

Modeled Impact of APOE4 Genotype on ARIA-E Incidence in Patients Treated With Lecanemab

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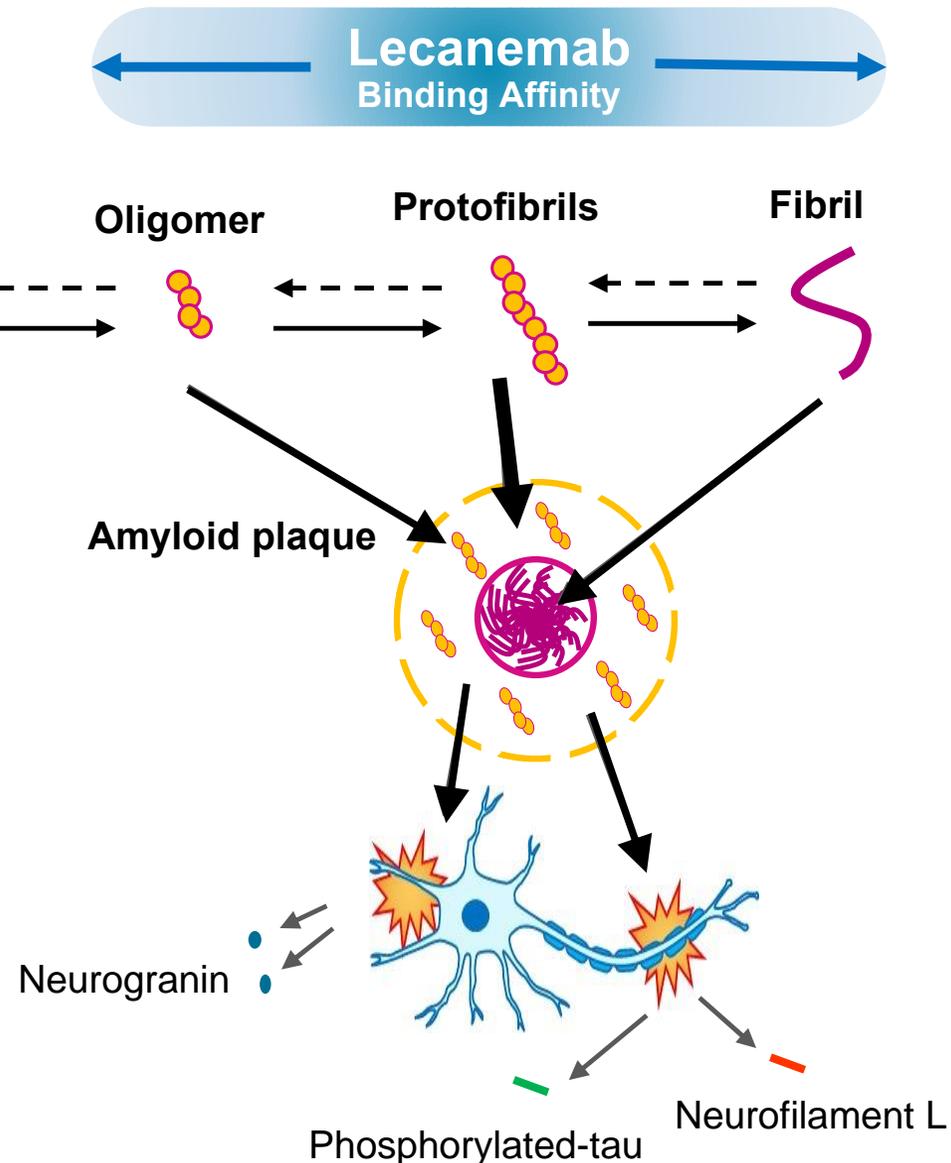
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Presenter Disclosures

- **Dr Reyderman is an employee of Eisai.**

Lecanemab: Anti-A β Protofibril Monoclonal Antibody

Selectively Targets A β Protofibrils



- Humanized immunoglobulin G1 (IgG1) monoclonal antibody
- Selectively binds to soluble A β aggregate species
 - >1000-fold selectivity for protofibrils over A β monomers (low affinity for A β monomer¹)
 - Preferential activity for A β protofibrils over fibrils (>10x)²⁻⁵
- Evaluated in an 856-patient placebo-controlled, randomized phase 2 study with an open-label extension (OLE)⁶
 - Reduction in brain amyloid accompanied by a consistent reduction of clinical decline
 - Well-tolerated with 9.9% ARIA-E at 10 mg/kg IV biweekly (~3% symptomatic)
 - Incidence of ARIA-E was lecanemab dose-dependent and greater in APOE4 carriers.

A β , amyloid-beta; ARIA-E, amyloid related imaging abnormalities - edema; IV, intravenous; p-tau, phosphorylated tau.

1. Tucker S, et al. *J Alzheimers Dis.* 2015;43(2):575-88. 2. Lord A, et al. *Neurobiol Dis.* 2009;36:425-34. 3. Sehlin D, et al. *PLoS One.* 2012;7:e32014. 4. Sehlin D, et al. *Neurodegener Dis.* 2011;8:117-23. 5. Logovinsky V, et al. *Alzheimer's Research & Therapy.* 2016;8:14. 6. Swanson et al. *Alzheimers Res Ther.* 2021;13(1):80.

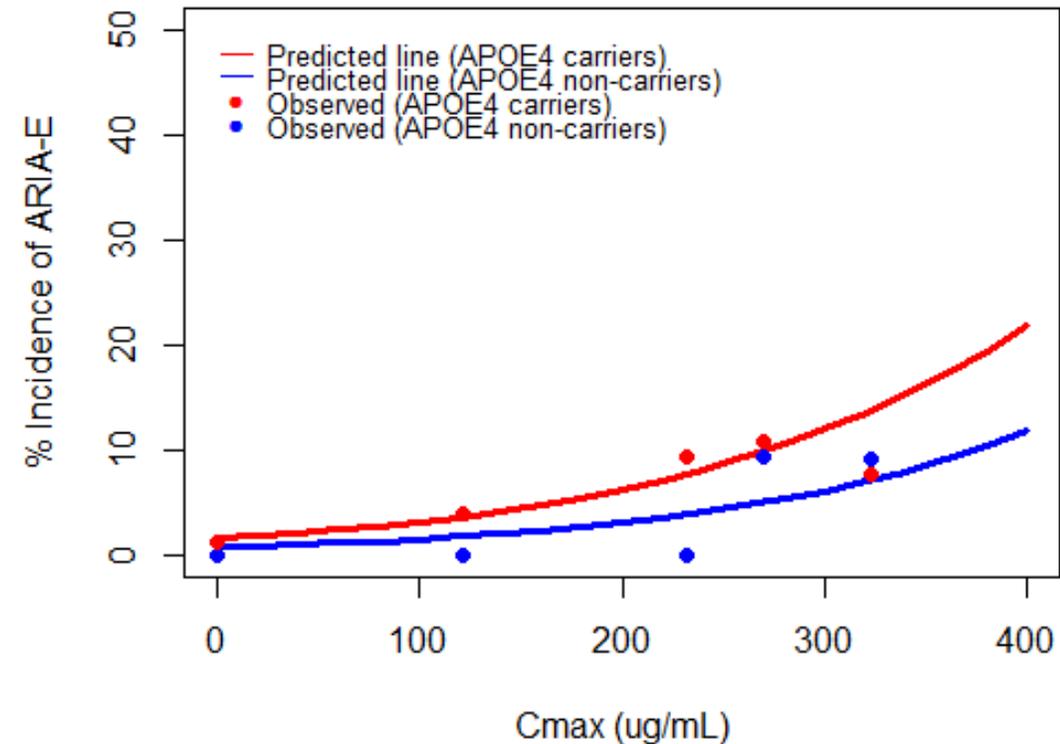
ARIA-E Exposure-Response Analysis

Effect of APOE4 Carrier Status

- Population PK model was used to derive $C_{max,ss}$ and $C_{ave,ss}$ for sequential Exposure-Response analysis of ARIA-E
- The incidence of ARIA-E as a function of lecanemab exposure was modeled using logit function
 - The exposure parameters, $C_{av,ss}$ and $C_{max,ss}$ were tested as linear function
 - Observed and predicted ARIA-E incidence attenuates over time
 - Log-hazard model was used as a base model

- Estimated baseline hazard was 26% lower for APOE4 non-carriers than carriers
- ARIA-E best correlated with C_{max} at steady-state

Model-Predicted % Incidence of ARIA-E vs. Lecanemab $C_{ss,max}^2$



Filled circles represent the observed proportion of subject with ARIA-E for each $C_{ss,max}$ quartile (1Q-4Q), plotted at the median $C_{ss,max}$ of each group. Solid line represents the model predicted line.

Analysis Objectives

- To evaluate APOE4 genotype defined as APOE4 homozygous carrier, heterozygous carrier, and APOE4 non-carrier as a covariate in the exposure-ARIA-E model
- To evaluate lecanemab exposure parameter ($C_{ss,max}$ or $C_{ss,ave}$) that is best predictor of ARIA-E incidence
- To test hypothesis that ARIA-E incidence in APOE4 heterozygous carriers and APOE4 noncarriers is similar

ARIA-E Exposure-Response Model :

Exposure Parameters and APOE4 Genotype Covariate Evaluation

- Inclusion of APOE4 genotype as a covariate in the model with 3 categories led to a statistically significant decrease in model objective function (OFV) ($P < 0.001$)
- Inclusion $C_{ss,ave}$ as a measure of lecanemab exposure resulted in larger OFV
 - $C_{ss,max}$ is a better predictor of ARIA-E compared to $C_{ss,ave}$
- Incidence of ARIA-E is similar in APOE4 heterozygous carriers and APOE4 noncarriers and distinct in APOE4 homozygous carriers

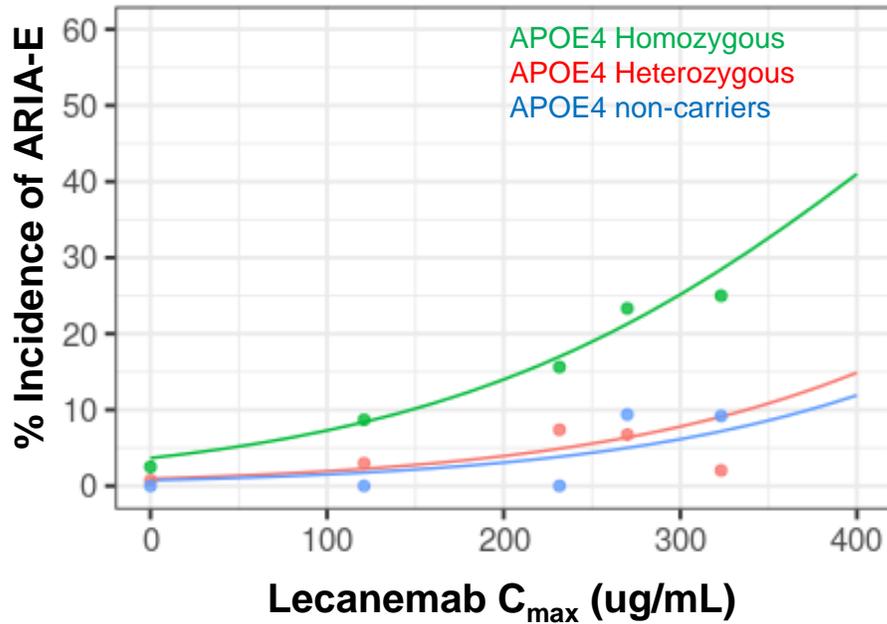
Model #	Model Description	OFV*
1	$C_{ss,max}$ <u>without</u> covariates	345.597
2	$C_{ss,max}$ with APOE4 genotype as <u>3 categories</u> (APOE4 noncarrier vs. hetero vs. homo)	326.584 ($P < 0.001$)
3	$C_{ss,ave}$ with APOE4 genotype as <u>3 categories</u> (APOE4 noncarrier vs. hetero vs. homo)	341.897
4	$C_{ss,max}$ with APOE4 genotype as <u>2 categories</u> (APOE4 homo vs others)	326.979 $P = 0.5297$ (vs. Model #2)

OFV = objective function value.

*defined as 2 categories, APOE4 homozygous carriers and other (APOE4 heterozygous carriers and APOE4 noncarriers)

ARIA-E Exposure-Response Model Predictions

Model Predicted and Observed ARIA-E vs. C_{max}



Filled circles represent the CORE observed proportion of subject with ARIA-E for each C_{max} quartile (1Q-4Q) and placebo, plotted at the median C_{max} of each group. Solid line represents the model predicted line.

- Model-predicted ARIA-E rates for APOE4 homozygous carriers treated with lecanemab 10 mg/kg biweekly were 22.5% (versus 6.8% in heterozygous or 5.4% in noncarriers)
 - Comparable to the observed 25% incidence in newly-treated homozygous carriers (1/4) in the OLE study
 - < 50% incidence in homozygous carriers (5/10) in the Core

Study 201 Core Model Predicted ARIA-E by C_{max}

Category	10 mg/kg Monthly (C_{max} =239 ug/mL)	10 mg/kg Biweekly (C_{max} =280 ug/mL)
APOE4+		
Homo	17.7%	22.5%
Hetero	5.1%	6.8%
APOE4-	4.0%	5.4%

Study 201 OLE Observed ARIA-E

Category	Newly Treated Core placebo* (N=45)
APOE4+	
Homo	1/4 (25.0%)
Hetero	3/27 (11.1%)
APOE4-	0/14 (0%)

*10mg/kg bi-weekly

Summary and Conclusions

- APOE4 genotype is a significant covariate in the exposure-ARIA-E model
- The probability of ARIA-E was best correlated with lecanemab maximum concentration at steady-state ($C_{ss,max}$)
- ARIA-E incidence is higher in APOE4 homozygous carriers and lower and similar in APOE4 noncarriers and APOE4 heterozygous carriers
- No statistically significant difference in ARIA-E incidence was found between non-carriers and heterozygous carriers
- Model-predicted ARIA-E rates for APOE4 homozygous carriers after lecanemab 10 mg/kg biweekly were 22.5% and comparable to observed 25% (1/4) ARIA-E incidence in newly-treated Core placebo APOE4 homozygous carriers in the OLE phase but lower than the 50% rate (5/10 observed in the Core phase)
- Limitation: under representation of APOE4 carriers in Core phase in 10 mg/kg biweekly dosing group. However:
 - The 10 mg/kg monthly has 90% APOE4 carriers (and similar C_{max} as 10 mg/kg biweekly)
 - 70% APOE4 carriers in OLE where rates are consistent, and all received 10 mg/kg IV biweekly
 - Kaplan-Meier estimates (not shown) take into account administrative terminations in estimating rate
 - DSMB reviews in Clarity AD and would require informing sponsor/FDA if greater risk
- Data from the larger Clarity AD study will be used to confirm these findings

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