Characterizing the APOE4/Trem2^{R47H} Mouse Model for MODEL-AD Late Onset Alzheimer's Disease

Harriet Williams¹ on behalf of the MODEL-AD consortium^{1,2,3,4} ¹The Jackson Laboratory, Bar Harbor, Maine, USA; ²Stark Neuroscience Research Institute, Indiana, USA; ⁴University of California, Irvine, California, USA; ⁴University, Irvine, California, Irvine, California, USA; ⁴University, Irvine, California, Ir

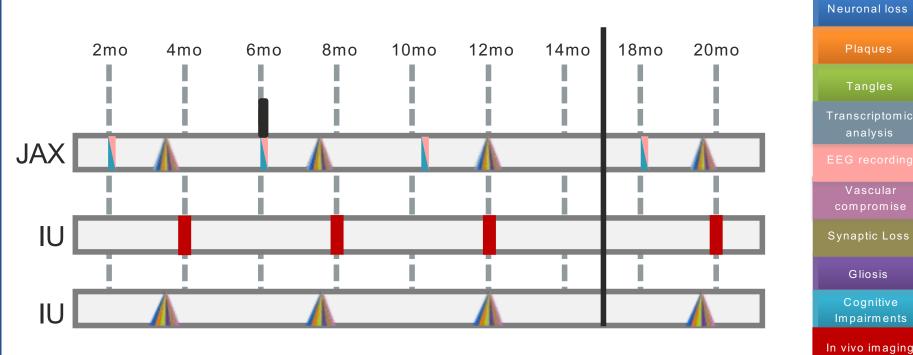
ABSTRACT

Alzheimer's disease (AD) is a debilitating neurodegenerative disorder. More than 46 million people are affected worldwide, with no effective treatment currently available. Research into fully understanding the disease has been fueled by animal models with familial AD (fAD) mutations, which accounts for 2-5% of the AD population. While these models have proved fruitful in understanding some of the baseline endophenotypes of AD, they have not been adequate to develop therapeutics or fully elucidate the processes leading to AD. One component of this lack of an effective therapeutic is due to the inability to fully reproduce an AD phenotype in these animals.

Here, the MODEL-AD consortium presents a new mouse model for Late Onset Alzheimer's disease (LOAD). We have developed a model carrying the two highest genetic risk factors for LOAD, the most common and strongest risk factor, APOE4, and the R47H allele of Trem2. To characterize this model, we have looked at three time points to understand the aging phenotype of these APOE4/Trem2*R47H mice.

At these three time points, mice were exposed to a battery of behavioral assays, continuous activity monitoring, frailty assessment, open field, grip consisting spontaneous alternation, rotarod, delayed spatial novelty, and episodic strenath. memory. Once completed, brains are removed and assessed. We are primarily looking for hallmark signs of AD, such as beta amyloid plaques and neurofibrillary tangles, as of neuroinflammation, vascular compromise, and neuronal morphology/counts

The development of this new model will enable us to gain a deeper understanding into the two of the genetic factors contributing to LOAD. Our goal is that the new models will lead to better therapeutic testing as well as determining if any preventative actions can be taken to impede the onset of AD. For more information see www.model-ad.org

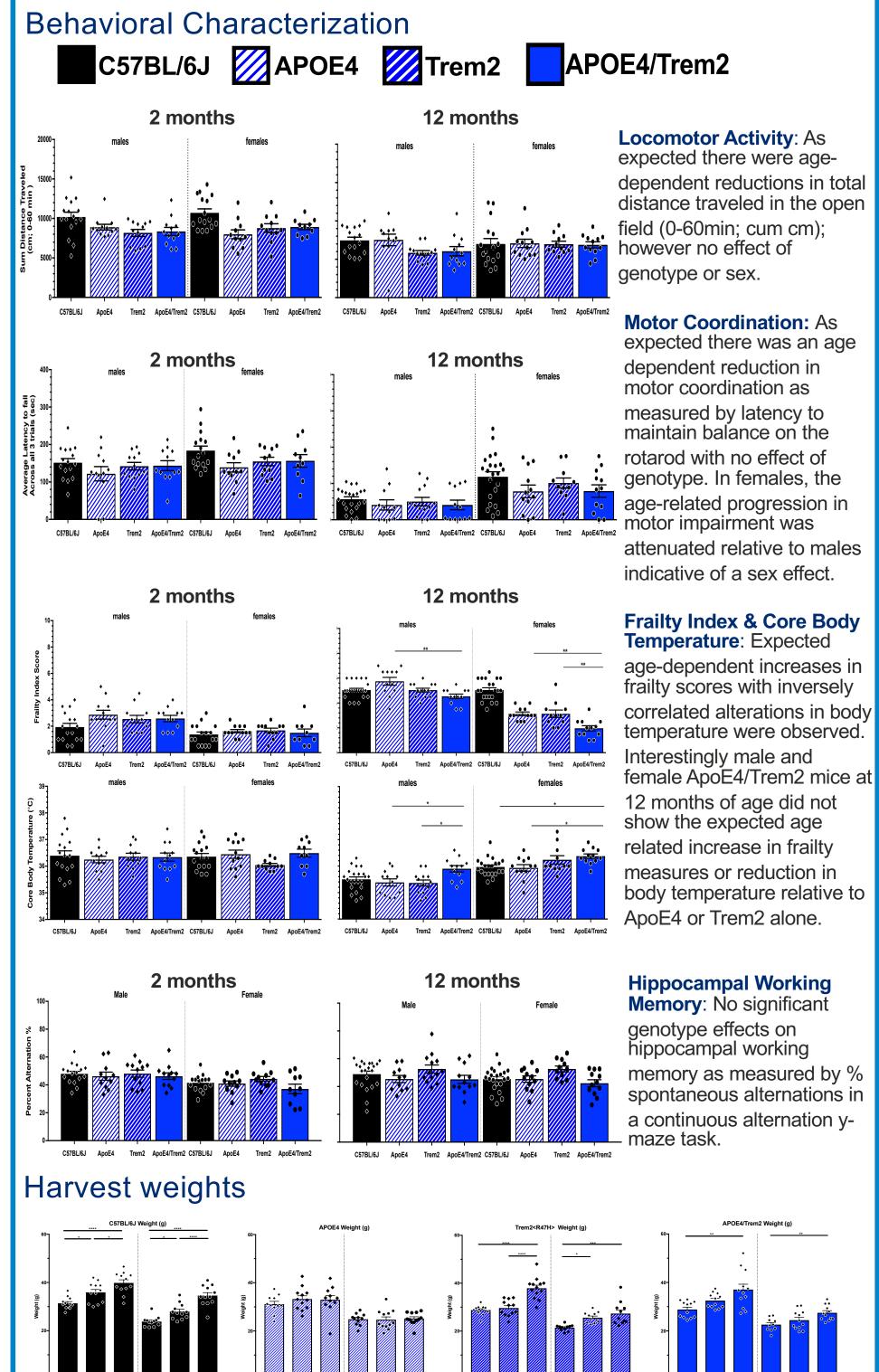


/letabolomics/Pro -eomics pilot

Deep phenotyping testing paradigm and histological analysis

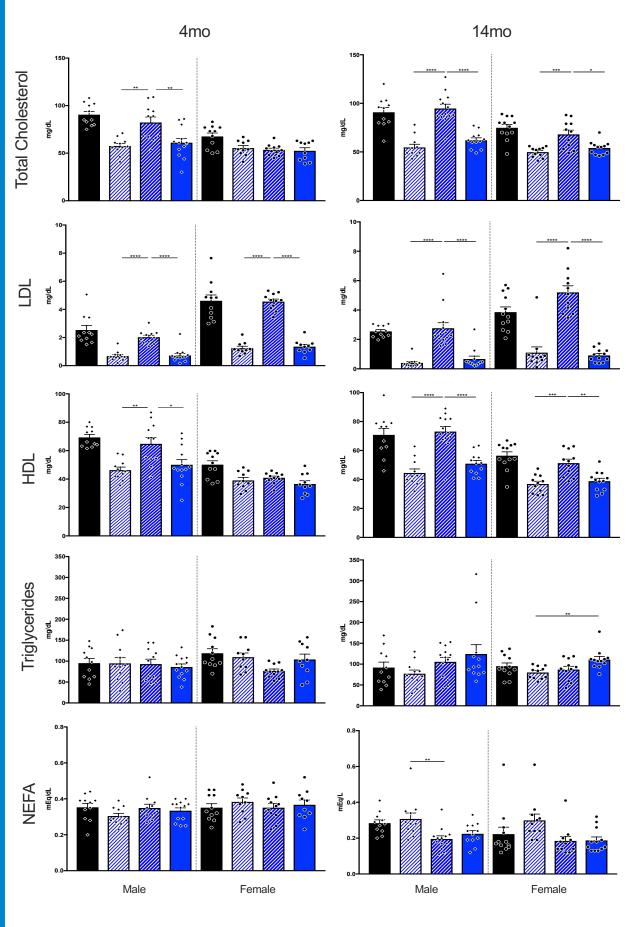
Further Information

- MODEL AD: <u>www.modelad.org</u>
- AMP-AD Knowledge Portal: <u>http://www.svnapse.org/ampad</u>
- JAX AD models: <u>https://www.iax.org/alzheimers</u>
- AlzForum research models: http://www.alzforum.org/research-models



At harvest, each mouse is weighed. As expected, C57BL/6J mice gain weight as they age. Whereas the APOE4 mice do not appear to gain weight through age.

Blood biochemistry

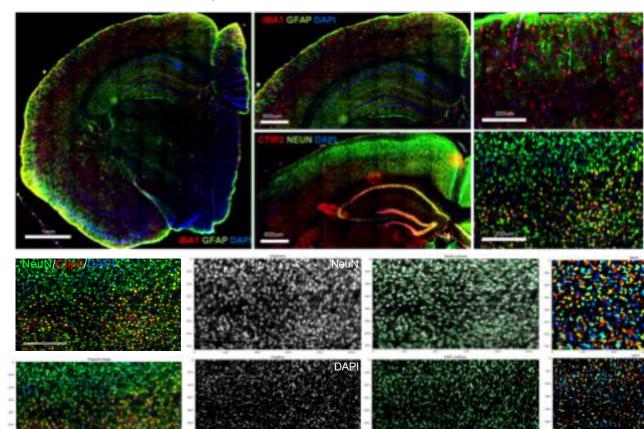


Histological imaging and analysis methods

Marker	Tar
Neun/Ctip2	Nei
GFAP/Iba1	Rea
X-34/Lamp1/Iba1	Pla
CD31/Fibrin/Iba1	Ves
LFB/Cresyl Violet	Gro
AT8/H&E	рТа

APOE4/Trem2 Weight (g)

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eurons/Excitatory neurons active astrocytes/microglia-monocytes/Astrocytes aques/Dystrophic neurites/Microglia-monocytes ssels/Vascular leakage/Microglia-monocytes oss morphology and myelin

Non-fasted blood was taken a harvest. Serum was separated and profiled for various cholesterol fractions.

Total cholesterol: Levels were seen to be significantly reduced in ApoE4 carrying mice in both males and females. This reductions was seen at 4 and 14mo.

LDL: LDL levels were significantl reduced in the ApoE4 carrying mice at both time points.

HDL: A significant reduction in HDL is seen in male mice at 4mo. At 14mo, a significant reduction of HDL was seen in both sexes.

Triglycerides: No significant differences were seen in triglyceride levels at 4mo of age. At 14mo of age, triglyceride levels were significantly increased in comparison to ApoE4 in females

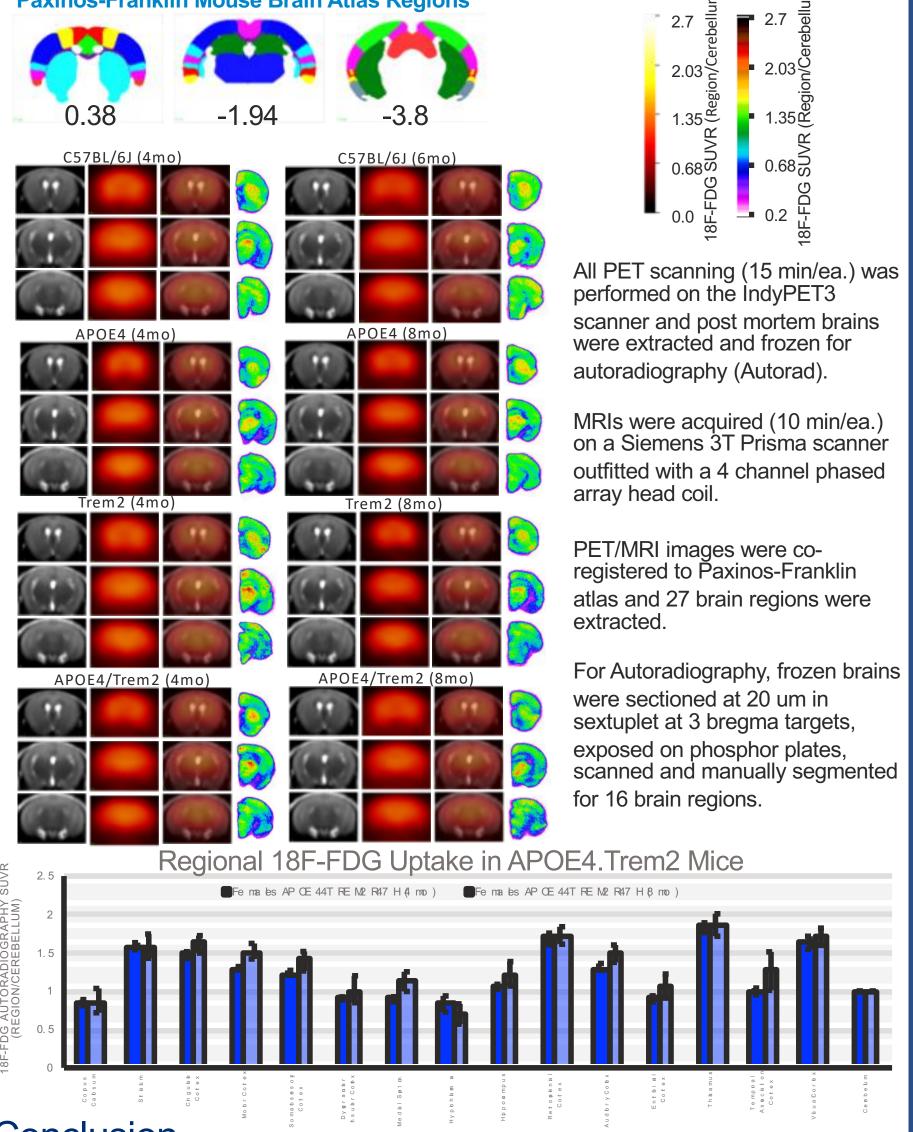
Non-Essential Fatty Acids (NEFA): At 4mo of age levels remained consistent across all genotypes. At 14mo Trem2 males had significantly reduced NEFA levels in comparison to ApoE4

Upon harvest, brains are removed, weighed, and hemispheres are separated. One hemisphere is fixed. cryopreserved, and sectioned in series at 20µm. Sections are stained with various antibody combinations, as outlined in the **Conclusion** table. Images are acquired using the Aperio Versa. Specific regions are extracted from the images, and processed using Cell Profiler, to assess cell numbers. This will allow for identification of cell loss in new models.

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	# of Neun	827
	#not associated with DAPI	110
R. 19	Total NeuN associated	717

MR/PET: 18F-FDG to assess glucose uptake

Paxinos-Franklin Mouse Brain Atlas Regions



- Behavioral characterization revealed the expected age-dependent alterations across phenotypes with no effect of genotype up to 12 months of age which would be anticipated in a model of LOAD. Further testing at 18mo of age is in progress.
- While there are no gross statistical significances between regions at, and between, 4 and 8mo in PET and AutoRad, further analysis continues for additional time points.
- In vivo imaging, transcriptomic, histological, and biochemical analysis continues on all time points.

Acknowledgments

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Sage UCI University of California, Irvine