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## **Background and Objectives**

- Alzheimer's disease (AD) is a progressive neurodegenerative brain disease that gradually destroys memory and cognitive skills<sup>1</sup>.
- AD is the most common cause of dementia among older adults. An AD afflicted brain is shown to have accumulation of abnormal protein clumps (plaques) and tangled fibers (neurofibrillary tangles)<sup>1</sup>.
- Our objective was to develop a comprehensive quantitative systems pharmacology (QSP) model of (AD) pathologies to assess the impact of investigational treatments in support of drug development in this progressive neurodegenerative disease with a high unmet medical need.

## Methodology

The comprehensive QSP model based on ordinary differential equations (ODE) includes the two defining features of AD pathology: Aβ production and aggregation to form dense plaques, and tau hyperphosphorylation, aggregation, spreading, and formation of neurofibrillary tangles (NFTs)<sup>1</sup>. Detailed features of the model include:

- Regulated A\u00f340 and A\u00f342 production and secretion, including BACE1 and  $\gamma$ -secretase activity
- Aβ monomer aggregation into oligomers, fibrils, and plaques with mixed Aβ42 / Aβ40 composition
- Aβ clearance by protein degradation, receptor-mediated uptake, phagocytosis, active and passive transport
- Peripheral production of Aβ
- Tau production, hyperphosphorylation, aggregation, NFT formation, and extracellular spreading
- Hypothesized regulation of tau pathology by Aβ
- Active and passive transport of soluble Aβ and tau species between brain interstitial fluid (ISF), cerebrospinal fluid (CSF), and plasma
- Representation of both ApoE4 carrier and non-carrier status
- Antibody PK and binding to  $A\beta$  or tau species and consequent impact on A $\beta$  or tau pathology

**Software:** SimBiology (R2017b), a MATLAB<sup>®</sup> based application was used for the implementation of the model. Calibration and **Qualification:** Initial conditions and parameters were informed by literature and in-house preclinical and clinical data. Biomarkers and endpoints were compared to clinical data. Qualification was informed by Rosa's Model Qualification Method.<sup>2</sup>

**Biomarkers and Endpoints: (a)** Fluid biomarkers (Aβ and Tau), **(b)** Aβ PET SUVR, **(c)** Tau PET SUVR

**Therapies/Interventions: (a)** Aβ targeting agents: solanezumab, crenezumab, aducanumab and gantenerumab. (b) Tau targeting agents (anti-tau antibody)

Key Assumptions and Limitations: Neuronal cell population and protein production is assumed to be constant. Brain ISF is modeled as a single well-mixed compartment. The model does not attempt to translate biomarker dynamics to cognitive endpoints at this stage.

# A Quantitative Systems Pharmacology Model of **Alzheimer's Disease Pathology and Treatment Modalities**









**Figure 4** Simulated profiles of target neutralization in brain (interstitial fluid volume = 242 ml) after administration of solanezumab, crenezumab, aducanumab, and gantenerumab at their clinical doses as flat dosing **a.** unbound Aβ monomers; **b.** unbound A $\beta$  oligomers; **c.** unbound A $\beta$  plaques.

## **Patient Phenotypes**

The most abundant data for Aβ and tau come from post-mortem brains from deceased moderate to severe (mod/sev) AD patients. A prodromal AD phenotype was then developed representing an earlier version of the same AD patient. In addition, the model includes patients with different ApoE4 carrier status.





- facilitates a **quantitative assessment** of the development on **biomarker dynamics** via in-
- The model can be leveraged to evaluate the disease from a mechanistic perspective as it

### References

- . Masters CL, et al. Alzheimer's disease. Nature Reviews Disease Primers 1 15056 (2015) . Friedrich CM. A model qualification method for mechanistic physiological
- QSP models to support model-informed drug development. *CPT:PSP* 5 43-53. (2016)
- . Cummings JL, et al. ABBY: A phase 2 randomized trial of crenezumab in mild to moderate Alzheimer disease. *Neurology* 90 e1889-97. (2018) 4. Salloway SS, et al. Amyloid positron emission tomography and
- cerebrospinal fluid results from a crenezumab anti-amyloid-beta antibody double-blind, placebo-controlled, randomized phase II study in mild-tomoderate Alzheimer's disease (BLAZE). Alzheimer's Research & Therapy 10 96. (2018)
- 5. https://clinicaltrials.gov/ct2/show/NCT02353598 accessed 31 March 2019 6. Sevigny J, et al. The antibody aducanumab reduces Aβ plaques in Alzheimer's disease. Nature 537 50-6. (2016)
- 7. Ostrowitzki S, et al. A phase III randomized trial of gantenerumab in prodromal Alzheimer's disease. Alzheimer's Research & Therapy 9 95. (2017)
- . Klein G, *et al*. Gantenerumab reduces amyloid-β plaques in patients with prodromal to moderate Alzheimer's disease: a PET substudy interim analysis. *Alzheimer's Research & Therapy* 11 101. (2019)
- 9. Haeberlein SB, et al. EMERGE and ENGAGE Topline Results: Two Phase 3 Studies to Evaluate Aducanumab in Patients With Early Alzheimer's Disease. CTAD San Diego CA USA. (2019)
- 0.Sanabria-Bohorquez S, et al. Evaluation of [18F]GTP1 (Genentech tau probe 1) Extent and Load for assessing tau burden in Alzheimer's disease. Presented at HAI, 16-18 January 2019, Miami, FL, USA.