# Calorie Restriction Reduces the Influence of Glucoregulatory Dysfunction on Regional Brain Volume in Aged Rhesus Monkeys

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Insulin signaling dysregulation is related to neural atrophy in hippocampus and other areas affected by neurovascular and neurodegenerative disorders. It is not known if long-term calorie restriction (CR) can ameliorate this relationship through improved insulin signaling or if such an effect might influence task learning and performance. To model this hypothesis, magnetic resonance imaging was conducted on 27 CR and 17 control rhesus monkeys aged 19-31 years from a longitudinal study. Voxel-based regression analyses were used to associate insulin sensitivity with brain volume and microstructure cross-sectionally. Monkey motor assessment panel (mMAP) performance was used as a measure of task performance. CR improved glucoregulation parameters and related indices. Higher insulin sensitivity predicted more gray matter in parietal and frontal cortices across groups. An insulin sensitivity  $\times$  dietary condition interaction indicated that CR animals had more gray matter in hippocampus and other areas per unit increase relative to controls, suggesting a beneficial effect. Finally, bilateral hippocampal volume adjusted by insulin sensitivity, but not volume itself, was significantly associated with mMAP learning and performance. These results suggest that CR improves glucose regulation and may positively influence specific brain regions and at least motor task performance. Additional studies are warranted to validate these relationships. Diabetes 61:1036-1042, 2012

ower insulin sensitivity and reduced insulinmediated glucose uptake can adversely impact the brain. Glucoregulatory dysfunction is related to less gray matter (GM) volume cross-sectionally and longitudinally in medial temporal lobe, prefrontal cortex, and other areas impacted by normal aging, neurovascular disorders, and Alzheimer disease (AD). Such

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relationships are seen in both rhesus monkeys (1) and humans (2,3). Importantly, insulin-signaling dysfunction in AD patients can negatively influence brain volume in the absence of type 2 diabetes (3), suggesting that mild to moderate glucoregulatory perturbation may be detrimental over time. Studies in rodents show that medial temporal lobe and prefrontal cortex have dense insulin receptor staining and may rely on insulin signaling for optimal glucose uptake and utilization (2,4). AD is characterized by reduced insulin sensitivity, transcription of mitochondrial metabolism genes, and central glucose uptake (5,6). Lower insulin sensitivity may pathogenically affect the brain through microvascular disease (7), increased production of advanced glycation end products and free radicals (8,9), or lower cerebral blood flow and glucose transport (10).

Despite these relationships between insulin signaling and brain, studies vary widely in the glucoregulatory measures and brain areas that are assessed (11). It is therefore of interest to use voxel-wise analysis methods (12) to investigate where insulin sensitivity variation is associated with regional volume or tissue microstructure across the brain. Our group has previously reported on the longitudinal effects of calorie restriction (CR) regarding glucose regulation in aged rhesus macaques since middle age (13–15). CR led to improved glucose tolerance and higher insulin sensitivity, effects that may benefit areas like hippocampus and prefrontal cortex and mediate improved task learning and performance. This cohort therefore afforded a unique opportunity to look at the effects of CR on glucose regulation and its association with brain and behavior in a primate species.

In this study, an index of insulin sensitivity, S<sub>I</sub>, was derived from a glucose tolerance test. This measure represents the ability of insulin to promote glucose uptake and inhibit hepatic glucose production (13,16). Insulin sensitivity in the periphery and brain strongly correspond (17). We hypothesized that higher S<sub>I</sub> would predict more GM volume in brain areas with dense insulin receptor staining and that are affected by insulin signaling dysregulation (2,18–21). Furthermore, it has been well established that CR in nonhuman primates has several beneficial effects related to glucose regulation, such as improved vascular health, lower triglycerides, and other factors that delay age-associated pathogenesis (22). Thus, we tested the interaction between S<sub>I</sub> and dietary condition, to see if CR monkeys exhibited more volume or tissue density per S<sub>I</sub> unit increase versus controls beyond the association between S<sub>I</sub> and brain seen across both dietary groups. In other words, this interaction tested if there was a regionspecific beneficial effect for restricted animals. Several

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insulin signaling, glycation, and vascular measures were tested as potential mediating factors (1,8). Finally, due to hippocampus being susceptible to glucoregulatory dysfunction (3) and its role in AD, we performed a region of interest (ROI) analysis limited to that region. The ROI analysis, which was independent from the voxel-wise results (23), tested the extent to which predicted changes in hippocampus specific to S<sub>I</sub> were associated with visuomotor performance on the monkey motor assessment panel (mMAP) (24,25). The hippocampus is in part involved in learning new spatiomotor sequences (26).

# **RESEARCH DESIGN AND METHODS**

**Subjects.** Forty-four rhesus monkeys (Macaca mulatta) between 19 and 31 years of age were included in this study. Seventeen animals were fed ad libitum for approximately 8 h/day, whereas the remaining 27 subjects were fed 30% fewer calories relative to their own baseline intake. Length of CR diet was 12-17 years and initiated in middle age. Demographic data are available in Table 1. These monkeys were the remaining subjects of a longitudinal CR project begun at the Wisconsin National Primate Research Center in 1989. Details of the CR manipulation, housing, and husbandry have been described previously (13,27). Magnetic resonance imaging data acquisition and preprocessing. Scan session and image acquisition parameters have been detailed elsewhere (28). Briefly, animals were anesthetized while images were acquired using a General Electric 3.0 T scanner (GE Medical Systems, Milwaukee, WI). T1-weighted volumetric scans were used to determine regional volumes. Diffusion tensor imaging was acquired to quantitatively estimate coherence of white matter tracts by computing fractional anisotropy, axial diffusivity, and radial diffusivity, whereas microstructural density of brain parenchyma was determined using mean diffusivity (29). T1-weighted scan parameters were: repetition time = 8.772 ms; echo time = 1.876 ms; inversion time = 600 ms; flip angle =  $10^{\circ}$ ; number of excitations = 2; matrix =  $256 \times 224$ ; and field of view = 160 mm. A total of 124 coronal slices were acquired with a thickness of 0.7 mm and no gap, resulting in  $0.625 \times 0.625$  $\times$  0.7 mm voxels. For diffusion tensor imaging, a diffusion-weighted echo-planar imaging sequence with 12 gradients was used with the following parameters: repetition time/echo time = 10,000/77.2 ms; b = 816 s/mm<sup>2</sup>; number of excitations = 6; field of view = 140 mm; in-plane matrix =  $256 \times 256$ ; and slice thickness = 2.5 mm, no gap, and 35 slices. Voxels were resampled to  $0.5 \times 0.5 \times 0.5$ mm for T1-weighted and diffusion tensor imaging sequences. T2-weighted scans were acquired for global volume estimation and are described elsewhere (28).

**Imaging artifacts.** All images were inspected to locate scan artifacts that could adversely affect the voxel-wise analyses. Three diffusion tensor-imaging scans and one volumetric scan had artifact abnormalities (motion, respiratory, phase, other) or gross misregistration with the template brain that rendered data unusable. Six additional diffusion tensor-imaging scans (controls = 2, CR = 4) had one or more 1-to 2-mm abnormalities located in the white matter of the dorsal convexity that precluded analysis.

**Magnetic resonance imaging preprocessing.** Preprocessing of magnetic resonance images for use in voxel-wise analyses has been described previously (28,30). Briefly, T1-weighted images were segmented and normalized to 112RM-SL atlas space (28) with Diffeomorphic Anatomical Registration Through Exponentiated Lie algebra (31). Volumetric segments were smoothed using a 4-mm full width at half maximum Gaussian kernel. Fractional anisotropy, mean diffusivity, axial diffusivity, and radial diffusivity maps were computed using DTIFTT in FSL (www.fnrib.ox.ac.uk/fsl/). Diffusion tensorimaging measurements were aligned to the 112RM-SL atlas space in SPM and smoothed with a 4-mm smoothing kernel.

Glucoregulation and other physiological indices. We have previously described the frequently sampled glucose tolerance test procedure that provides data on basal and acute glucose and insulin signaling dynamics (13). Derived indices relevant to this report are described in Table 1. Glucoregulation measures were collected within 6 months of the magnetic resonance imaging (MRI) scan. Insulin sensitivity, or S<sub>I</sub>, was calculated using the Minimal Method Model (32). This measure provides an accurate quantification of  $S_{\rm L}$  as well as other aspects of glucose kinetics, and is far more practical than other methods like the glucose-clamp technique. Plasma glucose was measured using the glucose oxidase method (Yellow Springs Instruments, Yellow Springs, OH). Insulin levels were determined using double antibody radioimmunoassay (Millipore, Billerica, MA). The homeostatic assessment of insulin resistance (HOMA-IR) was calculated using basal glucose and basal insulin (33). HOMA-IR was calculated in order to see if basal insulin resistance explained variance beyond its converse construct, which would be insulin sensitivity measured by S<sub>I</sub>. Homocysteine was collected using methods previously described (30). Glucoregulatory data near the time of the MRI scan were not available for one CR and one control monkey.

**Glucoregulatory impairment.** Animals were classified by an expert (R.J.C.) as having normal, prediabetic/at-risk, or type 2 diabetes-like profiles using established criteria (15).

Anatomical region of interest: hippocampus. In order to independently assess (23) if glucoregulatory dysfunction might influence motor learning via  $S_I$  predicted variation in brain, an expert (A.A.W.) drew a mask on the 112RM-SL atlas to isolate the bilateral hippocampus using methods previously described (34). Mean GM volume was extracted from the ROI for all normalized monkey brains and was used in conjunction with  $S_I$  to predict performance on a motor task described next.

**Motor learning and performance.** It was of interest to see if the relationship between insulin signaling and hippocampal volume predicted changes in motor task learning and performance. To this end, our cohort has previously been tested on the mMAP (25), which required subjects to retrieve an appetitive stimulus from a flat platform, straight rod, or curved rod (24). An automated system recorded the number of seconds necessary to reach from the home cage to the first area of the affixed apparatus (reaction time), from the first to second area (coarse motor movement), and from the second area to a small receptacle that held the stimulus (fine motor movement). We used fine motor performance data from the most difficult task (i.e., curved rod), because CR animals acquire and perform this task more quickly than controls (25). Animals did not differ on simpler tasks. To test our hypothesis regarding insulin signaling and hippocampal volume adjusted or not adjusted by S<sub>I</sub>. The sample size for this analysis was 26 animals (C = 7; CR = 19).

Voxel-wise statistical analyses: structural MRI. To investigate regional brain associations of S<sub>I</sub>, multiple regression voxel-wise analyses were conducted in SPM8 (12). For the purposes of this report, this type of regression technique produces t-statistic, color-coded result maps that are the product of a regression model performed at every voxel in the brain for a given modality. Contiguous groups of voxels that attain statistical significance, called clusters, will thus overlap with and implicate different brain regions. In this study, volumetric or diffusion tensor-imaging scans were entered as the dependent variable. The independent variable was S<sub>I</sub>. Covariates included age at scan, sex, dietary condition, and the S<sub>t</sub> term when testing an interaction (see below). Analyses of volume additionally covaried a global index of either gray or white matter (28). Our primary analysis of interest was testing a  $S_{\rm I}$  imes dietary condition interaction term to see if CR monkeys showed more volume or microstructure per S<sub>I</sub> unit change beyond the association seen with S<sub>I</sub> alone, suggesting a further beneficial effect. The voxel and cluster level thresholds were set at P < 0.005 (uncorrected) and P < 0.05 (corrected). Type 1 error was minimized by first using an omnibus F-contrast (P < 0.05, uncorrected) for  $S_{I\!\!\prime}$  dietary condition, and  $S_{I}$   $\times$  dietary condition to mask subsequent contrasts, followed by Monte Carlo simulations to estimate cluster sizes that would occur due to chance (25,30). Clusters required 280 contiguous voxels to reach significance at P < 0.05 (corrected). Reported whole-brain cluster coordinates correspond to the Saleem-Logothetis atlas (35) and are displayed on the 112RM-SL template image (28). Standard rhesus monkey atlases were used to identify fibers (36) and subcortical structures (37).

Statistical analyses: brain-physiology and brain-behavior associations. Tests were conducted using SPSS 18.0 (SPSS Inc., Chicago, IL) at an  $\alpha$  of 0.05. Logarithmic transformations were used to adjust nonnormally distributed indices. ANOVA tested group differences for demographic and biological variables. Multiple regression was used to determine which basal and frequently sampled glucose tolerance test variables significantly explained error variance in the  $S_I$  association and interaction voxel-wise analyses. The first regression block included HOMA-IR to account for basal insulin regulation. The second block contained a priori variables of interest directly related to  $S_I$ : basal glucose and insulin, compensatory pancreatic sensitivity represented by Disposition Index (13), and glycosylated hemoglobin levels. The third block contained the vascular risk factor homocysteine.

Brain-behavior mediational models tested whether or not  $S_{\Gamma}$  predicted variation in brain was correlated with motor learning and performance. To avoid circular analysis (23), an independent anatomical ROI approach was used to derive mean image signal (e.g., mean GM volume) within the bilateral hippocampus.  $S_{I}$  was then linearly regressed onto the hippocampal signal. The Pearson's statistic was then used to correlate the predicted change in hippocampal volume due  $S_{I}$  with the mean number of seconds it took for a monkey to complete the fine motor portion of the curved rod mMAP task during initial acquisition (i.e., the learning phase) and when the animal reached proficiency (25).

# RESULTS

**Subject characteristics and biological indices.** See Table 1 for descriptive data and statistics. The mean age, sex composition, and total brain volume of the two dietary

#### TABLE 1

Demographics, total brain volume, and glucoregulatory values for control and CR monkeys

	Control	Range	CR	Range	P value
Demographics and global volume					
Age (years)	$23.84 \pm 2.79$	19 - 27	$24.32 \pm 2.77$	19-31	NS
Sex (male/female) $(n)$	7/10	NA	11/16	NA	NS
Body weight (kg)	$12.39 \pm 2.84$	9-19	$8.94 \pm 1.71$	6-12	0.001
Total brain volume (mm <sup>3</sup> )	$86.80 \pm 10.56$	67-105	$82.45 \pm 9.20$	56-99	NS
Physiological indices					
Basal glucose (mg/dL)	$82.56 \pm 41.77$	52-218	$60.78 \pm 7.44$	51-82	0.009
Basal insulin (µU/mL)	$47.81 \pm 50.47$	4-199	$19.26 \pm 17.93$	4-82	0.009
Disposition Index $([\Delta I_{(0-30)}/\Delta G_{(0-30)}]/I_b)$	$1,931 \pm 1,521$	138 - 5,705	$4,244 \pm 2,411$	106-12,483	0.001
$HbA_{1c}$ (%)	$10.48 \pm 4.41$	7-22	$8.59 \pm 1.56$	5-12	0.043
Homocysteine (µmol/L)	$8.16 \pm 3.30$	5-15	$9.88 \pm 5.09$	5-24	NS
HOMA-IR ( $[G_b \times I_b]/405$ )	$8.62 \pm 7.63$	0.51 - 28	$2.99 \pm 2.86$	0.50 - 12	0.001
$\mathrm{S_{I}}~(\mu\mathrm{U/mL} imes~10^{4}~\mathrm{min^{-1}})$	$3.28 \pm 2.31$	0.23 - 7	$7.44 \pm 4.11$	0.61 - 16	0.001

The unit of measurement, mean  $\pm$  SD, and range for a given variable is noted for control and then CR monkeys. Insulin sensitivity, as assessed by S<sub>I</sub>, quantitatively represents the capacity of insulin to stimulate glucose uptake and inhibit glucose production. Disposition Index represents pancreatic insulin sensitivity and is calculated as the ratio of insulin secretion to glucose uptake 0–30 min after glucose bolus as compared with prechallenge insulin levels. Basal glucose, basal insulin, and HOMA-IR are based on samples collected before glucose bolus administration. Glucoregulatory data besides basal measures near the time of scan were not available for one CR and one control monkey. ANCOVA was used to test for dietary group effects.

conditions did not differ. CR monkeys showed expected benefits in insulin signaling and related indices (22).

**Prediabetic and diabetic glucoregulatory impairment.** Six controls and zero CR monkeys were classified as having preclinical (n = 4) or diabetes-like glucoregulatory dysfunction (n = 2). Diagnosis was not a significant covariate in regression analyses.

**Regional GM:**  $S_I$  association. To examine the association of  $S_I$  on regional GM,  $S_I$  was regressed onto GM volume voxels across all subjects. As indicated in Fig. 1*A*–*C* (yellow-orange areas) and Table 2, higher  $S_I$  predicted more GM in motor and somatosensory cortices. Fig. 1*G* depicts this relationship by illustrating the correlation between  $S_I$  and the voxel with the highest (peak) *t*-statistic located in left primary motor cortex. The association of  $S_I$  and GM was comparable for both control ( $R^2 = 0.516$ ; P < 0.001) and CR ( $R^2 = 0.324$ ; P < 0.001) monkeys.

Regional GM: predictors of S<sub>I</sub> association. Measurements directly or indirectly related to glucoregulation that are described in Table 1 may elucidate potential mechanisms underlying the relationship between S<sub>I</sub> and regional GM. Therefore, follow-up multiple linear regression was conducted. The dependent variable was the predicted volume change in left motor cortex shown in Fig. 1G. Different regression models using stepwise, forward, or enter methods produced similar results (data not shown). The final adjusted regression model predicted 20.7% of the variance across both groups [F(6,35) = 4.170; P < 0.01]. Although the influence of S<sub>I</sub> on GM was not significantly mediated by HOMA-IR or basal glucose, increased basal insulin was modestly related to less GM ( $R^2 = 0.133$ ; P < 0.05). Higher levels of homocysteine ( $R^2 = 0.169$ ; P < 0.05) and glycosylated hemoglobin ( $R^2 = 0.148$ ; P < 0.05) were also associated with less frontal GM volume.

**Regional GM:**  $S_I \times$  **dietary condition.** Given the salubrious effects of CR on vascular health and metabolic indices related to improved glucose regulation (22), we tested for a similar beneficial effect on the brain in CR monkeys. Thus, this contrast examined if CR monkeys had more GM volume per unit increase in  $S_I$  compared with controls beyond the association with  $S_I$  alone, suggesting a protective effect. As shown in Table 2 and depicted in

Fig. 1A (purple areas), voxel clusters were present in bilateral anterior hippocampus and both inferior and middle temporal gyri. The interaction is represented in Fig. 1Husing the peak t-statistic voxel. CR monkeys showed more GM as  $S_I$  increased ( $R^2 = 0.159$ ; P < 0.05). For controls, however, higher S<sub>I</sub> unexpectedly corresponded to less GM  $(R^2 = 0.449; P < 0.01)$ . To validate this result, a mean index of  $S_I$  was calculated using data from all glucose tolerance tests since the start of the project in 1989, which was up to 14 assessments depending on the length of time an animal spent in the study. This index or area under the curve estimates resulted in clusters and graphs similar to those produced using the S<sub>I</sub> value nearest to the MRI scan date (data not shown). Additional significant brain areas found in the interaction analysis were caudal perirhinal, entorhinal, and parahippocampal cortices, insula, amygdala, temporal pole, anterior cingulate cortex, and orbital and medial prefrontal cortices (Fig. 1A-F).

**Regional GM: predictors of S<sub>I</sub> × dietary condition effect.** Multiple linear regression was simultaneously performed on each dietary condition group to detect mediators that might explain the S<sub>I</sub> × dietary condition interaction effect. The dependent variable was the predicted change in hippocampal GM depicted in Fig. 1*H*. The same regression model was used as in the S<sub>I</sub> association analysis (see RESEARCH DESIGN AND METHODS). Table 3 indicates the result. Fig. 2*A*–*C* shows that control monkeys had strong associations between HOMA-IR, basal insulin, or basal glucose and GM related to the interaction. Fig. 2*D*–*F* shows nonsignificant relationships for the same variables in CR monkeys.

 $S_I$  predicted variation in hippocampus GM and fine motor performance. We finally wished to test if predicted changes in hippocampal GM related to  $S_I$  were associated with mMAP fine motor performance during the curved rod task (24). An independent analysis was conducted using  $S_I$  measured at the time closest to scan in 7 controls and 19 CR monkeys that had successfully learned the task. As reported elsewhere (25), this task was chosen because CR monkeys performed it significantly more quickly during acquisition (CR: 4.84 s; control: 6.26 s) and after gaining proficiency (CR: 3.08 s; control: 3.73 s).





FIG. 1. The relationship between S<sub>I</sub> and regional GM volume across subjects and an  $S_I \times$  dietary condition interaction testing such an association for each dietary group.  $S_I$  near the time of scan was not available for one control and one CR monkey. Sixteen controls and 26 CR monkeys were analyzed. Higher S<sub>I</sub> for both control and CR monkeys (yellow-orange color) corresponded to more GM in parietal and frontal cortices (A-C). For the interaction, CR animals showed more GM per unit increase in S<sub>I</sub> compared with controls in hippocampus, temporal pole, agranular insula, striatum, prefrontal cortices, and anterior cingulate cortex (purple color, A-F). To illustrate these results, a representative voxel from each analysis was graphed, both for the association of  $S_I$  across groups (G) and the interaction between groups (H). A sagittal view of the brain depicts the location of the coronal slices in A-F along a posterior-anterior axis. Color bars and color maps represent t values. GM volume is depicted in arbitrary units (A.U.). Brains are oriented such that the left hemisphere is on the left side.

Unadjusted hippocampal volume was not related to motor learning ( $R^2 = 0.034$ ; not significant) or proficient ( $R^2 = 0.01$ ; not significant) mMAP performance. By contrast, when first adjusting hippocampal volume by S<sub>I</sub>, this brain measure significantly explained 12% of the variance for mMAP performance during the acquisition phase (P = 0.042) and 24.1% (P = 0.005) after monkeys became proficient at the task (25).

**Regional white matter, diffusion tensor imaging, and**  $S_I$  analyses. No clusters exceeded the minimum number of voxels required for type 1 error correction in white matter volume or diffusion tensor imaging modalities.

# DISCUSSION

We hypothesized that higher insulin sensitivity, indexed by  $S_I$ , would predict more GM or denser microstructure in brain regions that are influenced by insulin signaling and impacted by neurodegenerative and neurovascular disorders. Across CR and control subjects, higher  $S_I$  was associated with more GM volume in somatosensory and motor cortices. These areas have low insulin receptor density relative to medial temporal lobe and prefrontal cortex in rodents (38). Interestingly,  $S_I$  interacted with dietary condition, for which CR monkeys with higher  $S_I$ 

had significantly more GM in hippocampus, insula, prefrontal cortex, and other regions with a high density of insulin receptors. Higher  $S_I$  among controls was unexpectedly related to less GM in these regions. Models suggest that prediabetes or type 2 diabetes in controls did not influence this result. Similar relationships between impaired glucoregulation and brain volume have been seen in AD patients with no history of type 2 diabetes (3). As such, mild to moderate insulin signaling dysregulation may negatively affect key brain areas.

Although the relationship between S<sub>I</sub> and GM among controls in the interaction was unexpected, it is not likely due to assay error or a sudden change in glucoregulatory dynamics. A similar interaction result was found when using a mean S<sub>I</sub> index derived from every glucose tolerance test conducted since 1989. More importantly, an estimate of basal insulin resistance, HOMA-IR, corresponded to less hippocampal GM in controls, a result that has been observed in humans (21). No relationship was seen between S<sub>I</sub> and white matter volume or tissue microstructure, which may reflect the role of insulin-like growth factor signaling rather than insulin in mediating oligodendrocyte development and growth factor-mediated preservation (39). Although tissue microstructure is negatively impacted by type 2 diabetes (40), it may be due to vascular pathology or other consequences instead of antecedent decreases in insulin sensitivity.

Our results suggest that there is wide variation in how insulin signaling may affect energy metabolism and other functions in brain. The central action of insulin on reducing oxidative damage or maintaining synaptic plasticity appears to be area-specific due to receptor density and binding dynamics (2,17,38). For example, intracerebrovascular treatment of rats with low doses of streptozotocin, which is normally toxic to pancreatic insulin-secreting cells and creates a diabetes-like state, reduced downstream phosphatidylinositol-3 kinase activity primarily in hippocampus but to a much lesser extent in frontal cortex without affecting peripheral glucose regulation (41). Intracerebrovascular administration of insulin also affected adenosine 5'-triphosphate storage in hippocampus but not parietotemporal cortex (4).

By extension, the relatively moderate relationship between higher S<sub>1</sub> and more GM in motor and somatosensory areas may be due to indirect mechanisms attributed to higher insulin concentrations. Chronic peripheral hyperinsulinemia is typically characterized by hyperglycemia and breakdown of the epithelial vasculature. Levels of glycosylated hemoglobin and homocysteine, a biomarker for vascular health, were significant mediators of the association between S<sub>I</sub> and GM. In brain, it is conceivable that a microvascular insult combined with other neuropathologies might synergistically reduce perfusion and glucose transport into parenchyma, leading to damage (42). For example, streptozotocin alone in transgenic APP/ PS1 mice produced more advanced glycation end products in microvasculature and worse spatial performance; similar pathophysiological effects were also seen in human cerebromicrovascular cells exposed to streptozotocin and amvloid B (9).

The interaction of  $S_I$  and GM between dietary groups revealed several important findings related to insulin signaling and brain in CR versus control monkeys. Ameliorative effects may be due to the direct action of insulin, although CR may act through related mechanisms such as less atherogenic dyslipidemia, lower expression of

#### TABLE 2

Voxel-wise	results	for	$\mathbf{S}_{\mathbf{I}}$	and	$\mathbf{G}\mathbf{M}$	volume

•			Cluster size
Location	x, y, z	t	(voxels)
S <sub>I</sub> association			
L primary motor cortex	-12, 16, 36	4.38	7,538
L somatosensory areas 3a and 3b	-22, 16, 23	3.88	
L primary motor cortex	-3, 12, 38	3.86	
R primary motor cortex	18, 19, 30	3.20	2,007
$S_I \times$ dietary condition interaction			
L hippocampus	-17, 12, -2	5.37	3,025
L medial prefrontal cortex	-2, 44, 17	4.80	1,332
R ventral prefrontal cortex	6, 38, 20	3.14	
L dorsal inferotemporal cortex	-28, 6, 8	4.52	1,842
R hippocampus	14, 13, -1	3.85	1,827
L agranular temporal pole	-12, 24, 4	3.84	847
L medial belt of auditory cortex	-21, 20, 5	3.06	
R striatum	7, 31, 14	3.51	3,602
R agranular insula	14, 28, 5	3.37	
R orbital prefrontal cortex	1, 26, 10	3.29	
L primary visual cortex	-11, -20, 30	3.47	320
L ventral prefrontal cortex	-24, 32, 10	3.44	393

The relationship between an index of insulin sensitivity, S<sub>I</sub>, and GM volume across subjects (S<sub>I</sub> association) and between the control and CR groups (S $_{\rm I}$   $\times$  dietary condition interaction). For a given voxelwise analysis, a given term was regressed onto each GM voxel in the brain. The voxel threshold for statistical significance was P < 0.005. Type 1 error was minimized when considering groups of contiguous, statistically significant voxels called clusters in different brain regions. Specifically, using Monte Carlo simulations, clusters had to exceed a number of voxels (cluster size) such that clusters generated from the data had a P < 0.05 of occurring by chance. For larger clusters, results are displayed for the maximally significant voxel followed by voxels from nearby regions. Coordinates refer to the sagittal, coronal, and axial planes of the Saleem-Logothetis atlas for rhesus macaques. L, left; R, right.

proinflammatory cytokines, and fewer prothrombotic factors due to reduced insulin resistance (43). S<sub>I</sub> represents the capacity for insulin to facilitate glucose uptake and inhibit hepatic production. Lower SI would correspond

to a higher insulin secretion peak followed by prolonged hyperinsulinemia. In this vein, Burns and coworkers (3) found that a higher insulin area under the curve during a glucose tolerance test predicted more hippocampal GM in early AD without a history of type 2 diabetes, but not in nondemented participants. This compensatory response may reflect glucoregulatory and metabolic dysfunction that can occur in early AD (2,6). Similarly, mediational models suggest that control animals with high basal insulin, but relatively lower basal glucose levels, would benefit from reduced insulin sensitivity in areas that rely on insulin signaling for optimal neural function. Previous studies have shown that infusion of insulin or antagonism via streptozotocin affects adenosine 5'-triphosphate storage in hippocampus (4), dopamine metabolism in striatum (44), and 2-deoxy-D-[<sup>14</sup>C]glucose utilization in rat medial temporal lobe, striatum, and frontal cortex (18). All of these brain regions were implicated in the current study. Although this compensatory mechanism for rhesus monkey controls may benefit these areas, hyperinsulinemia and lower insulin sensitivity is harmful to skeletal muscle (45) and may also cause vascular damage (42).

CR monkeys showed the expected relationship between higher  $S_I$  and more GM in the  $S_I \times$  dietary condition interaction. This positive correlation was similar to CR results for the S<sub>I</sub> association analysis. We have previously established that CR in rhesus monkeys reduced basal glucose and insulin, lowered insulin resistance, and increased lean muscle intracellular receptor substrate-1 expression (14,46). Such effects for glucoregulatory indices were also found in the current sample. Either intermittent fasting or CR also lowers serum glucose and insulin, as well as protects hippocampal neurons from excitotoxic injury induced by kainite (47).

Finally, hippocampal GM volume adjusted by  $S_I$  was significantly correlated with learning and memory performance for the mMAP task, whereas there was no relationship with unadjusted volume. This analysis was independent from voxel-wise analyses to minimize circularity concerns (23). Both approaches are complementary and can elucidate potential relationships among

#### TABLE 3

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Significant predictors	of SI $\times$	dietary	condition	interaction	with	regional	$\mathbf{G}\mathbf{M}$

Dietary condition	Variable	В	SE b	β	t
	Constant	1.889	0.412		
	HOMA-IR	-0.013	0.004	-0.526*	-3.097
	DI	-3.515E-5	1.069E-5	$-0.286^{+}$	-3.284
Control	$I_{\rm b}$	0.402	0.067	0.877‡	5.990
	Ğ	-1.780	0.248	-0.969 <sup>+</sup>	-7.185
	HbA <sub>1c</sub>	1.000	0.132	0.794‡	7.589
	Homocysteine	0.011	0.009	0.198	1.194
	Constant	0.053	0.833		
	HOMA-IR	0.017	0.019	0.502	0.905
	DI	9.296E-6	9.018E-6	0.229	1.031
Calorie restriction	I <sub>b</sub>	-0.169	0.178	-0.539	-0.539
	G <sub>b</sub>	0.125	0.492	0.064	0.064
	HbA <sub>1c</sub>	-0.182	0.282	-0.150	-0.150
	Homocysteine	-0.001	0.005	-0.037	-0.156

Multiple regression was used to assess which glucoregulatory variables from a frequently sampled glucose tolerance test mediated the  $S_I \times$ dietary condition interaction effect. Specifically, using data from a representative voxel in left anterior hippocampus, glucoregulatory measures were regressed onto the predicted change in GM due to insulin sensitivity for control or CR monkeys. For the final control model, the adjusted  $R^2$  of the mediation model was 0.889 [F(4,15) = 25.137; P < 0.001]. For the final CR mediational model, the adjusted  $R^2$  was 0.142 [F(5,25) = 0.378; NS]. DI, Disposition Index;  $G_b$ , basal glucose; HbA<sub>1c</sub>, glycosylated hemoglobin;  $I_b$ , basal insulin. \* $P \le 0.05$ . †P < 0.01.  $\pm P \le 0.001$ .



FIG. 2. Partial regression plots depicting error variance of the  $S_I \times$  dietary condition interaction explained by the HOMA-IR, basal insulin, and basal glucose. Control (n = 16) and CR (n = 26) monkeys are, respectively, represented by red triangles and blue circles. The green arrows indicate a representative control subject across parameters. Among controls, increases in HOMA-IR and basal glucose predicted less GM, whereas higher insulin levels were associated with more GM. These relationships were not significant among CR monkeys. GM volume is depicted in arbitrary units (A.U.).

glucoregulatory dysfunction, brain, and behavior. As  $S_I$  increases during CR, there might be less hippocampal atrophy over time that could positively influence task learning and performance. A comparable increase in insulin sensitivity among controls, by contrast, may be detrimental to hippocampus and could negatively affect mMAP performance.

There are several limitations that should be noted. It is not yet clear if improved peripheral insulin signaling in primates reflects similar processes in the brain. It must also be established if the relationship between insulin sensitivity and GM primarily reflects glucose uptake dynamics or other functions of insulin. For controls, the discrepancy between HOMA-IR and S<sub>I</sub> findings for GM warrant caution in interpretation, although several measures of insulin sensitivity produced the same result. Given that CR monkeys have less age-related morbidity and mortality (15), there may also be a survivor bias that influenced the voxel-wise results. This bias may make current results more conservative, however, given that surviving controls are likely more resilient to age-related pathophysiology. Regarding mediational models, glucoregulatory variables are sometimes multicollinear and may affect the interpretation of coefficients but not the overall model. Although basal insulin and HOMA-IR were very highly correlated (data not shown), this relationship is sensible both physiologically and statistically. Finally, the existing data are cross-sectional, and causation cannot be inferred.

In summary, increased  $S_I$  among CR monkeys was associated with more GM in parietofrontal cortices, hippocampus, and other regions that vary in insulin receptor density and signaling. Among controls, higher  $S_I$  showed a positive relationship with GM volume in parietofrontal cortices with low insulin receptor density, but predicted less GM in structures and areas that have high receptor density. CR or CR mimetics may benefit some specific brain regions and aspects of task learning and performance. Nevertheless, additional studies are needed to validate and

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clarify the association between glucoregulatory dysfunction and GM volume.

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