## P005 / #785

## **POSTERS**

## HUMAN AMYLOID SEEDS AGGREGATE MORE EFFICIENT THAN SEEDS FROM OLD 3XTG-AD MICE

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Aims: Most age-associated neurodegenerative disorders involve the aggregation of specific proteins within the nervous system, as occurs in Alzheimer's disease (AD). Recent evidence indicates that A $\beta$  can misfold and aggregate into seeds that structurally corrupt native proteins, mimicking a prion-like process of template protein corruption or seeding. In fact, studies in animal models show that the injection of brain homogenates from AD patients or from aged APP-transgenic mice containing A $\beta$  aggregates, can induce some of the neuropathological hallmarks of AD. However, it is still unknown which A $\beta$ -misfolded species are most efficient in triggering the aggregation process. Here, we seek to perform a comparative study to determine whether A $\beta$  seeds from humans vs a familial AD line (the 3xTg-AD model) is more efficient to generate amyloid aggregates.

**Methods:** We employed histological and molecular approaches to determine amyloid level, species and aggregative capacity of brain homogenates from an AD patient (stage C for amyloid, from the Alzheimer's Disease Research Center at UCI) vs old-3xTg-AD mice (25-month-old). Such brain homogenates were injected into the hippocampus of 7-month-old 3xTg-AD mice and the mice were analyzed at 18 months of age.

**Results:** Our findings demonstrated that amyloid seeds from the human patient have more capacity to generate Aß plagues vs seeds from aged 3xTq-AD mice.

**Conclusions:** These results suggest that seeds from human patients seem to be more amyloidogenic than from aged 3xTg-AD mice. Thus, more profound understanding these factors will provide key insight on how amyloid pathology progress in AD.