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Individualized clinical management of patients at risk for Alzheimer's dementia

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Data Sharing Statement

All de-identified data related to the specific outcomes of this study are sharable with a written data-sharing and publication agreement, and will be available (along with data dictionary) three months after publication. Requests can be made to Pentara Corporation at shendrix@pentaracorp.com. The study protocol, statistical analysis plan, and informed consent form will be available upon publication. Requests for these documents can be made to rii9004@med.cornell.edu. Neuroimaging data (e.g., MRI, FDG-PET, Amyloid PET) will be unavailable until full publication of those results in the future.

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Summary

INTRODUCTION: Multi-domain intervention for Alzheimer's disease (AD) risk reduction is an emerging therapeutic paradigm.

METHODS: Patients were prescribed individually-tailored interventions (education/ pharmacologic/non-pharmacologic) and rated on compliance. Normal cognition/subjective cognitive decline/preclinical-AD were classified as *Prevention*. Mild cognitive impairment due to AD/mild-AD were classified as *Early Treatment*. Change from baseline to 18-months on the modified-Alzheimer's Prevention Cognitive Composite (primary outcome) was compared against matched historical control cohorts. Cognitive aging composite (CogAging), AD/cardiovascular risk-scales, and serum biomarkers were secondary outcomes.

RESULTS: 174 were assigned interventions (age 25–86). Higher-compliance Prevention improved more than both historical cohorts (*P*=.0012,*P*<.0001). Lower-compliance Prevention also improved more than both historical cohorts (*P*=.0088,*P*<.0055). Higher-compliance Early Treatment improved more than lower-compliance (*P*=.0007). Higher-compliance Early Treatment improved more than historical cohorts (*P*<.0001,*P*=.0428). Lower-compliance Early Treatment did not differ (*P*=.9820,*P*=.1115). Similar effects occurred for CogAging. AD/cardiovascular risk-scales and serum biomarkers improved.

DISCUSSION: Individualized multi-domain interventions may improve cognition and reduce AD/cardiovascular risk scores in patients at-risk for AD-dementia.

Keywords

Alzheimer's disease prevention; Multi-domain interventions; Alzheimer's Prevention Clinic; Personalized medicine; Preclinical Alzheimer's disease

Introduction

Late-life Alzheimer's disease (AD) develops over an extended preclinical period.^{1–4} Considering over 46 million people in the United States alone have preclinical AD, this pre-

dementia period offers a unique opportunity for early intervention to address modifiable risk. $_5$

Given the paucity of effective AD treatments, prevention or delay of dementia is essential. Further, AD drug trials may have been more successful if initiated earlier in the disease course.⁶ It is therefore important to evaluate the effectiveness of AD interventions across the disease spectrum, especially in at-risk individuals before clinically-evident decline.

Population-attributable risk models estimate that risk factor modification (e.g., hypertension, insulin resistance, physical inactivity, hearing loss, depression) may prevent up to one-third of AD cases.^{7,8} These targetable risk factors may influence AD pathological pathways (e.g., glucose hypometabolism, inflammation, oxidative stress, amyloid burden, trophic factors).^{8,9} The Finnish Geriatric Intervention Study to Prevent Cognitive Impairment and Disability (FINGER) study was the first large long-term randomized controlled trial (RCT) showing multi-domain interventions (nutrition/physical activity/cognitive training) can maintain cognitive function and reduce the risk of cognitive impairment among at-risk older adults from the general population.^{10,11} Other RCTs applying lifestyle modifications have demonstrated similar effects in mild cognitive impairment (MCI) participants and adults at-risk for cognitive decline.^{12,13} However, encouraging data from RCTs require translation to clinical practice, including verification of how patient compliance (or "dose response") affects outcomes.¹⁴

Considering the heterogeneity of AD pathology, the application of precision medicine allows for interventions that can be targeted for individual patients.^{12,15} The National Institutes of Health defines precision medicine as "an emerging approach for disease treatment and prevention that takes into account individual variability in genes, environment, and lifestyle for each person".¹⁶ An overall structure of how precision medicine may be achieved in the future will be through convergence of technological advances (e.g., big data, genomic sequencing, "-omics" technologies, systems biology, integrated disease modeling) as it is hypothesized that deconstructing the disease into multiple subsets that exist within a heterogeneous population, and tailoring therapies accordingly, may be preferentially effective based on individual biological make-up (protein-protein interactions, epigenetic modifications, metabolic pathways).^{17,18} A term that has been used to adapt this approach, using currently available clinical assessments in everyday practice,¹⁹ is *clinical precision* medicine, where medical history (e.g., lifestyle patterns, life-course events), physical/ neurological examination, anthropometrics, commercially-available blood biomarkers (including genetics), and cognitive assessments inform a multi-modal management plan.^{20,21} Patients are followed longitudinally to evaluate the effectiveness of, and further refine, personally-tailored interventions. In 2013, an Alzheimer's Prevention Clinic (APC) was established in New York, with research collaboration in Puerto Rico.^{21,22} APC's mission is to mitigate late-life AD dementia risk by applying individualized clinical management strategies toward primary, secondary, and tertiary AD prevention while simultaneously studying its comparative effectiveness (Figure S1).²³

In this proof-of-concept trial, we investigated effects of multi-domain evidence-based individually-tailored interventions on cognition, AD/cardiovascular risk scores, and AD-risk biomarkers in real-world clinical practice.^{22,24}

2. Methods

2.1. Study design and participants

In this prospective comparative effectiveness trial, all patients requesting an APC clinical consultation between March 12, 2015 and January 10, 2018 were initially screened via telephone (Figure S2) for participation to achieve a pre-specified sample of at least 150 participants with baseline and post-intervention assessments (powered to detect a 3.5-point difference [SD 6.5] on the primary outcome with 90% power and a sample size of 75 participants in each compliance group; See Figure S3 for study design, Appendix A for power calculation). Inclusion criteria assessed via initial telephone screen were a family history of AD and no/minimal cognitive complaints. Exclusion criteria assessed during an in-person evaluation included a diagnosis of moderate-to-severe AD dementia or other dementia; disorders affecting safe engagement in interventions (e.g., malignant disease, major depression, psychotic disorder); or coincident participation in another trial. Participants with a clinical diagnosis of MCI or early mild dementia with negative Amyloid neuroimaging were also excluded (n=7). See CONSORT diagram for additional details (Figure S2).

Institutional Review Board approval was obtained on February 16th, 2015 and patients were consented to participate in the Comparative Effectiveness Dementia & Alzheimer's Registry (Protocol #1408015423). See Appendix B for consent procedures.

2.2. Procedures

Participants underwent a comprehensive screening evaluation: detailed clinical history, physical examination, anthropometrics, blood biomarkers, apolipoprotein-e4 (*APOE*-e4) genotyping, and cognitive assessment (Table S1 and detailed in prior publication).²² Additional assessments were ordered in symptomatic patients (incorporating American Academy of Neurology Guidelines²⁵), when indicated. Amyloid positron emission tomography (PET) or cerebrospinal fluid biomarkers were used to confirm/exclude AD pathology in participants with a clinical diagnosis of MCI or early mild dementia. Participants were diagnosed as normal cognition, subjective cognitive decline (SCD), preclinical AD (PRE), MCI due to AD, or early mild AD dementia incorporating the 2011 National Institutes of Health and the Alzheimer's Association diagnostic criteria (Appendix C).^{1,22,26,27}

Enrolled participants were given individualized, multi-domain intervention recommendations informed by clinical and biomarker data (methods previously described)²², and received a mean of 21 recommendations by a neurologist or family nurse practitioner (Figure 1). Categories of recommendations included patient education/genetic counseling, pharmacological approaches (medications/vitamins/supplements), non-pharmacological approaches (customized recommendations for exercise, nutrition, vascular

risk, sleep, cognitive engagement, stress, general medical care), and others based on methods previously published.²² Longitudinal follow-up occurred every 6-months with continual refinement of interventions for each participant. Upon follow-up, each participant was assessed as "compliant" or "not compliant" with each individual recommendation. A compliance score was calculated as a percentage of recommendations adhered to on a scale of 1–10 (1 represents 0–10% of recommendations, etc.) as independently assessed by two clinicians based on patient report at the visit and patient Likert-scale ratings. Clinicians then assigned an overall compliance score by consensus before review of any follow-up data. Higher-compliance participants were pre-specified as following >60% of all recommendations given, versus lower-compliance participants (60%).²⁸

As an example of the application of the previously published method of an individualized clinical approach, a peri-menopausal 59-year-old woman (apolipoprotein E4 [APOE $\varepsilon 3/\varepsilon 4$] heterozygote) without subjective cognitive complaints and a past medical history of untreated "borderline" hypertension (~140s/80s), hyperlipidemia and abdominal obesity, elevated waist-to-hip ratio (.93), elevated visceral body-fat, insulin resistance, elevated homocysteine and normal (albeit sub-optimal) memory function received 25 individualized recommendations.²² These included patient education about potential risks/benefits of longterm hormone replacement therapy, genetic counseling, referral to a preventative cardiologist for blood pressure control (goal 120s/70s) and consideration of a coronary calcium scan for cardiovascular risk stratification, exercise counseling including a targeted amount/type of aerobic-versus-resistance training (geared for body-fat reduction), nutrition advice centered on Mediterranean-style diet (emphasis on fatty fish and extra-virgin olive oil consumption to address elevated LDL and low HDL-cholesterol), while limiting high-glycemic foods (considering insulin resistance) and optimizing B-complex (B12/folate/B6) vitamin intake (considering elevated homocysteine) and cocoa flavanols (considering insulin resistance, elevated blood pressure and lower-than-expected memory performance), as well as several other detailed recommendations such as sleep hygiene, cognitive engagement/training strategies, stress management, ongoing care with primary care physician (Figure 1), and information on AD prevention clinical trials which she may soon qualify for based on age/ genotype.²² An introductory course on AD prevention (10 lessons, 2+ hours of interactivemultimedia content) that has been shown to increase knowledge and willingness to participate in AD prevention clinical trials is also recommended via the online learning portal AlzU.org.²⁹ On follow-up, she was given a compliance score of 8 based on clinical consensus, and was thus classified as a higher-compliance Prevention participant (based on following 71-80% of the 25 recommendations).

Adverse events were recorded during each follow-up, with the treating clinician asking all participants whether they experienced any side effects/harm related to assigned interventions. Trial registered at ClinicalTrials.gov ().

2.3. Outcomes

The primary outcome was change in performance on the modified Alzheimer's Prevention Initiative Cognitive Composite (m-APCC) from baseline to 18-months.³⁰ Statistical comparisons were performed between higher- and lower-compliance groups within each

diagnostic classification and against matched historical control cohorts: National Alzheimer's Coordinating Center (NACC) and Rush University Memory and Aging Project (Rush) (Figure 2).

The original APCC was empirically determined to document progression of preclinical cognitive decline related to AD progression, and was selected due to its concurrent use in two AD prevention clinical trials (Alzheimer's Prevention Initiative Generation Program, Autosomal-Dominant AD Trial).^{31,32} Similar to other trials,^{33,34} we refined the APCC based on the selection of tests administered (Table S2 and prior publication of neuropsychological measures used in our clinic).²⁴ Tests comprising the m-APCC were selected to represent the same cognitive domains as those used in the APCC.²⁴

Secondary outcomes included changes on a composite of neuropsychological tests associated with non-pathological cognitive aging (CogAging, Appendix D), two AD risk scales (Australian National University–AD Risk Index [ANU-ADRI], Cardiovascular Risk Factors, Aging and Incidence of Dementia [CAIDE]), two cardiovascular risk scores (American College of Cardiology/American Heart Association [ACC/AHA], Multi-Ethnic Study of Atherosclerosis [MESA]), and risk biomarkers (Table S1).^{35–38}

See Table S1, S9–S12/Appendix E for exploratory outcomes/results.

2.4. Statistical Analyses

2.4.1. General—Participants were classified based on clinical diagnosis and level of compliance (Figures 2, S1). Two-sided *P*-values were used for all comparisons with no correction for multiplicity due to the a priori intent to investigate the primary outcome separately within the diagnosis groups. Secondary analyses may be considered hypothesis-generating and not confirmatory.

2.4.2. Mixed Model Repeated Measures (MMRM)—Change from baseline in all outcomes was analyzed at 6, 12, and 18 months for the Full Analysis Set (FAS) using MMRM that included all available data for participants with at least one follow-up visit. Least squares mean (LSMEANs) estimates at each visit were reported and groups were compared with least squares differences (LSDIFFs). The primary model included diagnostic classification (Prevention/Early Treatment) and compliance (Lower/Higher) with Diagnosis×Compliance interaction, as well as age, baseline score, baseline Mini-Mental State Examination, and visit. LSMEANS estimates from the Diagnosis×Compliance interaction are shown for the primary analysis. The interaction between quantitative compliance and diagnosis group was used to assess whether compliance affected diagnosis groups differently. SAS® V9.4 PROC MIXED was used.

2.4.3. Historical Comparison—NACC (n=38836) and Rush (n=3289) were the two data repositories used to derive comparisons (as neither cohort received therapeutic interventions). See Table S3 for demographic comparisons. Participants were matched for age and m-APCC score at baseline within diagnosis category (Appendix F). However, MCI diagnoses in each cohort were not amyloid-confirmed unlike our cohort. Since the NACC dataset had APOE genotype, additional analyses were performed in APOE4 carriers which

were matched as a proxy for increased likelihood of amyloid positivity and potentially more comparable rates of decline to our amyloid-confirmed participants.^{39,40} The Rush cohort included data from the Religious Orders Study, Memory and Aging Project, and Minority Aging Research Study.^{41,42} Since the youngest Rush participant was >50, only our participants aged 50+ were used for this comparison in addition to using age for matching.

2.4.4. Compliance Adjusted Model—Since participant characteristics may affect compliance levels, predictors of compliance were assessed by fitting a stepwise regression model, with compliance as the outcome variable, and including APOE-e4 carrier status, age, gender, diagnostic classification, baseline cognitive scores, baseline blood biomarkers, baseline biometrics, and baseline risk scores as predictors. To assess the specific impact of compliance, significant baseline predictors of compliance (at α <.05) were identified and corrected for as covariates in the adjusted MMRM, which also included a term for Baseline×Time interaction. This adjusted model is compared to the primary model in Table 1.

2.4.5. Exploratory Analyses—Change in each AD-risk biomarker was assessed for correlation with m-APCC and CogAging to assess whether biomarker improvements were associated with corresponding improvements in cognition.

3. Results

3.1. Disposition

202 patients were screened via telephone and were scheduled for an in-person evaluation. Of these, 10 scheduled a visit but did not come and 18 did not meet inclusion/exclusion criteria (7 excluded due to clinical diagnosis of MCI or early mild dementia with negative amyloid imaging, 8 due to clinical diagnosis of mild to moderate AD, 2 due to history of major depression, and 1 due to diagnosis/ongoing treatment of multiple myeloma. Of the remaining 174 patients (ages 25–86), all were assigned interventions (Table S4). 154 participants (88.5%) had at least one post-baseline assessment and were included in the FAS analysis (Figure S2). Study discontinuation rate was 22.1% at 12-months and 26.6% at 18-months (Figure S2/Table S4). Of those allocated to treatment, 24 (15.6%) discontinued because the treating physician left the practice (relocation), while 17 (11.0%) were lost to follow-up. See Table S4 for disposition at each time-point.

3.2. Demographics and Baseline Characteristics

Baseline characteristics are reported in Table 3/Appendix G. There were no differences at baseline between the 20 participants who were assigned interventions but did not follow-up compared to those with at least one post-baseline assessment (Table S5). Of those who followed-up, >20% were born outside the United States and over one-third reside outside the New York-metropolitan area. Higher- and lower-compliance early treatment participants exhibited significant differences in m-APCC and CogAging at baseline, with no differences between Prevention compliance groups.

Serum biomarkers differed between higher- and lower-compliance Early Treatment groups only for glycated hemoglobin (HbA1c), and none between Prevention groups. Biometric baseline values were similar between higher- and lower-compliance groups in Prevention and Early Treatment (Table 3).

3.3. MMRM for Primary Outcome - m-APCC

3.3.1. Compliance by Diagnosis Group (Prevention vs. Treatment) Interaction (Figure 3)—In Prevention participants, higher- and lower-compliance groups showed significant improvements by 4.6 (95% CI=3.09–6.19, P<.0001) and 4.5 (2.24–6.84, P=. 0002) points on the m-APCC, respectively. There was no difference between these groups (-2.79–2.61, P=.9488). In Early Treatment participants, the higher-compliance group increased by 4.8 points but this was not significant (-1.06–10.67, P=.1073). The lower-compliance Early Treatment group had significant worsening by 6.0 points (-10.83, -1.20, P=.0148). The difference between these groups (10.8 points) was significant (4.67–16.97, P=. 0007).

3.3.2. Historical Comparison for the Primary Outcome (Table 2)—The highercompliance Prevention group improved more than NACC by 3.1 (1.14–5.06, P=.0012) and Rush by 4.9 (2.55–7.25, P<.0001). The lower-compliance Prevention group improved more than NACC by 2.9 (0.74–5.06, P=.0088) and Rush by 4.0 (1.26–6.74, P=.0055). The highercompliance Early Treatment group improved more than NACC by 10.3 (5.99–14.61, P<. 0001) and Rush by 5.3 (0.20–10.40, P=.0428). Lower-compliance Early Treatment did not differ from NACC (P=.9820) or Rush (P=.1115).

See Table S6 for additional analyses matching our amyloid confirmed participants to enriched NACC participants who were APOE4 carriers.

See Figure S4 for additional details.

3.4. Adjustment for Baseline Factors Predictive of Compliance

3.4.1. Predictors of Compliance—The baseline compliance model identified three baseline parameters that significantly predicted compliance: baseline HbA1c (P<.0001), baseline ACC/AHA risk score (P<.0001), and baseline homocysteine (P=.0225). Each extra percentage of baseline HbA1c predicted a 32.5 percentage point increase in compliance on average. An increase of 10 points on the ACC/AHA risk scale predicted a 7 percentage point decrease in compliance on average. An increase of 1 µmol/L of homocysteine at baseline predicted a 2 percentage point increase in compliance on average. The interaction analysis for quantitative compliance and diagnosis resulted in a statistically significant interaction for compliance by diagnosis (p=0.0049) and compliance by diagnosis by visit (p=0.0003). Each extra point of compliance (complying with an additional 10% of recommendations) results in 0.06 point improvement in APCC at 18 months within the non-MCI group (p=0.8547), and 2.41 points of improvement in the MCI group (p=0.0003).

The adjusted model resulted in notably similar estimates of change on the m-APCC (see m-APCC[adjusted] in Table 1), suggesting that differences in m-APCC performance for lower-

and higher-compliance groups were not explained by baseline characteristics predictive of compliance or differing rates of progression depending on baseline scores.

3.5. Secondary endpoints

3.5.1. Cognitive Aging Changes (Non-AD Specific)—For Prevention participants, CogAging showed a mean improvement of 2.6 (0.6) years for the higher-compliance group (1.47–3.67, P<.0001) and 3.4 (0.8) years for the lower-compliance group (1.73–5.09, P<.0001) (difference=-0.8 (-2.84–1.16, P=.4069)). Early Treatment participants improved by 2.0 (2.3) years in CogAging for the higher-compliance group but the change was not significant (-2.48–6.48, P=.3786), and worsened by 5.9 (1.8) years for the lower-compliance group (2.3–9.48, P=.0015) (difference=7.9 (3.52–12.26, P=.0005)).

3.5.2. Risk Scales (Table 1 and Figure S5).—For ANU-ADRI at 6-months, Prevention decreased by 2.8 (0.5) points for higher-compliance (1.76-3.75, *P*<.0001) and decreased by 1.2 (0.6) for lower-compliance (0.01-2.35, *P*=.0480) (difference=1.6 [-0.01-3.15, *P*=.0508]). Early Treatment decreased by 5.9 (2.1) for higher-compliance (1.73-10.11, *P*=.0060) and decreased by 3.9 (1.7) for lower-compliance (0.52-7.27, *P*=.0240) (difference=2.0 [-0.87-4.92, *P*=.1695]).

For CAIDE at 18-months, Prevention decreased by 0.1 (0.1) points for higher-compliance (-0.14-0.25, *P*=.6053) and did not change 0.0 (0.1) for lower-compliance (-0.26-0.33, *P*=. 8247) (difference=0.0 [-0.33-0.37, *P*=.9177]). Early Treatment decreased by 0.9 (0.3) for higher-compliance (0.19–1.53, *P*=.0120) and decreased by 0.7 (0.3) for lower-compliance (0.14–1.35, *P*=.0170) (difference =0.1 [-0.59-0.83, *P*=.7389]).

For ACC/AHA cardiovascular at 18-months, Prevention decreased by 3.8 (0.4) points for higher-compliance (3.05–4.49, *P*<.0001), and decreased by 2.8 (0.4) for lower-compliance (2.06–3.60, *P*<.0001) (difference=0.9 [0.08–1.79, *P*=.0317]). Early Treatment decreased by 10.4 (3.0) for higher-compliance (4.54–16.30, *P*=.0006) and decreased by 13.0 (2.4) for lower-compliance (8.20–17.78, *P*<.0001) (difference=2.6 [-3.28-8.42, *P*=.3867]).

For MESA at 18-months, Prevention decreased by 1.7 (0.2) points for higher-compliance (1.39–1.99, P<.0001) and decreased by 1.4 (0.1) for lower-compliance (1.17–1.64, P<.0001) (difference=0.3 [0.04–0.61, P=.0891]). Early Treatment decreased by 2.7 (0.7) for higher-compliance (1.37–3.95, P<.0001) and decreased by 2.7 (1.0) for lower-compliance (0.73–4.68, P=.0076) (difference=0.1 [-1.73–1.86, P=.9557]).

3.5.3. Serum Risk Biomarkers—In Prevention participants, improvements were found in HDL-C (6.0, P < 0.0001), hs-CRP (-1.3, P < 0.0001), adiponectin (2.1, P < 0.0001) and 25hydroxy-vitamin D (4.5, P=0.0010). In Early Treatment participants, fibrinogen (-40.2, P=0.0269), homocysteine (-1.0, P=0.0416), HDL-C (10.0, P=0.0095), hs-CRP (-1.8, P=0.0006), adiponectin (5.1, P=0.0001) and Lp(a) Mass (14.6, P=0.0035) improved. No biomarker changes were significantly correlated with either change in m-APCC or change in CogAging across all patient groups, with the exception of cystatin C. A worsening in cystatin C of 0.1 point corresponded to greater improvement in CogAging by 1.2 years (P=0.0227). Table S7 shows the mean change in biomarkers from baseline to 18 months.

These changes were compared between the diagnostic groups and correlated with change in cognitive outcomes. See Table S7 for all secondary and exploratory biomarker endpoints.

3.6. Safety Analysis

No serious adverse events were reported. Intervention-related adverse events occurred in 9.1% of participants (5.9% Prevention, 20% Early Treatment) (Table S8), including gastrointestinal complaints, myalgia/arthralgia, ankle sprain, irritability, insomnia, lethargy, fatigue, somnolence, nightmares, and anxiety (each <2%).

4. Discussion

To our knowledge, this is the first empirical trial in a clinical setting indicating that individualized AD risk factor management may improve cognitive function which may be related to AD pathology. In addition, secondary analyses demonstrated that multi-domain tailored interventions may reduce calculated AD and cardiovascular risk scores across a broad range of ages and diagnostic classifications, and may potentially have a cognitiveaging-modifying effect on non-pathological age-related cognitive decline. Within the Early Treatment group, cognitive improvements were seen only in the higher-compliance group, suggesting that close adherence to the interventions is needed to derive benefit within the context of definitive AD pathology. However, cognitive improvements were seen in both the higher- and lower-compliance Prevention participants, with both compliance groups demonstrating improvements compared to historical cohorts. Further, our population was easily and quickly recruited from a real-world clinical setting and the interventions were well-tolerated, adding to the translational value for practice (Appendix J). Additionally, while socio-economic factors will differ among varied cohorts, intervention-related costs negatively impacted adherence in 7.1% of participants (Appendix K).

Because m-APCC measures AD-related cognitive change, improvements may have resulted from targeting risk factors that lead to AD pathogenesis; however, direct evidence of changes in these pathways was not obtained. Additional evidence from longitudinal volumetric magnetic resonance imaging, fludeoxyglucose-PET and amyloid-PET would demonstrate whether observed improvements were related to disease modification. Neuroimaging data is forthcoming from a brain imaging substudy (n=135) begun in 2018.⁹ Furthermore, longitudinal measurement of potential key AD-related biomarkers, such as those related to neuroinflammation and synaptic dysfunction, may be incorporated into future studies to investigate the direct effects on AD pathology.

Cognitive aging composites indicated that the estimated delay of cognitive decline may be approximately three years in Prevention participants and two years in the higher-compliance Early Treatment group. Improvements in cognitive decline related to non-pathological cognitive aging may potentially be linked to reducing vascular dementia risk and/or targeting other factors (e.g., synaptic plasticity, alterations in neuronal structure, dysfunction of neuronal networks).⁴³ However, due to lack of cognitive aging biomarkers, biological factors related to this potential response were not measured and are thus unclear. Treatment effects observed using both cognitive composite measures may suggest that treatment response is broad. Therefore, addressing risk factors that collectively impact overall health may assist in

mitigating age-related cognitive decline, along with other potential health benefits stemming from treating comorbidities (e.g., cardiovascular risk).

We observed improvements in several AD-risk biomarkers. In Prevention participants, improvements were found in HDL-C, hs-CRP, adiponectin, and 25-hydroxy-vitamin D. In Early Treatment participants, fibrinogen, homocysteine, HDL-C, hs-CRP, adiponectin, and Lp(a) improved. However, none of these correlated with improvement in cognition. Unexpectedly, a worsening of cystatin C of 0.1 point corresponded to an improvement in CogAging. One possible explanation for lack of correlations between biomarkers and cognition is that such relationships likely involve multiple biomarker changes that may vary by person, as well as varied baseline values within a broad spectrum of reference ranges. A bayesian hierarchical analysis also did not identify an individual biomarker or category of biomarkers that was primarily associated with observed cognitive changes. See Appendix H/I for discussion.

While further study incorporating a host of biomarkers pre- vs. post-intervention may help to inform causality, we observed changes in efficacy outcomes such as serum biormarkers, anthropometrics and risk scales (Figure S6). These changes, along with the comparison of compliance groups, correction of multiple covariates, and matched historical comparisons, may suggest that these findings were potentially driven by the prescribed interventions.

Traditionally, treatment trials have attempted to isolate one effect at a time for single interventions, but the complexity of AD may require targeting multiple mechanisms simultaneously to affect disease progression. Our initial evidence of broad effects across risk scales, and measurements of cognition and biomarkers changing in expected directions, suggest this approach warrants further rigorous study.

Our study has several limitations. Our key limitation is the lack of a concurrent, randomized control group. Two considerations led to this design. A true control group may not have been possible, since well-informed participants actively enrolled in an AD risk reduction study may seek out and make lifestyle and/or other behavioral changes that impact outcomes. Additionally, given the setting of a real-world clinical practice where patients seek AD risk reduction care for modifiable risk factors in a clinic outside of a traditional solely research environment, it would not be feasible to withold treatment from a non-intervention randomized control group.

The disadvantage of an uncontrolled study is that it is unclear whether observed effects are due to baseline characteristics of participants or other aspects of general study conduct unrelated to treatment. In an attempt to mitigate these effects, we corrected the model for baseline predictors of compliance by including them as covariates to better ensure the improved outcomes were not primarily due to baseline characteristics. We also used historical comparison cohorts with similar demographics and matched each participant based on age and baseline m-APCC. We used historical comparisons to also help account for study procedure effect, such as practice effects. Compared to these matched historical comparisons, participants demonstrated greater improvements at similar time-points. Because these historical comparison groups were not part of any intervention, a response

associated with intervention expectations may potentially explain part of the cognitive benefit that was observed in our study. However, the 18-month duration is longer than is usually expected for this type of effect. Furthermore, improvements found in lab biomarkers and AD and CV risk scales are less likely to be influenced by placebo effects.⁴⁴ Future studies which include randomized non-intervention groups would allow for more definitive conclusions.

Because few NACC participants and no Rush participants had available amyloid biomarkers, we were unable to match on confirmed AD pathophysiology. After updating the matching algorithm to include APOE4 positivity as an enrichment strategy for NACC participants, our cohort continued to show cognitive benefit. While the lack of amyloid biomarkers is an important limitation, we would have expected the rate of cognitive decline in historical subjects without amyloid-confirmation to be slower than a matched population with amyloid confirmation, resulting in a more conservative estimate of the intervention effect. Unexpectedly, when enriching for APOE4, we observed less decline in the enriched population. Another limitation stems from the study environment of a real-world clinical practice and the challenges of rating compliance. There is a paucity of evidence on how to use compliance in comparative effectiveness studies as an outcome to differentiate treatment effects. While some studies have defined high compliance as following two-thirds of prescribed recommendations, we selected 60% for two reasons.^{45,46} An initial motivation was that a cut-point of 60% led to a roughly even number of patients in higher- and lowercompliance categorizations when care was previously provided in the clinic (from 2013-14 prior to initiating the comparative effectiveness study). Further, we applied categorizations from a prior study quantifying compliance into 4 groups: noncompliant (compliant to treatment schedule less than 20% of the time), low (20% to 59% of the time), moderate (60% to 79% of the time), and high (80% of the time) compliance.²⁸ Based on this study framework, we divided our participants into higher (60%) and lower (<60%) compliance groups. See Appendix K/Table S13 for additional information.

It is important to note that because lower compliance is often related to disease severity, statistical corrections for baseline m-APCC, HbA1c, homocysteine, AHA/ACC, and age were applied to decrease the possibility of bias due to these potential confounders. Further, the separation of diagnosis and compliance groups was critical due to a strong compliance by diagnosis interaction effect.

While our sample size was modest and further stratification led to relatively small diagnostic and compliance groups, observed effects seen were of a large enough magnitude to still be detectable. Continued recruitment across additional sites globally (n=1000 planned) will allow for confirmation of these proof-of-concept results and more detailed analysis of patient subgroups (e.g., age, ethnicity), biomarkers, and intervention approaches. Expanded recruitment may enable deeper understanding of precision effects and more definitive conclusions, and allow assessment of the impact of medical comorbidities and concomitant medications.

Practice effects due to repeated cognitive test exposure is another potential concern. To mitigate this, we administered alternate test forms at each time-point and required that

participants complete simulated at-home cognitive assessments prior to baseline. This also primed participants to testing conditions/procedures and mitigated test anxiety in effort to reduce practice effects. Also, practice effects on cognitive measures tend to occur at briefer test-retest intervals than those involved in this study, and the comparison to historical cohorts who took related measures repeatedly demonstrated improvement beyond what can reasonably be explained by practice effects.⁴⁷

While the m-APCC was our primary outcome, there is no gold standard for which cognitive measures should be used (and how often), and the degree of benefit which should be expected.⁴⁸ Cognitively normal patients at-risk may have a lower ceiling for benefit as they do not yet manifest cognitive decline. As such, assessment scales cannot be easily repeated from prior treatment trials, and novel composite measures sensitive to pre-dementia decline may hold promise.^{24,48} Because the study was conducted in the real-world clinical setting and one of the treating clinicians left the practice due to geographical relocation, 24 participants (58.6%) were lost to follow-up for this reason. Future studies should consider safeguards to account for similar factors that can substantially influence discontinuation rate. However, because the major contributing factor to discontinuation would not be expected to be related to response to treatment, it may be less likely that loss of these patients introduces bias in our results.

Furthermore, while patients who seek risk reduction care tend to be highly motivated, this approach may not be as effective in patient populations with lower motivation. Further discussion about factors related to compliance are detailed in Appendix K. Also, despite the study's translational value, long-term effects are unknown. Longitudinal assessments are ongoing. Additionally, while the median age in our cohort was 61 years, and the mean age was 60 years, our cohort included a broad age-range due to younger, middle-aged, and older patient demand in a real-world clinical setting. Nevertheless, the majority of participants (~75%) were aged 50 years or older. Age was included as a continuous linear covariate in the primary model, and as such, all estimated changes were for an average aged person (60 years old). The Prevention group had 0.1 points less improvement per year of age, and the Early Treatment group had 0.2 points less improvement per year of age. As such, an older population may demonstrate less improvements in cognition and, similar to AD drug trials, this intervention may be more effective in younger and/or less impaired populations. Future studies are warranted to more deeply understand age effects of this intervention.

With a clear understanding of these limitations, we offer this framework as a tool for clinicians across a broad range of subspecialties, and clinician researchers, to approach patient care while further clarifying its effectiveness (see Figure S7 to visualize levels of personalization). Given the magnitude of disease, significant morbidity of late-life dementia, and growing interest in applying *preventative* neurology to clinical care, it is important to report these findings as larger studies are developed and while our own sample size grows. Overall, these results help extend prior RCT/observational findings into a clinic setting where individualized lifestyle modifications produced measurable benefits.

From a practical clinical perspective, individualized AD risk factor management maybe applied for care to tens of millions of patients at-risk for AD dementia.

Our study adds to the body of comparative effectiveness research by applying the same framework across distinct pre-dementia diagnostic classifications, and provides a reproducible model for future research.

Further study in a large, multi-site, international cohort study, merits consideration.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Research in Context

Systematic Review

Authors searched ClinicalTrials.gov and World Health Organization's International Clinical Trial Registry Platform to identify multidomain precision medicine intervention studies to delay cognitive decline in patients at-risk for AD. Search terms were "prevention of dementia OR prevention of Alzheimer's" and "precision medicine OR personalized medicine." While several randomized controlled trials utilizing multidomain interventions were found, no completed precision medicine studies were identified. One recruiting study investigating an individualized intervention () was found yet results are not available.

Interpretation

To our knowledge, this is the first empirical trial to demonstrate individualized multidomain interventions may improve cognitive function and reduce AD/ cardiovascular risk scores in patients at risk for AD dementia in real-world clinical practice.

Future directions

Given the paucity of treatments and extended preclinical period, focus on AD risk reduction is essential. This study provides a feasible framework for studying AD risk reduction in clinical practice. Further research on individualized multi-domain interventions is warranted in larger cohorts across sites globally.

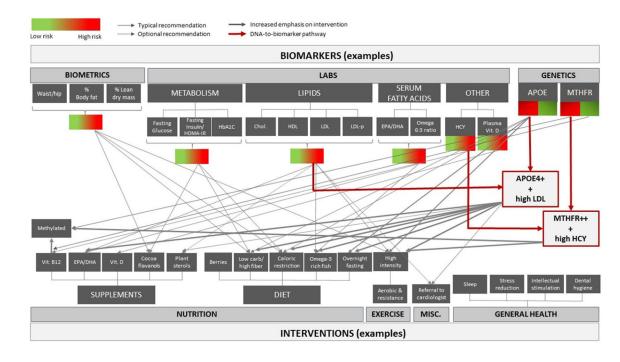


Figure 1: Example Biomarker to Intervention Paradigm

NOTE. Each data point collected during the initial clinical intake and evaluation, as well as at each follow-up visit, is used to inform which precision medicine interventions are recommended per participant.



Figure 2: Comparison Groups

NOTE. Participants were classified to reflect the different biological phases along the AD continuum (Figure S1) and level of compliance into one of the following four analysis groups: Higher-compliance Prevention, Lower-compliance Prevention, Higher-compliance Early Treatment, and Lower-compliance Early Treatment. Each group was compared to two matched historical control cohorts, NACC and Rush (n=38836 and n=3289, respectively)

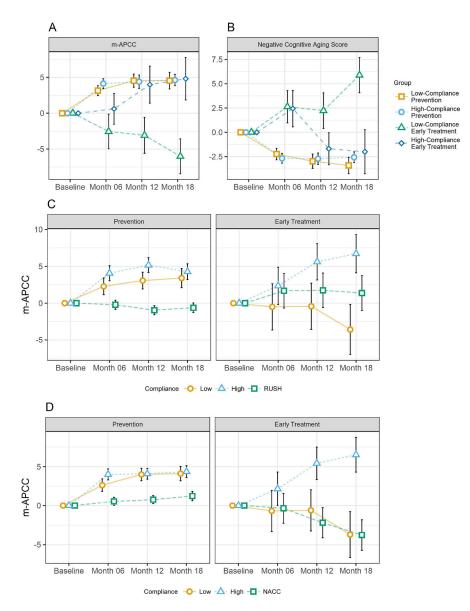


Figure 3: m-APCC(a) and Cognitive aging(b), NACC comparison(c), and Rush comparison(d) NOTE: A) Change from Baseline on the m-APCC at 18 months amongst the four diagnosis x compliance groups. B) Change from Baseline on the non-pathological CogAging composite at 18 months amongst the four diagnosis x compliance groups. C) Comparison of change in m-APCC between higher-compliance, lower-compliance, and Rush control (matched for baseline m-APCC and age). D) Comparison of change in m-APCC between higher-compliance, and NACC control (matched for baseline m-APCC and age).

Table 1:

Comparison of Prevention vs. Treatment Groups for Lower- vs. Higher-Compliance

Compliance	Visit	Lower-	P-val	Higher-	P-val	Higher- vs. Lower-	P-val
Prevention							
m-APCC	Mo. 6	3.2 (0.7)	< 0.0001	4.1 (0.7)	< 0.0001	1.0 (1.0)	0.3091
	Mo. 12	4.5 (0.9)	< 0.0001	4.4 (0.9)	< 0.0001	-0.1 (1.3)	0.9143
	Mo. 18	4.5 (1.2)	0.0002	4.6 (0.8)	< 0.0001	0.1 (1.4)	0.9488
m-APCC	Mo. 6	3.0 (0.7)	< 0.0001	4.0 (0.7)	< 0.0001	1.0 (0.9)	0.2863
(adjusted)	Mo. 12	4.4 (0.9)	< 0.0001	4.4 (1.0)	< 0.0001	0.0 (1.3)	0.9821
	Mo. 18	4.5 (1.2)	0.0002	4.7 (0.9)	< 0.0001	0.2 (1.3)	0.9022
CogAging	Mo. 6	-2.2 (0.6)	0.0002	-2.7 (0.5)	< 0.0001	-0.4 (0.8)	0.5693
	Mo. 12	-3.0 (0.7)	< 0.0001	-2.7 (0.6)	< 0.0001	0.3 (0.9)	0.7849
	Mo. 18	-3.4 (0.8)	< 0.0001	-2.6 (0.6)	< 0.0001	0.8 (1.0)	0.4069
ANU-ADRI	Mo. 6	-1.2 (0.6)	0.048	-2.8 (0.5)	< 0.0001	-1.6 (0.8)	0.0508
CAIDE	Mo. 18	0.0 (0.1)	0.8247	-0.1 (0.1)	0.6053	0.0 (0.2)	0.917
ACC/AHA	Mo. 18	-2.8 (0.4)	< 0.0001	-3.8 (0.4)	< 0.0001	-0.9 (0.4)	0.0317
MESA	Mo. 18	-1.4 (0.1)	< 0.0001	-1.7 (0.2)	< 0.0001	-0.3 (0.2)	0.0891
Early Treatm	ent						
m-APCC	Mo. 6	-2.5 (2.4)	0.2941	0.6 (2.1)	0.7782	3.2 (2.2)	0.1463
	Mo. 12	-3.1 (2.5)	0.2221	4.0 (2.6)	0.1253	7.1 (2.4)	0.0044
	Mo. 18	-6.0 (2.4)	0.0148	4.8 (3.0)	0.1073	10.8 (3.1)	0.0007
m-APCC	Mo. 6	-3.3 (2.4)	0.1726	0.0 (2.4)	0.9861	3.2 (2.2)	0.1365
(adjusted)	Mo. 12	-4.3 (2.6)	0.1057	3.2 (2.9)	0.2724	7.5 (2.5)	0.0037
	Mo. 18	-7.6 (3.1)	0.0140	3.9 (3.2)	0.2298	11.5 (3.5)	0.0007
CogAging	Mo. 6	2.6 (1.7)	0.1161	2.4 (1.9)	0.1946	-0.2 (1.5)	0.8973
	Mo. 12	2.2 (1.8)	0.2244	-1.7 (1.7)	0.3076	-3.9 (1.8)	0.0348
	Mo. 18	5.9 (1.8)	0.0015	-2.0 (2.3)	0.3786	-7.9 (2.2)	0.0005
ANU-ADRI	Mo. 6	-3.9 (1.7)	0.024	-5.9 (2.1)	0.0060	-2.0 (1.5)	0.1695
CAIDE	Mo. 18	-0.7 (0.3)	0.0170	-0.9 (0.3)	0.0120	-0.1 (0.4)	0.7389
ACC/AHA	Mo. 18	-13.0 (2.4)	< 0.0001	-10.4 (3.0)	0.0006	2.6 (3.0)	0.386
MESA	Mo. 18	-2.7 (1.0)	0.0076	-2.7 (0.7)	< 0.0001	0.1 (0.9)	0.955

Table 2:

m-APCC comparison to Historic Controls

			Change from Baseline Within Groups			Difference Between Groups in Change from Baseline		
Analysis	Visit	Statistic	Lower- Compliance	Higher- Compliance	Historic Control	Higher- vs. Lower- Compliance	Lower- vs. Historic	Higher- vs. Historic
Prevention	n							
NACC	Mo. 6	LSMean (SE)	2.6 (0.8)	4.0 (0.7)	0.6 (0.5)	1.4 (1.0)	2.1 (0.9)	3.5 (0.9)
		p-value	0.0011	< 0.0001	0.2975	0.1897	0.0284	< 0.0001
	Mo. 12	LSMean (SE)	4.0 (0.8)	4.1 (0.7)	0.8 (0.5)	0.1 (1.0)	3.2 (0.9)	3.3 (0.9)
		p-value	< 0.0001	< 0.0001	0.1397	0.9301	0.0008	0.0001
	Mo. 18	LSMean (SE)	4.1 (0.9)	4.4 (0.8)	1.2 (0.6)	0.2 (1.2)	2.9 (1.1)	3.1 (1.0)
		p-value	< 0.0001	< 0.0001	0.0385	0.8343	0.0088	0.0012
Rush	Mo. 6	LSMean (SE)	2.3 (1.1)	4.1 (1.0)	-0.2 (0.6)	1.8 (1.5)	2.5 (1.3)	4.3 (1.2)
		p-value	0.0456	< 0.0001	0.7273	0.2266	0.0518	0.0003
	Mo. 12	LSMean (SE)	3.1 (1.1)	5.2 (1.0)	-1 (0.6)	2.1 (1.5)	4.0 (1.3)	6.1 (1.2)
		p-value	0.0075	< 0.0001	0.125	0.152	0.0019	<0.0001
	Mo. 18	LSMean (SE)	3.4 (1.3)	4.3 (1.1)	-0.6 (0.7)	0.9 (1.6)	4 (1.4)	4.9 (1.2)
		p-value	0.0092	< 0.0001	0.343	0.5945	0.0055	< 0.0001
Early Trea	atment							
NACC	Mo. 6	LSMean (SE)	-0.7 (2.6)	2.2 (2.1)	-0.3 (1.9)	2.8 (2.8)	-0.3 (2.4)	2.5 (2.2)
		p-value	0.7957	0.3156	0.8571	0.3066	0.8877	0.2452
	Mo. 12	LSMean (SE)	-0.6 (2.6)	5.4 (2.1)	-2.2 (1.9)	6 (2.7)	1.6 (2.4)	7.6 (2.1)
		p-value	0.8186	0.0096	0.2533	0.029	0.5077	0.0004
	Mo. 18	LSMean (SE)	-3.7 (2.9)	6.5 (2.2)	-3.8 (1.9)	10.2 (3.1)	0.1 (2.7)	10.3 (2.2)
		p-value	0.2064	0.0036	0.0546	0.0011	0.9820	< 0.0001
Rush M	Mo. 6	LSMean (SE)	-0.5 (3.1)	2.4 (2.5)	1.7 (2.3)	2.9 (3.2)	-2.2 (2.8)	0.7 (2.5)
		p-value	0.8734	0.3529	0.4729	0.3798	0.4339	0.7917
	Mo. 12	LSMean (SE)	-0.4 (3.1)	5.6 (2.5)	1.7 (2.3)	6.0 (3.2)	-2.2 (2.8)	3.9 (2.5)
		p-value	0.8930	0.0235	0.4573	0.0617	0.4377	0.1203
	Mo. 18	LSMean (SE)	-3.6 (3.4)	6.7 (2.6)	1.4 (2.4)	10.3 (3.6)	-5.0 (3.1)	5.3 (2.6)
		p-value	0.2946	0.0102	0.5611	0.0041	0.1115	0.0428

Table 3:

Patient Demographics and Baseline Characteristics

NOTE: Denominator for percentages is the number of subjects with observed data for the variable within each category. SD=Standard Deviation, Min=Minimum, Max=Maximum. Due to missing data, sample sizes for the cognitive outcomes listed in Table 1 ranged from 110–119 in the prevention group and 32–34 in the treatment group at baseline.

		Prever	ntion	Early Treatment		
Variable	Subcategory or statistic	Lower- Compliance	Higher- Compliance	Lower- Compliance	Higher- Compliance	Total N=154
Gender	Female	37 (68.5%)	33 (50.8%)	8 (40%)	8 (53.3%)	86 (55.8%)
	Male	17 (31.5%)	32 (49.2%)	12 (60%)	7 (46.7%)	68 (44.2%)
Diagnosis	MCI			17 (85%)	15 (100%)	35 (22.7%)
	Mild AD			3 (15%)		
	Normal	35 (64.8%)	44 (67.7%)			79 (51.3%)
	Pre-clinical AD	2 (3.7%)	4 (6.2%)			6 (3.9%)
	Subjective Cognitive Decline	17 (31.5%)	17 (26.2%)			34 (22.1%)
Age Group		41 (75.9%)	40 (61.5%)		3 (20%)	84 (54.5%)
	Age Median (61)					
		13 (24.1%)	25 (38.5%)	20 (100%)	12 (80%)	70 (45.5%)
	Age>Median (61)					
APOE-e4 Group [*]	Heterozygotes	21 (39.6%)	25 (38.5%)	12 (60%)	3 (20%)	61 (39.9%)
	Homozygotes	4 (7.5%)	6 (9.2%)	3 (15%)	4 (26.7%)	17 (11.1%)
	Non-Carriers	28 (52.8%)	34 (52.3%)	5 (25%)	8 (53.3%)	75 (49%)
Race	White	46 (85.2%)	59 (90.8%)	16 (80%)	9 (60%)	130 (84.4%)
	Other	5 (9.3%)	4 (6.2%)	1 (5%)	3 (20%)	13 (8.4%)
	Missing	3 (5.6%)	2 (3.1%)	3 (15%)	3 (20%)	11 (7.1%)
Age	Mean (SD)	53.9 (11.9)	57.4 (11.4)	74.4 (6.3)	73.1 (8.2)	59.9 (13.2)
	Diff. (p-value) 3.67 (0.0906)		0906)	1.28 (0.6019)		
BMI	Mean (SD)	25.1 (3.8)	24.8 (3.5)	26.5 (4.5)	25.6 (4.2)	25.3 (3.9)
	Diff. (p-value)	0.26 (0.6971)		0.93 (0.5374)		
Education Level	Mean (SD)	15.9 (1.05)	16.1 (0.8)	15.3 (1.2)	15.7 (0.6)	15.9 (0.9)
	Diff. (p-value) 0.16 (0.5822)		0.33 (0.6779)			
Cognitive Scores						
m-APCC	Mean (SD)	72.1 (8.00)	71.62 (9.24)	42.03 (8.60)	54.98 (14.54)	65.50 (13.97)
	Diff. (p-value)	1.25 (0.4595)		12.95 (0.0035)		
Cognitive Aging	Mean (SD)	54.98 (6.46)	56.44 (6.63)	74.95 (7.75)	68.69 (9.56)	59.53 (9.97)
	Diff. (p-value)	1.47 (0.	2271)	6.26 (0.0400)		

		Prevention		Early Treatment			
Variable	Subcategory or statistic	Lower- Compliance	Higher- Compliance	Lower- Compliance	Higher- Compliance	Total N=154	
MMSE	Mean (SD)	29.56 (0.64)	29.39 (1.28)	26.80 (2.02)	28.07 (2.70)	28.9 (1.72	
	Diff. (p-value)	0.17 (0	.4050)	1.27 (0.	1255)		
Risk scores		•					
ACC	Mean (SD)	6.29 (8.58)	8.34 (8.68)	31.84(18.77)	25.17 (13.74)	12.3 (14.30	
	Diff. (p-value)	2.05 (0	.2024)	6.68 (0.			
ANU-ADRI	Mean (SD)	11.33 (9.36)	10.88 (8.64)	28.35 (12.73)	26.67 (10.15)	14.8 (11.82	
	Diff. (p-value)	(p-value) 0.46 (0.7829)		1.68 (0.	6765)		
	Mean (SD)	3.98 (2.43)	4.28 (2.48)	4.35 (1.81)	4.00 (2.07)	4.16 (2.34	
CAIDE	Diff. (p-value)	0.30 (0	.5155)	0.35 (0.	5985)		
MESA	Mean (SD)	2.58 (2.00)	3.87 (3.66)	9.65 (8.30)	8.07 (6.58)	4.58 (5.08	
	Diff. (p-value)	1.29 (0.0220)		1.58 (0.	5467)		
Biomarkers	•						
Cystatin C	Mean (SD)	0.79 (0.17)	0.81 (0.15)	0.94 (0.28)	0.95 (0.14)	0.83 (0.18	
	Diff. (p-value)	0.02 (0.4833)		0.02 (0.8493)			
Fibrinogen	Mean (SD)	333.19 (64.04)	319.36 (59.83)	382.17(73.98)	401.38 (92.35)	340.1 (71.77	
	Diff. (p-value)	13.83 (0.2329)		19.21 (0.5059)			
HbA1c	Mean (SD)	5.28 (0.35)	5.36 (0.33)	5.37 (0.26)	5.62 (0.30)	5.36 (0.34	
	Diff. (p-value)	0.08 (0.2130)		0.25 (0.0127)			
HDL Cholesterol	Mean (SD)	65.03 (15.62)	68.81 (21.04)	67.44 (32.54)	63.74 (22.08)	66.8 (21.21	
	Diff. (p-value)	3.78 (0.2764)		3.70 (0.7072)			
Homocysteine	Mean (SD)	9.58 (2.24)	9.72 (2.64)	10.57 (2.81)	10.06 (2.72)	9.82 (2.53	
	Diff. (p-value)	0.14 (0.7531)		0.51 (0.5947)			
HOMAIR	Mean (SD)	2.06 (1.66)	1.81 (1.24)	2.52 (2.18)	1.89 (1.57)	2.00 (1.56	
	Diff. (p-value)	0.26 (0.4109)		0.63 (0.4564)			
hs-CRP	Mean (SD)	1.67 (2.09)	1.58 (3.35)	4.37 (11.78)	6.39 (19.03)	2.44 (7.69	
	Diff. (p-value)	0.09 (0.8616)		2.02 (0.7010)			
	Mean (SD)	121.98 (42.63)	108.34 (37.44)	108.14(62.02)	125.33 (61.33)	114.7 (45.72	
LDL Cholesterol Direct	Diff. (p-value)	13.64 (0).0657)	17.19 (0.4207)			
Lp(a) mass	Mean (SD)	35.48 (38.18)	31.13 (42.16)	33.50 (24.86)	34.17 (23.45)	33.2 (37.63	
	Diff. (p-value)	4.36 (0.6813)		0.67 (0.9628)			
Triglycerides	Mean (SD)	88.57 (61.85)	75.53 (42.06)	85.46 (48.98)	107.46 (59.49)	84.5 (52.71	
	Diff. (p-value)	13.04 (0.1757)		22.0 (0.2388)			

		Prevention		Early Tre		
Variable	Subcategory or statistic	Lower- Compliance	Higher- Compliance	Lower- Compliance	Higher- Compliance	Total N=154
Vitamin D	Mean (SD)	38.97 (13.67)	42.00 (13.94)	36.93 (11.78)	40.88 (14.84)	40.17 (13.66)
	Diff. (p-value)	3.03 (0.	3.95 (0.2	3865)		
Biometrics/vital	signs					
	Mean (SD)	27.08 (6.96)	26.00 (7.54)	28.43 (6.30)	29.75 (5.87)	26.98 (7.09)
Body fat percentage	Diff. (p-value)	Diff. (p-value) 1.08 (0.4987)			1.32 (0.6271)	
	Mean (SD)	18.32 (2.22)	18.99 (2.07)	18.31 (2.22)	18.21 (1.29)	18.61 (2.09)
Dry lean mass percentage	Diff. (p-value)	0.67 (0.1400)		0.10 (0.8978)		
Waist-to-hip ratio	Mean (SD)	1.12 (0.08)	1.12 (0.10)	1.07 (0.11)	1.13 (0.16)	1.12 (0.10)
	Diff. (p-value)	0.00 (0.	7967)	0.06 (0.2		
Pulse	Mean (SD)	68.95 (9.60)	67.88 (11.89)	67.76 (10.95)	67.25 (7.21)	68.17 (10.52)
	Diff. (p-value)	1.07 (0.	6422)	0.51 (0.8880)		
Systolic blood pressure	Mean (SD)	122.80 (14.30)	119.20 (13.83)	136.00 (15.26)	130.42 (18.57)	123.89 (15.66)
	Diff. (p-value)	3.61 (0.	2236)	5.58 (0.3828)		
	Mean (SD)	73.22 (11.23)	70.41 (9.88)	74.47 (7.04)	70.33 (13.62)	71.93 (10.44)
Diastolic blood pressure	Diff. (p-value)	2.81 (0.	2057)	4.14 (0.2935)		

* One patient declined APOE testing.

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