



Sex Differences in Brain Disorders: Emerging Transcriptomic Evidence and Implications for Therapeutic Development

Nilüfer Ertekin-Taner, M.D., Ph.D.
Professor of Neurology and Neuroscience
Mayo Clinic Florida

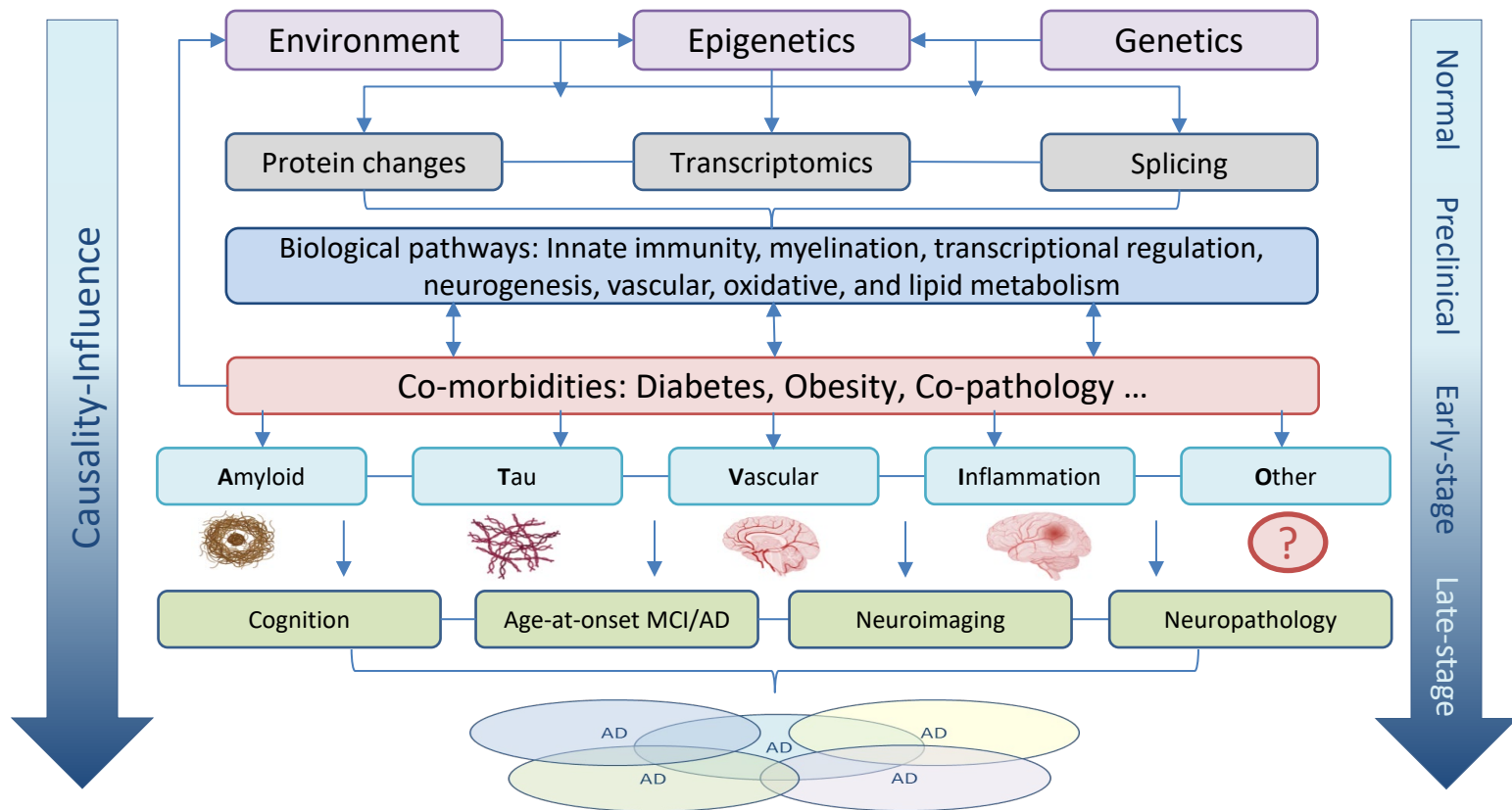
National Academy of Sciences, Engineering, Medicine
Part B: Neurodevelopmental and Neurodegenerative Disorders



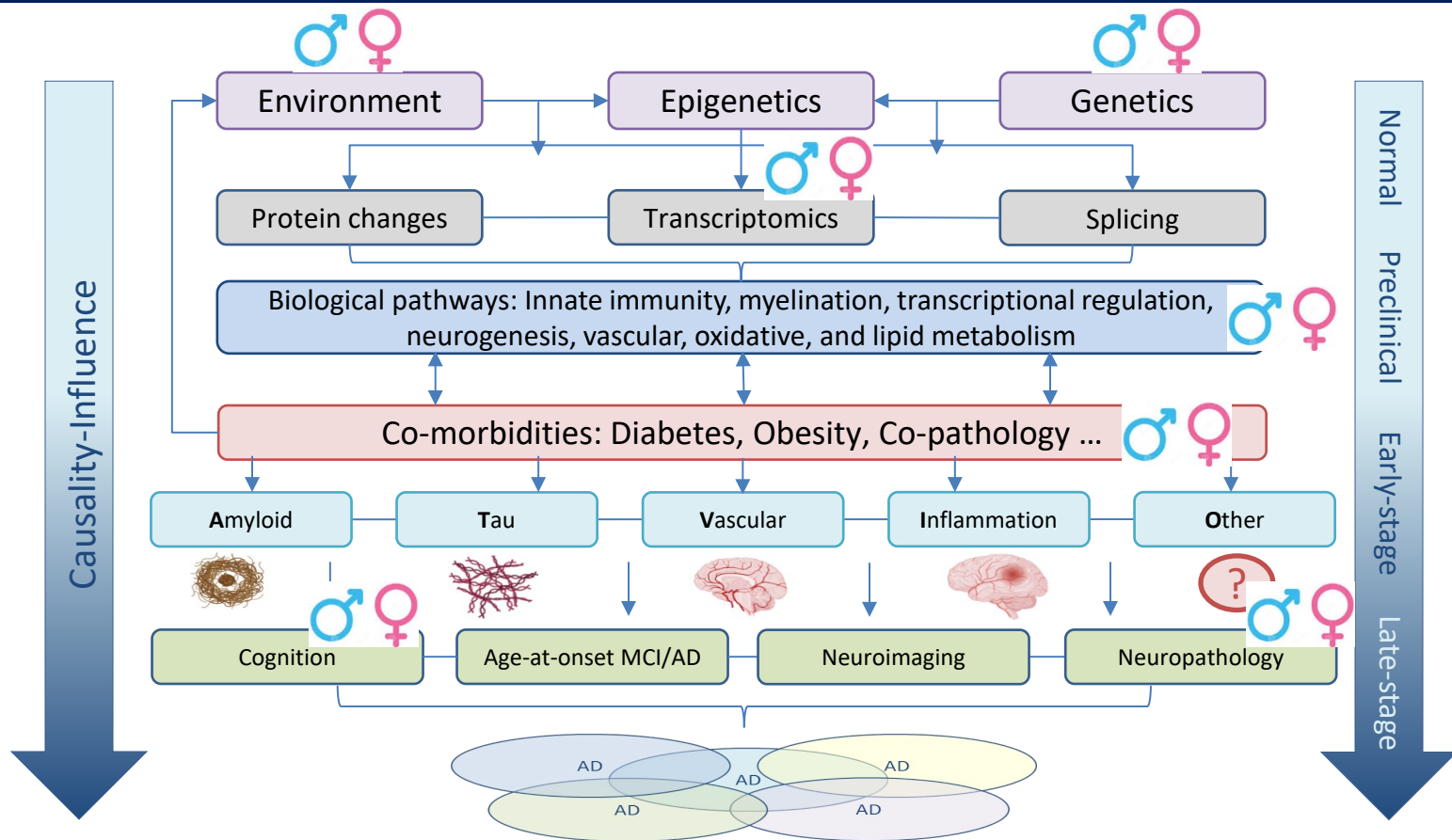
September 23rd, 2020



A (Simplified) Model of Neurodegenerative Diseases



A (Simplified) Model of Neurodegenerative Diseases



Evidence for Sex Differences in Alzheimer's Disease – “It's complicated”

- **Epidemiology:**

- Women have 2x lifetime risk of Alzheimer's disease (AD) than men.
- Higher prevalence in women-but not all geographical regions.
- Higher incidence in low/middle income countries for women.
- Suggested explanations: Lower education in women, survival bias against men (death due to cardiovascular disease higher in middle age), genetic, hormonal, but no definitive studies.

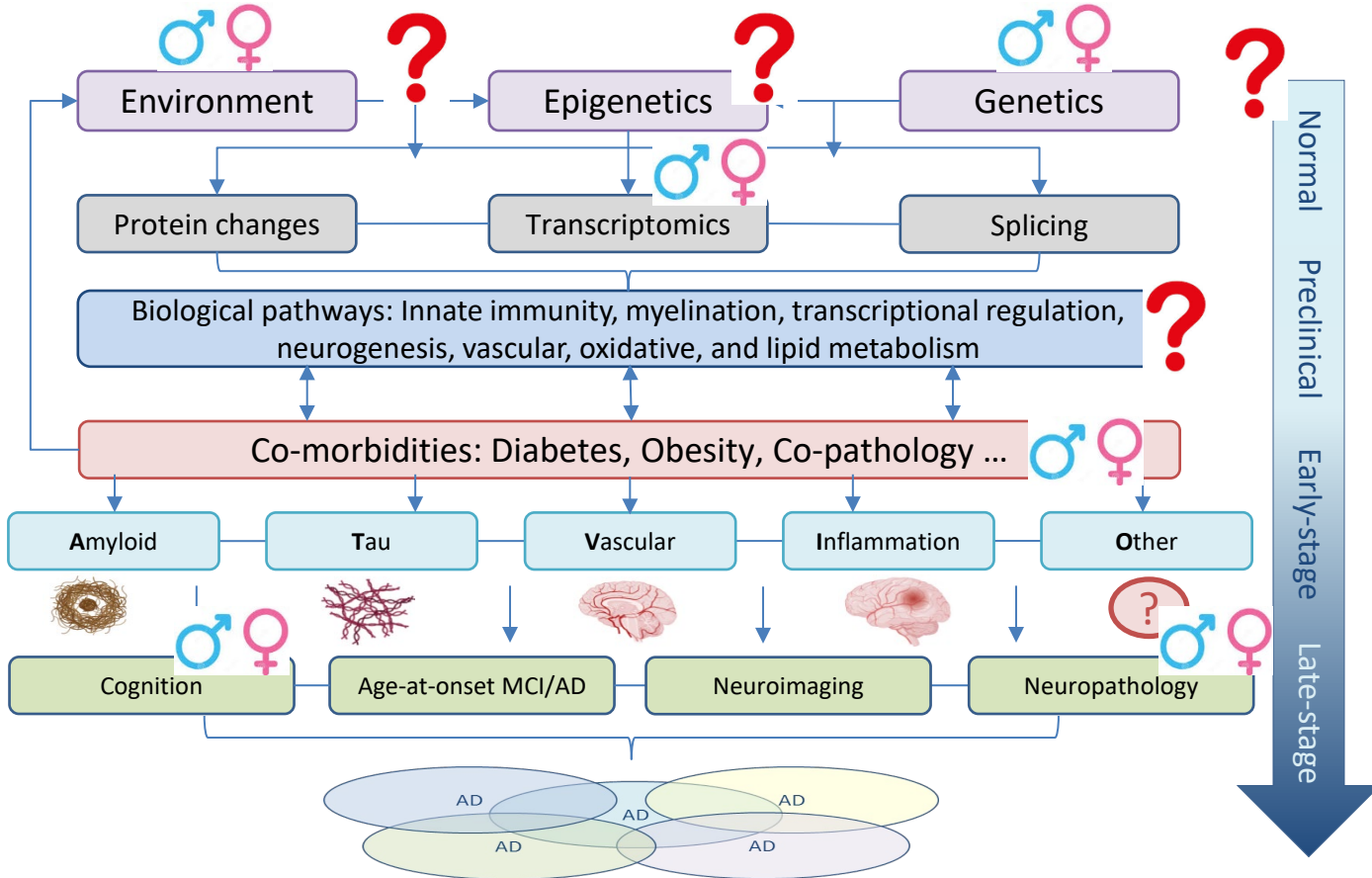
- **Symptomatology:**

- Women with AD show higher rates of cognitive decline.
- Some studies show greater rates of behavioral symptoms and dependence in women with AD.

- **Neuropathology:**

- Some but not overwhelming evidence for differences in tau or A β by neuropathology or biomarker studies (CSF, imaging).
- However, for given AD neuropathology or biomarker levels, women have higher odds of dementia and faster cognitive decline.

Sex Differences – Knowledge Gaps



- Effect of sex-differences in environment on epigenome -> transcriptome -> disease pathways.
- Impact of lifelong changes in hormonal status.
- Longitudinal changes in peripheral transcriptome -> disease progression/outcomes.
- Sex differences in multi-ethnic populations.

Sex Differences in Brain Disorders – Bridging the Gap to Therapies

- **Establish and enhance well-powered longitudinal cohorts:**
 - Peri-menopausal to post-menopausal aged women and age-matched men.
 - Repeated measures of clinical, cognitive, biomarker outcomes.
 - Peripheral (blood, saliva, CSF) multi-omics (transcriptome, epigenetic, proteome, metabolome) and genetic data.
 - Ideally matched brain tissue/data.
 - Enhance inclusion of underrepresented populations.
- **Integrative analyses:**
 - Effects of hormonal status, education, socio-economic factors on multi-omics and disease outcomes.
 - Role of genetic factors on multi-omics and (eQTL, mQTL, pQTL) disease outcomes.
 - Sex-stratified and sex-interaction analysis of all data (in addition to sex-adjusted analysis).
- **Model systems to study sex-differences:**
 - Investigate role of chromosomal sex, gonadal sex and hormones on disease-related outcomes, transcriptional and epigenetic changes in brain and peripheral tissue of model systems.
 - Study effects of chromosomal sex in iPSC-based neuronal and glial cellular and organoid models based on samples collected from male and female participants from multi-ethnic groups who represent the full disease spectrum.