## TECHNOLOGY AND DEMENTIA PRECONFERENCE



PODIUM PRESENTATION

## Early identification of participants at risk of cognitive decline: An online tool for patient prediction of cognitive decline

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## Abstract

Background: The success of therapeutic options for treatment of Alzheimer's disease (AD) and the growing emphasis for such treatment to commence in the preclinical phase makes it necessary to have robust empirical models of clinical disease progression to understand findings from clinical trials, allow clinicians to evaluate effects of new drugs, and to select individuals for future trials. Such models have been developed from relatively small samples, with incomplete data/substantial loss to follow-up. The ADOPIC consortium provides the largest complete AD natural history sample to date. We applied our sigmoid models of disease progression to cognitive data from ADOPIC, and developed an algorithm to predict cognitive change.

Method: We developed an online application (Prediction of Alzheimer's Disease Progression Tool, [PADPT]) to estimate decline in cognition over the course of AD within 2,861 ADOPIC participants (1,434 Cognitively unimpaired [CU], 342 MCI [Mild Cognitive Impairment], 211 AD, and 847 progressed from CU→MCI or MCI→AD), with  $\geq 1$  PET Amyloid (A $\beta$ ) scan and  $\geq 36$  months of cognitive assessment with the pre-clinical Alzheimer's cognitive composite (PACC). Individual participant slopes were defined according to age and gender using linear mixed effects models. A $\beta$ + (≥50CL) participant rates of decline vs their mean PACC score were investigated using quantile polynomial regression. Within PADPT, single visit information on age, clinical classification and APOE £4 allele status is entered to compute the estimated annual rate of decline. Candidate results are mapped to the sigmoid curve to determine their predicted rate of cognitive decline over the chosen time period.

Result: Quantile regression bands (QRB's) aligned with participant cognitive decline. Individuals with stable cognition clustered to the top right of the plot (Figure 1). Mean slopes for APOE ε4 allele status separated QRB's for CU and AD, but not for MCI groups. All participants allocated to cognitive decline groups (past the first sigmoid

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inflection point) demonstrated decline in PACC score that remained in the QRB's, regardless of initial clinical classification.

**Conclusion:** Used in conjunction with baseline profile information, the PADPT application can assist with clinical decision making both pre and post-treatment and also identify suitable clinical trial candidates who, untreated, will show subtle decline in cognition.

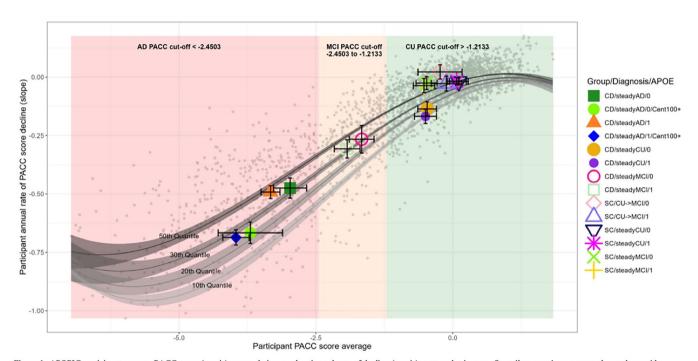


Figure 1: APOPIC participant average PACC score (x-axis) versus their annual estimated rate of decline (y-axis) scatter plot in grey. Quantile regression curves and error bars with coloured dots for participants with Centiloid>50 only. Profiles shown in legend for those with cognitive decline (classified as [CD]) and stable cognition (SC) groups, clinical diagnosis status: cognitively unimpaired (CU), mild cognitive unimpaired (MCI) and Alzheimer's disease (AD) participants as well as participants who progressed towards AD-dementia throughout the study (CU->MCI). APOE £4 allele non-carrier (0) and carrier (1) status. Error bars denote 95% confidence interval in both average PACC score and annual rate of decline. Based on these profiles, PACC cut-off values for a potential new candidate were estimated as CU to have a PACC score greater than -1.2133 (green box), MCI between -2.4503 to -1.2133 (orange box) and AD patient to have a PACC score lower than -2.4503 (red box). AD patients with Centiloid greater than 100 denoted as Cent100+.

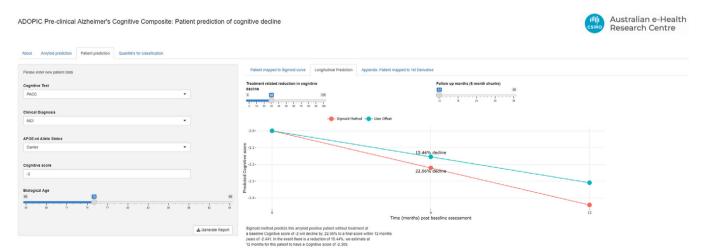


Figure 2: Screen shot of online web-based tool for pre-clinical Alzheimer's cognitive composite score (PACC) prediction. Suitable only for candidates with Centiloid 50 or more. Candidates clinical diagnosis, APOE ε4 allele status, baseline PACC score and age are entered on the bottom left, and using previous methods.