

Longitudinal Speech-based Cognitive Trajectories Of Csf Biomarker-positive Versus Biomarker-negative Individuals

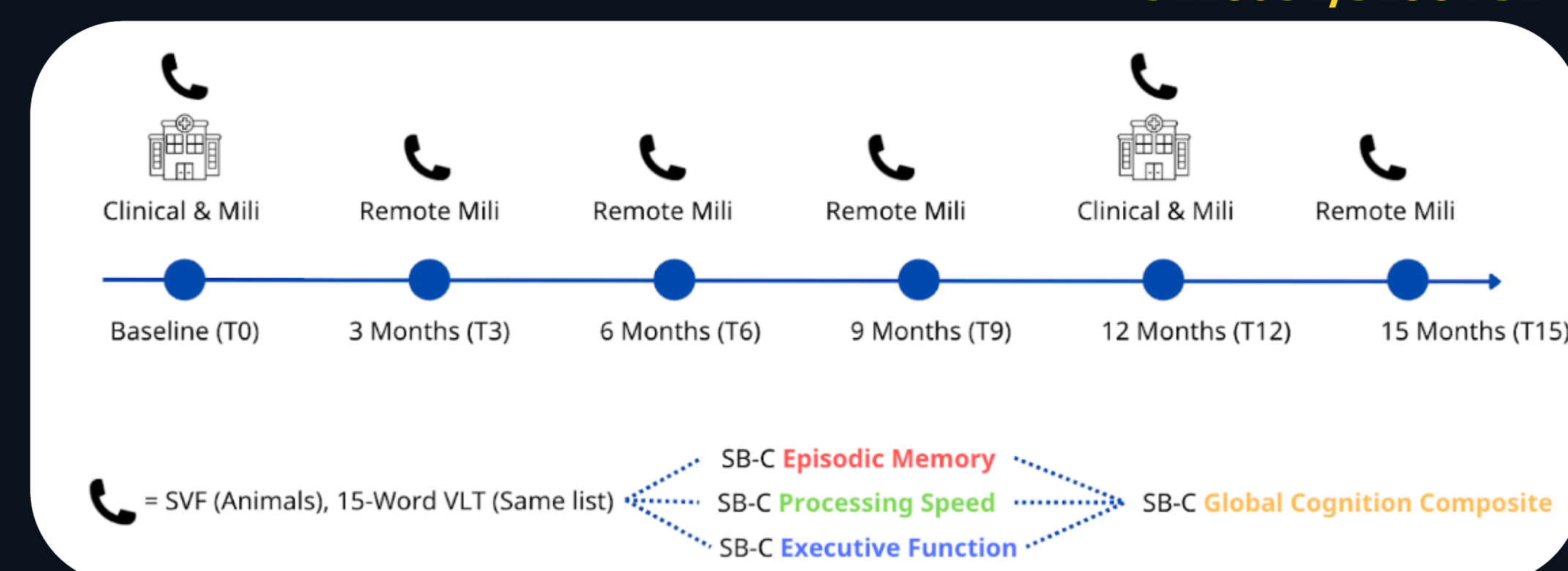
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Introduction

Cerebrospinal fluid (CSF) biomarkers provide evidence of Alzheimer's disease (AD) pathology, yet their relationship to cognitive change captured by digital tools remains unclear. Speech-based digital cognitive measures are sensitive to early impairment¹. Yet, whether they capture distinct longitudinal trajectories in biomarker-positive versus biomarker-negative individuals is unknown. Here, we examined whether **a speech-based global cognitive composite (SB-C) and its domain scores** captures longitudinal change as a function of CSF biomarker status.

Methods



Participants: $N = 50$ participants at baseline divided into biomarker positive ($N = 15$) and biomarker negative ($N = 35$) based on **CSF A β 42/P-tau 181**.

Statistics: Linear mixed-effects models examined longitudinal SB-C trajectories adjusting for covariates and participant-specific random effects. Estimated slopes and marginal means changes from baseline were derived. CSF was collected prior to speech; all biomarker-positive participants were included, while biomarker-negative participants were restricted to a CSF-to-baseline-speech gap ≤ 2.5 years.

Results

Longitudinally, there was a significant interaction between CSF-biomarker status x time for the **SB-C global** ($p = 0.03$) **and memory scores** ($p = 0.05$). **Biomarker-negative individuals improved faster over time** indicating **learning-related gains (Fig.1)**. No significant interaction was found for the SB-C executive function ($p = 0.16$) and processing speed scores ($p = 0.10$)

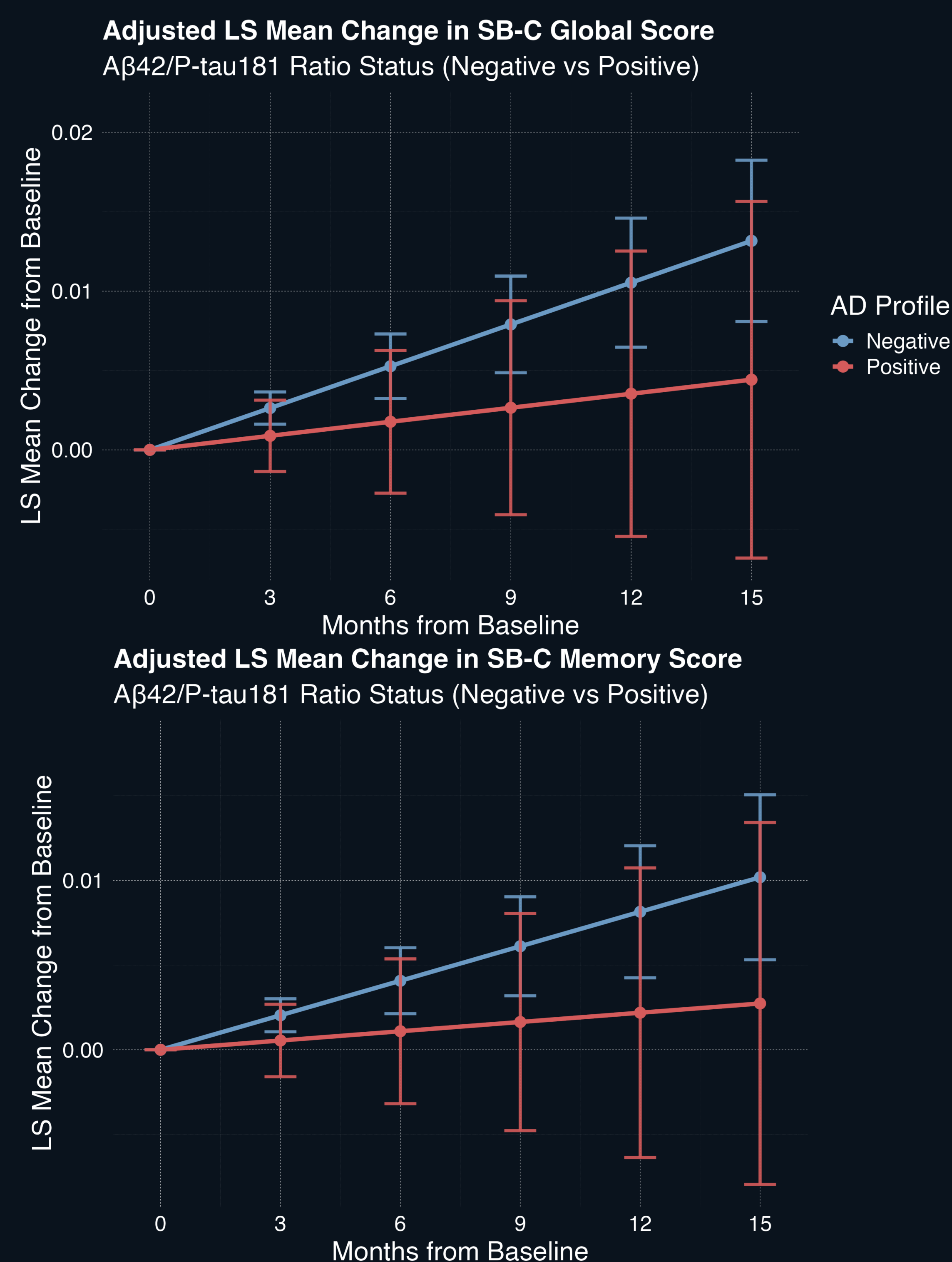


Figure 1. Adjusted LS Mean Changes from Baseline in SB-C Scores

Conclusion

Despite modest sample size and temporal separation between CSF collection and speech, **SB-C captured longitudinal patterns by CSF biomarker status**. Improvement confined to biomarker-negative individuals, potentially reflecting preserved learning/practice-related gains. Findings highlight the potential of speech-based markers in complementing biomarker-informed stratification and improving **longitudinal monitoring**.

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