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# DIVERGENT INFLUENCES OF CARDIOVASCULAR DISEASE RISK FACTOR DOMAINS ON COGNITION, GREY AND WHITE MATTER MORPHOLOGY

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**Objective:** Hypertension, diabetes, dyslipidemia, and obesity are associated with preclinical alterations in cognition and brain structure; however, this often comes from studies of comprehensive risk scores or single isolated factors. We examined associations of empirically-derived cardiovascular disease risk factor domains with cognition and brain structure.

**Methods:** 124 adults (age~59.8±13.1;41% African American;50% women) underwent neuropsychological and cardiovascular assessments and structural MRI. Principal component analysis of 9 cardiovascular disease risk factors resulted in a four-component solution representing 1-cholesterol, 2-glucose dysregulation, 3-metabolic dysregulation, and 4-blood pressure. Separate linear regression models for learning, memory, executive functioning and attention/information processing were performed with all components entered at once, adjusting for age, sex, and education. MRI analyses included whole brain cortical thickness and tractbased fractional anisotropy adjusted for age and sex.

**Results:** Higher blood pressure associated with poorer learning (B=-0.19;p=0.019), memory (B=-0.22;p=0.005), and executive functioning performance (B=-0.14;p=0.031), and lower cortical thickness within the right lateral occipital lobe. Elevated glucose dysregulation associated with poorer attention/information processing performance (B=-0.21;p=0.006) and lower fractional anisotropy in the right inferior and bilateral superior longitudinal fasciculi. Cholesterol was associated with higher cortical thickness within left caudal middle frontal cortex. Metabolic dysfunction was positively associated with right superior parietal lobe, left inferior parietal lobe and left precuneus cortical thickness.

**Conclusions:** Cardiovascular domains associated with distinct cognitive, grey, and white matter alterations and distinct age groups. Future longitudinal studies may assist in identifying vulnerability profiles that may be most important for individuals with multiple cardiovascular disease risk factors.

**Key Words:** cardiovascular disease risk factors, blood pressure, glucose, metabolic, cortical thickness, diffusion tensor imaging; white matter integrity

Acronyms: CVD-RF = cardiovascular disease risk factors; BMI = body mass index; DTI = diffusion tensor imaging; FA = fractional anisotropy; HC = healthy controls; SBP = systolic blood pressure; DBP = diastolic blood pressure; Chol = cholesterol; GluDys = glucose dysregulation; MetabDys = metabolic dysregulation; BP = blood pressure; HbA1c = hemoglobin A1c; PCA = principal component analysis; LRN = learning; MEM = memory; EF = executive function; AIP = attention and information processing; CVLT-II = California Verbal Learning Test-II; WMS-III = Wechsler Memory Scales-III; TMT = Trail Making Test; WAIS-III = Wechsler Adult Intelligence Scale-III; GLM = general linear model; FWHM = full width half maximum; QDEC = Query, Design, Estimate, Contrast; TFCE = threshold free cluster enhancement; ILF = inferior longitudinal fasciculus; SLF = superior longitudinal fasciculi

# Introduction

Cardiovascular disease risk factors (CVD-RFs) including hypertension, dyslipidemia, diabetes, and obesity increase the likelihood of stroke, heart failure, and mortality (1). CVD-RFs also have a deleterious impact on cognition and brain aging. In fact, all are independent risk factors for cognitive decline and dementia (2,3). Given their frequent comorbidity, CVD-RFs representing distinct pathophysiological processes including hypertension-related reductions in vasodilatory capacity of cerebral arterioles (4) or increases in CNS inflammatory responses secondary to diabetes (5) are often clustered together into a single risk score when investigating brain/behavior relationships (6). These scores have contributed significantly to our understanding that, even in mid-life, higher overall CVD-RF burden is associated with increased risk for dementia in later life (3). While risk scores are advantageous for examining cumulative burden, they present limitations for understanding contributions of distinct CVD-RF domains (7). Advancing our understanding of not only the comorbid impact but also the distinct contributions CVD-RF domains have on brain structure and cognition may assist in identifying 'vulnerability profiles' that may be monitored in affected individuals.

CVD-RF domains may have diverse and potentially antagonistic impact on gray matter morphology. Elevations in systolic and diastolic blood pressure have been associated with diffuse cortical thinning of the frontal, temporal, and occipital lobes, whereas higher cholesterol levels including total and LDL cholesterol and triglycerides have been associated with increased cortical thickness within frontal, temporal, and parietal lobes (8–11). In contrast, glucose dysregulation (as measured by such metrics as fasting glucose, hemoglobin A1c and diabetes diagnosis) and body mass index (BMI) have been weakly and inconsistently related to cortical thickness with results primarily involving frontal, anterior cingulate, temporal, and occipital regions (9–12). These investigations often looked at particular CVD-RFs and/or diagnoses in isolation (8,11,12), very few took a CVD-RF domain approach to investigate blood pressure, short and longer-term glucose control, 'good' and 'bad' cholesterol, etc. within one study (9,10). Furthermore, given that CVD-RFs predict cognitive decline (2,3) it is unclear how their divergent and often mixed relationship (8–12) with cortical morphology relates to cognition.

CVD-RF domains also have heterogeneous associations with white matter integrity as measured by diffusion tensor imaging (DTI). The most widely used metric of white matter integrity (13) is DTI-derived fractional anisotropy (FA). FA alterations have been extensively linked with CVD-RFs. Individuals with hypertension and diabetes have been shown to have lower FA than controls, with hypertension preferentially affecting posterior brain regions and diabetes impacting more anterior brain regions (14,15). BMI has been positively and negatively associated with regional FA (16,17) as have serum cholesterol levels (18,19). While lower FA has long been associated with cognitive decline in healthy aging (20); fewer studies have explored the role of these CVD-RF-associated FA alterations on cognition. Lower FA has been associated with poorer memory, executive function, and processing speed in both diabetes and untreated hypertension (21,22); further research incorporating more CVD-RF domains is warranted.

Our study aim was to examine the association of multiple CVD-RF domains with cognition and brain structure. We examined the associations of CVD-RF profiles with cognition, grey and white matter in a sample of ethnically diverse, community-dwelling adults. Exploring the relationship of CVD-RFs to cognition in conjunction with grey and white matter integrity may provide insight (23) into the associations of brain/behavior and CVD-RFs (20). We first examined CVD-RFs and cognition, hypothesizing that higher (systolic and diastolic) blood pressure and glucose dysregulation measured using glucose and hemoglobin A1c would be associated with poorer memory, executive function, and attention/information processing. We

then examined associations of CVD-RFs on regional grey and white matter, hypothesizing that higher blood pressure would be associated with lower cortical thickness, whereas higher cholesterol measures would be associated with greater cortical thickness. Furthermore, we hypothesized that elevated blood pressure would be associated with diminished FA in posterior brain regions, whereas increased glucose dysregulation would be associated with lower FA in anterior regions. Finally, we assessed whether CVD-RF related grey and white matter alterations correlated with CVD-RF associated cognitive profiles.

# Methods

#### **Participants**

Healthy controls (HCs) were identified from a larger program of research investigating aging, diabetes and depression approved by the University of Illinois at Chicago Institutional Review board and conducted in accordance with the Declaration of Helsinki. HCs provided written informed consent at enrollment. As previously described (6), HCs, aged 30 years and older (range=30-89), were recruited from the community between 10/2009-08/2012 and underwent an initial telephone screening to rule-out neurological and/or Axis 1 disorders, head injury or loss of consciousness, substance abuse/dependence, and/or psychotropic medication use. Upon successful completion of the telephone screen, study staff including board eligible/certified psychiatrists conducted a more detailed in person screening. HC status was confirmed if: 1) Mini-Mental State Examination (24) scores were  $\geq$ 24, 2) there was an absence of current and/or past psychiatric symptoms as assessed by the Structured Clinical Interview for DSM-IV and 3) Hamilton Depression Rating Scale scores were <8.

Of 124 eligible participants (Table 1), 43 (34.7%) met criteria for diabetes (non-fasting glucose $\geq$ 200 mg/dl or anti-diabetic mediation use) (25), a rate higher than the Illinois state average (23.8%) (26) and likely attributable to active recruitment of individuals with diabetes in

the larger research. Seventy participants were hypertensive (systolic blood pressure [SBP] $\geq$ 140 mmHg, diastolic blood pressure [DBP] $\geq$ 90 mmHg, or antihypertensive medication use) (27), 54 met criteria for obesity (BMI $\geq$ 30 kg/m<sup>2</sup>) (28), 40 had hypercholesterolemia (total cholesterol $\geq$ 240 mg/dl or lipid lowering medication use) (29), and 27 did not met criteria for any of these conditions. Forty-two participants were on anti-diabetic medications, 50 were on antihypertensives, and 36 were on lipid lowering medications.

#### **CVD-RF** Assessment

One reading of brachial SBP and DBP were obtained with the participant sitting upright using an automated device (Welch Allyn, model 300, Welch Allyn Inc., Arden, NC) and appropriate cuff size. Weight in kilograms and height in centimeters were collected to calculate BMI. A non-fasting blood sample was obtained by venipuncture to determine plasma concentrations of glucose, total cholesterol, HDL-cholesterol, LDL-cholesterol, triglycerides, and hemoglobin A1c (HbA1c). Cholesterol and HbA1c values have been documented to be minimally affected by normal food consumption (30).

In order to allow for the consideration of CVD-RF domains without treating any one CVD-RF in isolation at the expense of another, we took our lead from the literature on studies of biological determinants of cerebrovascular health (8,9). As suggested by the literature (8,9), CVD-RF metrics of interest (SBP, DBP, BMI, glucose, total cholesterol, inverted HDLcholesterol, LDL-cholesterol, triglycerides, and HbA1c) were submitted to a PCA with varimax rotation to empirically determine the CVD-RF metrics that had high shared variance (e.g., SBP and DBP; glucose and HbA1c) and to control for multiple comparisons.

Similar to previous research using this technique (8,9), our PCA resulted in four orthogonal factors that explained 78.0% variance (minimum eigenvalue=1; Table 2) representing Cholesterol (Chol; Component #1), Glucose Dysregulation (GluDys; Component #2), Metabolic

Dysregulation (MetabDys; Component #3), and Blood Pressure (BP; Component #4). While consistent with prior research (9), our MetabDsy loadings were more robust, likely due to the omission of previously documented (9) low-loading creatine from our analyses. PCA marker variables guided the creation of domain scores to represent CVD-RFs with the highest shared variance per Component (e.g., Component 4: SBP and DBP; Component 2: glucose and HbA1c) (31). Raw scores for the highest loadings per Component (Table 2) were transformed into standardized z-scores based on the study sample's mean and standard deviation and averaged to create BP, GluDys, MetabDys, and Chol domain scores.

#### Neuropsychological Assessment

Conducted by a trained research assistant and described in detail elsewhere (6), cognitive domains were calculated by averaging z-scores within each construct tested: (1) Learning (LRN): immediate recall from the California Verbal Learning Test-II (CVLT-II) Trials 1-5 (32) and the Wechsler Memory Scales-III (WMS-III) Logical Memory-I and Visual Reproduction-I (33); (2) Memory (MEM): CVLT-II long delay free recall (32), WMS-III Logical Memory-II and Visual Reproduction-II (33); (3) Executive Function (EF): Trail Making Test (TMT) Part B time to completion (34) (reversed), Delis-Kaplan Executive Function System category switching total (35), Stroop Interference score (36), Wechsler Adult Intelligence Scale-III (WAIS-III) Digit Span Backwards raw score (37), Self-Ordered Pointing Task total errors (38) (reversed); (4) Attention and Information Processing (AIP): TMT Part A time to completion (34) (reversed), Stroop Color and Word raw scores (36), WAIS-III Digit-Symbol Coding raw score (37). Cronbach's alphas revealed adequate internal consistency (LRN=0.75, MEM=0.75, EF=0.73, AIP=0.84).

#### MRI Acquisition

Whole brain MRI data in the axial plane were collected on a Philips Achieva 3T scanner (Philips Medical Systems, the Netherlands) using an 8-channel sensitivity encoding head coil. Anatomical image data were collected using MPRAGE (FOV=240mm, contiguous slices=134, TR/TE=8.4/3.9ms, flip angle=8°, voxel size= $1.1 \times 1.1 \times 1.1$ mm<sup>3</sup>). DTI was obtained aligned to the anterior commissure-posterior commissure line, using a single-shot spin-echo echo-planar imaging sequence (FOV=240mm, contiguous slices=67, TR/TE=6,994/71ms, flip angle=90°, voxel size= $0.83 \times 0.83 \times 2.2$ mm<sup>3</sup>, 32 isotropically distributed diffusion-weighted directions with b=700s/mm<sup>2</sup>, and one B<sub>0</sub> images with no diffusion sensitization). Scanning time was reduced to ~4 minutes using a SENSE parallel imaging technique with a factor of 2.5.

Of the 124 subjects with available cognitive and cardiovascular data, 86 had T<sub>1</sub> images. Reasons for unavailable T<sub>1</sub> data included: claustrophobia=11; BMI=4; metallic implants=3; scanner problems=19; poor image quality=1. Twenty-four did not complete DTI due to time constraints and another 9 were removed for poor quality images (DTI n=53). Individuals with available MRI data were older than those without MRI data ( $61.7\pm12.6$  versus  $55.6\pm13.2$ ; p=.017); groups did not differ in education or sex.

#### Image Processing

All MRI data was visually examined for quality. This inspection included an examination for artifacts from movement or other causes, such as space-occupying or other focal lesions, including stroke and other gross neuroanatomical abnormalities. As outlined above, 10 participants were excluded based on this inspection.

Cortical thickness quantification was conducted in FreeSurfer Version 5.3.0 (<u>http://surfer.nmr.mgh.harvard.edu/</u>). Following motion correction and skull stripping, thickness measurements were derived at each vertex along the reconstructed surface by computing the shortest distance between the outer cortical (pial) surface and grey/white matter border (39,40). These measurements were mapped onto each participant's inflated cortical surface and checked for quality control by viewing individual images in tkmedit and tksurfer. A spherical averaging procedure was implemented to average thickness measurements across participants.

DTI analysis, implemented in FSL v4.1.9 (http://www.fmrib.ox.ac.uk/fsl), used FMRIB's Diffusion Toolbox and Tract-Based Spatial Statistics (41). FA images underwent motion correction and skull stripping, normalization to the 1mm<sup>3</sup> MNI152 stereotaxic space by implementing a nonlinear registration of each participant's FA image to the FSL FA template and then applying an affine transformation to the MN1152 template – images were then visually inspected for errors in processing and/or distortion. Normalized FA images were averaged to create a mean FA map. A mean FA skeleton was derived by applying a 0.2 threshold (41) and images were then visually inspected to ensure that the major tracts were well aligned to the skeleton. Each participant's FA data was projected onto the group skeleton to identify the highest local FA value and then applied to the skeleton.

#### **Statistical Analysis**

Associations between CVD-RF and cognition domains were assessed using IBM SPSS Statistics 22 (IBM Corp, Armonk, NY). Separate linear regression models ( $p \le 0.05$ ) assessed each cognitive domain score individually (4 models) with all CVD-RF domain scores entered at once, adjusting for age, sex, and education. Additional adjustments for race, smoking status (current smoker: yes/no) and treatment effects (presence/absence of medication use for hypertension, diabetes and/or hypercholesterolemia) were also considered. Given the age range of our sample (30-89 years), we explored age as a moderator variable by adding an interaction term between significant CVD-RF domains and age as appropriate. The relationship between individual CVD-RF domains and cortical thickness was evaluated using a general linear model (GLM) at each vertex in the cortical mantle adjusting for age and sex using the Query, Design, Estimate, Contrast (QDEC) interface in FreeSurfer (www.surfer.nmr.mgh.harvard.edu; 42). Maps were smoothed with a 15mm full width half maximum (FWHM) Gaussian kernel, a more conservative threshold than prior relevant studies (9,10). Correction for multiple comparisons was performed with a cluster-wise procedure using Monte Carlo Null-z simulation method within the QDEC processing stream (43). For each model, 10,000 iterations of the stimulation were performed using a threshold of 1.3 (p<0.05). Mean cortical thickness in identified clusters was extracted within QDEC. In supplementary regression analyses, associations between individual regions of interest (ROIs) derived from the FreeSurfer atlas and CVD-RF domains as well as individual CVD-RF variables were assessed adjusting for age and sex.

Associations between CVD-RF domains and FA, adjusting for age and sex, were assessed using regression models adjusted for age and sex. Five thousand permutations were performed and cluster-based thresholding correction was implemented with threshold free cluster enhancement (TFCE) (44). Statistical maps were then thresholded at p<0.001 and clusters sizes larger than 100 voxels were identified. The anatomical location of significant clusters was identified using JHU ICBM-DTI-81 White Matter Labels and JHU White Matter Tractography (45) and mean FA values were extracted using FSL.

Due to the truncated MRI sample, we examined the association between CVD-RF related morphological alterations, cortical thickness and FA clusters significantly associated with CVD-RFs, defined and extracted as described above, with their respective CVD-RF cognitive associates using two-sided Pearson's correlations ( $p \le 0.05$ ) in SPSS. We also ran more stringent mediation (46) and indirect mediation analyses with bootstrapping using Preacher and Hayes methods (47,48).

#### Results

*Cognition*– In separate cognitive models with all cardiovascular domains (BP, GluDys, Chol and MetabDys) entered simultaneously adjusting for age, sex and education, only BP domain scores were significantly and negatively associated with LRN (B=-0.19,SE=0.08;p=0.019) and MEM (B=-0.22,SE=0.07;p=0.005) performance. In contrast, BP as well as GluDys domain scores negatively associated with EF (BP: B=-0.14,SE=0.06;p=0.031; GluDys: B=-

0.11,SE=0.06;p=0.058); however, only the BP domain score reached significance (Table 3). Only GluDys domain scores significantly and negatively associated with poorer AIP (B=-0.21,SE=0.07;p=0.006). No associations were observed for Chol or MetabDys. Results remained unchanged with further adjustments for race, smoking status and treatment effects except the BP and EF association was no longer significant.

Exploring age as a moderator variable resulted in a significant age X BP interaction for LRN (p=0.027) and MEM (p=0.014) only; there were no significant age x GluDys interactions (all p-values $\geq$ 0.65). After adjusting for all covariates including race, smoking status and treatment effects, only the age X BP interaction for MEM remained significant (p=0.048). A follow-up Johnson-Neyman plot revealed a cut-point of significance occurring for the conditional effect of BP on MEM such that individuals <64.4 years old showed a significant association between BP and MEM (all p-values $\leq$ 0.050) while individuals  $\geq$ 64.4 years of age did not (all p-values $\geq$ 0.072).

*Cortical Thickness*– Higher BP domain scores were associated with lower cortical thickness in the right lateral occipital lobe controlling for age and sex (Figure 1a). No significant associations between cortical thickness and GluDys were observed (Table 4). Chol domain scores positively

associated with cortical thickness within the left caudal medial frontal cortex after adjustment for age and sex (Figure 1b). Higher MetabDys domain scores were associated with higher cortical thickness measures in the right superior parietal lobe, left inferior parietal lobe, and left precuneus after adjustments for age and sex (Figure 1b). Additional adjustments for race, smoking status and treatment effects did not alter these results. For completeness, FreeSurfer cortical thickness ROIs as analyzed individually is represented in Supplemental Table I with regressions between individual CVD-RFs that comprised each domain score and results of these individual analyses shown in Supplemental Table II.

An initially significant age X BP interaction for the right lateral occipital lobe adjusting for age and sex (p<0.001) did not remain significant after adding race, smoking status and treatment effects to the model (p=0.791). Exploring age as a moderator variable did not reveal any significant age X Chol or age X MetabDys interactions regardless of adjustments (all pvalues $\geq$ 0.110).

*DTI tractography*–After adjustment for age and sex, higher GluDys domain scores associated with lower FA in the right inferior longitudinal fasciculus (ILF) and bilateral superior longitudinal fasciculi (SLF; Figure 2). No other CVD-RF domain scores associated with FA in models adjusted for age and sex. Additional adjustments for race, smoking status and treatment effects did not alter our results.

Exploring age as a moderator variable revealed significant age X GluDys interactions for all identified white matter tracts listed when only age and sex were in the model (all pvalues≥0.044). Only the age X GluDys interaction in the right SLF model remained significant after adding race, smoking status and treatment effects to the model (p=0.001). Johnson-Neyman plots revealed a cut-point of significance occurring for the conditional effect of GluDys on right SLF FA with individuals <56.4 years of age showing a significant association between GluDys and FA in the right SLF (all p-values $\leq 0.050$ ) and individuals  $\geq 56.4$  years of age not showing this association.

*Neuroimaging and Cognition*–We extracted the cortical thickness and FA cluster measures associated with CVD-RF domain scores to determine whether these clusters significantly correlated with their relevant CVD-RF cognitive associates. For example, LRN, MEM and EF correlates to BP were subjected to two-tailed Pearson's correlations with the lateral occipital cortical thickness cluster (also associated with BP); no cognitive domains correlated with this thickness measure. When considering the white matter tracts significantly correlated with GluDys, EF (a trending correlation with GluDys) was positively associated with right ILF FA values [r(df=51)=0.30,p=0.027] and bilateral SLF values [left: r(df=51)=0.32,p=0.020; right: r(df=51)=0.33,p=0.014] while AIP scores significantly related to GluDys positively correlated with right SLF FA values [r(df=51)=0.27,p=0.048]. Indirect mediation analyses with bootstrapping did not corroborate these results, i.e., no mediation or indirect pathway results were significant.

#### Discussion

The overarching message from this research is that CVD-RF domains had more divergent than convergent influences on cognition and brain structure. For example, BP and GluDys domains were more strongly associated with cognition than Chol and MetabDys domains. Additionally, BP, Chol and MetabDys were associated with cortical thickness either negatively (BP) or positively (Chol and MetabDys), while only GluDys related to (lower) FA. Similarly, only GluDys-related FA alterations associated with poorer cognitive functioning suggesting this was the most consistent of our brain/behavior relationship findings. While these results may highlight the unique information derived by the examination of empirically derived CVD-RF domains, when combined with our results using age as a moderator variable, this study highlights the importance of considering how age may influence brain/behavior associates of CVD-RFs – an area of increased interest across multiple CVD-RFs (49–51).

Only BP and GluDys emerged as salient associates of cognition. Specifically, higher BP associated with poorer LRN, MEM, and EF while greater GluDys associated with poorer AIP and poorer EF (albeit a non-significant trend). These findings are consistent with a systematic review of the literature that showed hypertension and diabetes display a consistent detrimental impact on cognition whereas obesity and dyslipidemia have weak and variable effects (52). This may be due to the age of participants across studies, i.e., mid- versus late-life (52), and/or the grouping of obesity and dyslipidemia – variables increasingly seen to be distinct (8). Our results also indicate that continuous blood pressure and glucose measurements are negatively associated with cognition regardless of a diagnosis of hypertension and/or diabetes. Given that continuous measures incorporate pre-clinical levels of disease known to be detrimental to brain health (17,53), our results may be early support for such recent recommendations as lowing the 'optimal' SBP to <120mmHg (54). While further research is needed to fully determine the inflection points beyond which cognition is negatively impacted by BP or glucose, this study and others (55) suggest focusing on particular age ranges, e.g., individuals less than age 65 when considering BP, may facilitate this work.

Cardiovascular domains had distinct and competing influences on cortical thickness. Higher BP related to lower cortical thickness in posterior regions, while higher Chol was related to greater cortical thickness in more anterior regions. Higher MetabDys also associated with greater cortical thickness, however, in more posterior, e.g., parietal regions. While cholesterol and metabolic-related results regarding cortical thickness may not be intuitive, other studies report similar findings (8,9,56). In fact, these studies advocate that BMI and lipids may not be consistently related to each other but instead, may be independent CVD-RFs (8), they may also require a non-linear approach to understanding their brain-behavior relationships. Additionally, grey matter measures vary in the literature, introducing differences between cortical thickness and grey matter volume that may influence CVD-RF associations, e.g., cortical thickness is more vulnerable to environmental exposures while both measures have distinct genetic influences. Genetic factors link cholesterol levels in the peripheral and central nervous systems; this may differentially impact associations made between genetic regulation of circulating cholesterol and alterations in grey as well as white matter morphology (18) further influencing our results.

Elevations in BP were negatively associated with cortical thickness in the lateral occipital cortex, in line with studies demonstrating that hypertension induces cortical degradation in posterior regions including the parietal and occipital lobe, areas typically preserved in healthy aging (15). This may be due, in part, to the fact that BP elevations can diminish blood brain barrier integrity, which disrupts nutrient delivery and propagates neuronal damage, potentially resulting in cortical thinning (53,57). Furthermore, chronic elevations in BP can reduce the vasodilatory capacity of the cerebral arterioles by increasing media thickness and narrowing the lumen (58). Subsequent reductions in cerebral blood flow may result in hypoperfusion and induce neuronal damage and grey matter atrophy (42).

GluDys was the only factor associated with TBSS-derived FA values and FA subsequently correlated with the cognitive domains shown to be related to GluDys; the only white matter findings related to CVD-RFs and their relevant CVD-RF cognitive associates. Some GluDys-related white matter associates were driven by younger participants, thus, the wide 30-89 age range of our sample may have precluded replication of previously reported BP and FA associations (15). In contrast, and consistent with other studies (14,17), higher GluDys was associated with lower FA in the right ILF and SLF, bilaterally – key white matter tracts connecting posterior to more anterior regions of brain to facilitate higher-level cognitive functions. Elevations in glucose dysregulation may impact FA by propagating pro-inflammatory cytokines (59) capable of crossing the blood brain barrier (60), and instigating a local CNS inflammatory response characterized by activation of microglial cells (61). Activated microglia, in turn, may impede the development and migration of oligodendrocyte progenitor cells (62), disrupting myelin production and diminishing white matter integrity as well as the successful relay of information to perform executive functioning and attention/information processing tasks. Future work focusing on the role of CVD-RFs, particularly glucose, on myelin degradation may assist in determining the underlying etiology of our findings (63).

While this study examined a broad range of cardiovascular, cognitive, and brain morphology factors, limitations exist. Consistent with prior literature (8,9), we utilized PCA to create CVD-RF domains consisting of factors with high shared variance. While this approach is efficacious for creating empirically defined groups of highly similar variables, reducing redundancy, and diminishing multiple comparisons, BMI loaded onto both the GluDys and MetabDys domains – albeit with lower values than any other variable loading in the PCA. This may suggest that BMI is either a multidimensional variable and/or one that did not exert as strong a contribution as other CVD-RFs in our PCA. Future studies should incorporate more direct assessments of adiposity such as those provided by dual-energy x-ray absorptiometry scanning. Our cross-sectional design precludes determination of causation, making longitudinal studies necessary. Additionally, larger sample sizes will be necessary to empirically examine the interplay between cortical thickness and white matter alterations (23). The neuroimaging and cognition mediation analyses and assessment of indirect effects may have suffered from a truncated neuroimaging sample making our call for larger sample sizes even more germane. These analyses may also have suffered from a limited number of cognitive domains, i.e., a

visuospatial/visuoperceptual component may have related to BP and its lateral occipital associates.

DTI-derived FA reflects diverse morphological changes such as alterations in myelination, axonal density, inflammation, and edema (13). While FA may be divided into axial and radial diffusivity, many argue that these measures are not only influenced by inflammation and edema, they are highly correlated and may not reflect axonal loss (i.e., axial diffusivity) and myelin degradation (i.e., radial diffusivity) in isolation (13,63,64). Neuroimaging techniques including multi-component relaxometry (64), which eliminate the influence of inflammation and edema, may provide greater insight into underlying etiological factors driving white matter associates to CVD-RFs including white matter perfusion and demyelination (63). Future studies should compare the various microstructural metrics for axonal loss and myelin degradation both to each other as well as to CVD-RF profiles to better understand the underlying neuroanatomy driving the results of this study. Contrary to previous studies (15), elevations in BP were not related to FA. This may be due, in part, to the fact that we only obtained one BP recording as opposed to the recommended minimum of two (65) that would have reduced variability inherent in a single measurement. Having said that, previous studies show that early elevations in BP may impact the cortical mantle with white matter damage emerging later as chronic hypoperfusion ensues (53); unfortunately we did not have the sample size to divide our group by age-decades to investigate this possibility. Finally, although cluster-based thresholding procedures to adjust for multiple comparisons are standard in the literature, these procedures may still result in an inflated Type-I error (66).

Similar to their impact on the peripheral nervous system (67), CVD-RFs may induce perturbations across multiple physiological pathways, resulting in distinct yet phenotypic brain alterations that may ultimately contribute to pathological aging and dementia. Given that CVD- RFs are modifiable, and optimal control recommendations continue to gravitate toward a more conservative approach (54), understanding their distinct and overlapping associations on brain aging in early, mid-, and late-life as distinct time periods may help highlight the beneficial effects of lowering levels of cardiovascular risk at particular points across the lifespan.

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# **Figure Legends**

Figure 1. Cortical thickness associations between (A) BP and right lateral occipital; (B) Chol and left caudal medial frontal; (C) MetabDys and left inferior parietal; (D) MetabDys and left precuneus; (E) MetabDys and right superior parietal NOTE: Whole brain cortical thickness results derived from FreeSurfer. Maps were smoothed with a 15mm full width half maximum Gaussian kernel and correction for multiple comparisons was performed using the Monte Carlo Null-z simulation (p<0.05). All analyses have been adjusted for age and sex. Red indicates negative associations with cortical thickness and yellow indicates positive associations with cortical thickness.

Figure 2. TBSS results in red overlayed on the mean white matter tract skeleton (in green) derived from FSL using non-parametric permutation testing and cluster-based thresholding correction (p<0.001, cluster size>100 voxels). Significant cluster voxels associated with GluDys and fractional anisotropy adjusted for age and sex are depicted by the arrows pointing to the right inferior longitudinal fasciculus (ILF) and the right and left superior longitudinal fasciculus (SLF).

Table I: Sample characteristics

	Cognitive	Neuroimaging	p-value
	Sample (n=124)	Sample (n=86)	
DEMOGRAPHICS			
Age, years	59.8 <u>+</u> 13.1	61.7 <u>+</u> 12.6	0.31
Education, years	15.6 <u>+</u> 2.7	15.9 <u>+</u> 2.8	0.55
Sex (M:F)	62:62	43:43	0.56
Race (Caucasian:African	62:51:11	51:28:7	0.69
American:Other)			
MMSE	28.9 <u>+</u> 1.2	29.0 <u>+</u> 1.1	0.42
CARDIOVASCULAR RISK			p-value
FACTORS			
Systolic blood pressure, mmHg	132.4 <u>+</u> 15.2	133.1 <u>+</u> 13.7	0.64
Diastolic blood pressure, mmHg	79.6 <u>+</u> 9.5	80.3 <u>+</u> 10.0	0.61
Body mass index, kg/m <sup>2</sup>	29.9 <u>+</u> 7.7	28.2 <u>+</u> 6.1	0.18
Total cholesterol, mg/dl	186.6 <u>+</u> 44.6	184.8 <u>+</u> 42.6	0.95
LDL cholesterol, mg/dl	103.2 <u>+</u> 39.1	99.2 <u>+</u> 34.9	0.62
HDL cholesterol, mg/dl	57.7 <u>+</u> 19.1	61.0 <u>+</u> 19.6	0.17
Triglycerides, mg/dl	128.2 <u>+</u> 67.0	122.3 <u>+</u> 64.9	0.52
Glucose, mg/dl	103.4 <u>+</u> 29.5	101.3 <u>+</u> 23.3	0.94
HbA1c, %	6.3 <u>+</u> 1.2	6.1 <u>+</u> 0.9	0.41

Note: All p>0.05. All values represent mean<u>+</u>standard deviation unless otherwise noted.

M:F=male:female, MMSE=Mini-Mental State Examination, HbA1c=hemoglobin A1c

	Component 1	Component 2	Component 3	Component 4
Total	0.990	-0.049	-0.052	-0.042
Cholesterol				
LDL Cholesterol	0.954	0.030	0.045	0.075
Glucose	0.001	0.902	0.014	-0.121
Hemoglobin	-0.063	0.889	0.239	0.080
A1c				
Triglycerides	0.123	0.002	0.828	-0.140
HDL	-0.269	0.175	0.794	0.151
Cholesterol				
Body Mass	0.124	0.422	0.521	0.122
Index				
Systolic Blood	-0.046	-0.022	0.020	0.854
Pressure				
Diastolic Blood	0.075	0.008	0.019	0.855
Pressure				
Eigenvalue	2.367	1.982	1.544	1.125

# Table II: Varimax rotated factor matrix from PCA

Note: Minimum eigenvalue extraction=1.

Table III: Regression results for the association between cardiovascular risk factor and cognitive domains

	Learning	Memory	Executive	Attention/
			Function	Information
				Processing
	β, SE, (p-	β, SE, (p-	β, SE, (p-	β, SE, (p-
	value)	value)	value)	value)
Cholesterol	-0.008	0.046	0.011	0.031
	0.071	0.068	0.057	0.070
	(0.91)	(0.50)	(0.84)	(0.66)
Glucose	-0.088	-0.076	-0.114	-0.209
	0.075	0.071	0.060	0.074
	(0.24)	(0.29)	(0.058)	(0.006)*
Metabolic	-0.115	-0.116	0.017	0.089
	0.094	0.090	0.075	0.093
	(0.22)	(0.20)	(0.82)	(0.34)
Blood Pressure	-0.193	-0.224	-0.142	-0.153
	0.081	0.077	0.065	0.080
	(0.019)*	(0.005)*	(0.031)*	(0.060)
1	1	1	1	1

Note: \*p<0.050, analyses were adjusted for age, education, and sex.

Table IV: Significant clusters of cortical thickness associated with cardiovascular risk factor domain scores

	Region	Cluster Size	Max Z	X, Y, Z
		(mm <sup>2</sup> )		Talairach
				Coordinates
Cholesterol	Left Caudal	1450.9	3.377	-38.3, 9.4, 49.3
	Middle Frontal			
Metabolic	Right Superior	7117.6	3.899	22.8, -40.9, 57.8
	Parietal			
	Left Inferior	4817.1	3.222	-39.3, -70.0,
	Parietal			34.6
	Left Precuneus	1969.9	3.812	-6.8, -59.5, 41.8
Blood Pressure	Right Lateral	1699.7	-3.644	43.8, -77.8, -6.2
	Occipital			

NOTE: Whole brain results adjusted for age and sex derived from Freesurfer. Maps were smoothed with a 15mm FWHM Gaussian kernel and corrected for multiple comparisons using the Monte Carlo Null-z simulation (p<0.050).

Supplementary Table I: Multiple linear regression results for the association between cardiovascular risk factor component scores and cortical thickness regions of interest

	Cholesterol β, SE, (p-value)	Glucose β, SE, (p-value)	Metabolic β, SE, (p-value)	Blood Pressure β, SE, (p-value)
Caudal Anterior Cingulate L	-0.023, 0.017, (0.19)	0.000, 0.017, (0.99)	-0.014, 0.021, (0.50)	0.004, 0.020, (0.84)
R	-0.010, 0.016, (0.51)	0.001, 0.016, (0.96)	0.006, 0.019, (0.73)	-0.001, 0.018, (0.94)
Caudal Middle Frontal L	0.018, 0.006, (0.003)*	-0.010, 0.006, (0.12)	-0.005, 0.008, (0.47)	0.012, 0.007, (0.092)
R	0.005, 0.006, (0.39)	-0.009, 0.006, (0.11)	-0.004, 0.007, (0.59)	0.010, 0.006, (0.10)
Cuneus L	0.003, 0.006, (0.61)	0.000, 0.006, (0.98)	0.006, 0.007, (0.41)	0.003, 0.007, (0.67)
R	-0.001, 0.006. (0.90)	0.000, 0.006. (0.99)	0.002, 0.007, (0.83)	-0.003, 0.007, (0.65)
Entorhinal L	0.018, 0.018, (0.30)	-0.014, 0.018, (0.42)	0.001, 0.022, (0.95)	0.002, 0.020, (0.92)
R	-0.010, 0.022, (0.67)	0.002, 0.022, (0.94)	-0.038, 0.027, (0.16)	0.033, 0.025, (0.19)
Fusiform L	0.008, 0.006, (0.20)	-0.004, 0.006, (0.54)	-0.005, 0.008, (0.48)	0.001, 0.007, (0.87)
R	0.012, 0.007, (0.10)	-0.007, 0.007, (0.36)	-0.008, 0.009, (0.36)	-0.012, 0.008, (0.13)
Inferior Parietal L	0.000, 0.006, (0.97)	0.012. 0.005, (0.036)*	0.011, 0.007, (0.095)	-0.030, 0.006, (0.11)
R	0.007, 0.006, (0.22)	0.001, 0.006, (0.81)	-0.002, 0.007, (0.81)	-0.007, 0.007, (0.31)
Inferior Temporal L	0.010, 0.007, (0.15)	-0.007, 0.007, (0.30)	-0.014, 0.008, (0.076)	-0.006, 0.008, (0.43)
R	-0.003, 0.007, (0.67)	0.001, 0.007, (0.89)	-0.009, 0.009, (0.32)	-0.015, 0.008, (0.072)
Lateral Occipital L	0.006, 0.006, (0.29)	0.008, 0.006, (0.17)	0.005, 0.007, (0.50)	-0.006, 0.007, (0.40)
R	0.006, 0.006, (0.34)	0.007, 0.006, (0.27)	-0.006, 0.007, (0.42)	-0.041, 0.007, (0.041)*
Lateral Orbitofrontal L	0.008, 0.007, (0.28)	-0.011, 0.007, (0.14)	-0.036, 0.008, (<0.001)*	0.005, 0.008, (0.53)
R	0.000, 0.007, (0.99)	-0.009, 0.007, (0.24)	-0.016, 0.009, (0.068)	0.007, 0.008, (0.43)
Lingual L	0.001, 0.005, (0.82)	-0.005, 0.005, (0.34)	0.000, 0.006, (0.99)	-0.001, 0.006, (0.90)
R	0.004, 0.006, (0.53)	0.000, 0.006, (0.93)	-0.003, 0.007, (0.69)	0.005, 0.007, (0.48)
Medial Orbitofrontal L	0.003, 0.009, (0.78)	-0.007, 0.009, (0.46)	-0.025, 0.011, (0.023)*	0.000, 0.010, (0.99)
R	-0.009, 0.011, (0.42)	-0.016, 0.011, (0.16)	-0.035, 0.013, (0.007)*	-0.006, 0.012, (0.614)
Middle Temporal L	0.002, 0.006, (0.75)	-0.010, 0.006, (0.088)	-0.018, 0.007, (0.009)*	-0.012, 0.006, (0.068)
R	0.002, 0.006, (0.70)	-0.003, 0.006, (0.66)	-0.011, 0.008, (0.17)	-0.015, 0.007, (0.039)*
Parahippocampal L	0.025, 0.017, (0.14)	-0.023, 0.017, (0.17)	-0.032, 0.020, (0.12)	-0.024, 0.019, (0.21)
R	0.008, 0.015, (0.62)	-0.014, 0.015, (0.37)	-0.020, 0.019, (0.28)	-0.009, 0.017, (0.59)
Parsopercularis L	-0.001, 0.006, (0.86)	-0.012, 0.005, (0.029)*	-0.011, 0.007, (0.095)	0.001, 0.006, (0.93)
R	0.009, 0.006, (0.17)	-0.006, 0.006, (0.32)	-0.009, 0.008, (0.26)	-0.005, 0.007, (0.49)

Parsorbitalis L	-0.006, 0.010, (0.53)	-0.005, 0.010, (0.65)	-0.031, 0.012, (0.009)*	0.002, 0.011, (0.89)
R	-0.014, 0.009, (0.14)	0.002, 0.010, (0.87)	-0.007, 0.012, (0.55)	0.013, 0.011, (0.22)
Parstriangularis L	0.005, 0.007, (0.54)	-0.008, 0.007, (0.30)	-0.014, 0.009, (0.12)	0.009, 0.008, (0.30)
R	-0.002, 0.008, (0.80)	-0.012, 0.008, (0.13)	-0.014, 0.009, (0.14)	-0.003, 0.009, (0.72)
Posterior Cingulate L	0.007, 0.007, (0.32)	0.000, 0.007, (0.96)	0.002, 0.008, (0.80)	-0.006, 0.008, (0.41)
R	-0.001, 0.007, (0.88)	0.011, 0.007, (0.13)	0.005, 0.009, (0.60)	-0.009, 0.008, (0.29)
Precuneus L	0.004, 0.005, (0.48)	0.006, 0.005, (0.28)	0.011, 0.006, (0.078)	0.001, 0.006, (0.88)
R	0.006, 0.006, (0.27)	0.009, 0.006, (0.11)	0.003, 0.007, (0.70)	-0.004, 0.006. (0.54)
Rostral Anterior Cingulate L	0.006, 0.015, (0.71)	-0.013, 0.015, (0.40)	-0.020, 0.018, (0.27)	-0.005, 0.017, (0.77)
R	0.019, 0.014, (0.18)	-0.024, 0.014, (0.091)	-0.024, 0.017, (0.17)	-0.002, 0.016, (0.90)
Rostral Middle Frontal L	-0.002, 0.005, (0.66)	-0.003, 0.005, (0.60)	-0.008, 0.006, (0.22)	0.005, 0.006, (0.41)
R	-0.001, 0.005, (0.86)	-0.005, 0.005, (0.37)	-0.002, 0.006, (0.78)	0.007, 0.006, (0.21)
Superior Frontal L	0.003, 0.005, (0.59)	0.003, 0.005, (0.53)	0.006. 0.006, (0.37)	0.003, 0.006, (0.65)
R	0.010, 0.005, (0.053)	0.001, 0.005, (0.90)	0.002, 0.006, (0.70)	0.013, 0.006, (0.026)*
Superior Parietal L	0.006, 0.006, (0.35)	0.009, 0.006, (0.14)	0.014, 0.007, (0.061)	-0.004, 0.007, (0.60)
R	0.006, 0.006, (0.28)	0.008, 0.006, (0.16)	0.016, 0.007, (0.015)*	-0.001, 0.006, (0.93)
Superior Temporal L	0.004, 0.007, (0.59)	-0.011, (0.007), (0.12)	-0.006, 0.008, (0.49)	-0.015, 0.008, (0.061)
R	0.006, 0.007, (0.39)	-0.012, 0.007, (0.12)	-0.001, 0.009, (0.88)	-0.019, 0.008, (0.027)*

Note: L=left hemisphere, R=right hemisphere, \*p<0.05, All analyses have been adjusted for age, sex

Supplementary Table II: Association between individual cardiovascular disease risk factor components and cortical thickness linear regression results where p was  $\leq 0.050$ 

# a) Cholesterol Domain

	Total Cholesterol	LDL Cholesterol
	β, SE, (p-value)	β, SE, (p-value)
L. Caudal Middle Frontal	0.002, 0.006, (0.001)*	0.015, 0.006, (0.013)*

Note: \*p<0.05, All analyses have been adjusted for age, sex

#### b) Glucose domain

	Glucose	HbA1c
	β, SE, (p-value)	β, SE, (p-value)
L. Inferior Parietal	0.011, 0.005, (0.033)*	0.009, 0.005, (0.090)
L. Parsopercularis	-0.010, 0.005, (0.045)*	-0.010, 0.005,(0.048)*

Note: \*p<0.05, All analyses have been adjusted for age, sex

# c) Metabolic domain

	Triglycerides	HDL Cholesterol	BMI
	$\beta$ , SE, (p-value)	$\beta$ , SE, (p-value)	β, SE, (p-value)
L. Lateral Orbitofrontal	-0.024, 0.006,(<0.001)*	0.026, 0.007,(<0.001)*	-0.016, 0.007, (0.014)*
L. Medial Orbitofrontal	-0.016, 0.008, (0.052)	0.025, 0.009, (0.007)*	-0.006, 0.008, (0.45)
R. Medial Orbitofrontal	-0.025, 0.010, (0.013)*	0.033, 0.011, (0.003)*	-0.009, 0.010, (0.37)
L. Middle Temporal	-0.009, 0.005, (0.090)	0.009, 0.006, (0.13)	-0.015, 0.005, (0.004)*
L. Parsorbitalis	-0.020, 0.009, (0.030)*	0.023, 0.010, (0.025)*	-0.015, 0.009, (0.098)
R. Superior Parietal	0.008, 0.005, (0.11)	-0.013, 0.006,(0.021)*	0.009, 0.005, (0.068)

Note: \*p<0.05, All analyses have been adjusted for age, sex

# d) Blood Pressure (BP) domain

	Systolic BP	Diastolic BP
	β, SE, (p-value)	$\beta$ , SE, (p-value)
R. Lateral Occipital	-0.014, 0.006, (0.017)*	-0.007, 0.006, (0.23)
R. Middle Temporal	-0.015, 0.006, (0.15)*	-0.008, 0.006, (0.23)
R. Superior Frontal	0.011, 0.005, (0.029)*	0.008, 0.005, (0.093)
R. Superior Temporal	-0.011, 0.007, (0.14)	-0.017, 0.007,(0.017)*

Note: \*p<0.050, All analyses have been adjusted for age, sex

Supplementary Table III: Multiple linear regression results for the association between cardiovascular risk factor component scores and diffusion tensor imaging regions of interest

	Cholesterol β, SE, (p-value)	Glucose β, SE, (p-value)	Metabolic β, SE, (p-value)	Blood Pressure $\beta$ , SE, (p-value)
Superior Longitudinal	0.002, 0.001, (0.013)*	-0.002, 0.001, (0.014)*	0.000, 0.001, (0.99)	0.000, 0.001, (0.99)
Fasciculus				
Superior Longitudinal	0.002, 0.001, (0.033)*	-0.003, 0.001, (0.013)*	0.000, 0.001, (0.82)	0.000, 0.001, (0.91)
Fasciculus Temporal				
Cingulum Hippocampus	0.001, 0.001, (0.37)	-0.002, 0.001, (0.17)	-0.002, 0.002, (0.16)	0.000, 0.001, (0.82)
Inferior Fronto-Occipital	0.002, 0.001, (0.089)	-0.003, 0.001, (0.007)*	0.000, 0.001, (0.92)	0.002, 0.001, (0.24)
Fasciculus				
Inferior Longitudinal	0.002, 0.001, (0.23)	-0.003, 0.001, (0.087)	0.000, 0.004, (0.97)	0.002, 0.002, (0.15)
Fasciculus				
Anterior Thalamic Radiation	0.001, 0.001. (0.45)	-0.001, 0.001, (0.16)	0.002, 0.001, (0.13)	0.001, 0.001, (0.15)
Uncinate Fasciculus	0.003, 0.002, (0.12)	-0.003, 0.002, (0.13)	-0.001, 0.002, (0.73)	-0.001, 0.002, (0.51)

Note: \*p<0.05, All analyses have been adjusted for age, sex

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Figure 1. Cortical thickness associations between (A) BP and right lateral occipital; (B) Chol and left caudal medial frontal; (C) MetabDys and left inferior parietal; (D) MetabDys and left precuneus; (E) MetabDys and right superior parietal NOTE: Whole brain cortical thickness results derived from FreeSurfer. Maps were smoothed with a 15mm full width half maximum Gaussian kernel and correction for multiple comparisons was performed using the Monte Carlo Null-z simulation (p<0.05). All analyses have been adjusted for age and sex. Red indicates negative associations with cortical thickness and yellow indicates positive associations with cortical thickness.



Figure 2. TBSS results in red overlayed on the mean white matter tract skeleton (in green) derived from FSL using non-parametric permutation testing and cluster-based thresholding correction (p<0.001, cluster size>100 voxels). Significant cluster voxels associated with GluDys and fractional anisotropy adjusted for age and sex are depicted by the arrows pointing to the right inferior longitudinal fasciculus (ILF) and the right and left superior longitudinal fasciculus (SLF).

