

## Scope of Amyloid Formation Inhibitor Development Targeting $\beta$ -Secretase

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Alzheimer's disease (AD) is a neurodegenerative disease characterized by progressive dementia accompanied by extensive neuritic plaque in brain caused by deposition of amyloid fibrils consisting of amyloid  $\beta$  peptides ( $A\beta$ ). Two kinds of proteases cleave the one transmembrane precursor protein (APP) consisting of about 700 amino acid residues, and produce  $A\beta$ .  $A\beta$  is located in the extracellular and transmembrane region in APP.  $\beta$ -Secretase cleaves the N-terminal part of  $A\beta$ , and then  $\gamma$ -secretase cleaves C-terminal part. The typical isoforms of  $A\beta$  are  $A\beta$ 40 with C-terminal Val and  $A\beta$ 42 with 2 amino acids (Ile-Ala) extended at C-terminal.  $A\beta$ 42 shows higher aggregating tendency than  $A\beta$ 40. Since the aggregated  $A\beta$ 42 entangles  $A\beta$ 40 and forms amyloid fibrils, accumulation of  $A\beta$ 42 is considered to be deeply involved in onset of AD.  $\alpha$ -Secretase cleaves APP at the different site from  $\beta$ -secretase and inhibits the production of  $A\beta$ .

In 1999, the structure of BACE1 ( $\beta$ -APP cleaving enzyme) was determined by cloning and shown to be a transmembrane glycoprotein and an aspartic protease, which drastically accelerated the inhibitor studies. The  $A\beta$  hypothesis that the accumulation of  $A\beta$  triggers the onset of AD may be established if drugs to inhibit the production of  $A\beta$  are developed for therapeutics of AD. At present, BACE1 inhibitors are designed based on substrate transition-state concept and demonstrated that they inhibit  $A\beta$  production in vivo.