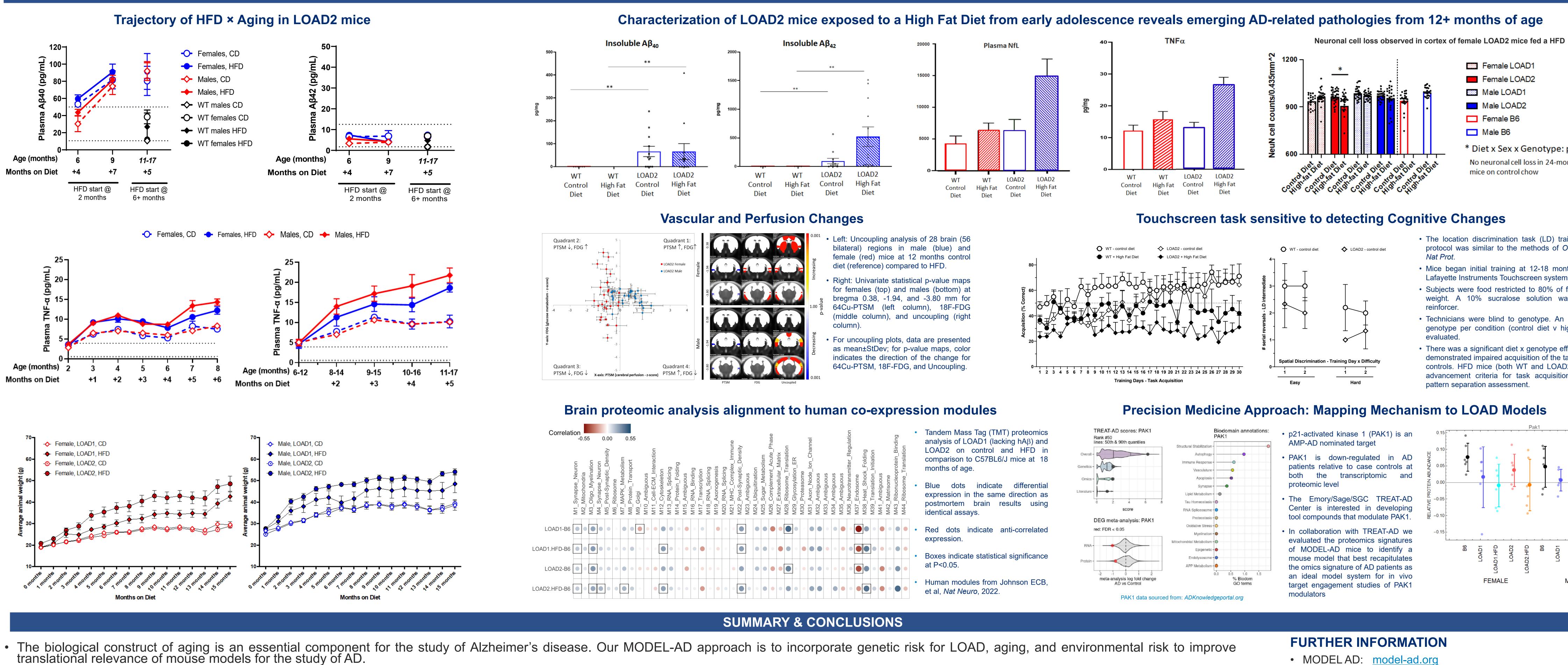
Aging × Genetics × Environment: Characterization of Precision Disease Models for Preclinical Testing for Late-onset Alzheimer's disease

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- present with sporadic late onset AD (LOAD).



INTRODUCTION

• The ability to effectively translate therapeutic efficacy from the bench to clinical success for Alzheimer's disease (AD) has been hampered in part due to limited recapitulation of the complexity of the disease in animal models. While analogous AD risk mutations have been engineered into animal models and have dominated the research field, these have primarily been familial, early onset risk alleles which do not capture the risk for AD for the majority of patients that

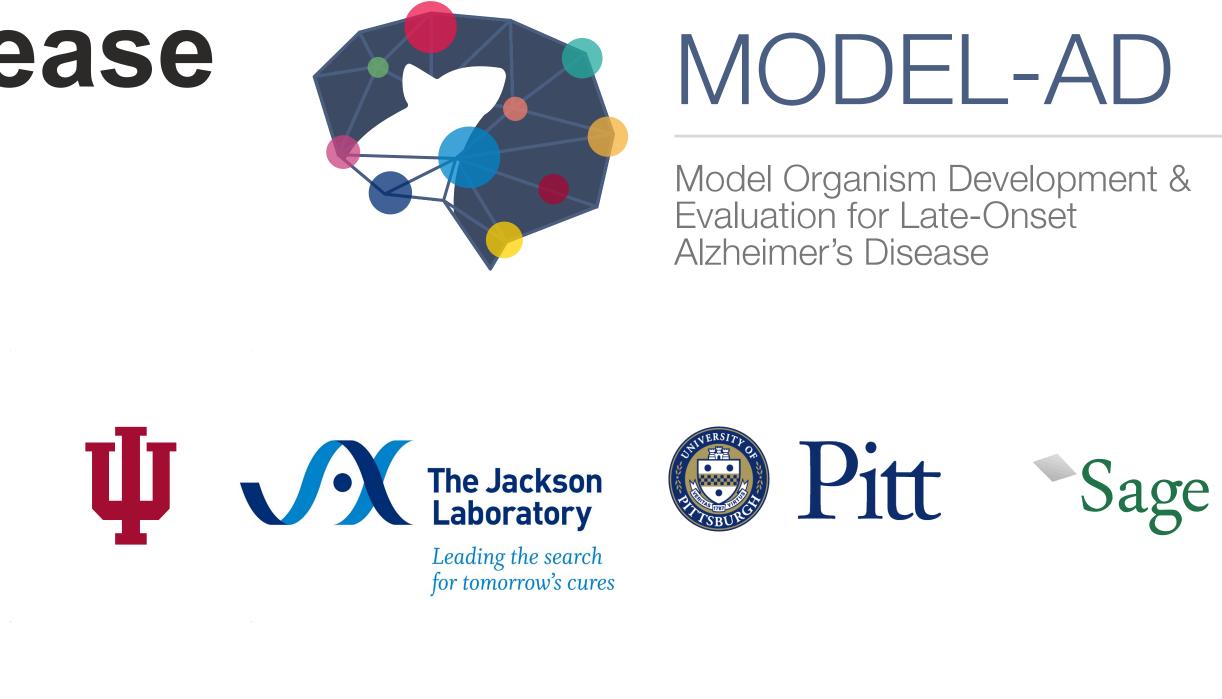
• The IU/JAX/PITT MODEL-AD consortium is focused on developing mouse models with genetic risk variants associated with LOAD, in combination with environmental risk factors and aging to enable improved translation. The present studies aimed to characterize mice expressing humanized Aβ in combination with multiple genetic risk factors (APOE4 and the R47H risk variant in the Trem2 gene; LOAD2) and aged in the presence of a high-fat, high-sugar diet (HFD).

 LOAD2 mice exposed to HFD from adolescence (LOAD2+HFD) demonstrated aging changes relative to LOAD2 mice in the absence of HFD, including presentation of insoluble Aβ42 in brain and plasma, and
increased inflammatory cytokines. 12-month aged LOAD2+HFD mice also demonstrated increased NfL in CSF, as well as vascular and perfusion changes as measured by MRI. By 18 months, LOAD2+HFD mice
demonstrated reductions in hippocampal neurons as well as cognitive impairment relative to LOAD2 mice in the absence of HFD on a translational touchscreen task. Intriguingly, gene expression profiles and
proteomic signatures of aged LOAD2+HFD mice aligned with 'omics signatures of AD patients in the absence of core neuritic plaques, which were not detected up to 24 months of age. • Mice with genetic risk for LOAD coupled with environmental risk factors demonstrate aging-dependent changes in line with a spectrum and trajectory of features of clinical LOAD. From a precision medicine approach, our MODEL-AD Preclinical Testing Core and our TREAT-AD colleagues are prioritizing LOAD2+HFD mice as an important model system for evaluating the therapeutic potential of non-amyloid targeting therapeutics as well as for prophylactic interventions initiated prior to significant amyloid accumulation.

- (adolescent HFD, aHFD) as well as beginning later in life (middle aged, at 6-12 months; mHFD).

RESULTS

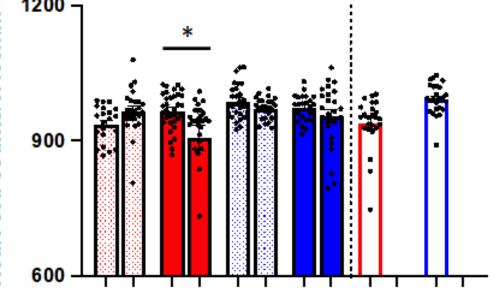
Precision Medicine Approach: Mapping Mechanism to LOAD Models



METHODS

• Initially, alleles were created using CRISPR to knock-in risk variants to the mouse locus on a sensitized C57BL/6J genetic background expressing APOE4 and Trem2^{R47H} ("LOAD1"; see Kotredes et al, 2021, *PMID: 34707490*). Later models were made on the same background with a humanized Aβ1-42 region ("LOAD2"). The LOAD2 mice are congenic on a C57BL/6J background and available at The Jackson Laboratory without restrictions (JAX stock# 030670).

• To investigate the role of high fat diet (HFD) exposure on exacerbating the manifestation of AD related pathologies on aging and genetic risk, we provided HFD ad libitum (Research Diets, Inc. catalog# D12451i; 45% fat) using two separate paradigms: beginning at 2 months of age following weaning



Female LOAD2 Male LOAD1 Male LOAD2 Female B6 🗖 Male B6

Female LOAD1

* Diet x Sex x Genotype: p=0.0277 No neuronal cell loss in 24-month LOAD2 mice on control chow

• Mice began initial training at 12-18 months of age using

• Subjects were food restricted to 80% of free-feeding body

• Technicians were blind to genotype. An n=4 per sex per

• There was a significant diet x genotype effect. LOAD2 mice

demonstrated impaired acquisition of the task relative to WT

controls. HFD mice (both WT and LOAD2) failed to meet

genotype per condition (control diet v high fat diet) were

weight. A 10% sucralose solution was used as the

Lafayette Instruments Touchscreen systems.

• The location discrimination task (LD) training and testing protocol was similar to the methods of *Oomen et al 2013*

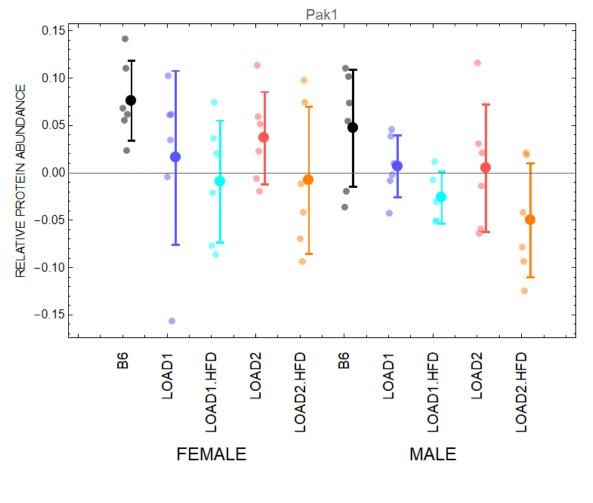
advancement criteria for task acquisition to advance to pattern separation assessment.

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evaluated.

- p21-activated kinase 1 (PAK1) is an
- PAK1 is down-regulated in AD patients relative to case controls at both the transcriptomic and
- The Emory/Sage/SGC TREAT-AD Center is interested in developing tool compounds that modulate PAK1.
- In collaboration with TREAT-AD we evaluated the proteomics signatures of MODEL-AD mice to identify a mouse model that best recapitulates the omics signature of AD patients as an ideal model system for in vivo target engagement studies of PAK1



ACKNOWLEDGEMENTS

TREAT-AD: treatad.org

MODEL-AD was established with funding from The National Institute on Aging (U54 AG054345).

AD Knowledge portal: <u>adknowledgeportal.org</u>

• AD Data Explorer: modeladexplorer.org