

MEETING ABSTRACTS

BUTYRYLCHOLINESTERASE GENETIC POLYMORPHISM AND NEUROIMAGING BIOMARKERS IN ALZHEIMER'S DISEASE

DeBay, Drew R.¹; Maxwell, Selena¹; Luke, David¹; Fisk John D.²; Burrell, Steve³; Bowen Chris V.³; Song, Xiaowei; Black, Sandra E.⁵; Darvesh, Sultan^{1,6}

¹ Dalhousie University Departments of Medical Neuroscience, Halifax NS, Canada

² Dalhousie University Departments of Psychiatry, Halifax NS, Canada

³ Dalhousie University Departments of Radiology, Halifax NS, Canada

⁴ ImageTech Laboratory, Surrey Memorial Hospital, Health Science and Innovation, Fraser Health Authority, Surrey, BC, Canada

⁵ University of Toronto & Sunnybrook Health Sciences Centre Department of Medicine (Neurology), Toronto, Ont., Canada

⁶ Dalhousie University Departments of Medicine (Neurology), Halifax NS, Canada

Objective: The influence of butyrylcholinesterase (BChE) genetic polymorphism in Alzheimer's (AD) remains controversial. *BCHE-K* and *BCHE-A* genetic variants cause reduction of BChE, an enzyme implicated in AD. Some studies have reported a protective effect of *BCHE-K*, others suggest increased AD risk, particularly when associated with *APOE4*. We utilized a candidate gene-driven analyses to determine the effects of *BCHE-K* and *BCHE-A* on AD biomarkers using ADNI data (<http://adni.loni.ucla.edu/>).

Methods: Participants were genotyped for *BCHE-K* (615) and *BCHE-A* (785), each stratified into control (C), MCI or AD groups. MRI, ¹⁸F-FDG and amyloid-PET were assessed. ANCOVA compared main effects of i)diagnosis, ii)^{BCHE-K}, iii)*BCHE-A* and iv)*APOE4* status on each biomarker with age, education and sex as covariates.

Results: The allelic frequency was 20.8%, 4.6% and 26.5% for *BCHE-K*, *BCHE-A* and *APOE4*. For MRI, main effects for diagnosis were significant ($p < 0.0001$), with reduction in whole-brain and selected regional volumes (7-27%, $p \leq 6 \times 10^{-6}$) in AD vs. C. For 18FDG-PET, the main effect for diagnosis was significant ($p = 5 \times 10^{-9}$), with 14% decrease in metabolism in AD vs. C ($p = 7 \times 10^{-10}$). For amyloid-PET, the main effects for diagnosis and *APOE4* status were significant ($p = 0.034$; $p = 3 \times 10^{-6}$), with 12% increase in retention in AD vs. C ($p = 0.023$) and 16% increase among carriers of at least one *APOE4* allele vs. non-carriers ($p = 8 \times 10^{-6}$). No significant effects of these biomarkers were observed due to *BCHE-K* or *BCHE-A* status ($p \geq 0.209$).

Conclusions: These data suggest *BCHE-K* or *BCHE-A* may not significantly effect structural, metabolic or molecular AD biomarkers. Further ROI/voxel-wise analyses are warranted to uncover potential regional changes among AD *BCHE* variants.

Keywords: Alzheimer's disease; butyrylcholinesterase genetic variants; neuroimaging; amyloid-PET; FDG-PET