

## The $\beta$ -stranded A $\beta$ 29-42 peptide is bound to membrane

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CBMN, FSAGX

The amyloid- $\beta$  1-40 and 1-42 peptides (A $\beta$ ) are together known to be the major component of senile plaques in Alzheimer's disease (AD). Nevertheless, the mechanisms of A $\beta$  1-40 and A $\beta$  1-42 toxicity are still poorly understood. But there is increasing evidence to show interactions between the carboxy-terminus of A $\beta$  peptides and membranes. Molecular modeling has allowed us to advance that A $\beta$  29-42 is a tilted peptide: it penetrates membrane at an angle of  $70^\circ$  as an  $\alpha$ -helix. To investigate this hypothesis, we have studied the A $\beta$  29-42 peptide together with model membranes using solid-state NMR experiments. Density centrifugation analyses confirmed the binding of the peptide at a 2.5 % peptide to lipid molar ratio.  $^{13}\text{C}$  MAS NMR experiments revealed that A $\beta$  29-42 presents mainly an  $\alpha$ -helix in HFP and a  $\beta$ -strand conformation in the presence of membrane. The study of membrane dynamics using static  $^2\text{H}$  NMR, with chain labeled lipids and Alanine methyl labeled peptide did not show any modification of the lipid and peptide dynamics upon binding. However  $^{31}\text{P}$  low speed MAS spectrum has indicated a significant broadening of PC and PE resonances, indicating interaction with the lipid's polar heads. These results suggest that A $\beta$  29-42 adopts a  $\beta$ -strand conformation when bound to lipids and does not penetrate membrane.