

The Metabolic Syndrome and Cognitive Performance: The Northern Manhattan Study

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Key Words

Cognitive performance · Cognitive impairment · Vascular dementia · Vascular cognitive impairment · Cerebrovascular disorders · Metabolic syndrome

Abstract

Background: The metabolic syndrome (MetS) is a risk factor for diabetes, stroke, myocardial infarction, and increased mortality, and has been associated with cognition in some populations. We hypothesized that MetS would be associated with lower Mini-Mental State Examination (MMSE) scores in a multi-ethnic population, and that MetS is a better predictor of cognition than its individual components or diabetes. **Methods:** We conducted a cross-sectional analysis among 3,150 stroke-free participants. MetS was defined by the modified National Cholesterol Education Program guidelines-Adult Treatment Panel III (NCEP-ATPIII) criteria. Linear regression and polytomous logistic regression estimated the association between MMSE score and MetS, its individual components, diabetes, and inflammatory biomarkers. **Results:** MetS was inversely associated with MMSE score (unadjusted $\beta = -0.67$; 95% CI $-0.92, -0.41$). Adjusting for potential confounders, MetS was associated with lower

MMSE score (adjusted $\beta = -0.24$; 95% CI $-0.47, -0.01$), but its individual components and diabetes were not. Those with MetS were more likely to have an MMSE score of <18 than a score of ≥ 24 (adjusted OR = 1.94; 95% CI 1.26, 3.01). There was an interaction between MetS and race-ethnicity, such that MetS was associated with lower MMSE score among non-Hispanic whites and Hispanics but not non-Hispanic blacks. **Conclusions:** MetS was associated with lower cognition in a multi-ethnic population. Further studies of the effect of MetS on cognition are warranted, and should account for demographic differences. Copyright © 2011 S. Karger AG, Basel

Introduction

The metabolic syndrome (MetS) is a risk factor for diabetes, stroke, myocardial infarction [1], and increased mortality [2, 3]. Previous studies demonstrated an association between cardiovascular risk factors (including hypertension and diabetes) and cognitive aging [4]; additionally, insulin resistance and abnormal cholesterol metabolism have been connected to memory loss and the development of Alzheimer disease and vascular dementia [5, 6].

The relationship between MetS and cognition, however, remains uncertain, and appears to depend in part on demographics and levels of inflammatory biomarkers. Moreover, it is unclear if MetS is a better predictor of cognitive performance than its individual components or diabetes alone. Few studies, however, have looked separately at the MetS components individually and as a whole [7, 8]. Previous studies evaluated non-Hispanic whites, and only one study assessed a Hispanic population composed of Mexican-Americans [9]. None of the previous studies evaluated the cognitive effects of MetS among Caribbean Hispanics, a group for whom MetS is quite prevalent [10].

The purpose of this study was to investigate the relationship between MetS and cognitive performance using the Mini-Mental State Examination (MMSE) [11] in a multi-ethnic cohort. We sought, furthermore, to determine if MetS is a better predictor of cognitive performance than its individual components or diabetes, and if inflammatory biomarkers modify this association.

Subjects and Methods

Cohort Selection and Evaluation

The cohort for this analysis was derived from the participants enrolled in the Northern Manhattan Study (NOMAS) identified by random-digit dialing and recruited for an in-person assessment, as previously described [10]. Baseline cognitive assessment was performed using the 30-point MMSE in English and Spanish, depending on the participant's language spoken at home. The study was approved by the Institutional Review Board at Columbia University and the University of Miami and participants provided informed consent.

Metabolic Syndrome

MetS was defined by the modified National Cholesterol Education Program guidelines-Adult Treatment Panel III (NCEP-ATPIII) criteria represented by the presence of 3 or more of the 5 components linked to insulin resistance: (1) waist circumference ≥ 102 cm in men and ≥ 88 cm in women; (2) triglycerides ≥ 150 mg/dl; (3) high-density lipoprotein (HDL) < 40 mg/dl in men and < 50 mg/dl in women; (4) fasting plasma glucose ≥ 100 mg/dl, and (5) systolic blood pressure ≥ 130 mm Hg, or diastolic blood pressure ≥ 85 mm Hg. Blood pressure was based on the average of two measurements, or a subject self-reported history of hypertension, as previously described [10].

Covariates

Risk factors, behavioral and sociodemographic data that might influence cognition were collected as described in previous publications [10].

Education was defined based on whether or not high school was completed. Health insurance status was dichotomized to Medicaid/no insurance versus private insurance/Medicare, as previously described [12]. Moderate alcohol intake was defined as currently drinking > 1 drink/month and ≤ 2 drinks/day [13].

Smoking was defined as current smokers. Physical activity was defined by the amount of leisure activity engaged in during the 10 days prior to assessment. Social support was assessed based on marital status and knowing three or more people well enough to visit with in their homes [14].

Laboratory Assessments

Fasting blood specimens were drawn at baseline into serum tubes and spun within one hour at 3000 g and 4°C for 20 min and immediately frozen at -70°C . HDL and triglyceride levels were measured using an automated spectrometer (Hitachi 705, Boehringer, Mannheim, Germany) as described previously [12]. Inflammatory marker levels were measured in batched samples and assays were performed blinded to MMSE. Serum amyloid A (SAA) and high-sensitivity C-reactive protein (hsCRP) were measured using the BNII nephelometric assay system (Dade-Behring, Deerfield, Ill., USA) [15]. Interleukin-6 (IL-6) and tumor necrosis factor receptor 1 (TNFR1) were measured by enzyme-linked immunosorbent assays according to manufacturer instructions (BioSource International, Camarillo, Calif., USA), as previously described [16].

Statistical Analyses

Baseline characteristics were compared in relation to the MetS presence using χ^2 tests for proportions, t tests for continuous variables and Wilcoxon rank test for medians. Linear regressions were conducted to calculate β and 95% CIs for MMSE scores as a continuous measure, and polytomous logistic regressions with canonical link to calculate odds ratio (OR) and 95% CI for categorized MMSE. We analyzed MMSE as a continuous outcome, as well as according to categories based on previously defined MMSE thresholds [17] – MMSE score ≥ 24 (normal, reference), MMSE score 18–23 (mild to moderate cognitive impairment), and MMSE score < 18 (severe cognitive impairment) – in order to facilitate clinical interpretation and to permit comparison with previous studies.

We used MetS as a main predictor, and the five individual components of MetS and diabetes as secondary predictors of interest.

Unadjusted and adjusted models for sociodemographic factors (age, sex, race-ethnicity, education) and additional risk factors (moderate alcohol consumption, smoking, social support, marriage status and physical activity) were constructed. Covariates were included in the models based on a priori hypotheses.

Interactions between MetS and sociodemographic factors were tested and stratified models were constructed as indicated. Further models incorporating SAA, hsCRP, TNFR1 and IL-6 were constructed to assess for independent effects of these markers, as well as interactions with MetS on MMSE score. Analyses were conducted using SAS v9.1.3 (SAS Institute, Cary, N.C., USA) and statistical significance defined as $p \leq 0.05$.

Results

Cohort Description

Both data on MetS and MMSE were available for 3,150 participants. Their baseline characteristics are shown in table 1. Mean age was 69.0 ± 10.3 years. There were 37% men, 53% Hispanics, 21% non-Hispanic whites and 24% non-Hispanic blacks.

Table 1. Descriptive analysis

Characteristics	Overall	MetS		p value ^a
		no	yes	
n (%)	3,150	1,649 (52.3)	1,501 (47.7)	
Sociodemographic characteristics				
Mean age \pm SD, years	69.0 \pm 10.3	69.4 \pm 11.0	68.6 \pm 9.4	0.04
Male	1,180 (37.5)	701 (22.3)	479 (15.2)	<0.0001
Race and ethnicity ^b				
Non-Hispanic white	660 (21.0)	383 (23.2)	277 (18.5)	ref.
Non-Hispanic black	762 (24.2)	447 (27.1)	315 (21.0)	0.81
Hispanic	1,656 (52.6)	765 (46.4)	891 (59.4)	<0.0001
Other risk factors				
High school graduate	1,450 (46.0)	851 (27.0)	598 (19.0)	<0.0001
Medicaid or no insurance	1,361 (43.2)	617 (19.6)	744 (23.6)	<0.0001
Married	1,005 (31.9)	537 (17.0)	468 (14.9)	0.40
3 friends or more	2,684 (85.2)	1,402 (44.5)	1,282 (40.7)	0.76
Any physical activity	1,834 (58.2)	1,033 (32.8)	801 (25.4)	<0.0001
Mild/moderate alcohol intake	1,031 (32.7)	615 (19.5)	416 (13.2)	<0.0001
Current smoker	548 (17.4)	295 (9.4)	253 (8.0)	0.55
Former smoker	1,126 (35.7)	582 (18.5)	544 (17.3)	0.74
Never smoker	1,475 (46.8)	772 (24.5)	703 (22.3)	ref.
MetS				
Glucose \geq 100 mg/dl ^c	1,095 (34.8)	230 (7.3)	865 (27.5)	<0.0001
BP \geq 130/85 mm Hg, or HTN ^d	2,659 (84.4)	1,214 (38.5)	1,445 (45.9)	<0.0001
Abdominal obesity ^e	1,329 (42.2)	318 (10.1)	1,011 (32.1)	<0.0001
Low HDL ^f	1,702 (54.0)	440 (14.0)	1,262 (40.0)	<0.0001
Triglycerides \geq 150 mg/dl	931 (29.5)	117 (3.7)	814 (25.8)	<0.0001
History of diabetes	686 (21.8)	164 (5.2)	522 (16.6)	<0.0001
History of hypertension	2,318 (73.6)	1,023 (32.5)	1,295 (41.1)	<0.0001
Cognitive performance				
Median MMSE score (IQR)	27 (24–29)	27 (25–29)	27 (24–29)	<0.0001
Non-Hispanic whites	28 (27–30)	29 (27–30)	28 (26.5–29)	0.0003
Non-Hispanic blacks	27 (25–29)	27 (25–29)	28 (25–29)	0.1031
Hispanics	26 (23–28)	27 (24–28)	26 (23–28)	0.0007

Values denote numbers with percentages in parentheses, unless otherwise indicated.

^a t test for continuous and χ^2 for dichotomous variables. ^b Other race/ethnicity had 72 participants representing 2.3% of the sample. ^c History of hypertension. ^d Systolic BP \geq 130 mm Hg, diastolic \geq 85 mm Hg; also includes history of hypertension (HTN). ^e Abdominal obesity defined as waist $>$ 89.41 cm (35.2 inches) for women or $>$ 103.63 cm (40.8 inches) for men. ^f HDL $<$ 50 mg/dl for women or $<$ 40 mg/dl for men.

IQR = Interquartile range.

The MetS prevalence in our cohort was 47.6%, and was similar among non-Hispanic whites and non-Hispanic blacks (41.9 and 41.3%, respectively), but greater among Hispanics (53.8%; $p < 0.0001$). The median MMSE score was 27 (IQR: 24–29), and non-Hispanic blacks and Hispanics had lower MMSE scores than non-Hispanic whites ($p < 0.0001$; table 1).

MetS and MMSE Score

Those with MetS had lower cognitive performance scores than those without MetS (unadjusted $\beta = -0.67$;

95% CI $-0.92, -0.41$). The association persisted after adjusting for age, sex, race-ethnicity, education, alcohol consumption, smoking, social support, marriage status, and physical activity ($\beta = -0.24$; 95% CI $-0.47, -0.01$; table 2). To put our estimates regarding MetS in perspective, we examined differences in MMSE score by age. Scores were lower by 0.096 (SE = 0.006) points per 1 year increase in age. Thus, having MetS was associated with the equivalent of 2.5 years of cognitive aging in this cohort.

When we used categorized MMSE scores as an outcome, compared to those without MetS, those with MetS

Table 2. MetS, individual components, diabetes and MMSE score

	Δ MMSE		
	model 1	model 2	model 3
MetS	-0.67 (-0.92, -0.41)	-0.28 (-0.51, -0.05)	-0.24 (-0.47, -0.01)
Adjusted for MetS individual components (abdominal obesity, blood glucose, blood pressure, triglycerides and HDL)			
Abdominal obesity	-0.20 (-0.47, 0.07)	-0.04 (-0.29, 0.21)	-0.01 (-0.26, 0.24)
Blood pressure \geq 130/85 mm Hg	-0.97 (-1.34, -0.61)	-0.07 (-0.40, 0.26)	-0.06 (-0.38, 0.27)
Triglycerides \geq 150 mg/dl	0.02 (-0.28, 0.33)	0.03 (-0.24, 0.30)	0.04 (-0.23, 0.30)
Low HDL	-0.38 (-0.66, -0.10)	-0.19 (-0.44, 0.06)	-0.16 (-0.42, 0.09)
Blood glucose \geq 100 mg/dl	-0.21 (-0.49, 0.06)	-0.11 (-0.36, 0.13)	-0.07 (-0.32, 0.17)
Adjusted for waist, systolic blood pressure, diastolic blood pressure, triglycerides, HDL and blood glucose			
Waist per inch	0.008 (-0.016, 0.033)	0.006 (-0.019, 0.030)	0.023 (-0.005, 0.050)
Systolic blood pressure per mm Hg	0.004 (-0.003, 0.011)	0.004 (-0.003, 0.012)	-0.017 (-0.025, -0.010)
Diastolic blood pressure per mm Hg	-0.011 (-0.024, 0.003)	-0.011 (-0.025, 0.002)	0.004 (-0.011, 0.018)
Triglycerides per mg/dl	0.001 (-0.001, 0.002)	0.001 (-0.001, 0.002)	0.001 (-0.001, 0.003)
HDL per mg/dl	0.010 (0.001, 0.020)	0.012 (0.002, 0.021)	0.016 (0.006, 0.026)
Blood glucose per mg/dl	-0.001 (-0.004, 0.001)	-0.001 (-0.004, 0.001)	-0.004 (-0.006, -0.001)
Adjusted for MetS individual components (abdominal obesity, blood pressure, triglycerides and HDL) except blood glucose			
Diabetes	-0.42 (-0.74, -0.10)	-0.17 (-0.45, 0.11)	-0.12 (-0.40, 0.17)

Δ MMSE = Difference in MMSE score; 95% CI given in parentheses.

Model 1: Unadjusted; model 2: adjusted for sociodemographics (age, sex, race/ethnicity, insurance and education); model 3: adjusted for sociodemographics, alcohol consumption, smoking, social support and physical activity.

Table 3. Mets and categories of MMSE scores

	OR (95% CI)		
	model 1	model 2	model 3
MMSE score <18 vs. \geq 24	2.25 (1.53-3.33)	1.93 (1.26-2.96)	1.94 (1.26-3.01)
MMSE score <18 vs. 18-23	1.71 (1.12-2.59)	1.73 (1.11-2.70)	1.77 (1.13-2.78)
MMSE score 18-23 vs. \geq 24	1.32 (1.10-1.59)	1.11 (0.90-1.37)	1.10 (0.89-1.35)

Model 1: Unadjusted; model 2: adjusted for sociodemographics (age, sex, race/ethnicity, insurance and education); model 3: adjusted for sociodemographics, alcohol consumption, smoking, social support and physical activity.

were more likely to have an MMSE score of <18 than a score of \geq 24 (adjusted OR = 1.94; 95% CI 1.26, 3.01). In unadjusted models, those with MetS also were more likely to have an MMSE score of 18-23 than a score of \geq 24, but this association was no longer significant after adjusting for potential confounders (adjusted OR = 1.10; 95% CI 0.89, 1.35; table 3).

MetS and MMSE Score by Race-Ethnicity

There was evidence of an interaction between MetS and race-ethnicity for MMSE scores such that the MMSE score was lower for those with MetS among non-Hispan-

ic whites but not non-Hispanic blacks (p for interaction = 0.01). In stratified models, MetS was associated with lower MMSE score among Hispanics (adjusted β = -0.41; 95% CI -0.72, -0.10) and non-Hispanic whites (adjusted β = -0.51; 95% CI -1.01, -0.01), but not among non-Hispanic blacks (adjusted β = 0.38; 95% CI -0.09, 0.84; table 4).

We did not find interactions by age, sex, education, or medical insurance status.

MetS Components, Diabetes, and MMSE Score

When evaluating the effects of each of the individual components of MetS, only two of the five components

Table 4. MetS and MMSE score stratified by race-ethnicity

Race-ethnicity	Change in MMSE score (95% CI)*
Hispanics	-0.41 (-0.72, -0.10)
Non-Hispanic whites	-0.51 (-1.01, -0.01)
Non-Hispanic blacks	0.38 (-0.09, 0.84)

* Adjusted for age, sex, education, insurance status, moderate alcohol consumption, smoking, social support and physical activity.

were associated with MMSE score in univariate analyses: high blood pressure ($\beta = -0.97$; 95% CI $-1.34, -0.61$) and low HDL ($\beta = -0.38$; 95% CI $-0.66, -0.10$). After adjusting for potential confounders, however, none of the individual components were associated with MMSE score (table 2). When analyzed as a continuous measure, however, HDL remained associated with MMSE score after fully adjusting for potential confounders (adjusted $\beta = 0.01$ per mg/dl HDL; $p = 0.03$), though other MetS components were not.

Diabetes was inversely associated with MMSE score, and this association remained after adjusting for individual components of MetS. After further adjusting for other potential confounders, however, the relationship was no longer significant (adjusted $\beta = -0.12$; 95% CI $-0.40, 0.17$; table 2).

Inflammation

Inflammatory biomarkers were available in a subsample of our cohort (IL-6: $n = 1,664$, hsCRP: $n = 2,219$, SAA: $n = 2,145$ and TNFR1: $n = 1,847$). TNFR1 was independently associated with MMSE score (adjusted $\beta = -0.14$ per 1 ng/ml; 95% CI $-0.23, -0.05$); however, other inflammatory markers were not associated with MMSE score. MetS was associated with MMSE score even after adjusting for inflammatory markers and there were no interactions between inflammatory markers and MetS on MMSE score.

Discussion

In this multi-ethnic, urban, stroke-free, community-based cohort, we found an inverse association between MetS and MMSE score, independent of other confounding factors. This association was limited to Hispanics and non-Hispanic whites, but did not vary by sex, age, education, medical insurance or inflammatory biomarkers.

These findings have important public health implications. In the past two decades in the United States, the age-adjusted prevalence of MetS among adults increased from 23.7% (1988–1994) [18] to 34.4% (2003–2006) [19]. Based on the Census Projection of the US Population [20], in 2010, 77 million individuals above 20 years old have MetS, reaching 85 million by the year 2020. The difference in cognitive performance attributable to MetS might be small on an individual basis; however, due to the high MetS prevalence, the burden of impaired cognition due to the MetS may be substantial in the overall population.

Our findings showed similar results and magnitude confirming previous studies of the association between MetS and lower cognitive performance [21–24], and extend these findings to a predominantly Hispanic population in which the MetS prevalence is high. Hispanics have been traditionally understudied, although they are the fastest growing population in the US, they have the highest MetS prevalence, and together with non-Hispanic blacks, may be at a higher risk of dementia than non-Hispanic whites [25]. Studies of the relationship between vascular risk factors and cognition among Hispanics are therefore needed.

The negative effect of MetS on cognition has however not been observed in all populations. A European study [26] demonstrated better cognitive performance among women greater than 80 years of age with MetS. The LEIDEN 85-PLUS study [27] failed to demonstrate a difference in cognitive performance between individuals with and without MetS, and found slower cognitive decline among individuals over 85 years of age with MetS. The WHICAP study, which studied a similar demographic population, but older than our cohort (median 76 years of age), failed to demonstrate an association between MetS and incident dementia. However, it showed an association between diabetes and hyperinsulinemia, and increased incidence of Alzheimer disease and vascular dementia [28]. These results may be explained by survival bias, however, it is possible that people who live to old age with MetS either have a milder case or are not as affected by the syndrome.

The main hypothesis by which MetS is thought to affect cognition is through insulin resistance that can lead to cerebral small vessel disease. Both diabetes and hyperinsulinemia have been associated with increased risk of developing Alzheimer disease, vascular dementia [29] and cognitive decline. In addition, MetS is also associated with leukoaraiosis, a risk factor for worse cognitive performance [30] and cognitive decline [31].

There could also be direct independent effects of each of the MetS components themselves. Hypertension [4] and hyperlipidemia [32] are reported to increase the risk

of dementia and/or cognitive decline. Advanced age has been associated with a greater prevalence of MetS and its components, which could further lead to an increase in cognitive dysfunction [26]. Similarly, diabetes has been associated with lower cognition [33], although we did not find an independent effect of diabetes after adjusting for other risk factors. Our results, moreover, support the utility of the concept of MetS as a syndrome, and suggest that considering the risk factors together may be more informative with regard to cognitive performance than each individual component. We did, however, find that HDL independently contributed to lower score on the MMSE, even after adjusting for other risk factors, though other risk factors did not. The relationship of lipids to cognition deserves further study. We did not consider the use of medications for lipids or blood pressure control because they were not part of the ATPIII definition of MetS.

Our models provide some support for the role of elevated inflammatory biomarkers in predicting poor cognitive performance. Elevated TNFR1 levels were associated with lower MMSE score in our cohort, but we did not confirm that the relationship between MetS and MMSE score was mediated through inflammation, as suggested by other studies [34]. Elevation of TNFR1 has also been associated with decreased memory performance [35] and it may be a reflection of the inflammatory mechanisms operative in atherosclerotic disease as it has been shown to be associated, in our cohort and others, with increased carotid plaque [36], left ventricular hypertrophy [16] and vascular dementia [37].

The reason for the apparent lack of effect of MetS among non-Hispanic blacks is uncertain. It is notable in this regard, however, that investigators in the Health ABC study [38] also failed to demonstrate decreased cognitive performance among African Americans with MetS, though they did demonstrate greater cognitive decline over time. It is possible that the absence of an effect of MetS on MMSE score among non-Hispanic blacks relates to a lower baseline MMSE score among non-Hispanic blacks. The MMSE has a substantial ceiling effect, however, such that there is more discrimination among individuals at lower scores than higher scores, making this explanation less likely. It is also possible, however, that our study and the other available studies did not have enough power to show an effect among non-Hispanic blacks alone. Alternatively, the results are the result of chance. Future studies on the race-ethnic variability of MetS in relation to cognition are warranted.

This study's strengths are the multi-ethnic cohort, including Caribbean Hispanics, a largely understudied

population with a high burden of MetS; a large sample size; and the ability to control for multiple potential confounders. We addressed the MMSE in a continuous and categorical way, as well as the effects of diabetes, individual components of the MetS, and inflammation.

Our study also has limitations. First, the cross-sectional design does not allow us to make causal inferences. The MMSE has been demonstrated to have significant performance variability due to age, education, social status, ethnicity and language [17, 39]. The MMSE thresholds utilized may have lower sensitivity and specificity among individuals with lower levels of education [40]. These cutoff scores are not very well documented in Hispanic elderly, however, because few studies have clinically validated the suspicion of cognitive impairment in Hispanics [17]. The majority of Hispanic participants in our study are of Caribbean origin; therefore our results may or may not be generalized to other Hispanic populations. Apolipoprotein genotype E4 was not routinely available in study participants, and therefore was not included in our models.

Conclusion

In summary, this study provides evidence of an inverse association between MetS and cognitive performance. This effect was found among Hispanics and non-Hispanic whites, but not among non-Hispanic blacks. MetS as a cluster of risk factors appears to better translate the association with poor cognitive performance than its individual components or diabetes alone.

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