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Establishing Biological Plausibility for Cognitive Frailty

by

Lana Jean Sargent

A dissertation submitted to the faculty of the Medical University of South Carolina in partial fulfillment of the requirements for the degree of Doctor of Philosophy in the College of Nursing.

July/2017

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TABLE OF CONTENTS

ACKNOWLEDGEMENTS	iv
LIST OF TABLES	vi
LIST OF FIGURES	vi
CHAPTERS	
ABSTRACT	1
INTRODUCTION	2
MANUSCRIPT 1:.....	13
Assessing the Current State of Cognitive Frailty: Measurement Properties.....	13
MANUSCRIPT 2.....	21
Determining Biological Factors for Cognitive Frailty: A Systematic Review.....	21
MANUSCRIPT 3:.....	41
Establishing Biological Plausibility for Cognitive Frailty: A Population Predictive Model	41
MANUSCRIPT 4:.....	71
Anticholinergic Burden is a Predictor of Cognitive Decline, Physical Frailty and Cognitive Frailty	71
SUMMARY	90
APPENDICES	94
MANUSCRIPT 1: Supplemental documents	94
MANUSCRIPT 2: Supplemental documents	107
MANUSCRIPT 3: Supplementary Methods, Statistical and Genomic Analyses	144
MANUSCRIPT 4: Anticholinergic Burden Scale.....	179
IRB APPROVAL.....	182
TRAINING AIM DOCUMENT	183

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LIST OF TABLES

Manuscript 1

Table 1.Operational definitions of Cognitive Frailty 15

Table 2.Use of markers for Cognitive Frailty..... 17

Manuscript 2

Table I. Cognitive decline biomarkers by category and frequency 34

Table II. Frailty biomarkers by category and frequency..... 35

Table III. Cognitive frailty biomarkers by category and frequency 36

Table IV. Serum and genetic correlations by phenotype 37

Manuscript 3

Table I. Genomic features by phenotype model I..... 61

Table II. Genomic features by phenotype model II..... 61

Table III. Protein and clinical features by phenotype model I..... 62

Table IV. Protein and clinical marker features by phenotype model II..... 64

Manuscript 4

Table 1. Characteristics of participants by phenotype..... 85

Table 2. Distribution of anticholinergic burden score by phenotype and difference between health control and phenotype 86

Table 3. Generalized linear regression results: association between anticholinergic burden and phenotypes 87

Table 4. Ordinal regression results: association between anticholinergic burden and phenotype..... 87

LIST OF FIGURES

Figure1. Complex systems theory for Cognitive Frailty..... 11

Manuscript 1

Figure I. PRISMA flow diagram of study selection and citation analysis..... 14

Manuscript 2

Figure I. PRISMA flow diagram of study selection and citation analysis⁶ 32

Figure II. Systematic review publication date range 33

Manuscript 3

Figure 1. Study approach workflow diagram 51

Manuscript 4 (NA)

ABSTRACT

Cognitive frailty is considered a potentially reversible age-related condition characterized by the simultaneous presence of both physical frailty and cognitive decline. The concept of cognitive frailty existing in older adults is indisputable, although the mechanisms and the directional relationship behind the dynamic association remain unexplained. Mechanisms have been suggested, often linking cognitive frailty to cognitive impairment or as a component of frailty but without an understanding of the biological bases for these associations we cannot not move forward with intervention trials.

This dissertation examines the biological mechanisms for cognitive frailty. The study is the first to use a large number of protein and genetic markers identified by a systematic review to define the underlying pathology for cognitive frailty. We use an innovative Boosted trees machine learning technique for developing a population based predictive model. Xgboost is based in boosted trees and provides more efficient and accurate predictive modeling with large datasets and a rapid / robust framework for feature selection. Statistical modeling is used to design, test, and validate an accurate method for and identifying and classifying the features that predict individuals with cognitive frailty. The tree boosting model is used for the evaluation of multiple variables simultaneously and provides a high predictive value with low bias.

The results presented within this dissertation create a foundation of understanding for a new aging condition and encourage translational research focused on the detection and prevention of cognitive frailty.

INTRODUCTION

“I forget what I was trying to say, one word or another gets in the way of the word I meant to use. Nothing stays. So I say something else, I compensate....are these the words I meant to say? But wait, are these the words I meant to say? These words migrate, they refuse to stay in place. This is my new life, my new way, I forget what I was trying to say.” Sherman Alexie.

Caregivers of patients with cognitive decline and patients themselves will suggest that their symptoms for memory loss and changes in physical function came long before they received a diagnosis by their provider. A report on the economic implications of cognitive decline estimates in 2015 there are 5.1 million individuals(1,2). With the aging “baby boomer” generation the trajectory that individuals will exhibit cognitive decline will be 13.5 million by the year 2050 in the United States(1,2). Efforts to unravel the mechanisms for cognitive decline have led to the recognition of a unique cluster of individuals who present with the simultaneous presences of both physical frailty and cognitive impairment without dementia(3). Both cognitive decline and physical frailty independently lead to increased disability, falls, mortality, an increase in health service need, and high direct/indirect costs to healthcare, often long-term care and hospitalization(4,5). Individuals with physical frailty and cognitive impairment may have a higher risk for disability than individuals with isolated physical frailty or cognitive impairment. Yet, historically, most research groups have excluded older adults with cognitive impairment from frailty studies(4). The International Consensus Group organized by the International Academy on Nutrition and Aging (I.A.N.A) and the

International Association of Gerontology and Geriatrics (I.A.G.G) convened in 2013 to identify related domains of physical frailty and cognition and termed the phenomenon “cognitive frailty”(3).

Establishing a model to detect cognitive frailty

The Institute of Medicine Report on Cognitive Aging described a need to develop an operational definition of cognitive frailty for use in research, clinical detection, and public health surveillance(6). A model for detecting cognitive frailty could provide practitioners with the tools needed for early detection and secondary prevention. Currently, the instrumental assessments for cognitive frailty are time-consuming, expensive, and require extensive training, and the clinical translation properties are not clear(3). The translation of the cognitive frailty construct into the clinical setting is limited by the lack of consensus on an operational definition and considerable heterogeneity and complexity in the diagnostic criteria. The primary purpose of this research was to create a population predictive model to gain a more in-depth understanding of the underlying biological mechanisms for cognitive frailty as currently defined by the International Consensus Group in 2013. This dissertation focuses on defining the shared mechanisms for physical frailty and cognitive impairment and establishing a model for determining the presence of risk factors that may predict cognitive frailty in the clinical setting. The model will advance the development of an operational definition by determining whether the potential risk factors at present may predict cognitive frailty in the clinical setting.

Mechanisms behind cognitive frailty

The mechanisms and the directional relationship behind the dynamic association of physical frailty and cognitive impairment or cognitive frailty remain unexplained.

Pathological events leading to cognitive frailty years before the onset of cognitive decline may be marked by epigenetic modifications that influence memory-associated gene transcription. However, to date, no investigators have simultaneously characterized the trajectory of cognitive decline and physical function, underlying cellular events that include physiological factors, and epigenetic modifications. The results presented here will further explicate the shared mechanisms, including putative biomarkers for physical frailty and cognitive impairment to enhance our understanding of the shared neuropathology in a secondary data analysis. Such an understanding will lead to intervention studies focused on preventing disability and mortality, decreasing health service use, and improving health outcomes for older adults.

OPERATIONAL DEFINITIONS

The extent to which we can predict cognitive frailty using biomarkers depends on the accuracy that our behavioral markers have on early identification. Screening for the detection of cognitive decline (i.e. neuropsychological) and frailty is determined by the identification tools for defining individuals with cognitive frailty. Individuals with cognitive frailty present with a unique neuropsychological profile, scoring worse on executive and attention tests with individuals having 3 or more of the frailty criteria being more impaired than individuals with only 1 of the frailty criteria(7). This dissertation focused on markers for early detection therefore, definitions used to

establish phenotype sub-groups in this study were structured to detect early cognitive decline including pre-frail individuals using neuropsychological testing focused on executive and attention memory domains. The definitions used are as follows:

Cognitive decline – mild neurocognitive disorders

Evidence of modest cognitive decline from a previous level of performance in one or more cognitive domains (complex attention, executive function, learning and memory, language, perceptual motor, or social cognition) with a modest impairment in cognitive performance by standardized neuropsychological testing or clinical assessment in absence of a diagnosis of dementia(8,9).

Frailty

The operational definition for frailty is defined as a clinical syndrome condition including 3 out of the 5 criteria related a physical phenotype including: 1) weak muscle strength (grip strength), 2) slow gait speed, 3) unintentional weight loss, 4) exhaustion and low physical activity(4). Pre-frailty includes 1 or 2 of the criteria is present, identifying a sub-group of individuals potentially progressing to frailty(4).

Cognitive frailty

The International Consensus Group (I.A.N.A. /I.A.G.G.) report is an acknowledgment of the need to focus research efforts on a clinical condition characterized by the co-occurrence of physical frailty and cognitive impairment, in absence of overt dementia diagnosis or underlying neurological conditions(3). The cognitive frailty construct is considered a heterogeneous clinical syndrome in older adults with evidence of: 1)

physical frailty and cognitive impairment (Clinical Dementia Rating score of 0.5); and 2) exclusion of a clinical diagnosis of Alzheimer's Disease or other dementia(3).

Details on the cut-off scores used to define the phenotypes are explained in further detail in manuscripts 3 and 4.

INNOVATION

An important innovation in this study was the use of machine learning (ML) statistical modeling to build a predictive model for cognitive frailty while further defining the unique features for cognitive decline and frailty. We use Boosted trees, a machine learning technique for supervised learning, these are ensembles of regression trees, similar to decision trees and are used for prediction or classification. Xgboost is based in boosted trees and provides more efficient and accurate predictive modeling with large datasets and a rapid / robust framework for feature selection. Statistical modeling is used to design, test, and validate an accurate method for classifying patients into phenotypic outcomes. The tree boosting model for the evaluation of multiple variables simultaneously provides a high predictive value with low bias. The second innovation in this study is the defining of putative biomarkers related to cognitive frailty leading to a better understanding of the interrelated neuropathology between physical frailty and cognitive impairment. The study is the first to use a large number of protein and genetic markers (n=289) identified by a systematic review to define the underlying pathology for cognitive frailty.

Impact of Proposed Research

Developing and validating a model for the detection and classification of cognitive frailty will improve the ability to detect patients with a potentially reversible cognitive and physical decline. Identification of biomarkers and an understanding of the physiological and genetic factors for cognitive frailty will help distinguish between changes related to normal aging, irreversible pathological process, and specific neurological diseases that may be reversible(6). The findings will encourage new research and may lead to effective interventions for the prevention and treatment of cognitive and physical decline in an aging population.

THEORETICAL FRAMEWORK

This dissertation used Complex Systems Theory as a primary theoretical framework. Complex Systems Theory (CTS) is an approach to science that involves multiple factors that interact nonlinearly to form a dynamic set of relationships leading to physiological change(10). Based in the tradition of ontology, CTS can identify the grouping together of the mechanistic elements of biology and the heuristic elements of philosophy to model the linkages that create a complex concept such as cognitive frailty. Biological mechanisms, proteins or gene expression and their patterns of interaction are inherently complex systems about which numerous empirical data exist (in this case within population databases) that are “dynamic and transformational” vs. inductive assumptions (11,12). Computational methods developed in bioinformatics are uniquely designed to analyze and interpret large amounts of biological data. This dissertation

created a theoretical framework based on the modeling of complex systems using bioinformatics (figure 1).

SPECIFIC AIMS

This dissertation consists of four manuscripts; 1) an integrative review assessing the measurement properties for cognitive frailty, 2) a systematic review exploring the biological factors for cognitive frailty, 3) a population based modeling study establishing biological plausibility for cognitive frailty, and 4) additional analysis of a unique feature from the modeling study and potential epigenetic factor for cognitive frailty; anticholinergic burden's association with cognitive decline, physical frailty, and cognitive frailty.

Aim 1. To determine associations between putative biomarkers and cognitive frailty as currently defined by the International Consensus Group in 2013 using a focused secondary analysis of the InCHIANTI study dataset.

1a. Establish a predictive model using statistical methodologies using an integrative approach to precisely define and predict cognitive frailty based on overlapping risk factors for frailty and cognitive decline.

1b. Establish a relationship among measurable physiological, clinical factors, and the development of cognitive frailty.

1c. Establish associations between physical frailty and cognitive parameters (i.e., losses in specific types of memory and mental acuity).

Manuscript 1 includes a comprehensive review of the measurement tools for defining the phenotype cognitive frailty. Manuscript 2 includes a large systematic review

of the potential putative clinical, protein, and genetic biomarkers for cognitive frailty. The markers identified in this comprehensive review were used as predictors in the population modeling study. Manuscript 3, is the population based predictive model analysis. Findings from the model study resulted in anticholinergic burden as a unique predictor of cognitive decline, frailty, and cognitive frailty. Considering anticholinergic medication burden could be a potentially reversible cause for cognitive frailty additional analyses was completed which resulted in manuscript 4.

Aim 2. To determine associations between genetic biomarkers; single-nucleotide polymorphisms (SNPs) to explain the phenotypic variance for cognitive frailty using a focused secondary analysis of the InCHIANTI study dataset.

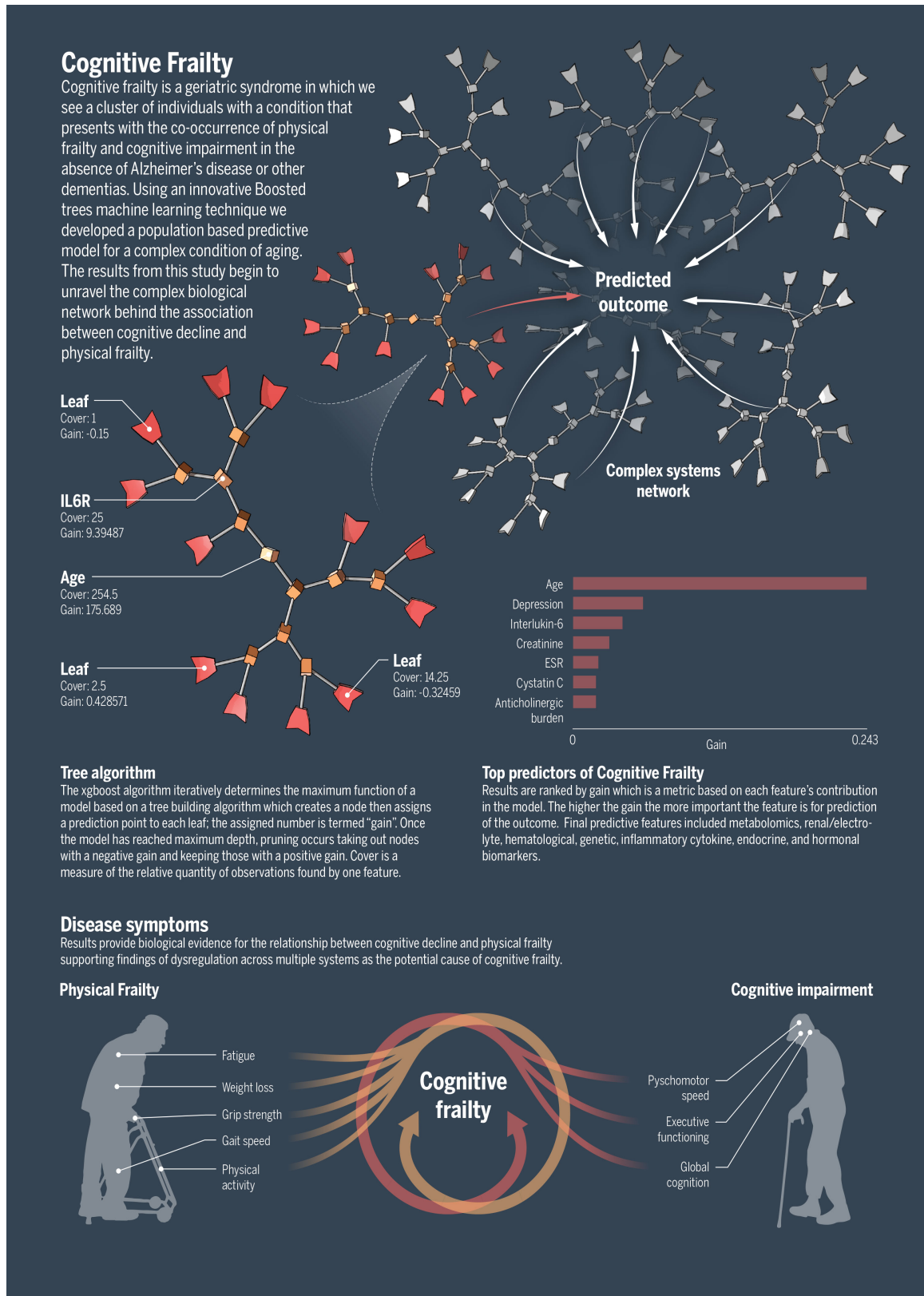
Manuscript 3 includes analyses of genetic biomarkers (SNPs) and highlights the variance seen for individuals with cognitive frailty compared to unique genetic predictors of cognitive decline and frailty alone.

Training Aim3. Acquire the necessary training, expertise, and knowledge to accomplish aims 1 and 2. Goal 1: Apply advanced statistical methods; Goal 2: Develop neuropsychiatric assessment skills.

Due to the innovative statistical modeling and bioinformatics utilized in this dissertation, additional training was needed beyond the standard Doctoral in Philosophy in Nursing Science curriculum to build knowledge and achieve stated aims. I completed the bioinformatics 101 seminar series which included training on: high-throughput technology, high-throughput sequencing data types and public data repositories, DNA and RNA-seq applications and analyses, ChIP-seq applications and analyses, and

pathway and functional enrichment analysis methods. The bioinformatics certificate is included in the supplemental documents. Additionally, I attended conference training on Health Measures, which included training on the NIH neurophysiological, and physical measures toolbox and Patient-Reported Outcomes Measurement Information Systems (PROMIS) measures.

Figure1. Complex systems theory for Cognitive Frailty



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ASSESSING THE CURRENT STATE OF COGNITIVE FRAILITY: MEASUREMENT PROPERTIES

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Abstract: *Background:* Currently, an estimated 25-30% of people ages 85 or older have dementia, with a projected 115 million people worldwide living with dementia by 2050. With this worldwide phenomenon fast approaching, early detection of at-risk older adults and development of interventions focused on preventing loss in quality of life are increasingly important. A new construct defined by the International Consensus Group (I.A.N.A./I.A.G.G) as «cognitive frailty» combines domains of physical frailty with cognitive impairment and provides a framework for research that may provide a means to identify individuals with cognitive impairment caused by nonneurodegenerative conditions. Using the integrative review method of Whittemore and Knafl, 2005 this study examines and appraises the optimal measures for detecting cognitive frailty in clinical populations of older adults. *Methods:* The integrative review was conducted using PubMed, CINAHL, Web of Science, PsycInfo, and ProQuest Dissertations & Theses. From the total 185 articles retrieved, review of titles and key words were conducted. Following the initial review, 168 articles did not meet the inclusion criteria for association of frailty and cognition. Of the 18 fulltext articles reviewed, 11 articles met the inclusion criteria; these articles were reviewed in-depth to determine validity and reliability of the cognitive frailty measures. *Results:* Predictive validity was established by the studies reviewed in four main areas: frailty and type of dementia MCI (OR 7.4, 95% CI 4.2-13.2), vascular dementia (OR 6.7, 95% CI 1.6-27.4) and Alzheimer's dementia (OR 3.2, 95% CI 1.7-6.2), frailty and vascular dementia (VaAD) is further supported by the rate of change in frailty x macroinfarcts ($r = 0.032, p < 0.001$); frailty and the individual domains of cognitive function established with the relationship of neurocognitive speed and change in cognition using regression coefficients; individual components of frailty and individual domains of cognitive function associations included slow gait and executive function ($\beta -0.20, p < 0.008$), attention ($\beta -0.25 p < 0.008$), processing speed ($\beta -0.16, p < 0.008$), word recall ($\beta -0.18, p = 0.02$), and logical memory ($\beta = 0.04, p = 0.04$). Weak grip was predictive for changes in executive function ($\beta -0.16, p = 0.008$). Physical activity was associated with changes in executive function ($\beta = -0.18, p = 0.02$) and word recall ($\beta = 0.17, p = 0.02$), individual components of frailty and global cognitive function were found in several studies which included grip strength ($r = -0.51, p < 0.001$), gait speed ($r = -0.067, p < 0.001$), and exhaustion ($\beta -0.18, p < 0.008$). *Conclusions:* This paper presents the first-known review of the measurement properties for the cognitive frailty construct since the published results from the International Consensus Group (I.A.N.A./I.A.G.G). Evidence presented in this review continues to support the link between physical frailty and cognition with developing validity to support distinct relationships between components of physical frailty and cognitive decline. Results call attention to inconsistencies in reporting of reliability, validity, and heterogeneity in the measurements and operational definition for cognitive frailty. Further research is needed to establish an operational definition and develop psychometrically appropriate clinical measures to construct an understanding of the relationship between physical frailty and cognitive decline.

Key words: Cognitive decline, physical frailty, measurements, cognitive frailty

Introduction

With the number of individuals ages 80 and older on the rise, the burden of dementia is expected to be one of the most daunting and costly consequences of longer life expectancies. Early detection of at-risk older adults and the development of interventions focused on preventing loss in quality of life are increasingly more important. Diagnosing dementia, especially in the early stages of the disease is difficult; many cases go undiagnosed even in the intermediate or more advanced stages (1). This is partly because dementia is a complex condition that cannot be attributed to a single functional or cognitive domain and the need to better understand the underlying neuropathology contributing to non-aging related cognitive

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impairment (2-4).

The relationship between physical frailty and cognitive impairment has become increasingly more apparent with recent studies suggesting that the two are interrelated. Efforts focused on understanding the relationship may provide a means to identify individuals with cognitive impairment caused by non-neurodegenerative conditions which might be reversible (2, 3). Although, frailty and cognitive impairment have been shown to be related, both constructs have long been studied separately (3). To address this gap, the International Consensus Group organized by the International Academy on Nutrition and Aging (I.A.N.A) and the International Association of Gerontology and Geriatrics (I.A.G.G) convened on April 16th, 2013 in an effort to identify domains of physical frailty and

ASSESSING THE CURRENT STATE OF COGNITIVE FRAILITY: MEASUREMENT PROPERTIES

cognition. Additionally, the consensus group recommended formal assessments based on studies that supported findings of an association between progressive physical frailty and cognitive impairment in older adults. The new construct called cognitive frailty (3), extends the physical frailty construct with a formal cognitive assessment and a comprehensive assessment of depressive symptoms.

The construct cognitive frailty, will provide new opportunities for research, assist in further defining cognitive impairment related to physical causes, and promote interventions that lead to improved quality of life in older adults. Multiple studies have been conducted to develop clinical screening tools for the detection of cognitive and functional decline independently, with many clinical screening instruments available to clinicians. However, the optimal measures or combination of measures to accurately detect cognitive frailty in the clinical setting is unclear (3). As researchers attempt to deconstruct the relationship between physical frailty and cognitive impairment, the emphasis must be placed on evaluating the strength of the psychometric tests used to evaluate the new construct. The purpose of this integrative review was to examine the literature to determine progress in the establishment of validity and reliability for the measurement of cognitive frailty.

Operational and Theoretical Definitions

Establishing a comprehensive understanding of the new construct cognitive frailty requires a critical review of what is known about the consensus on operational definitions and tools used to study frailty and cognitive impairment individually.

Frailty

The first definition of frailty was proposed in 1988 (6), but since that time the international community has come to no consensus on a definition of the term or an assessment tool to measure the condition (7). The International (I.A.N.A.) Task Force on Frailty identified 17 cohort-based definitions, all using different frailty assessment tools. More recently, Rodríguez-Mañas et al, 2013 attempted to achieve consensus for an operational definition using a Delphi process, which resulted in consensus on the value of screening for physical frailty in the following six domains: physical performance, including gait speed and mobility, nutritional status, mental health, and cognition. Because there is still a need to identify a specific combination of clinical and laboratory biomarkers for a diagnosis, an operational definition was not recommended (8). Even though consensus has not been reached regarding an operational definition of frailty, the theoretical definition, which is generally agreed upon, describes frailty as a multidimensional geriatric syndrome with increased vulnerability to stressors as a result of reduced capacity of different physiological systems with adverse health outcomes that include falls, disability, hospitalizations, and mortality (7, 9, 10).

The criteria used to identify frailty often depend on the operational definition. The commonly-known criterion is the “phenotypic” definition developed by the work completed in the Cardiovascular Health Study (CHS) (5, 11). The CHS phenotype includes decline in lean body mass, strength, endurance, balance, walking performance, and low activity (5). It allows for a continuous scoring system versus a nominal system because it can capture the multidimensional nature of frailty. The components have concurrent and predictive validity with hazard ratios (HR) ranging from 1.82-4.46 ($p < 0.05$) for outcomes that include incident disease, hospitalization, falls, disability and mortality in community-dwelling older adults (5). Additionally, the CHS model has positive predictive validity (PPV) in detection of physical limitations. The Edmonton Frail Scale (EFS) includes evaluation of the social support domain and has been validated with non-specialists with no formal training in geriatric care (12). Construct validity for the EFS for detection of physical performance was statistically significant ($r = -0.58$, $p = 0.006$, $n=21$) along with inter-rater reliability ($k = 0.77$, $p = 0.0001$) and internal consistency (Cronbach $\alpha = 0.62$) (12). However, the use of the EFS for the detection of cognitive impairment ($r = -0.005$, $p = 0.801$, $n=30$) was not statistically significant (12).

Other validated frailty instruments with unique operational definitions have been described in the literature: the Frailty Index (FI), Clinical Frailty Scale, Study of Osteoporotic Fractures (SOF), SPPB (gait speed, repeated chair stands, and tandem balance tests) validated in the Established Population for Epidemiologic Studies of the Elderly (EPESSE), and Tilburg Frailty Indicator (TFI) which includes three frailty domains (physical, psychological and social) (13–16). Several frailty assessment tools are time consuming, not practical except for research purposes, and have slightly different measurement properties. The literature reflects the lack of consensus and ongoing debate about how to operationalize a definition for frailty (17).

Cognitive Impairment

The theoretical and operational definition for the progressive loss of memory unrelated to the normal aging process has been controversial. Mild cognitive impairment (MCI) was first proposed by Petersen et al, 1999 then revised with the International Working Group on Mild Cognitive Impairment (19). MCI is the most commonly used term to describe a progressive measurable change in memory that differs from healthy aging adults. The recommended criteria for MCI is self and/or informant report of memory impairment and/or evidence of decline over time on objective tasks with preserved activities of daily living, and minimal impairment in complex instrumental functions with no diagnosis of dementia (19). Resulting from the research on MCI the Diagnostic Statistical Manual-5 (DSM-5) included a category of neurocognitive disorder and distinguishes between mild (mNCD) and major (mNCD) neurocognitive disorders to describe the heterogeneity

ASSESSING THE CURRENT STATE OF COGNITIVE FRAILTY: MEASUREMENT PROPERTIES

these articles were reviewed in-depth to determine validity and reliability of the cognitive frailty measures.

Data extraction, was used to identify the psychometric properties based on the measurements provided in the article or if the criteria could be found in the original longitudinal study as referenced by the author. The level of evidence was appraised for each study using the Center for Evidence Based Medicine Levels of Evidence (23). Studies were evaluated with a systematic approach and rated based on their strength of evidence. The operational definitions for both frailty and cognition were reported separately to highlight the combination of tools being used to study the relationship between physical frailty and cognition and report on measurement properties and significant findings. A framework, presented in Table 1, was developed to report the operational definition criteria being used for cognitive frailty based on impairment in the physiological domains defined by The Interventions on Frailty Working Group: mobility, balance, muscle strength, motor processing, nutrition (often operationalized as nutritional status or weight change/sarcopenia), cognition, endurance (including feelings of fatigue and exhaustion), and physical activity (24). Cognition was further defined in the framework based on the use of neuropsychiatric testing and/or a clinical cognitive assessment tool (i.e. MMSE or CDR) in the operational definition. To accompany these results, and to help with replication of the work, the search strategy and data extraction results have been made available online.

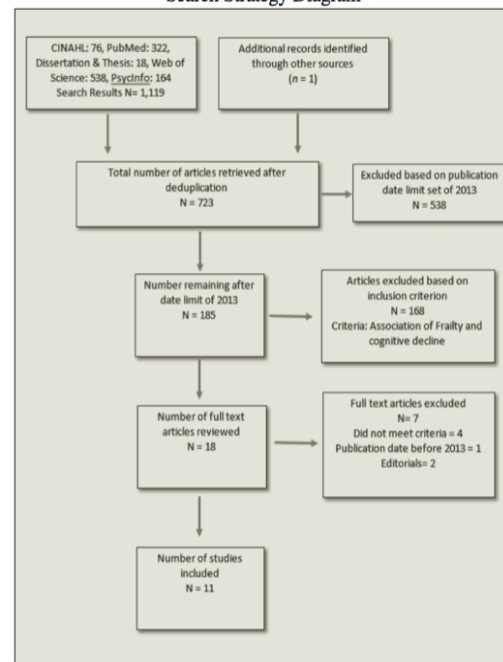
Results

The association between physical frailty and cognitive decline was established in cross-sectional and longitudinal studies before the International Consensus Group (I.A.N.A./I.A.G.G) proposed the definition of cognitive frailty in 2013 (25). Additionally, evidence presented in this review supports the link between physical frailty and cognitive decline with developing validity to support distinct relationships between components of physical frailty and cognition in community-dwelling older adults. Table 2 presents a comparison of the screening tools used by the ten studies included in this review and those proposed by the International Consensus Group (I.A.N.A./I.A.G.G) as a framework for evaluating the development and validation of an operational definition for cognitive frailty.

None of the researchers explicitly described using a theoretical framework; however, all the studies discussed components of cognitive frailty in relation to the International Consensus Group's (I.A.N.A./I.A.G.G) proposed definition. All 11 studies examined the correlation of physical frailty and cognitive impairment. Additionally, six studies examined rate of change in frailty scores in association to rate of deterioration of cognitive scores. Participants were non-demented at baseline in all but two studies, including baseline amnesic Mild Cognitive Impairment (aMCI) and a

probable/possible diagnosis of dementia (26, 27). Although several studies reported baseline cognitive status, scores were not always considered in the statistical model. This finding may be important because baseline cognition can decrease the association between frailty and all dementia outcomes; association between frailty and dementia was stronger with higher baseline scores (HR 1.78, 95% CI 1.14-2.78) than those with lower baseline cognitive scores (HR 0.79, 95% CI 0.50-1.26 p value for interaction = 0.02) (28).

Figure 1
Search Strategy Diagram



Cross-sectional studies

Six cross-sectional studies examined the association of frailty and cognitive decline using a modified CHS criterion (5). Functional status evaluations were added in several studies (26, 29, 30) and co-morbidities, age, gender, BMI, and depression were often considered in the covariate analysis (26, 27, 31). The cross-sectional studies relied on clinical evaluations including MMSE, executive tests, gait speed, grip strength, weight loss, and psychological markers (Table 2). Few of the studies used biomarkers, and only one used imaging in the operational definition (30).

Cohort study

One cohort study examined the associations between frailty and cognitive decline over 12 months (32). The study used

Table 1
Operational Definitions of Cognitive Frailty

Reference	Mobility/ Gait Speed	Strength	Balance	Motor Pro- cessing	Nutrition/ Weight loss	Endurance/ Fatigue	Physical Activity	Neuropsy- chiatric Testing	Clinical Cognitive Assessment Tool [†]
Shimada et al. 2013	X	X			X	X	X	X	X
Kulmala et al. 2014	X	X			X	X	X		X
Buchman et al. 2014	X	X			X	X		X	X
Rolfson et al. 2013*	X	X	X	X	X	X		X	X
Oosterveld et al. 2014	X	X			X	X	X	X	X
McGough et al. 2013	X	X			X		X	X	X
Alencar et al. 2013	X	X			X	X	X	X	X
Gray et al. 2013	X	X			X	X	X	X	X
Solfrizzi et al. 2013	X	X	X	X	X	X	X		X
Robertson et al. 2014	X	X			X	X	X	X	X
Han et al. 2014	X	X			X	X	X		X

*Rolfson et al. (2013) used 3 operational definitions: CHS, Edmonton Frail Scale, and Frailty Index; † Clinical Cognitive Assessment Tool was defined as use of any of the following: MMSE, MoCA, CDR, ADAS-Cog or CASI

the CHS criterion (5) with the addition of a functional status evaluation and tested the MMSE and Clinical Dementia Rating Scale (CDR). The study did not control for chronic diseases or depression. Additionally total sample size (n=182) was small, affecting power for individual classifications of frailty (non-frail n=43, pre-frail n=104, frail n=35) (30).

Longitudinal studies

Results from four longitudinal studies were published after 2013. A modified CHS criterion (5) was used in three of the studies. One study used more than one frailty instrument to determine if the relationship between neurocognitive speed (NCS) and frailty was affected by how frailty was operationalized (33). The use of biomarkers, clinical markers, and imaging varied among studies. The use of biomarkers and imaging was more commonly used in the longitudinal studies than cohort and cross-sectional studies (Table 2). Functional status evaluation was added in one study (34) and co-morbidities were considered in the analysis for all of the studies.

Validity

For all the studies in this review, criterion validity was examined for performance of the operationalization of various cognitive frailty measurements. Predictive and discriminant validity was commonly reported as odds ratio (OR) or hazard ratio (HR); two studies used Pearson correlations and multiple linear regression models to establish associations between components of physical frailty and cognitive function. Predictive validity was established by investigating frailty and rate of change in cognition or correlation of frailty and cognitive decline. Discriminant validity was established by analyzing the relationship between measures of frailty (frail, pre-frail, and robust) and type of dementia (MCI, clinically diagnosed dementia, vascular dementia, and Alzheimer's) (26, 28, 30, 32). All of the studies evaluated community-dwelling older adults for which the CHS frailty measures are validated (5). Only one study compared more than one operational definition of frailty: CHS, FI, and EFS (33). Heterogeneity was present in the objective measures, and the terminology-specific language for the components of the CHS frailty construct often varied from the validated CHS criteria (5).

ASSESSING THE CURRENT STATE OF COGNITIVE FRAILTY: MEASUREMENT PROPERTIES

Heterogeneity was present in the objective measures for cognitive assessment and neuropsychiatric testing. Two studies assessed global cognition with the MMSE (30, 34), four used the MMSE and domain specific neuropsychiatric testing (26, 29, 32, 33), three used only domain neuropsychiatric testing (27, 28, 35), and one assessed global cognition with both the MMSE and MoCA with domain specific neuropsychiatric testing (36). The Cognitive Dementia Rating scale (CDR) had no predictive validity with evidence of no difference between frailty and cognition (relative risk = 2.1; $p = 0.393$) (32). The National Center for Geriatrics and Gerontology-Functional Assessment tool (NCGG-FAT) had good test-retest reliability with moderate to high external validity (Person $r = 0.496$ to 0.842). The MMSE continues to be the most commonly used clinical cognitive assessment tool for operationalizing cognitive frailty (25); concurrent validity (Pearson $r = 0.776$; $p < 0.001$) and reliability test-retest (Person $r = 0.827$; $p = 0.001$) (37) with neuropsychiatric testing predictive and discriminate validity is established by the rate of change in MMSE and CHS frailty criterion (32).

Predictive validity was established in four main areas: 1) frailty and type of dementia: MCI (OR 2.0; $p = <0.001$) and (OR 7.4, 95% CI 4.2-13.2) (29, 30); vascular dementia (OR 6.7, 95% CI 1.6-27.4) and (HR 2.68, 95% CI 1.16-7.17) (30, 34); and Alzheimer's dementia (OR 3.2, 95% CI 1.7-6.2), (HR 1.08, 95% CI 0.74-1.57), and (HR 0.62, 95% CI 0.20-1.89) (28, 30, 34). The relationship between frailty and vascular dementia (VaAD) is further supported by the rate of change in frailty x macroinfarcts ($r = 0.032$, $p < 0.001$) (35). Evidence of convergent validity exists between dementia and non-dementia types with findings to support the associations between frailty and non-Alzheimer's dementia (OR 2.57, 95% CI 1.08-6.11).

2) Frailty and the individual domains of cognitive function was identified by evaluating the relationship of neurocognitive speed and change in cognition using regression coefficients (33) and evaluation of the MMSE subdomains. Individual domains of cognitive function were found to be gender specific (31). Predictive validity was dependent on the frailty operational definition; Frailty Index (FI) and NCS (OR 0.87, 95% CI 0.81-0.95) compared to the modified CHS and EFS which found no correlation with neurocognitive speed (33).

3) Individual components of frailty and individual domains of cognitive function associations included slow gait and executive function ($\beta -0.20$), attention ($\beta -0.25$), processing speed ($\beta -0.16$) (36), word recall ($\beta -0.18$, $p = 0.02$), and logical memory ($\beta = 0.04$, $p = 0.04$) (27). Weak grip was predictive for changes in executive function ($\beta -0.16$, $p = 0.008$) (27). Physical activity was associated with changes in executive function ($\beta = -0.18$, $p = 0.02$) and word recall ($\beta = 0.17$, $p = 0.02$) (27).

4) Individual components of frailty and global cognitive function were found in several studies (27, 28, 34-36). Individual components included grip strength ($r = -0.51$, $p < 0.001$), gait speed ($r = -0.067$, $p < 0.001$) (35), and exhaustion

($\beta -0.18$) (36) were predictive for changes in global cognition.

Psychological markers were frequently used for the assessment of endurance, fatigue, or depression. However, variability existed in the type of assessment scale used and how the psychological marker was operationalized. Psychological markers were typically used to either assess endurance for fatigue in the CHS criteria (29, 35) or considered as a covariate in the statistical analysis (27, 28, 32, 34). Variability in the psychological markers can be seen in Table 2 and online material.

Reliability

Due to the heterogeneity in the objective measures for frailty, reliability was not consistently examined for cognitive frailty. The limited reliability and variability in the operational measurements used for the CHS frailty criteria add challenges to establishing an operational definition for cognitive frailty. Motor performance was the only measurement for which validity and reliability was established (34).

Feasibility

Instrumental assessments for cognitive frailty are currently time-consuming, expensive, require extensive training, and the clinical translation properties are not clear. The addition of biomarkers and imaging potentiates the complexity of the feasibility for measures and complicates the process for detection of cognitive frailty in the clinical setting.

Discussion

The findings from this review continue to support evidence for the association between physical frailty and cognitive decline. However, while cross-sectional studies have detected a relationship, further studies are needed to determine causal pathways (38). Studies continue to use different combinations of measurement instruments for cognitive frailty, but are measuring similar domains of physical frailty and cognition. Based on the findings in this review the CHF criteria with measures of mobility/gait speed, strength, nutrition/weight loss, endurance/fatigue, and physical activity, neuropsychiatric testing and a cognitive assessment tool was the most common operational definition (Table 1). Further testing of the cognitive frailty construct should attempt to provide validity and reliability for objective measures and scales which are based on self-report. Self-report scales must prove to be stable over time (test-retest reliability), and those administered by several individuals need to exhibit good inter-rater reliability. Additionally, inclusion of a theoretical framework will provide a structure for generating cumulative knowledge on which interventions can be based.

Studies are starting to deconstruct the relationship between the components of physical frailty and cognitive decline. Unravelling of the complex cognitive frailty construct will refine the operational definition and improve an

Table 2
Use of biological, clinical, and imaging markers for cognitive frailty: International Consensus Group (I.A.N./I.A.G.G.)

	Shimada et al. 2013	Kubota et al. 2014	Buchanan et al. 2014	Roffson et al. 2013	Oosterveld et al. 2014	McCough et al. 2013	Alencar et al. 2013	Gray et al. 2013	Sofrizzi et al. 2013	Robertson et al. 2014	Han et al. 2014
Biomarkers											
Inflammatory markers (e.g. CRP, IL-6)			X								
Beta-amyloid protein (Aβ)		X									
aPOEε4 genotype			X				X				
Anemia											
Serum albumin									X		
Cholesterol									Xβ		
Vitamin D status											
Clinical markers											
MMSE	X	X		X	X	X	X		X	X	X
Executive tests	X		X		X	X		X		X	
ADAS-Cog					X	X					
CDR					X	X				X	
MoCA											
Gait speed	X	X	X	X	X	X	X	X	X	X	X
Hand grip strength	X	X	X	X	X	X	X	X	X	X	X
Weight loss	X	X	X	X	X	X	X	X	X	X	X
Psychological marker: GDS	X®		X£	X€		X§	Xφ	XΩ	X*		X§
Actigraphy											
Imaging											
Dual energy											
X-ray absorptiometry scans (DEXA)											
Cerebral Computed tomography		X						X	X		
Cerebral Magnetic resonance imaging		X						X	X		
Functional MRI											
Diffusion tensor imaging (DTI)											
Tractography											
Electrophysiological methods											
Cognitive evoked potentials											

Y: CT scan; MRI and laboratory tests (not specified) were used to make a diagnosis of vascular dementia, Alzheimer's disease, Lewy bodies, and dementia related to other medical causes; ®: Partial GDS scale; £: Psychological marker evaluated with two questions from the Center for Epidemiologic Studies; €: Psychological marker evaluated with the Edmonston Frail Scale; §: GDS-15 scale; φ: GDS-15 scale and Cornell Depression Scale; Ω: Psychological marker evaluated with Center for Epidemiologic Studies Depression; * GDS 30 scale; β: Reported in original study.

ASSESSING THE CURRENT STATE OF COGNITIVE FRAILITY: MEASUREMENT PROPERTIES

understanding of the clinical distinction between cognitive impairment due to physical frailty and an isolated neurological condition. Disentangling the association between frailty and cognitive decline requires the use of convergent validity to determine if the cognitive frailty construct is able to distinguish among between different types of dementia (e.g., Vascular, Alzheimer's, Lewy Body, and Parkinson's dementia) (27). The association of cognitive decline and frailty may be responsible for part of the heterogeneity in the presentation of dementia. Movement toward evaluating specific domains of cognitive impairment such as executive functioning and psychomotor speed versus a global assessment of dementia will facilitate an understanding of the implications for cognitive frailty. However, the current lack of validity and reliability of a cognitive frailty operational definition means that it is not possible to recommend translation of measures to detect the presence of risk factors that may predict cognitive frailty in the clinical settings.

A limitation of this review was the exclusion of studies that did not address the cognitive frailty construct. In the future, a review of the literature focused on individual physical function measures may identify other markers associated with cognitive impairment. Further research with epidemiological and population based studies that includes diverse ethnic and social economic groups will help establish a better understanding of the prevalence of cognitive frailty. The majority of studies in this review either did not report ethnicity or the sample included a high proportion of white (88%-99%) females (58%-80%). Only two studies provided a population-based estimate of cognitive frailty with samples of 5,104 Japanese (29) and 4,649 Irish community-dwelling older adults (36). Understanding how demographics effect the measurement of cognitive frailty are important since psychometric tools may be effected by populations which have higher rates frailty, comorbidity, cardiovascular disease, poorer health, decreased access to care, and low education and income (5). Inclusion of chronic diseases, such as depression and cardiovascular disease, as a part of the study design is an important part of describing other factors that may contribute to cognitive frailty over time. Additionally, adjustment for the presence of apolipoprotein (APOE) $\epsilon 4$ alleles and other biomarkers (e.g. inflammatory makers, beta-amyloid protein) could help describe the pathophysiological mechanisms.

The early detection of cognitive decline emphasizes a promising focus for the development of preventive and therapeutic interventions. Current studies suggest the importance in understanding both constructs separately as a way to deconstruct dissociable components, describe common pathologies, and develop a single operational definition which would allow for targeted interventions. Ensuring validity and reliability in the measures used is paramount if providers are to identify individuals at risk for pathological non-normal aging changes and develop interventions to improve the quality of life of older adults. Further research is needed to establish an operational definition for cognitive frailty, develop a better

understanding of the directional relationship between physical frailty and cognitive impairment, gender differences, and identify biomarkers to assist with detection of diagnosis and disease progression.

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Conflict of interest: The authors have no conflict of interests to report

Ethical Standards: To reduce bias in this rigorous review the authors adhered to the Whittemkore & Knafli, 2005 and PRISMA guidelines. This study did not use human subjects.

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MANUSCRIPT 2:

Determining Biological Factors for Cognitive Frailty: A Systematic Review

Abstract: On April 16th, 2013 the International Consensus Group (I.A.N.A/I.A.G.G) formally defined the novel phenotype cognitive frailty; a condition characterized by the co-occurrence of physical frailty and cognitive impairment. We hypothesize that there are biological factors to describe the interconnection between physical frailty and cognitive impairment. This systematic review focuses on identifying the shared measurable biological and genomic mechanisms for physical frailty and cognitive decline. Two independent reviewers assessed the eligibility of each report based on predefined inclusion criteria to ensure interrater reliability; a third reviewer resolved conflicting assessments. The review was conducted using PubMed, Embase, Scopus, Web of Science, LILACS, Gene Indexer, and GWAS Central. Findings resulted in 1232 abstracts for full review, 327 articles were included in the final review. Data extraction identified a correlation between 16 distinct inflammatory and protein markers with biomarker-related gene expression for cognitive frailty. Meaningful findings were identified in the relationship between protein and genetic markers found for both cognitive decline and physical frailty. This systematic review presents the first known findings of the underlying biological characteristics for cognitive frailty providing evidence for converging pathophysiological pathways.

Introduction

In the past century, scientific research has been driven by molecular science with the common goal of identifying a single group of biological or genetic mechanisms as the

cause of disease. We now understand that the mechanisms underlying disease processes are multi-factorial and system based. A multi-system physiological disease requires a systems approach to precision research especially with older adults who have variable trajectories to the aging process with multiple co-morbidities. Efforts to unravel this complexity start with understanding the unique biological factors for a cluster of individuals presenting with similar symptoms and trajectories. Cognitive frailty can be considered a unique geriatric phenomenon in which we see a cluster of individuals with a condition which simultaneously presents with both physical frailty and cognitive impairment¹. The International Consensus Group organized by the International Academy on Nutrition and Aging (I.A.N.A) and the International Association of Gerontology and Geriatrics (I.A.G.G) convened in 2013 to identify related domains of physical frailty and cognition. The new construct called “cognitive frailty” is defined by the presence of physical frailty and cognitive impairment in the absence of Alzheimer’s disease or other dementias¹. The mechanisms and the directional relationship behind the dynamic association of these two constructs remains unexplained. There exists strong evidence for the association of frailty and cognitive decline with suggestion for pathophysiological mechanisms which are shared by both clinical manifestations². Although, some research has been conducted on the association between physical function and cognitive decline there is still no comprehensive list or understanding of the underlying mechanisms for cognitive frailty. Therefore, to further develop an understanding of cognitive frailty, it is critical that the operational definition explore both clinical and biological markers for cognitive decline and physical frailty.

Identification of a measurable cellular, biochemical, or molecular markers for cognitive frailty has not been identified. Because both cognitive decline and physical frailty are large heterogeneous conditions it may not be possible to identify one biomarker to measure both cognitive decline and frailty. The use of one or more biomarkers specific to both constructs will improve our understanding of the association^{3,4}. It is possible that the underlying biological mechanisms for cognitive frailty are at the intersect between cognitive decline and physical frailty or cognitive frailty may contain some of its own unique markers of disease.

Some evidence exists to support inflammatory biomarkers (neuroinflammatory cytokines) such as C-reactive protein (CRP) and Interleukin-6 (IL-6) as antecedent biomarkers since they are associated with frailty and cognitive decline^{1,3}. The complicated use of inflammatory biomarkers, such as CRP, for detection of disease is that they can be detected in other co-morbid diseases found in older adults (i.e. cardiovascular disease, rheumatologic disease). Wilson, Finch, and Cohen (2002) completed a review exploring over 30 neuroinflammatory cytokines and their findings indicate the potential for detection of cognitive decline and evidence for associated improvement of cognition with targeted interventions to reduce the production of specific neuroinflammatory cytokine markers⁵. Finally, genetic factors associated with cognitive frailty have not been fully explored. There have been several genome-wide association studies (GWAS) and candidate gene studies for cognitive decline with only more recent studies exploring the genetic mechanisms for frailty.

Methods

Search strategy

In this review, we followed the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines⁶. A systematic review of the literature was performed using the following online databases: PubMed, Embase, Scopus, Web of Science, LILACS, Gene Indexer, and GWAS Central. For reproducibility, we have provided the PubMed search strategy in the supplementary appendix (Figure I). Databases were searched from the start date of the database to 22 December, 2015. An update of the searches was performed prior to the data extraction phase on 26 May, 2016 to identify any new publications. In addition to database searching, articles were hand-pulled from references and identified through other sources.

Inclusion and exclusion criteria

Studies that included information on biomarkers or genetic markers for dementia, physical frailty, or cognitive frailty were included. Reviews, animal studies, imaging biomarkers, and case studies were excluded. Studies on a geriatric population, aged 65 and older, were included. Articles about other disease states such as cancer, Multiple Sclerosis, Down syndrome, Parkinson's disease, human immunodeficiency virus (HIV), and Huntington's disease were excluded. Articles published in English were included.

Study appraisals

A multi-step approach was used to evaluate relevant articles using Covidence, a web-based software platform selected by Cochrane Reviews that organizes and streamlines the systematic review process⁷. Figure I shows the stages (PRISMA) for retrieving the

studies for inclusion and extraction. We conducted a review of the titles and abstracts of all the papers identified through database searching and hand pulling from references lists. Three reviewers participated in this step and each article was reviewed by two reviewers (LS and AS) to ensure interrater reliability. A third reviewer (SH) resolved conflicting assessments. A fourth reviewer (EA) was available for additional arbitration however their services were not required. From 5942 articles identified, titles and/or abstracts reporting on information pertaining to biomarkers or genetic markers for cognitive decline, physical frailty, or cognitive frailty was included. 1232 potential relevant articles were chosen for closer review, two reviewers with appropriate subject expertise (LS and AS) assessed the full-text of the articles for relevancy. 327 full-text articles reporting on the relevant topic met inclusion/exclusion criteria and 899 articles were excluded. Reviewer disagreements were addressed in regular meetings and resolved. A final 327 articles were included in this systematic review.

Extraction

The analysis for this paper was generated using Qualtrics software, Version 9.2017 of Qualtrics (Copyright © [2017] Qualtrics. Qualtrics and all other Qualtrics product or service names are registered trademarks or trademarks of Qualtrics, Provo, UT, USA. <http://www.qualtrics.com>.) The survey created in Qualtrics (Qualtrics, Provo, UT) ensured consistency in reporting of biological markers limiting open text boxes, consistent categorizing of biomarkers by clinical, genetic, and fluid markers in the following categories: inflammatory/immunity, protein, metabolomics, oxidative stress. The database assigned each biomarker unique numeric code (i.e. IL6-3, CRP-27). When

data entry was complete, the final data frame was exported from Qualtrix and an analysis was carried out using R V. 3.2.1. R is free, open-source software that provides many statistical and graphic techniques. R packages used included 'MASS' and 'ggplot2'^{8,9}.

We did not complete a formal method of assessment for the quality of the studies with a meta-analysis given that the goal of this review is to identify potential putative markers for a new phenotype "cognitive frailty". Level of evidence was appraised for longitudinal, observational (cohort, cross Sectional, case-control studies), and randomized clinical trials (RCTs) using the Center for Evidence Based Medicine Levels of Evidence¹⁰. Additionally, there are limited (RCTs) for frailty and none for cognitive frailty. We do provide a compressive list of the principle results, study design, and detail list of genetic findings correlated to one of the following phenotypes: cognitive decline, frailty, and cognitive frailty. The markers extracted for correlation to cognitive frailty were identified by the reviews to be studies that explored both frailty and cognitive decline in the same study.

Findings and discussion

A total of 327 articles were used to extract the clinical, genetic, and protein markers for three phenotypes: cognitive decline, physical frailty, and cognitive frailty. Date ranges for the studies are shown in Figure II. Studies were reviewed in the following categories 39 genetic studies: 9 GWAS and 30 candidate gene studies, 279 biological protein studies, 9 medication risk studies. Additional study designs included observational (Cohort, cross sectional, and case-control studies), longitudinal, RCT and In Vitro studies.

For the 13 studies that included both a longitudinal and observational (Cohort, cross sectional, and case-control studies) study design we extracted markers from both study designs. The studies were categorized by phenotype: cognitive decline (n= 243), frailty (n= 72), and cognitive frailty (n= 11). Phenotypes were further defined by the type of cognitive decline (i.e. Alzheimer's disease, mild cognitive impairment) and component of frailty (i.e. gait, sarcopenia, grip strength, physical activity) as stated in the study or a combination both was considered cognitive frailty. The supplementary appendix (table I) shows the clinical and biomarkers extracted from 288 articles. Tables I-III show the biomarkers extracted by phenotype in the following categories: clinical, inflammatory/immunity, laboratory, protein, metabolomics, and oxidative stress. Additionally, a summation or frequency in which the biomarker occurred out of the 288 articles is shown by phenotype.

Clinical markers

Although, clinical markers were not a part of the search strategy several of the studies reported clinical findings associated with cognitive decline, physical frailty, and cognitive frailty. Demographics such as increasing age were a factor for all phenotypes, lower education and income were factors for individuals with cognitive decline and frailty. Other clinical markers included: measures of cardiovascular disease, elevated blood pressure, multiple co-morbidities, changes in body mass index (BMI), and alcohol intake. One of the most interesting clinical findings was an association between medications and all phenotypes. These included hypertension, benzodiazepine, anticholinergic, and psychoactive medications. Two categories of hypertensive medications beta-blockers

(i.e. metoprolol and atenolol) and angiotensin-converting enzyme (ACE) inhibitors were found to have the most significant effect on cognitive decline^{11,12}. Additionally, there was a significant interaction between ACE inhibitor use and carriers of *ApoE4* (odds ratio: 20.9, 95% CI 3.08-140.95, $p = .002$)¹². Anticholinergic burden was found to be associated with cognitive decline and physical frailty. An interaction was found between *ApoE4* carriers and anticholinergic medications with users having the lowest cognitive scores. Irrespective of *ApoE4* status, drugs with high anticholinergic properties were associated with cognitive and physical decline^{11,13-16}. Methods for measuring medication burden varied significantly between studies making it difficult compare study results.

Inflammatory/Immunity markers

There were 16 neuroinflammatory cytokine markers associated with cognitive decline and frailty. These included: elevated levels of IL6, CRP, tumor necrosis factor (TNF-alpha), uric acid, IL1-beta, erythrocyte sedimentation rate (ESR), cortisol/dehydroepiandrosterone ratio, IL1RA, CD8, IL6R, TNF-a receptor I (TNFR1), cortisol, homocysteine, fibrinogen, and beta 2-microglobulin (B2M). Additionally, all the neuroinflammatory markers associated with cognitive frailty were associated with either cognitive decline or frailty. These neuroinflammatory cytokines were found to be associated with cognitive decline and frailty in cross-sectional and longitudinal studies suggesting that these markers could be both early and persistent markers. The presence of the hypothalamic-pituitary-adrenal (HPA) axis hormones such as dehydroepiandrosterone can interact with inflammatory markers to influence disease.

This relationship should be explored further with clinical markers such as gender and body mass index.

Laboratory markers

Twenty laboratory markers are associated with both phenotypes and include:

Nutritional markers: low levels of vitamin D, total albumin, and selenium;

Cardiovascular/endocrine markers: elevated total cholesterol, triglycerides, LDL, insulin like growth factor protein (IGF-1), glucose, insulin resistance, HbA1c; Hematology/renal markers: elevated creatinine, creatinine clearance, blood urea nitrogen (BUN), white blood cells (WBC); and decreased hemoglobin, hematocrit, cobalamin deficiency (B12), and increased methylmalonic acid (MMA), and hormonal marker: low levels of total testosterone associated with decreased lean muscle mass and cognitive decline. These markers combined with endocrine and immune markers suggest changes to the cellular immune system and HPA axis that are related to cognitive and physical decline.

Additionally, several studies included these markers and the inflammatory/immune markers as a composite score and found an increased risk for developing cognitive decline, frailty, and mortality¹⁷⁻²².

Protein markers

Several of the protein markers were measured by cerebrospinal fluid (CSF) and included known biomarkers associated with the neurofibrillary tangles involved in the pathogenesis of neurodegenerative diseases such as Alzheimer's disease and frontotemporal dementia²³. None of these markers (i.e. p-tau, A β -42) have been studied in frailty. Three markers measured by serum/plasma were associated with both

cognitive decline and frailty, these included: sirtuin 1 and cystatin C. The down regulation of Sirtuin 1 has been reported to be involved in the pathway that controls the expression of A β peptide through *ADAM10*²⁴. Concentrations of sirtuin 1 decline with age but the decline was found to be more significant in individuals with cognitive decline and frailty compared to age matched healthy individuals^{24,25}. Additionally, cystatin C has been thought to bind to soluble A β preventing accumulation in the brain²⁶. Decreased serum cystatin C has been associated with higher risk for cognitive decline and gait speed decline^{27,28}.

Metabolomics and oxidative stress markers

No metabolomics markers were found to be related to cognitive frailty. Two oxidative stress markers were associated, these included: malondialdehyde (MDA) and protein carbonyls. MDA and protein carbonyls are well established oxidative biomarkers and are considered to be a good measure of systemic oxidative stress²⁹. Both are associated with frailty and cognitive decline but not predictive of the development or progression of disease^{29,30}.

Genetic

The supplementary appendix table II shows a complete list of genetic markers identified by phenotype. Three genes were found to be associated with cognitive decline and frailty in candidate gene studies: *IL6* rs1800796, *TNF* rs1800629, and *COMT* with different SNPs, rs4680 for cognitive decline and rs4646316 for frailty. *IL6* and *TNF* have corresponding serum markers that are associated with both phenotypes (see inflammatory/immunity markers)³¹⁻³⁴.

There are 12 serum biomarker and gene correlations, these are shown in table IV.

Further evaluation is need to determine if there is a direct correlation between gene expression and serum marker function.

Conclusions

It has previously been postulated that a dysregulation across multiple systems may be the potential cause for both cognitive and physical decline^{18,19,21}. The results from this systematic review provide evidence for a biological association between cognitive decline and physical frailty. The potential in identifying a unique biomarker that is the key to a specific molecular or cellular event is enticing but considering the complexity and individual variability to aging we need to consider the possibility that these interactions are non-linear. Several studies presented here have taken various approaches to combining biomarkers using method such as allostatic load index, physiologic dysfunction scores, principle components analysis (PCA), and serum protein based algorithms (random forest methods) to yield a more accurate understanding in the relationship between biomarkers and detection of disease^{18,19,21,22}. Future research should focus approaches that could include multiple markers of disease to build an accurate model for the detection of cognitive frailty. Finding should be reproducible and validated before translating into clinical practice. Integrating multiple biomarkers has potential to help us better understand the complex physiological interactions. Such validated models for disease detection will be invaluable in the prevention and early detection of diseases unique to aging.

Figure I. PRISMA flow diagram of study selection and citation analysis⁶

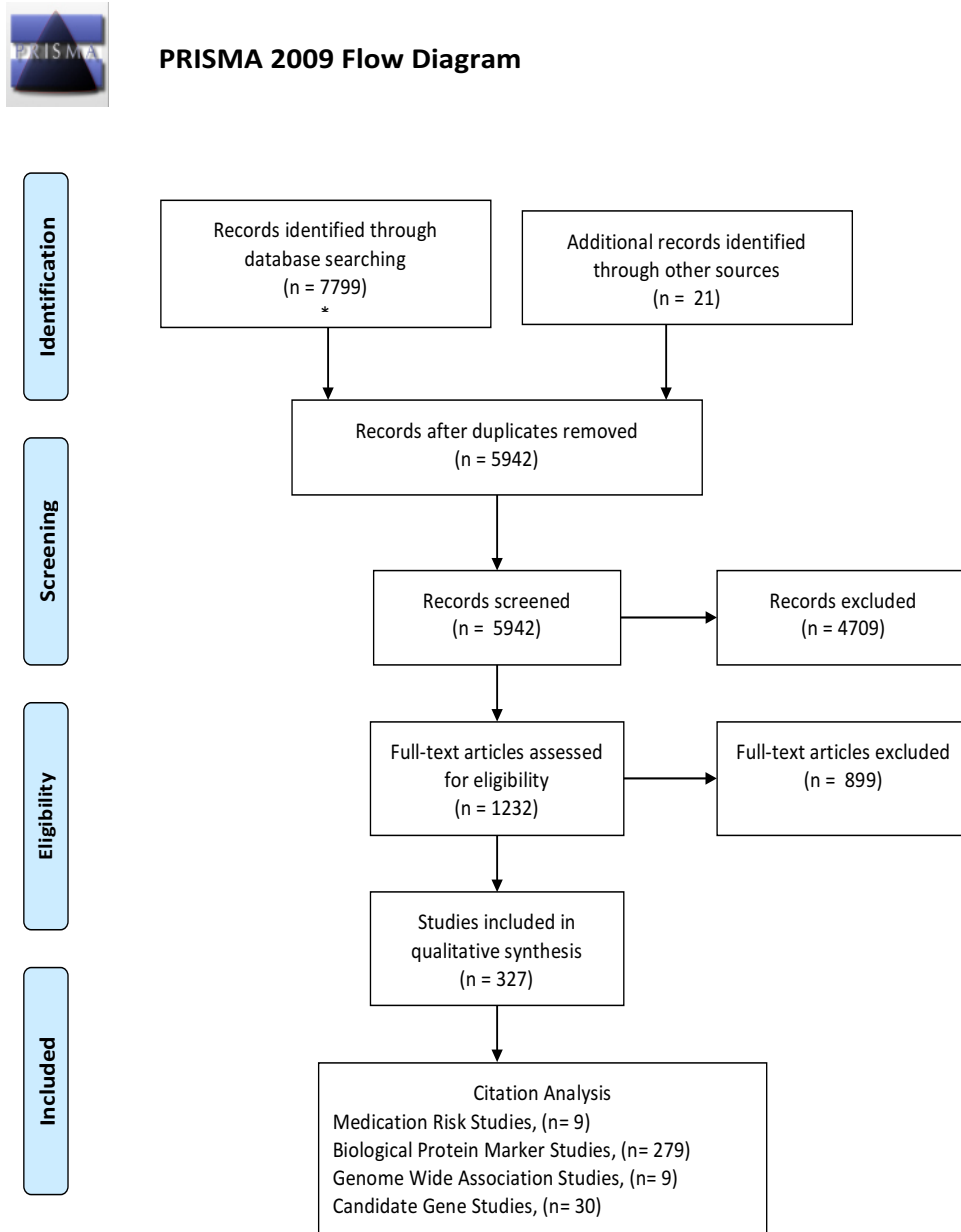


Figure II. Systematic review publication date range

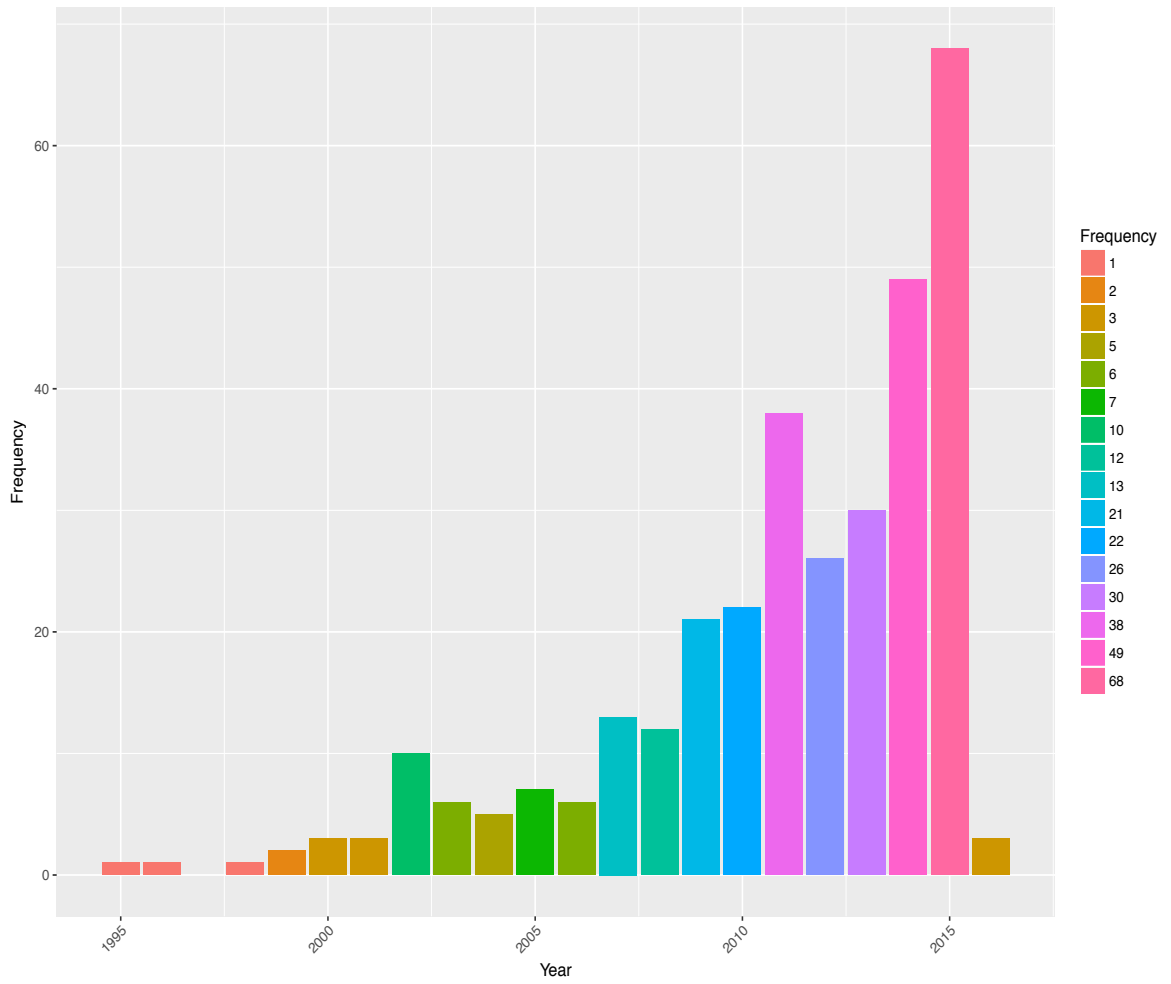


Table I. Cognitive decline biomarkers by category and frequency

1.Frequency	1.Inflammatory/Immunity Markers	2.Frequency	2.Laboratory Markers	3.Frequency	3.Protein Markers	4.Frequency	4.Metabolomic Markers	5.Frequency	5.Oxidative Stress Markers	6.Frequency	6.Clinical Markers
19	C-reactive protein	9	Composite Score (multiple markers)	37	Aβ 1-40/t-tau ratio	2	Docosahexaenoic acid (DHA)	6	F2-isoprostanes/isoprostanes	6	Elevated blood pressure
17	IL6	8	Albumin	32	Aβ42	1	Sphingolipid-SM(d18:1/18:0)	2	Choline plasmalogen(PlsCho)	5	Anticholinergic medications
9	Homocysteine	7	Olfactory marker	26	Aβ 1-42	1	sphingomyelin [SM(39:1)]	2	Glutathione peroxidase (GSH-Px)	5	Change in Body Mass Index
8	Tumor necrosis factor (TNF-alpha)	5	Ocular marker	19	PhosphoTau181 (P-tau181)	1	SM/ceramide ratio	2	Oxidative stress markers /Total antioxidant stat	2	Alcohol intake
5	YKL-40 (neuroinflammation or Chitinase-3 Chi3L3)	4	Folate	13	P-tau	1	SM/ceramide ratio	1	Peroxisomal b-oxidation levels	2	Elevated systolic pressure
5	Cortisol	3	Creatinine	8	Aβ1-42/ Aβ1-40 ratio	1	PC aa 36:1 Glycerophospholipids	1	Ethanolamin plasmalogen (PlsEtn)	2	Elevated diastolic pressure
4	IL8	3	Cobalamin deficiency (B12)	7	Aβ 1-40	1	PC aa 32:0 Glycerophospholipids	1	PlsCho + PlsEtn	2	Increase Waist Circ/Waist-to-hip
3	TNF-α receptor I (TNFR1)	3	Platelet distribution width (PDW)	7	P-tau231	1	PC 16:0/20:5 phosphatidylcholine	1	PlsCho/PlsEtn Ratio	1	Change in resting heart rate
3	Fibrinogen	2	Nutrient biomarker patterns (NBP)	5	Aβ40	1	PC 16:0/22:6 phosphatidylcholine	1	Plasmalogen	1	Cardiovascular disease
3	Uric Acid	2	Blood urea nitrogen (BUN)	5	P-tau181/Aβ42	1	PC 18:0/22:6 phosphatidylcholine	1	Protein carbonyls	1	Benzodiazepine medications
3	Monocyte chemotactic Protein-2 (MCP-2)	2	Methylmalonic acid (MMA)	5	Cystatin C	1	Ceramides C16:0	1	Malondialdehyde (MDA)	1	Hypertensive medications
3	Resistin	2	Glucose	4	t-tau/ Aβ42	1	Ceramides C20:0	1		1	Angiotension converting enzyme medications
2	IL10	2	Insulin resistance (IR-HOMA)	3	Aβ 1-42/t-tau ratio	1	Ceramides C22:0	2		2	Psychoactive medications
2	IL1beta	2	Lipids: Triglycerides	3	Apolipoprotein A-1 (ApoA1)	1	Ceramides C24:0	1		1	Low level of education
2	IL17E	2	Lipids: LDL cholesterol	3	Brain derived neurotrophic factor (BDNF)	1	Ceramides C26:0	1		1	Instrumental activities of daily living (IADL)
2	Clusterin	2	Lipids: HDL cholesterol	3	Lysosomal-associated membrane protein 1 (LAM)	1	Stearoyl	1		1	Activities of daily living (ADLs)
2	TNF-α receptor II (TNFR2)	2	Free Testosterone	2	sAβ/APP ratio	1	Eicosapentaenoic acid (EPA)				
2	Macrophage Migration Inhibitory Factor (MIF)	2	Insulin like growth factor protein (IGF-1)	2	Aβ42/ Aβ40						
2	Cortisol/Dehydroepiandrosterone ratio	2	Insulin like growth factor protein (IGF-2)	2	Neurofilament light chain (NFL)						
2	Vascular cell adhesion molecule 1 (VCAM1)	2	Insulin like growth factor protein Binding Protein (2	Apolipoprotein A-1 (ApoA2)						
2	Soluble receptor for advanced glycation end products (2	Insulin like growth factor protein Binding Protein (2	Complement factor H (CFH) protein 1						
2	Plasma Pentraxin 3 (PTX3)	2	Anemia	2	Chromogranin A (CgA)						
2	alpha 2-macroglobulin (A2M)	2	Hemoglobin	2	Visinin-like protein-1 (VLIP-1)						
2	Adiponectin	2	Polyunsaturated fatty acids (O3PUFAs)/ n-6/n-3 ra	2	β-secretase (BACE-1)						
1	IL1	2	alpha-1-antichymotrypsin (ACT)	2	Ubiquitin						
1	IL6R	2	Vascular endothelial growth factor (VEGF)	2	Heat shock protein 70						
1	IL13	1	Peroxidase	2	Epidermal growth factor (EGF)						
1	IL1RA	1	Creatinine Clearance	2	Pancreatic peptide (PP)						
1	IL7	1	N-acetylaspartate (NAA)/creatinine (Cr)	1	Soluble amyloid β protein (sAβ)						
1	IL12p70	1	Methylcitric acid (MCA)	1	Amyloid β precursor protein (APP)						
1	D-dimer	1	Holotranscobalamin (holoTC)	1	Aβ 1-42/p-tau ratio						
1	Procalcitonin	1	Glycohemoglobin (HbA1c)	1	P-tau231/Aβ42/40 ratio						
1	Erythrocyte sedimentation rate (ESR)	1	Lipids: Total Cholesterol	1	T-tau/Aβ42/40 ratio						
1	GlycA	1	24S-hydroxycholesterol	1	Apolipoprotein C2						
1	Macrophage inflammatory protein 1-alpha (MIP 1α)	1	Aspartate transaminase (AST)	1	Apolipoprotein H						
1	Plasminogen activator inhibitor (PAI-1)	1	Gamma glutamyl transferase (GGT)	1	ApoB/ApoA1 ratio						
1	Serum Amyloid A	1	Total Testosterone	1	A1AcidG						
1	Fibrinogen gamma-chain	1	Total Bilirubin	1	Transthyretin (TTR)						
1	Neural cell adhesion molecule (NCAM)	1	Vitamin E	1	Ceruloplasmin						
1	Adhesion molecule soluble intercellular adhesion mole	1	Vitamin D	1	Cathepsin D						
1	Soluble receptor for advanced glycation end products (1	Vitamin C	1	Glycogen synthase kinase-3 (GSK3-α)						
1	Neutrophil/Lymphocyte ratio	1	Beta-Carotene	1	Neuronal Cell Adhesion Molecule (NrcAM)						
1	Monocyte chemotactic protein-1 (MCP-1)	1	Calcium	1	Axl receptor tyrosine kinase (AXL)						
1	CD40	1	Nitrate+Nitrate3	1	VLIP-1/Abeta1-42						
1	IgG2	1	Selenium	1	Sirtuin/SIRT1						
1	IgA	1	Hematocrit	1	Aβ/β-actin						
1	P-selectin	1	Mean platelet volume (MPV)	1	α-secretase (ADAM10)						
1	Matrix Metalloproteinase-10 (MMP-10)	1	Transferrin	1	Rab3						
1	Chemokine receptor 2 (CCR2) (protein2 list)	1	Haptoglobin	1	Rab7						
1	Beta 2-microglobulin (B2M)	1	White blood cells (WBC)	1	Early Endosome Marker (EEA1)						
1	FAS ligand belongs to TNF family	1	Total Urinary polyphenols (TUPs)	1	Lysosomal-associated membrane protein 2 (LAMP-2)						
1	CD8	1	Alpha-1-antitrypsin (alpha1-AT)	1	Microtubule-associated protein 1A/1B-light chain 3 (LC3)						
		1	Lactoferrin (LTF)	1	Phospholipase A2 (PLA2)						
		1	N-terminal pro b-type natriuretic peptide (NT-proβ	1	Carcinoembryonic antigen						
		1	Luteinizing hormone (LH)	1	Osteoprotegerin (OPG)						
				1	Neruogranin (NGRN)						
				1	Cellular prion protein (PrPc)						
				1	Kidney Injury Molecule (KIM-1)						
				1	Growth-regulated alpha protein (GRO-α)						
				1	Eotaxin-3						
				1	Unfolded p53						
				1	P-t181p/Ab1-42 ratio						

Table II. Frailty biomarkers by category and frequency

1.Frequency	1.Inflammatory/Immunity Markers	2.Frequency	2.Laboratory Markers	3.Frequency	3.Protein Markers	4.Frequency	4.Metabolomic Markers	5.Frequency	5.Oxidative Stress Markers	6.Frequency	6.Clinical Markers
33	IL6	10	Vitamin D	2	Propeptide of type I procollagen (PINP)	1	X12063	2	Serum 8-hydroxy-2-deoxyguanosine (8-OHdG)	4	Cardiovascular disease
24	C-reactive protein	6	Albumin	2	C-terminal telopeptide of type-1 collagen (Beta CTX)	1	Urate	1	Protein carbonyls	3	Increase Waist Circ/Waist-to-hip
6	Tumor necrosis factor (TNF-alpha)	5	Composite Score (multiple markers)	2	Extracellular heat shock protein (eHsp) 72	1	Mannose	1	thol level (TTL)	2	Calibrated Protein intake
6	Uric Acid	5	Lipids: Total Cholesterol	1	Cystatin C	1	Myostatin	1	Derivate of reactive oxygen metabolites (d-ROM)	1	Increased falls
5	Fibrinogen	5	Insulin like growth factor protein (IGF-1)	1	Cytomegalovirus			1	Malondialdehyde (MDA)	1	Alcohol intake
4	IL1beta	5	Parathyroid hormone (PTH)	1	C-terminal Agrin Fragment (CAF)					1	Change in Body Mass Index
3	Erythrocyte sedimentation rate (ESR)	5	White blood cells (WBC)	1	Sirtuin 1					1	More than 2 chronic diseases
3	Cortisol/Dehydroepiandrosterone ratio	4	Insulin resistance (IR-HOMA)	1	Sirtuin 2					1	Anticholinergic medications
2	IL1RA	3	Creatinine	1	Sirtuin 3						
2	Motif chemokine 10/ Interferon-gamma (CXCL-10/IFN-gama)	3	Glycohemoglobin (HbA1c)	1	Complement component protein (C1q)						
2	CD8	3	Hemoglobin	1	Klotho						
2	Dehydroepiandrosterone sulphate (DHEAS)	3	Lymphocytes	1	Lipoplysaccharide bining protein (LBP)						
1	IL2	2	Estimated glomerular filtration rate (eGFR)								
1	IL6R	2	Cobalamin deficiency (B12)								
1	IL18	2	Methylmalonic acid (MMA)								
1	TNF-a receptor I (TNFR1)	2	Carotenoids								
1	Cortisol	2	Lipids: Triglycerides								
1	Homocysteine	2	Neutrophils								
1	Beta 2-microglobulin (B2M)	2	Follistatin								
		2	Von Willebrand Factor Vllc								
		1	Creatinine Clearance								
		1	Glucose								
		1	Lipids: LDL cholesterol								
		1	Free thyroxine, FT4								
		1	Phytohemagglutinin								
		1	Pokeweed mitogen								
		1	Total Testosterone								
		1	Estrogen								
		1	Vitamin B6								
		1	Selenium								
		1	Blood urea nitrogen (BUN)								
		1	Thyroid stimulating hormone, TSH								
		1	Free thyroxine, FT3								
		1	Anemia								
		1	Hematocrit								
		1	Monocytes								
		1	Cystathionine								
		1	Ratio-Zinc/Copper								
		1	Total Urinary polyphenols (TUPs)								
		1	Total dietary polyphenols (TDPs)								
		1	alpha-tocopherol								
		1	alpha-1-antichymotrypsin (ACT)								
		1	Von Willebrand Factor Vlllc								

Table III. Cognitive frailty biomarkers by category and frequency

1.Frequency	1.Inflammatory/Immunity Markers	2.Frequency	2.Laboratory Markers	3.Frequency	3.Protein Markers	6.Frequency	6.Clinical Markers
6	C-reactive protein	2	Creatinine Clearance	1	Apolipoprotein A-I (ApoA1)	4	Change in Body Mass Index
4	IL6	2	Cobalamin deficiency (B12)	1	Prostaglandin F2-alpha	2	More than 2 chronic diseases
2	Dehydroepiandrosterone sulphate (DHEAS)	2	Insulin like growth factor protein (IGF-1)	1	Apolipoprotein B	1	Income
1	IL8	2	Vitamin D			1	Low level of education
1	IL1beta	2	White blood cells (WBC)			1	Alcohol intake
1	IL1alpha	1	Albumin			1	Elevated blood pressure
1	Fibrinogen	1	Creatinine			1	Elevated systolic pressure
1	CD8	1	Glucose			1	Cardiovascular disease
1	Homocysteine	1	Glycohemoglobin (HbA1c)			1	Psychoactive medications
1	Cortisol	1	Lipids: LDL cholesterol			1	Depression
		1	Alpha-linolenic acid			1	Instrumental activities of daily living (IADL)
		1	Anemia			1	Activities of daily living (ADL)
		1	Sodium				
		1	Phosphate				
		1	Polyunsaturated fatty acids (O3PUFAs)/ n-6/n-3 ratio				
		1	Hematocrit				
		1	Hemoglobin				
		1	Mean corpuscular volume (MCV)				
		1	Red blood cells (RBC)				
		1	White blood cells (WBC)				
		1	Lymphocytes				
		1	Monocytes				
		1	Neutrophils				
		1	Urate				
		1	Glucose				
		1	Total protein				
		1	Alanine aminotransferase (ALT)				
		1	Calcium				
		1	Lipids: Triglycerides				
		1	Lipids: Total Cholesterol				
		1	Free thyroxine, fT4				
		1	Ferritin,				
		1	Lipids: HDL cholesterol				
		1	Free thyroxine, fT3				
		1	N-terminal pro b-type natriuretic peptide (NT-proBNP)				

Table IV. Serum and genetic correlations by phenotype

Serum biomarker	Phenotype associated with serum biomarker	Genetic biomarker	Phenotype associated with genetic biomarker
Vitamin D (25(OH)D)	Frailty and cognitive decline	VDR (Vitamin D receptor)	Sarcopenia
Cystatin C	Frailty and cognitive decline	CST3 (cystatin)	Cognitive decline
Chemokine receptor 2 (CCR2)	Cognitive decline	CCL2	Cognitive decline
Myostatin	Frailty	MSTN (myostatin)	Sarcopenia
Klotho	Frailty	<i>KLOTHO</i>	Cognitive function
IL-6	Frailty and cognitive decline	IL-6	Sarcopenia and cognitive decline
TNF-alpha	Frailty and cognitive decline	TNF-alpha	Sarcopenia, frailty, and cognitive decline
IL-6R	Frailty and cognitive decline	IL-6R	Cognitive decline
CRP	Frailty and cognitive decline	AP2A2 (trait CRP), USP50 (trait CRP)	Cognitive decline
IL-1 β	Frailty and cognitive decline	IL-1 β	Cognitive decline
IL-18	Frailty	IL-18	Frailty
IL-12p70	Cognitive decline	IL-12A	Frailty
Brain derived neurotrophic factor (BDNF)	Cognitive decline	BDNFval66Met	Cognitive decline

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MANUSCRIPT 3:

Establishing Biological Plausibility for Cognitive Frailty: A Population Predictive Model

Abstract:

Background: This study aims to create a population predictive model to gain a more in-depth understanding of the underlying biological mechanisms for cognitive frailty as currently defined by the International Consensus Group in 2013. **Methods:** Data were from the InCHIANTI study, collected at baseline from 1998-2000. This group is a representative sample (n=1,453) of a population of white European origin from two small towns in Tuscany, Italy. To build our model, we used biomarkers with implications for clinical research and practice; a total of 132 putative SNPs and 155 protein biomarkers were identified from a systematic review (manuscript 2). We used a tree boosting model, Extreme Gradient Boosting (xgboost), a machine learning technique for supervised learning. **Results:** We developed two predictive models with high accuracy, AUCs for Model I is 0.877 (95% CI 0.825-0.903) and 0.864 (95% CI 0.804-0.899) for Model II. Results provide biological evidence for the relationship between cognitive decline and physical frailty supporting findings of dysregulation across multiple systems as the potential cause of cognitive frailty. One of the top predictors for cognitive frailty included anticholinergic burden with the presents of *SLCO1B1* rs4363657 (TMT-A $\beta = .20$, TMT-B $\beta = .38$). **Conclusions:** The results from this study establish a foundation for an understanding of the underlying biological mechanisms for the relationship between cognitive decline and physical frailty.

Introduction

The relationship between the phenotypes physical frailty and cognitive decline has been established in epidemiological studies. Both are associated with higher rates of disability, falls, mortality, an increase in health service need, and high direct/indirect costs to healthcare from long-term care and hospitalization¹⁻⁶. Evidence exists to support a longitudinal bidirectional relationship between physical function and cognitive decline; finding that associations between physical functioning and consequent cognitive decline are similar to associations with individuals with cognitive decline and consequent physical functioning⁷. These findings support an a priori hypothesis for shared biological mechanisms that underlie the association of physical and cognitive decline.

Although physical and cognitive impairment have been shown to be related, both phenotypes have long been studied separately⁴. To address this gap, the International Consensus Group organized by the International Academy on Nutrition and Aging (I.A.N.A) and the International Association of Gerontology and Geriatrics (I.A.G.G) convened in 2013 to identify related domains of physical frailty and cognition. The new construct called “cognitive frailty” is defined by the presence of physical frailty and cognitive impairment in the absence of Alzheimer’s disease or other dementias⁴. The International Consensus Group (I.A.N.A. /I.A.G.G.) report is an acknowledgment of the need to focus research efforts on a clinical condition characterized by the occurrence of physical frailty and cognitive impairment, in the absence of overt dementia diagnosis or underlying neurological conditions⁴. The cognitive frailty construct is considered a

heterogeneous clinical syndrome in older adults with evidence of: 1) physical frailty and cognitive impairment; and 2) exclusion of a clinical diagnosis of Alzheimer's Disease or other dementia⁴.

The introduction of this new phenotype demonstrates evidence for cognitive frailty as a subgroup of cognitive decline and physical frailty. Genetic risk factors and biological markers may be unique to individuals who present with cognitive frailty in contrast to those with isolated cognitive or physical decline. A model for detecting cognitive frailty could provide practitioners with the tools needed for early detection and secondary prevention for individuals with cognitive frailty. Currently, the instrumental assessments for cognitive frailty are time-consuming, expensive, require extensive training, and the clinical translation of these assessments is not clear⁴. Translating the cognitive frailty construct into the clinical setting is limited by the lack of consensus on an operational definition and considerable heterogeneity in the diagnostic criteria⁸. An understanding of the biomarkers that define cognitive frailty will help distinguish between changes related to normal aging, irreversible pathological process, and specific neurological diseases that may be reversible⁹. The strength in understanding the biological underpinnings of cognitive frailty is the ability to provide early detection and accurate diagnosis.

The primary purpose of this research was to create a population predictive model to gain a more in-depth understanding of the underlying biological mechanisms for cognitive frailty as currently defined by the International Consensus Group in 2013. This paper focuses on defining the shared mechanisms for physical frailty and cognitive

impairment and establishing a model for determining the presence of risk factors that may predict cognitive frailty in the clinical setting. An important innovation in this study was the use of machine learning (ML) statistical modeling to define the differences between the following groups: cognitive decline, physical frailty and cognitive frailty. The study builds an algorithmic classifier for cognitive frailty with candidate factors identified by a systematic review (results published elsewhere). Notably, the identification of unique biomarkers may also serve to group patients by underlying pathophysiologic processes and further refine the assignment to a clinical diagnostic category. Such precision in the determination of genetic and biological biomarkers related to cognitive frailty will lead to a better understanding of the interrelated pathology between physical frailty and cognitive impairment and, ultimately, to early detection and targeted interventions focused on the prevention of cognitive and functional disabilities.

Methods

Study Population

Figure 1 shows a summary of our workflow, further details on phenotypes and the list of biomarkers are available in the supplementary appendix. Clinical, protein, and genetic biomarker samples were from participants of the InCHIANTI study, collected at baseline from 1998-2000. This group is a representative sample (n=1,453) of the population of white European origin from two small towns in Tuscany, Italy. The primary aim of the InCHIANTI study to evaluate function and mobility in older community-dwelling individuals. A detailed description of the study design, data collection, and sampling

procedure are published elsewhere¹⁰. This secondary study was approved by the ethics committee at *Centre de recherché Clinique du CHUS*, project #547.

Predictive Measures

The International Consensus Group's (I.A.N.A. /I.A.G.G.) list of potential biomarkers is not meant to be complete, accurate, or exhaustive⁴. Since an exhaustive list of biomarkers is not present in the literature; we used a systematic review to identify factors associated with cognitive decline, physical frailty, and cognitive frailty based on the current operational definitions (Sargent et al., 2018). We searched the following online databases: PubMed, Embase, Scopus, Web of Science, LILACS, Gene Indexer, and GWAS Central. Databases were searched from the start date of the database to 22 December, 2015. An update of the searches was performed prior to the data extraction phase on 26 May, 2016 to identify any new publications. The systematic review resulted in 327 articles for the final synthesis, identifying 456 predictive protein and genetic biomarkers. A total of 289 variables identified from the systematic review were available in the InCHIANTI database. Variables were removed if there was > 12% missing data, resulting in 132 putative SNPs and 155 protein biomarkers. To build our model, we used protein markers with implications for clinical research and practice, and completed genetic risk score estimates (i.e. the cumulative genetic risk burden estimated from SNPs of interest, or GRS) before including the individual single nucleotide polymorphisms (SNPs) in the final models. Many of the protein markers included in our model are used clinically for detection of disease; therefore we organized the results by using the clinical designation identified by clinical pathology laboratories. The categories

include inflammation/immunity, nutrient, lipid metabolism, metabolomics, renal/electrolyte, hematology/liver, endocrine/hormones, and clinical features. Known predictive clinical features identified repeatedly in the systematic review were age, depression, gender, and level of education. Baseline diagnosis of dementia was included in the models for frailty and cognitive frailty. Additionally, systematic review identified a group of medications, specifically anticholinergic medications, as a risk for cognitive and physical decline^{11,12}. Anticholinergic burden was calculated using the Anticholinergic Cognitive Burden Scale (ACB) and examined as a predictor for all phenotypes.

Outcome Measures

Neuropsychological tests include the Mini-Mental State Examination (MMSE) as a test of general cognition and the Trail Making Test, Part A and B (TMT). Psychomotor speed is assessed using the TMT-A, scoring based on time in seconds to completion with a score range of 0 to 300 seconds¹³. The executive functioning domain was assessed using the TMT-B (any individual scoring 300-600 seconds were included as 300)¹³. TMT, part A and B cut off scores are based off of established norms for mild neurocognitive disorders¹⁴. Normative data for time to complete the TMT tests in seconds was stratified by age and education¹⁵. Additionally, the neuropsychological profile for individuals with cognitive frailty is different from those with frailty or cognitive decline alone with recent findings of lower performance on TMT tests, scoring worse on executive and attention domains¹⁶. The CES-D self-report scale (0-60) is used to measure depressive symptoms. Reliability, validity, and factor structure have been similar across a diverse demographic

and the scale has been used extensively in epidemiologic studies for depression and physical function¹⁷.

Frailty measures included the number of frailty symptoms for subjects ≥ 65 years of age. Frailty as defined by the cardiovascular health study (CHS), allows for a continuous scoring system versus a nominal system because it can capture the multidimensional nature of frailty¹⁸. The InCHIANTI criteria for frailty defined unintentional weight loss as losing weight not related to diet, classified the values of body mass index, strength, walking speed and height based on all subjects ≥ 65 years and used two questions of the CES-D for the definition of exhaustion.

In this study two models of cognitive frailty were developed, because conceptually the models need to cover variables of physical frailty and cognitive decline for populations seen in geriatric and primary care centers with implications for future clinical research and translation into practice. Primary care has a key role in early identification of cognitive and physical decline. The MMSE, despite known limitations for the diagnosis of dementia, has retained popularity in the primary care setting with increased use for screening and diagnosis and is recommended by the Alzheimer's Society¹⁹. *Model I* defines cognitive decline and cognitive frailty with the use of criteria from the MMSE while *Model II* defines these phenotypes with participants who have completed the MMSE with additional Trail Making Tests, Part A and B²⁰⁻²². In this study frailty was characterized by individuals with one or more of the frailty criteria, including pre-frail and frail as one group¹. Cognitive frailty is defined as individuals with cognitive decline and one or more of the frailty criteria¹⁶.

Statistical Analysis

The supplementary appendix includes additional details of the statistical methods, beginning with detail about model development in the InCHIANTI dataset, which we used to train and test the initial model, internal validation, and calibration of the model. Evidence supports the use of tree boosting models using Extreme Gradient Boosting (xgboost) in R, statistical software, as an effective method for building a reproducible predictive model for the detection of a complex heterogeneous phenotype with large numbers of potential biomarkers^{23,24}. Boosted trees, a machine learning technique for supervised learning, are ensembles of regression trees, similar to decision trees and are used for prediction or classification. Xgboost is based in boosted trees and provides more efficient and accurate predictive modeling with large datasets and a rapid / robust framework for feature selection. Statistical modeling is used to design, test, and validate an accurate method for classifying patients into phenotypic outcomes.

The tree boosting model for the evaluation of multiple variables simultaneously provides a high predictive value with low bias. Additionally, parameters are set to prevent over fitting for the models. The data were randomly divided, two thirds was assigned to the training cohort, and one third was assigned to the validation cohort. One of the features that is central to xgboost is its ability to combine multiple trees or “weak predictors” to reach maximum prediction performance while reducing bias. This approach uses large amounts of data from different aspects of clinical, genetic, and biomarker research, strengthening the models’ generalizability and classification power. Xgboost iteratively re-weighs the variables, taking a weighted majority; the parameters

identified after pruning comprised the final predictive model²⁵. None of the candidate features in the models are used in the diagnosis of cognitive decline, physical frailty, or cognitive frailty. This standard technique prevents circularity, overestimation, and over fitting for both the models generated. Parameters for the model include: max depth = "10", nthread = "12", nrounds = 5-200, objective = "binary:logistic", evaluation metric = "auc", silent = "1", gamma = default = "0" to control the number of trees, and eta default = "0.3" to prevent over fitting. We used the default setting for all other parameters which can be found in the xgboost 0.6 documentation²⁴.

To evaluate the models, we used the evaluation metric area under the receiver operating curve (AUC). AUC were calculated from each model and used to determine discrimination of participants with cognitive frailty (case), cognitive decline (case), and physical frailty (case) from healthy individuals (control) in the training cohort. An AUC of 0.5 was considered chance, > 0.8 informative, and > 0.9 clinically relevant.

The xgboost algorithm iteratively determines the maximum function of a model based on a tree building algorithm (quadratic problem) which creates a node then assigns a prediction point to each leaf; the assigned number is termed "gain". Once the model has reached maximum depth, pruning occurs by taking out the nodes with a negative gain and keeping those with a positive gain. Results from the population predictive model are ranked by gain which is a metric based on each feature's contribution in the model. When comparing top features to other features in the model, the higher the gain the more important the feature is for prediction of the outcome. Cover is a measure of the relative quantity of observations found by one feature and frequency is the percentage

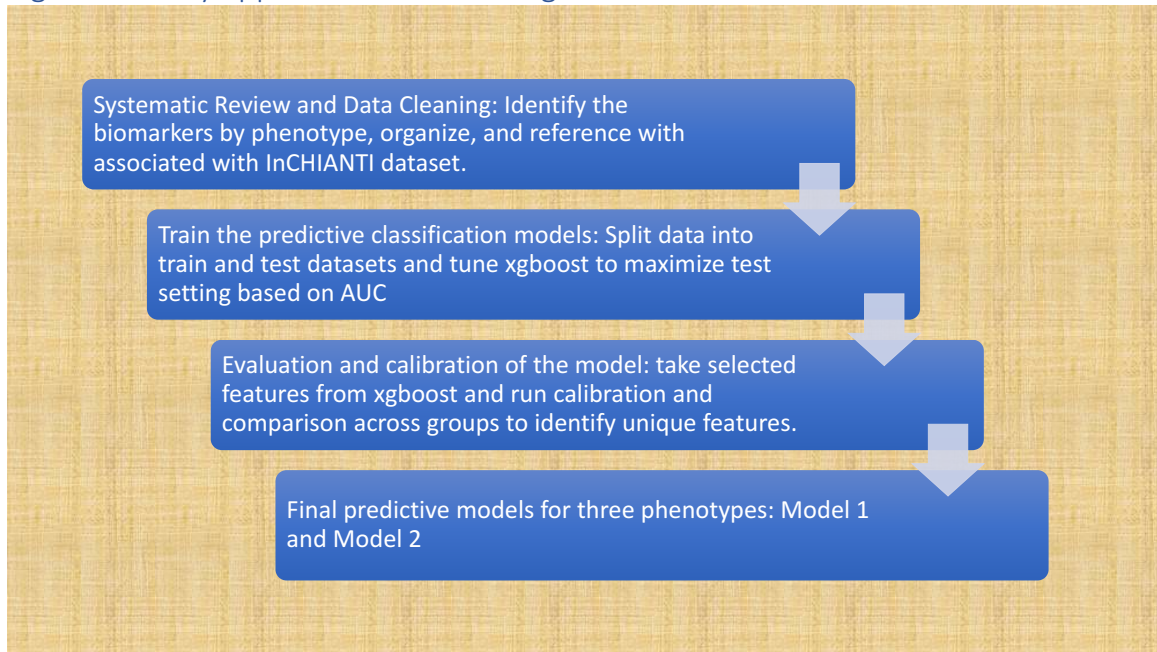
representing the relative number of time a feature is used in the trees of the model²⁴.

Gain is the most relevant metric to interpreting the rank and importance of each feature.

A case-control design is used to study genome wide variations between participants with cognitive frailty (case) and those with only cognitive decline (control), only physical frailty (control), and healthy individuals (control). Univariate analysis, *t*-tests for continuous and chi-squared tests for binomial traits, were used to determine the significance of the predictor. We used logistic regression for case-control analyses under additive allele dosage. To evaluate additive effects of SNPs, a positive regression coefficient means that each copy of the allele of interest increases the risk for the cognitive frailty phenotype^{26,27}. The appendix includes further details and results about the generation of the genetic data and creation of the GRS from 132 genetic risk factors implicated in one or more studies from the systematic review. Our study used the high-performance computational capabilities of the Biowulf Linux cluster at the National Institutes of Health (Bethesda, MD, USA) in the and genotypic data from the InCHIANTI study.

The final models identified features that were predictive of cognitive frailty with unique features for cognitive decline and physical frailty. Mechanisms that contribute to the development of cognitive frailty were determined by evaluation of fluid biomarkers and genome wide genetic variability as a predictor of the development and persistence of cognitive frailty.

Figure 1. Study approach workflow diagram



Note: Profile of model development and validation workflow. Blue boxes indicate steps of the workflow specific to the InCHIANTI data set.

Results

A total of 1,453 adults participated, 1,326 provided blood samples at baseline.

Participants had a mean age of 69 years (S.D.=15.7), 56% were female and 44% were male, and completed a secondary level of education. All participants completed the MMSE, 369 participants scored ≤ 23 ($M=25$, S.D.=5.1), 525 scored ≥ 78 on the TMT-A ($n=1,240$), and 634 scored ≥ 106 on the TMT-B ($n=1,057$).

The supplementary appendix (tables IV-IX) contains the tables for final predictive model features ranked by gain. The results show predictive features for cognitive frailty when measured using the MMSE (Model 1) and TMT part A and B (Model II) with unique features for cognitive decline and physical frailty in both models. Bivariate results for clinical, genomic, and protein biomarkers are shown in the appendix (tables X - XVIII).

For discrimination of participants with cognitive frailty from healthy controls, the AUC of Model I is 0.877 (95% CI 0.825-0.903) and 0.864 (95% CI 0.804-0.899) for Model II. Parameter estimates for each predictive factor and associated descriptive statistics were evaluated to provide biological insight into the underpinnings of the classification algorithm. Next, we carried out calibration tests for all possible values between 5-200 groups and evaluated the distribution of the test statistics per subgrouping. We noted a normal distribution of AUCs across all iterations, with no statistically significant deviation from the expected values in any group, suggesting good model fit. Both models showed high accuracy with AUCs ranging from 0.808-0.877 for model I and 0.831-0.864 model II within the framework of the calibration tests.

Demographic features and anticholinergic burden results are shown in Table 5-6 and significant differences between healthy control and phenotype are shown in Table 10 of the supplementary appendix. Gender was a predictor for all three phenotypes in Model I but not a predictor in Model II. There were more females than males with cognitive decline for all three phenotypes in both models. Baseline diagnosis of dementia, while found to be a predictor in Model I for frailty and cognitive frailty was not a predictor in Model II. Anticholinergic burden (ACB) was a predictor for all three phenotypes in both models with larger ACB mean scores for those with cognitive decline, frailty, and cognitive frailty. In Model II, anticholinergic burden had a significant effect on both psychomotor speed (TMT-A) and executive functioning (TMT-B) for all three phenotypes. Anticholinergic burden was found to be one of the top predictors for all phenotypes in model I and II. Detailed analyses for anticholinergic burden are described

elsewhere and included in the results tables of this manuscript (Sargent et al., 2018 in manuscript 4).

Genomic results

Table 1 and 2 shows the comparison of genomic features by phenotype for Model I and Model II respectively.

Model I

Ten genes were predictive of cognitive frailty measured by the MMSE and CHS criteria; four genes are unique to cognitive frailty: (*BIN1*) rs7561528 allele A ($\beta = -.04$), *ACE* rs4968782 allele G ($\beta = .10$), and *WTAPP1* rs603050 allele G ($\beta = -.14$), *MTRR* rs1801394 allele G ($\beta = .80$) and six overlap with features associated with cognitive decline and frailty: *IL6* rs1800796 allele C ($\beta = .25$), (*ACOT11*) rs12752888 allele C ($\beta = -.47$), *DAB1* rs1539053 allele A ($\beta = .51$), (*MMP3*) rs948399 allele C ($\beta = .41$), *CD33* rs3865444 allele A ($\beta = .62$), and *UBR5* rs7840202 allele C ($\beta = -.15$). Of these markers five showed a significant difference between control and cognitive frailty: (*ACOT11*) rs12752888 ($p = .001$), *DAB1* rs1539053 ($p = .01$), (*MMP3*) rs948399 ($p = .01$), *CD33* rs3865444 ($p = .03$), and *MTRR* rs1801394 ($p = .001$).

Four SNPs were uniquely associated with frailty: *CNTN5* rs10501927 allele G ($\beta = -.10$), *WTAPP1* rs11225434 allele C ($\beta = .10$), *SORL1* rs4935774 allele C ($\beta = .04$), and *CREBBP* rs129968 allele A ($\beta = .10$) Eight SNPs are unique to cognitive decline *BTRC* rs10883631 allele G ($\beta = .11$), *TOMM40* rs2075650 allele G ($\beta = .10$), *IL6R* rs2228145 allele C ($\beta = -.31$), *USP50* rs3131609 allele C ($\beta = .10$), *COMT* rs4646316 allele T ($\beta = -.62$), *AP2A2* rs7396366 allele C ($\beta = .10$), *KLOTHO* rs9527025 allele C ($\beta = .20$).

Model II

Individual variants were predictive for psychomotor speed (TMT-A) and executive functioning domain (TMT-B). Significant differences between control and disease are shown in appendix (tables XVI - XVIII).

Twenty-one genes were predictive of cognitive frailty measured by TMT and CHS criteria in model II; eight are unique to cognitive frailty *ACE* rs4316 allele T (TMT-A β = -.07, TMT-B β = -.06), *ACE* rs1800764 allele C (TMT-A β = .06, TMT-B β = .06), *EPHA1* rs11771145 allele A (TMT-A β = -.10, TMT-B β = .13), *CREBBP* rs129968 allele A (TMT-A β = .05, TMT-B β = .03), *TNF* rs1800629 allele A (TMT-A β = .15, TMT-B β = .10), *IL18* rs360722 allele A (TMT-A β = .05, TMT-B β = -.02), *WTAPP1* rs603050 allele T (TMT-A β = -.21, TMT-B β = -.10), and *SELP* rs6131 allele T (TMT-A β = -.07, TMT-B β = -.03).

Thirteen of the cognitive frailty genetic features overlap with variants from cognitive decline and frailty: (*MMP3*) rs948399 allele C (TMT-A β = .29, TMT-B β = 0.02), (*ACOT11*) rs12752888 allele C (TMT-A β = -.34, TMT-B β = -.37), *APOE* rs429358 allele C (TMT-A β = -.23, TMT-B β = -.59), *SLCO1B1* rs4363657 allele C (TMT-A β = .20, TMT-B β = .38), *TOMM40* rs8106922 allele G (TMT-A β = -.31, TMT-B β = .09), *CNTN5* rs10501927 allele G (TMT-A β = -.11, TMT-B β = -.06), *SORL1* rs1614735 allele G (TMT-A β = .02, TMT-B β = .07), *IL1-beta* rs16944 allele A (TMT-A β = -.01, TMT-B β = -.13), *ACE* rs4343 allele A (TMT-A β = -.02, TMT-B β = -.02), (*SSB*) rs11894266 allele C (TMT-A β = -.05, TMT-B β = -.06), *UBR5* rs7840202 allele C (TMT-A β = -.06, TMT-B β = -.05), *MAPT* rs3785880 allele G (TMT-A β = -.06, TMT-B β = -.05), *BTRC* rs10883631 allele G (TMT-A β = -.01, TMT-B β = .01).

Of these markers five showed a significant difference between control and cognitive frailty for psychomotor speed or executive functioning: (*ACOT11*) rs12752888 allele C (TMT-A, $p = .01$, TMT-B $p = .02$), *APOE* rs429358 allele C (TMT-B, $p = .01$), *SLCO1B1* rs4363657 allele C (TMT-B, $p = .02$), *TOMM40* rs8106922 allele G (TMT-A, $p = .05$), (*MMP3*) rs948399 allele C (TMT-A, $p = .05$).

Frailty has one unique SNP: *NECTIN2* rs6859 allele A (TMT-A $\beta = -.02$, TMT-B $\beta = -.07$). and cognitive decline has eleven unique SNPs: *KCNU1* rs1157242 allele T (TMT-A $\beta = .13$, TMT-B $\beta = .44$), *SORL1* rs1133174 allele A (TMT-A $\beta = .05$, TMT-B $\beta = .02$), *KLOTHO* rs1207568 allele A (TMT-A $\beta = -.05$, TMT-B $\beta = -.18$), *GCKR* rs1260326 allele C (TMT-A $\beta = .02$, TMT-B $\beta = .08$), *COMT* rs4680 allele A (TMT-A $\beta = -.02$, TMT-B $\beta = .06$), *SORL1* rs4935774 allele C (TMT-A $\beta = .11$, TMT-B $\beta = .05$), *ATM* rs611646 allele T (TMT-A $\beta = .08$, TMT-B $\beta = .04$), *MS4A4E* rs676309 allele C (TMT-A $\beta = -.07$, TMT-B $\beta = -.17$), *SLC2A9* rs737267 allele T (TMT-A $\beta = .10$, TMT-B $\beta = -.08$), *TCN2* rs740234 allele G (TMT-A $\beta = -.02$, TMT-B $\beta = -.10$), (*BIN1*) rs744373 allele G (TMT-A $\beta = .01$, TMT-B $\beta = -.15$). Cognitive decline and frailty have three shared SNPs that were not features for cognitive frailty *PRNP* rs1799990 allele G (TMT-A $\beta = .45$, TMT-B $\beta = .30$), *CR1* rs3818361 allele A (TMT-A $\beta = .20$, TMT-B $\beta = .14$), and *ABCA7* rs4147929 allele A (TMT-A $\beta = .02$, TMT-B $\beta = .03$).

Protein biomarker results

Tables III and IV shows a comparison of the protein markers by category and phenotype. Significant differences between control and cognitive frailty are shown in the supplementary appendix (Tables XI-XVIII). The results show a mean difference in the laboratory value between healthy controls and those with cognitive decline, physical

frailty, and cognitive frailty. In Model I and Model II, all phenotypes share features in all categories and each phenotype has unique features. Cognitive frailty in Model I has seven unique features transforming growth factor B1 and fatty acid 22:0 with a mean increase in cystatin C ($p < 0.0001$), decrease serum calcium ($p = .0004$), increase serum creatinine ($p = .02$), increase urine nitrites ($p = .02$), increase soluble transferrin receptor ($p = .01$) for individuals with cognitive frailty compared to healthy controls. Cognitive frailty (Model I) shared 70 of the 91 features with frailty and 53 of the 93 protein fluid biomarkers features with cognitive decline. Cognitive frailty in Model II had only two unique features; urine glucose and serum IGF binding protein; IGF binding protein is decreased in individuals with cognitive frailty for psychomotor speed ($p = .0001$) and executive functioning ($p = .0004$). Cognitive frailty (Model 2) shared 70 of the 90 features with frailty and 82 of the 125 protein fluid biomarkers features with cognitive decline.

Discussion

In this study, we developed two models using xgboost for the prediction of cognitive frailty and further defined the association between cognitive decline and frailty. Both models have a larger population of women with older age being associated with cognitive frailty. Anticholinergic burden was highly predictive of cognitive frailty and is found as a unique predictive feature of frailty and cognitive decline in both models.

Genomic results suggest that Model I and Model II are measuring different variants. Model I has unique genomic features *DAB1* rs1539053 allele A, *CD33* rs3865444 allele A, and *MTRR* rs1801394 allele G, as predictive of cognitive frailty. *CD33* has putative functions in the immune system involved in processes at the cell membrane with links

to greater cell surface expression of monocytes and is considered an Alzheimer's disease susceptibility loci²⁸. *DAB1* is required for the organization of multiple neuronal types in the cerebral cortex and is important for normal cognitive function^{29,30}. *MTRR* rs1801394 is a marker for vitamin B12 in a pathway with methylmalonic acid (MMA) levels³¹. Lower serum MMA leads to higher serum lipids and higher homocysteine levels potentially leading to reduced energy metabolism³¹. All three of these protein markers were found in the cognitive frailty model I. Additionally, *MTRR* has been linked to 2-4 times greater odd of being frail.

One of the interesting genomic findings was *SLCO1B1* rs4363657 allele C that is predictive of frailty and cognitive frailty in Model II. The *SLCO1B1* has been associated with X12063 which is a metabolite, both are associated as markers of lean muscle mass loss³². Additionally, *SLOCO1B1* has been linked to drug metabolism specifically, higher blood concentrations of statins³³. *SLOCO1B1* is essential for the hepatic uptake and the C variant is associated with reduced *OATP1B1* activity. *OATP1B1* can facilitate drug uptake and at the blood-brain barrier may affect the distribution of drugs into the central nervous system³⁴. The association with anticholinergic metabolism and *SLOCO1B1* has not been explored. Variants in model I and II included *MMP3* and (*ACOT11*). *MMP3* rs948399 allele C is predictive of frailty and cognitive decline and (*ACOT11*), rs12752888 allele C is a member of the acyl-CoA thioesterase family that catalyzes the conversion of activated fatty acids³⁵. In this study (*ACOT11*) rs12752888 allele C was found to have a protective effect. (*ACOT11*) rs12752888 has not been studied in individuals with physical frailty or cognitive frailty previously.

Protein marker results show a relationship between neuroinflammatory cytokines and cognitive frailty. Neuroinflammatory cytokines (nonantibody proteins) have a role in the neuroimmunoendocrine processes and have been postulated to be related to cognition due to their ability to penetrate the blood-brain barrier and affect the central nervous system¹. This study found elevated levels of neuroinflammatory cytokines with interleukins IL1, IL6, IL6R, and tumor necrosis factors (TNF) as predictive features for cognitive frailty in both models along with associated genetic markers: *IL6* rs1800796, *IL6R* rs2228145, *TNF* rs1800629, and *IL1-beta* rs16944. Additionally, participants with cognitive frailty had higher levels of resistin ($p < .0001$) compared to controls in both models; resistin regulates IL-6, TNF, and hs-CRP². Both fibrinogen and advanced glycation end product (AGE) ($p < .0001$) were both found to be elevated showing a link to oxidative stress and high levels of alpha-2 globulin (A2M) ($p < .0001$). A2M is considered a protease inhibitor cytokine transporter linked to Alzheimer's disease was found in participants with cognitive frailty³. Several studies have shown a relationship between many of these neuroinflammatory markers and cognitive and physical decline^{5,6}. In this study, we found many of these markers to be predictive for both cognitive decline and physical frailty. Additionally, these patterns of neuroinflammatory cytokines have been found in the InCHIANTI study to be associated with other complex chronic disease highlighting comorbidity as a confounding factor⁴.

Dehydroepiandrosterone sulfate (DHEA) was found to be low for those with cognitive frailty when compared to control ($p < 0.001$). DHEA has been found to inhibit IL-6 providing a connection between endocrine and immune function. Another interesting

finding is the connection between nutrition and cognitive frailty with low fatty acid levels and high levels of c-terminal telopeptide of type-1 collagen I (PINP) and parathyroid hormone (PTH). Both PINP and PTH have been linked to low levels of vitamin D which was a finding in this study for participants with cognitive frailty⁷. Methylnalonic acid (MMA) is linked to vitamin B12 and high levels of homocysteine found in both models ($p < .0001$) in addition, *MTRR* rs1801394 is associated with the same pathway. Serum MMA has been linked to both cognitive performance and increased risk for frailty^{8,9}.

Metabolomic (ceramides C16:0, C20:0, C20:5, C22:0, C24:0) markers were found in both models, some markers were found to be elevated and others low for participants with cognitive frailty. Since this study evaluated individuals with early cognitive decline at a single time point it is possible that serum ceramides varied according to the timing and onset of memory impairment and need to be explored further^{10,11}.

Cognitive frailty model I (n=101) and II (n=110) feature comparison show a difference in some biomarkers however, there were 66 shared biomarkers; 58 protein, 4 genomic, and 4 clinical markers. Some differences in the model features suggest lack of concordance between the clinical measures MMSE and TMT part A and B. These observations highlight the fact that pathways between clinical decision tools and precision science are not strictly linear in nature. When comparing models I and II for all phenotypes less variability with fewer unique features and more shared mechanisms.

There are several potential genomic and protein biomarker interactions, which are not fully explored in this manuscript. We did not attempt to complete a comprehensive

pathway analysis for the variables in the predictive models. The exploratory nature of this work will encourage new research into understanding these pathways. The study included a small homogenous sample with large numbers of biomarkers creating limitations for translation into clinical research. Additionally, the study was retrospective using existing data. Future research should be directed towards understanding the potentially reversible cause of cognitive frailty, validating the models in epidemiological data with more diverse demographic groups, and exploring the predictive features in prospective studies.

Conclusion

The results from this study support the use of an innovative Boosted trees machine learning technique in developing a population based predictive model for a complex condition of aging, cognitive frailty. Results provide biological evidence for the relationship between cognitive decline and physical frailty supporting findings of dysregulation across multiple systems as the potential cause of cognitive frailty. The results from this study begin to unravel the complex biological network behind the association between cognitive decline and physical frailty.

Table I. Genomic features by phenotype model I

SNP	Associated Allele	Chromosome	Gene	Cognitive Decline	Frailty Genomic	Cognitive Frailty
rs10883631	G	10	BTRC	X		
rs12752888	C	1	ACOT11/LOC105378734	X		X
rs1539053	A	1	DAB1	X	X	X
rs1800796	C	7	IL6	X		X
rs2075650	G	19	TOMM40	X		
rs2228145	C	1	IL6R	X		
rs3131609	C	15	USP50	X		
rs4646316	T	22	COMT	X		
rs7396366	C	11	AP2A2	X		
rs948399	C	11	MMP3	X	X	X
rs9527025	C	13	Klotho	X		
rs10501927	G	11	CNTN5		X	
rs11225434	C	11	WTAPP1		X	
rs129968	A	16	CREBBP		X	
rs3865444	A	19	CD33		X	X
rs4935774	C	11	SORL1		X	
rs7840202	C	8	UBR5		X	X
rs1801394	G	5	MTRR			X
rs4968782	G	17	ACE			X
rs603050	T	11	WTAPP1			X
rs7561528	A	2	BIN1/LOC105373605			X

Note: bold text indicates the closes gene

Table II. Genomic features by phenotype model II

SNP	Associated Allele	Chromosome	Gene	Cognitive Decline	Frailty Genomic	Cognitive Frailty
rs10501927	G	11	CNTN5	X	X	X
rs1133174	A	11	SORL1	X		
rs1157242	T	8	KCNU1	X		
rs1207568	A	13	Klotho	X		
rs1260326	C	2	GCKR	X		
rs12752888	C	1	ACOT11/LOC105378734	X	X	X
rs1614735	G	11	SORL1	X		X
rs16944	A	2	IL-1beta	X		X
rs1799990	G	20	PRNP	X	X	
rs3818361	A	1	CR1	X	X	
rs4147929	A	19	ABCA7	X	X	
rs4343	A	1	ACE	X		X
rs4680	A	22	COMT	X		
rs4935774	C	11	SORL1	X		
rs611646	T	11	ATM	X		
rs676309	C	11	MS4A4E	X		
rs737267	T	4	SLC2A9	X		
rs740234	G	22	TCN2	X		
rs744373	G	2	BIN1	X		
rs948399	C	11	MMP3	X	X	X
rs429358	C	19	APOE		X	X
rs11894266	C	2	SSB		X	X
rs8106922	G	19	TOMM40		X	X
rs7840202	C	8	UBR5		X	X
rs3785880	G	17	MAPT		X	X
rs10883631	G	10	BTRC		X	X
rs4363657	C	12	SLCO1B1		X	X
rs6859	A	19	NECTIN2		X	
rs11771145	A	7	EPHA1			X
rs129968	A	16	CREBBP			X
rs1800629	A	6	TNF			X
rs1800764	C	17	ACE			X
rs360722	A	11	IL-18			X
rs4316	T	17	ACE			X
rs603050	T	11	WTAPP1			X
rs6131	T	1	SELP			X

Note: bold text indicates the closes gene

Table III. Protein and clinical features by phenotype model I

	Cognitive Decline	Frailty	Cognitive Frailty
Clinical Features			
Age	X	X	X
Anticholinergic Burden	X	X	X
Depression	X	X	X
Gender	X	X	X
Level of Education	X	X	X
Baseline Diagnosis of Dementia		X	X
Inflammatory/Immunity			
24-hour urinary cortisol (Åµg/24 hours)	X	X	X
Urinary cortisol (Åµg/mL)	X		X
Adiponectin via RIA (Åµg/mL)	X		X
Alpha-1 globulin (%)	X		X
Alpha-2 globulin (%)	X		X
Alpha-2-macroglobulin (mg/dL)	X		X
Cortisol:DHEAS ratio (based on nmols)	X	X	X
Dehydroepiandrosterone sulfate (Åµg/dL)	X	X	X
Fibrinogen (mg/dL)	X		X
Homocysteine via FPIA analysis (Åµmol/L)	X	X	X
Interleukin-10 via ELISA (pg/mL)	X		X
Interleukin-12 via Bio-Plex (pg/mL)	X	X	X
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	X		
Interleukin-6 via ELISA ultrasensitive (pg/mL)	X	X	
Resistin via EIA (ng/mL)	X	X	X
Serum cortisol (Åµg/dL)	X	X	X
Soluble IL-6 receptor via ELISA (ng/mL)	X	X	
Soluble TNF-a receptor I via quantitative sandwich EIA (pg/mL)	X	X	X
TNF-related apoptosis-inducing ligand (pg/mL)	X	X	X
Uric acid (mg/dL)	X		X
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)		X	X
Beta globulins (%)		X	X
C-reactive protein - high sensitivity (Åµg/mL)		X	X
Endogenous secretory receptor for AGEs (ng/mL)		X	X
Erythrocyte sedimentation rate (ESR) (mm/hour)		X	X
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)		X	X
Interleukin-1B via ELISA (pg/mL)		X	X
Interleukin-8 via Bio-Plex (pg/mL)		X	
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)		X	X
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)		X	X
Soluble CD14 via ELISA (ng/mL)		X	X
Soluble TNF-a receptor II via quantitative sandwich EIA (pg/mL)		X	X
Tumor necrosis factor-a via multiplex technology (pg/mL)		X	X
Cystatin C (mg/L)			X
Transforming growth factor-B1 (pg/mL)			X
Renal/Electrolyte			
24-hour urinary creatinine (mg/24 hours)	X	X	X
Blood urea nitrogen (mg/dL)	X	X	X
Creatine phosphokinase (U/L)	X	X	X
Creatinine clearance, 24-hr urine (mL/minute)	X	X	
Cystatin C (mg/L)	X	X	
Urinary Ca (mmol/L)	X	X	X
Urinary Na (mmol/L)	X	X	
24-hour urinary cortisol (Åµg/24 hours)		X	
Na+ (mEq/L)		X	
Urinary creatinine (mg/dL)		X	X
Urine proteins (mg/dL)		X	X
Ca++ (mg/dL)			X
Serum creatinine (mg/dL)			X
Urine nitrites			X
Nutrient Biomarker			
Albumin (%)	X	X	X
Beta-carotene via high performance liquid chromatography (Åµmol/L)	X	X	X
Lycopene via high performance liquid chromatography (Åµmol/L)	X	X	
Omega-3 plasma fatty acid weight (mg/L)	X	X	X
Omega-6 plasma fatty acid weight (mg/L)	X	X	X
Omega-6 fatty acids as % of total fatty acid area		X	X
Ratio of Omega-6:Omega-3 as % of total fatty acid area	X		
Ratio of Omega-6:Omega-3 as % of total fatty acid mols	X		
Total proteins (g/dL)	X		
Vitamin E alpha tocopherol, high performance liquid chromatography, (Åµmol/L)	X	X	X
Vitamin B6 via high performance liquid chromatography (ng/mL)		X	X
Vitamin E gamma tocopherol, high performance liquid chromatography, (Åµmol/L)		X	X

	Cognitive Decline	Frailty	Cognitive Frailty
Hematology/Liver			
Ferritin (ng/mL)	X	X	X
Folate via RIA (ng/mL)	X	X	
Gamma glutamyl transferase (U/L)	X	X	X
GPT (also known as ALT) (U/L)	X	X	
Lymphocytes (n, K/ÅµL)	X		
MCH concentration (MCHC) (g/dL)	X	X	X
Mean corpuscular hemoglobin (MCH) (pg)	X		
Mean corpuscular volume (MCV)	X		
Methylmalonic acid MMA (Åµmol/L)"	X		
Monocytes (%)	X	X	X
Red blood cells (RBC) (n, millions/ÅµL)	X		
Red cell distribution width (RDW) (%)	X	X	X
Vitamin B12 via RIA (pg/mL)	X	X	X
White blood cells (WBC) (n, K/ÅµL)	X	X	X
Hematocrit (%)		X	
Hemoglobin (g/dL)		X	X
Lymphocytes (%)		X	X
Mean corpuscular volume (MCV) (fL)		X	X
Mean platelet volume (MPV) (fL)		X	X
Methylmalonic acid, MMA (Åµmol/L)		X	
Monocytes (n, K/ÅµL)		X	
Neutrophils (%)		X	
Neutrophils (n, K/ÅµL)		X	
Retinol via high performance liquid chromatography (Åµmol/L)		X	X
Soluble transferrin receptor (nmol/L)			X
Lipid Metabolism			
Lipids: HDL cholesterol (mg/dL)	X	X	X
Lipids: total cholesterol (mg/dL)	X	X	
Lipids: triglycerides (mg/dL)	X		
Lipoprotein(a) (mg/dL)	X	X	
Lipids: LDL cholesterol (mg/dL)		X	X
Metabolomics(plasma lipids)			
Fatty acid C16:0 (palmitic) area	X		X
Fatty acid C16:0 as % of total fatty acid area	X	X	X
Fatty acid C16:0 as % of total fatty acid weight	X	X	X
Fatty acid C16:0 (Åµmol/L)		X	
Fatty acid C20:0 (arachidic) area	X		X
Fatty acid C20:0 as % of total fatty acid weight	X	X	
Fatty acid C20:0 weight (mg/L)	X	X	X
Fatty acid C20:0 as % of total fatty acid area		X	X
Fatty acid C20:5 n-3 as % of total fatty acid weight	X		X
Fatty acid C20:5 n-3 weight (mg/L)	X	X	
Fatty acid C20:5 n-3 as % of total fatty acid area		X	X
Fatty acid C22:0 (behenic) area	X		X
Fatty acid C22:0 weight (mg/L)	X	X	
Fatty acid C22:0 as % of total fatty acid area			X
Fatty acid C24:0 (lignoceric) area	X	X	
Fatty acid C24:0 as % of total fatty acid weight	X	X	X
Fatty acid C24:0 as % of total fatty acid area		X	X
Fatty acid C24:0 weight (mg/L)		X	
Endocrine/Hormones			
Blood glucose (mg/dL)	X	X	
C-terminal telopeptide of type-1 collagen (ng/mL)	X	X	X
Estradiol via radioimmunoassay (pg/mL)	X	X	
Free thyroxine, fT4 (ng/dL)	X	X	X
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	X		X
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	X	X	X
Plasma insulin via RIA (mIU/L)	X	X	X
Thyroid stimulating hormone, TSH (mIU/L)	X	X	X
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)		X	X
Free testosterone (ng/dL), Vermeulen		X	
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)		X	X
Total testosterone (ng/mL)		X	X

Table IV. Protein and clinical marker features by phenotype model II

	Cognitive Decline	Frailty	Cognitive Frailty
Clinical Features			
Age	X	X	X
Anticholinergic Burden	X	X	X
Depression	X	X	X
Level of Education	X		X
Inflammatory/Immunity			
24-hour urinary cortisol ($\hat{A}\mu\text{g}/24$ hours)	X	X	X
Adiponectin via RIA ($\hat{A}\mu\text{g}/\text{mL}$)	X	X	X
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	X	X	X
Endogenous secretory receptor for AGEs (ng/mL)	X	X	X
Alpha-1 globulin (%)	X	X	X
Alpha-2 globulin (%)	X		
Alpha-2-macroglobulin (mg/dL)	X	X	X
Beta globulins (%)	X		
C-reactive protein - high sensitivity ($\hat{A}\mu\text{g}/\text{mL}$)	X	X	X
C-reactive protein - low sensitivity ($\hat{A}\mu\text{g}/\text{mL}$)	X		
Cortisol:DHEAS ratio (based on nmols)	X	X	
Dehydroepiandrosterone sulfate ($\hat{A}\mu\text{g}/\text{dL}$)	X	X	X
Erythrocyte sedimentation rate (ESR) (mm/hour)	X	X	X
Fibrinogen (mg/dL)	X		X
Homocysteine via FPIA analysis ($\hat{A}\mu\text{mol}/\text{L}$)	X	X	X
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	X	X	X
Interleukin-10 via ELISA (pg/mL)	X	X	
Interleukin-12 via Bio-Plex (pg/mL)	X		X
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	X	X	X
Interleukin-1B via ELISA (pg/mL)	X	X	X
Interleukin-6 via ELISA ultrasensitive (pg/mL)	X	X	X
Interleukin-8 via Bio-Plex (pg/mL)	X	X	X
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	X	X	
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	X	X	X
Resistin via EIA (ng/mL)	X	X	X
Retinol via high performance liquid chromatography ($\hat{A}\mu\text{mol}/\text{L}$)	X		
Serum cortisol ($\hat{A}\mu\text{g}/\text{dL}$)	X	X	X
Soluble CD14 via ELISA (ng/mL)	X	X	X
Soluble IL-6 receptor via ELISA (ng/mL)	X	X	
IL-6 high-sensitivity ELISA calculated from ELISA ultrasensitive (pg/mL)		X	
Soluble TNF-a receptor I via quantitative sandwich EIA (pg/mL)	X	X	X
Soluble TNF-a receptor II via quantitative sandwich EIA (pg/mL)	X	X	X
TNF-related apoptosis-inducing ligand (pg/mL)	X	X	X
Transforming growth factor-B1 (pg/mL)	X	X	
Tumor necrosis factor-a via multiplex technology (pg/mL)	X	X	X
Uric acid (mg/dL)	X	X	X
Urinary cortisol ($\hat{A}\mu\text{g}/\text{mL}$)	X	X	X
Renal/Electrolyte			
24-hour urinary creatinine (mg/24 hours)	X	X	X
Blood urea nitrogen (mg/dL)	X	X	X
Ca ⁺⁺ (mg/dL)	X		
Urinary Ca (mmol/L)	X	X	X
Creatine phosphokinase (U/L)	X	X	X
Creatinine clearance, 24-hr urine (mL/minute)	X	X	X
Cystatin C (mg/L)	X		X
Na ⁺ (mEq/L)	X		
Serum creatinine (mg/dL)	X	X	X
Urinary creatinine (mg/dL)	X		X
Urinary Na (mmol/L)	X	X	X
Urine hemoglobin (mg/dL)	X	X	
Urine proteins (mg/dL)	X	X	X

Nutrient Biomarker	Cognitive Decline	Frailty	Cognitive Frailty
Albumin (%)	X		X
Beta-carotene via high performance liquid chromatography (Åµmol/L)	X	X	X
Lycopene via high performance liquid chromatography (Åµmol/L)	X		X
Omega-3 fatty acids as % of total fatty acid area	X	X	X
Omega-3 fatty acids as % of total fatty acid weight	X	X	
Omega-3 plasma fatty acid weight (mg/L)	X	X	
Omega-6 fatty acids as % of total fatty acid area	X	X	
Omega-6 fatty acids as % of total fatty acid mols	X		
Omega-6 fatty acids as % of total fatty acid weight	X		X
Omega-6 plasma fatty acid weight (mg/L)	X	X	X
Ratio of Omega-6:Omega-3 as % of total fatty acid area	X	X	X
Ratio of Omega-6:Omega-3 as % of total fatty acid mols	X		
Ratio of Omega-6:Omega-3 as % of total fatty acid weight	X	X	
Total proteins (g/dL)	X		
Vitamin B6 via high performance liquid chromatography (ng/mL)	X	X	X
Vitamin E alpha tocopherol, high performance liquid chromatography, (Åµmol/L)	X	X	X
Vitamin E gamma tocopherol, high performance liquid chromatography, (Åµmol/L)	X	X	X
Hematology/Liver			
AST (U/L)	X		
Ferritin (ng/mL)	X	X	
Gamma glutamyl transferase (U/L)	X	X	X
GPT (also known as ALT) (U/L)	X	X	
Hematocrit (%)	X		X
Hemoglobin (g/dL)	X		
Lymphocytes (%)	X	X	X
Lymphocytes (n, K/ÅµL)	X		X
MCH concentration (MCHC) (g/dL)	X	X	X
Mean corpuscular hemoglobin (MCH) (pg)	X		X
Mean corpuscular volume (MCV) (fL)	X	X	
Methylmalonic acid, MMA (Åµmol/L)	X	X	X
Monocytes (%)	X	X	X
Monocytes (n, K/ÅµL)	X	X	X
Neutrophils (%)	X		X
Neutrophils (n, K/ÅµL)	X	X	
Red blood cells (RBC) (n, millions/ÅµL)	X	X	
Red cell distribution width (RDW) (%)	X		
Soluble transferrin receptor (nmol/L)	X		
Vitamin B12 via RIA (pg/mL)	X	X	X
White blood cells (WBC) (n, K/ÅµL)	X	X	X
Folate via RIA (ng/mL)	X	X	X
Lipid Metabolism			
Lipids: HDL cholesterol (mg/dL)	X	X	X
Lipids: LDL cholesterol (mg/dL)	X	X	X
Lipids: total cholesterol (mg/dL)	X	X	X
Lipoprotein(a) (mg/dL)	X	X	X
Metabolomics(plasma lipids)			
Fatty acid C16:0 as % of total fatty acid area	X	X	X
Fatty acid C16:0 as % of total fatty acid weight	X	X	
Fatty acid C16:0 weight (mg/L)	X		X
Fatty acid C16:0 (palmitic) area		X	X
Fatty acid C20:0 (arachidic) area	X		
Fatty acid C20:0 as % of total fatty acid area	X	X	
Fatty acid C20:0 as % of total fatty acid mols	X		
Fatty acid C20:0 as % of total fatty acid weight	X		
Fatty acid C20:0 weight (mg/L)	X	X	
Fatty acid C20:5 n-3 as % of total fatty acid area	X	X	X
Fatty acid C20:5 n-3 as % of total fatty acid weight	X		
Fatty acid C20:5 n-3 cis (eicosapentaenoic, EPA) area	X	X	
Fatty acid C20:5 n-3 weight (mg/L)		X	X
Fatty acid C22:0 (behenic) area	X	X	X
Fatty acid C22:0 as % of total fatty acid area	X	X	
Fatty acid C22:0 as % of total fatty acid weight	X	X	
Fatty acid C22:0 weight (mg/L)	X		X
Fatty acid C24:0 (lignoceric) area	X		
Fatty acid C24:0 as % of total fatty acid weight	X	X	
Fatty acid C24:0 as % of total fatty acid area		X	
Fatty acid C24:0 weight (mg/L)		X	

Endocrine/Hormones	Cognitive Decline	Frailty	Cognitive Frailty
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	X	X	X
Blood glucose (mg/dL)	X	X	X
Urine glucose (mg/dL)			X
C-terminal telopeptide of type-1 collagen (ng/mL)	X	X	X
Estradiol via radioimmunoassay (pg/mL)	X	X	X
Free testosterone (ng/dL), Vermeulen	X	X	X
Total testosterone (ng/mL)	X	X	X
Free thyroxine, fT4 (ng/dL)	X	X	
IGF binding protein-3, serum, immunoradiometric assay (ng/mL)	X		
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***			X
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	X	X	X
Plasma insulin via RIA (mIU/L)	X	X	X
Thyroid stimulating hormone, TSH (mIU/L)	X		X
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	X	X	X

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MANUSCRIPT 4:

Anticholinergic Burden is a Predictor of Cognitive Decline, Physical Frailty and Cognitive Frailty

Abstract:

OBJECTIVES: To investigate whether anticholinergic burden scores are associated with three phenotypes; cognitive decline, physical frailty and cognitive frailty.

DESIGN: Retrospective cohort study.

SETTING: InCHIANTI study, Chianti geographic area of Tuscany, Italy.

PARTICIPANTS: Population of 1,453 adults aged 20-102 years.

MEASUREMENTS: Anticholinergic burden was calculated using the Anticholinergic Cognitive Burden Scale (ACB); neuropsychological tests included the Mini-Mental Status Examination and Trail Making Test A and B (TMT); frailty is defined by the Cardiovascular Heart Study, and cognitive frailty is defined by the International Consensus Group (I.A.N.A/ I.A.G.G). Anticholinergic burden was examined as a predictor for all phenotypes using logistic and ordinal regression models adjusting for covariates.

RESULTS: Anticholinergic burden is associated with cognitive decline, frailty, and cognitive frailty. The odds of having cognitive decline increased by 1.21 points (95% CI = 1.06-1.37, $p < .001$), the odds of being frail increased by 1.33 (95% CI = 1.18-1.50, $p < .001$), and the odds of cognitive frailty increased by 1.36 (95% CI = 1.21-1.54, $p < .001$). Population modeling results indicated the ACB score as one of the stronger predictors for cognitive decline, physical frailty and cognitive frailty with areas under the receiver operating curve of 0.88 and 0.86 respectively. Anticholinergic burden association with cognitive decline as measured by TMT adjusted for covariates was not significant; in

contrast the relationships of ACB with cognitive frailty measured by the TMT-A and TMT-B were statistically significant (both $p < .001$).

CONCLUSION: Our data support a relationship between anticholinergic burden and cognitive decline, further strengthen the association with physical frailty and provide new evidence for an association with cognitive frailty.

Key words: anticholinergic; burden; frailty; cognition; cognitive frailty, xgboost models

INTRODUCTION

The burden of multiple diseases perpetuates the increased consumption of medications.

Older adults are especially susceptible to polypharmacy and medication adverse risks due to declines in physiological reserve, reduced liver and kidney function required to metabolize medications and increased central nervous system sensitivity to medications¹. A decline in physiologic reserve coupled with the use of anticholinergic medicines increases the risk for impaired functional and cognitive performance²⁻⁵.

Anticholinergic medications block the neurotransmitter acetylcholine in the central and peripheral nervous system, selectively blocking acetylcholine from binding to the muscarinic receptors in the brain^{6,7}. Additionally, there is growing evidence that anticholinergic affect older adults in greater proportion due to the ability of these medications to permeate the blood-brain barrier^{2,8}. Anticholinergic burden is considered to be the cumulative effect on an individual taking one or more medications with anticholinergic activity confounded by age-related pharmacokinetic and pharmacodynamic changes^{1,5,6}. Higher anticholinergic burden can occur with specific medications known to have high anticholinergic activity or with an accumulation of

medications with low, medium, and high anticholinergic burden^{9,10}. An increase in circulating anticholinergic activity causes inhibition of acetylcholine transmission to the central nervous system suggesting a cholinergic deficit that is hypothesized to be involved in causing impaired cognitive and motor function¹¹. There are substantial differences in methods for measuring anticholinergic burden and no standard or consensus on how to quantify burden. Systematic reviews on the current anticholinergic burden scales have all shown an association between higher anticholinergic burden and adverse outcomes; cohort studies have mainly focused on cognitive and physical outcomes^{5,9}.

Less understood is the effect anticholinergic burden has on physical frailty⁵. Although there is evidence to support the relationship between physical function and higher anticholinergic burden, the methods for measuring physical functioning have focused on activities of daily living (ADLs) and instrumental activities of daily living (IADLs) without controlling for confounding health factors contributing to the outcome^{5,9}. Changes in ADLs and IDLs can be affected by multiple psychosocial and physiological factors that are not a direct measure of disease. A recent study found a significant association of anticholinergic burden with gait and impaired balance measured by the timed-up and go(TUG), functional reach(FR), and grip strength(GS) assessments¹². Frailty as defined by the Cardiovascular Heart Study (CHS) is a disease process and a non-normal process of aging¹³. The CHS frailty phenotype includes decline in lean body mass, strength, endurance, balance, walking performance, and low activity¹³. Additionally, there is growing evidence for a shared relationship between cognitive decline and physical

frailty¹⁴⁻¹⁶. The International Consensus Group organized by the International Academy on Nutrition and Aging (I.A.N.A) and the International Association of Gerontology and Geriatrics (I.A.G.G) which convened in 2013 to identify related domains of physical frailty and cognition, termed this relationship as “cognitive frailty”¹⁵.

Studies thus far have primarily used the Mini-Mental State Examination (MMSE) to measure cognitive decline which as a composite test does not capture distinct areas of cognitive function such as processing speed, attention, psychomotor speed, abstraction, flexibility, ability to execute and modify a plan of action¹⁷. The goal of this study was to use logistic and ordinal regression models to determine the relationship between anticholinergic burden and three phenotypes: cognitive decline defined by the MMSE and Trail Making Tests, part A and B, physical frailty, and cognitive frailty. Additionally, we included anticholinergic burden in a separate population based predictive model study to determine if anticholinergic burden is predictive of cognitive decline, frailty, and cognitive frailty. The population predictive model incorporates additional measures of disease such as protein and genomic biomarkers thereby evaluating ACB with confounding disease processes (Sargent et al., 2018 in preparation).

METHODS

Data

The subjects in the present study were participants in *Invecchiare in Chianti* (Aging in Chianti, “InCHIANTI Study”). InCHIANTI was a prospective population based study of 1,453 adults aged 20-102 randomly selected from two towns in Tuscany, Italy using a multistage stratified sampling at baseline from 1998 to 2000¹⁸. All aspects of the

InCHIANTI research were approved by the ethics committees at the institutions responsible for data collection, and this secondary study was approved by the ethics committee at *Centre de recherché Clinique du CHUS*, project #547. During the initial InCHIANTI baseline 90-minute interview, information was collected on demographic and clinical characteristics for the three phenotypes and baseline medications taken regularly in the prior 15 days to determine anticholinergic burden. The name of the drug, preparation and dosage were collected from medication boxes or bottles including over the counter vitamins, food supplements, sleeping pills, or laxatives. Initial medication information was converted from the brand name to the active ingredient.

Measures

For the current study, a total of 2,883 baseline medications were used to analyze the anticholinergic burden effect on 1,155 individuals ≥ 65 years of age with cognitive decline, physical frailty, and cognitive frailty. Currently, there are 7 expert-based anticholinergic rating scales for which quantification of the tool is based on expert opinion, and published data, and includes both genders with a mean age of 65 years or older^{4,9}. The Anticholinergic Cognitive Burden (ACB) scale is the most validated scale for evaluating adverse health outcomes including cognitive and physical function^{4,10}. The anticholinergic properties of each medication were quantified using the ACB scale based on each drug's serum anticholinergic activity¹⁹. To determine ACB scores, each participants' medications were assigned points (0, 1, 2, 3) according to the published 2012 update and summed for a total anticholinergic burden score. Higher scores indicate higher anticholinergic properties. An example of medications with ACB scores

include: Amitriptyline = 3, Amantadine = 2, and Atenolol = 1. The ACB scale has identified medications with anticholinergic properties that have correlated with a 0.33-point decline in the MMSE score over 2 years²⁰. The neuropsychological tests included the MMSE as a test of general cognition and Trail Making Test, part A and B (TMT). The TMT testing was included to further explore distinct areas of cognitive function. TMT-A is used to assess psychomotor speed; scoring is based on time in seconds to completion with a score range of 0 to 300 seconds²¹. TMT-B is used to assess the executive functioning domain (any individual time over the limit of 300-600 seconds was included as 300)²¹. Normative data for time to complete the TMT tests in seconds is stratified by age and education²². Additionally, the neuropsychological profile for individuals with cognitive frailty is considered to be different from those with frailty or cognitive decline alone with recent findings of lower performance on TMT tests^{22,23}. The Center for Epidemiologic Studies Depression Scale (CES-D) self-report scale was used to measure depressive symptoms. The CES-D has been used extensively in epidemiologic studies for depression and physical function displaying similar reliability, validity, and factor structure across a diverse demographic²⁴.

Frailty measures included the number of frailty symptoms with performance test data. Frailty as defined by the cardiovascular health study (CHS), allows for a continuous scoring system versus a nominal system because it can capture the multidimensional nature of frailty¹⁴. The components have concurrent and predictive validity with hazard ratios (HR) ranging from 1.82-4.46 ($p < 0.05$) for outcomes that include incident disease, hospitalization, falls, disability and mortality in community-dwelling older adults¹³. The

InCHIANTI criteria for frailty defined unintentional weight loss as losing weight not related to diet, classified the values of body mass index, strength, walking speed and height based on all subjects ≥ 65 years and used two questions of the CES-D for the definition of exhaustion.

Phenotypic Classification

The MMSE score and the TMT part A and B was used to define two phenotypic classifications for cognitive decline and cognitive frailty. All participants completed the MMSE to define cognitive decline and cognitive frailty. Absence of cognitive decline is defined as a score of 24-30 on the education adjusted MMSE²⁵⁻²⁷. Frailty is characterized by individuals with one or more of the Frailty criteria¹³. Cognitive frailty is defined as individuals with cognitive decline and one or more of the frailty criteria²³.

- Robust with no physical frailty and absence of cognitive decline
- Robust with no physical frailty with cognitive decline (MMSE = ≤ 23)
- Frail (≥ 1 criterion) and absence of cognitive decline
- Frail (≥ 1 criterion) and cognitive decline (MMSE = ≤ 23)

Additional phenotypic classification included mild, moderate, or severe disease defined by the MMSE to characterize 24-30 as normal cognition, a score of 23-18 as moderate cognitive decline (combined mild and moderate degree of impairment), and a score ≤ 17 as cognitive impairment^{25,26}. Frailty is characterized by the CHS criteria cut offs and cognitive frailty is defined as individuals with both criteria¹³.

- Robust with no physical frailty and absence of cognitive decline
- Robust with no physical frailty with mild cognitive decline (MMSE = 18-23)
- Robust with no physical frailty with cognitive impairment (MMSE = ≤ 17)
- Pre-frail (1-2 criteria) and absence of cognitive decline
- Frail (≥ 3 criteria) and absence of cognitive decline

- Pre-frail (1-2 criteria) and with mild cognitive decline (MMSE = 18-23)
- Frail (≥ 3 criteria) and with mild cognitive decline (MMSE = 18-23)
- Pre-frail (1-2 criteria) and cognitive impairment (MMSE = ≤ 17)
- Frail (≥ 3 criteria) and cognitive impairment (MMSE = ≤ 17)

Additional neuropsychological testing (TMT-A and B) was used to define cognitive decline and as part of the definition of cognitive frailty²³. TMT-A and B cut off scores for cognitive decline are based on cut off norms established by Ashendorf et al., 2008.

- Robust with no physical frailty and absence of cognitive decline
- Robust with no physical frailty with Cognitive Decline (both Trail A ≥ 78 and Trail B ≥ 106)
- Frail (≥ 1 criterion) and Cognitive Decline (both Trail A ≥ 78 and Trail B ≥ 106)
- Frail (≥ 1 criterion) and Cognitive Decline (both Trail A ≥ 78 and Trail B ≥ 106)

Numbers of participants were insufficient for statistical analysis to include cognitive decline or cognitive frailty categorized into levels of mild, moderate, and severe phenotype with the TMT.

Statistical Analyses

We used logistic and ordinal regression to investigate the relationship between anticholinergic burden and all three outcomes. Covariates were selected to control for potential confounding effects. Demographic covariates included gender, age, and level of education. Disease processes considered as confounders included baseline diagnosis of: baseline dementia (n=82), vascular dementia (n=41), depression (n=412), and Parkinson's disease (n=16) and were included in the models as binary covariates.

In addition to the logistic and ordinal regression, ACB score was included in separate population based predictive model analyses with 298 additional predictors; these

included protein, clinical, and genetic markers of disease. Modeling of the dynamic interactions between confounding disease processes determined the strength of the relationship and predictive value for anticholinergic burden and disease outcome. Predictive modeling via ensemble learning using xgboost allowed for better accuracy by building multiple models, each of which learns to improve upon the errors of a prior model producing a final model that reflects the complex interactions between biological processes (i.e., protein and genetic biomarkers) on cognitive decline and frailty. Parameters for the xgboost model included a stepsize eta of = "0.3", rounds = 5-200, max depth = "10", nthread = "12", objective = "binary:logistic", evaluation metric = "auc", gamma = default = "0" to control the number of trees and prevent overfitting²⁸. Details on the population predictive model results and statistical methods beginning with model development in the InCHIANTI dataset used to train and test classifiers, complete internal validation, and calibration of the model are available in a separate publication (Sargent et al., 2018 in preparation). Bivariate analyses included non-parametric Kruskal-Wallis t-tests to assess differences between groups; medians and maximum quantiles are reported for healthy controls and three phenotypes. Next, Bonferroni correction was conducted to adjusted for multiple comparisons; adjusted p-values are reported. All statistical analyses were carried out using R V. 3.2.1.. R packages included 'glm2'-Fitting Generalized Linear Models, 'Ordinal'-Regression Models for Ordinal Data, and 'xgboost'-Extreme Gradient Boosting²⁸⁻³⁰.

RESULTS

Medication data was complete for 1,155 participants; table 1 describes the characteristics of the participants by phenotype and the percent of individuals with a total daily ACB score, which ranged from 0-9. Distribution of anticholinergic burden score by phenotype and differences between health control and phenotype are shown in Table 2. Tables displaying results for the top predictive features from the xgboost predictive modeling study are published elsewhere (Sargent et al., 2018 in preparation)

There was a significant association between anticholinergic burden and cognitive decline ($p = 0.02$), frailty ($p < .001$) and cognitive frailty ($p < .001$). Additionally, the odds of having cognitive decline increased by 1.21 points (95% CI = 1.06-1.37, $p < .001$), the odds of being frail increased by 1.33 (95% CI = 1.18-1.50, $p < .001$), and odds of cognitive frailty increased by 1.36 (95% CI = 1.21-1.54, $p < .001$). Model fit for all three phenotypes using the Wald chi-square test statistic was associated with a p-value of $< .001$, indicating that the overall effect rank was significant. Logistic and ordinal regression results are presented in Table 3 and 4. Results from the population predictive model are ranked by gain, which is a metric based on each feature's contribution in the model. When comparing top features to other features in the model, the greater the gain the more important the feature is for prediction of the outcome. Anticholinergic burden was the top 4% predictor out of 105, 14% of 101, and 70% of 93 selected features during the classifier build, with AUCs ranging from 0.81-0.88 for the outcomes frailty, cognitive frailty, and cognitive decline respectively measured with the MMSE (Sargent et al., 2018 in preparation).

Similarly, there was a significant association found between ACB score and cognitive decline when measured with the TMT-A and TMT-B without adjusting for covariates. When including the covariates age, gender, and baseline dementia individually in the models with only ACB score for TMT-B or age and gender for TMT-A, anticholinergic burden was no longer significant. Additionally, this was true when covariate-by-ACB interaction terms were included; none of the interaction terms was statistically significant (all $p > 0.2$). There was a significant association found between ACB score and cognitive frailty, as measured with TMT-A ($p = 0.007$) and TMT-B ($p < .001$). Model fit for cognitive frailty TMT-A and TMT-B using the Wald chi-square test statistic was associated with a p-value of $< .001$. Logistic regression results for cognitive decline and cognitive frailty measured with TMT are shown in Table 3. In the population predictive modeling results, anticholinergic burden was the top 32% of 149 and 40% of 110 predictors, with AUCs ranging from 0.86-0.83 for the outcomes cognitive decline and cognitive frailty respectively measured with the TMT-A and B (Sargent et al., 2018 in preparation).

DISCUSSION

Participants for all phenotypes were older with a greater proportion of females; few completed a high school education. Participants with cognitive decline, frailty, and cognitive frailty took more medications than individuals without these phenotypes. There were smaller numbers of participants with an ACB score > 4 with most scores above zero clustered between 1-4; suggesting that an ACB score of 1-4 range is sufficient to show association.

Logistic and ordinal regression results found in this study continue to support a relationship between anticholinergic burden and cognitive decline, further strengthen the association with physical frailty, and provide new evidence for an association with cognitive frailty. The population predictive model results with xgboost, showed anticholinergic burden to be a significant predictor for all three phenotypes (Sargent et al., 2018 in preparation).

Although frailty and cognitive decline have been shown to be related, both diseases have long been studied separately. The findings from this study provide the first evidence for a relationship between anticholinergic burden and cognitive frailty, affecting both cognitive speed and executive functioning. The study results show a relationship between anticholinergic burden and cognitive decline when measured with the MMSE but no relationship was observed when cognitive decline was measured with the TMT-A and TMT-B unless cognitive frailty was present. Another study found lower executive function composite scores on the Wechsler Memory Scale-Revised, Logical Memory Immediate Recall, and TMT-B test in a small sample (n=402) of individuals taking anticholinergic medications over 1 year with additional findings of increased brain atrophy and clinical decline³¹. Additionally, previous studies have shown a relationship between anticholinergic burden and transitions between frailty states and increased mortality for individuals who were robust at baseline; with every unit increase in burden being associated with a 73% risk of transition from robust to pre-frail. Further these studies showed that anticholinergic burden is associated with poor mobility, functional decline, psychomotor slowing, and falls^{5,12,32}.

A limitation of the study is that this was a secondary analysis of existing data. As such, the medications are from an international database and represent a specific population of individuals and do not consider potential differences in prescribing patterns throughout the world. Additionally, confounding may be a factor; for which it becomes difficult to distinguish between the effects of the medications and the disease process. Therefore, further research with adequately powered randomized controlled trials or prospective cohort studies with follow up periods in the clinical setting are needed to distinguish medication effect from disease progression. These findings highlight the need for longitudinal studies focused on understanding which domains of memory are affected.

Future research should focus on methods for detecting high risk individuals in the clinical setting, the relationship between Apolipoprotein E ϵ 4 and anticholinergic medications, and whether anticholinergic medications are a modifiable risk factor for the prevention of cognitive decline and physical frailty. Identification of reversible causes for cognitive and physical impairment is critical for the aging population.

Clinicians need to be aware of these findings and review cumulative anticholinergic burden in robust and vulnerable individuals and minimize the overall anticholinergic burden before symptoms of cognitive and physical decline are detectable. Until a better understanding of the implications that these findings have in the clinical setting, caution must be applied since medications with anticholinergic effects are used to treat many chronic diseases, such as congestive heart failure and hypertension. These findings

encourage new research and may lead to effective interventions for the prevention and treatment of cognitive and physical decline in an aging population.

CONCLUSION

Anticholinergic burden is associated with both cognitive decline and physical frailty.

Efforts to better understand the epigenetic effects, sum dose effect, and identify

individuals in clinical settings who may require anticholinergic medication

discontinuation are important next steps to prevent anticholinergic burden induced

outcomes.

Table 1. Characteristics of participants by phenotype

	Cognitive Decline (MMSE)	Frailty (CHS)	Cognitive Frailty (MMSE)	Cognitive Decline (TMT-A)	Cognitive Decline (TMT-B)	Cognitive Frailty (TMT-A)	Cognitive Frailty (TMT-B)
Phenotype (n)	(n=369)	(n=595)	(n=257)	(n=525)	(n=634)	(n=302)	(n=325)
Age, mean(SD)	80 (8.7)	78 (7.9)	82 (7.4)	76 (7.7)	72 (9.0)	78 (7.4)	76 (6.9)
Gender, %							
Male (n)	24.0 (120)	42.8 (214)	31.9 (82)	37.1 (195)	41.9 (266)	35.1 (106)	36.0 (117)
Female (n)	37.6 (249)	58.2 (381)	68.1 (175)	62.9 (330)	58.0 (368)	64.9 (196)	64.0 (208)
Education, %							
No Education	56.9 (210)	39.3 (234)	58.8 (151)	42.3 (222)	25.4 (161)	46.4 (140)	30.8 (100)
Elementary - Secondary	39.6 (146)	52.4 (312)	37.7 (97)	53.1 (279)	66.2 (420)	49.3 (149)	61.5 (200)
≥ High School	1.4 (5)	7.1 (42)	1.9 (5)	3.2 (17)	7.6 (48)	3.3 (10)	7.4 (24)
Medication use							
Number of drugs							
0 meds	73	83	34	107	141	35	51
1 to 4	228	305	169	334	408	201	208
5 to 7	56	100	45	70	73	53	56
≥ 8	12	23	9	14	12	13	10
mean(SD)							
Control	2.18 (2.01)	1.75 (1.76)	2.15 (2.02)	1.95 (1.87)	1.77 (1.73)	1.85 (1.82)	1.68 (1.66)
Phenotype	2.69 (2.19)	2.89 (2.21)	3.00 (2.16)	2.44 (2.12)	2.23 (2.02)	3.01 (2.20)	2.79 (2.19)
p-value*	<.001	<.001	<.001	<.001	<.006	<.001	<.001

Notes: SD = standard deviation, * two tailed t-Test with means and SD

Table 2. Distribution of anticholinergic burden score by phenotype and difference between health control and phenotype

% (n)	Cognitive Decline	Frailty	Cognitive Frailty	Cognitive Decline		Cognitive Frailty	
	MMSE (n=296)	CHS (n=512)	MMSE (223)	Trail A (n=418)	Trail B (n=493)	Trail A (n=267)	Trail B (n=274)
ACB							
0	47.0% (139)	51.0% (261)	42.2% (94)	57.9% (242)	62.9%(310)	50.2% (134)	55.5% (152)
1	23.6% (70)	22.9% (117)	25.1% (56)	20.6% (86)	20.1% (99)	22.5% (60)	21.2% (58)
2	14.5% (43)	11.9% (61)	16.1% (36)	10.8% (45)	7.9% (39)	13.1% (35)	9.9% (27)
3	10.1% (30)	8.8% (45)	11.2% (25)	6.7% (28)	5.5% (27)	8.2% (22)	7.7% (21)
4	2.7% (8)	3% (16)	3.1% (7)	2.4% (10)	2.4% (12)	3.4% (9)	3.6% (10)
5	1.0% (3)	1.4% (7)	.9% (2)	1.0% (4)	1.0% (5)	1.5% (4)	1.8% (5)
6	.7% (2)	.8% (4)	.9% (2)	.5% (2)	.2% (1)	.7% (2)	.4% (1)
9	.3% (1)	.2% (1)	.4% (1)	.2% (1)	(0)	.4% (1)	(0)
Control	0[6]	0[5]	0[6]	0[5]	0[4]	0[5]	0[4]
Phenotype	1[9]	0[9]	1[9]	0[9]	0[6]	0[9]	0[6]
p-value*	<.001	<.001	<.001	<.001	.042	<.001	<.001

Table 3. Generalized linear regression results: association between anticholinergic burden and phenotypes

Phenotype	(n)	Beta Coef	Std. Error	95%CI	p-value
Cognitive Decline (MMSE)	375	0.21	0.07	0.08-0.36	.004
Frailty (CHS)	595	0.31	0.07	0.17-45	<.001
Cognitive Frailty (MMSE)	257	0.26	0.08	0.11-0.41	<.001
Cognitive Decline (Trail A)	545	0.20	0.14	0.14-0.11	.14
Cognitive Decline (Trail B)	703	0.21	0.14	0.10-.47	.12
Cognitive Frailty (Trail A)	302	0.27	0.08	0.11-.43	<.001
Cognitive Frailty (Trail B)	325	0.38	0.09	0.19-0.57	<.001

Table 4. Ordinal regression results: association between anticholinergic burden and phenotype

Models	Phenotypes (MMSE & CHS)	n
1	Cognition	
	Cognitive Decline	501
	Cognitive Impairment	101
2	Frailty	
	Frail	88
	Pre-frail	507
3	Cognitive Frailty	
	Cognitive Decline & Frail	55
	Cognitive Decline & Pre-frail	217
	Cognitive Impaired & Frail	11
	Cognitive Impaired & Pre-frail	76

Models	Phenotype	Beta Coef	Std. Error	Odds Ratio	95%CI	p-value
1	Cognition	0.19	0.07	1.21	1.07-1.37	<.001
2	Frailty	0.29	0.06	1.33	1.87-1.50	<.001
3	Cognitive Frailty	0.31	0.06	1.36	1.21-1.54	<.001

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SUMMARY

This dissertation consists of four manuscripts; 1) an integrative review of the measurements for cognitive frailty, 2) a systematic review of the clinical and biological markers for cognitive decline and physical frailty, 3) an innovative population predictive model analyses establishing biological plausibility for cognitive frailty, 4) and a new finding of anticholinergic burden as a predictor of frailty and cognitive frailty. The results from this study establish a foundation for an understanding of the underlying biological mechanisms for the relationship between cognitive decline and physical frailty and found anticholinergic burden as one of the top predictors for frailty and cognitive frailty. In seeking to explore the importance and applicability of these results it is critical that others continue to replicate the model results. To accompany manuscript 3, help with replication and extension of this work, the code has been made publically available for the population predictive model.

Implications

The results from this dissertation have several implications for future research and have a potential for translation into practice. Through the lens of Complex Systems Theory, this dissertation begins to unravel the complexity behind a geriatric syndrome providing biological plausibility to cognitive frailty. Geriatric syndromes such as cognitive frailty are highly multifactorial and variable across the aging spectrum lending themselves to new ways of investigation. As Bryne (1998) notes: Not only can the complex not always be derived, even in principle from the less complex,... we can often

only understand the simpler [cognitive frailty] in terms of its origins in the more complex (p. 16). By using the framework of complex systems theory and an innovative Boosted trees machine learning technique (xgboost) we determined key biological mechanism for a dysregulation across multiple systems as the potential cause for cognitive frailty. The future to understanding complex geriatric syndrome should include a systems approach by using highly accurate statistical modeling to identify measurable markers. There were multiple biological associations determined by the study results that should be investigated further. One of the interesting findings is anticholinergic burden in conjunction with the association of *SLCO1B1* as predictors for cognitive frailty. *SLO1B1* is an important pharmacokinetic gene that is involved in the removal of drug compounds and transport of drug metabolites at the blood-brain barrier(1). It has been implicated as a marker of lean muscle mass loss and may affect the distribution of drugs into the central nervous system(1,2).

Limitations

The limitations of the dissertation research included the use of a small homogenous sample with large numbers of biomarkers creating limitations for translation into clinical research. Additionally, the study was retrospective using existing data. The analyses used a randomly assigned training subset to validate the model within a relatively homogenous InCHIANTI cohort. Additionally, no external validation of the model was completed. The model would be strengthened by external validation in a in a mixed ethnic and demographic age range. Through the process of completing this dissertation I have gained invaluable expertise in statically modeling of a large dataset and have

learned skills in the field of bioinformatics. The dissertation required me to learn bash and R coding, along with learning how to manipulate genetic data in PLINK.

Future research

There are several areas for future research based on this dissertation work. There is a need to test and validate the model in a second more ethnically diverse population before translation into clinical practice. Further investigate anticholinergic burden as an epigenetic cause of cognitive frailty by exploring the relationship between putative genetic markers discovered in the model analyses (i.e. *SLCO1B1* and *COMT*). Some of these findings can be translated into clinical studies. Research focusing on methods for detecting high-risk individuals in the clinical setting and descriptive studies to understand the scope and effect of cognitive frailty are needed. Intervention studies are essential to understanding the role of nutrition and/or physical activities have on neuroinflammatory cytokines and other system markers for cognitive frail individual's progression. Additionally, further work can be done on whether anticholinergic medications are a modifiable risk factor for the prevention of cognitive frailty. Identification of reversible causes for cognitive and physical impairment is critical for the aging population.

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APPENDICES

MANUSCRIPT 1: Supplemental documents

Appendix A. Search Strategies – Conducted January 30, 2015

Database	Added Filters	Time Period	Terms	Results
PubMed	English, Human	1983-Present	(((((("Frailty"[TIAB] OR "Frail"[TIAB] OR "Physical Frailty"[TIAB] OR "Cognitive Frailty"[TIAB])) OR "Frail Elderly"[Mesh])) AND ((Alzheimer*[TIAB] OR Presenile Dementia*[TIAB] OR Senile Dementia*[TIAB] OR Mild Cognitive Impairment*[TIAB] OR Mild Neurocognitive Disorder*[TIAB] OR Mild Neurocognitive Disorder*[TIAB] OR Early Dementia*[TIAB] OR Early Onset Dementia*[TIAB] OR Cognitive Decline[TIAB] OR Mild Cognitive Impairment*[TIAB]))	322
CINAHL	English, Human	1992-Present	(MH "Alzheimer's Disease") OR ("Alzheimer*" OR "Presenile Dementia*" OR "Senile Dementia*" OR "Mild Cognitive Impairment*" OR "Mild Neurocognitive Disorder*" OR "Mild Neurocognitive Disorder*" OR "Early Dementia*" OR "Early Onset Dementia*" OR "Cognitive Decline" OR "Mild Cognitive Impairment*") AND (MH "Frailty Syndrome") AND "Frailty" OR "Frail" OR "Physical Frailty" OR "Cognitive Frailty"	76
PsycInfo	None	2005-Present	(Title:(("Frailty" OR "Frail" OR "Physical Frailty" OR "Cognitive Frailty") OR Abstract: ("Frailty" OR "Frail" OR "Physical Frailty" OR "Cognitive Frailty")) AND ((Index Terms: ("Cognitive Impairment")) OR Title: ("Alzheimer*" OR "Presenile Dementia*" OR "Senile Dementia*" OR "Mild Cognitive Impairment*" OR "Mild Neurocognitive Disorder*" OR "Mild Neurocognitive Disorder*" OR "Early Dementia*" OR "Early Onset Dementia*" OR "Cognitive Decline" OR "Mild Cognitive Impairment*") OR Abstract: ("Alzheimer*" OR "Presenile Dementia*" OR "Senile Dementia*" OR "Mild Cognitive Impairment*" OR "Mild Neurocognitive Disorder*" OR "Mild Neurocognitive Disorder*" OR "Early Dementia*" OR "Early Onset Dementia*" OR "Cognitive Decline" OR "Mild Cognitive Impairment*"))	164
Dissertation & Thesis	None	1984-Present	All ("Alzheimer*" OR "Presenile Dementia*" OR "Senile Dementia*" OR "Mild Cognitive Impairment*" OR "Mild Neurocognitive Disorder*" OR "Mild Neurocognitive Disorder*" OR "Early Dementia*" OR "Early Onset Dementia*" OR "Cognitive Decline" OR "Mild Cognitive Impairment*") AND all ("Frailty" OR "Frail" OR "Physical Frailty" OR "Cognitive Frailty")	18
Web of Science	English	1991-Present	(("Alzheimer*" OR "Presenile Dementia*" OR "Senile Dementia*" OR "Mild Cognitive Impairment*" OR "Mild Neurocognitive Disorder*" OR "Mild Neurocognitive Disorder*" OR "Early Dementia*" OR "Early Onset Dementia*" OR "Cognitive Decline" OR "Mild Cognitive Impairment*")) AND TOPIC: (Frailty OR Frail OR Physical Frailty OR Cognitive Frailty)	560

Table 3. Data Extraction and Measurement Properties

Author & Title	Theoretical Framework	Population Assessed	Frailty Assessment Instruments	Cognitive Assessment Instruments	Reliability	Validity	Feasibility	Principal Results	Level of Evidence*
Shimada et al. 2013 Combined Prevalence of Frailty and Mild Cognitive Impairment (MCI) in a Population of Elderly Japanese People	Indices of cognitive frailty were discussed	Country: Japan N= 5104 56 and older non-demented persons enrolled in Obu Study of Health Promotion for the Elderly (OSHPE)-community dwelling Exclusion criteria: history of Parkinson disease, stroke, or MMSE <18	CHS criteria: frailty phenotype defined by 3 or more of the 5 domains: Mobility: timed walk of 2.4 meter (3.2 feet) (cut off <1.0 m/s) Strength: Grip strength dynamometer (cut off male: <26 kg, female: <17kg) Physical activity: Self-report (no tool listed) Endurance: self-report & included questions from GDS* Nutrition: self-reported weight loss in previous 2 years	MCI criteria: Subjective memory complaint, cognitive impairment, no functional dependency and no clinical criteria for dementia MMSE (cut off <23 impaired) (Folstein, Folstein, and McHugh 1975) National Center for Geriatrics and Gerontology-Functional Assessment Tool (NCGG-FAT)	§ NCGG-FAT – test-retest reliability (ICC = 0.764 to 0.942)	Frailty and MCI Odds Ratio (OR) (2.0, 95% CI 1.5-2.5 p <0.01) Reported values from original study: § NCGG-FAT – External validity (Pearson r = 0.496 to 0.842)	Time intensive to measure both domains MMSE- 11 questions; 5-10 minutes to perform NCGG-FAT-effective for assessing multidimensional cognitive screening; easily administered using tablet technology; instructions on display, training to use tool is limited; knowledge of neuropsychiatric measures not extensive, with a battery of neuropsychiatric test completed in 20-30 min	Frailty is strongly associated with cognitive impairment Additional Findings: Increasing age and Frailty p for trend <0.01, MCI p < 0.05 Education associated with frailty p <0.01, MCI p <0.01 Frailty higher in women than men p <0.05, MCI no differences for gender	1b Cross-sectional study

<p>Kulmala et al. 2014 Association between Frailty and Dementia: A Population-Based Study</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: Finland N=781</p> <p>76-100 years, mean age 82 non-demented community dwelling</p> <p>Population based sample from the Geriatric Multidisciplinary Strategy for the Good Care of the Elderly (GeMS)</p>	<p>CHS criteria: frailty phenotype defined by 3 or more of the 5 domains:</p> <p>Slowness- timed maximal 10-meter (32.8 feet) walking test (no cut off mentioned)</p> <p>Weakness-grip strength dynamometer (highest of 2 measurements used)</p> <p>Low physical activity- Grimby scale</p> <p>Poor endurance and energy- self-report question</p> <p>Shrinking/sarcopenia- weight loss >5% over previous year</p>	<p>MMSE (cut off <25 impaired) (Folstein, Folstein, and McHugh 1975)</p>	<p>Not reported</p>	<p>Age & gender-adjust models support these findings</p> <p>Frail, pre-frail, & robust associated percentages with clinically diagnosed dementia: (52%, 19% and 11%, p < 0.01); vascular dementia (9%, 3%, and 1% p = 0.001); and Alzheimer's (30, 15, and 9%, p <0.001)</p> <p>Frailty & cognitive impairment (OR 7.4, 95% CI 4.2-13.2)</p> <p>Frailty & clinically diagnosed dementia (OR 6.5%, 95% CI 3.6-11.8)</p> <p>Frailty & vascular dementia (OR 6.7, 95% CI 1.6-27.4)</p> <p>Frailty & Alzheimer's (OR 3.2, 95% CI 1.7-6.2)</p>	<p>Time intensive to measure both domains, clinical diagnosis, and imaging can be expensive</p> <p>Clinical translation properties for detection of cognitive frailty unclear</p>	<p>Frailty is associated with cognitive impairment</p> <p>Frail individuals were 7.4 times more likely to have cognitive impairment, 6.5 times more likely to have clinically diagnosed dementia; 6.7 times more likely to have vascular dementia, and over 3.2 times more likely to have Alzheimer's disease than those who were robust</p>	<p>1a</p> <p>Cross-sectional study</p>
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<p>Buchman et al. 2014 Brain pathology contributes to simultaneous change in physical frailty and cognition in old age</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: U.S. N=2167 Religious Order Study (ROS) and Memory and Aging Project</p>	<p>CHS criteria: Physical frailty Grip strength: dynamometer Gait: time to walk 8 feet Body composition was based on body mass index (BMI). Fatigue: two questions derived from a modified version of the Center for Epidemiologic Studies–Depression Scale</p>	<p>19 cognitive tests scored and reviewed by neuropsychologist (Wilson et al. 2002) Five cognitive domains: episodic memory, semantic memory, working memory, perceptual speed, and visuospatial ability</p>	<p>Not reported</p>	<p>Slope measures for physical frailty and cognition (N = 1,794, 82.8%) Frailty and cognition controlling for number of chronic health conditions (r= -0.708, p<0.001); demographic variables/race (r= -0.68, p < 0.001) Gait and cognition (r= -0.67, p<0.001) Grip strength and cognition (r= - 0.51, p<0.001) BMI and cognition (r = -0.17 p=0.003) Association of brain pathologies with rates of change of frailty and cognition (r = -0.708, p <0.001)</p>	<p>Time and resource intensive Clinical translation of the cognitive frailty construct are unclear</p>	<p>Strong linear relationship between rates of change in frailty and cognition Relationship between frailty and cognition remained when controlling for demographic variables and race and number of chronic disease Strongest correlation between gait speed and cognition Presence of macroinfarcts, AD pathology, and nigral neuronal loss were each with rapid progression frailty and cognitive decline</p>	<p>1b Population-based, longitudinal study (10 years)</p>
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<p>Rolfson et al. 2013 An assessment of neurocognitive speed in relation to frailty</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: Canada N=164 Mean age 74</p> <p>Baseline cohort of community based older adults; Non-demented population from the Oxford Project to Investigate Memory and Aging (OPTIMA)</p>	<p>Modified CHS criteria: Weight loss: "Have you lost a lot of weight in the last six months?" ("some change" or "considerable change")</p> <p>Subjective exhaustion: "Do you find you have recently lost energy and it is harder to get things done?"</p> <p>Physical activity: Immobility as defined by either (a) informant history (Does he or she have trouble getting about...? "some difficulty" or "great difficulty") or (b) physical examination suggesting the need for a mobility aid or another person.</p> <p>Slow walking speed: Physical Examination evidence of slow ambulation</p> <p>Weakness: Physical Examination evidence of suboptimal arm or leg power (Grade 4/5 or less) Frailty Index (FI): 70/83 items used</p> <p>Modified Edmonton Frail Scale (EFS): 5 items used in analysis (number of medications, depression, weight loss, urinary incontinence and clock drawing test)</p>	<p>MMSE (cut off not mentioned)</p> <p>Neurocognitive speed (NCS) cut off <18 pattern</p> <p>Comparison test (PCT) <11 Letter comparison test (LCT) <7</p>	<p>Not reported</p>	<p>NCS (DV) and MMSE, FI, EFS (IVs): (OR 1.19, 95% CI 1.04-1.36, p = 0.012); OR 0.87, 95% CI 0.81-0.95, p=0.001), OR: 0.94, 95% CI 0.59-1.49, p=0.779)</p> <p>Modified EFS and NCS: (OR 0.94, 95% CI 0.59-1.49, p= 0.779)</p>	<p>Time and resource intensive</p> <p>Clinical translation properties unclear</p>	<p>Strong correlation between NCS and frailty.</p> <p>Association was evident with FI and NCS</p> <p>NCS was not associated with a modified CHS or modified EFS.</p> <p>Modified CHS was significant when the MMSE was taken out</p>	<p>1b</p> <p>Population-based, longitudinal study (3 years)</p>
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<p>Oosterveld et al. 2014 The influence of co-morbidity and frailty on the clinical manifestation of patients with Alzheimer's disease</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: Netherlands N=213 Clinical Course of Cognition and Comorbidity-Dementia Study (4C-Dementia study) 46-93 years old; mean 75 with probable (n=193) or possible (n=20) diagnosis of Alzheimer's Disease</p>	<p>Modified CHS: Scoring range from 0-5; 3 or higher = frail; 2 = pre-frail Measurement details were not listed; based on definition of Fried: Weight loss Activity level Emotion/energy level Grip strength Gait velocity: 15 feet walk test</p>	<p>Baseline measure: MMSE score ≥ 10, CDR^B score 0.5-2 (0.5- very mild, 1- mild, 2-moderate) Study neuropsychological test domains: episodic memory, working memory, executive functioning, mental speed, perception, and verbal fluency</p>	<p>Not reported</p>	<p>Frailty association with poorer cognitive performance Able to distinguish between frail and non-frail patients with Alzheimer's Disease (AD) ($\beta = -0.31$, $P < 0.001$)</p>	<p>Time and resource intensive Clinical translation properties unclear</p>	<p>Higher frailty score was highly correlated with poorer cognitive performance and poorer clinical manifestations of Alzheimer's disease Association between co-morbidity, frailty, and clinical manifestation of Alzheimer's disease</p>	<p>1b Cross-sectional Population-based study</p>
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<p>McGough et al. 2013 Dimensions of physical frailty and cognitive function in older adults with amnesic mild cognitive impairment</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: U.S. N= 201</p> <p>Analysis of baseline data from the Resources and Activities for Life-Long Independence (RALLI) Study</p> <p>70 and older, sedentary, and classified as having amnesic-MCI</p>	<p>Modified CHS criteria:</p> <p>Physical slowness: gait speed calculating energy expenditure (MET levels) cut off: < 383 Kcals/week men & <270 Kcals/week women</p> <p>Physical activity: Self-report using the Physical Activity Scale for the Elderly (PASE)</p> <p>Strength: Grip strength – cut off points stratified by sex and BMI</p> <p>Gait speed: 8-foot timed walk (best of two) – cut off stratified by sex and height</p> <p>Weight loss – assessed as a covariate, BMI calculated using baseline height/weight</p>	<p>Baseline Neuropsychological testing: MMSE, Wechsler Memory Scale-Revised (WMS-R), Logical memory (LM) I & II, CDR^f</p> <p>Study neuropsychological tests: severity measured with ADAS-Cog Attention and executive function: Trail Making A & B (TMT-A)</p> <p>Memory: WMS-R Logical Memory I (LM1), Word recall sub-item on ADAS-Cog</p> <p>*GDS: depression screening</p>	<p>Not reported</p>	<p>Reported on adjusted measures:</p> <p>Gait speed and cognitive function: ADAS-Cog ($\beta = -0.19$, $p < 0.008$) Executive function: TMT-A ($\beta = -0.23$, $p = 0.001$) TMT-B ($\beta = -0.20$, $p = 0.006$), Word Recall ($\beta = -0.18$, $p = 0.02$) and LM1 ($\beta = 0.14$, $p = 0.04$)</p> <p>Grip strength and attention: TMT-A ($\beta = -0.16$, $p = 0.008$)</p> <p>Physical activity and executive function ($\beta = -0.18$, $p < 0.02$) and word recall ($\beta = 0.17$, $p = 0.02$)</p>	<p>Time and resource intensive</p> <p>Clinical translation properties unclear</p>	<p>Slower gait speed was associated with elevated severity of cognitive impairment</p> <p>Gait speed associated with individual cognitive domains: attention, executive function, word recall, & memory</p> <p>Physical activity associated with the individual cognitive domain of executive function</p> <p>Grip strength associated with the individual cognitive domain of attention</p> <p>Grip strength not associated with severity of cognitive impairment</p>	<p>1b Cross-sectional study Baseline data from RTC</p>
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Alencar, et al. 2013 Frailty and cognitive impairment among community-dwelling elderly	Indices of cognitive frailty were discussed	Country: Brazil N= 182 Community-dwelling 65 years or older; with and without cognitive impairment Exclusion criteria: bed-ridden, restricted to wheelchair, terminal stage, hearing or vision impairment that would affect testing, stroke, severe stage Parkinson's disease, severe dementia (grade 3 on CDR ^ε)	CHS criterion: frailty phenotype defined by 3 or more of the 5 domains: Weight loss: Unintentional weight loss ≥ 4.5 kg Strength: Grip strength (adjusted gender & BMI) Fatigue: Two questions on the Center for Epidemiologic Studies Depression scale Slowness: time in seconds to walk 4.6 meters (14.8 feet-adjusted gender & BMI) Physical activity: Short version – Minnesota Leisure Time Activity Questionnaire Nutritional status: BMI with cut off: <22kg underweight; ≥22kg and ≤ 27kg ideal range; >27kg overweight Functional status: Katz scale: basic activities of daily living (BADL) instrumental activities of daily living (IADLs), Advanced activities of daily living	Cognitive function assessed in two-stage sequential testing: MMSE Cut off: 17/18 illiterate participants, 20/21 1-4 yrs of school, 23/24 5-8yrs of school, 25/26 9+ yrs of school (Nitrini and Caramelli 2007) When MMSE positive for cognitive changes then Brief Cognitive Screening Battery (BCSB) was completed ^ε CDR used for classification for degree of dementia: score 0.5-2 (0.5- very mild, 1-mild, 2-moderate, 3-severe) *GDS-15 – depression symptoms in individuals without cognitive impairment and Cornell Depression Scale in Dementia for individuals with cognitive impairment	Not reported	Mean difference at baseline MMSE - 12 months MMSE for non-frail, pre-frail, frail: (1.31, 0.49, 0.77 p = 0.005) Change in CDR baseline - 12 months CDR for non-frail, pre-frail, frail: (4 n=43, 17 n=104, & 7 n=35 p=0.393) Relative risk (RR) with ^ε CDR non-frail, pre-frail, frail: (RR = 1.0, 1.7 95% CI 0.63-0.49, 2.1 95% CI 0.68-6.7, p = 0.393) Relative risk with MMSE non-frail, pre-frail, frail: (RR = 1.0, 3.5 95% CI 1.51-8.4, 4.6 95% CI 1.9-11.2)	Time and resource intensive Clinical translation properties unclear	Risk of incidence and rates of progression for frailty and cognitive were significant when using the MMSE but not with the CDR	1b Prospective cohort study (12 months)
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<p>Gray et al. 2013 Frailty and incident dementia</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: U.S. N=2619</p> <p>From the Adult Changes in Thought (ACT) study</p> <p>65 and older without dementia at baseline</p> <p>Exclusion criteria: History of stroke, Parkinson's disease, or any component of frailty missing</p>	<p>CHS criteria: ≥3 = frail, 1-2 = pre-frail, 0 not frail</p> <p>Weakness: grip strength-average of 3 attempts (cut off by sex and BMI)</p> <p>Slowness: walking speed-10 foot walk – 2 walks average time</p> <p>Physical activity: self-report based on type of activity and length of time</p> <p>Weight loss: loss of 7.5% of body weight since previous visit</p> <p>Exhaustion: 10-item Center for Epidemiological Studies Depression (CES-D)</p>	<p>Cognitive Abilities Screening Instrument (CASI; 13) 40 item, 100-point global cognitive functioning test (cut off 86)</p> <p>1-hr neurocognitive battery: clock drawing, verbal fluency, Mattis Dementia Rating Scale, Boston naming, verbal paired associations and recall, logical memory and recall, Word List Memory, Constructional Praxis and recall, Trails A and B, and Information and Comprehension subtest items</p>	<p>Not reported</p>	<p>Frailty and all cause dementia Hazard Ratio (HR) (1.20, 95% CI 0.85-1.69)</p> <p>Frailty and Alzheimer's (HR 1.08, 95% CI 0.74-1.57)</p> <p>Frailty and non-Alzheimer's (HR 2.57, 95% CI 1.08-6.11)</p> <p>Gait speed and Alzheimer's disease (AD): (HR 2.13, 95% CI 1.09-4.16) CASI: Sensitivity 95.6%; specificity 92.0%</p>	<p>Time and resource intensive</p> <p>Clinical translation properties unclear</p>	<p>Frailty is associated with a 2.57 fold increase risk for non-AD dementia</p> <p>Individual frailty components were not significantly related to risk for dementia or AD.</p> <p>Slow gait speed was the only significantly related component to increased risk for non-AD dementia.</p>	<p>1b</p> <p>Population-based, longitudinal study (mean 6.5 years)</p>
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Solfrizzi et al. 2013 Frailty syndrome and the risk of vascular dementia: the Italian Longitudinal Study on Aging	Indices of cognitive frailty were discussed	Country: Italy N=2581 From the Italian Longitudinal Study on Aging (ILSA) 65-84 years; no dementia at baseline Exclusion: severe sensorial deficit, bedridden, use of wheelchair, dizziness, severe osteoarthritis, Parkinson's disease, or stroke	Modified CHS criteria: Weight loss: Unintentional weight loss > 5kg in past year (additional question: "Do you think that your clothes are wide?" Exhaustion: *GDS score \geq 10 and negative answer to the question: "Do you feel full of energy?" Weakness: Negative chair stand test: Inability to stand from a chair unaided, or without using the arms (standardized by sex and body mass index) Slowness: Time \geq 7 seconds spent to walk 5m (standardized by sex and height) Physical activity: structured questionnaire developed in the CHIANTI Study (Patel et al. 2006) Levels of physical activity in the past year. ADL and IADL tasks & *GDS item "Do you practice physical activity?" Motor performance: six tests: 3 explored dynamic balance and coordination; 3 assessed static balance	MMSE score of > 15 were considered to make plausible *GDS scores	Motor performance Intra-observer reliability: Dynamic balance: timed & counted tests 0.071 Tandem gain errors: 0.80 reaction time & .089-0.96 for chair stand, rapid step ups, standing on one leg, step length, and walking speed Intra-observer agreement; 0.63 gait – 0.82 abnormal turn ADL & IADL intra-observer agreement Cohen's Kappa=0.80 Inter-observer agreement; gait: 0.38 step asymmetry – 0.82 abnormal turn	Frailty association with overall dementia (HR 1.85, 95% CI 1.01-3.40) Frailty association with AD (HR 0.62, 95% CI 0.20-1.89) Frailty association with vascular dementia (HR 2.68, 95% CI 1.16-7.17)	Time and resource intensive Clinical translation properties unclear	Frailty syndrome at baseline was associated with a greater risk of developing overall dementia with strong associations with vascular dementia Relationship between grip strength and risk of AD	1b Population-based, longitudinal study (3.9 years)
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<p>Robertson et al. 2014 Cognitive function in the prefrailty and frailty syndrome</p>	<p>Indices of cognitive frailty were discussed</p>	<p>Country: Republic of Ireland N=4,649</p> <p>Adults 50 and older</p> <p>Exclusion: stroke, Parkinson's disease, taking antidepressants, or severe cognitive impairment MMSE <18</p>	<p>CHS criteria:</p> <p>Poor grip strength: Two readings from dominant hand – mean strength</p> <p>Slow gait speed: GAITRite portable electronic walkway system (16-foot) walkway with extra 2.5 m at each end for acceleration/deceleration</p> <p>Low levels of physical activity: short form International Physical Activity Questionnaire Kcals per week</p> <p>Unintentional weight loss survey: "In the past year, have you lost 10lbs or more in weight when you were not trying to."</p> <p>Exhaustion: Used 2 items from the 20-item Center for Epidemiological Studies Depression (CES-D)</p>	<p>Global cognition MMSE & MoCA</p> <p>Executive function: visual reasoning, color trails Test B, & verbal fluency</p> <p>Memory: visual recall, visual recognition, immediate, & self-rated</p> <p>Attention: color trails Test A & sustained attention to response task</p> <p>Processing speed: Cognitive reaction time</p>	<p>Not reported</p>	<p>Components of frailty and domains of cognitive function: Exhaustion & global cognition (β- 0.18, $p < 0.008$)</p> <p>Slow gait & executive function (β- 0.20, $p < 0.008$), attention (β - 0.25, $p < 0.008$), and processing speed (β- -0.16, $p < 0.008$)</p> <p>Weak grip & global cognition (β- 0.26, $p < 0.008$) and executive function (β-0.14, $p < 0.008$)</p>	<p>Time and resource intensive</p> <p>Clinical translation properties unclear</p>	<p>Cognitive function is related to pre-frailty and frailty</p> <p>Gait speed and grip strength were associated with executive function, processing speed, and attention. These results were further validated with evidence that frail individuals had lower cognitive scores than pre-frail and robust.</p>	<p>1b Cross-sectional study</p>
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Han et al. 2014 Association of cognitive impairment with frailty in community-dwelling older adults	Indices of cognitive frailty were discussed	Country: South Korea N=10,388 Adults 65 and older, data from the 2008 Living Profiles of Older People Survey	CHS criteria: Weight loss: Unintentional weight loss of more than 5kg in past 6 months Exhaustion: Self-reported fatigue or depressive symptoms Low Physical Activity: Defined as energy expenditure due to physical activity in the lowest quintile in the last week Gait speed: Slowest quintile for the 2.5-m-walk speed (adjusted for height by gender) Grip strength: Lowest quintile (based on gender's body mass index)	Modified version of the Korean version of the MMSE (MMSE-KC) Cognitive impairment was defined as > 1.5 SD below age, gender, and education-specific mean scores	Not reported	MMSE-KC associated with higher odds of pre-frail (OR=1.27, 95% CI 1.04-1.55 in men; OR=1.25, 95% CI 1.02-1.53 in women) and frail (OR = 1.81, 95% CI 1.25-2.60 in men; OR = 1.69, 95% CI 1.25-2.30 in women) Higher cognitive function specific domains scores associated with lower likelihood of frailty by gender: Men: attention (OR =0.72, 95% CI 0.58-0.89), recall (OR = 0.89, 95% CI 0.80-0.98), judgement (OR = 0.80, 95% CI 0.64-0.99) Women: language repetition (OR = 0.83 95% CI 0.73-0.95) and visual construction (OR = 0.82 95% CI 0.70-0.96)	Time intensive to measure both domains Clinical translation properties unclear	Cognitive impairment is associated with pre-frail and frailty Higher scores in specific domains of cognitive function were identified as having a lower association with frailty; several were gender specific	1b Cross-sectional study
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* OCEBM Levels of Evidence Working Group. "The Oxford Levels of Evidence 2". Oxford Centre for Evidence-Based Medicine. <http://www.cebm.net/?s=levels+of+evidence>.

§ Results reported from original study

¥ Operational definition terms were reported in the table as written in the original study

* Geriatric Depression Scale (GDS)

£ Clinical Dementia Rating (CDR)

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MANUSCRIPT 2: Supplemental documents

Table I. Clinical and biomarkers results

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Abdullah et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	213	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-40								
(Aberg et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	80	Cognitive Decline Only	Alzheimer's disease, MCI		Insulin like growth factor protein (IGF-2)	Insulin like growth factor protein Binding Protein (IGFBP-2)							
(Adamis et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	142	Cognition Decline & Frailty	General cognitive decline	Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Insulin like growth factor protein (IGF-1)								
(Adriaensens et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	415	Cognition Decline & Frailty	General cognitive decline	Physical Function	IL-6		IL-1 beta CRP/hs-CRP	IL-1 Alpha IL-8				Alcohol intake	Low level education
(Aguilar et al., 2014)	1a	Observational (Cohort, Cross Sectional, Case-Control Studies)	348	Cognitive Decline Only	Alzheimer's disease, MCI		ApoE-4 single allele								
(Albrecht et al., 2015)	1b	Longitudinal Study	214	Cognitive Decline Only	MCI progression to AD		ApoE-4 single allele								
(Albrecht et al., 2015)	1a	Longitudinal Study	1112	Cognitive Decline Only	MCI progression to AD		ApoE-4 single allele								
(Alcolea et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	226	Cognitive Decline Only	MCI		Aβeta-42	t-tau		YKL-40 (neuroinflammation)					
(Al-Turki, Boston, McKirdy, & Barker, 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	470	Cognitive Decline Only	Alzheimer's disease		Body mass index	Alcohol intake		p-tau or Chitinase-3 Ch3L3					
(Alvarez-Rios et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	592	Frailty (pre-frail & frail)		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Propeptide of type I procollagen (PINP)								
(Andersson et al., 2008)	1b	Longitudinal Study	40	Cognitive Decline Only	MCI progression to AD		p-tau								
(Annweiler, Bataille, et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	43	Frailty only		Fatigue,Sarcopenia,Physical Activity	Beta 2-microglobulin (B2M)								
(Annweiler, Schott, et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1190	Cognition Decline & Frailty	General cognitive decline	Physical Function	Vitamin D (25(OH)D)	Chronic Disease 2 or more		Depression					
(Apostolova et al., 2015)	1a	Observational (Cohort, Cross Sectional, Case-Control Studies)	400	Cognitive Decline Only	Alzheimer's disease		IL-6	Clusterin	ApoE-genotype	Brain derived neurotrophic factor (BDNF)					
(S. Li, Okonkwo, Albert, & Wang, 2013)	1a	Observational (Cohort, Cross Sectional, Case-Control Studies)	139	Cognitive Decline Only	MCI progression to AD		ApoE-4 two alleles	Aβeta 1-42		P-tau181/Aβeta-42	IL-13				TNF-alpha
(Armstrong et al., 2014)		In Vitro	50	Cognitive Decline Only	Alzheimer's disease		Rab7	Rab7	EEA1	LAMP-1	LAMP-2				LC3
(Aschenbrenner et al., 2015)	1b	Longitudinal Study	238	Cognitive Decline Only	MCI progression to AD		Aβeta-42	t-tau							
(Ashton et al., 2015)	1a	Longitudinal Study	78	Cognitive Decline Only	Alzheimer's disease		alpha 2-macroglobulin (A2M)	Fibrinogen gamma-chain (FGG)	Complement factor H (CFH) protein 1						
(Atti et al., 2006)	1b	Longitudinal Study	1139	Cognitive Decline Only	General cognitive decline		Anemia								
(Auyeung et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1489	Frailty only		Gait,Sarcopenia,Grip Strength	Total Testosterone (TT)	Estradiol/Estrogen							
(Baldeiras et al., 2010)	1b	Longitudinal Study	70	Cognitive Decline Only	MCI progression to AD		Uric Acid	malondialdehyde (MDA)	Vitamin E						
(Bambo et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	56	Cognitive Decline Only	Alzheimer's disease		Ocular measures								
(Bambo et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	114	Cognitive Decline Only	Alzheimer's disease		Ocular measures								
(Bartali et al., 2006)	1b	Longitudinal Study	643	Frailty only		Physical Function	Vitamin B6	Cobalamin deficiency (B12)	Selenium						
(D. Baylis et al., 2013)	1b	Longitudinal Study	254	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	WBC	ESR	Neutrophils	Monocytes DHEAS	Lymphocytes	Albumin	T4	DHEAS	Cortison/DHEAS ratio
(Daniel Baylis et al., 2014)	1b	Longitudinal Study	367	Frailty only		Sarcopenia,Grip Strength	IL-1beta	IL-6	Cortisol	(dehydroepiandrosterone sulphate)	Cortisol/DHEAS ratio				
(Beasley et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	24417	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Calibrated protein intake - Food Frequency Questionnaire (FFQ)								
(Beasley et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	134961	Frailty only		Grip Strength,Physical Function	Calibrated protein intake - Food Frequency Questionnaire (FFQ)								
(Beauchet et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	934	Cognition Decline & Frailty	General cognitive decline	Gait,Grip Strength	Medication (Psychoactive drugs)	BMI			BMI				
(Berr, Balansard, Arnaud, Rousset, & Alperovitch, 2000)	1b	Longitudinal Study	1166	Cognitive Decline Only	General cognitive decline		Oxidative stress markers /Total antioxidant status (TAS)	Selenium							
(Bertens, Knol, Scheltens, & Visser, 2015)	1b	Longitudinal Study	284	Cognitive Decline Only	MCI progression to AD		Aβeta 1-42	t-tau							
(Blain et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	220	Frailty only		Gait,Sarcopenia,Grip Strength,Physical Function	BMI	Creatinine	IL-6	CRP/hs-CRP	Insulin like growth factor protein (IGF-1)	Vit D	Total cholesterol		LDL
(K Blennow et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	572	Cognitive Decline Only	Alzheimer's disease		Aβeta1-42/ Aβeta1-40 ratio	Aβeta 1-42	Aβeta 1-40 phosphoTau181 (P-tau181)						
(Kaj Blennow et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	53	Cognitive Decline Only	Alzheimer's disease		t-tau Amyloid precursor protein (APP)	Aβeta-42							
(Borroni et al., 2004)	1b	Longitudinal Study	48	Cognitive Decline Only	MCI progression to AD		Composite Score: Pyridoxal 5-phosphate (B6), Thiamin (B1), Riboflavin (B2), Folate (B9) Ascorbic acid (vitamin C), -Tocopherol (vitamin E), Cobalamin (B12), 25-Hydroxyvitamin D	Composite Score: trans linoleic acid (18:2w-6), Trans- linoleic acid (18:2w6f)	Composite Score: Arachidonic acid (20:4- linoleic acid 6), gamma-Linolenic acid (18:3w6)						
(Bowman et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	104	Cognitive Decline Only	Alzheimer's disease		ApoE-4 two alleles	ApoE-4 single allele	P-tau231/Aβeta-42/40 ratio	T-tau/Aβeta-42/40 ratio					
(Bretsky et al., 1999)	1b	Longitudinal Study	195	Cognitive Decline Only	Alzheimer's disease										
(Brys et al., 2009)	1b	Longitudinal Study	66	Cognitive Decline Only	MCI progression to AD		P-tau31 Cellular prion protein (PrPc)	BMI	Alcohol intake						
(Breitling, Müller, Stegmaier, Kliegel, & Brenner, 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1322	Cognitive Decline Only	Alzheimer's disease										

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Gattaz, Forlenza, Talib, Barbosa, & Bottino, 2004)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	49	Cognitive Decline Only	Alzheimer's disease, MCI		Phospholipase A2 (PLA2)								
(Ghidoni et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	185	Cognitive Decline Only	Alzheimer's disease, MCI										
	1b	Longitudinal Study	59	Cognitive Decline Only	MCI progression to AD		Cystatin C	ApoE-4 single allele							
(Ge Li et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	315	Cognitive Decline Only	Alzheimer's disease		isoprostanes/isoprostanes	Aβeta 1-42							
	1b	Longitudinal Study	158	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-42	F2-isoprostanes/isoprostanes							
(Glodzik-Sobanska et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	78	Cognitive Decline Only	Alzheimer's disease		P-tau231								
(Goetzl et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	84	Cognitive Decline Only	Temporal Dementia (FTD)		Cathepsin D	LAMP-1	Ubiquitin	HSP70					
	1b	Longitudinal Study	60	Cognitive Decline Only	Alzheimer's disease, MCI		LAMP-1	Ubiquitin	Insulin resistance (IR-HOMA)						
(Gomes-Marcos et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1284	Frailty only	Alzheimer's disease	Physical Activity	Fibrinogen	CRP/hs-CRP		Creatinine		Glycohemoglobin (HbA1c)	Triglyceride	Hemoglobin	
(Growdon et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	215	Cognitive Decline Only	Alzheimer's disease		Olfactory measures								
(Gruenewald, Seeman, Karlamangla, & Sarkisian, 2009)	1b	Longitudinal Study	803	Frailty only	Alzheimer's disease, MCI	Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Composite Score: Systolic BP, Diastolic BP, HDL, total/HDL ratio, glycosylated hemoglobin, waist-hip ratio, dehydroepiandrosterone sulfate, urinary cortisol, urinary norepinephrine, urinary epinephrine, fibrinogen, c-reactive protein, IL-6	ApoE-4 single allele							
(Gupta et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1112	Cognitive Decline Only	Alzheimer's disease, MCI										
							Composite Score: C-reactive protein (CRP) and serum amyloid A (SAA), cytokines such as tumor necrosis factor alpha (TNF-α), interleukin-6 (IL-6), and interleukin-8 (IL-8), the enzyme myeloperoxidase (MPO), and the adhesion molecule 1 (sICAM-1), soluble intercellular adhesion molecule-1 (sICAM-1) CRP, TNF-α, IL-6, IL-8, SAA, MPO, sICAM-1		Composite Score: von Willebrand factor (vWF), soluble vascular cell adhesion molecule 1 (sVCAM-1), soluble endothelial selectin (sE-selectin), soluble thrombomodulin (sTM), and sICAM-1						
(Heringa et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	363	Cognitive Decline Only	General cognitive decline										
(Hohman, Bell, & Jefferson, 2015)	1b	Longitudinal Study	279	Cognitive Decline Only	Alzheimer's disease		Vascular endothelial growth factor (VEGF)								
(Howard, Ferrucci, & Sun, 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	672	Cognition Decline & Frailty	General cognitive decline	Grip Strength	vascular endothelial growth factor (VEGF)								
(Hsu, Cumming, Naganathan, Blyth, & Handelsman, 2014)	1b	Longitudinal Study	955	Cognitive Decline Only	General cognitive decline		Total Testosterone (TT)	Free Testosterone (fTT)							
(Llaw et al., 2016)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	205	Frailty only	Alzheimer's disease, MCI	Gait	Follistatin								
(Hye et al., 2014)	1b	Longitudinal Study	1148	Cognitive Decline Only	Alzheimer's disease, MCI		CRP/hs-CRP	ApoE-4 single allele	Complement factor H (CFH) protein 1	Neural cell adhesion molecule (NCAM)		Aβeta-40			
(Hochstrasser, Ehrlich, Marksteiner, Sperner-Unterwegar, & Humpel, 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	103	Cognitive Decline Only	Alzheimer's disease, MCI		Epidermal growth factor (EGF)	metalloproteinases (MMP-2)							
(Hendrickson et al., 2015)	1b	Longitudinal Study	176	Cognitive Decline Only	Alzheimer's disease		t-tau	p-tau							
(Hessen et al., 2015)	1b	Longitudinal Study	122	Cognitive Decline Only	MCI progression to AD		t-tau								
(Henrik Zetterberg et al., 2008)	1b	Longitudinal Study	87	Cognitive Decline Only	Alzheimer's, MCI progression to AD		βeta-secretase (BACE-1)								
(Inglés et al., 2014)	1b	Longitudinal Study	742	Frailty only	Alzheimer's disease, MCI	Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	malondialdehyde (MDA)	Protein carbonyls							
(Jefferson et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1926	Cognitive Decline Only	General cognitive decline										
(Jagielski et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	27971	Cognitive Decline Only	General cognitive decline		Glucose (FBG) or Insulin level (OGTT)	CRP/hs-CRP							
(Hendrickson et al., 2015)	1b	Longitudinal Study	1677	Cognition Decline & Frailty	Alzheimer's disease	Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	CRP/hs-CRP	IL-6							
(Barnett et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	144	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-42								
(Gomar et al., 2011)	1b	Longitudinal Study	116	Cognitive Decline Only	MCI progression to AD		t-tau	Aβeta 1-42	Aβeta1-42/t-tau ratio						
(Kanai, Matsubara, Isoe, & Utakami, 1998)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	60	Cognitive Decline Only	Alzheimer's disease, MCI		Insulin like growth factor protein (IGF-1)	factor protein Binding Protein (IGFBP-3)							
(Kanai et al., 1998)	1b	Longitudinal Study	236	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-42/ Aβeta 1-40 ratio	t-tau							
(Kantarci et al., 2007)	1b	Longitudinal Study	197	Cognitive Decline Only	Alzheimer's disease		N-acetylaspartate (NAA)/creatinine (Cr)								
(Gruenewald et al., 2009)	1b	Longitudinal Study	756	Cognitive Decline Only	General cognitive decline		Cortisol								
(Kelly et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	226	Cognitive Decline Only	General cognitive decline		Ocular measures								
(Kester et al., 2015)	1b	Longitudinal Study	163	Cognitive Decline Only	MCI progression to AD		Neruogranin (NGRN)	Aβeta-42		t-tau		phosphoTau181 (P-tau181)			
(Kester et al., 2012)	1b	Longitudinal Study	154	Cognitive Decline Only	Alzheimer's, MCI, MCI progression to AD		F2-isoprostanes/isoprostanes								
(Kester et al., 2015)	1b	Longitudinal Study	163	Cognitive Decline Only	MCI progression to AD		(neuroinflammation or Chitinase-3 Ch3L3)	Visinin-like protein-1 (VILIP-1)							
(Simpson et al., 2016)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	59	Cognitive Decline Only	Alzheimer's disease, MCI		IL-8	IL-8							
(Kim et al., 2011)	1b	Longitudinal Study	70	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-42	Aβeta 1-40	ApoE-ε3/ε4 single allele						
(Kleinschmidt et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	94	Cognitive Decline Only	Alzheimer's, MCI, General cognitive decline		Aβeta 1-42/ Aβeta 1-40 ratio	Aβeta 1-42							
(Koal, Klavins, Seppi, Kemmler, & Humpel, 2015)	NA	In Vitro	100	Cognitive Decline Only	Alzheimer's disease		Aβeta 42	t-tau	phosphoTau181 (P-tau181)			Sphingolipid-SM(d18:1/18:0)	PC aa 36:1	Glycerophospholipids	

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Kobrosly, Seplaki, Jones, & van Wijngaarden, 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	4511	Cognitive Decline Only	General cognitive decline		Composite: systolic and diastolic blood pressure, waist-to-hip ratio, glycohemoglobin, albumin, creatinine clearance, total cholesterol, triglycerides, WBC, resting heart rate, CRP								
(Kravitz, Corrada, & Kawas, 2009)	1b	Longitudinal Study	305	Cognitive Decline Only	Alzheimer's disease	Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity		Sirtuin 1	Sirtuin 2	Sirtuin 3					
(Kumar et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	200	Frailty only					Blood pressure	BUN (blood urine nitrogen)		Creatinine	Albumin (ALB)		
(Kuyumcu et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	416	Cognitive Decline Only	Alzheimer's disease			Neutrophil/Lymphocyte ratio							
(Lafaille-Magnan et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	73	Cognitive Decline Only	General cognitive decline			Olfactory measures							
(Laske et al., 2011)	1b	Longitudinal Study	40	Cognitive Decline Only	Alzheimer's, General cognitive decline			Brain derived neurotrophic factor (BDNF)							
(Licastro, Davis, Polazzi, Rossi, & Cucinotta, 1996)	1b	Longitudinal Study	40	Cognitive Decline Only	Alzheimer's disease		CRP/hs-CRP	Alpha-1-antitrypsin (alpha1-AT)	alpha 2-macroglobulin (A2M)		Ceruloplasmin	Acid glycoprotein	Alpha-1-antitrypsin (alpha1-AT)	Trasferin	
(Licastro et al., 2001)	1b	Longitudinal Study	28	Cognitive Decline Only	Alzheimer's, Vascular Dementia			antichymotrypsin (ACT)	Glutathione peroxidase (GSH-Px)						
(Li et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	174	Cognitive Decline Only	Alzheimer's disease, MCI		Lactoferrin (LTF)	t-tau/Aβeta-42	ApoE-4 single allele						
(Lilddalle et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	839	Cognitive Decline Only	General cognitive decline		Holo-transcobalamin (holoTC)	Cobalamin deficiency (B12)	Methylmalonic acid (MMA)	Homocysteine (Hcy)					
(Zuliani et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	140	Cognitive Decline Only	Late-Onset Alzheimer Disease (LOAD), Vascular			IL-6							
(Zubenko, Hughes III, & Stiffler, 2001)	1b	Longitudinal Study	325	Cognitive Decline Only	Alzheimer's disease		ApoE-4 two alleles								
(H Zetterberg et al., 2007)	1b	Longitudinal Study	100	Cognitive Decline Only	MCI progression to AD			t-tau	phosphoTau181 (p-tau181)	Aβeta 1-42					
(A. M. Zelisko, D. R. Kerwin, 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	89	Cognitive Decline Only	Alzheimer's disease			Circumference/waist-to-hip ratio	Adiponectin						
(Zamroziewicz, Paul, Rubin, & Barbey, 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	40	Cognitive Decline Only	General cognitive decline		BMI								
(S. X. Leng et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	696	Frailty only			Polysaturated fatty acids (O3PUFAs)/ n-6/n-3 ratio								
(Sean X. Leng et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	133	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity		Insulin like growth factor protein (IGF-1)		WBC					
(Liu et al., 2014)	1b	Longitudinal Study	230	Frailty only											
(Locascio et al., 2008)	1b	Longitudinal Study	122	Cognitive Decline Only	General cognitive decline			IL-6R	IL-6						
(Lopez et al., 2008)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	274	Cognitive Decline Only	MCI progression to AD			eGFR	Cystatin C						
(Luchsinger et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	327	Cognitive Decline Only	Alzheimer's disease			Aβeta-40	Aβeta-42	CRP/hs-CRP					
(Luis et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	60	Cognitive Decline Only	Alzheimer's disease, MCI			Aβeta 1-40	Abeta1-42	Cystatin C					
(Maeba, Nishimukai, Sakasegawa, Sugimori, & Hara, 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	440	Cognitive Decline Only	Alzheimer's disease, MCI			Homocysteine (Hcy)	Aβeta-42						
(Mancinella et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	201	Cognitive Decline Only	General cognitive decline			Aβeta-40	Aβeta-42	Aβeta-42/Aβeta-40					
(Marksteiner & Humpel, 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	138	Cognitive Decline Only	Alzheimer's disease, MCI			Peroxidase (POD)	IL-6	HDL (low/increased)			ApoA1	ApoA2	ApoC2
(Martin-Ruiz et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	852	Cognition Decline & Frailty	General cognitive decline			Choline	Ethanolamin	PLsCho + PlsEtn	PLsCho/PlsEtn Ratio				
(Matteini et al., 2008)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	700	Frailty only				plasmalogen (PlsCho)	plasmalogen (PlsEtn)	Fibrinogen					
(Niklas Mattsson et al., 2009)	1a	Observational (Cohort, Cross Sectional, Case-Control Studies)	1583	Cognitive Decline Only	Alzheimer's,MCI,MCI progression to AD			CRP/hs-CRP	CRP/hs-CRP						
(Yin, Fan, Lin, Xu, & Zhang, 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	114	Cognitive Decline Only	MCI			glycogen synthase kinase-3 (GSK-3alpha)							
(Yavuz et al., 2008)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	290	Cognitive Decline Only	Alzheimer's disease			Blood pressure, hematocrit, hemoglobin, MCV, RBC,WBC, lymphocytes, monocytes, neutrophils, sodium, phosphate, urate, creatinine, glucose, total protein, ALT, Albumin, calcium, HbA1c, TG, HDL, LDL, TC, ApoA1, Cortisol, ApoB, Free T3, Free T4, hsCRP, NT-pro BNP, Ferritin,							
(Yano et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	210	Cognitive Decline Only	General cognitive decline			Grip Strength,Physical Function	Homocysteine, Vit B12, Vit D, IL6, F2 alpha, CD8						
(Yarasheski, Bhasin, Sinha-Hikim, Pak-Loduca, & Gonzalez-Cadavid, 2002)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	95	Frailty only				Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Homocysteine (Hcy)	Cystathionine	Cobalamin deficiency (B12)	Methylmalonic acid (MMA)	Carotenoids		
(Yang et al., 2011)	1b	Longitudinal Study	820	Cognitive Decline Only	Alzheimer's,MCI,MCI progression to AD										
(S. H. Wu et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1005	Frailty only											
(L. C. Wu, Shiesh, Kuo, & Lin, 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	90	Frailty only											
(Wolfsgruber et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	245	Cognitive Decline Only	MCI										
(Wolfsgruber et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	245	Cognitive Decline Only	Alzheimer's disease										

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Windham et al., 2014)	1b	Longitudinal Study	1857	Cognitive Decline Only	General cognitive decline		Tumor necrosis factor receptor 2 (TNFR2) YKL-40	Tumor necrosis factor receptor 1 (TNFR1)	CRP/hs-CRP	IL-6					
(Wildsmith, Schauer, Kaur, Mathews, & Honigberg, 2013)	1b	Longitudinal Study	66	Cognitive Decline Only	General cognitive decline		(neuroinflammation of Chitinase-3 Ch3L3)								
(Wikby et al., 2005)	1b	Longitudinal Study	240	Cognitive Decline Only	General cognitive decline		CD8	IL-2	IL-6	Persistent viral infection					
(Westin, Buchhave, Minthon, Janciauskiene, & Hansson, 2011)	1b	Longitudinal Study	149	Cognitive Decline Only	General cognitive decline		Chemokine receptor 2 (CCR2)	t-tau							
(Weise et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	54	Cognitive Decline Only	Alzheimer's disease		Aβeta1-42	TNF-alpha	IL-6						
(Watanabe et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	131	Frailty only		Sarcopenia	C1q	Mean platelet volume (MPV)							
(R. Wang, Jin, Li, & Liang, 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	360	Cognitive Decline Only	Alzheimer's disease, MCI		Platelet distribution width (PDW)	phosphoTau181 (P-tau181)							
(L. Wang et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	207	Cognitive Decline Only	Alzheimer's disease		Aβeta-42	ApoE-4 single allele	Aβeta 1-42	phosphoTau181 (P-tau181)	Aβeta 1-42/t-tau ratio				
(L. Wang et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	188	Cognitive Decline Only	Alzheimer's disease		Aβeta-42								
(Madison, Shaw, Jack, & Weiner, 2010)	1b	Longitudinal Study	600	Cognitive Decline Only	MCI progression to AD		P-tau181/Aβeta-42	ApoE-4 single allele	Aβeta 1-42	phosphoTau181 (P-tau181)	Aβeta 1-42/t-tau ratio				
(N Mattsson et al., 2013)	1b	Longitudinal Study	46	Cognitive Decline Only	Alzheimer's disease		Angiotensin converting enzyme (ACE)	Chromogranin A (CgA)	Axl receptor tyrosine kinase (AXL)	metalloproteinases (MMP-2)	Aβeta-42	t-tau	p-tau		
(Niklas Mattsson et al., 2015)	1b	Longitudinal Study	35	Cognitive Decline Only	Alzheimer's, MCI progression to AD		Aβeta-42	ApoE-2 single allele	ApoE-4 single allele						
(Meng et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1131	Frailty only		Gait,Sarcopenia,Grip Strength,Physical Activity,Physical Function	CRP/hs-CRP								
(Mielke, Bandaru, et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	100	Cognitive Decline Only	Alzheimer's disease		Ceramides C16:0	Ceramides C20:0	Stearoyl sphingomyelin [SM(39:1)]						
(Mielke et al., 2011)	1b	Longitudinal Study	120	Cognitive Decline Only	Alzheimer's disease		DHSM/DHCer ratio	SM/ceramide ratio							
(Mielke, Haughey, et al., 2010)	1b	Longitudinal Study	63	Cognitive Decline Only	Alzheimer's disease, MCI		Ceramides C22:0 F2-isoprostanes/fsoprostanes	Ceramides C26:0							
(De Leon et al., 2006)	1b	Longitudinal Study	16	Cognitive Decline Only	MCI progression to AD			Aβeta-40	P-tau231						
(Mocchegiani et al., 2012)	1b	Longitudinal Study	346	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Ratio-Zinc /Copper (Czr)	IL-6	Albumin (ALB)	Blood urea nitrogen	Total Cholesterol	CRP			
(Hessen et al., 2015)	1b	Longitudinal Study	122	Cognitive Decline Only	MCI progression to AD		t-tau								
(Moore et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	24	Cognitive Decline Only	General cognitive decline		IL-6	CRP/hs-CRP	Waist circumference/waist-to-hip ratio						
(Moreno et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	120	Frailty only		Gait,Physical Activity	CRP/hs-CRP	Insulin resistance (IR-HOMA)							
(Thambisetty, Metter, et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	197	Cognitive Decline Only	Alzheimer's disease		Glucose (FBG) or Insulin level (OGTT)								
(Muldoon et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	280	Cognitive Decline Only	General cognitive decline		Docosahexaenoic acid (DHA)								
(Muzembo et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	86	Frailty only		Grip Strength	Serum 8-hydroxy-2-deoxyguanosine (8-OHdG)								
(A. Ng, Jion, Zainal, & Kandiah, 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	64	Cognitive Decline Only	General cognitive decline		Creatinine								
(Noble et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1331	Cognitive Decline Only	General cognitive decline		CRP/hs-CRP	CRP/hs-CRP							
(T. P. Ng, Niti, Feng, Xua, & Yap, 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1654	Cognitive Decline Only	General cognitive decline		Albumin (ALB)	Albumin (ALB)							
	1b	Longitudinal Study	1654	Cognitive Decline Only	General cognitive decline		Albumin (ALB)	Albumin (ALB)							
(Nurk et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	2195	Cognitive Decline Only	General cognitive decline		Choline	Cobalamin deficiency (B12)	Methylmalonic acid (MMA)						
(O'Bryant, Waring, et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	366	Cognitive Decline Only	General cognitive decline		plasmalogen(PisCho) CRP/hs-CRP								
	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	366	Cognitive Decline Only	General cognitive decline		Composite Score: 10 (macrophage inflammatory protein 1, eotaxin 1, tumor necrosis factor-alpha, fibrinogen, interleukin 5 [IL-5], IL-7, IL-10, C-reactive protein, monocyte chemoattractant protein 1, and von Willebrand factor)								
(O'Bryant, Waring, et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	400	Cognitive Decline Only	Alzheimer's disease			Extracellular heat shock protein (eHsp)							
(Ogawa et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	665	Frailty only		Gait,Sarcopenia,Grip Strength	IL-6	72	TNF-alpha						
	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	665	Frailty only		Gait,Sarcopenia,Grip Strength	Composite Score: metabolites: three amino acids (glutamic acid, alanine, and aspartic acid), one non-esterified fatty acid (22:6n-3, DHA), one bile acid (deoxycholic acid), one phosphatidylethanolamine [PE(36:4)], and one sphingomyelin [SM(39:1)]								
(Olazaran et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	251	Cognitive Decline Only	Alzheimer's disease, MCI										
(Olazaran et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	57	Frailty only		Gait,Sarcopenia,Physical Function	IL-6								
(Forlenza et al., 2010)	1b	Longitudinal Study	258	Cognitive Decline Only	MCI progression to AD		Aβeta-42	t-tau	p-tau	ApoE-4 single allele					
(Öztürk et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	330	Cognitive Decline Only	Alzheimer's disease		Platelet distribution width (PDW)	ESR	CRP/hs-CRP	Albumin (ALB)	LDL				
(Pabst et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	940	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Vitamin D (25(OH)D)								
(Papassotiropoulos et al., 2000)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	53	Cognitive Decline Only	General cognitive decline		24S-hydroxycholesterol	ApoE-4 single allele							
(Buchhave et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	45	Cognitive Decline Only	MCI progression to AD		t-tau	Aβeta-42							
	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	45	Cognitive Decline Only	MCI progression to AD		Composite Score: adipometabolic profile (AMP) and albumin, triglycerides, homocysteine, folate, total cholesterol								
(Perna, Guido, Grassi, & Rondanelli, 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	290	Frailty only		Sarcopenia									

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Perrin et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	292	Cognitive Decline Only	General cognitive decline		YKL-40 (neuroinflammation or Chitinase-3 Ch3L3)	Transthyretin (TTR)		NrCAM Chromogranin A (CgA)					
(Pirttilä et al., 1998) (P. et al., 2013)	1b 1b	Longitudinal Study Longitudinal Study	25 396	Cognitive Decline Only Cognitive Decline Only	MCI progression to AD Alzheimer's disease		ApoE-4 single allele Aβeta-42 Medication (ACE inhibitor)	sAβeta/APP ratio ApoE-4 single allele		Soluble amyloid βeta protein (sAβeta)					
(Qiu et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	355	Cognitive Decline Only	Alzheimer's disease		F2- isoprostanol/isoprostanol								
(Quinn et al., 2004)	1b	Longitudinal Study	40	Cognitive Decline Only	Alzheimer's disease										
(Quintino-Santos et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1480	Cognitive Decline Only	General cognitive decline		ApoE-4 single allele								
(Rabassa et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	652	Cognitive Decline Only	General cognitive decline		Total Urinary polyphenols (TUPs)								
(Rasgon et al., 2011) (Rembach et al., 2014)	2b 1b	Observational (Cohort, Cross Sectional, Case-Control Studies) Longitudinal Study	50 1112	Cognitive Decline Only Cognitive Decline Only	General cognitive decline MCI		Insulin resistance (IR-HOMA) Aβeta 1-42	Aβeta 1-40	Aβeta 1-40						
(Reuben, Judd-Hamilton, Harris, & Seeman, 2003)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	870	Frailty only		Physical Activity	IL-6	CRP/hs-CRP							
(Revel et al., 2015) (Riemenschneider et al., 2002)	1b 1b	Longitudinal Study Longitudinal Study	97 28	Cognitive Decline Only Cognitive Decline Only	General cognitive decline Alzheimer's disease, MCI		Glutathione peroxidase (GS4-Px) t-tau Composite Score: Allostatic load+ HDL/TC ratio, Triglycerides, A1c, fibrinogen, C-reactive protein, waist-to-hip ratio, systolic and diastolic blood pressure, and lung function (PEF). (all 9 belong to the highest 25% indicating health risk) composite allostatic load score. (inflammation, cardiovascular, metabolic, body fat, and respiratory)			Aβeta-42					
(Read & Grundy, 2014) (Rgler, Wichart, & Jellinger, 2001) (Ruiz et al., 2013) (Såmgård et al., 2010)	1b 2b 2b 1b	Longitudinal Study Observational (Cohort, Cross Sectional, Case-Control Studies) Observational (Cohort, Cross Sectional, Case-Control Studies) Longitudinal Study	6132 170 140 142	Frailty only Cognitive Decline Only Cognitive Decline Only Cognitive Decline Only	Frailty only Alzheimer's disease Alzheimer's disease, MCI Alzheimer's disease	Gait, Physical Function	t-tau Diastolic pressure t-tau Waist	Aβeta-42 Hematocrit p-tau	IL-6 IgA	ApoE-4 single allele Creatinine	Homocysteine (Hcy)	Urea	Uric acid		
(Sanada et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1488	Frailty only		Sarcopenia, Grip Strength	Circumference/waist-to-hip ratio DHEAS (dehydroepiandrosterone sulphate)	Glycohemoglobin (HbA1c)							
(Sanders et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	989	Cognition Decline & Frailty	General cognitive decline	Gait, Grip Strength	plasma desmosterol-to-cholesterol ratio (DES/CHO) plasma desmosterol-to-cholesterol ratio (DES/CHO)								
(Sato et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	401	Cognitive Decline Only	Alzheimer's disease		Derivate of reactive oxygen metabolites (d-ROM)								
(Saum et al., 2015) (Egll et al., 2015)	2b 1a	Observational (Cohort, Cross Sectional, Case-Control Studies) Longitudinal Study	2518 36	Frailty only Cognitive Decline Only	MCI progression to AD	Fatigue, Gait, Sarcopenia, Grip Strength, Physical Activity	Aβeta 1-42	CRP/hs-CRP	thol level (TTL)						
(Schaap, Pluijm, Deeg, & Visser, 2006) (Schofield, Ebrahimi, Jones, Bateman, & Murray, 2012) (Von Arnim et al., 2012)	1b 2b 2b	Longitudinal Study Observational (Cohort, Cross Sectional, Case-Control Studies) Observational (Cohort, Cross Sectional, Case-Control Studies)	986 56 232	Frailty only Cognitive Decline Only Cognitive Decline Only	Frailty only MCI progression to AD General cognitive decline	Sarcopenia, Grip Strength	IL-6	CRP/hs-CRP	alpha-1-antichymotrypsin (ACT)						
(Von Arnim et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	232	Cognitive Decline Only	General cognitive decline		Olfactory measures Vitamin C Tumor necrosis factor receptor 1 (TNFR1)	Beta-Carotene							
(Vieira et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	3150	Cognitive Decline Only	General cognitive decline			Blood pressure	HDL (low/increased)						
(S. Vestergaard et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1055	Frailty only		Fatigue, Gait, Sarcopenia, Grip Strength, Physical Activity	IL-6	CRP/hs-CRP	ft3						
(P. F. Vestergaard et al., 2014) (Verghese et al., 2011)	2b 2b	Observational (Cohort, Cross Sectional, Case-Control Studies) Observational (Cohort, Cross Sectional, Case-Control Studies)	150 333	Frailty only Frailty only		Sarcopenia, Grip Strength Gait	Insulin like growth factor protein (IGF-1) IL-6								
(Velayudhan, Pritchard, Powell, Proitsi, & Lovestone, 2013) (L. Van Den Ingh, A. Ahmed, 2011)	2b 2b	Observational (Cohort, Cross Sectional, Case-Control Studies) Observational (Cohort, Cross Sectional, Case-Control Studies)	57 254	Cognitive Decline Only Cognitive Decline Only	Alzheimer's disease General cognitive decline		Olfactory measures Total bilirubin								
(van den Boogaard et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	100	Cognitive Decline Only	General cognitive decline		TNF-alpha	IL-6	IL-8	Macrophage Migration Inhibitory Factor (MIF)			IL-1RA		
(Urpi-Sarda et al., 2015) (Umegaki et al., 2000)	2b 2b	Observational (Cohort, Cross Sectional, Case-Control Studies) Observational (Cohort, Cross Sectional, Case-Control Studies)	811 66	Frailty only Cognitive Decline Only	MCI	Fatigue, Gait, Sarcopenia, Grip Strength, Physical Activity	Total dietary polyphenols (TDPs) Cortisol	Total Urinary polyphenols (TUPs)	IL-6	CRP/hs-CRP	Total Cholesterol				
(Turana et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	109	Cognitive Decline Only	General cognitive decline		Ocular measures Plasminogen activator inhibitor (PAI-1) Cystatin C	Olfactory measures Serum amyloid A (SAA)	CRP/hs-CRP	TNF-alpha	IL-1 beta				
(Trollor et al., 2011) (Sundelof et al., 2008)	1a 1b	Observational (Cohort, Cross Sectional, Case-Control Studies) Longitudinal Study	710 761	Cognitive Decline Only Cognitive Decline Only	MCI MCI progression to AD										

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Uchida et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	411	Cognitive Decline Only	Alzheimer's disease, MCI		ApoA1	Complement 3	Transthyretin (TTR)						
	1b	Longitudinal Study	35	Cognitive Decline Only	MCI progression to AD		Transthyretin (TTR)	ApoE-genotype							
(Sunderland et al., 2003)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	203	Cognitive Decline Only	Alzheimer's disease		Aβeta 1-42	t-tau							
(Toledo, Xie, Trojanowski, & Shaw, 2013)	1b	Longitudinal Study	142	Cognitive Decline Only	Alzheimer's disease, MCI		Aβeta 1-42	t-tau	phosphoTau181 (P-tau181)						
(Thuot et al., 2010)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	23	Cognitive Decline Only	Alzheimer's disease, MCI		Insulin resistance (IR-HOMA)								
(Satizabal, Zhu, Mazoyer, Dufouil, & Tzourio, 2012)	1b	Longitudinal Study	1841	Cognitive Decline Only	General cognitive decline		IL-6	CRP/hs-CRP							
(Schoonenboom et al., 2005)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	39	Cognitive Decline Only	General cognitive decline		Aβeta-42	t-tau							
(Schram et al., 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	4365	Cognitive Decline Only	General cognitive decline		CRP/hs-CRP	IL-6							
	1b	Longitudinal Study	4365	Cognitive Decline Only	General cognitive decline		Composite Score: IL-6 and APOE e4								
(Semba et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	804	Frailty only		Sarcopenia, Grip Strength	Klotho								
(Seppälä et al., 2011)	1b	Longitudinal Study	131	Cognitive Decline Only	MCI progression to AD		Aβeta-42	t-tau	phosphoTau181 (P-tau181)						
		Observational (Cohort, Cross Sectional, Case-Control Studies)	131	Cognitive Decline Only	MCI		phosphoTau181 (P-tau181)	Aβeta-42							
(Shinkai et al., 1995)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	52	Frailty only		Physical Activity	Phytohemagglutinin	Pokeweed mitogen		IL-2	CXCL-10/IFN-gama		IL-4		
		Observational (Cohort, Cross Sectional, Case-Control Studies)	52	Frailty only			Brain derived neurotrophic factor (BDNF)								
(Shimada et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	827	Cognitive Decline Only	MCI										
(Singh-Manoux, 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	5217	Cognitive Decline Only	General cognitive decline		IL-6	CRP/hs-CRP							
	1b	Longitudinal Study	5217	Cognitive Decline Only	General cognitive decline		IL-6								
(Simpson et al., 2016)		In Vitro	107	Cognitive Decline Only	General cognitive decline		PC 16:0/20:5	PC 16:0/22:6	PC 18:0/22:6						
(Colbert et al., 2004)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	3075	Frailty only		Physical Activity	phosphatidylcholine	phosphatidylcholine	phosphatidylcholine						
(Snider et al., 2009)	1b	Longitudinal Study	49	Cognitive Decline Only	MCI progression to AD		IL-6	CRP/hs-CRP							
(Sohrabi et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	144	Cognitive Decline Only	General cognitive decline		Aβeta-42	t-tau	p-tau						
(F. Song et al., 2012)	1a	Longitudinal Study	664	Cognitive Decline Only	MCI progression to AD		Olfactory measures								
(I.-U. Song, Chung, Kim, & Maeng, 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	538	Cognitive Decline Only	Alzheimer's disease		ApoA1	ApoA2	ApoH	ApoB/ApoA1 ratio					
		Observational (Cohort, Cross Sectional, Case-Control Studies)	538	Cognitive Decline Only	Alzheimer's disease		CRP/hs-CRP								
(Spiegel et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	115	Cognitive Decline Only	Alzheimer's disease		phosphoTau181 (P-tau181)		P-tau231	phosphoTau181 (P-tau181)					
(Stricker et al., 2012)	1b	Longitudinal Study	342	Cognitive Decline Only	Alzheimer's disease, MCI		Aβeta-42								
(M. Soundararajan et al., 2011)	1b	Randomized control study	100	Cognitive Decline Only	General cognitive decline		Neurofilament light chain (NFL)								
(Skillbäck, Zetterberg, Blennow, & Mattsson, 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	5542	Cognitive Decline Only	Alzheimer's disease		Neurofilament light chain (NFL)								
(Teunissen et al., 2003)	1b	Longitudinal Study	144	Cognitive Decline Only	General cognitive decline		Homocysteine (Hcy)								
(Stomrud, Minthon, Zetterberg, Blennow, & Hansson, 2015)	1b	Longitudinal Study	44	Cognitive Decline Only	MCI progression to AD		Aβeta-42								
(Tapiola et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	123	Cognitive Decline Only	Alzheimer's disease		t-tau	Aβeta-42	alpha-secretase (ADAM10)	Ah/h-actin	APP ratio				
(Tang, Hynan, Baskin, & Rosenberg, 2006)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	41	Cognitive Decline Only	Alzheimer's disease	Fatigue, Gait, Sarcopenia, Grip Strength, Physical Activity	βeta-secretase (BACE-1)	Parathyroid hormone (PTH)							
(Tajar et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1504	Frailty only		Gait, Grip Strength, Physical Activity, Physical Function	Vitamin D (25(OH)D)	Lipopolysaccharide							
(Stehle et al., 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	59	Frailty only			binning protein (LBP)	CRP/hs-CRP	TNF-alpha	Tumor necrosis factor receptor 1 (TNFR1)					
		Observational (Cohort, Cross Sectional, Case-Control Studies)	59	Frailty only							YKL-40				
(Sutphen et al., 2015)	1b	Longitudinal Study	169	Cognitive Decline Only	General cognitive decline		Aβeta-42	t-tau	phosphoTau181 (P-tau181)	Visinin-like protein-1 (VILIP-1) or Chitinase-3 Ch3L3	(neuroinflammation)				
(Stanga, Lanni, Sinforiani, Mazzini, & Racchi, 2012)	1b	Longitudinal Study	67	Cognitive Decline Only	MCI progression to AD		Unfolded p53								
(Tay et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	44	Frailty only		Sarcopenia	IL-6	WBC	Albumin (ALB)						
(Allard, Artero, & Ritchie, 2003)	1b	Longitudinal Study	372	Cognitive Decline Only	General cognitive decline		Medication (Psychoactive drugs)								
(Bernhard T. Baune et al., 2008)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	369	Cognitive Decline Only	General cognitive decline		IL-8	IL-1 βeta							
(Gray et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	3434	Cognitive Decline Only	Alzheimer's disease										
		Observational (Cohort, Cross Sectional, Case-Control Studies)	3434	Cognitive Decline Only	Alzheimer's disease		Medication (Anticholinergic)								
(Fox et al., 2011)	1a	Observational (Cohort, Cross Sectional, Case-Control Studies)	13004	Cognitive Decline Only	General cognitive decline										

Citation	Level of Evidence	Type of study design	total (n)	Phenotype	Type of cognitive decline	Component of frailty	Biomarker - 1	Biomarker - 2	Biomarker - 3	Biomarker - 4	Biomarker - 5	Biomarker - 6	Biomarker - 7	Biomarker - 8	Biomarker - 9
(Ferrucci et al., 2002)	1b	Longitudinal Study	620	Frailty only		Gait,Sarcopenia,Grip Strength,Physical Function	IL-6								
(Boxer, Dauser, Walsh, Hager, & Kenny, 2008)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	60	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Vitamin D (25(OH)D)	Cortisol/DHEAS ratio	CRP/hs-CRP	IL-6	Parathyroid hormone (PTH)				
Butchart, Birch, Bassily, Wolfe, & Holmes, 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	94	Cognitive Decline Only	Alzheimer's disease		Free Testosterone (cFT)	LH	TNF-alpha						
(Beavers, Beavers, Serra, Bowden, & Wilson, 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	7544	Frailty only		Sarcopenia	Uric Acid								
(Barzilay, 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	3141	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Glucose (FBG) or Insulin level (OGTT)	WBC	CRP/hs-CRP		Von Willebrand Factor				
(Kizilarslanoglu et al., 2015)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	69	Cognitive Decline Only	Alzheimer's disease		Resistin				Fibrinogen	Vllc			
(Schmaltz et al., 2005)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	724	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	IL-6	Cytomegalovirus (CMV)							
(Roubenoff et al., 2003)	1b	Longitudinal Study	403	Frailty only		Sarcopenia	TNF-alpha	IL-6	Insulin like growth factor protein (IGF-1)						
(Kumar et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	93	Cognitive Decline Only	Alzheimer's disease, MCI		Sirtuin/SIRT1								
(S. Leng, Chaves, Koenig, & Walston, 2002)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	30	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	IL-6	Hemoglobin	Hematocrit						
(S X Leng, Xue, Tian, Walston, & Fried, 2007)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	558	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	WBC	IL-6							
(Levine & Crimmins, 2012)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	2287	Frailty only		Sarcopenia,Physical Activity,Physical Function	CRP/hs-CRP	HOMA							
(Liaw et al., 2016)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	205	Frailty only		Gait,Grip Strength	Follistatin Medication								
(Paterniti, Dufouil, & Alperovitch, 2002)	1b	Longitudinal Study	1389	Cognitive Decline Only	MCI		(Benzodiazepine)								
(Puts, Visser, Twisk, Deeg, & Lips, 2005)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1271	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Vitamin D (25(OH)D)								
	1b	Longitudinal Study	885	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Vitamin D (25(OH)D)	CRP/hs-CRP							
(Paterniti et al., 2002)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	32	Frailty only		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	IL-6	CXCL-10/ IFN-gama Medication							
(Uusvaara et al., 2009)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	295	Cognitive Decline Only	General cognitive decline		(Anticholinergic)	(Anticholinergic)							
(Visser et al., 2002)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	3075	Frailty only		Sarcopenia,Grip Strength	IL-6	IL-6							
(Wichmann et al., 2014)	1b	Longitudinal Study	1947	Cognitive Decline Only	General cognitive decline		IL-6	CRP/hs-CRP							
(Wilson, Cohen, & Pieper, 2003)	1b	Longitudinal Study	1752	Cognitive Decline Only	General cognitive decline		D-dimer	IL-6R			Vascular cell adhesion molecule 1 (VCAM1)				
(Yaffe et al., 2008)	1b	Longitudinal Study	3030	Cognitive Decline Only	General cognitive decline		Cystatin C Medication								
(Retrospective et al., 2011)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	134	Cognitive Decline Only	General cognitive decline		(Anticholinergic)								
(Lancôt et al., 2014)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	131	Cognitive Decline Only	General cognitive decline		(Anticholinergic)	Medication (Hypertensive drug use)							
(Sharma et al., 2016)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1315	Cognitive Decline Only	General cognitive decline		Adiponectin	(PTX3)							
(Mooijaart et al., 2013)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	5653	Cognitive Decline Only	General cognitive decline		IL-6								
(Herukka et al., 2011)	1b	Longitudinal Study	123	Cognitive Decline Only	Alzheimer's disease		p-tau	Aβeta-42							
(Jansen et al., 2016)	2b	Observational (Cohort, Cross Sectional, Case-Control Studies)	1705	Frailty (pre-frail & frail)		Fatigue,Gait,Sarcopenia,Grip Strength,Physical Activity	Medication (Anticholinergic)								

* OCEBM Levels of Evidence Working Group. "The Oxford Levels of Evidence 2". Oxford Centre for Evidence-Based Medicine. <http://www.cebm.net/?s=levels+of+evidence>.

Table II. Genetic studies for cognitive decline and frailty

Citation	Study country	Type of Study	Total (n)	Primary focus of study	Gene	SNP	Chromosome	Effect/Minor allele
(A.A. et al., 2014)	Australia	Candidate Gene Study	292	Cognitive Decline	SORL1	rs2298813	11	G
				Cognitive Decline	SORL1	rs4935774	11	T
				Cognitive Decline	SORL1	rs1133174	11	G
(Chibnik et al., 2011)	US	Candidate Gene Study	1666	Cognitive Decline	CR1	rs6656401	1	A
				Cognitive Decline	PICALM	rs7110631	11	G
(Choi et al., 2003)	Korea	Candidate Gene Study	13667	Cognitive Decline	ApoE-genotype	rs429358_rs7412	19	e4
(Dixon et al., 2014)	Canada	Candidate Gene Study	237	Cognitive Decline	ApoE-genotype	rs429358_rs7412	19	e4
				Cognitive Decline	COMT	rs4680	22	G
(Erten-Lyons, Jacobson, Kramer, Grupe, & Kaye, (Fiocco et al., 2010)	US	Candidate Gene Study	243	Cognitive Decline	FAS	RS1468063	10	T
	US	Candidate Gene Study	2840	Cognitive Decline	COMT	rs4680	22	G
(Goh et al., 2015)	Singapore	Candidate Gene Study	27	Cognitive Decline	TOMM40	rs10524523	19	T
(Green et al., 2014)	US	Candidate Gene Study	160	Cognitive Decline	ApoE-genotype	rs429358 or rs7412	19	e4
				Cognitive Decline	CLU	rs11136000	8	C
(Lillenes et al., 2011)	Norway	Candidate Gene Study	1066	Cognitive Decline	OGG1	rs1052133	3	G
				Cognitive Decline	OGG1	rs1052133	3	G
				Cognitive Decline	APEX1	rs1048945	14	C
				Cognitive Decline	MUTYH	rs3219484	1	C
(Wang et al., 2015)	Australia	Candidate Gene Study	619	Cognitive Decline	EPHA1	rs11771145	7	A
(Schmidt, Wolff, Von Ahsen, & Zerr, 2012)	Germany	Candidate Gene Study	40	Cognitive Decline	CST3	rs1064039	20	T
				Cognitive Decline	EXOC3L2 or XTP7	rs597668	19	C
				Cognitive Decline	ApoE-e4 two alleles	rs429358_rs7412	19	e4/4
				Cognitive Decline	BIN1	rs744373	2	G
(Thambisetty et al., (Hohman, Koran, & Thornton-Wells, 2014)	US	Candidate Gene Study	57	Cognitive Decline	CR1	rs3818361	1	A
(Hollingworth et al.,	Australia	Genome Wide Association Studies (GWAS)	374	Cognitive Decline	POT1	rs4728019	7	A
	ADGC-Multi-center study	Genome Wide Association Studies (GWAS)	30172	Cognitive Decline	ABCA7	rs3764650	19	G
				Cognitive Decline	MS4A6A	rs610932	11	T
				Cognitive Decline	MS4A4E	rs670139	11	T
				Cognitive Decline	CD2AP	rs9349407	6	C
				Cognitive Decline	CD33	rs3865444	19	A
				Cognitive Decline	EPHA1	rs11767557	7	C
(Kauwe et al., 2014)	US	Genome Wide Association Studies (GWAS)	840	Cognitive Decline	IL-6R	rs4845622	1	C
				Cognitive Decline	IL-6R	rs61812598	1	G
				Cognitive Decline	IL-6R	rs4129267	1	T
				Cognitive Decline	IL-6R	rs2228145	1	C
				Cognitive Decline	IL-6R	rs2229238	1	T
				Cognitive Decline	TDRD10	rs3811448	1	G
				Cognitive Decline	CCL2	rs2228467	3	C
				Cognitive Decline	CCL4	rs6808835	3	G
				Cognitive Decline	CCL4	rs6762266	3	C
				Cognitive Decline	CCL4/LOC102724297	rs11575821	3	G
				Cognitive Decline	CCL4/LOC102724297	rs113263161	3	G
				Cognitive Decline	CCL4	rs11574428	3	T
				Cognitive Decline	CCL4	rs3092960	3	G
				Cognitive Decline	CCL4	rs6441977	3	G
				Cognitive Decline	MMP3	rs573521	11	A
				Cognitive Decline	MMP3	rs645419	11	A
				Cognitive Decline	MMP3	rs679620	11	T
				Cognitive Decline	WTAPP1	rs7926920	11	A
				Cognitive Decline	WTAPP1	rs11225434	11	C
				Cognitive Decline	MMP3	rs948399	11	T
				Cognitive Decline	WTAPP1	rs495366	11	A
				Cognitive Decline	MMP3	rs650108	11	A
				Cognitive Decline	WTAPP1	rs603050	11	T
				Cognitive Decline	ACE	rs4968782	17	G
				Cognitive Decline	ACE	rs4459609	17	C
				Cognitive Decline	ACE	rs4316	17	C
				Cognitive Decline	ACE	rs4343	17	G
(Hu et al., 2011)	US	Genome Wide Association Studies (GWAS)	1605	Cognitive Decline	UBR5	rs7840202	8	C
				Cognitive Decline	PARP6	rs11637611	15	C
				Cognitive Decline	ACOT11/LOC105378734	rs12752888	1	C
				Cognitive Decline	MYO9A	rs3784313	15	G
				Cognitive Decline	MYO9A	rs2957734	15	A
				Cognitive Decline	MYO9A	rs4777466	15	T
				Cognitive Decline	MYO9A	rs7175373	15	C
				Cognitive Decline	MYO9A	rs1481862	15	C
				Cognitive Decline	MYO9A	rs7497104	15	T
				Cognitive Decline	MYO9A	rs2929525	15	C
				Cognitive Decline	MYO9A	rs2306489	15	T
(Desikan et al., 2016)	IGAP-Multi-center study	Candidate Gene Study	143878	Cognitive Decline	AP2A2	rs7396366	11	C
				Cognitive Decline	USP50	rs1311609	15	C
				Cognitive Decline	TSPOAP1	rs2526378	17	G
				Cognitive Decline	HS35T1/LOC107986178	rs13113697	4	T
(Feulner et al., 2010)	Germany	Genome Wide Association Studies (GWAS)	970	Cognitive Decline	ECHDC3	rs7920721	10	A
				Cognitive Decline	MAPT	rs1467967	17	A
				Cognitive Decline	MAPT	rs3785880	17	G
				Cognitive Decline	KANSL1	rs6503454	17	G
				Cognitive Decline	MAPT-AS1	rs1158660	17	A
				Cognitive Decline	PCK1	rs17411904	20	C
				Cognitive Decline	LMNA	rs9919256	1	A
				Cognitive Decline	LMNA	rs11578696	1	G
				Cognitive Decline	LMNA	rs915179	1	A
				Cognitive Decline	LMNA	rs12128066	1	C
				Cognitive Decline	SEMA4A	rs12401573	1	C
				Cognitive Decline	LIPA	rs12780342	10	T
				Cognitive Decline	PGBD1	rs9461448	6	G
				Cognitive Decline	ZSCAN31	rs7772827	6	C
				Cognitive Decline	PGBD1	rs1320879	8	A
				Cognitive Decline	CH25H	rs17117126	10	A
				Cognitive Decline	ApoE-genotype	rs405509	19	T
				Cognitive Decline	TOMM40	rs8106922	19	G
				Cognitive Decline	TOMM40	rs2075650	19	G
				Cognitive Decline	TOMM40	rs157580	19	G
				Cognitive Decline	APOC1	rs439401	19	T
				Cognitive Decline	APOC2	rs5167	19	G
				Cognitive Decline	SORL1	rs4935774	11	T
				Cognitive Decline	SORL1	rs1614735	11	T
				Cognitive Decline	SORL1	rs12576704	11	G
				Cognitive Decline	SORL1	rs10502262	11	T
				Cognitive Decline	SORL1	rs3781835	11	G
				Cognitive Decline	TDRD10	rs3811448	1	A
				Cognitive Decline	TDRD10	rs7556449	1	G
				Cognitive Decline	UBE2Q1	rs7543174	1	C
				Cognitive Decline	ADAR	rs9427097	1	G
(Del-Aguila et al., 2015)	US	Candidate Gene Study	3476	Cognitive Decline	ABCA7	rs4147929	19	A
(Corneveaux et al.,	US, UK	Candidate Gene Study	1600	Cognitive Decline	CR1	rs6656401	1	A
				Cognitive Decline	LOC651924	rs6907175	6	A
				Cognitive Decline	CLU	rs11136000	8	T
				Cognitive Decline	PICALM	rs541458	11	C
				Cognitive Decline	GAB2	rs10793294	11	C
				Cognitive Decline	ACE	rs1800764	17	C
				Cognitive Decline	CST3	rs1064039	20	T

Citation	Study country	Type of Study	Total (n)	Primary focus of study	Gene	SNP	Chromosome	Effect/Minor allele
(Baune et al., 2008)	Australia	Candidate Gene Study	369	Cognitive Decline	IL-1beta	rs16944	2	G
						rs1800796	7	G
(Lambert et al., 2009)	France	Candidate Gene Study	7275	Cognitive Decline	TNF	rs1800629	6	A
						rs6656401	1	A
(Lim et al., 2015)	Australia	Candidate Gene Study	333	Cognitive Decline	CLU	rs11136000	8	T
(Reitz et al., 2013)	ADGC-Multi-center study	Genome Wide Association Studies (GWAS)	5896	Cognitive Decline	BDNF	rs6265	11	T
(Mooijaart et al., 2013)	US	Candidate Gene Study	5804	Cognitive Decline	BIN1	rs55636820	2	G
						rs115550680	19	G
(Forlenza et al., 2010)	Brazil	Candidate Gene Study	258	Cognitive Decline	EPHA1	rs6973770	7	G
(Vounou et al., 2012)	Australia	Genome Wide Association Studies (GWAS)	475	Cognitive Decline	CR1	rs9429784	1	G
(Mooijaart et al., 2013)	US	Candidate Gene Study	5804	Cognitive Decline	CD33(rsq)	rs114282264	19	G
						rs1729941	2	C
(Forlenza et al., 2010)	Brazil	Candidate Gene Study	258	Cognitive Decline	IL-6	rs1729941	2	C
(Vounou et al., 2012)	Australia	Genome Wide Association Studies (GWAS)	475	Cognitive Decline	ApoE-e4 single allele	rs429358_rs7412	19	e4
					ApoE-genotype	rs429358_rs7412	19	e4
					TOMM40	rs2075650	19	e4
					BZW1	rs3815501	2	G
						rs11132507	4	T
						rs11132508	4	C
					MIR924HG	rs1681052	18	T
						rs7761213	6	T
						rs17345545	1	C
					PDZD2	rs13340334	5	C
						rs17103124	14	T
						rs8025706	15	T
					FAM171B	rs12185469	18	A
					YES1	rs12185470	18	T
					YES1	rs10766003	11	G
					TEAD1	rs1503659	4	C
						rs913587	9	A
					KDM4C	rs17380902	2	C
						rs17686103	5	C
					LINC01019	rs785232	2	T
					C2orf88	rs4771473	13	T
						rs11740943	5	T
						rs7536709	1	T
					YES1	rs17516202	18	G
					TOMM40	rs157580	19	G
					MEF2D	rs1750304	1	A
					MEF2D	rs1171560	1	G
						rs9263844	6	T
						rs9263846	6	G
					MTRF1	rs7999394	13	G
					MTRF1	rs3794328	13	C
						rs11590365	1	A
						rs11204949	1	C
						rs11204971	1	G
					FLG	rs12405278	1	A
						rs215340	12	A
						rs7603289	2	C
					OSTF1	rs11144246	9	A
					MICA	rs6910087	6	T
						rs4685279	3	C
(Vounou et al., 2012)	Australia	Genome Wide Association Studies (GWAS)	475	Cognitive Decline	ARHGEF10	rs3824139	8	A
						rs6932730	6	C
					MEF2D	rs1750304	1	A
					MEF2D	rs1171560	1	A
						rs9263969	6	T
						rs6700106	1	A
						rs795342	12	A
						rs10026499	4	A
						rs7979925	12	C
						rs2325	10	T
						rs7944761	11	C
						rs9501132	6	T
						rs215347	12	G
					USP13	rs2268939	3	A
					KAZ	rs6429696	1	C
						rs11215380	11	A
					ADCY2	rs727432	5	T
						rs11783329	8	A
					MAML2	rs7114756	11	T
						rs17309585	8	T
						rs10491327	5	T
					PDE1C	rs12534148	7	A
					MYO3B	rs2883782	2	T
						rs2798062	9	A
						rs10934170	3	T
						rs17826780	4	T
						rs7843577	8	T
					TOMM40	rs2075650	19	G
						rs1405443	7	A
					RBFOX1	rs758491	16	C
						rs914166	21	T
						rs11150643	16	C
					COX7A2L	rs1981664	2	A
					COX7A2L	rs10206058	2	G
					PAPPA	rs717963	9	A
						rs10509839	10	A
					RGS6	rs763732	14	T
						rs6884345	5	C
						rs11242336	5	C
					ANK3	rs10994250	10	A
					ANK3	rs10821707	10	G
						rs3912887	4	A
						rs419867	21	A
						rs2837900	21	C
						rs2837902	21	A
					SORBS2	rs13132552	4	T
						rs12633719	3	C
						rs9522088	13	T
						rs885339	13	A
						rs2381958	5	A
						rs10041184	5	A
						rs4265409	1	C
					ANTXR1	rs7584948	2	G

Citation	Study country	Type of Study	Total (n)	Pimary focus of study	Gene	SNP	Chromosome	Effect/Minor allele
(Vounou et al., 2012)	Australia	Genome Wide Association Studies (GWAS)	475	Cognitive Decline	TENM4	rs501435	11	C
						rs1001684	5	A
						rs1257687	14	A
						rs7336788	13	G
						rs10065570	5	T
						rs6783007	3	T
						rs10070362	5	C
						rs9522086	13	T
						rs11946115	4	T
						rs7734346	5	T
						rs10155062	5	A
						rs1289501	11	C
						rs11949577	4	A
						rs13436090	5	A
						rs17370295	3	A
						rs3760961	19	G
						rs2965069	7	T
						rs478090	11	C
						rs2965245	19	A
						rs962492	5	C
						rs7963861	12	T
						rs705837	4	T
						rs11856999	15	T
						rs7653663	3	A
						rs12597064	16	G
						rs633398	4	C
						rs631271	4	A
						rs1529442	5	A
						rs6864491	5	A
						rs10445932	2	C
						rs885120	5	G
						rs12236788	9	A
rs2075650	19	G						
rs11136000	8	G						
rs3851179	11	T						
rs157580	19	T						
rs6859	19	G						
rs8106922	19	A						
rs11894266	2	G						
rs610932	11	T						
rs10501927	11	T						
rs9446432	6	G						
rs7561528	2	C						
rs744373	2	A						
rs662196	11	G						
rs583791	11	C						
rs676309	11	C						
rs1157242	8	C						
rs1539053	1	T						
rs11827375	11	G						
rs1408077	1	A						
rs9384428	6	A						
rs6701713	1	C						
rs3818361	1	A						
rs562020	13	G						
rs398655	13	G						
rs398655/rs562020	13	C						
rs2283368	13	AG						
rs9526984	13	C						
rs9536314	13	G						
rs9527024	13	G						
rs648202	13	A						
rs9536314	13	T						
rs9527025	13	C						
rs1207568	13	A						
rs1799990	20	G						
rs4646450	7	A						
rs4363657	12	T						
rs737267	4	T						
rs1260326	2	T						
(Mekil, Nazroo, Marshall, Kumari, & Patel et al., 2014)	UK	Candidate Gene Study	3160	Frailty phenotype	TNF	rs1800629	6	A
						rs1566729	11	T
						rs2047812	11	A
						rs1566728	11	C
						rs611646	11	T
						rs4646316	22	A
						rs731236	12	G
						rs121913168	12	T
						rs397515373	2	ClinVar
						rs1800796	7	G
rs361525	6	A						
rs28362304	2	T						
rs949963	2	T						
rs1544468	22	G						
rs2267163	22	T						
rs1801394	5	G						
rs5744256	11	C						
rs543810	11	C						
rs1293344	11	G						
(Matteini et al., 2008)	US	Candidate Gene Study	326	Frailty phenotype	IL-18	rs360722	11	A
						rs4679868	3	A
						rs9852519	3	A
						rs1799986	12	T
						rs6131	1	T
(Frayling et al., 2007)	Italy	Candidate Gene Study	1671	Physical Function	IL18	rs1770449	1	A
						rs10925235	1	G
						rs2297967	1	A
						rs10802569	1	A
						rs4659725	1	G
						rs1050993	1	C
						rs7567647	2	A
						rs129968	16	A
						rs3769827	2	A
						rs6747918	2	G
rs2037815	2	G						
rs6745051	2	G						
rs2287396	14	C						
rs2929408	3	T						
rs2833383	21	A						
rs1400657	2	T						
rs740234	22	C						
rs10883642	10	G						
rs10883631	10	A						
rs2227729	17	A						

Figure I. PubMed search strategy

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(("Frailty"[TIAB] OR "Frail"[TIAB] OR "Physical Frailty"[TIAB] OR "Frail Elderly"[Mesh] OR "Sarcopenia"[Mesh] OR "Muscle Weakness"[Mesh] OR "hand strength"[Mesh] OR "motor activity"[Mesh] OR "weight loss"[Mesh] OR "fatigue"[Mesh] OR "lassitude"[tiab] OR "motor activity"[tiab] OR "motor activities"[tiab] OR "physical activities"[tiab] OR "locomotor activity"[tiab] OR "locomotor activities"[tiab] OR "hand strength"[tiab] OR "grip"[tiab] OR "grips"[tiab] OR "grasp"[tiab] OR "grasps"[tiab] OR "gait speed"[tiab] OR "grip strength"[tiab] OR "physical activity"[tiab] OR "weight loss"[tiab] OR "fatigue"[tiab] OR "sarcopenia"[tiab] OR "tiredness"[tiab] OR "muscular weakness"[tiab])  
  
OR  
  
("Alzheimer Disease"[Mesh] OR "Dementia"[Mesh] OR "Mild Cognitive Impairment"[Mesh] OR "Cognition Disorders"[Mesh] OR "Alzheimer"[tiab] OR "Alzheimers"[tiab] OR "Alzheimer's"[tiab] OR "presenile dementia"[tiab] OR "senile dementia"[tiab] OR "cognitive impairment"[tiab] OR "cognitive impairments"[tiab] OR "neurocognitive disorder"[tiab] OR "neurocognitive disorders"[tiab] OR "dementia"[tiab] OR "dementias"[tiab] OR "cognitive decline"[tiab] OR "cognitive declines"[tiab] OR "cognition disorder"[tiab] OR "cognition disorders"[tiab])  
  
OR  
  
("cognitive frailty"[tiab])  
  
AND  
  
(("Biomarkers "[Mesh] OR "biological markers"[tiab] OR "biological marker"[tiab] OR "biologic markers"[tiab] OR "biologic marker"[tiab] OR "biomarkers"[tiab] OR "biomarker"[tiab] OR "clinical markers"[tiab] OR "clinical marker"[tiab] OR "Immunologic markers"[tiab] OR "immunologic marker"[tiab] OR "immune marker"[tiab] OR "immune markers"[tiab] OR "viral markers"[tiab] OR "viral marker"[tiab] OR "serum markers"[tiab] OR "serum marker"[tiab] OR "surrogate endpoints"[tiab] OR "surrogate endpoints"[tiab] OR "surrogate end points"[tiab] OR "surrogate end point"[tiab] OR "surrogate markers"[tiab] OR "surrogate marker"[tiab] OR "biochemical markers"[tiab] OR "biochemical marker"[tiab] OR "laboratory markers"[tiab] OR "laboratory marker"[tiab] OR "disease marker"[tiab] OR "disease markers"[tiab] )  
  
OR  
  
("Genetic markers"[Mesh] OR "genetic markers"[tiab] OR "genetic marker"[tiab] OR "DNA markers"[tiab] OR "DNA marker"[tiab] OR "Chromosome marker"[tiab] OR "Chromosome markers")  
  
OR  
  
("Genome-Wide Association Study"[Mesh] OR "genome wide association"[tiab] OR "whole genome association"[tiab] OR "GWAS"[tiab] OR "candidate gene study"[tiab] OR "candidate gene studies"[tiab]))  
  
AND  
  
("Clinical Trials as Topic"[Mesh] OR "Clinical Trial" [Publication Type] OR "Randomized Controlled Trial" [Publication Type] OR "Longitudinal Studies"[Mesh] OR "Random Allocation"[Mesh] OR "Cross-Sectional Studies"[Mesh] OR "clinical trial"[tiab] OR "clinical trials"[tiab] OR "randomized controlled"[tiab] OR "randomised controlled"[tiab] OR "random allocation"[tiab] OR "cross sectional study"[tiab] OR "cross sectional studies"[tiab] OR "cross sectional analysis"[tiab] OR "cross sectional analyses"[tiab] OR "longitudinal study"[tiab] OR "longitudinal studies"[tiab] OR "cross sectional survey"[tiab] OR "cross sectional surveys"[tiab] OR "prevalence study"[tiab] OR "prevalence studies"[tiab] OR "randomization"[tiab] OR "randomisation"[tiab] OR "cross-sectional research"[tiab] OR "cross-sectional design"[tiab] OR "Genome-Wide Association Study"[Mesh] OR "genome wide association"[tiab] OR "whole genome association"[tiab] OR "GWAS"[tiab] OR "candidate gene study"[tiab] OR "candidate gene studies"[tiab])
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Reproducibility

In seeking to explore the importance and applicability of these results it is critical that others continue to replicate model results before they can be used in the clinical setting. To accompany this report, help with replication and extension of our work, the code has been made publically available for model I and model II online.

Database

The subjects in the present study were participants in *Invecchiare in Chianti* (Aging in Chianti, “InCHIANTI Study”). InCHIANTI is a prospective population based study of 1,453 adults aged 20-102 randomly selected from two towns in Tuscany, Italy using a multistage stratified sampling at baseline from 1998 to 2000¹. All aspects of the InCHIANTI research were approved by the ethics committees at the institutions responsible for data collection.

Definitions used to establish phenotype sub-groups in this study

Cognitive decline – mild neurocognitive disorders

Evidence of modest cognitive decline from a previous level of performance in one or more cognitive domains (complex attention, executive function, learning and memory, language, perceptual motor, or social cognition) with a modest impairment in cognitive performance by standardized neuropsychological testing or clinical assessment in absence of a diagnosis of dementia^{2,34}.

Frailty

The operational definition for frailty is defined as a clinical syndrome condition including 3 out of the 5 criteria related a physical phenotype including: 1) weak muscle strength (grip strength), 2) slow gait speed, 3) unintentional weight loss, 4) exhaustion and low physical activity⁵. Pre-frailty includes 1 or 2 of the criteria is present, identifying a sub-group of individuals potentially progressing to frailty⁵.

Cognitive Frailty

A syndrome in older adults with evidence of both physical frailty and cognitive impairment without a clinical diagnosis of Alzheimer’s Disease or other dementia⁶.

Phenotypic classification for this study

Model I

Participants with an MMSE normal cognition 24-30 and cognitive decline $\leq 23^{7-9}$. In this study frailty is characterized by individuals with one or more of the frailty criterion⁵. Cognitive frailty is defined as individuals with cognitive decline and one or more of frailty criterion¹⁰.

- Robust with no physical frailty and absence of cognitive decline
- Robust with no physical frailty with cognitive (MMSE = ≤ 23)
- Frail (≥ 1 criteria) and absence of cognitive decline
- Frail (≥ 1 criteria) and cognitive decline (MMSE = ≤ 23)

Model II

Participants that completed the MMSE with additional neuropsychiatric testing Trail Making Test, Part A and B (TMT) to define cognitive decline and cognitive frailty^{10,11}. TMT cut off scores for cognitive decline are based on cut off norms established by Ashendorf et al., 2008.

- Robust with no physical frailty and absence of cognitive decline
- Robust with no physical frailty with cognitive decline (Trail A ≥ 78 , Trail B ≥ 106)
- Frail (≥ 1 criteria) and cognitive decline (Trail A ≥ 78 , Trail B ≥ 106)
- Frail (≥ 1 criteria) and cognitive decline (Trail A ≥ 78 , Trail B ≥ 106)

Laboratory assay methods

At the baseline survey, most of the participants performed 24-hour urine collection early in the morning mid-stream sample urine for the routine examination. Total urinary polyphenols were measured at the Department of Food Science and Technology, School of Pharmacy, University of Barcelona, Spain. Prior to blood collection all participants consumed a diet free of meat and fish. Participants donated fasting blood samples for routine blood examinations. Blood collection was performed with the standard procedure method to prevent red cell hemolysis. The blood collection included two sets of collection tubes: one for routine tests and second for collecting specimens including serum, plasma, DNA for the biological bank. All routine blood tests, performed in the Laboratory of Clinical Chemistry and Microbiological Assays, Annunziata Hospital in Florence, Italy. Plasma fatty acids (FAs) were measured by the Section of Gerontology and Geriatrics, Department of Clinical and Experimental Medicine, Perugia, Italy.

The technique used was gas chromatography with a fused silica capillary column to achieve the optimum separation of the different fatty acids.

Software for analyses

All statistical analyses were carried out using R V. 3.2.1. R is free, open-source software that provides many statistical and graphic techniques. R packages used included 'glm2'-Fitting Generalized Linear Models, 'Ordinal'-Regression Models for Ordinal Data, and 'xgboost'-Extreme Gradient Boosting¹²⁻¹⁴. The software package PLINK, an analysis toolset was used for the management of genotype data and basic associating testing^{15,16}.

Model generation

The predictive genetic and laboratory biomarkers were identified in a comprehensive systematic review and analyzed using an Extreme Gradient Boosting (xgboost) in R¹⁴. While boosting was initially developed for machine learning, 'xgboost' in R is based in boosted trees. Xgboost is an open source tool and a variant of the gradient boosting machine and uses a tree based model. Xgboost is used in this study for a supervised learning problem where the variables identified from the systematic review are used to predict three phenotypes cognitive decline, physical frailty, and cognitive frailty.

Evaluation of the model

With the use of any predictive model in machine learning there is a chance for inflated risk of capitalizing on chance features (over fitting) in the data. Over fitting of the integrative model was mitigated in two ways: 1) having a distinct training and validation process for the model and 2) using xgb in R which has a built-in parameter settings for selection to reduce poor predictive performance. *Internal validation*: A randomly assigned training subset was used to validate the model within the InCHIANTI cohort *in silico* (via simulation).

Calibration of the model

Parameter estimates for each predictive factor and associated descriptive statistics was evaluated to provide biological insight into the underpinnings of the classification algorithm. We first evaluated the calibration by partitioning the data into 5, 10, 20, 30, 40, 50, 75, 100 and 200 groups and then ran the calibration test. Next, we repeated tests for all possible values between 5-200 groups and evaluated the distribution of the test statistic. The best prediction

thresholds were determined using AUC, 87.7% for Model I and 86.4% for Model II. Population predictive features by phenotype ranked by gain for Model I are presented in Tables 4-6 and Model II Tables 7-9.

Genetic Data

Genotypic data was generated at the National Institute on Aging's Laboratory of Neurogenetics. Samples of genomic DNA extracted from leukocytes¹⁷. Genotypic data used for the model were extracted out of the binary Plink files from the InCHIANTI database. SNPs which could not be identified in the binary files were extracted from genotype imputed files, genotype imputation was completed with Minimac (V2). The SNPs included meet the following standard: per variant and per sample missingness < 5%, European ancestry, MAF < 0.001 and a r_{sq} < 0.3. Additionally, Samples were filtered for 95% or greater genotyping call rate, no ancestry outliers, and no sex discrepancies.

Supplementary Data Table I: Laboratory values as they appear in the InCHIANTI Datasets by Clinical Category

Inflammatory/Immunity	Nutrient Biomarker	Lipid Metabolism
BL Uric acid (mg/dL)	BL Omega-3 fatty acids as % of total fatty acid area	BL Lipids: total cholesterol (mg/dL)
BL Urinary cortisol (μ g/mL)	BL Omega-3 plasma fatty acid weight (mg/L)	BL Lipids: HDL cholesterol (mg/dL)
BL 24-hour urinary cortisol (μ g/24 hours)	BL Omega-3 fatty acids as % of total fatty acid weight	BL Lipids: triglycerides (mg/dL)
BL C-reactive protein - low sensitivity (μ g/mL)	BL Omega-3 fatty acids as % of total fatty acid mols	BL Lipids: LDL cholesterol (mg/dL)
BL C-reactive protein - high sensitivity (μ g/mL)	BL Omega-6 fatty acids as % of total fatty acid area	BL Lipoprotein(a) (mg/dL)
BL Interleukin-6 via ELISA ultrasensitive (pg/mL)	BL Omega-6 plasma fatty acid weight (mg/L)	
BL IL-6 high-sensitivity ELISA calculated from ELISA ultrasensitive (pg/mL)	BL Omega-6 fatty acids as % of total fatty acid weight	Metabolomics(plasma lipids)
BL Soluble IL-6 receptor via ELISA (ng/mL)	BL Omega-6 fatty acids as % of total fatty acid mols	BL Fatty acid C16:0 (palmitic) area
BL Interleukin-10 via ELISA (pg/mL)	BL Ratio of Omega-6:Omega-3 as % of total fatty acid area	BL Fatty acid C16:0 (palmitic) area
BL Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	BL Ratio of Omega-6:Omega-3 as % of total fatty acid weight	BL Fatty acid C16:0 as % of total fatty acid area
BL Interleukin-1B via ELISA (pg/mL)	BL Ratio of Omega-6:Omega-3 as % of total fatty acid mols	BL Fatty acid C16:0 weight (mg/L)

BL Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	BL Vitamin B6 via high performance liquid chromatography (ng/mL)	BL Fatty acid C16:0 as % of total fatty acid weight
BL Transforming growth factor-B1 (pg/mL)	BL Vitamin B6 via high performance liquid chromatography (nmol/L)	BL Fatty acid C16:0 (µmol/L)
BL Tumor necrosis factor-α via multiplex technology (pg/mL)	BL Vitamin E gamma tocopherol, high performance liquid chromatography (µmol/L)	BL Fatty acid C16:0 as % of total fatty acid mols
BL Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	BL Vitamin E alpha tocopherol, high performance liquid chromatography (µmol/L)	BL Fatty acid C20:0 (arachidic) area
BL Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	BL Vitamin E gamma tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	BL Fatty acid C20:0 as % of total fatty acid area
BL TNF-related apoptosis-inducing ligand (pg/mL)	BL Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	BL Fatty acid C20:0 weight (mg/L)
BL Interleukin-8 via Bio-Plex (pg/mL)	BL Beta-carotene via high performance liquid chromatography (µmol/L)	BL Fatty acid C20:0 as % of total fatty acid weight
BL Interleukin-12 via Bio-Plex (pg/mL)	BL Lycopene via high performance liquid chromatography (µmol/L)	BL Fatty acid C20:0 (µmol/L)
BL Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	BL Total proteins (g/dL)	BL Fatty acid C20:0 as % of total fatty acid mols
BL Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	BL Albumin (%)	BL Fatty acid C20:5 n-3 cis (eicosapentaenoic, EPA) area
BL Serum cortisol (µg/dL)		BL Fatty acid C20:5 n-3 as % of total fatty acid area
BL Serum cortisol (nmol/L)		BL Fatty acid C20:5 n-3 weight (mg/L)
BL Dehydroepiandrosterone sulfate (µg/dL)		BL Fatty acid C20:5 n-3 as % of total fatty acid weight
BL Dehydroepiandrosterone sulfate (nmol/L)		BL Fatty acid C20:5 n-3 (µmol/L)
BL Cortisol:DHEAS ratio (based on nmols)		BL Fatty acid C20:5 n-3 as % of total fatty acid mols
BL Soluble CD14 via ELISA (ng/mL)		BL Fatty acid C22:0 (behenic) area
BL Fibrinogen (mg/dL)		BL Fatty acid C22:0 as % of total fatty acid area
BL Erythrocyte sedimentation rate (ESR) (mm/hour)		BL Fatty acid C22:0 weight (mg/L)

BL Homocysteine via FPIA analysis ($\mu\text{mol/L}$)		BL Fatty acid C22:0 as % of total fatty acid weight
BL Resistin via EIA (ng/mL)-		BL Fatty acid C22:0 ($\mu\text{mol/L}$)
BL Adiponectin via RIA ($\mu\text{g/mL}$)- (metabolic function)		BL Fatty acid C22:0 as % of total fatty acid mols
BL Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)		BL Fatty acid C24:0 (lignoceric) area
BL Alpha-1 globulin (%)		BL Fatty acid C24:0 as % of total fatty acid area
BL Alpha-2 globulin (%)		BL Fatty acid C24:0 weight (mg/L)
BL Alpha-2-macroglobulin (mg/dL)		BL Fatty acid C24:0 as % of total fatty acid weight
BL Beta globulins (%)		BL Fatty acid C24:0 ($\mu\text{mol/L}$)
BL Endogenous secretory receptor for AGEs (ng/mL)		BL Fatty acid C24:0 as % of total fatty acid mols
Renal/Electrolyte	Hematology/Liver	Endocrine/Hormones
BL Na ⁺ (mEq/L)	BL White blood cells (WBC) (n, $\text{K}/\mu\text{L}$)	BL Blood glucose (mg/dL)
BL Ca ⁺⁺ (mg/dL)	BL Neutrophils (n, $\text{K}/\mu\text{L}$)	BL 25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)
BL Urinary creatinine (mg/dL)	BL Lymphocytes (n, $\text{K}/\mu\text{L}$)	BL Parathyroid hormone, two-site immunoradiometric assay (pg/mL)
BL 24-hour urinary creatinine ($\text{mg}/24$ hours)	BL Monocytes (n, $\text{K}/\mu\text{L}$)	BL Thyroid stimulating hormone, TSH (mIU/L)
BL Creatinine clearance, 24-hr urine (mL/minute)	BL Neutrophils (%)	BL Free thyroxine, fT4 (ng/dL)
BL Urinary Ca (mmol/L)	BL Lymphocytes (%)	BL Plasma insulin via RIA (mIU/L)
BL Urinary Na (mmol/L)	BL Monocytes (%)	BL Total testosterone (ng/mL)
BL Urine glucose (mg/dL)	BL Red blood cells (RBC) (n, millions/ μL)	BL Total testosterone (nmol/L)
BL Urine proteins (mg/dL)	BL Hemoglobin (g/dL)	BL Free testosterone (ng/dL), Vermeulen
BL Urine hemoglobin (mg/dL)	BL Hematocrit (%)	BL Free testosterone (nmol/L), Vermeulen
BL Urine ketones (mg/dL)	BL Mean corpuscular volume (MCV) (fL)	BL Estradiol via radioimmunoassay (pg/mL)
BL Urine bilirubin (mg/dL)	BL Mean corpuscular hemoglobin (MCH) (pg)	BL Estradiol via radioimmunoassay (nmol/L)
BL Urine urobilinogen (mg/dL)	BL MCH concentration (MCHC) (g/dL)	BL C-terminal telopeptide of type-1 collagen (ng/mL)

BL Urine nitrites	BL Red cell distribution width (RDW) (%)	BL Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)-(IGFBP1)
BL Serum creatinine (mg/dL)	BL Mean platelet volume (MPV) (fL)	BL IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***
BL Blood urea nitrogen (mg/dL)	BL Ferritin (ng/mL)	BL IGF binding protein-3, serum, immunoradiometric assay (nmol/L)
BL Creatine phosphokinase (U/L)	BL Folate via RIA (ng/mL)	
BL Cystatin C (mg/L)	BL Folate via RIA (nmol/L)	
	BL Vitamin B12 via RIA (pg/mL)	
	BL Vitamin B12 via RIA (pmol/L)	
	BL Methylmalonic acid(methylmalonic aciduria), MMA (μmol/L)	
	BL Soluble transferrin receptor (nmol/L)	
	BL Soluble transferrin receptor (mg/L)	
	BL GOT (also known as AST) (U/L)	
	BL GPT (also known as ALT) (U/L)	
	BL Gamma glutamyl transferase (U/L)	
	BL Retinol via high performance liquid chromatography (μmol/L)	

Supplementary Data Table II: Variants included in the Genomic Risk Score GRS calculations and individual effect estimates of single variants for predictive modeling. Phenotype association is based on the findings from the systematic review and the relationship found between variant and disease outcome.

Variant Name-Allele	Allele Frequency (%)	Gene/Closest RefSeq Gene	Variant Detail-dbSNP	Phenotype Association
rs1048945_C	1.3	APEX1	rs1048945 C/G Ancestral: G Minor: C	Cognition
rs1052133_G	20.6	OGG1	rs1052133 C/G Ancestral: C Minor: G	Cognition
rs1064039_T	19.0	CST3	rs1064039 A/G Ancestral: G Minor: T	Cognition
rs10793294_C	21.7	GAB2	rs10793294 A/C Ancestral: G Minor: C	Cognition
rs10883631_G	48.4	BTRC	rs10883631 A/G Ancestral: G Minor: A	Frail
rs10883642_G	48.4	BTRC	rs10883642 A/G Ancestral: A Minor: A	Frail
rs11225434_C	47.9	WTAPP1	rs11225434 C/T Ancestral: T Minor: C	Cognition
rs113263161_A	10.4	CCRL2/LOC102724297	rs113263161 A/G Ancestral: G Minor: A	Cognition
rs1133174_A	41.0	SORL1	rs1133174 A/G Ancestral: G Minor: A	Cognition
rs11574428_A	10.2	CCRL2	rs11574428 A/T Ancestral: T Minor: A	Cognition
rs11575821_A	11.4	CCRL2/LOC102724297	rs11575821 A/G Ancestral: G Minor: A	Cognition
rs1207568_A	19.4	KLOTHO	rs1207568 C/T Ancestral: C Minor: A	Cognition
rs13113697_T	27.2	HS3ST1/LOC107986178	rs13113697 G/T Ancestral: G Minor: T	Cognition
rs1468063_T	12.4	FAS	rs1468063 A/G Ancestral: G Minor: T	Cognition
rs1566728_C	14.1	PTPRJ	rs1566728 A/G Ancestral: G Minor: C	Frail
rs16944_A	33.4	IL1B	rs16944 A/G Ancestral: A Minor: A	Cognition
rs1799990_G	30.9	PRNP	rs1799990 A/G Ancestral: A Minor: G	Cognition
rs1800629_A	12.3	TNF	rs1800629 A/G Ancestral: G Minor: A	Cog/Frail
rs1800764_C	47.6	ACE	rs1800764 C/T Ancestral: C Minor: T	Cognition
rs1800796_C	5.0	IL6	rs1800796 C/G Ancestral: G Minor: C	Cog/Frail
rs1801394_G	43.9	MTRR	rs1801394 A/G Ancestral: A Minor: G	Frail
rs2047812_A	14.8	PTPRJ	rs2047812 C/T Ancestral: C Minor: A	Frail
rs2227729_G	7.5	VTN	rs2227729 C/T Ancestral: C Minor: G	Frail
rs2228145_C	38.0	IL6-R	rs2228145 A/C/T Ancestral: A Minor: C	Cognition
rs2228467_C	8.2	CCL4	rs2228467 C/T Ancestral: T Minor: C	Cognition
rs2229238_T	16.9	IL6-R	rs2229238 C/T Ancestral: C Minor: T	Cognition
rs2267163_T	36.5	TCN2	rs2267163 C/T Ancestral: C Minor: T	Frail
rs2283368_C	12.3	KLOTHO	rs2283368 C/T Ancestral: T Minor: C	Cognition
rs2465481_A	47.0	GNAI1	rs2465481 C/T Ancestral: C Minor: A	Cognition
rs2714465_G	45.0	GNAI1	rs2714465 A/G Ancestral: A Minor: G	Cognition
rs3092960_A	10.7	CCR2	rs3092960 A/G Ancestral: G Minor: A	Cognition
rs3131609_C	32.8	USP50	rs3131609 A/G Ancestral: A Minor: C	Cognition
rs360722_A	16.9	IL18	rs360722 C/T Ancestral: T Minor: A	Frail
rs3865444_A	27.1	CD33	rs3865444 G/T Ancestral: G Minor: A	Cognition
rs4147929_A	19.3	ABCA7	rs4147929 A/G Ancestral: G Minor: A	Cognition
rs429358_C	6.9	APOE	rs429358 C/T Ancestral: C Minor: C	Cognition
rs4316_T	38.1	ACE	rs4316 C/T Ancestral: C Minor: T	Cognition
rs4845622_C	38.6	IL6R	rs4845622 A/C Ancestral: A Minor: C	Cognition
rs4968782_G	41.0	ACE	rs4968782 A/G Ancestral: G Minor: G	Cognition
rs55636820_A	6.0	BIN1	rs55636820 A/G Ancestral: G Minor: A	Cognition
rs562020_A	34.6	KLOTHO	rs562020 C/T Ancestral: T Minor: A	Cognition
rs573521_A	47.2	MMP3	rs573521 C/T Ancestral: C Minor: A	Cognition
rs5744256_G	18.3	IL18	rs5744256 C/T Ancestral: T Minor: G	Frail
rs603050_T	31.3	WTAPP1	rs603050 A/G Ancestral: G Minor: T	Cognition
rs611646_T	48.6	ATM	rs611646 A/T Ancestral: A Minor: A	Frail

rs61812598_A	37.9	IL6-R	rs61812598 A/G Ancestral: G Minor: A	Cognition
rs6441977_A	10.2	CCRL2	rs6441977 A/G Ancestral: G Minor: A	Cognition
rs650108_A	30.1	MMP3	rs650108 A/G Ancestral: G Minor: A	Cognition
rs6762266_C	10.4	CCRL2	rs6762266 C/T Ancestral: T Minor: C	Cognition
rs679620_T	46.7	MMP3	rs679620 A/G Ancestral: G Minor: T	Cognition
rs6808835_T	10.5	CCRL2	rs6808835 G/T Ancestral: T Minor: T	Cognition
rs7110631_C	31.2	PICALM	rs7110631 C/G Ancestral: G Minor: C	Cognition
rs7396366_C	36.0	AP2A2	rs7396366 G/T Ancestral: T Minor: C	Cognition
rs7412_T	6.6	APOE	rs7412 C/T Ancestral: C Minor: T	Cognition
rs7497104_T	28.6	MYO9A	rs7497104 C/T Ancestral: T Minor: T	Cognition
rs7926920_A	46.9	WTAPP1	rs7926920 A/G Ancestral: G Minor: A	Cognition
rs9267487_C	6.5	DDX39B	rs9267487 C/T Ancestral: T Minor: C	Frail
rs9349407_C	24.5	CD2AP	rs9349407 C/G Ancestral: G Minor: C	Cognition
rs948399_C	26.9	MMP3	rs948399 C/T Ancestral: T Minor: C	Cognition
rs9527025_C	14.8	KLOTHO	rs9527025 C/G Ancestral: C Minor: C	Cognition
rs3219484_T	3.8	MUTYH	rs3219484_A/G Ancestral: G Minor: T	Cognition
rs12752888_C	26.8	ACOT11/LOC105378734	rs12752888 C/T Ancestral: T Minor: C	Cognition
rs1539053_A	45.6	DAB1	rs1539053 C/T Ancestral: T Minor: G	Cognition
rs3811448_A	19.3	TDRD10	rs3811448 A/G Ancestral: A Minor: A	Cognition
rs4129267_T	37.9	IL6-R	rs4129267 C/T Ancestral: C Minor: T	Cognition
rs915179_G	36.0	LMNA	rs915179 A/G Ancestral: G Minor: A	Cognition
rs9919256_A	13.7	LMNA	rs9919256 A/G Ancestral: A Minor: A	Cognition
rs6131_T	19.4	SELP	rs6131 A/G Ancestral: A Minor: T	Frail
rs3818361_A	19.5	CR1	rs3818361 C/T Ancestral: C Minor: A	Cognition
rs1260326_C	46.3	GCKR	rs1260326 C/T Ancestral: C Minor: T	Frail
rs744373_G	28.2	BIN1	rs744373 C/T Ancestral: T Minor: G	Cognition
rs7561528_A	31.2	BIN1/LOC105373605	rs7561528 A/G Ancestral: A Minor: A	Cognition
rs11894266_C	43.5	SSB	rs11894266 C/T Ancestral: C Minor: T	Cognition
rs6747918_A	49.2	CASP8	rs6747918 A/G Ancestral: A Minor: A	Frail
rs2929408_A	22.4	KAT2B	rs2929408 G/T Ancestral: G Minor: A	Frail
rs737267_T	25.6	SLC2A9	rs737267 A/G/T Ancestral: G Minor: T	Frail
rs9461448_G	4.7	PGBD1	rs9461448 G/T Ancestral: T Minor: G	Cognition
rs9446432_C	8.2	C6orf155	rs9446432 C/T Ancestral: T Minor: C	Cognition
rs9384428_C	32.5	MIR1202/LOC101928923	rs9384428 C/T Ancestral: T Minor: C	Cognition
rs4646450_A	16.4	CYP3A5	rs4646450 C/T Ancestral: T Minor: A	Frail
rs11767557_C	16.8	EPHA1-AS1	rs11767557 C/T Ancestral: T Minor: C	Cognition
rs11771145_A	32.9	EPHA1-AS1	rs11771145 A/G Ancestral: A Minor: A	Cognition
rs11136000_T	39.0	CLU	rs11136000 C/T Ancestral: T Minor: T	Cognition
rs1157242_T	16.2	KCNU1	rs1157242 A/G Ancestral: G Minor: T	Cognition
rs7840202_C	29.9	UBR5	rs7840202 A/C Ancestral: C Minor: C	Cognition
rs7920721_G	39.4	ECHDC3	rs7920721 A/G Ancestral: A Minor: G	Cognition
rs7905675_A	34.9	TFAM	rs7905675 A/G Ancestral: A Minor: G	Cognition
rs17117126_G	9.5	CH25H	rs17117126 A/G Ancestral: G Minor: G	Cognition
rs6265_T	21.6	BDNF	rs6265 A/G Ancestral: G Minor: T	Cognition
rs1566729_T	14.1	PTPRJ	rs1566729 A/G Ancestral: G Minor: T	Frail
rs583791_C	49.5	MS4A6A	rs583791 A/G Ancestral: G Minor: C	Cognition
rs610932_T	48.5	MS4A6A	rs610932 A/C Ancestral: A Minor: T	Cognition
rs662196_C	49.6	MS4A6A	rs662196 A/G Ancestral: G Minor: C	Cognition
rs670139_T	31.2	MS4A4E	rs670139 A/C/T Ancestral: C Minor: T	Cognition
rs676309_C	31.1	MS4A4E	rs676309 A/G Ancestral: A Minor: C	Cognition

rs11827375_A	10.5	C11orf30	rs11827375 A/G Ancestral: G Minor: A	Cognition
rs3851179_T	36.0	PICALM	rs3851179 A/G Ancestral: G Minor: T	Cognition
rs541458_C	31.6	PICALM	rs541458 C/T Ancestral: T Minor: C	Cognition
rs10501927_G	23.6	CNTN5	rs10501927 G/T Ancestral: T Minor: G	Cognition
rs495366_A	30.1	WTAPP1	rs495366 A/G Ancestral: G Minor: A	Cognition
rs645419_A	46.7	MMP3	rs645419 A/G Ancestral: G Minor: A	Cognition
rs10502262_T	27.7	SORL1	rs10502262 A/G Ancestral: G Minor: T	Cognition
rs1614735_G	47.6	SORL1	rs1614735 G/T Ancestral: T Minor: G	Cognition
rs2298813_A	4.0	SORL1	rs2298813 A/G Ancestral: G Minor: A	Cognition
rs3781835_A	2.3	SORL1	rs3781835 A/G Ancestral: G Minor: A	Cognition
rs4935774_C	20.5	SORL1	rs4935774 C/T Ancestral: C Minor: C	Cognition
rs4363657_C	15.2	SLCO1B1	rs4363657 C/T Ancestral: T Minor: C	Frail
rs1799986_T	17.4	LRP1	rs1799986 A/C/T Ancestral: C Minor: T	Frail
rs398655_C	45.0	KLOTHO	rs398655 G/T Ancestral: G Minor: A	Cognition
rs648202_T	13.7	KLOTHO	rs648202 C/T Ancestral: C Minor: T	Cognition
rs9526984_G	7.4	KLOTHO	rs9526984 A/G Ancestral: A Minor: G	Cognition
rs9527024_A	14.8	KLOTHO	rs9527024 A/G Ancestral: A Minor: A	Cognition
rs9536314_G	14.7	KLOTHO	rs9536314 A/G/T Ancestral: T Minor: G	Cognition
rs2287396_T	17.7	GSTZ1	rs2287396 C/T Ancestral: C Minor: T	Frail
rs7175373_C	29.1	MYO9A	rs7175373 A/C/G Ancestral: C Minor: C	Cognition
rs129968_A	39.8	CREBBP	rs129968 A/G Ancestral: A Minor: G	Frail
rs3785880_G	39.8	MAPT	rs3785880 G/T Ancestral: T Minor: G	Cognition
rs2526378_G	46.8	TSPOAP1	rs2526378 C/T Ancestral: C Minor: A	Cognition
rs4343_A	40.1	ACE	rs4343 A/G Ancestral: A Minor: G	Cognition
rs4459609_C	40.9	ACE	rs4459609 A/C Ancestral: A Minor: C	Cognition
rs3764650_G	11.8	ABCA7	rs3764650 G/T Ancestral: T Minor: G	Cognition
rs157580_G	39.1	TOMM40	rs157580 A/G Ancestral: G Minor: G	Cognition
rs2075650_G	7.5	TOMM40	rs2075650 A/G Ancestral: G Minor: G	Cognition
rs405509_T	42.8	APOE	rs405509 A/C Ancestral: C Minor: T	Cognition
rs597668_C	12.0	EXOC3L2	rs597668 C/T Ancestral: C Minor: C	Cognition
rs6859_A	38.8	NECTIN2	rs6859 A/G Ancestral: G Minor: A	Cognition
rs8106922_G	44.8	TOMM40	rs8106922 A/G Ancestral: A Minor: G	Cognition
rs17411904_C	7.7	PCK1	rs17411904 C/T Ancestral: T Minor: C	Cognition
rs2833383_T	27.9	TIAM1	rs2833383 C/T Ancestral: C Minor: T	Frail
rs4646316_T	27.7	COMT	rs4646316 C/T Ancestral: C Minor: T	Frail
rs4680_A	46.4	COMT	rs4680 C/T Ancestral: G Minor: A	Cognition
rs740234_G	24.2	TCN2	rs740234 C/T Ancestral: T Minor: G	Frail

Notes: *Proxy SNP, Cog/Frail – variant was found for both phenotypes in the systematic review, bold text indicates the closest gene

Genetic risk scores

One hundred and thirty-one variants were catalogued from a large systematic review and used to construct genetic risk scores for three models. All variants were used to create an all risk score (n=132), variants related to the phenotypes cognitive decline and physical frailty constructed cognitive risk scores (n=105) and frailty risk scores (n=27). Risk scores were calculated by summation of the number of risk alleles across all the variants divided by the number of SNPs in the score to obtain an average number of risk alleles per locus. After the scaled risk allele counts were summed and divided by the number of loci, they were transformed into Z scores. Z score transformation assists in communicating the effect estimates with the Z corresponding to a single standard deviation from the control mean genetic risk for the phenotypes. All risk scores were calculated using PLINK. R V. 3.2.1 was used to fit multinomial and logistic regression models using standard covariates and risk scores as predictors of cognitive decline, physical frailty, and cognitive frailty as the outcome variable. Stepwise backward and forward selection using AIC and p values facilitated the best fit models.

Supplementary Table III:

Model I Genetic risk scores – Population predictive model features by phenotype

Phenotype (n)		All Risk Scores	Cognition Risk Scores	Frail Risk Scores
Cognitive Decline MMSE (369)	p	.1286	.0659	.8768
	β	.12	.15	-.01
	SE	.08	.08	.08
Frail CHS (595)	p	.0488	.0401	.6509
	β	0.14	.14	.03
	SE	0.07	.07	.07
Cognitive frailty MMSE (257)	p	.0455	.0479	.7775
	β	0.19	.19	-0.03
	SE	0.10	.10	.09

Model II Genetic risk scores – Population predictive model features by phenotype

Phenotype (n)		All Risk Scores	Cognition Risk Scores	Frail Risk Scores
Cognitive Decline Trail B (634)	p	.6097	.5959	.4440
	β	.05	.05	-.07
	SE	.09	.09	.09
Cognitive Decline Trail A (525)	p	.0351	.0370	.3274
	β	.16	.16	.07
	SE	.08	.07	.07
Cognitive Frailty Trail B (325)	p	.2082	.1992	.7394
	β	.11	.11	.03
	SE	.08	.09	.08
Cognitive Frailty Trail A (302)	p	.6298	.4242	.2734
	β	.04	.06	-.08
	SE	.08	.08	.08

Table IV: Cognitive Decline Features Model I

Cognitive Decline Features	Gain	Cover	Frequency	Importance
Age	0.247016911	0.117594993	0.048543689	0.247016911
Level of Education	0.160608946	0.187763494	0.097087379	0.160608946
TNF-related apoptosis-inducing ligand	0.025564877	0.029595722	0.029126214	0.025564877
24-hour urinary creatinine	0.02142342	0.016832239	0.014563107	0.02142342
Fatty acid C16:0 as % of total fatty acid area	0.018881415	0.036076992	0.024271845	0.018881415
Dx Depression	0.018852494	0.021616819	0.019417476	0.018852494
Cystatin C	0.017786637	0.035118001	0.024271845	0.017786637
Dehydroepiandrosterone sulfate	0.01624112	0.0178982	0.014563107	0.01624112
Adiponectin via RIA	0.0154553	0.018357822	0.019417476	0.0154553
Beta-carotene via high performance liquid chromatography (µmol/L)	0.015327296	0.014660641	0.019417476	0.015327296
Mean corpuscular volume (MCV)	0.013822196	0.015698515	0.024271845	0.013822196
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	0.012997192	0.007511319	0.019417476	0.012997192
Free thyroxine fT4 (ng/dL)	0.012865081	0.006865344	0.014563107	0.012865081
Fibrinogen (mg/dL)	0.010709775	0.023801966	0.014563107	0.010709775
Lymphocytes	0.010046812	0.008887038	0.014563107	0.010046812
Red cell distribution width (RDW) (%)	0.008983628	0.003876676	0.009708738	0.008983628
Interleukin-12	0.008595815	0.002439156	0.009708738	0.008595815
Fatty acid C16:0 (palmitic) area	0.008348097	0.00299276	0.009708738	0.008348097
Fatty acid C20:0 (arachidic) area	0.00819782	0.01193878	0.014563107	0.00819782
Lipids: HDL cholesterol	0.007781061	0.013694142	0.019417476	0.007781061
Fatty acid C20:5 n-3 as % of total fatty acid weight	0.007714811	0.016204393	0.009708738	0.007714811
Ferritin	0.007209597	0.010436238	0.009708738	0.007209597
Gender	0.007151353	0.00641112	0.009708738	0.007151353
Fatty acid C24:0 as % of total fatty acid weight	0.007083429	0.010963022	0.014563107	0.007083429
24-hour urinary cortisol (µg/24 hours)	0.006833969	0.003491305	0.009708738	0.006833969
Creatinine clearance 24-hr urine	0.006443681	0.007022278	0.009708738	0.006443681
Fatty acid C20:0 weight (mg/L)	0.006025751	0.005788322	0.009708738	0.006025751
Vitamin E gamma tocopherol high performance liquid chromatography	0.005902007	0.00573116	0.009708738	0.005902007
Soluble IL-6 receptor via ELISA	0.005647251	0.002059633	0.004854369	0.005647251
Cortisol:DHEAS ratio	0.005261354	0.019818409	0.014563107	0.005261354
Methylmalonic acid MMA (µmol/L)"	0.005213763	0.001448349	0.009708738	0.005213763
Resistin via EIA	0.00521251	0.006015127	0.009708738	0.00521251
Plasma insulin via RIA	0.005080217	0.003755668	0.009708738	0.005080217
Creatine phosphokinase	0.004950723	0.001190354	0.004854369	0.004950723
Homocysteine via FPIA analysis	0.004917852	0.007184674	0.004854369	0.004917852
Interleukin-10 via ELISA	0.004745208	0.00376727	0.004854369	0.004745208
Fatty acid C24:0 (lignoceric) area	0.004584681	0.00330742	0.004854369	0.004584681
Red blood cells	0.004528429	0.011768391	0.009708738	0.004528429
Fatty acid C20:5 n-3 weight (mg/L)	0.004501496	0.001118337	0.004854369	0.004501496
Estradiol via radioimmunoassay (pg/mL)	0.00425931	0.000629884	0.004854369	0.00425931
Vitamin B12 via RIA	0.004252471	0.018127967	0.009708738	0.004252471
BL Omega-3 plasma fatty acid weight (mg/L)	0.004221882	0.002369331	0.004854369	0.004221882
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.004149518	0.014263272	0.009708738	0.004149518
Fatty acid C24:0 as % of total fatty acid area	0.004102069	0.005203763	0.004854369	0.004102069
Ratio of Omega-6:Omega-3 as % of total fatty acid area	0.004049393	0.008024251	0.009708738	0.004049393

Urinary Na	0.003793497	0.000890377	0.009708738	0.003793497
Alpha-2-macroglobulin	0.003641408	0.01779343	0.009708738	0.003641408
Lipids: triglycerides (mg/dL)	0.003635412	0.000947582	0.009708738	0.003635412
rs3131609_C	0.003539956	0.002923127	0.004854369	0.003539956
rs1800796_C	0.003487553	0.001642754	0.004854369	0.003487553
Lycopene via high performance liquid chromatography	0.003366475	0.001272687	0.004854369	0.003366475
Soluble TNF-a receptor I via quantitative sandwich EIA	0.003259691	0.001869279	0.004854369	0.003259691
Albumin (%)	0.003252256	0.000885554	0.004854369	0.003252256
MCH concentration (MCHC) (g/dL)	0.003207126	0.001501542	0.004854369	0.003207126
C-terminal telopeptide of type-1 collagen	0.003132828	0.001238391	0.004854369	0.003132828
Alpha-1 globulin	0.003093762	0.001077661	0.004854369	0.003093762
Alpha-2 globulin (%)	0.002957393	0.001951717	0.004854369	0.002957393
Urinary cortisol	0.002954808	0.005640344	0.004854369	0.002954808
Lipoprotein(a)	0.002822909	0.004583778	0.004854369	0.002822909
BL Blood glucose (mg/dL)	0.002796243	0.005340663	0.009708738	0.002796243
Anticollnergetic Burden	0.002789313	0.009621728	0.004854369	0.002789313
rs2228145_C	0.002741554	0.001278252	0.004854369	0.002741554
BL Ratio of Omega-6:Omega-3 as % of total fatty acid mols	0.002656076	0.004937615	0.004854369	0.002656076
Blood urea nitrogen	0.002558617	0.00897141	0.004854369	0.002558617
Parathyroid hormone two-site immunoradiometric assay "	0.002550438	0.000968729	0.004854369	0.002550438
Serum cortisol	0.00249517	0.002893942	0.004854369	0.00249517
Lipids: total cholesterol	0.002460557	0.001170389	0.004854369	0.002460557
Fatty acid C22:0 (behenic) area	0.002339895	0.00187473	0.004854369	0.002339895
Vitamin E alpha tocopherol high performance liquid chromatography	0.002198081	0.005823143	0.004854369	0.002198081
Urinary Ca	0.002164721	0.000730392	0.004854369	0.002164721
Folate via RIA	0.002113781	0.00106923	0.004854369	0.002113781
Monocytes (%)	0.00199266	0.000547857	0.004854369	0.00199266
Total proteins (g/dL)	0.001944932	0.007589282	0.004854369	0.001944932
rs948399_C	0.001742443	0.001020202	0.004854369	0.001742443
Omega-6 plasma fatty acid weight (mg/L)	0.001653381	0.001473014	0.004854369	0.001653381
rs10883631_G	0.001571027	0.000810486	0.004854369	0.001571027
White blood cells (WBC)	0.001509285	0.000889183	0.004854369	0.001509285
ALT	0.001401955	0.000216533	0.004854369	0.001401955
Fatty acid C20:0 as % of total fatty acid weight	0.001362604	0.007402723	0.004854369	0.001362604
Interleukin-18 via ELISA ultrasensitive using plasma	0.001317785	0.000289673	0.004854369	0.001317785
rs7396366_C	0.001163588	0.000555404	0.004854369	0.001163588
Gamma glutamyl transferase	0.000800672	0.002452777	0.004854369	0.000800672
Fatty acid C22:0	0.000561923	0.000672233	0.004854369	0.000561923
Fatty acid C16:0 as % of total fatty acid weight	0.000554415	0.000510348	0.004854369	0.000554415
Uric acid	0.000537078	0.000186889	0.004854369	0.000537078
rs2075650_G	0.000487925	0.000597468	0.004854369	0.000487925
Thyroid stimulating hormone	0.000404183	0.000235127	0.004854369	0.000404183
rs4646316_T	0.000302482	0.00026136	0.004854369	0.000302482
Mean corpuscular hemoglobin	0.000184601	0.002295018	0.004854369	0.000184601
Interleukin-6 via ELISA ultrasensitive	6.12E-05	0.000353843	0.004854369	6.12E-05

Table V: Frailty Features Model I

Frailty Features	Gain	Cover	Frequency	Importance
Depression	0.098775745	0.084652562	0.027559055	0.098775745
Creatinine clearance, 24-hr urine (mL/minute)	0.052445936	0.033303749	0.011811024	0.052445936
Age	0.049474709	0.022604179	0.019685039	0.049474709
Anticholinergic Burden	0.030906413	0.030358874	0.023622047	0.030906413
Homocysteine via FPIA analysis (μmol/L)	0.024715965	0.023600409	0.007874016	0.024715965
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	0.023717251	0.022258754	0.015748031	0.023717251
Vitamin B6 via high performance liquid chromatography (ng/mL)	0.022823838	0.024226924	0.015748031	0.022823838
Interleukin-6 via ELISA ultrasensitive (pg/mL)	0.021999328	0.031654032	0.015748031	0.021999328
Dehydroepiandrosterone sulfate (μg/dL)	0.021750399	0.00821273	0.019685039	0.021750399
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	0.021744382	0.029219322	0.019685039	0.021744382
Erythrocyte sedimentation rate (ESR) (mm/hour)	0.019839941	0.028859155	0.019685039	0.019839941
Endogenous secretory receptor for AGEs (ng/mL)	0.018672872	0.019026448	0.023622047	0.018672872
24-hour urinary creatinine (mg/24 hours)	0.017065385	0.008624125	0.019685039	0.017065385
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	0.016999208	0.005822401	0.011811024	0.016999208
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.01625036	0.019029454	0.015748031	0.01625036
Lipoprotein(a) (mg/dL)	0.015535882	0.02537913	0.019685039	0.015535882
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (μmol/L)	0.015019838	0.023585828	0.015748031	0.015019838
Vitamin B12 via RIA (pg/mL)	0.014433258	0.012129297	0.019685039	0.014433258
Vitamin E gamma tocopherol, high performance liquid chromatography (μmol/L)	0.014300271	0.007085453	0.011811024	0.014300271
Folate via RIA (ng/mL)	0.014001884	0.011108339	0.011811024	0.014001884
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	0.013772642	0.017304651	0.015748031	0.013772642
Interleukin-1B via ELISA (pg/mL)	0.013205983	0.030090209	0.015748031	0.013205983
Beta-carotene via high performance liquid chromatography (μmol/L)	0.013116604	0.008019449	0.011811024	0.013116604
Creatine phosphokinase (U/L)	0.012919658	0.015168335	0.007874016	0.012919658
Plasma insulin via RIA (mIU/L)	0.011798838	0.011456837	0.011811024	0.011798838
Retinol via high performance liquid chromatography (μmol/L)	0.011423155	0.00461384	0.011811024	0.011423155
Methylmalonic acid, MMA (μmol/L)	0.011339303	0.008264588	0.015748031	0.011339303
Omega-6 fatty acids as % of total fatty acid area	0.011008005	0.004272542	0.019685039	0.011008005
Monocytes (n, K/μL)	0.010487533	0.018913621	0.011811024	0.010487533
Lipids: LDL cholesterol (mg/dL)	0.009591423	0.003739115	0.011811024	0.009591423
Tumor necrosis factor-α via multiplex technology (pg/mL)	0.009463133	0.006942978	0.011811024	0.009463133
Urinary Na (mmol/L)	0.009315233	0.004721585	0.011811024	0.009315233
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	0.009129694	0.013434801	0.015748031	0.009129694
Urinary Ca (mmol/L)	0.009022724	0.012591808	0.015748031	0.009022724
C-terminal telopeptide of type-1 collagen (ng/mL)	0.008609358	0.007960146	0.007874016	0.008609358
Interleukin-8 via Bio-Plex (pg/mL)	0.008566678	0.00594986	0.003937008	0.008566678
Fatty acid C24:0 as % of total fatty acid area	0.008095749	0.012731259	0.011811024	0.008095749
TNF-related apoptosis-inducing ligand (pg/mL)	0.007711265	0.003574424	0.011811024	0.007711265
Free testosterone (ng/dL), Vermeulen	0.007578292	0.011968667	0.015748031	0.007578292
Cystatin C (mg/L)	0.006550153	0.000712653	0.003937008	0.006550153
Na+ (mEq/L)	0.006516226	0.007030191	0.003937008	0.006516226
Monocytes (%)	0.00639573	0.01284515	0.011811024	0.00639573
Hematocrit (%)	0.00623512	0.006186332	0.003937008	0.00623512
24-hour urinary cortisol (μg/24 hours)	0.006090747	0.010514246	0.007874016	0.006090747
Interleukin-12 via Bio-Plex (pg/mL)	0.006015217	0.004092294	0.007874016	0.006015217
Blood glucose (mg/dL)	0.005694126	0.001798266	0.007874016	0.005694126
Soluble CD14 via ELISA (ng/mL)	0.0055483	0.001419796	0.003937008	0.0055483
Soluble IL-6 receptor via ELISA (ng/mL)	0.005477014	0.003230929	0.007874016	0.005477014
Fatty acid C24:0 as % of total fatty acid weight	0.005404971	0.002740247	0.003937008	0.005404971
Total testosterone (ng/mL)	0.005367844	0.01031343	0.003937008	0.005367844
rs948399_C	0.005345514	0.002189977	0.003937008	0.005345514

Urine proteins (mg/dL)	0.005280709	0.01350518	0.007874016	0.005280709
Neutrophils (n, K/ μ L)	0.005181902	0.000457207	0.003937008	0.005181902
Fatty acid C20:0 as % of total fatty acid weight	0.005052286	0.000994181	0.003937008	0.005052286
Serum cortisol (μ g/dL)	0.005017158	0.027936629	0.011811024	0.005017158
Level of Education	0.00493252	0.005284448	0.003937008	0.00493252
Red cell distribution width (RDW) (%)	0.004763014	0.005996977	0.003937008	0.004763014
Vitamin E gamma tocopherol, high performance liquid chromatography, assay #2 (μ mol/L)	0.004502593	0.002631694	0.003937008	0.004502593
Blood urea nitrogen (mg/dL)	0.004480007	0.006420977	0.011811024	0.004480007
Thyroid stimulating hormone, TSH (mIU/L)	0.004433716	0.00184381	0.007874016	0.004433716
rs10501927_G	0.004414743	0.009323168	0.007874016	0.004414743
Lipids: HDL cholesterol (mg/dL)	0.004247467	0.002294261	0.003937008	0.004247467
rs129968_A	0.004192118	0.003263095	0.007874016	0.004192118
Resistin via EIA (ng/mL)	0.0041092	0.006075858	0.003937008	0.0041092
Gamma glutamyl transferase (U/L)	0.004070941	0.011275891	0.007874016	0.004070941
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	0.004052691	0.001761794	0.003937008	0.004052691
Baseline diagnosis of Dementia	0.003942136	0.019120532	0.007874016	0.003942136
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	0.003824567	0.000791652	0.007874016	0.003824567
Urinary creatinine (mg/dL)	0.003755628	0.000396059	0.007874016	0.003755628
Ferritin (ng/mL)	0.003742311	0.008416166	0.007874016	0.003742311
C-reactive protein - high sensitivity (μ g/mL)	0.003565832	0.002077369	0.003937008	0.003565832
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	0.003520041	0.002588955	0.003937008	0.003520041
Lymphocytes (%)	0.003478691	0.000977976	0.003937008	0.003478691
Fatty acid C16:0 as % of total fatty acid weight	0.003475484	0.004796531	0.003937008	0.003475484
rs11225434_C	0.003330988	0.004787995	0.003937008	0.003330988
Fatty acid C16:0 (palmitic) area	0.003293404	0.000936887	0.003937008	0.003293404
Neutrophils (%)	0.003128871	0.00067737	0.003937008	0.003128871
Fatty acid C20:5 n-3 weight (mg/L)	0.003123782	0.001971299	0.003937008	0.003123782
Fatty acid C20:0 as % of total fatty acid area	0.003010934	0.002094985	0.003937008	0.003010934
GPT (also known as ALT) (U/L)	0.002937445	0.00592063	0.003937008	0.002937445
Albumin (%)	0.002854574	0.002198538	0.003937008	0.002854574
Mean platelet volume (MPV) (fL)	0.002770872	0.000419397	0.003937008	0.002770872
rs1539053_A	0.002756129	0.012007864	0.003937008	0.002756129
Cortisol:DHEAS ratio (based on nmols)	0.002566654	0.000898211	0.003937008	0.002566654
MCH concentration (MCHC) (g/dL)	0.002565159	0.010052336	0.003937008	0.002565159
Free thyroxine, fT4 (ng/dL)	0.002443825	0.007244682	0.007874016	0.002443825
Beta globulins (%)	0.002269127	0.000431623	0.003937008	0.002269127
Lipids: total cholesterol (mg/dL)	0.002175655	0.002408026	0.003937008	0.002175655
Fatty acid C20:0 weight (mg/L)	0.002119043	0.001718137	0.003937008	0.002119043
Estradiol via radioimmunoassay (pg/mL)	0.002044564	0.000316794	0.003937008	0.002044564
Fatty acid C22:0 weight (mg/L)	0.001960255	0.000352188	0.003937008	0.001960255
Lycopene via high performance liquid chromatography (μ mol/L)	0.001838456	0.002966245	0.003937008	0.001838456
Fatty acid C16:0 as % of total fatty acid area	0.001816434	0.010717187	0.003937008	0.001816434
Omega-6 plasma fatty acid weight (mg/L)	0.001762988	0.001987793	0.003937008	0.001762988
rs7840202_C	0.001405255	0.001059102	0.003937008	0.001405255
Hemoglobin (g/dL)	0.001237461	0.000333737	0.003937008	0.001237461
Gender	0.001217717	0.001564023	0.003937008	0.001217717
Omega-3 plasma fatty acid weight (mg/L)	0.001079396	0.000479765	0.007874016	0.001079396
Fatty acid C20:5 n-3 as % of total fatty acid area	0.001022163	0.000592968	0.003937008	0.001022163
White blood cells (WBC) (n, K/ μ L)	0.001016876	0.000724366	0.003937008	0.001016876
Fatty acid C24:0 (lignoceric) area	0.000965197	0.000281381	0.003937008	0.000965197
Fatty acid C24:0 weight (mg/L)	0.000951989	0.000337432	0.003937008	0.000951989
rs3865444_A	0.00059402	0.000268936	0.003937008	0.00059402
rs4935774_C	0.000287175	0.000250477	0.003937008	0.000287175
Mean corpuscular volume (MCV) (fL)	0.000204245	0.00046439	0.003937008	0.000204245

Table VI: Cognitive Frailty Features Model I

Cognitive Frailty Features	Gain	Cover	Frequency	Importance
Age	0.226782261	0.171557774	0.059322034	0.226782261
Baseline Diagnosis of Dementia	0.099398955	0.098462848	0.029661017	0.099398955
Level of Education	0.044233154	0.075160958	0.029661017	0.044233154
Depression	0.034553704	0.038157217	0.029661017	0.034553704
TNF-related apoptosis-inducing ligand (pg/mL)	0.03034655	0.028129602	0.033898305	0.03034655
24-hour urinary creatinine (mg/24 hours)	0.025460108	0.014329518	0.008474576	0.025460108
Fatty acid C24:0 as % of total fatty acid area	0.023009514	0.008399373	0.012711864	0.023009514
Fibrinogen (mg/dL)	0.015823506	0.009692466	0.021186441	0.015823506
24-hour urinary cortisol (µg/24 hours)	0.015266068	0.037498193	0.021186441	0.015266068
Lipids: HDL cholesterol (mg/dL)	0.014715469	0.011976521	0.016949153	0.014715469
Transforming growth factor-B1 (pg/mL)	0.014096962	0.022255665	0.016949153	0.014096962
Urinary cortisol (µg/mL)	0.014020579	0.027768583	0.021186441	0.014020579
Cystatin C (mg/L)	0.012966575	0.012313314	0.012711864	0.012966575
Blood urea nitrogen (mg/dL)	0.012798018	0.015478663	0.016949153	0.012798018
Anticholinergic Burden	0.012377409	0.015077936	0.029661017	0.012377409
Gender	0.011802004	0.013718017	0.021186441	0.011802004
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	0.011677247	0.005056691	0.016949153	0.011677247
Erythrocyte sedimentation rate (ESR) (mm/hour)	0.011479619	0.027685275	0.016949153	0.011479619
Creatine phosphokinase (U/L)	0.011252254	0.009828549	0.008474576	0.011252254
Serum cortisol (µg/dL)	0.009997371	0.006957488	0.012711864	0.009997371
Omega-6 fatty acids as % of total fatty acid area	0.009927552	0.003755798	0.004237288	0.009927552
Dehydroepiandrosterone sulfate (µg/dL)	0.009699038	0.004094407	0.008474576	0.009699038
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.009690317	0.011910741	0.008474576	0.009690317
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	0.008586559	0.00937805	0.016949153	0.008586559
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	0.008545621	0.011291322	0.016949153	0.008545621
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	0.007953886	0.004648823	0.012711864	0.007953886
Vitamin B6 via high performance liquid chromatography (ng/mL)	0.007871375	0.004486658	0.012711864	0.007871375
Soluble CD14 via ELISA (ng/mL)	0.00781868	0.006603146	0.008474576	0.00781868
Uric acid (mg/dL)	0.007707399	0.004887846	0.012711864	0.007707399
Fatty acid C20:0 as % of total fatty acid area	0.007346802	0.001933709	0.008474576	0.007346802
Thyroid stimulating hormone, TSH (mIU/L)	0.007266114	0.005429717	0.012711864	0.007266114
C-terminal telopeptide of type-1 collagen (ng/mL)	0.007200276	0.009389035	0.004237288	0.007200276
Urine proteins (mg/dL)	0.007174622	0.011835413	0.008474576	0.007174622
Total testosterone (ng/mL)	0.006692034	0.004245555	0.008474576	0.006692034
Resistin via EIA (ng/mL)	0.006665635	0.003066128	0.012711864	0.006665635
Hemoglobin (g/dL)	0.006538294	0.001687461	0.004237288	0.006538294
Gamma glutamyl transferase (U/L)	0.006461435	0.002914706	0.004237288	0.006461435
Fatty acid C24:0 as % of total fatty acid weight	0.006316549	0.006171081	0.008474576	0.006316549
Free thyroxine, FT4 (ng/dL)	0.006171355	0.00694256	0.008474576	0.006171355
Fatty acid C20:0 weight (mg/L)	0.006114046	0.003017798	0.008474576	0.006114046
Red cell distribution width (RDW) (%)	0.006079822	0.00257699	0.008474576	0.006079822
Cortisol:DHEAS ratio (based on nmols)	0.005840558	0.010952017	0.012711864	0.005840558
Vitamin E gamma tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	0.005830235	0.004808919	0.012711864	0.005830235
Monocytes (%)	0.005572667	0.003896981	0.008474576	0.005572667
rs1800796_C	0.005375181	0.001349969	0.004237288	0.005375181
MCH concentration (MCHC) (g/dL)	0.005308074	0.00983547	0.004237288	0.005308074
Fatty acid C22:0 (behenic) area	0.004956023	0.0045059	0.012711864	0.004956023
Vitamin E alpha tocopherol, high performance liquid chromatography (µmol/L)	0.004726676	0.010909678	0.008474576	0.004726676
Urine nitrites	0.004714047	0.002983739	0.004237288	0.004714047
Fatty acid C20:5 n-3 as % of total fatty acid weight	0.004676963	0.00545449	0.008474576	0.004676963

Interleukin-10 via ELISA (pg/mL)	0.004582369	0.003833224	0.008474576	0.004582369
rs7561528_A	0.004441844	0.002401485	0.004237288	0.004441844
Fatty acid C22:0 as % of total fatty acid area	0.004203919	0.001526901	0.004237288	0.004203919
Homocysteine via FPIA analysis (μmol/L)	0.004184592	0.001820163	0.004237288	0.004184592
Beta-carotene via high performance liquid chromatography (μmol/L)	0.004176619	0.001453662	0.008474576	0.004176619
Ferritin (ng/mL)	0.0041346	0.005952456	0.008474576	0.0041346
Plasma insulin via RIA (mIU/L)	0.004084085	0.005935122	0.012711864	0.004084085
Vitamin B12 via RIA (pg/mL)	0.00402366	0.00333813	0.004237288	0.00402366
Alpha-1 globulin (%)	0.003832509	0.014051952	0.004237288	0.003832509
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	0.003743268	0.00657122	0.008474576	0.003743268
Alpha-2 globulin (%)	0.00373557	0.004502258	0.004237288	0.00373557
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	0.003691119	0.002535915	0.008474576	0.003691119
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	0.003628467	0.007143239	0.004237288	0.003628467
C-reactive protein - high sensitivity (μg/mL)	0.003595036	0.004072452	0.004237288	0.003595036
rs3865444_A	0.003568828	0.006992619	0.004237288	0.003568828
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	0.003553408	0.003953066	0.008474576	0.003553408
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	0.003538457	0.00071027	0.008474576	0.003538457
rs12752888_C	0.003529126	0.003210877	0.004237288	0.003529126
rs1801394_G	0.002941323	0.009906123	0.004237288	0.002941323
Serum creatinine (mg/dL)	0.002671886	0.001135792	0.004237288	0.002671886
rs7840202_C	0.002603431	0.000660589	0.004237288	0.002603431
Endogenous secretory receptor for AGEs (ng/mL)	0.002571665	0.003716514	0.004237288	0.002571665
Soluble transferrin receptor (nmol/L)	0.002553661	0.003976923	0.004237288	0.002553661
Fatty acid C16:0 as % of total fatty acid weight	0.002533683	0.003256162	0.004237288	0.002533683
Retinol via high performance liquid chromatography (μmol/L)	0.002487013	0.004645502	0.004237288	0.002487013
Adiponectin via RIA (μg/mL)	0.002423759	0.003373917	0.004237288	0.002423759
Ca ⁺⁺ (mg/dL)	0.002412787	0.009697184	0.004237288	0.002412787
Alpha-2-macroglobulin (mg/dL)	0.002206422	0.003378556	0.004237288	0.002206422
Urinary Ca (mmol/L)	0.002203996	0.001244438	0.004237288	0.002203996
Interleukin-1B via ELISA (pg/mL)	0.002202815	0.001684749	0.004237288	0.002202815
Omega-6 fatty acids as % of total fatty acid mols	0.002083401	0.005710026	0.004237288	0.002083401
Beta globulins (%)	0.00198861	0.002508954	0.004237288	0.00198861
Fatty acid C20:5 n-3 as % of total fatty acid area	0.001750099	0.001527257	0.004237288	0.001750099
rs1539053_A	0.001651772	0.002032121	0.004237288	0.001651772
rs603050_T	0.001603176	0.000828142	0.004237288	0.001603176
Albumin (%)	0.001497865	0.00144821	0.004237288	0.001497865
Fatty acid C20:0 (arachidic) area	0.00142038	0.000965866	0.004237288	0.00142038
Lymphocytes (%)	0.001375674	0.000826441	0.008474576	0.001375674
Tumor necrosis factor-α via multiplex technology (pg/mL)	0.001189543	0.000626886	0.004237288	0.001189543
Mean corpuscular volume (MCV) (fL)	0.001108848	0.002522049	0.004237288	0.001108848
Fatty acid C16:0 (palmitic) area	0.001087023	0.000250191	0.004237288	0.001087023
rs948399_C	0.001081045	0.006526954	0.004237288	0.001081045
Fatty acid C16:0 as % of total fatty acid area	0.001062891	0.00384092	0.004237288	0.001062891
White blood cells (WBC) (n, K/μL)	0.001029476	0.000265338	0.004237288	0.001029476
Urinary creatinine (mg/dL)	0.001010878	0.001412657	0.004237288	0.001010878
Lipids: LDL cholesterol (mg/dL)	0.000972468	0.004258281	0.004237288	0.000972468
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	0.000969549	0.002657776	0.008474576	0.000969549
Omega-3 plasma fatty acid weight (mg/L)	0.000945649	0.000533488	0.004237288	0.000945649
Interleukin-12 via Bio-Plex (pg/mL)	0.000902174	0.000602753	0.004237288	0.000902174
Mean platelet volume (MPV) (fL)	0.00059614	0.000315899	0.004237288	0.00059614
rs4968782_G	0.00026173	0.000367413	0.004237288	0.00026173

Table VII: Cognitive Decline Features Model II

Cognitive Decline Features	Gain	Cover	Frequency	Importance
Age	0.337620007	0.169876149	0.04730832	0.337620007
Level of Education	0.101396229	0.107996945	0.042414356	0.101396229
Soluble IL-6 receptor via ELISA (ng/mL)	0.036437613	0.034064328	0.019575856	0.036437613
Retinol via high performance liquid chromatography (µmol/L)	0.02218011	0.02573877	0.020391517	0.02218011
Hemoglobin (g/dL)	0.014452739	0.007164163	0.005709625	0.014452739
Alpha-2 globulin (%)	0.012001659	0.007818561	0.008972268	0.012001659
Albumin (%)	0.0113596	0.015724044	0.01141925	0.0113596
Fatty acid C22:0 as % of total fatty acid area	0.011304629	0.009233553	0.008972268	0.011304629
Soluble CD14 via ELISA (ng/mL)	0.010711498	0.009622826	0.010603589	0.010711498
White blood cells (WBC) (n, K/µL)	0.010337767	0.006623209	0.009787928	0.010337767
Free thyroxine, FT4 (ng/dL)	0.010186247	0.012097542	0.018760196	0.010186247
Alpha-2-macroglobulin (mg/dL)	0.010031357	0.006651148	0.006525285	0.010031357
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	0.009701129	0.010461974	0.01223491	0.009701129
Soluble transferrin receptor (nmol/L)	0.008957121	0.005636462	0.004893964	0.008957121
Tumor necrosis factor-α via multiplex technology (pg/mL)	0.008815898	0.004477882	0.009787928	0.008815898
IGF binding protein-3, serum, immunoradiometric assay (ng/mL)	0.008096051	0.007201284	0.01141925	0.008096051
Neutrophils (n, K/µL)	0.007846614	0.013026383	0.003262643	0.007846614
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	0.007495273	0.010668779	0.013050571	0.007495273
Monocytes (n, K/µL)	0.007009487	0.008960452	0.008156607	0.007009487
Total testosterone (ng/mL)	0.006935307	0.004804623	0.005709625	0.006935307
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	0.006418083	0.010770744	0.015497553	0.006418083
Lycopene via high performance liquid chromatography (µmol/L)	0.006368592	0.013556497	0.010603589	0.006368592
Red cell distribution width (RDW) (%)	0.00632215	0.007376994	0.004078303	0.00632215
Urinary Na (mmol/L)	0.006308561	0.004860064	0.010603589	0.006308561
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.006234605	0.015158455	0.01223491	0.006234605
Estradiol via radioimmunoassay (pg/mL)	0.006087853	0.010731789	0.010603589	0.006087853
Interleukin-8 via Bio-Plex (pg/mL)	0.005942379	0.010463931	0.004078303	0.005942379
Urinary Ca (mmol/L)	0.005933618	0.013130706	0.013866232	0.005933618
Mean corpuscular volume (MCV) (fL)	0.005816721	0.005494248	0.007340946	0.005816721
MCH concentration (MCHC) (g/dL)	0.005676189	0.016226196	0.009787928	0.005676189
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	0.005299693	0.002967166	0.004893964	0.005299693
Neutrophils (%)	0.005259012	0.011038932	0.007340946	0.005259012
Omega-6 fatty acids as % of total fatty acid area	0.005235156	0.008265979	0.009787928	0.005235156
Fatty acid C22:0 (behenic) area	0.005221169	0.002244553	0.005709625	0.005221169
Fibrinogen (mg/dL)	0.005135355	0.013681843	0.015497553	0.005135355
Resistin via EIA (ng/mL)	0.00507799	0.004782631	0.01141925	0.00507799
Endogenous secretory receptor for AGEs (ng/mL)	0.005063804	0.005090446	0.01223491	0.005063804
Na+ (mEq/L)	0.005060928	0.003545462	0.003262643	0.005060928
Lipids: total cholesterol (mg/dL)	0.00494213	0.004189768	0.003262643	0.00494213
C-reactive protein - high sensitivity (µg/mL)	0.004882981	0.007480991	0.003262643	0.004882981
Vitamin E alpha tocopherol	0.004726757	0.003329939	0.006525285	0.004726757
Cystatin C (mg/L)	0.004380661	0.009384469	0.008972268	0.004380661
Parathyroid hormone	0.004375232	0.011901958	0.015497553	0.004375232
Adiponectin via RIA (µg/mL)	0.004335486	0.008986282	0.016313214	0.004335486
Urinary cortisol (µg/mL)	0.004259415	0.004101516	0.004893964	0.004259415
Plasma insulin via RIA (mIU/L)	0.004230209	0.004566037	0.008972268	0.004230209
Blood glucose (mg/dL)	0.004019193	0.00527928	0.007340946	0.004019193
Fatty acid C24:0 (lignoceric) area	0.003943775	0.003301283	0.003262643	0.003943775
C-terminal telopeptide of type-1 collagen (ng/mL)	0.003927989	0.006010915	0.013050571	0.003927989
24-hour urinary cortisol (µg/24 hours)	0.003839175	0.00320971	0.004078303	0.003839175
Lymphocytes (%)	0.003810109	0.005980394	0.008156607	0.003810109
Homocysteine via FPIA analysis (µmol/L)	0.003723506	0.008958335	0.005709625	0.003723506
Folate via RIA (ng/mL)	0.003721675	0.007480933	0.01223491	0.003721675
Ratio of Omega-6:Omega-3 as % of total fatty acid weight	0.003719853	0.00421514	0.004893964	0.003719853
Ca++ (mg/dL)	0.003651681	0.000474762	0.001631321	0.003651681
GPT (also known as ALT) (U/L)	0.003632801	0.004977336	0.007340946	0.003632801
24-hour urinary creatinine (mg/24 hours)	0.003451352	0.00780884	0.007340946	0.003451352
Anticholinergic Burden Scale Sum Score	0.00332339	0.002536376	0.002446982	0.00332339
Interleukin-10 via ELISA (pg/mL)	0.003322959	0.001514995	0.003262643	0.003322959
Beta globulins (%)	0.003321289	0.001896441	0.002446982	0.003321289
Creatinine clearance, 24-hr urine (mL/minute)	0.003264735	0.006994998	0.004893964	0.003264735
Interleukin-12 via Bio-Plex (pg/mL)	0.003192173	0.002789774	0.005709625	0.003192173
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	0.003185556	0.002444962	0.004893964	0.003185556
Vitamin B12 via RIA (pg/mL)	0.003166109	0.006439686	0.008156607	0.003166109
Urinary creatinine (mg/dL)	0.003122104	0.004013699	0.004893964	0.003122104
Transforming growth factor-B1 (pg/mL)	0.003012441	0.005744855	0.006525285	0.003012441
Fatty acid C24:0 as % of total fatty acid weight	0.002794238	0.005796356	0.003262643	0.002794238
Thyroid stimulating hormone, TSH (mIU/L)	0.002728823	0.008640367	0.01223491	0.002728823
Dehydroepiandrosterone sulfate (µg/dL)	0.002672337	0.003760759	0.004078303	0.002672337
Blood urea nitrogen (mg/dL)	0.002618936	0.003559652	0.01223491	0.002618936

Lipoprotein(a) (mg/dL)	0.002613825	0.004184359	0.006525285	0.002613825
Gamma glutamyl transferase (U/L)	0.002507317	0.001113761	0.001631321	0.002507317
Omega-3 fatty acids as % of total fatty acid weight	0.002446855	0.002262325	0.004078303	0.002446855
Serum creatinine (mg/dL)	0.002434934	0.003678538	0.004893964	0.002434934
Monocytes (%)	0.002412472	0.001491166	0.002446982	0.002412472
Serum cortisol (µg/dL)	0.002348164	0.00222644	0.004893964	0.002348164
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	0.002246116	0.00345916	0.008156607	0.002246116
Fatty acid C16:0 weight (mg/L)	0.002240052	0.002564482	0.004078303	0.002240052
Fatty acid C16:0 as % of total fatty acid weight	0.00223029	0.003813884	0.008156607	0.00223029
Fatty acid C20:5 n-3 as % of total fatty acid area	0.002230142	0.001064622	0.001631321	0.002230142
Methylmalonic acid, MMA (µmol/L)	0.002226268	0.002950204	0.002446982	0.002226268
Omega-3 plasma fatty acid weight (mg/L)	0.002058344	0.001317004	0.002446982	0.002058344
Ferritin (ng/mL)	0.00196076	0.005336786	0.008972268	0.00196076
Fatty acid C22:0 weight (mg/L)	0.001816603	0.00405515	0.005709625	0.001816603
Vitamin E gamma tocopherol	0.001806414	0.002203634	0.004893964	0.001806414
Lymphocytes (n, K/µL)	0.001690487	0.001349849	0.002446982	0.001690487
Lipids: LDL cholesterol (mg/dL)	0.00168208	0.005742559	0.004078303	0.00168208
Omega-6 plasma fatty acid weight (mg/L)	0.001668354	0.001010453	0.003262643	0.001668354
Interleukin-6 via ELISA ultrasensitive (pg/mL)	0.001545979	0.003147402	0.007340946	0.001545979
Creatine phosphokinase (U/L)	0.00148699	0.003371184	0.006525285	0.00148699
AST (U/L)	0.00148228	0.001280547	0.003262643	0.00148228
Red blood cells (RBC) (n, millions/µL)	0.001465528	0.003944097	0.002446982	0.001465528
Interleukin-1B via ELISA (pg/mL)	0.001462355	0.004043732	0.003262643	0.001462355
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	0.001341605	0.000218363	0.000815661	0.001341605
Fatty acid C20:5 n-3 as % of total fatty acid weight	0.001325055	0.003336324	0.006525285	0.001325055
Fatty acid C22:0 as % of total fatty acid weight	0.001245406	0.004586598	0.004893964	0.001245406
Vitamin B6 via high performance liquid chromatography (ng/mL)	0.001214394	0.002876213	0.003262643	0.001214394
Lipids: HDL cholesterol (mg/dL)	0.001198349	0.002421287	0.001631321	0.001198349
Urine hemoglobin (mg/dL)	0.001125041	0.002413289	0.000815661	0.001125041
Urine proteins (mg/dL)	0.001117313	0.001020134	0.000815661	0.001117313
Vitamin E gamma tocopherol	0.001085563	0.004068181	0.004893964	0.001085563
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	0.001080915	0.002484724	0.002446982	0.001080915
Total proteins (g/dL)	0.001012553	0.001464147	0.001631321	0.001012553
Free testosterone (ng/dL), Vermeulen	0.000911295	0.003313265	0.004893964	0.000911295
Hematocrit (%)	0.0008604	0.001047362	0.002446982	0.0008604
Uric acid (mg/dL)	0.000829065	0.000289542	0.000815661	0.000829065
Fatty acid C20:0 as % of total fatty acid area	0.000757176	0.000224922	0.000815661	0.000757176
Fatty acid C20:5 n-3 cis (eicosapentaenoic, EPA) area	0.000723203	0.00149168	0.003262643	0.000723203
C-reactive protein - low sensitivity (µg/mL)	0.000710255	0.001478066	0.002446982	0.000710255
TNF-related apoptosis-inducing ligand (pg/mL)	0.000646151	0.00201871	0.006525285	0.000646151
Omega-3 fatty acids as % of total fatty acid area	0.000637288	0.001485081	0.002446982	0.000637288
Fatty acid C20:0 weight (mg/L)	0.000636213	0.000729121	0.002446982	0.000636213
Fatty acid C20:0 (arachidic) area	0.00062639	0.002906608	0.000815661	0.00062639
Erythrocyte sedimentation rate (ESR) (mm/hour)	0.000622845	0.001423501	0.003262643	0.000622845
Omega-6 fatty acids as % of total fatty acid weight	0.000620856	0.00168309	0.005709625	0.000620856
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	0.000596805	0.001070486	0.003262643	0.000596805
Depression	0.000535185	0.001586769	0.001631321	0.000535185
Cortisol:DHEAS ratio (based on nmols)	0.000500719	0.001411308	0.004078303	0.000500719
Fatty acid C16:0 as % of total fatty acid weight	0.000453894	0.000722058	0.001631321	0.000453894
Beta-carotene via high performance liquid chromatography (µmol/L)	0.000433366	0.001824619	0.004078303	0.000433366
Mean corpuscular hemoglobin (MCH) (pg)	0.000407813	0.001502518	0.002446982	0.000407813
Vitamin E alpha tocopherol	0.000407016	0.00132765	0.004078303	0.000407016
Alpha-1 globulin (%)	0.00028578	0.001049279	0.002446982	0.00028578
Ratio of Omega-6:Omega-3 as % of total fatty acid mols	0.000218333	0.001446424	0.004893964	0.000218333
Fatty acid C20:0 as % of total fatty acid mols	0.000178967	0.001169142	0.003262643	0.000178967
Fatty acid C16:0 as % of total fatty acid area	0.000166057	0.000634616	0.002446982	0.000166057
Fatty acid C20:0 as % of total fatty acid weight	6.18E-05	0.000575459	0.002446982	6.18E-05
Omega-6 fatty acids as % of total fatty acid mols	5.70E-05	0.001159185	0.002446982	5.70E-05
Ratio of Omega-6:Omega-3 as % of total fatty acid area	1.80E-05	0.000249955	0.000815661	1.80E-05

Table VIII: Frailty Features Model II

Frailty Features	Gain	Cover	Frequency	Importance
Depression	0.087477094	0.037432424	0.023728814	0.087477094
Age	0.050692047	0.012070795	0.010169492	0.050692047
Creatinine clearance, 24-hr urine (mL/minute)	0.037570335	0.008241562	0.006779661	0.037570335
Anticholinergic Burden Scale Sum Score	0.031644308	0.019787206	0.013559322	0.031644308
Homocysteine via FPIA analysis (μmol/L)	0.030538058	0.023689405	0.016949153	0.030538058
Vitamin B6 via high performance liquid chromatography (ng/mL)	0.024418035	0.013747038	0.016949153	0.024418035
Interleukin-6 via ELISA ultrasensitive (pg/mL)	0.023153468	0.019395502	0.016949153	0.023153468
Erythrocyte sedimentation rate (ESR) (mm/hour)	0.021774327	0.012878682	0.010169492	0.021774327
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	0.020875442	0.028967385	0.027118644	0.020875442
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	0.020207291	0.029445221	0.020338983	0.020207291
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.019045838	0.026328877	0.023728814	0.019045838
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	0.017850277	0.010389129	0.013559322	0.017850277
Interleukin-8 via Bio-Plex (pg/mL)	0.016988136	0.00439415	0.006779661	0.016988136
24-hour urinary cortisol (μg/24 hours)	0.016449453	0.033101985	0.023728814	0.016449453
24-hour urinary creatinine (mg/24 hours)	0.014868273	0.007683043	0.010169492	0.014868273
Cortisol:DHEAS ratio (based on nmols)	0.014535409	0.009961332	0.013559322	0.014535409
Interleukin-1B via ELISA (pg/mL)	0.014350009	0.023361053	0.016949153	0.014350009
Creatine phosphokinase (U/L)	0.013873065	0.004025311	0.006779661	0.013873065
Plasma insulin via RIA (mIU/L)	0.013636196	0.00598363	0.013559322	0.013636196
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (μmol/L)	0.013177049	0.013905765	0.013559322	0.013177049
Vitamin E gamma tocopherol, high performance liquid chromatography, assay #2 (μmol/L)	0.012842565	0.022567257	0.016949153	0.012842565
Methylmalonic acid, MMA (μmol/L)	0.012496487	0.017167754	0.013559322	0.012496487
C-terminal telopeptide of type-1 collagen (ng/mL)	0.012409377	0.007676331	0.010169492	0.012409377
Fatty acid C20:5 n-3 weight (mg/L)	0.011973673	0.009092363	0.013559322	0.011973673
Blood urea nitrogen (mg/dL)	0.0116946	0.010837487	0.016949153	0.0116946
Dehydroepiandrosterone sulfate (μg/dL)	0.011520092	0.002702369	0.010169492	0.011520092
Serum cortisol (μg/dL)	0.011018876	0.022298275	0.020338983	0.011018876
Ratio of Omega-6:Omega-3 as % of total fatty acid weight	0.010754702	0.001495996	0.010169492	0.010754702
Endogenous secretory receptor for AGEs (ng/mL)	0.010218501	0.01710008	0.016949153	0.010218501
Vitamin E gamma tocopherol, high performance liquid chromatography (μmol/L)	0.010084578	0.010687575	0.010169492	0.010084578
Free testosterone (ng/dL), Vermeulen	0.009711035	0.018664758	0.016949153	0.009711035
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	0.009156656	0.00691564	0.006779661	0.009156656
Monocytes (%)	0.008981439	0.021490956	0.016949153	0.008981439
C-reactive protein - high sensitivity (μg/mL)	0.008807614	0.009176731	0.010169492	0.008807614
Urine proteins (mg/dL)	0.008690624	0.009293653	0.006779661	0.008690624
Lipoprotein(a) (mg/dL)	0.008365481	0.014865471	0.010169492	0.008365481
Vitamin E alpha tocopherol, high performance liquid chromatography (μmol/L)	0.0083467	0.001745162	0.003389831	0.0083467
Lipids: LDL cholesterol (mg/dL)	0.008345016	0.014511828	0.013559322	0.008345016
Tumor necrosis factor-α via multiplex technology (pg/mL)	0.008263742	0.006171901	0.013559322	0.008263742
rs429358_C	0.008207264	0.009874568	0.006779661	0.008207264
Mean corpuscular volume (MCV) (fL)	0.007813068	0.002890419	0.003389831	0.007813068
White blood cells (WBC) (n, K/μL)	0.007684036	0.010274828	0.010169492	0.007684036
Lymphocytes (%)	0.007590389	0.000963582	0.006779661	0.007590389
rs10501927_G	0.007400781	0.00947889	0.010169492	0.007400781
Fatty acid C22:0 as % of total fatty acid area	0.007284487	0.008147218	0.006779661	0.007284487
Ferritin (ng/mL)	0.007123548	0.00835951	0.013559322	0.007123548
Urinary Na (mmol/L)	0.007073396	0.019248543	0.013559322	0.007073396
Folate via RIA (ng/mL)	0.007011844	0.015035268	0.010169492	0.007011844
Red blood cells (RBC) (n, millions/μL)	0.006799328	0.006376907	0.003389831	0.006799328
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	0.006789192	0.001887906	0.006779661	0.006789192
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	0.006415482	0.012690498	0.010169492	0.006415482
Fatty acid C20:0 as % of total fatty acid area	0.006179054	0.001341804	0.003389831	0.006179054
Transforming growth factor-B1 (pg/mL)	0.00593631	0.010002974	0.010169492	0.00593631
Beta-carotene via high performance liquid chromatography (μmol/L)	0.005883288	0.004352394	0.010169492	0.005883288
Free thyroxine, fT4 (ng/dL)	0.005714929	0.010081288	0.010169492	0.005714929

Monocytes (n, K/ μ L)	0.005607768	0.017694614	0.013559322	0.005607768
rs12752888_C	0.005524297	0.001255391	0.003389831	0.005524297
Fatty acid C16:0 (palmitic) area	0.005491614	0.009704169	0.010169492	0.005491614
IL-6 high-sensitivity ELISA calculated from ELISA ultrasensitive (pg/mL)	0.005425693	0.006151266	0.003389831	0.005425693
TNF-related apoptosis-inducing ligand (pg/mL)	0.005359504	0.00793391	0.006779661	0.005359504
Fatty acid C22:0 (behenic) area	0.004905328	0.003961571	0.006779661	0.004905328
Serum creatinine (mg/dL)	0.004786974	0.000934218	0.006779661	0.004786974
Macrophage inflammatory protein-1b via Bio-Plex (pg/mL)	0.004784017	0.012336797	0.010169492	0.004784017
Fatty acid C16:0 as % of total fatty acid area	0.004516237	0.004702992	0.003389831	0.004516237
Vitamin B12 via RIA (pg/mL)	0.004424143	0.000391277	0.003389831	0.004424143
Fatty acid C16:0 as % of total fatty acid weight	0.004402626	0.00403443	0.006779661	0.004402626
Estradiol via radioimmunoassay (pg/mL)	0.004344044	0.000553	0.003389831	0.004344044
rs1799990_G	0.004294903	0.000615221	0.003389831	0.004294903
Interleukin-10 via ELISA (pg/mL)	0.004255737	0.000180399	0.003389831	0.004255737
Urinary Ca (mmol/L)	0.004248745	0.010166493	0.006779661	0.004248745
Omega-3 plasma fatty acid weight (mg/L)	0.004240931	0.010351219	0.006779661	0.004240931
Alpha-2-macroglobulin (mg/dL)	0.004052975	0.005200807	0.006779661	0.004052975
Alpha-1 globulin (%)	0.003887647	0.010139885	0.006779661	0.003887647
Soluble CD14 via ELISA (ng/mL)	0.003845678	0.005222931	0.003389831	0.003845678
Fatty acid C20:5 n-3 as % of total fatty acid area	0.003796235	0.008663351	0.010169492	0.003796235
MCH concentration (MCHC) (g/dL)	0.003719857	0.005707609	0.006779661	0.003719857
rs11894266_C	0.003698922	0.00444944	0.006779661	0.003698922
rs8106922_G	0.003649847	0.000344203	0.003389831	0.003649847
Total testosterone (ng/mL)	0.003388364	0.006891703	0.006779661	0.003388364
rs7840202_C	0.003264864	0.003877332	0.003389831	0.003264864
Ratio of Omega-6:Omega-3 as % of total fatty acid area	0.003164728	0.009068617	0.006779661	0.003164728
Gamma glutamyl transferase (U/L)	0.003062083	0.012510653	0.010169492	0.003062083
Lipids: HDL cholesterol (mg/dL)	0.003057754	0.008728878	0.006779661	0.003057754
Resistin via EIA (ng/mL)	0.003006318	0.004709943	0.006779661	0.003006318
Urine hemoglobin (mg/dL)	0.002983008	0.004156007	0.003389831	0.002983008
Lipids: total cholesterol (mg/dL)	0.002916458	0.009319371	0.006779661	0.002916458
Adiponectin via RIA (μ g/mL)	0.00276724	0.008715665	0.006779661	0.00276724
Uric acid (mg/dL)	0.002622806	0.001158878	0.003389831	0.002622806
Fatty acid C22:0 as % of total fatty acid weight	0.002467219	0.000332762	0.003389831	0.002467219
rs3785880_G	0.002427357	0.000734247	0.003389831	0.002427357
rs10883631_G	0.002364875	0.000173967	0.006779661	0.002364875
Omega-3 fatty acids as % of total fatty acid area	0.002326994	0.0085397	0.006779661	0.002326994
Omega-6 plasma fatty acid weight (mg/L)	0.00219113	0.00264264	0.003389831	0.00219113
Fatty acid C24:0 weight (mg/L)	0.002189504	0.001294336	0.003389831	0.002189504
Soluble IL-6 receptor via ELISA (ng/mL)	0.002001984	0.003757795	0.003389831	0.002001984
Omega-6 fatty acids as % of total fatty acid area	0.001960985	0.00316459	0.003389831	0.001960985
rs4363657_C	0.001848912	0.005068245	0.003389831	0.001848912
ALT (U/L)	0.001744514	0.003480211	0.003389831	0.001744514
rs6859_A	0.001731615	0.000162633	0.003389831	0.001731615
Omega-3 fatty acids as % of total fatty acid weight	0.001710078	0.004212545	0.003389831	0.001710078
Urinary cortisol (μ g/mL)	0.001597598	0.000305797	0.003389831	0.001597598
Fatty acid C20:0 weight (mg/L)	0.001571151	0.000274045	0.003389831	0.001571151
rs948399_C	0.001532092	0.004343178	0.003389831	0.001532092
Blood glucose (mg/dL)	0.001363833	0.004740806	0.003389831	0.001363833
Fatty acid C24:0 as % of total fatty acid weight	0.001273097	0.003990736	0.003389831	0.001273097
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	0.000967469	0.003538632	0.003389831	0.000967469
rs4147929_A	0.000892763	0.004034426	0.003389831	0.000892763
Fatty acid C24:0 as % of total fatty acid area	0.000868838	0.004156007	0.003389831	0.000868838
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	0.000847489	0.004086658	0.003389831	0.000847489
Fatty acid C20:5 n-3 cis (eicosapentaenoic, EPA) area	0.000807547	0.003996284	0.003389831	0.000807547
Neutrophils (n, K/ μ L)	0.000572539	0.000112781	0.003389831	0.000572539

Table IX: Cognitive Frailty Features Model II

Cognitive Frailty Features	Gain	Cover	Frequency	Importance
Age	0.243224429	0.110510017	0.056140351	0.243224429
Depression	0.058485822	0.040044719	0.028070175	0.058485822
Interleukin-6 via ELISA ultrasensitive (pg/mL)	0.041790268	0.019616729	0.014035088	0.041790268
Creatinine clearance, 24-hr urine (mL/minute)	0.030831483	0.018261243	0.014035088	0.030831483
Erythrocyte sedimentation rate (ESR) (mm/hour)	0.02106198	0.017790494	0.014035088	0.02106198
Creatine phosphokinase (U/L)	0.019570487	0.025241618	0.021052632	0.019570487
C-terminal telopeptide of type-1 collagen (ng/mL)	0.019104806	0.01358605	0.014035088	0.019104806
Level of Education	0.018254792	0.013650884	0.010526316	0.018254792
Cystatin C (mg/L)	0.016738529	0.018435368	0.028070175	0.016738529
MCH concentration (MCHC) (g/dL)	0.016397158	0.008585323	0.010526316	0.016397158
Vitamin B6 via high performance liquid chromatography (ng/mL)	0.016303861	0.020917254	0.01754386	0.016303861
Plasma insulin via RIA (mIU/L)	0.01479562	0.015709937	0.01754386	0.01479562
Tumor necrosis factor- α via multiplex technology (pg/mL)	0.014217085	0.018035692	0.01754386	0.014217085
C-reactive protein - high sensitivity (μ g/mL)	0.014035509	0.015325407	0.01754386	0.014035509
Total testosterone (ng/mL)	0.013534893	0.018254719	0.010526316	0.013534893
Vitamin E alpha tocopherol, high performance liquid chromatography (μ mol/L)	0.013064167	0.012399389	0.010526316	0.013064167
Vitamin E gamma tocopherol, high performance liquid chromatography (μ mol/L)	0.013060202	0.01874998	0.021052632	0.013060202
Albumin (%)	0.012957738	0.006378732	0.010526316	0.012957738
rs4343_A	0.012850391	0.014645507	0.010526316	0.012850391
Dehydroepiandrosterone sulfate (μ g/dL)	0.012658594	0.011407339	0.007017544	0.012658594
Blood glucose (mg/dL)	0.012532317	0.010288275	0.014035088	0.012532317
Vitamin E gamma tocopherol, high performance liquid chromatography, assay #2 (μ mol/L)	0.011033869	0.022001512	0.021052632	0.011033869
Serum creatinine (mg/dL)	0.010983672	0.023256757	0.01754386	0.010983672
Monocytes (n, K/ μ L)	0.010578729	0.02257873	0.01754386	0.010578729
Vitamin B12 via RIA (pg/mL)	0.010033777	0.01388142	0.01754386	0.010033777
Alpha-2-macroglobulin (mg/dL)	0.009632353	0.015012622	0.010526316	0.009632353
Monocytes (%)	0.009258662	0.020772466	0.014035088	0.009258662
Soluble TNF- α receptor II via quantitative sandwich EIA (pg/mL)	0.009249448	0.006819346	0.010526316	0.009249448
Soluble TNF- α receptor I via quantitative sandwich EIA (pg/mL)	0.008910896	0.011338502	0.007017544	0.008910896
Fatty acid C16:0 (palmitic) area	0.008774872	0.012353205	0.01754386	0.008774872
Folate via RIA (ng/mL)	0.008433119	0.009100673	0.01754386	0.008433119
Fatty acid C20:5 n-3 weight (mg/L)	0.008258345	0.012186001	0.010526316	0.008258345
24-hour urinary creatinine (mg/24 hours)	0.008193473	0.007541586	0.014035088	0.008193473
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	0.007746372	0.010792335	0.014035088	0.007746372
Anticholinergic Burden Scale Sum Score	0.007620497	0.004940168	0.003508772	0.007620497
Interleukin-1B via ELISA (pg/mL)	0.007547503	0.007646187	0.014035088	0.007547503
Omega-6 fatty acids as % of total fatty acid weight	0.007517237	0.005039685	0.007017544	0.007517237
Ratio of Omega-6:Omega-3 as % of total fatty acid area	0.007500432	0.00741643	0.010526316	0.007500432
TNF-related apoptosis-inducing ligand (pg/mL)	0.00719579	0.004106597	0.014035088	0.00719579
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	0.007054061	0.014543542	0.010526316	0.007054061
White blood cells (WBC) (n, K/ μ L)	0.006973812	0.008888854	0.01754386	0.006973812
24-hour urinary cortisol (μ g/24 hours)	0.00650171	0.007939441	0.010526316	0.00650171
Serum cortisol (μ g/dL)	0.00583663	0.007164907	0.014035088	0.00583663
Soluble CD14 via ELISA (ng/mL)	0.005597229	0.008704836	0.010526316	0.005597229
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (μ mol/L)	0.005576786	0.006297576	0.007017544	0.005576786
Urinary creatinine (mg/dL)	0.005467574	0.004446002	0.003508772	0.005467574
Mean corpuscular hemoglobin (MCH) (pg)	0.005181663	0.013591552	0.010526316	0.005181663
Homocysteine via FPIA analysis (μ mol/L)	0.005058175	0.01368253	0.014035088	0.005058175
Urinary Na (mmol/L)	0.005040676	0.00568806	0.003508772	0.005040676
Alpha-1 globulin (%)	0.004905499	0.00071443	0.003508772	0.004905499
Fatty acid C20:5 n-3 as % of total fatty acid area	0.004842838	0.004365281	0.007017544	0.004842838
rs11894266_C	0.004815596	0.001626576	0.003508772	0.004815596
rs429358_C	0.004217372	0.009503585	0.007017544	0.004217372
Lymphocytes (%)	0.00400406	0.005602415	0.003508772	0.00400406

Neutrophils (%)	0.00388351	0.007716045	0.010526316	0.00388351
Lymphocytes (n, K/ μ L)	0.003737902	0.004870851	0.007017544	0.003737902
Fibrinogen (mg/dL)	0.003655515	0.008378834	0.010526316	0.003655515
Omega-6 plasma fatty acid weight (mg/L)	0.00365299	0.002054096	0.003508772	0.00365299
Hematocrit (%)	0.003643905	0.003572639	0.003508772	0.003643905
rs129968_A	0.003602199	0.006056578	0.003508772	0.003602199
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	0.003578744	0.00854427	0.010526316	0.003578744
Lipids: LDL cholesterol (mg/dL)	0.003552118	0.005060933	0.003508772	0.003552118
rs129968_A	0.003366369	0.001166338	0.003508772	0.003366369
rs129968_A	0.003136504	0.008586573	0.007017544	0.003136504
Interleukin-8 via Bio-Plex (pg/mL)	0.003044247	0.001130557	0.007017544	0.003044247
Adiponectin via RIA (μ g/mL)	0.003007961	0.003092498	0.003508772	0.003007961
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	0.002996903	0.008428031	0.007017544	0.002996903
Omega-3 fatty acids as % of total fatty acid area	0.002933676	0.009319042	0.007017544	0.002933676
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	0.002864988	0.008290595	0.007017544	0.002864988
rs3785880_G	0.002812	0.002084713	0.003508772	0.002812
Fatty acid C16:0 weight (mg/L)	0.002756752	0.008702471	0.007017544	0.002756752
rs3785880_G	0.002755476	0.00205785	0.003508772	0.002755476
Urine proteins (mg/dL)	0.002722263	0.005708104	0.003508772	0.002722263
Uric acid (mg/dL)	0.002571323	0.005440035	0.003508772	0.002571323
Monocyte chemoattractant protein-1 via Bio-Plex (pg/mL)	0.002567023	0.00851028	0.007017544	0.002567023
rs1800629_A	0.002566224	0.000827461	0.003508772	0.002566224
rs7840202_C	0.002544047	0.000872785	0.003508772	0.002544047
Beta-carotene via high performance liquid chromatography (μ mol/L)	0.002486527	0.002014956	0.003508772	0.002486527
rs360722_A	0.002412795	0.002982585	0.003508772	0.002412795
rs12752888_C	0.002354584	0.003431046	0.003508772	0.002354584
Resistin via EIA (ng/mL)	0.00233496	0.005168944	0.007017544	0.00233496
Lipoprotein(a) (mg/dL)	0.002292647	0.003466989	0.003508772	0.002292647
Fatty acid C16:0 as % of total fatty acid area	0.002120433	0.000759578	0.003508772	0.002120433
Urine glucose (mg/dL)	0.002063151	0.00532065	0.003508772	0.002063151
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	0.00204864	0.002395909	0.003508772	0.00204864
Fatty acid C22:0 weight (mg/L)	0.001854603	0.003613198	0.003508772	0.001854603
Blood urea nitrogen (mg/dL)	0.001848406	0.00356917	0.007017544	0.001848406
Lycopene via high performance liquid chromatography (μ mol/L)	0.001844727	0.004465825	0.003508772	0.001844727
Interleukin-12 via Bio-Plex (pg/mL)	0.001789151	0.000398635	0.003508772	0.001789151
Endogenous secretory receptor for AGEs (ng/mL)	0.001712395	0.000343816	0.003508772	0.001712395
Gamma glutamyl transferase (U/L)	0.001699071	0.000303829	0.003508772	0.001699071
Lipids: HDL cholesterol (mg/dL)	0.001657368	0.003377608	0.007017544	0.001657368
Estradiol via radioimmunoassay (pg/mL)	0.001620834	0.002680514	0.003508772	0.001620834
Lipids: total cholesterol (mg/dL)	0.001591306	0.003105553	0.003508772	0.001591306
Thyroid stimulating hormone, TSH (mIU/L)	0.001578132	0.00108992	0.003508772	0.001578132
Urinary Ca (mmol/L)	0.001387468	0.004556416	0.003508772	0.001387468
Methylmalonic acid, MMA (μ mol/L)	0.00137796	0.001595891	0.003508772	0.00137796
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	0.001353188	0.004284569	0.007017544	0.001353188
rs16944_A	0.001347871	0.006524396	0.007017544	0.001347871
rs1614735_G	0.001347489	0.004511853	0.003508772	0.001347489
Fatty acid C22:0 (behenic) area	0.0011975	0.000379567	0.003508772	0.0011975
rs6131_T	0.001162675	0.004222359	0.003508772	0.001162675
rs8106922_G	0.001103625	0.003949645	0.003508772	0.001103625
Urinary cortisol (μ g/mL)	0.001072581	0.004113434	0.003508772	0.001072581
rs10501927_G	0.001051906	0.004078632	0.003508772	0.001051906
rs4363657_C	0.000844935	0.003497748	0.003508772	0.000844935
Free testosterone (ng/dL), Vermeulen	0.00078235	0.003440932	0.003508772	0.00078235
rs10883631_G	0.000627947	0.003206086	0.003508772	0.000627947
rs11771145_A	0.000341898	0.000144683	0.003508772	0.000341898
rs948399_C	0.00025933	0.000123066	0.003508772	0.00025933

Table X. Clinical features by healthy control and phenotype

Model 1	Cognitive Decline		Frailty		Cognitive Frailty					
	mean(SD)	p-value	mean(SD)	p-value	mean(SD)	p-value				
Age										
Control	65(15.7)		72(6.2)		73(6.4)					
Phenotype	80(8.7)	<0.0001	78(7.9)	<0.0001	82(7.4)	<0.0001				
Anicholinergic Burden										
Control	2.18 (2.01)		1.75 (1.76)		2.15 (2.02)					
Phenotype	2.69 (2.19)	<0.0001	2.89 (2.21)	<0.0001	3.00 (2.16)	<0.0001				
Gender	(n)		(n)		(n)					
Healthy Control(M/F)	521/557		286/274		418/480					
Phenotype(M/F)	121/254	<0.0001	214/381	<0.0001	82/175	<0.0001				
Depression										
Control	272		91		250					
Phenotype	140	<0.0001	269	<0.0001	110	<0.0001				
Baseline Dementia										
Control			12		12					
Phenotype			70	<0.0001	70	<0.0001				
Model 2	Cognitive Decline				Frailty		Cognitive Frailty			
	TrailA		TrailB				TrailA		TrailB	
Age	mean(SD)	p-value	mean(SD)	p-value	mean(SD)	p-value	mean(SD)	p-value	mean(SD)	p-value
Control	61(16.4)		52(17.4)		72(6.2)		64(15.6)		61(16.2)	
Phenotype	76(7.7)	<0.0001	72(9.0)	<0.0001	78(7.9)	<0.0001	78(7.4)	<0.0001	76(6.9)	<0.0001
Anicholinergic Burden										
Control	1.95 (1.87)		1.77 (1.73)		1.75 (1.76)		1.85 (1.82)		1.68 (1.66)	
Phenotype	2.44 (2.12)	<0.0011	2.23 (2.02)	0.042	2.89 (2.21)	<0.0001	3.01 (2.20)	<0.0001	2.79 (2.19)	<0.0001
Depression	(n)		(n)		(n)		(n)		(n)	
Control	135		52		91		188		120	
Phenotype	339	<0.0001	220	<0.0001	269	<0.0001	151	<0.0001	152	<0.0001

Table XI. Genomic univariate results Model I

Gene/Phenotype	Chromosome	SNP-allele	Allele Count	Estimate	Std. Error	z value	Pr(> z)
Cognitive Decline							
ACOT11	1	rs12752888_C	1	-0.48	0.15	-3.30	0.001
DAB1	1	rs1539053_A	1	0.33	0.16	1.99	0.05
DAB1	1	rs1539053_A	2	0.45	0.19	2.29	0.02
COMT	22	rs4646316_T	2	-0.62	0.29	-2.11	0.04
IL6R	1	rs2228145_C	1	-0.31	0.15	-2.13	0.03
Frailty							
MMP3	11	rs948399_C	2	0.60	0.30	2.01	0.05
Cognitive Frailty							
ACOT11	1	rs12752888_C	1	-0.47	0.18	-2.67	0.01
DAB1	1	rs1539053_A	1	0.51	0.20	2.58	0.01
MMP3	11	rs948399_C	1	0.41	0.17	2.46	0.01
MTRR	5	rs1801394_G	2	0.80	0.23	3.48	0.001
CD33	19	rs3865444_A	2	0.62	0.28	2.24	0.03

Note: bold text indicates the closes gene

Table XII. Genomic univariate results Model II

Gene/Phenotype	Neurocognitive Test	Chromosome	SNP-allele	Allele Count	Estimate	Std. Error	z value	Pr(> z)
Cognitive Decline								
ACOT11	Trail B	1	rs12752888_C	2	-0.58	0.27	-2.12	0.03
ACOT11	Trail A	1	rs12752888_C	1	-0.25	0.13	-1.96	0.05
KCNU1	Trail B	8	rs1157242_T	1	0.47	0.16	2.90	0.004
PRNP	Trail B	20	rs1799990_G	1	0.30	0.15	2.10	0.04
PRNP	Trail A	20	rs1799990_G	2	0.45	0.22	2.06	0.04
BIN1	Trail B	2	rs744373_G	1	-0.31	0.14	-2.16	0.03
Frailty								
NECTIN2		19	rs6859_A	1	0.33	0.14	2.34	0.02
ABCA7		19	rs4147929_A	2	-0.27	0.14	-1.96	0.05
APOE		19	rs429358_C	1	-0.45	0.19	-2.27	0.02
SLCO1B1		12	rs4363657_C	1	0.38	0.14	2.57	0.01
MMP3		11	rs948399_C	2	0.60	0.29	2.01	0.04
Cognitive Frailty								
ACOT11	Trail B	1	rs12752888_C	1	-0.37	0.15	-2.46	0.01
ACOT11	Trail A	1	rs12752888_C	1	-0.34	0.15	-2.28	0.02
APOE	Trail B	19	rs429358_C	1	-0.59	0.23	-2.54	0.01
SLCO1B1	Trail B	12	rs4363657_C	1	0.38	0.16	2.39	0.02
MMP3	Trail A	11	rs948399_C	1	0.29	0.15	2.00	0.05
TOMM40	Trail A	19	rs8106922_G	1	-0.31	0.16	-1.92	0.05

Note: bold text indicates the closes gene

Table XIII. Difference between health control and cognitive decline results Model I

Cognitive Decline Model1	Control Mean	SD	Cognitive Mean	SD	t -test	Corrected p-value
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	56.26	36.69	39.66	29.96	<0.0001	<0.0001
Adiponectin via RIA (µg/mL)	12.50	8.79	17.15	12.21	<0.0001	<0.0001
Albumin (%)	59.58	3.40	57.43	3.86	<0.0001	<0.0001
Alpha-2-macroglobulin (mg/dL)	203.26	66.61	222.27	66.26	<0.0001	<0.0001
Alpha-1 globulin (%)	2.54	0.39	2.79	0.48	<0.0001	<0.0001
Alpha-2 globulin (%)	11.06	1.28	11.59	1.46	<0.0001	<0.0001
Blood urea nitrogen (mg/dL)	32.98	9.09	39.03	17.24	<0.0001	<0.0001
Fatty acid C20:5 n-3 weight (mg/L)	20.16	8.93	17.85	6.99	<0.0001	<0.0001
Fatty acid C20:5 n-3 as % of total fatty acid weight	0.63	0.22	0.56	0.18	<0.0001	<0.0001
Creatinine clearance, 24-hr urine (mL/minute)	86.84	30.09	66.91	25.91	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)	33.95	0.98	33.47	1.15	<0.0001	<0.0001
Creatine phosphokinase (U/L)	108.00	89.65	85.68	58.45	<0.0001	<0.0001
C-terminal telopeptide of type-1 collagen (ng/mL)	0.46	0.23	0.62	0.39	<0.0001	<0.0001
Cystatin C (mg/L)	0.93	0.26	1.16	0.46	<0.0001	<0.0001
Dehydroepiandrosterone sulfate (µg/dL)	115.68	96.75	72.89	64.01	<0.0001	<0.0001
Estradiol via radioimmunoassay (pg/mL)	13.46	17.95	8.90	6.13	<0.0001	<0.0001
Fibrinogen (mg/dL)	341.17	73.84	378.87	76.32	<0.0001	<0.0001
Free thyroxine, fT4 (ng/dL)	1.42	0.31	1.53	0.45	<0.0001	<0.0001
ALT (U/L)	21.19	14.29	17.22	9.37	<0.0001	<0.0001
Red blood cells (RBC) (n, millions/µL)	4.56	0.41	4.35	0.48	<0.0001	<0.0001
Homocysteine via FPIA analysis (µmol/L)	14.59	6.43	17.62	7.69	<0.0001	<0.0001
Red cell distribution width (RDW) (%)	13.54	0.95	14.01	1.23	<0.0001	<0.0001
Methylmalonic acid, MMA (µmol/L)	0.10	0.03	0.11	0.03	<0.0001	<0.0001
Omega-3 plasma fatty acid weight (mg/L)	110.63	41.96	98.98	37.76	<0.0001	<0.0001
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	23.69	17.54	31.58	24.54	<0.0001	<0.0001
Resistin via EIA (ng/mL)	3.78	1.84	4.62	2.57	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	1310.62	578.43	1842.17	1068.12	<0.0001	<0.0001
Urinary Ca (mmol/L)	2.43	1.65	1.97	1.55	<0.0001	<0.0001
24-hour urinary creatinine (mg/24 hours)	1058.67	372.66	825.55	326.16	<0.0001	<0.0001
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)	388.48	148.94	429.37	175.28	0.0002	0.0003
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	4397.72	1104.44	4122.44	1097.82	0.0002	0.0003
Omega-6 plasma fatty acid weight (mg/L)	1060.54	234.87	998.91	256.10	0.0003	0.0004
Lymphocytes (n, K/µL)	1.94	0.65	1.79	0.63	0.0005	0.0007
TNF-related apoptosis-inducing ligand (pg/mL)	75.80	40.87	69.69	23.55	0.001	0.002
Interleukin-6 via ELISA ultrasensitive (pg/mL)	1.76	2.07	3.04	7.1	0.002	0.002
Cortisol:DHEAS ratio (nmols)	0.28	0.71	0.53	1.81	0.002	0.002
Ratio of Omega-6:Omega-3 as % of total fatty acid area	16.38	5.05	17.51	6.06	0.005	0.005
Beta-carotene via high performance liquid chromatography (µmol/L)	0.43	0.28	0.38	0.23	0.009	0.011
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	33.68	7.32	32.33	8.31	0.011	0.012
Ratio of Omega-6:Omega-3 as % of total fatty acid mols	11.52	3.34	12.17	4.11	0.016	0.018
Uric acid (mg/dL)	5.03	1.35	5.27	1.65	0.019	0.021
Mean corpuscular volume (MCV) (fL)	90.04	4.65	90.76	5.22	0.03	0.031
Serum cortisol (µg/dL)	13.62	5.00	13.02	4.32	0.039	0.039

Table XIV. Difference between healthy control and frailty results Model I

Frailty Model 1	Control Mean	SD	Frailty Mean	SD	t -test	Corrected p-value
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	54.93	34.51	43.53	35.76	<0.0001	<0.0001
Albumin (%)	59.18	3.38	57.96	3.73	<0.0001	<0.0001
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	34.44	7.65	32.65	7.39	<0.0001	<0.0001
Blood urea nitrogen (mg/dL)	33.79	7.44	37.5	15.92	<0.0001	<0.0001
Creatinine clearance, 24-hr urine (mL/minute)	81.09	24.06	70.00	26.43	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)	33.90	1.02	33.56	1.05	<0.0001	<0.0001
Creatine phosphokinase (U/L)	104.22	61.69	86.84	55.12	<0.0001	<0.0001
C-terminal telopeptide of type-1 collagen (ng/mL)	0.47	0.23	0.58	0.35	<0.0001	<0.0001
Cystatin C (mg/L)	0.97	0.19	1.13	0.42	<0.0001	<0.0001
Homocysteine via FPIA analysis (µmol/L)	14.97	5.70	17.31	8.12	<0.0001	<0.0001
Red cell distribution width (RDW) (%)	13.62	0.93	13.89	1.16	<0.0001	<0.0001
Interleukin-6 via ELISA ultrasensitive (pg/mL)	1.66	1.75	2.92	5.74	<0.0001	<0.0001
Omega-6 plasma fatty acid weight (mg/L)	1069.54	249.81	1005.32	234.97	<0.0001	<0.0001
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	24.06	19.79	30.54	22.59	<0.0001	<0.0001
Resistin via EIA (ng/mL)	3.72	1.67	4.36	2.48	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	1343.02	429.61	1780.92	979.8	<0.0001	<0.0001
24-hour urinary creatinine (mg/24 hours)	1020.45	334.7	860.38	323.47	<0.0001	<0.0001
C-reactive protein - high sensitivity (µg/mL)	4.06	5.99	6.79	11.93	<0.0001	<0.0001
Free testosterone (ng/dL), Vermeulen	2.41	2.22	1.72	1.9	<0.0001	<0.0001
Hemoglobin (g/dL)	13.99	1.25	13.43	1.51	<0.0001	<0.0001
Hematocrit (%)	41.25	3.23	39.96	3.95	<0.0001	<0.0001
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	142.73	85.5	177.97	159.09	<0.0001	<0.0001
Neutrophils (n, K/µL)	3.59	1.18	3.90	1.31	<0.0001	<0.0001
Lymphocytes (%)	31.42	7.87	29.5	8.23	<0.0001	<0.0001
Total testosterone (ng/mL)	2.58	2.09	1.91	1.89	<0.0001	<0.0001
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	2625.69	612.55	3053.98	958.87	<0.0001	<0.0001
Erythrocyte sedimentation rate (ESR) (mm/hour)	17.72	14.75	25.45	21.55	<0.0001	<0.0001
Dehydroepiandrosterone sulfate (µg/dL)	91.21	69.26	75.51	63.29	0.0002	0.0003
Folate via RIA (ng/mL)	3.50	2.12	3.03	1.88	0.0002	0.0003
Free thyroxine, FT4 (ng/dL)	1.43	0.29	1.51	0.41	0.0002	0.0003
Neutrophils (%)	59.52	8.49	61.48	8.52	0.0002	0.0003
Soluble CD14 via ELISA (ng/mL)	1724.25	315.92	1810.47	383.4	0.0002	0.0003
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	122.04	54.74	109.52	53.64	0.0002	0.0003
TNF-related apoptosis-inducing ligand (pg/mL)	79.52	54.09	70.44	20.08	0.0005	0.0008
Endogenous secretory receptor for AGEs (ng/mL)	0.43	0.19	0.48	0.27	0.0005	0.0008
Omega-6 fatty acids as % of total fatty acid area	30.16	4.16	29.17	4.57	0.0005	0.0008
Lipids: LDL cholesterol (mg/dL)	139.09	35.77	132.56	32.7	0.0022	0.0032
Urinary creatinine (mg/dL)	73.94	35.12	67.37	31.9	0.0023	0.0033
Omega-3 plasma fatty acid weight (mg/L)	110.92	44.27	102.85	37.76	0.003	0.004
Lipids: total cholesterol (mg/dL)	220.84	40.73	213.53	38.74	0.0031	0.0042
Urinary Ca (mmol/L)	2.35	1.65	2.04	1.58	0.0036	0.0047
White blood cells (WBC) (n, K/µL)	6.01	1.56	6.29	1.63	0.0037	0.0048
Vitamin B6 via high performance liquid chromatography (ng/mL)	7.47	6.61	6.09	9.08	0.0057	0.0072
ALT (U/L)	20.47	11.99	18.43	12.05	0.0062	0.0076
Lycopene via high performance liquid chromatography (µmol/L)	0.71	0.34	0.65	0.34	0.0081	0.0097
Fatty acid C20:5 n-3 weight (mg/L)	20.46	9.87	18.95	7.51	0.0088	0.0103
Retinol via high performance liquid chromatography (µmol/L)	1.97	0.50	1.88	0.54	0.0103	0.0118
Urinary Na (mmol/L)	96.75	46.4	89.89	39.48	0.0153	0.0172
24-hour urinary cortisol (µg/24 hours)	105.33	52.21	95.94	73.57	0.0231	0.0255
Urine proteins (mg/dL)	0.73	7.61	1.92	8.98	0.0292	0.0315
Fatty acid C24:0 weight (mg/L)	4.66	4.51	4.05	4.11	0.0316	0.0331
Fatty acid C16:0 as % of total fatty acid weight	22.38	2.36	22.72	2.48	0.0319	0.0331
Fatty acid C16:0 as % of total fatty acid area	24.66	2.36	24.99	2.47	0.0408	0.0416
Fatty acid C20:5 n-3 as % of total fatty acid area	0.47	0.21	0.44	0.19	0.0471	0.0471

Table XV. Difference between healthy control and cognitive frailty Model I

Cognitive Frailty Model 1	Control Mean	SD	Cognitive Frailty Mean	SD	t -test	Corrected p-value
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	52.59	36.24	35.7	29.34	<0.0001	<0.0001
Adiponectin via RIA (µg/mL)	13.24	9.5	17.84	12.39	<0.0001	<0.0001
Albumin (%)	58.98	0.38	56.96	4.01	<0.0001	<0.0001
Alpha-1 globulin (%)	2.59	0.39	2.86	0.51	<0.0001	<0.0001
Alpha-2 globulin (%)	11.21	1.25	11.71	1.55	<0.0001	<0.0001
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	34.18	7.33	31.05	7.93	<0.0001	<0.0001
Blood urea nitrogen (mg/dL)	37.14	9.44	41.67	19.73	<0.0001	<0.0001
Fatty acid C20:5 n-3 as % of total fatty acid weight	0.63	0.23	0.55	0.17	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)	33.84	1	33.3	1.11	<0.0001	<0.0001
Creatine phosphokinase (U/L)	99.49	59.53	79.37	54.47	<0.0001	<0.0001
C-terminal telopeptide of type-1 collagen (ng/mL)	0.49	0.25	0.68	0.41	<0.0001	<0.0001
Cystatin C (mg/L)	0.99	0.26	1.26	0.51	<0.0001	<0.0001
Dehydroepiandrosterone sulfate (µg/dL)	87.58	67.99	66.59	58.9	<0.0001	<0.0001
Fibrinogen (mg/dL)	351.8	72.83	388.15	80.03	<0.0001	<0.0001
Homocysteine via FPIA analysis (µmol/L)	15.46	6.66	18.84	8.18	<0.0001	<0.0001
Red cell distribution width (RDW) (%)	13.66	0.94	14.15	1.31	<0.0001	<0.0001
Omega-3 plasma fatty acid weight (mg/L)	109.63	42.53	96.43	34.25	<0.0001	<0.0001
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	25.32	18.84	35.26	28.42	<0.0001	<0.0001
Resistin via EIA (ng/mL)	3.81	1.86	4.94	2.82	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	1430.03	579.89	2091.58	82.89	<0.0001	<0.0001
TNF-related apoptosis-inducing ligand (pg/mL)	77.35	44.29	65.53	19.93	<0.0001	<0.0001
24-hour urinary creatinine (mg/24 hours)	979.14	333.91	767.17	306.4	<0.0001	<0.0001
Fatty acid C20:5 n-3 as % of total fatty acid area	0.47	0.21	0.4	0.16	<0.0001	<0.0001
Lipids: LDL cholesterol (mg/dL)	138	33.95	127.04	34.78	<0.0001	<0.0001
Hemoglobin (g/dL)	13.9	1.29	12.95	1.6	<0.0001	<0.0001
Omega-6 fatty acids as % of total fatty acid area	29.98	4.23	28.41	4.77	<0.0001	<0.0001
Soluble CD14 via ELISA (ng/mL)	1741.7	334.78	1870.97	406.93	<0.0001	<0.0001
Total testosterone (ng/mL)	2.37	2.06	1.74	1.75	<0.0001	<0.0001
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	119.35	54.96	101.45	50.44	<0.0001	<0.0001
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	2709.69	709.84	3362.15	1054.91	<0.0001	<0.0001
Erythrocyte sedimentation rate (ESR) (mm/hour)	19.3	16.32	30.9	24.75	<0.0001	<0.0001
Vitamin E alpha tocopherol, high performance liquid chromatography (µmol/L)	30.7	8.31	27.17	8.37	<0.0001	<0.0001
Omega-6 fatty acids as % of total fatty acid mols	31.76	4.32	30.18	4.85	<0.0001	<0.0001
Lymphocytes (%)	30.92	8.02	28.56	8.17	0.0002	0.0003
Urine nitrites	0.1	0.42	0.32	0.71	0.0002	0.0003
Urinary Ca (mmol/L)	2.28	1.64	1.83	1.47	0.0004	0.0006
Ca++ (mg/dL)	9.46	0.45	9.32	0.5	0.0004	0.0006
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	151.95	111.77	194.04	178.49	0.0011	0.0016
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***	4279.38	1121.16	4009.81	1077.64	0.0018	0.0025
C-reactive protein - high sensitivity (µg/mL)	4.81	8.05	7.91	13.73	0.0018	0.0025
Free thyroxine, FT4 (ng/dL)	1.45	0.31	1.56	0.5	0.002	0.003
Beta-carotene via high performance liquid chromatography (µmol/L)	0.43	0.27	0.37	0.24	0.0039	0.0052
Beta globulins (%)	11.94	1.18	12.25	1.55	0.0065	0.0085
White blood cells (WBC) (n, K/µL)	6.08	1.55	6.44	1.76	0.007	0.0089
Mean platelet volume (MPV) (fL)	11.14	0.97	10.94	1	0.0079	0.0097
Fatty acid C16:0 as % of total fatty acid weight	22.44	2.36	22.98	2.62	0.008	0.0097
Uric acid (mg/dL)	5.13	1.37	5.47	1.76	0.009	0.0107
Soluble transferrin receptor (nmol/L)	16.66	5.65	18.3	8.56	0.0097	0.0113
Advanced glycation endproduct (AGE): Carboxymethyl-lysine (ng/mL)	361.53	105.78	390.73	152.77	0.0107	0.0122
Fatty acid C16:0 as % of total fatty acid area	24.42	2.36	25.24	2.61	0.0112	0.0125
Alpha-2-macroglobulin (mg/dL)	210.52	68.3	223.74	68.06	0.0122	0.0134
Serum creatinine (mg/dL)	0.92	0.19	0.98	0.38	0.0217	0.0234
Urine proteins (mg/dL)	0.98	7.78	2.8	10.35	0.0333	0.0352
Fatty acid C20:0 weight (mg/L)	2.87	2.84	2.52	1.94	0.0412	0.0427
Plasma insulin via RIA (mIU/L)	11.47	6.05	10.5	6.27	0.0429	0.0437
Lipids: HDL cholesterol (mg/dL)	56.27	14.72	53.8	16.43	0.0466	0.0466

Table XVI. Difference between healthy control and cognitive decline Model II

Cognitive Decline Model II	Control		Cognitive		Corrected		Control		Cognitive		Corrected			
	TrailA	Mean	SD	Mean	SD	t -test	p-value	TrailB	Mean	SD	Mean	SD	t -test	p-value
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)		60.15	35.34	45.59	35.96	<0.0001	<0.0001		65.04	35.67	50.09	35.12	<0.0001	<0.0001
Adiponectin via RIA (µg/mL)		12.02	8.69	15.3	11.07	<0.0001	<0.0001		11.26	8.43	14.02	10.52	<0.0001	<0.0001
Albumin (%)		60.08	3.24	58.34	3.47	<0.0001	<0.0001		60.9	31.4	58.86	3.32	<0.0001	<0.0001
Alpha-2-macroglobulin (mg/dL)		198.81	63.42	216.06	70.31	<0.0001	<0.0001		192.94	59.16	211.62	69.68	<0.0001	<0.0001
Alpha-1 globulin (%)		2.5	0.38	2.67	0.41	<0.0001	<0.0001		2.46	0.39	2.6	0.39	<0.0001	<0.0001
Alpha-2 globulin (%)		10.94	1.24	11.35	1.31	<0.0001	<0.0001		10.66	1.25	11.24	1.25	<0.0001	<0.0001
Beta globulins (%)		11.66	1.24	12.04	1.31	<0.0001	<0.0001		11.43	1.28	11.97	1.27	<0.0001	<0.0001
Blood urea nitrogen (mg/dL)		32.23	7.68	35.75	12.41	<0.0001	<0.0001		30.77	7.1	34.43	10.32	<0.0001	<0.0001
Fatty acid C16:0 weight (mg/L)		NA	NA	NA	NA	NA	NA		704.19	201.01	733.25	183.76	0.0341	0.0366
Fatty acid C20:5 n-3 as % of total fatty acid area		0.49	0.21	0.45	0.18	0.0011	0.0015		NA	NA	NA	NA	NA	NA
Fatty acid C20:5 n-3 weight (mg/L)		20.55	9.62	19.22	7.22	0.0097	0.0112		NA	NA	NA	NA	NA	NA
Fatty acid C20:5 n-3 as % of total fatty acid weight		0.64	0.24	0.59	0.19	0.0011	0.0015		NA	NA	NA	NA	NA	NA
Fatty acid C24:0 as % of total fatty acid weight		0.15	0.13	0.13	0.14	0.016	0.0176		0.16	0.14	0.14	0.13	0.0071	0.0085
Ca++ (mg/dL)		9.49	0.44	9.42	0.45	0.0093	0.0109		NA	NA	NA	NA	NA	NA
Creatinine clearance, 24-hr urine (mL/minute)		92.61	30.4	72.83	25.51	<0.0001	<0.0001		99.97	31.39	79.81	26.95	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)		34.09	0.94	33.64	1.02	<0.0001	<0.0001		34.26	0.93	33.8	0.99	<0.0001	<0.0001
Mean corpuscular hemoglobin (MCH) (pg)		30.67	1.74	30.45	1.96	0.0368	0.0385		NA	NA	NA	NA	NA	NA
Cortisol:DHEAS ratio (nmols)		0.25	0.58	0.43	1.54	0.0118	0.0133		0.23	0.61	0.45	1.25	0.0437	0.046
Creatine phosphokinase (U/L)		113.55	100.7	95.32	58.92	<0.0001	<0.0001		125.29	128.15	99.28	58.23	0.0004	0.0006
C-reactive protein - high sensitivity (µg/mL)		3.89	5.78	5.87	11.73	0.0004	0.0006		3.18	5.17	5	8.07	<0.0001	<0.0001
C-terminal telopeptide of type-1 collagen (ng/mL)		0.43	0.2	0.54	0.3	<0.0001	<0.0001		0.41	0.19	0.48	0.26	<0.0001	<0.0001
Cystatin C (mg/L)		0.89	0.2	1.06	0.35	<0.0001	<0.0001		0.84	0.19	0.99	0.28	<0.0001	<0.0001
Dehydroepiandrosterone sulfate (µg/dL)		124.75	101.28	85.52	70.91	<0.0001	<0.0001		153.71	115.09	91.51	72.62	<0.0001	<0.0001
Endogenous secretory receptor for AGEs (ng/mL)		0.43	0.21	0.46	0.22	0.0382	0.0393		NA	NA	NA	NA	NA	NA
Estradiol via radioimmunoassay (pg/mL)		14.79	17.95	9.25	6.72	<0.0001	<0.0001		18.56	21.94	9.89	0.34	<0.0001	<0.0001
Fibrinogen (mg/dL)		334.82	71.84	361.82	75.38	<0.0001	<0.0001		320.51	70.23	353.39	73.21	<0.0001	<0.0001
Free testosterone (ng/dL), Vermeulen		3.04	2.98	1.73	1.91	<0.0001	<0.0001		3.5	3.3	2.13	2.27	<0.0001	<0.0001
Free thyroxine, fT4 (ng/dL)		1.39	0.29	1.48	0.37	<0.0001	<0.0001		1.39	0.27	1.44	0.33	0.0081	0.0096
Blood glucose (mg/dL)		91.66	23.24	97.13	28.81	0.0004	0.0006		87.99	18.83	96.84	28.52	<0.0001	<0.0001
ALT (U/L)		21.76	13.91	18.88	11.88	0.0001	0.0001		NA	NA	NA	NA	NA	NA
Red blood cells (RBC) (n, millions/µL)		4.58	0.39	4.47	0.43	<0.0001	<0.0001		NA	NA	NA	NA	NA	NA
BL Hemoglobin (g/dL)		14.03	1.28	13.58	1.31	<0.0001	<0.0001		14.05	1.32	13.82	1.27	0.0056	0.007
BL Hematocrit (%)		41.13	3.28	40.36	3.5	<0.0001	<0.0001		NA	NA	NA	NA	NA	NA
Homocysteine via FPIA analysis (µmol/L)		13.91	5.44	16.26	7.09	<0.0001	<0.0001		13.39	5.22	15.17	6.22	<0.0001	<0.0001
Red cell distribution width (RDW) (%)		13.44	0.89	13.81	1	<0.0001	<0.0001		13.33	0.87	13.68	0.97	<0.0001	<0.0001
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)		NA	NA	NA	NA	NA	NA		365.8	143.93	399.41	151.46	0.0005	0.0007
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)		146.07	101.47	164.11	128.81	0.0078	0.0093		135.45	82.99	161.09	130.16	0.0001	<0.0001
Interleukin-6 via ELISA ultrasensitive (pg/mL)		1.55	1.82	2.47	5.54	0.0002	0.0003		1.31	1.84	1.97	2.18	<0.0001	<0.0001
Plasma insulin via RIA (mIU/L)		10.57	6.24	22.45	6.24	0.0161	0.0176		9.93	6.45	11.27	6.19	0.0016	0.0021
Lipoprotein(a) (mg/dL)		NA	NA	NA	NA	NA	NA		19.14	22.85	22.22	25.42	0.0495	0.0495
Methylmalonic acid, MMA (µmol/L)		0.1	0.03	0.11	0.03	0.0012	0.0016		0.1	0.03	0.11	0.03	0.0002	0.0003
Lymphocytes (n, K/µL)		1.96	0.65	1.85	0.65	0.0007	0.001		2.04	0.65	1.9	0.67	0.0012	0.0016
Na+ (mEq/L)		141.6	2.35	142.045	2.63	0.0019	0.0024		141.51	2.28	141.93	2.52	0.0064	0.0079

Cognitive Decline Model II	TrailA	Control	SD	Cognitive	SD	t -test	Corrected	TrailB	Control	SD	Cognitive	SD	t -test	Corrected
		Mean		Mean			p-value		Mean		Mean			p-value
Omega-3 fatty acids as % of total fatty acid area		2.09	0.62	1.88	0.57	<0.0001	<0.0001		2.16	0.67	1.97	0.59	<0.0001	<0.0001
Omega-3 plasma fatty acid weight (mg/L)		113.81	43.61	104.23	37.54	<0.0001	<0.0001		NA	NA	NA	NA	NA	NA
Ratio of Omega-6:Omega-3 as % of total fatty acid weight		3.53	0.98	3.21	0.93	<0.0001	<0.0001		3.64	1.04	3.34	0.94	<0.0001	<0.0001
Ratio of Omega-6:Omega-3 as % of total fatty acid weight		10.34	2.95	10.7	3.19	0.0479	0.0486		NA	NA	NA	NA	NA	NA
Omega-6 fatty acids as % of total fatty acid area		31.41	4.39	29.3	4.38	<0.0001	<0.0001		32.26	4.3	30.02	4.35	<0.0001	<0.0001
Omega-6 fatty acids as % of total fatty acid mols		33.15	4.43	331.08	4.46	<0.0001	<0.0001		33.99	0.25	31.8	4.42	<0.0001	<0.0001
Omega-6 plasma fatty acid weight (mg/L)		1082.37	243.69	1028.01	223.72	0.0001	0.0001		NA	NA	NA	NA	NA	NA
Omega-6 fatty acids as % of total fatty acid weight		34.05	4.43	31.98	4.47	<0.0001	<0.0001		34.89	4.31	32.7	4.43	<0.0001	<0.0001
Lymphocytes (%)		31.98	8.19	30.58	8.03	0.0028	0.0034		32.9	8.42	31.03	8.09	0.0007	0.001
Neutrophils (%)		59.12	8.81	60.47	8.4	0.0067	0.0081		58.2	9.19	60.03	8.52	0.002	0.0026
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)		22.24	17.65	27.87	19.24	<0.0001	<0.0001		22.33	22.59	25.03	14.54	0.0466	0.0474
Resistin via EIA (ng/mL)		3.71	1.72	4.13	2.23	0.0007	0.001		3.59	1.65	3.89	1.95	0.0126	0.0143
Retinol via high performance liquid chromatography (µmol/L)		1.97	0.48	1.88	0.49	0.0005	0.0007		1.99	0.47	1.91	0.49	0.0105	0.0121
Soluble CD14 via ELISA (ng/mL)		1651.2	339.4	1781.38	335.52	<0.0001	<0.0001		1595.57	318.07	1733.72	340.59	<0.0001	<0.0001
Soluble transferrin receptor (nmol/L)		15.98	5.49	16.99	5.48	0.0016	0.002		NA	NA	NA	NA	NA	NA
Total testosterone (ng/mL)		2.76	2.2	1.94	1.92	<0.0001	<0.0001		2.97	2.29	2.17	1.99	<0.0001	<0.0001
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)		147.92	72.25	109.88	51.83	<0.0001	<0.0001		164.2	78.49	121.24	57.78	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)		1200.34	443.28	1594.19	785.34	<0.0001	<0.0001		1101.48	441.51	1418.13	603.9	<0.0001	<0.0001
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)		2416.55	657.28	2869.39	827.95	<0.0001	<0.0001		2267.59	638.78	2695.09	747.26	<0.0001	<0.0001
Thyroid stimulating hormone, TSH (mIU/L)		1.66	2.24	2.36	7	0.0357	0.0379		NA	NA	NA	NA	NA	NA
Urinary Ca (mmol/L)		2.57	1.74	2.09	1.47	<0.0001	<0.0001		2.62	1.56	2.31	1.66	0.0054	0.0069
24-hour urinary cortisol (µg/24 hours)		108.88	55.55	100.63	68.22	0.0273	0.0294		111.52	50.52	102.66	58.56	0.0145	0.0161
Urinary cortisol (µg/mL)		NA	NA	NA	NA	NA	NA		0.08	0.06	0.07	0.05	0.0201	0.022
24-hour urinary creatinine (mg/24 hours)		1132.16	384.36	884.66	304.86	<0.0001	<0.0001		1211	383.23	977.63	348.15	<0.0001	<0.0001
Urinary creatinine (mg/dL)		81.52	39.44	67.67	32.17	<0.0001	<0.0001		88.75	42.01	71.27	34.64	<0.0001	<0.0001
Urinary Na (mmol/L)		101.12	45.54	92.5	41.92	0.0011	0.0015		106.71	46.11	93.3	41.77	<0.0001	<0.0001
Erythrocyte sedimentation rate (ESR) (mm/hour)		16.98	15.83	22.82	19.49	<0.0001	<0.0001		15.18	14.53	20.48	18.15	<0.0001	<0.0001
Mean corpuscular volume (MCV) (fL)		89.96	4.47	90.49	4.86	0.0492	0.0492		89.6	4.55	90.21	4.74	0.0458	0.0474
Vitamin B6 via high performance liquid chromatography (ng/mL)		8.32	5.57	6.56	6.64	<0.0001	<0.0001		9.34	5.91	6.93	6.24	<0.0001	<0.0001
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***		4497.55	1077.25	4229.28	1103.01	<0.0001	<0.0001		4595.71	993.38	4347.15	1145.3	0.0005	0.0007

Table XVII. Difference between healthy control and frailty Model II

Frailty Model II	Control Mean	SD	Frailty Mean	SD	t -test	Corrected p-value
Vitamin E alpha tocopherol, high performance liquid chromatography (µmol/L)	30.99	8.29	29.00	8.46	<0.0001	<0.0001
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)	34.44	7.65	32.65	7.39	<0.0001	<0.0001
Lymphocytes (%)	31.42	7.88	29.50	8.23	<0.0001	<0.0001
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)	54.93	34.5	43.52	35.77	<0.0001	<0.0001
Alpha-1 globulin (%)	2.57	0.36	2.72	0.47	<0.0001	<0.0001
Blood urea nitrogen (mg/dL)	33.79	7.44	37.5	15.92	<0.0001	<0.0001
Creatinine clearance, 24-hr urine (mL/minute)	81.09	24.06	70.00	26.43	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)	33.9	1.02	33.56	1.05	<0.0001	<0.0001
Creatine phosphokinase (U/L)	104.23	61.69	86.84	55.12	<0.0001	<0.0001
C-reactive protein - high sensitivity (µg/mL)	4.06	5.99	6.79	11.93	<0.0001	<0.0001
C-terminal telopeptide of type-1 collagen (ng/mL)	0.47	0.23	0.58	0.35	<0.0001	<0.0001
Free testosterone (ng/dL), Vermeulen	2.41	2.22	1.72	1.9	<0.0001	<0.0001
Red blood cells (RBC) (n, millions/µL)	4.57	0.38	4.42	0.48	<0.0001	<0.0001
Homocysteine via FPIA analysis (µmol/L)	14.97	5.70	17.32	8.12	<0.0001	<0.0001
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)	142.73	85.50	177.97	159.09	<0.0001	<0.0001
Interleukin-6 via ELISA ultrasensitive (pg/mL)	1.67	1.75	2.92	5.74	<0.0001	<0.0001
Neutrophils (n, K/µL)	3.60	1.18	3.90	1.31	<0.0001	<0.0001
Omega-6 plasma fatty acid weight (mg/L)	1069.54	249.81	1005.32	234.97	<0.0001	<0.0001
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)	24.06	19.79	30.55	22.59	<0.0001	<0.0001
Resistin via EIA (ng/mL)	3.72	1.67	4.36	2.48	<0.0001	<0.0001
Total testosterone (ng/mL)	2.58	2.09	1.91	1.89	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)	1343.02	429.62	1780.92	979.8	<0.0001	<0.0001
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)	2625.69	612.55	3053.98	958.87	<0.0001	<0.0001
24-hour urinary creatinine (mg/24 hours)	1020.45	334.70	860.38	323.47	<0.0001	<0.0001
Erythrocyte sedimentation rate (ESR) (mm/hour)	17.72	14.75	25.45	21.55	<0.0001	<0.0001
IL-6 high-sensitivity ELISA calculated from ELISA ultrasensitive (pg/mL)	3.11	2.00	4.23	2.82	<0.0001	<0.0001
Alpha-2-macroglobulin (mg/dL)	205.18	66.26	221.01	69.64	0.0002	0.0003
Dehydroepiandrosterone sulfate (µg/dL)	91.21	69.26	75.51	63.3	0.0002	0.0003
Folate via RIA (ng/mL)	3.50	2.12	3.03	1.88	0.0002	0.0003
Free thyroxine, fT4 (ng/dL)	1.43	0.30	1.51	0.41	0.0002	0.0003
Soluble CD14 via ELISA (ng/mL)	1724.25	315.92	1810.47	383.4	0.0002	0.0003
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)	122.04	54.74	109.52	53.63	0.0002	0.0003
Endogenous secretory receptor for AGEs (ng/mL)	0.43	0.19	0.48	0.27	0.0005	0.0008
Omega-6 fatty acids as % of total fatty acid area	30.16	4.16	29.17	4.57	0.0005	0.0008
TNF-related apoptosis-inducing ligand (pg/mL)	79.52	54.09	70.44	20.08	0.0005	0.0008
Lipids: LDL cholesterol (mg/dL)	139.01	35.77	132.56	32.70	0.0022	0.0034
Omega-3 plasma fatty acid weight (mg/L)	110.92	44.27	102.85	37.76	0.0029	0.0043
Lipids: total cholesterol (mg/dL)	220.84	40.73	213.53	38.74	0.0031	0.0045
Urinary Ca (mmol/L)	2.35	1.65	2.04	1.58	0.0036	0.0051
White blood cells (WBC) (n, K/µL)	6.01	1.56	6.3	1.63	0.0037	0.0051
Vitamin B6 via high performance liquid chromatography (ng/mL)	7.47	6.61	6.09	9.08	0.0057	0.0076
ALT (U/L)	20.48	11.99	18.44	12.05	0.0062	0.0081
Fatty acid C20:5 n-3 weight (mg/L)	20.46	9.87	18.95	7.51	0.0088	0.0113
Adiponectin via RIA (µg/mL)	13.31	9.72	15.05	10.83	0.0094	0.0118
Omega-3 fatty acids as % of total fatty acid area	1.99	0.62	1.89	0.59	0.0141	0.0172
Urinary Na (mmol/L)	96.75	46.40	89.89	39.48	0.0153	0.0183
Uric acid (mg/dL)	5.09	1.29	5.30	1.60	0.0175	0.0205
24-hour urinary cortisol (µg/24 hours)	105.33	52.21	95.94	73.57	0.0231	0.0265
Fatty acid C20:5 n-3 cis (eicosapentaenoic, EPA) area	79.94	51.57	73.05	43.36	0.0275	0.0309
Urine proteins (mg/dL)	0.73	7.61	1.93	8.98	0.0292	0.0321
Fatty acid C24:0 weight (mg/L)	4.65	4.51	4.05	4.11	0.0316	0.0337
Fatty acid C16:0 as % of total fatty acid weight	22.38	2.36	22.72	2.48	0.0319	0.0337
Fatty acid C16:0 as % of total fatty acid area	24.66	2.36	24.98	2.46	0.0408	0.0423
Omega-3 fatty acids as % of total fatty acid weight	3.35	0.97	3.23	0.95	0.0457	0.0465
Fatty acid C20:5 n-3 as % of total fatty acid area	0.47	0.22	0.44	0.19	0.0471	0.0471

Table XVIII. Difference between healthy control and cognitive frailty Model II

	TrailA	Control Mean	SD	Cognitive Frailty Mean	SD	t -test	Corrected p-value	TrailB	Control Mean	SD	Cognitive Frailty Mean	SD	t -test	Corrected p-value
Cognitive Frailty Model II														
25(OH)-D (25-hydroxyvitamin D) via RIA (nmol/L)		57.82	36.01	40.97	34.34	<0.0001	<0.0001		58.33	33.73	47.92	40.02	<0.0001	<0.0001
Adiponectin via RIA (µg/mL)		12.66	9.11	15.85	11.71	<0.0001	<0.0001		12.41	9.13	14.66	11.45	0.0028	0.0034
Albumin (%)		59.71	3.34	58.09	3.51	<0.0001	<0.0001		59.99	3.27	58.52	3.45	<0.0001	<0.0001
Alpha-2-macroglobulin (mg/dL)		201.68	64.3	221.07	73.22	<0.0001	<0.0001		197.37	61.88	223.54	74.06	<0.0001	<0.0001
Alpha-1 globulin (%)		2.53	0.38	2.69	0.43	<0.0001	<0.0001		2.51	0.39	2.64	0.43	<0.0001	<0.0001
Vitamin E alpha tocopherol, high performance liquid chromatography (µmol/L)		30.47	8.24	29.01	8.39	0.0092	0.01		NA	NA	NA	NA	NA	NA
Vitamin E alpha tocopherol, high performance liquid chromatography, assay #2 (µmol/L)		33.91	7.54	32.86	7.43	0.0367	0.0367		NA	NA	NA	NA	NA	NA
Blood urea nitrogen (mg/dL)		32.61	7.77	37.42	14.90	<0.0001	<0.0001		32.12	7.20	0.27	13.06	<0.0001	<0.0001
Fatty acid C20:5 n-3 as % of total fatty acid area		0.48	0.20	0.19	0.19	0.027	0.0275		NA	NA	NA	NA	NA	NA
Creatinine clearance, 24-hr urine (mL/minute)		88.69	29.71	68.68	25.64	<0.0001	<0.0001		9.48	30.34	74.67	25.66	<0.0001	<0.0001
MCH concentration (MCHC) (g/dL)		34.01	1.00	33.54	0.96	<0.0001	<0.0001		34.07	1.00	33.67	0.96	<0.0001	<0.0001
Serum cortisol (µg/dL)		NA	NA	NA	NA	NA	NA		13.67	5.06	12.73	4.37	0.0026	0.0033
Creatine phosphokinase (U/L)		110.89	92.60	88.84	53.96	<0.0001	<0.0001		115.10	99.11	91.46	54.64	<0.0001	<0.0001
C-reactive protein - high sensitivity (µg/mL)		4.01	6.14	7.11	14.37	0.0004	0.0005		3.77	6.02	5.83	9.44	0.0004	0.0006
C-terminal telopeptide of type-1 collagen (ng/mL)		0.44	0.21	0.59	0.35	<0.0001	<0.0001		0.43	0.20	0.53	0.31	<0.0001	<0.0001
Cystatin C (mg/L)		0.91	0.21	1.13	0.42	<0.0001	<0.0001		0.88	0.19	1.07	0.35	<0.0001	<0.0001
Dehydroepiandrosterone sulfate (µg/dL)		116.98	95.79	77.99	67.71	<0.0001	<0.0001		125.8	100.15	80.97	67.13	<0.0001	<0.0001
Endogenous secretory receptor for AGEs (ng/mL)		0.43	0.2	0.48	0.25	0.0077	0.0083		0.43	0.18	0.48	0.27	0.0086	0.0094
Estradiol via radioimmunoassay (pg/mL)		13.29	15.99	9.52	7.18	<0.0001	<0.0001		14.30	17.34	9.35	6.52	<0.0001	<0.0001
Fibrinogen (mg/dL)		339.88	72.16	367.95	78.12	<0.0001	<0.0001		334.56	73.08	360.54	72.38	<0.0001	<0.0001
Folate via RIA (ng/mL)		3.41	2.14	3.02	1.70	0.0013	0.0017		3.43	2.14	3.07	1.91	0.0078	0.0087
Free testosterone (ng/dL), Vermeulen		2.72	2.80	1.67	1.92	<0.0001	<0.0001		2.95	2.96	1.74	1.89	<0.0001	<0.0001
Blood glucose (mg/dL)		NA	NA	NA	NA	NA	NA		92.18	24.40	97.77	29.05	0.0028	0.0034
Hematocrit (%)		41.02	3.28	40.06	3.64	<0.0001	<0.0001		41.07	3.23	40.51	3.48	0.0147	0.0153
Homocysteine via FPIA analysis (µmol/L)		14.16	5.57	17.39	7.78	<0.0001	<0.0001		13.88	5.59	16.17	6.47	<0.0001	<0.0001
Interleukin-18 via ELISA ultrasensitive using plasma (pg/mL)		386.20	149.66	411.61	156.28	0.0146	0.0154		382.19	150.56	402.02	147.03	0.0478	0.0478
Interleukin-1 receptor antagonist via ELISA ultrasensitive (pg/mL)		146.60	97.39	177.16	154.61	0.0015	0.0019		142.80	95.94	174.54	153.13	0.0007	0.001
Interleukin-6 via ELISA ultrasensitive (pg/mL)		1.57	1.81	3.14	7.22	0.0003	0.0004		1.46	1.77	2.42	2.57	<0.0001	<0.0001
Lycopene via high performance liquid chromatography (µmol/L)		0.71	0.34	0.65	0.31	0.0042	0.005		NA	NA	NA	NA	NA	NA
Lymphocytes (n, K/µL)		1.95	0.64	1.83	0.67	0.0073	0.0085		1.98	0.65	1.87	0.69	0.0184	0.0188
Omega-3 fatty acids as % of total fatty acid area		2.04	0.62	1.88	0.56	0.0002	0.0003		2.07	0.63	1.96	0.60	0.0049	0.0056
Omega-6 plasma fatty acid weight (mg/L)		1069.85	241.60	1022.35	216.50	0.0024	0.0029		1086.09	239.82	1034.56	223.86	0.0016	0.0021
Omega-6 fatty acids as % of total fatty acid weight		33.57	4.45	31.76	4.64	<0.0001	<0.0001		33.93	4.42	32.32	4.52	<0.0001	<0.0001
Lymphocytes (%)		31.96	8.07	29.53	8.13	<0.0001	<0.0001		32.45	8.16	29.87	8.16	<0.0001	<0.0001
Neutrophils (%)		59.15	8.64	61.46	8.47	<0.0001	<0.0001		58.64	8.83	61.19	8.42	<0.0001	<0.0001
Parathyroid hormone, two-site immunoradiometric assay (pg/mL)		22.68	16.32	31.12	23.23	<0.0001	<0.0001		22.36	17.44	28.13	17.47	<0.0001	<0.0001
Resistin via EIA (ng/mL)		3.75	1.87	4.33	2.19	<0.0001	<0.0001		3.67	1.66	4.05	2.23	0.0094	0.01
Soluble CD14 via ELISA (ng/mL)		1670.14	331.90	1824.72	386.26	<0.0001	<0.0001		1653.78	323.41	1760.93	361.97	<0.0001	<0.0001
Total testosterone (ng/mL)		2.57	2.17	1.85	1.87	<0.0001	<0.0001		2.68	2.20	1.87	1.84	<0.0001	<0.0001
Total insulin-like growth factor-1, serum, immunoradiometric assay (ng/mL)		139.07	69.69	106.54	49.24	<0.0001	<0.0001		145.42	71.16	113.21	55.61	<0.0001	<0.0001
Soluble TNF-α receptor I via quantitative sandwich EIA (pg/mL)		1248.81	471.88	1763.25	914.92	<0.0001	<0.0001		1191.01	432.04	1592.49	741.76	<0.0001	<0.0001
Soluble TNF-α receptor II via quantitative sandwich EIA (pg/mL)		2473.75	654.70	3059.1	924.41	<0.0001	<0.0001		2399.93	623.03	2903.60	862.17	<0.0001	<0.0001
TNF-related apoptosis-inducing ligand (pg/mL)		76.51	42.76	71.47	19.74	0.0064	0.0074		NA	NA	NA	NA	NA	NA
Urinary Ca (mmol/L)		2.50	1.74	1.90	1.18	<0.0001	<0.0001		2.52	1.63	2.15	1.64	0.0001	0.0001
24-hour urinary cortisol (µg/24 hours)		108.98	32.17	93.34	57.87	0.0001	0.0001		109.66	50.09	96.01	67.41	0.0018	0.0023
Urinary cortisol (µg/mL)		1082.84	374.84	833.83	294.48	<0.0001	<0.0001		1119.54	381.66	902.09	314.87	<0.0001	<0.0001
24-hour urinary creatinine (mg/24 hours)		1082.84	374.84	833.83	294.48	<0.0001	<0.0001		1119.54	381.66	902.09	314.87	<0.0001	<0.0001
Urinary creatinine (mg/dL)		78.09	38.10	66.78	32.12	<0.0001	<0.0001		80.74	39.52	68.38	33.02	<0.0001	<0.0001
Urinary Na (mmol/L)		99.50	45.59	90.34	38.51	0.0014	0.0018		101.58	44.78	88.47	39.39	<0.0001	<0.0001
Uric acid (mg/dL)		4.98	1.28	5.22	1.56	0.0148	0.0154		4.93	1.29	5.19	1.36	0.0036	0.0042
Erythrocyte sedimentation rate (ESR) (mm/hour)		17.59	15.96	25.67	21.42	<0.0001	<0.0001		16.67	14.84	23.32	20.66	<0.0001	<0.0001
Vitamin B6 via high performance liquid chromatography (ng/mL)		8.12	6.21	5.75	5.47	<0.0001	<0.0001		8.48	6.28	6.08	5.79	<0.0001	<0.0001
IGF binding protein-3, serum, immunoradiometric assay (ng/mL) ***corrected***		4452.55	1077.41	4158.20	1124.84	0.0001	0.0001		4517.72	1060.85	4238.12	1166.19	0.0004	0.0006

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Drugs with ACB Score of 1

Generic Name	Brand Name
Alimemazine	Theralen™
Alverine	Spasmonal™
Alprazolam	Xanax™
Aripiprazole	Abilify™
Asenapine	Saphris™
Atenolol	Tenormin™
Bupropion	Wellbutrin™, Zyban™
Captopril	Capoten™
Cetirizine	Zyrtec™
Chlorthalidone	Diuril™, Hygroton™
Cimetidine	Tagamet™
Cildinium	Librax™
Clorazepate	Tranxene™
Codeine	Contin™
Colchicine	Colcrys™
Desloratadine	Clarinetx™
Diazepam	Valium™
Digoxin	Lanoxin™
Dipyridamole	Persantine™
Disopyramide	Norpace™
Fentanyl	Duregesic™, Actiq™
Furosemide	Lasix™
Fluvoxamine	Luvox™
Haloperidol	Haldol™
Hyalalazine	Aprisoline™
Hydrocortisone	Cortef™, Cortaid™
Iloperidone	Fenapt™
Isosorbide	Isordil™, Ismo™
Levocetirizine	Xyzal™
Loperamide	Immodium™, others
Loratadine	Claritin™
Metoprolol	Lopressor™, Toprol™
Morphine	MS Contin™, Avinza™
Nifedipine	Procardia™, Adalat™
Paliperidone	Invega™
Prednisone	Deltasone™, Sterapred™
Quinidine	Quinaglute™
Ranitidine	Zantac™
Risperidone	Risperdal™
Theophylline	Theodur™, Uniphyll™
Trazodone	Desyrel™
Triamterene	Dyrenium™
Venlafaxine	Effexor™
Warfarin	Coumadin™

Drugs with ACB Score of 2

Generic Name	Brand Name
Amantadine	Symmetrel™
Belladonna	Multiple
Carbamazepine	Tegretol™
Cyclobenzaprine	Flexeril™
Cyproheptadine	Pericort™
Loxapine	Loxitane™
Meperidine	Demerol™
Methotrimeprazine	Levoprome™
Molindone	Moban™
Nefopam	Nefogestic™
Oxcarbazepine	Trileptal™
Pimozide	Orap™

Drugs with ACB Score of 3

Generic Name	Brand Name
Amiripityline	Elavil™
Amoxapine	Asendin™
Atropine	Sal-Tropine™
Benztropine	Cogentin™
Brompheniramine	Dimetapp™
Carinoxamine	Histex™, Carbihist™
Chlorpheniramine	Chlor-Trimeton™
Chlorpromazine	Thorazine™
Clemastine	Tavist™
Clomipramine	Anafranil™
Clozapine	Clozaril™
Darifenacin	Enablex™
Desipramine	Norpramin™
Dicyclomine	Bentyl™
Dimenhydrinate	Dramamine™, others
Diphenhydramine	Benadryl™, others
Doxepin	Sinequan™
Doxylamine	Unisom™, others
Fesoterodine	Toviaz™
Flavoxate	Urispas™
Hydroxyzine	Atarax™, Vistaril™
Hyoscyamine	Anaspaz™, Levsin™
Imipramine	Tofranil™
Mecizine	Antivert™
Methocarbamol	Robaxin™
Nortriptyline	Pamelor™
Olanzapine	Zyprexa™
Orphenadrine	Norflex™
Oxybutynin	Ditropan™
Paroxetine	Paxil™
Perphenazine	Trilafon™
Promethazine	Phenergan™
Propantheline	Pro-Banthine™
Propiverine	Detrol™
Quetiapine	Seroquel™
Scopolamine	Transderm Scop™
Solifenacin	Vesicare™
Thioridazine	Mellaril™
Tolterodine	Detrol™
Trifluoperazine	Stelazine™
Trihexyphenidyl	Artane™
Trimipramine	Surmontil™
Trospium	Sanctura™

Categorical Scoring:

- Possible anticholinergics include those listed with a score of 1; Definite anticholinergics include those listed with a score of 2 or 3
- Numerical Scoring:
 - Add the score contributed to each selected medication in each scoring category
 - Add the number of possible or definite Anticholinergic medications

Notes:

- Each definite anticholinergic may increase the risk of cognitive impairment by 46% over 6 years.³
- For each on point increase in the ACB total score, a decline in MMSE score of 0.33 points over 2 years has been suggested.⁴
- Additionally, each one point increase in the ACB total score has been correlated with a 26% increase in the risk of death.⁴

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Medications Reviewed in 2012 Update

Medications Added with Score of 1:	Medications Added with Score of 2:
Aripiprazole (Abilify™)	Neopam (Nefogest™)
Asenapine (Saphris™)	
Cetirizine (Zyrtec™)	Medications Added with Score of 3:
Clidinium (Librax™)	Doxylamine (Unisom™, others)
Desloratadine (Clarinetx™)	Fesoterodine (Toviaz™)
Iloperidone (Fenapt™)	Propiverine (Detrunorm™)
Levocetirizine (Xyzal™)	Solifenacin (Vesicare™)
Loratadine (Claritin™)	Trospium (Sanctura™)
Paliperidone (Invega™)	
Venlafaxine (Effexor™)	

Medications Reviewed But NOT Added:
Fexofenadine (Allegra™)
Gabapentin (Neurontin™)
Topiramate (Topamax™)
Levetiracetam (Keppra™)
Tamoxifen (Nolvadex™)
Nizatidine (Axiid™)
Duloxetine (Cymbalta™)

Criteria for Categorization:

Score of 1: Evidence from in vitro data that chemical entity has antagonist activity at muscarinic receptor.

Score of 2: Evidence from literature, prescriber's information, or expert opinion of clinical anticholinergic effect.

Score of 3: Evidence from literature, expert opinion, or prescribers information that medication may cause delirium.

Aging Brain Care

ANTICHOLINERGIC COGNITIVE BURDEN SCALE

2012 Update

Developed by the Aging Brain Program
of the Indiana University Center for
Aging Research



Complete References:

1. Boustani MA, Campbell NL, Munger S, Maidment J, Fox GC. Impact of anticholinergics on the aging brain: a review and practical application. *Aging Health*. 2008;4(3):311-320.
2. Campbell N, Boustani M, Limbil T, et al. The cognitive impact of anticholinergics: a clinical review. *Clinical Interventions in Aging*. 2009;4(1):225-233.
3. Campbell N, Boustani M, Lane K, et al. Use of anticholinergics and the risk of cognitive impairment in an African-American population. *Neurology*. 2010;75:152-159.
4. Fox C, Richardson K, Maidment J, et al. Anticholinergic medication use and cognitive impairment in the older population: the Medical Research Council Cognitive Function and Ageing Study. *Journal of the American Geriatric Society*. 2011; 59(8): 1477-1483.
5. Cai X, Campbell N, Khan B, Callahan C, Boustani M. Long-term anticholinergic use and the aging brain. *Alzheimers Dementia*. 2012; epub ahead of print.

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Use of the Anti-Cholinergic Burden (ACB) Scale may only be in accordance with the Terms of Use for the ACB Scale which are available at <http://www.agingbraincare.org/tools/abc-anticholinergic-cognitive-burden-scale>.

To request permission for use, contact us at acb@agingbraincare.org.

Anticholinergic Burden Scale script with instructions for research assistant and/or participant permission to use instrument from author

Anticholinergic Burden Scale Permission

We do not have a formal letter. You can use the following email:

You have permission to use the Anticholinergic Cognitive Burden Scale for your dissertation related work including both research and educational purposes.

Malaz

Malaz Boustani, MD, MPH

IRB APPROVAL



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Office of Research Subjects Protection
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TO: Patricia Slattum
CC: Lana Sargent

FROM: VCU IRB Panel B

RE: Patricia Slattum ; IRB [HM20006652](#) Predicting cognitive frailty: a population modeling study

On 2/3/2016 the referenced research study **qualified for exemption** according to 45 CFR 46.101(b), category 4.

The information found in the electronic version of this study's smart form and uploaded documents now represents the currently approved study, documents, and HIPAA pathway (if applicable). You may access this information by clicking the Study Number above.

If you have any questions, please contact the Office of Research Subjects Protection (ORSP) or the IRB reviewer(s) assigned to this study.

- The reviewer(s) assigned to your study will be listed in the History tab and on the study workspace. Click on their name to see their contact information.

Attachment – Conditions of Exempt Approval **Conditions of Exempt Approval:**

In order to comply with federal regulations, industry standards, and the terms of this approval, the investigator must (as applicable):

1. Conduct the research as described in and required by the Protocol.
2. Provide non-English speaking patients with a translation of the approved Consent Form in the research participant's first language. The Panel must approve the translation.
3. The following changes to the protocol **must be** submitted to the IRB panel for review and approval before the changes are instituted. Changes that do not meet these criteria do not have to be submitted to the IRB. If there is a question about whether a change must be sent to the IRB please call the ORSP for clarification.

THESE CHANGES MUST BE SUBMITTED:

- Change in principal investigator
- Any change that increases the risk to the participant

TRAINING AIM DOCUMENT

