19q13 [50 Mb])	LOAD: risk association ( $\epsilon 4$ -allele)	a) $\uparrow$ A $\beta$ aggregation b) $\downarrow$ A $\beta$ clearance
<i>APP</i> (21q21 [26 Mb])	EOFAD: AA-change ( $n = 16$ mutations <sup>b</sup> ) LOAD: mostly neg. association findings	a) $\uparrow$ A $\beta_{42}$ /A $\beta_{40}$ ratio b) $\uparrow$ A $\beta$ generation/A $\beta$ aggregation
<i>PSEN1</i> (14q24 [73 Mb])	EOFAD: AA-change ( $n = 140$ mutations <sup>b</sup> ) LOAD: pos./neg. association findings	$\uparrow$ A $\beta_{42}$ /A $\beta_{40}$ ratio
<i>PSEN2</i> (14q42 [223 Mb])	EOFAD: AA-change ( $n = 10$ mutations <sup>b</sup> ) LOAD: mostly neg. association findings	$\uparrow$ A $\beta_{42}$ /A $\beta_{40}$ ratio

<sup>a</sup>"Mb" = million base-pairs, "EOFAD" = early-onset familial AD, "LOAD" = late-onset AD.

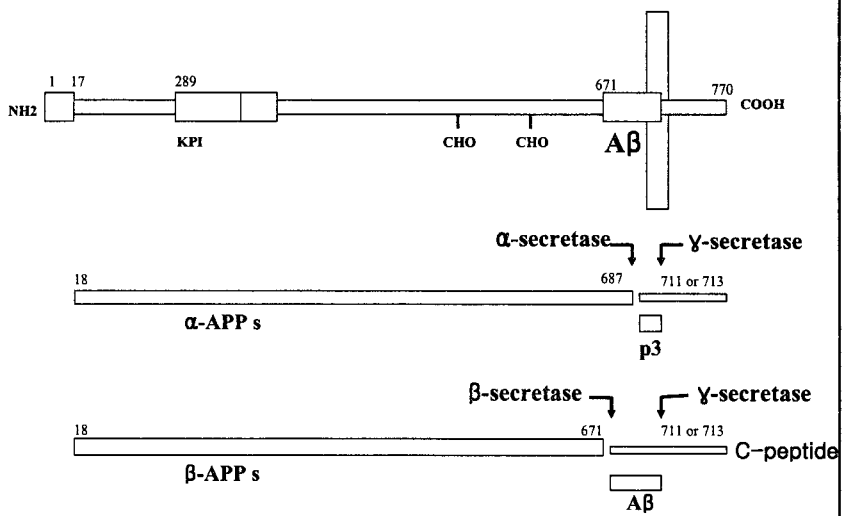
<sup>a</sup>Location according to "UCSC Human Genome Browser," May 2004 assembly (URL: <http://genome.ucsc.edu/cgi-bin/hgGateway>).

<sup>b</sup>According to the "Alzheimer's Disease Mutation Database" (URL: <http://molgen-www.uia.ac.be/ADMutations/>).

CELL 120:545-555 (2005)

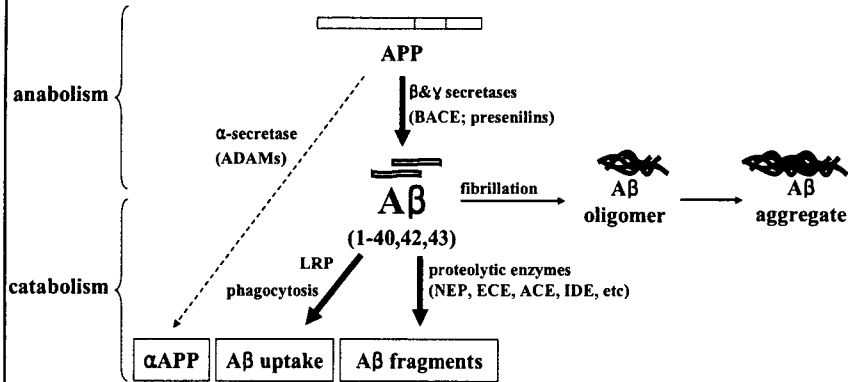
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## Amyloid precursor protein (APP) processing



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## Amyloid peptide metabolism



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## Proteolytic enzymes involved in degradation of the Alzheimer's amyloid peptide

### Neprilysin family

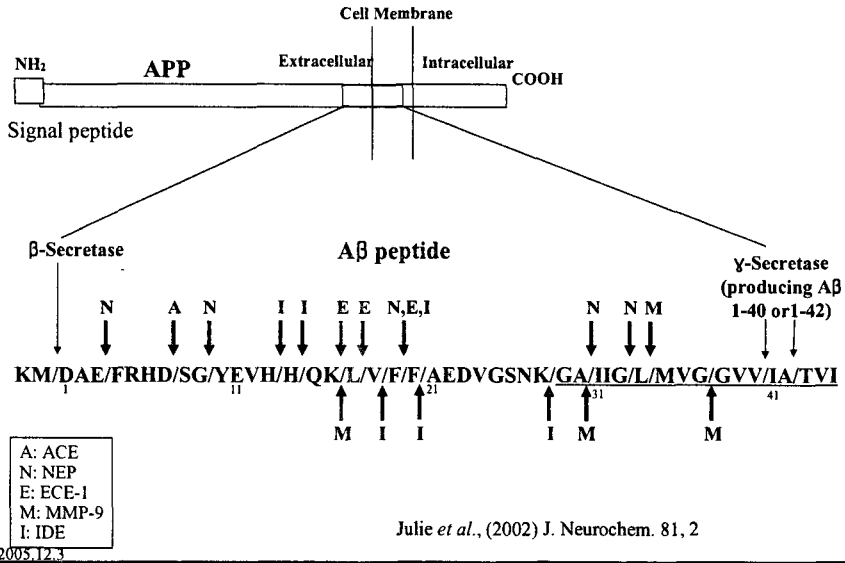
- neprilysin (NEP)
- endothelin-converting enzyme (ECE-1)
- angiotensin-converting enzyme (ACE)

### Insulin-degrading enzyme family

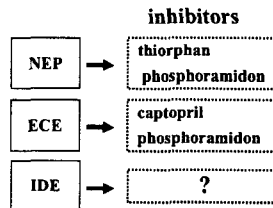
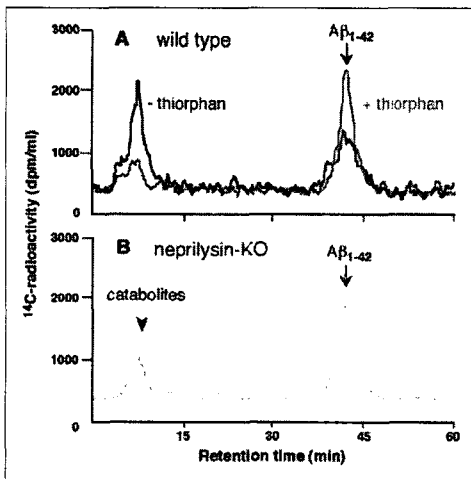
- insulin-degrading enzyme (IDE)

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### Peptidases responsible for Aβ cleavage



### Metabolic Regulation of Brain Aβ by Neprilysin



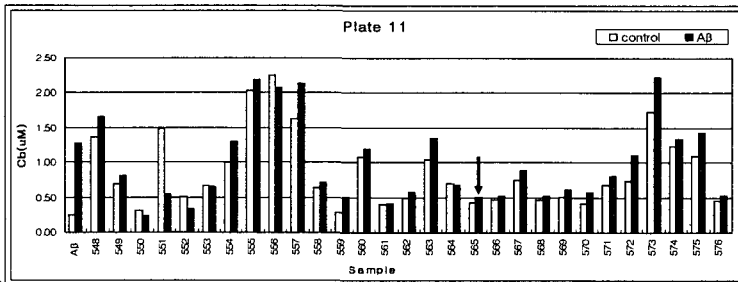
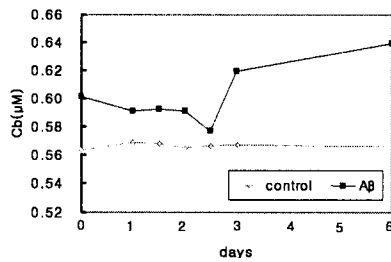
Saito group., *Science* (2001) 292:1550-1552

### Screening of microbial secretion product

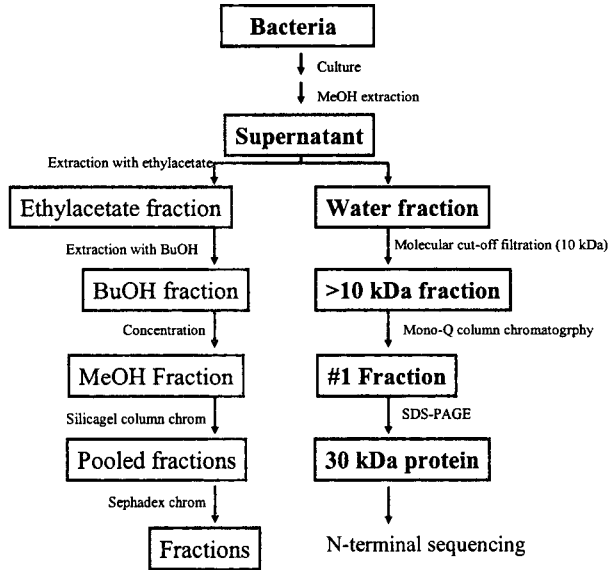
- *Identification of the candidate compounds from microbacteria, fungi and seaweeds libraries that are able to block Aβ-aggregation*

### Amyloid peptide aggregation assay

1. Congo red Assay  
 $Cb[M] = (A_{540}/25.295) - (A_{480}/46.306)$
2. Thioflavin T assay
3. Observe the pallet formation following centrifugation

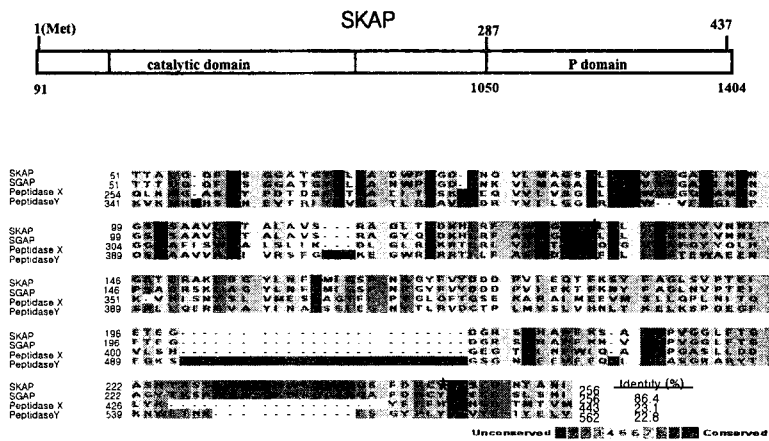


## Purification steps



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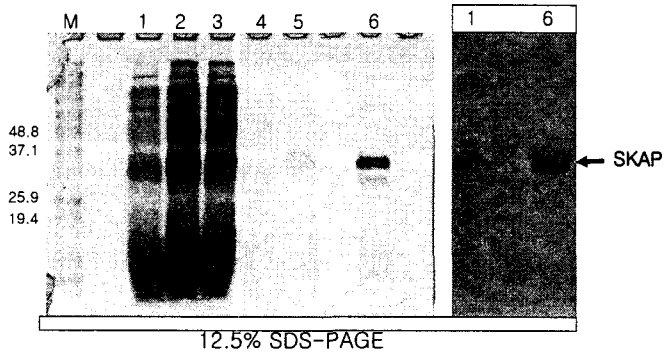
## Amino acid sequence comparison with other proteases



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### Preparation of the recombinant protein, SKAP30kDa, and identification with anti-SKAP antibody

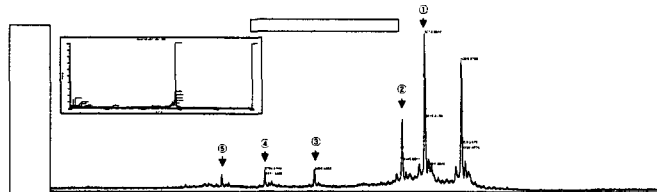


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### Amyloid peptide<sub>40,42</sub> cleavage by SKAP: MALDI-ToF analysis

(A) A $\beta$ <sub>40</sub> + SKAP, 3 h incubation



(B) A $\beta$ <sub>42</sub> + SKAP, 3 h incubation

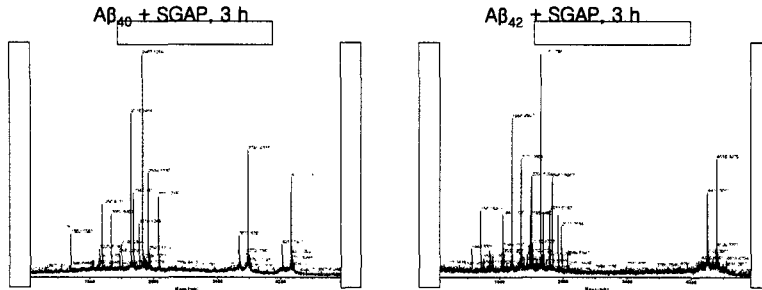


100  $\mu$ M of A $\beta$ ; 0.05  $\mu$ M SGAP

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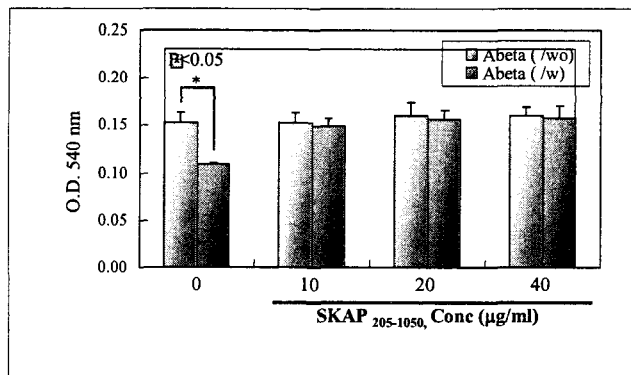
### Amyloid peptide<sub>40,42</sub> cleavage by SGAP



100  $\mu$ M of A $\beta$  : 0.05  $\mu$ M SGAP

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### Inhibition of A $\beta$ -induced neurotoxicity by SKAP



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## Summary

- ❑ We purified and cloned a gene (SKAP) from *Streptomyces* sp. (#90565) whose product is able to cleave amyloid peptide.

*Coding region=1314 bp, 437 amino acid, M.W.=45,209*

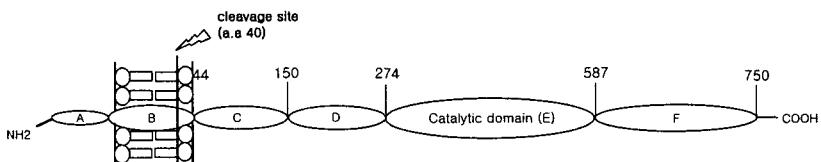
- ❑ Recombinant SKAP protein (30 kDa) blocked both fibrillation of amyloid peptide and reduced amyloid peptide-induced neuronal toxicity.

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## Peptidase X

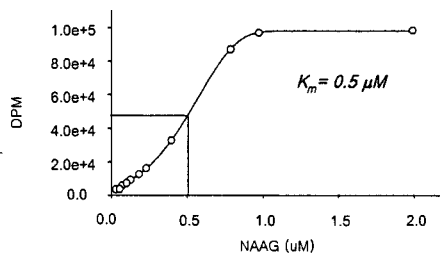
- ❑ 90-100 kDa transmembrane protein
- ❑ Located on neuronal and glial surfaces
- ❑ Tissue specificity: in various tissues

Schematic domain structure of human peptidase X



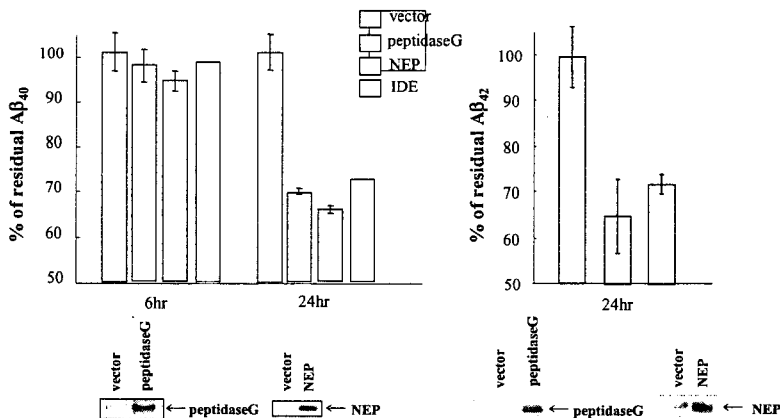
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### Preparation of recombinant peptidase G and measurement of its activity

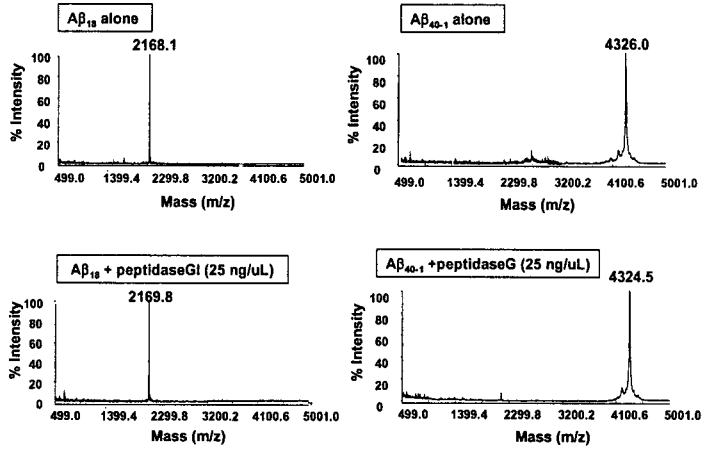


### $\text{A}\beta_{40}$ degradation by peptidaseX, NEP, IDE:

Transient transfection

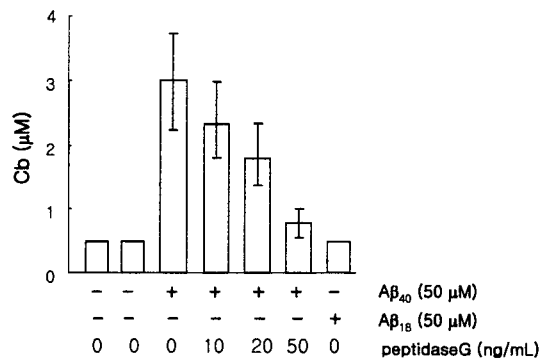


### $A\beta_{40-1}$ and $A\beta_{1-18}$ are not cleaved by peptidase G.



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### $A\beta_{40}$ aggregation is reduced by recombinant peptidase G protein



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### Neuronal cell death induced by A $\beta$ <sub>40</sub> decreased by incubation with peptidase G



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### Summary

- Both synthetic and endogenous A $\beta$  are degraded by peptidase G.
- Both A $\beta$ <sub>40</sub> and 42 are cleaved by peptidase G.
- Peptidase G cleaves A $\beta$ <sub>40</sub> into small fragments (A $\beta$ <sub>18</sub>) which lacks aggregation property and are not toxic to neuron.
- Peptidase G seems to degrade multimeric A $\beta$  more efficiently than monomeric A $\beta$ .
- Peptidase G protects neurons from toxicity induced by A $\beta$  by cleaving it into smaller fragments.
- Thus, dis-regulation of peptidase G could contribute amyloid deposit found in AD brain.

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**Ongoing project with aminopeptidase G and its future application to AD therapy**

- Treatment of AD mouse model (APP and presenilin) with peptidase G inhibitor: increases the both Ab40 and 42 level.
- Gene delivery of peptidaseG into the brain of AD model mouse
- Identification of the pharmacological agents that increase the endogenous peptidaseG activity
- Identification of the transcriptional factors or the signaling pathways that stimulate transcription of the peptidaseG gene

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***Thank you!!***

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