Fructose has been implicated in the pathogenesis of obesity and type 2 diabetes. In contrast to glucose, CNS delivery of fructose in rodents promotes feeding behavior. However, because circulating plasma fructose levels are exceedingly low, it remains unclear to what extent fructose crosses the blood-brain barrier to exert CNS effects. To determine whether fructose can be endogenously generated from glucose via the polyol pathway (glucose → sorbitol → fructose) in human brain, 8 healthy subjects (4 women/4 men; age, 28.8 ± 6.2 years; BMI, 23.4 ± 2.6; HbA1C, 4.9% ± 0.2%) underwent $^1$H magnetic resonance spectroscopy scanning to measure intracerebral glucose and fructose levels during a 4-hour hyperglycemic clamp (plasma glucose, 220 mg/dl). Using mixed-effects regression model analysis, intracerebral glucose rose significantly over time and differed from baseline at 20 to 230 minutes. Intracerebral fructose levels also rose over time, differing from baseline at 30 to 230 minutes. The changes in intracerebral fructose were related to changes in intracerebral glucose but not to plasma fructose levels. Our findings suggest that the polyol pathway contributes to endogenous CNS production of fructose and that the effects of fructose in the CNS may extend beyond its direct dietary consumption.
The human brain produces fructose from glucose

Janice J. Hwang,1 Lihong Jiang,2 Muhammad Hamza,1 Feng Dai,3 Renata Belfort-DeAguiar,1 Gary Cline,1 Douglas L. Rothman,2 Graeme Mason,2 and Robert S. Sherwin1

1Section of Endocrinology and 2Department of Radiology & Biomedical Imaging, Yale School of Medicine, New Haven, Connecticut, USA. 3Yale Center for Analytical Sciences, Yale School of Public Health, New Haven, Connecticut, USA.

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Introduction

Fructose has generated much attention as a potential driver of the US obesity and diabetes epidemics (1). In the brain, fructose, in contrast to glucose, appears to promote feeding behavior in rodents (2), and in humans, acute ingestion of fructose or glucose causes different patterns of cerebral blood flow changes in brain regions that influence eating behavior (3, 4). Because glucose enters the brain via passage across the blood-brain barrier (BBB) (5, 6), it is generally assumed that the CNS effects of fructose are mediated by peripheral fructose crossing the BBB as well (7). However, because of its efficient metabolism by the liver, extrahepatic circulating levels of fructose are exceedingly low. Thus, the extent to which peripheral fructose can cross the BBB to generate its CNS effects remains uncertain.

In this study, we use magnetic resonance spectroscopy (MRS) scanning, which allows for noninvasive direct measurements of metabolite concentrations to test an alternate mechanism, namely that fructose is produced endogenously within the human brain from glucose via the polyol pathway (glucose → sorbitol → fructose). The polyol pathway is an accessory glucose pathway that bypasses glycolysis (8, 9), and in the setting of hyperglycemia, up to 30% of glucose can be metabolized via this alternate pathway (10, 11). The polyol pathway is present throughout the human body, including the brain (12, 13), and it is responsible for the 20-fold higher levels of fructose observed in cerebrospinal fluid compared with blood (14).

In addition to fructose generation, flux through the polyol pathway also generates oxidative stress via production of NADH as well as because aldose reductase competes with glutathione reductase for NADPH, resulting in decreased glutathione (15). It is noteworthy that, in the periphery, this excessive oxidative stress has been implicated in the development of peripheral neuropathy (16), diabetic retinopathy and cataracts (17–20), nephropathy (21), fatty liver (22), and macrovascular disease (23) as well as platelet dysfunction (24, 25). In the brain, however, its role remains unclear. Thus, the current proof-of-concept study examines whether peripheral hyperglycemia can drive intracerebral production of fructose via the polyol pathway in lean, healthy individuals.
Results
Baseline fasting plasma glucose was 4.87 ± 0.15 mmol/l and rapidly increased to approximately 12 mmol/l during the hyperglycemic clamp (P < 0.0001) (Figure 1A). Fasting levels of plasma fructose and sorbitol were much lower at baseline (0.008 ± 0.001 mmol/l and 0.008 ± 0.003 mmol/l, respectively), and plasma fructose levels rose slightly (peak 0.0145 mmol/l at 180 min) (Figure 1B). No differences in plasma sorbitol levels were detected throughout (Figure 1B).

Baseline spectra were subtracted from spectra obtained every 10 minutes during the course of the glucose infusion in order to generate the difference spectra (Figure 2A), which were then compared with the reference spectra for glucose, fructose, and glucose and fructose obtained under identical scanning parameters (Figure 2B). Compared with baseline, intracerebral glucose levels rose significantly over time, beginning at 10 minutes (P = 0.0004) through the end of the study (P < 0.0001). Moreover, brain fructose levels also rose significantly compared with baseline, beginning at 20 minutes (P = 0.007) and extending through the end of the study (P < 0.01) (Figure 3A). Each of the 8 subjects had increases in intracerebral glucose (Figure 3B) and intracerebral fructose (Figure 3C). The magnitude of the change in intracerebral fructose concentration (obtained from times 60–180 minutes) was about 30%–35% of the change in intracerebral glucose (Figure 3A). Furthermore, the change in brain fructose levels was associated with the change in brain glucose levels (β coefficient = 0.18, SEM 0.03, P < 0.0001) but not plasma fructose levels (P = 0.36).

Discussion
The current data demonstrate that fructose is rapidly generated in the human brain during hyperglycemia. Our findings are consistent with those of a prior study in rats using 19F-MRS, which showed that, following i.v. 3-fluoro-3-deoxy-D-glucose infusion, there was a rise in intracerebral 3-fluoro-3-deoxy-D-sorbitol and 3-fluoro-3-deoxy-D-fructose. Furthermore, the production of sorbitol and fructose was blocked by sorbinil, an inhibitor of the rate-limiting enzyme, aldose reductase (26). Together with the previous finding that fructose and sorbitol are present in human cerebrospinal fluid at markedly higher concentrations than in the periphery (12, 14, 27), the present observations suggest that endogenous human brain fructose production may be a key underappreciated factor modulating ingested sugar’s effect on the brain. While it is possible that plasma fructose may cross the BBB, we observed that the rise of intracerebral fructose occurred much earlier (by 20 minutes) than the modest increase in peripheral fructose levels (at 180 minutes). Furthermore, in rats, a fructose infusion raising plasma fructose nearly 140-fold only increased brain fructose concentrations by 2-fold (3). Taken together, it is highly unlikely that plasma fructose crossing into the brain significantly contributed to the rising brain fructose levels.

In the present study, the change in brain fructose was about 30% of the change in brain glucose (~0.5 mmol/l for fructose, 1.5 mmol/l for glucose). When considered with our recent finding that plasma fructose levels are only 0.1% of plasma glucose levels, compared with CSF in which fructose is 6%–7% of glucose (28), these findings suggest that relative changes in brain fructose may be greater than relative changes in brain glucose levels, potentially making fructose a highly sensitive neuronal signal for carbohydrate ingestion. Interestingly, in Drosophila, any nutritious carbohydrate ingestion results in increased hemolymph fructose levels (through the polyol pathway), which in turn acts on a nutrient sensor in the fruit fly.
Whether fructose plays a similar role in modulating nutrient sensing in the mammalian or human brain requires further investigation. This proof-of-concept study has some limitations. First, using $^1$H MRS scanning, we were only able to measure changes in concentration and not absolute concentrations of fructose in the human brain. In addition, because the chemical spectrum for sorbitol overlaps completely with the spectra for glucose and fructose, we were unable to determine changes in sorbitol levels, which may have contributed to the difference spectrum intensity. Furthermore, in order to maximize sensitivity of signal detection, we scanned the occipital lobe and thus were not able to determine whether known regional variability in glucose utilization translates to differences in polyol pathway activity. Finally, our study design assumes that insulin does not regulate the polyol pathway. While animal studies have suggested that the polyol pathway activity is purely substrate dependent and not insulin dependent (30, 31), we did not directly control for the effects of insulin (or other hormones and metabolites, such as lactate; ref. 32) that conceivably might have indirect effects on the polyol pathway.

In light of the current obesity and diabetes epidemics, dietary fructose has garnered attention as a potential driver of these epidemics in part because of the high prevalence of high-fructose sweeteners (typically fructose/glucose ratios are 55:45 to 60:40) in the US diet. However, fructose is almost never consumed in isolation, and our finding that glucose levels drive brain fructose levels suggests that any sugar consumption that raises circulating glucose may in large part mediate fructose’s effects in the brain. Why the human brain produces fructose at such high concentrations compared with the plasma remains unclear. In humans, Glut 5, the principal fructose transporter (33), is present predominantly on microglial cells (34, 35), raising the possibility that endogenous fructose production in the brain may alter neuronal and glial interactions.

A growing body of evidence suggests that chronic hyperglycemia leads to many adverse effects on brain function, particularly neurovascular disorders and cognitive impairment (36, 37). However, the underlying mechanisms behind these associations remain unclear. The current data demonstrating that fructose is produced in the human brain in response to hyperglycemia via the polyol pathway activity as well as evidence for the presence of aldose reductase, the rate-limiting enzyme in the polyol pathway, throughout the brain (29). Whether fructose plays a similar role in modulating nutrient sensing in the mammalian or human brain requires further investigation.

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### Figure 2. Intracerebral fructose and glucose levels.

(A) Difference spectrum procedure showing for data from 1 subject. Blue spectra were obtained at baseline; red spectra were obtained at 10 minutes; black spectra is the difference between the red and blue spectra. (B) In vivo difference spectrum (final difference spectra) from 1 subject compared with reference spectra for fructose plus glucose, pure fructose, and pure glucose obtained under in vivo conditions of temperature 37°C, pH 7.4, and ionic strength of 150 mM. The fructose and glucose solution spectra were added, with their relative intensities adjusted to match their relative concentrations measured in vivo. The spectral regions plotted from the subject have been extended to include the region from ~3 to ~7 ppm where only noise is present in order to show the signal-to-noise ratio of the fructose peak. The arrow denotes the unique fructose peak at 4.0–4.1 ppm used to distinguish fructose from glucose.
human brain, particularly in the cerebral cortex, basal ganglia, and hippocampus (11), may have particular implications for disordered eating behavior and disordered cognition in patients with diabetes.

**Methods**

**Participants.** Eight healthy participants with normal weight were recruited with flyers and advertisements from the greater New Haven area (Table 1). Exclusion criteria included contraindications to entering an MRI, medication use (other than contraceptives for women), smoking history, illicit drug, or recent steroid use. Women who were breastfeeding, seeking pregnancy, or shown to be pregnant by urine test were also excluded.

**Experimental protocol.** Participants arrived at 7 AM following an overnight fast. One i.v. catheter was inserted into a distal arm for blood sampling. A second i.v. was inserted into the contralateral arm for 20% dextrose infusion that was adjusted every 5–10 minutes to achieve and maintain plasma glucose at 12 mmol/l. Additional blood samples were obtained for measurement of plasma sorbitol and fructose at baseline and 30, 60, 120, 180, and 240 minutes during hyperglycemia. The primary endpoint was change in intracerebral glucose and fructose levels. Secondary endpoints included changes in systemic sorbitol and fructose levels.

**1H MRS scanning.** Participants were positioned supine in a 4.0 T whole-body magnet interfaced to a Bruker AVANCE spectrometer (Bruker Instruments), with the head immobilized with foam inserts on top of a radiofrequency probe with a 1H circular coil as previously described (6, 38). After tuning, calibration, and acquisition of scout images for anatomical localization, intracerebral glucose and fructose concentration signals were obtained using stimulated echo acquisition mode localization (39) in 30 × 20 × 30 mm³ voxels in the occipital lobe for 20 minutes at baseline and then every 10 minutes over 4 hours of hyperglycemia. The sequence parameters were TR = 2,000 ms; TE/TM = 15 ms/10 ms; bandwidth = 5,000 Hz; sampling points = 2,048. Spectra were acquired with B0-lock and retrospective frequency adjustment for motion correction.

Although there is some overlap in their chemical spectra, fructose has unique peaks located at 4.0–4.1 ppm, which were used to distinguish it from glucose (Human Metabolome Database v3.6; refs. 40, 41). Baseline spectra were subtracted to eliminate overlap from other brain metabolites not directly affected by glucose levels. Changes in metabolite levels were measured by peak integration referenced to total voxel water signal and creatine, in the nondifferenced spectra, to determine concentration. Glucose integrals were obtained by integrating the large peak from 3.32 to 3.54 ppm and scaled back total intensity from 3.10 to 3.96 ppm in the free glucose spectrum (1:2.9), which contains 11 protons. Fructose integrals were obtained by integrating the peak from 3.96 to 4.07 ppm and scaled back total intensity from 3.54 to 4.20 ppm from the free fructose spectrum (1:4.8), which contains 12 protons. The concentrations of glucose and fructose were then compared with the creatine CH3 peak in the in vivo spectrum, which was assumed to be 10 mmol/kg and which contains 3 protons. Reference solutions of glucose (100 mM), sorbitol (100 mM), and fructose (100 mM) were

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**Table 1. Participant characteristics**

| Characteristic  | Mean (SD) |
|----------------|-----------|
| Age (yr)       | 28.8 (6.2) |
| Sex (male/female) | 4/4        |
| BMI (kg/m²)    | 23.4 (2.6) |
| HbA1C (%)      | 4.9 (0.2)  |
prepared in 100 mM phosphate buffer (pH 7.4) and measured after preheating the reference solutions to 37°C. Spectra were acquired using the same pulse sequence and parameters as used in vivo. Sorbitol peaks overlapped with glucose and fructose. While the peak pattern in the difference spectra was most consistent with glucose and fructose, we cannot rule out some degree of sorbitol contribution.

**Laboratory analysis.** Plasma glucose levels were measured via glucose oxidase (YSI Inc.). Plasma sorbitol and fructose were measured using gas chromatography–tandem mass spectrometry.

**Statistics.** The analyses of every repeatedly measured variable, including intracerebral glucose and fructose and plasma metabolite levels, were performed using the mixed-effects regression model method, taking in account within-subject correlation of repeated measures using a prespecified compound symmetry covariance matrix. Age, gender, and BMI were adjusted as covariates (i.e., as fixed effects). The association of intracerebral fructose with intracerebral glucose levels was also modeled using the mixed-effects regression model, in which the glucose level was included as a covariate in addition to age, gender, and BMI. The association of intracerebral fructose level with plasma fructose level was modeled in a similar manner. All the analyses were performed using SAS, version 9.4. A 2-sided \( P \) value of less than 0.05 was considered to be statistically significant.

**Study approval.** The Yale University Human Investigation Committee (New Haven, Connecticut, USA) approved the protocol, and all participants provided written informed consent prior to study participation.

**Author contributions**
JJH, LJ, and RSS had full access to all of the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. JJH, LJ, DLR, GM, and RSS conceived of and designed the study. JJH, LJ, MH, RBD, and GC acquired data. JJH, LJ, MH, FD, RBD, DLR, GM, and RSS analyzed and interpreted data. JJH, LJ, MH, RBD, GC, DLR, GM, and RSS wrote the manuscript. JJH, LJ, and FD performed statistical analysis.

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Address correspondence to: Robert Sherwin or Janice Hwang, The Anlyan Center, TAC 141S, PO Box 208020, New Haven, Connecticut 06520, USA. Phone: 203.785.6222; Email: Robert.Sherwin@yale.edu (R.S. Sherwin); Janice.Hwang@yale.edu (J.J. Hwang).

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