

Presenilin, γ -secretase and the search for pathogenic triggers in Alzheimer's disease

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Mutations that cause early-onset familial Alzheimer's disease (FAD) are in the substrate and enzyme that produce the amyloid β -peptide ($A\beta$), with the 42-residue form ($A\beta_{42}$) pathologically depositing as cerebral plaques. FAD mutations in the amyloid precursor protein (APP) and presenilin, the catalytic component of the γ -secretase complex, alter processive proteolysis of APP substrate to skew the proportion of longer, aggregation-prone $A\beta$ peptides and increase the $A\beta_{42}/A\beta_{40}$ ratio, long considered important for triggering pathogenesis. However, not all FAD mutations increase this ratio, and the large majority decrease overall proteolytic activity, resulting in reduced $A\beta$ production. We recently reported evidence that FAD mutations commonly lead to reduction in early cleavage steps in the processive proteolysis of APP substrate by γ -secretase due to stalling of enzyme-substrate (E-S) complexes (Devkota *et al.*, *Cell Reports*, 2024; Arafi *et al.*, *eLife*, 2025). Development of a *C. elegans* model system for FAD led to the finding that stalled and stabilized γ -secretase E-S complexes are sufficient to induce age-dependent synaptic loss and reduced lifespan, independent of $A\beta$ production. Thus, the stalled process—not the products—of γ -secretase proteolytic processing can cause synaptic degeneration. These findings raise the question of whether stalled E-S complexes—not $A\beta_{42}$ —is the primary disease driver in FAD. New unpublished results in support of the “stalled complex hypothesis” and its relevance to the pathogenesis of late-onset sporadic AD will be presented.

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