The continuous delivery of glucose to the brain is critically important to the maintenance of normal metabolic function. However, elucidation of the hormonal regulation of in vivo cerebral glucose metabolism in humans has been limited by the lack of direct, noninvasive methods with which to measure brain glucose. In this study, we sought to directly examine the effect of insulin on glucose concentrations and rates of glucose transport/metabolism in human brain using $^1$H-magnetic resonance spectroscopy at 4 Tesla. Seven subjects participated in paired hyperglycemic (16.3 ± 0.3 mmol/l) clamp studies performed with and without insulin. Brain glucose remained constant throughout (5.3 ± 0.3 μmol/g wet wt when serum insulin = 16 ± 7 pmol/l vs. 5.5 ± 0.3 μmol/g wet wt when serum insulin = 668 ± 81 pmol/l, P = NS). Glucose concentrations in gray matter–rich occipital cortex and white matter–rich periventricular tissue were then simultaneously measured in clamps, where plasma glucose ranged from 4.4 to 24.5 mmol/l and insulin was infused at 0.5 mU · kg$^{-1}$ · min$^{-1}$. The relationship between plasma and brain glucose was linear in both regions. Reversible Michaelis-Menten kinetics fit these data best, and no differences were found in the kinetic constants calculated for each region. These data support the hypothesis that the majority of cerebral glucose uptake/metabolism is an insulin-independent process in humans. *Diabetes* 50:2203–2209, 2001

The brain relies on the continuous delivery of glucose via the blood to maintain normal metabolic function. How glucose delivery into the central nervous system is regulated and how that delivery is altered by different metabolic conditions in the living human have been difficult to ascertain. In particular, the role of insulin in the regulation of cerebral glucose metabolism has been difficult to directly assess. Although evidence acquired using both in vitro and in vivo approaches support and refute the hypothesis that insulin regulates the entry of glucose into brain tissue (1–9), studies performed in living animals have been limited by their inability to directly measure cerebral glucose concentrations.

The brain is composed of both gray and white matter. Both rely on glucose for the maintenance of normal function, but the rates at which they metabolize glucose have been found to be different (9,10). Whether these differences in glucose metabolism equate to differences in cerebral glucose concentrations and whether insulin differentially regulates regional cerebral glucose metabolism have not been directly examined in humans because, until recently, direct methods to measure brain glucose concentrations in healthy subjects have not been feasible. Invasive methods to measure intracerebral glucose concentrations, such as the use of an implanted equilibrium dialysis probe, have been reported in abstract form (11), but such a technique is not acceptable in the study of normal human physiology. Indirect methods to quantitate brain glucose concentrations, such as those based on measuring the difference between glucose concentrations in the arterial circulation and the venous effluent of the brain (12), have provided important insights into understanding the regulation of brain-glucose uptake but are limited by the model assumptions used for data analysis (13). Positron emission tomography (PET) has also been used to effectively measure regional rates of cerebral glucose metabolism (14), but this method is limited by frequent reliance on a non-native glucose analog, an inability to directly measure metabolite concentration, and difficulty separating signals derived from different chemical species. PET also has difficulty obtaining serial studies from the same subject due to the radiation risks posed by the isotope used in the study (14).

Our group has recently developed magnetic spectroscopic methods that allow us to directly and noninvasively measure brain glucose concentrations in living humans during periods of tightly controlled plasma glucose (15,16). We used these methods to examine the hormonal regulation of glucose concentration in both the gray and white matter tissue of healthy humans. Our studies were designed to test the hypothesis that cerebral glucose transport/metabolism is largely an insulin-independent process. We also sought to determine whether gray and white matter glucose concentrations and metabolic rates differ under steady-state conditions.

RESEARCH DESIGN AND METHODS

Healthy subjects were recruited from the students and staff at the University of Minnesota. Before participation, subjects provided informed consent as governed by the Institutional Review Board at the University of Minnesota. Magnetic resonance spectroscopy (MRS) was used to measure brain glucose

From the 1Department of Medicine, University of Minnesota Medical School, Minneapolis, Minnesota, and the 2Department of Radiology, University of Minnesota. Before participation, subjects provided informed consent as governed by the Institutional Review Board at the University of Minnesota. Magnetic resonance spectroscopy (MRS) was used to measure brain glucose.
INSULIN AND BRAIN GLUCOSE TRANSPORT

**MRS**

<table>
<thead>
<tr>
<th>Period I</th>
<th>Period II</th>
</tr>
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<tbody>
<tr>
<td>Insulin</td>
<td>Glucose</td>
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</table>

**Minutes**

**FIG. 1.** Protocol for experiment 1. Somatostatin infusion was begun at time zero and advanced to a rate of 0.16 μg·kg⁻¹·min⁻¹ for 30 min and then maintained throughout the study. Glucose infusion was begun at minute 30 to maintain a constant plasma glucose concentration of 16.7 mmol/l. **H-MRS** was performed from minute 60 to 90 (period I) for later calculation of brain glucose concentration in the absence of insulin. Insulin was begun at 2.0 μU·kg⁻¹·min⁻¹ at minute 90. **H-MRS** was performed from minute 150 to 189 (period II) for later calculation of brain glucose concentration in the presence of insulin.

Concentrations. During the spectroscopic studies, plasma concentrations of both glucose and insulin were controlled at a constant target level using the glucose/insulin clamp technique. We performed two sets of experiments to address our study hypothesis.

**Experiment 1.** Subjects came to the Center for Magnetic Resonance Research in the morning after an overnight fast. In preparation for the study, an intravenous catheter was placed retrograde into one foot for the acquisition of blood samples and two additional intravenous catheters were placed into the upper extremities for the delivery of somatostatin, insulin, and glucose. The leg used for blood sampling was wrapped in heated towels and hot packs to arterialize the venous blood (17). After obtaining baseline samples for glucose and ketones, a somatostatin infusion was begun and advanced to a rate of 0.16 μg·kg⁻¹·min⁻¹ for 30 min to suppress endogenous insulin secretion (18). Once the target somatostatin infusion rate had been achieved, glucose (50% dextrose) was infused as necessary to achieve and maintain the plasma glucose concentration at 16.7 mmol/l. Beginning with the initiation of the somatostatin infusion, blood samples were obtained every 5 min for determination of plasma glucose concentration (using a Beckman Autoanalyzer). Additional samples were obtained every 30 min for later determination of serum insulin concentrations by radioimmunoassay (19). At the end of the first 60 min of somatostatin infusion, subjects were placed into the 4 Tesla magnet for examination by **H-MRS** in the absence of insulin (period I). When the collection of the spectral data was complete, subjects were removed from the magnet, given an intravenous infusion of insulin at a rate of 2.0 μU·kg⁻¹·min⁻¹, and maintained at a plasma glucose concentration of 16.7 mmol/l. The insulin infusion rate was selected to bring subjects to a serum insulin concentration in the high physiological range. After 1 h, subjects were placed back into the magnet and spectral data were collected in the presence of insulin (period II). Figure 1 details the protocol followed for experiment 1.

**Experiment 2.** For these experiments, subjects also reported to the Center for Magnetic Resonance Research in the morning after an overnight fast. The protocol as detailed for experiment 1 was followed, except that at time 0, insulin was begun at a rate of 0.5 μU·kg⁻¹·min⁻¹. The insulin infusion rate was selected to bring subjects to a serum insulin concentration in the physiological range. After 45 min, subjects were either maintained at euglycemia or given a bolus injection of dextrose (50% in water) to bring them to target glycemia (5–30 mmol/l). Plasma glucose was then clamped at target glycemia by the infusion of dextrose (50% in water), and subjects were placed in the magnet for data acquisition. After an additional 45 min, some subjects were given an additional bolus injection of dextrose to bring them to a second hyperglycemic target.

MRS. Experiments were performed on a 4 Tesla Siemens/Oxford magnet interfaced to a Varian console as previously described (16,20). Subjects were placed supine on a bed above the surface coil and their heads were held in place by cushions. To minimize exposure to gradient noise, all subjects wore earplugs. A quadrature transmit/receive RF coil consisting of two 14-cm diameter single-turn coils was used (21). A 16–27 ml nominal volume of gray matter–rich occipital cortex was selected for study for both experiments 1 and 2. For experiment 2, an additional 16 ml nominal volume of periventricular white matter–rich tissue was also selected for study. Localization of the signals was based on T₂-weighted MDEFT (22) or FSE (23) imaging. Localized shimming was performed using FASTMAP (24), which has shown consistent 7- to 8-Hz linewidths for the water resonance at 4 Tesla. For the measurements, we used TE = 4–20 ms, TM = 33 ms, TR = 4.5 (25). Water suppression was accomplished by the application of four 25-ms Gaussian pulses (26) or by applying a series of RF pulses according to the VAPOR scheme (27). In either case, we verified that water suppression did not affect the signal by ≥5% outside ±0.4 ppm. Outer volume saturation was achieved in slices adjacent to the volume of interest using hyperbolic secant pulses with variable RF power (28). Free induction decays were averaged over a time period of ~1 min, stored separately in memory, and then corrected for small frequency changes and averaged over at least 10 min. In each case, the plasma glucose concentrations were maintained at steady state for at least 20 min before spectral data were acquired. For experiment 1, data were collected over a minimum of 10 min while glycemia was clamped at the target level. For experiment 2, data were acquired in alternating mode simultaneously from the occipital and the periventricular regions over a minimum of 20 min, during which glycemia was clamped at target level.

Localized proton nuclear magnetic resonance (NMR) signals, obtained by the methods described above, were zero filled and apodized with 2-Hz exponential linebroadening. Peak areas were quantified using peak-fitting software supplied by the spectrometer manufacturer as previously described (15,16). Quantification of the glucose peak at 5.23 ppm was performed as previously described and validated by comparison with measurements made using 13C-MRS in our laboratory (16). In brief, the area under the glucose peak at 5.23 ppm was calculated relative to the area under the creatine methyl resonance at 3.04 ppm. For gray matter–rich regions, the concentration of creatine was set to 10 μmol/g wet wt, based on cortical concentrations of 9.6 μmol/g wet wt (29) and on contributions of 1 μmol/g wet wt γ-aminobutyric acid and 1–2 μmol/g wet wt glutathione in this region of the brain (30). The creatine concentration in the white matter–rich voxels was calculated to that in the gray matter using the relative requirements in RF power as a means to correct for the small differences in RF coil sensitivity. The half-volume RF coil design we used (21) resulted in the 90° RF power setting, varying by 1–2 dB, which corresponds to a signal correction between 12 and 25%. The creatine concentration was 7.3 ± 0.9 μmol/g wet wt (mean ± SD) for the white matter–rich areas, which suggests that the majority of our voxel contained white matter according to Hetherington et al. (31); this conclusion was further supported by the approximately twofold higher choline to creatine ratio.

**Data analysis.** Data are means ± SE. To minimize the potential effects of intersubject variation and modifications to magnetic resonance methodology, all data acquired during experiment 1 were evaluated based on intrasubject paired comparison, using observations obtained during the same study session. The data acquired during experiment 2 were examined using the reversible Michaelis-Menten model as previously described (16). This model assumes that trans membrane glucose influences the affinity for cis membrane glucose transport (product inhibition). Differences between groups were detected using Student’s t test.

**RESULTS**

**Experiment 1.** Seven healthy subjects (four women and three men) were recruited for participation in our investigation. Their mean age was 30 ± 2 years, and their mean BMI was 21.8 ± 0.2 kg/m². During period I, when somatostatin and glucose were infused, the serum insulin concentration was 16 ± 7 pmol/l and the plasma glucose concentration was maintained at 16.5 ± 0.5 mmol/l. During period II, when somatostatin, glucose, and insulin were infused, serum insulin rose to 668 ± 81 pmol/l and plasma glucose remained stable at 16.0 ± 0.4 mmol/l.

The glucose signal at 5.23 ppm in the 1H spectrum was easily resolved under the hyperglycemic conditions of our study (Fig. 2). No difference was found between the brain glucose concentrations measured in the absence and presence of insulin. During period I, when insulin secretion was suppressed, brain glucose concentration was 5.3 ± 0.3 μmol/g wet wt, whereas during period II, when insulin was infused at 2.0 μU·kg⁻¹·min⁻¹, brain glucose was 5.5 ± 0.5 μmol/g wet wt (P = NS) (Fig. 3). The 95%
The confidence interval for the difference in brain glucose concentration measured during periods I and II was 0.1–0.9 μmol/g wet wt, and the standard deviation of this difference was 0.2 μmol/g wet wt. With this variability around the mean, our sample size of seven provided an 80% likelihood of identifying a difference of 0.3 μmol/g wet wt between the brain glucose concentration measured in the absence and presence of insulin and a 90% likelihood of detecting a difference >0.4 μmol/g wet wt.

Although the analysis was based on paired statistics and was therefore considered to be independent of interindividual differences, we sought to further evaluate our findings by generating in vivo difference spectra for each subject. These difference spectra were created by subtracting the ¹H spectrum acquired in the absence of insulin from that acquired in the presence of insulin. In every subject, the resulting spectrum was a flat line, as shown in Fig. 2, suggesting that any difference in brain glucose signal acquired under the two experimental conditions was below the noise level of the instrument.

**Experiment 2.** Twenty healthy subjects were recruited for participation in this experiment. Their mean age was 35 ± 2 years, and their mean BMI was 26.1 kg/m². Brain glucose concentrations were measured at serum glucose concentrations ranging from 4.4 to 24.5 mmol/l. Serum insulin concentrations averaged 108 ± 6 pmol/l during these experiments. Figure 4 shows representative spectra acquired simultaneously from both gray and white matter-rich regions in a single subject.

The relationship between plasma glucose concentration and brain glucose concentration in both the gray and white matter-rich areas was observed to be linear (Fig. 5).
Reversible Michaelis-Menten kinetics fit these data best, as we previously found in experiments in which serum insulin concentrations were suppressed by the infusion of somatostatin (16). The kinetics constants calculated for the occipital cortex from these experiments with insulin were indistinguishable from the constants calculated from the data previously obtained during the experiments without insulin (Table 1).

In 10 experiments, data were simultaneously acquired from both the gray and white matter–rich regions during stable glycemia. As shown in Table 2, the values calculated for the white matter–rich periventricular tissue were significantly lower than the values calculated for the gray matter–rich occipital cortex ($P < 0.01$). However, when the data were corrected for the small difference in water content between gray (0.83 ml/g) and white (0.75 ml/g) matter (32), the average ratio between gray and white matter glucose concentration was $1.04 \pm 0.04$, which is no longer significantly different from 1.0, as judged from fitting $y = ax$. No statistically significant differences were found between kinetic constants calculated for these separate regions of cerebral tissue (Table 1).

**DISCUSSION**

The purpose of these experiments was to determine whether insulin plays a significant role in the regulation of cerebral glucose transport/metabolism in healthy human subjects. To address this question, we used the novel technique of high-field $^1$H-MRS and, for the first time, directly and noninvasively examined the effect of this hormone on the concentration of native glucose in the living human brain. We found that the infusion of insulin was without significant effect on the in vivo glucose concentration measured in the gray matter–rich occipital cortex of normal human volunteers. We further observed that the glucose transport kinetics calculated from data acquired from the occipital cortex in the presence of insulin were not different from those calculated previously in the absence of insulin. Taken together, these findings support the hypothesis that cerebral glucose transport/metabolism are largely an insulin-independent process.

The role of insulin in the regulation of brain-glucose metabolism has long been an area of controversy. Although most investigators have concluded that insulin was without effect on glucose transport across the blood-brain barrier (3–6,9), Hertz et al. (1) reported that insulin increased glucose transport in humans and Namba et al. (8) made similar observations in rats. However, a recent re-evaluation by Knudsen et al. (33) of the data from Hertz et al. (1) suggested that insulin may not truly have an effect on glucose transport/metabolism in humans, a conclusion that is also supported by the report of Cranston et al. (7). Interestingly, both insulin receptors (34–36) and insulin-sensitive GLUT4 (37–41) have been found at the blood-brain barrier, and insulin has been shown to be an important modulator of the autonomic response to hypoglycemia (42) and of feeding behavior (43). Insulin crosses the blood-brain barrier via receptor-mediated transcytosis. Despite the obvious importance of the hormone in regulating some cerebral functions, our observations, particularly the finding that the in vivo difference spectrum created by subtracting the data collected in the presence of insulin from that collected in its absence is a flat line, provide support for the hypothesis that insulin does not alter glucose content or any other brain metabolites detectable by $^1$H-NMR spectroscopy. Hence, insulin ap-

| Table 1: Kinetic constants calculated using reversible Michaelis-Menten model |
|-----------------------------------|-----------------|-----------------|
| White matter–rich region          | $K_i$ (mmol/l)  | $T_{max}/CMR_{glc}$ |
| in the presence of insulin        | 1.96 ± 2.45    | 2.15 ± 0.25     |
| Gray matter–rich cortex           | 0.98 ± 2.13    | 2.24 ± 0.23     |
| in the presence of insulin        | 0.6 ± 2.0      | 2.3 ± 0.2       |
| Gray matter–rich cortex           |                |                 |
| in the absence of insulin*        |                |                 |

Data are means ± SE. *Data previously published (16).
pears to be without effect on glucose transport across the 

blood-brain barrier when metabolism remains unchanged.

In our first experiment, we used a paired comparison to 
determine whether insulin altered intracerebral glucose 
concentrations. With a sample size of seven, this study had 
sufficient power to detect a difference as small as 0.36 
\(\mu\)mol/g wet wt, with 90% likelihood. Such a value repre-
sents 7% of the mean brain glucose concentration mea-
sured during our experiments and is unlikely to be of 

major physiological significance. Thus, although a small 
effect of insulin on glucose transport across the blood-

brain barrier cannot be completely eliminated with our 
data, such as an effect on the slow glycogen metabolism in 

brain (44), we feel confident that this hormone does not play a clinically relevant role in the regulation of brain-

Glucose concentrations in the brain are achieved 

through a careful balance between glucose entering the 

brain across the blood-brain barrier and glucose leaving 

the brain through metabolism. For glucose concentration 
to remain constant under steady-state conditions, the rate 
of glucose influx must equal the rate of glucose efflux. 

Recent investigations using PET have demonstrated that 

insulin was without effect on the metabolic rate of glucose in 

the cerebral cortex of humans (7,9). These observations, 
coupled with the data presented in the present study, provide strong evidence that insulin is without effect on the kinetics of glucose transport across the blood-brain barrier. Future investigation into the role of insulin in the regulation of intracerebral-glucose metabol-

ism will therefore be of interest. Particularly intriguing are the observations of Nelson et al. (45), who reported that insulin administration before death was shown to increase the amount of glycogen present in the post-
mortem rat brain. However, the micromolar glycogen 

concentrations that have recently been measured in anes-

thetized rats in vivo (44) suggest that the contribution of 
glycogen to overall cerebral glucose metabolism must be small.

In the second experiment, we measured the kinetic 

constants for glucose transport across the blood-brain 

barrier for both gray and white matter–rich regions. Using 

the reversible Michaelis-Menten model, the resulting ki-
netic constants were not statistically significantly differ-
et, implying that insulin has similar effects on glucose uptake/metabolism in both gray and white matter glucose uptake/metabolism. Interestingly, we observed that the steady-state concentrations of the white matter–rich periventricular tissue were significantly lower than simulta-

neously measured concentrations in the gray matter–

rich occipital cortex. This quantification (\(\mu\)mol/g wet wt) was performed on a per gram of wet weight basis, and the ratio between gray and white matter was 1.4 ± 0.2. However, the results do not imply that a significant chemical glucose concentration gradient exists between white and gray matter. When expressing the glucose concentrations on a per milliliter of brain water basis, i.e., when correcting for the small difference in water content 

between gray (0.83 ml/g) and white (0.75 ml/g) matter (32), the average ratio between gray and white matter glucose concentration (\(\mu\)mol/ml brain water) was 1.1 ± 0.1, which is no longer significantly different from 1.0. There-


fore, we conclude that the chemical concentration gradi-
et between gray and white matter is not significant. 

Because glucose transport in the brain occurs by facili-
tated diffusion, this implies that there is no net glucose mass transfer between gray and white matter. Other investigators have consistently measured the metabolic rate of glucose in white matter to be significantly lower than that in gray matter (9,10), which implies that there are

<table>
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<th>Subject</th>
<th>Plasma glucose (mmol/l)</th>
<th>Gray matter glucose ((\mu)mol/g wet wt)</th>
<th>White matter glucose ((\mu)mol/g wet wt)</th>
<th>Gray matter glucose: white matter glucose ((\mu)mol/ml brain water)*</th>
<th>White matter glucose ((\mu)mol/ml brain water)†</th>
<th>Gray matter glucose: white matter glucose ((\mu)mol/ml brain water)†</th>
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Mean ± SE

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Data are means ± SE. *Assumed gray matter water content = 0.83 ml/g (31); †assumed white matter content = 0.75 ml/g (31).
marked differences in $T_{\text{max}}$ between these tissues. This difference can be considered to also reflect the density of the transporter at the blood-brain barrier (the permeability surface area product); therefore, it is related to the blood volume in gray and white matter. Thus, it appears that brain glucose content is heavily regulated in normal brain tissue, underscoring the important role it plays in normal brain function.

The study of in vivo cerebral glucose metabolism in humans has been limited by the inability to directly and noninvasively measure brain glucose concentrations. Techniques such as the indicator dilution method (12) and PET have provided important insights into the regulation of cerebral glucose transport and metabolism in humans but are by their very nature indirect and based on assumptions that are difficult to experimentally confirm. The microdialysis method allows direct measurement of extracellular brain glucose, which appears to be equal to intracellular glucose concentrations (46), but may become inaccurate over time because of the scarring that occurs in the tissue surrounding the probe. In addition, the invasiveness of the technique precludes its application to general investigation of healthy volunteers. $^{13}$C-MRS at low field has offered a solution to the limitations presented by these indirect or invasive techniques, but it requires the infusion of $^{13}$C-glucose to achieve sufficient signal-to-noise to allow quantification of brain glucose content (47). $^1$H-MRS does not require the infusion of substrate and uniquely allows for the direct study of native glucose in brain tissue. However, at lower field strengths, the resolution of the glucose signal in the $^1$H spectrum has been difficult. Most investigators working at lower field strengths have relied on a glucose signal of 3.43 ppm for quantification purposes (48–50). However, this peak overlaps with myo-inositol and does not easily allow for precise quantification (15,51).

The peak at 5.23 ppm is free of spectral overlap with other metabolic species (15) and can be well resolved for quantification using a field strength of 4 Tesla and the methods of shimming used in the current study.

One limitation of $^1$H-MRS is the difficulty that can be encountered in measuring brain glucose content under hypoglycemic conditions. However, if long acquisition times are used so that the signal-to-noise ratio can be optimized, data sufficient for accurate quantification of glucose can be acquired. Another limitation is that the glucose content measured in a region reflects the glucose present in both cerebral tissue and the intravascular space. However, since cerebral blood volume is minimal (on the order of 2–4% of total volume) (52–54), the contribution of blood glucose to the measurement of brain glucose is small and without effect on the calculation of glucose transport kinetics across the blood-brain barrier. At a plasma concentration of 15 mmol/L, the contribution of blood to the glucose concentration measured in gray matter would be $\sim 0.6$ mmol/L or $\sim 10\%$ of the total glucose concentration measured in a volume of cerebral tissue. When the data are adjusted for the contribution of intravascular glucose to the overall glucose concentration in the brain, the conclusions drawn from the data are unchanged. Consequently, $^1$H-MRS at high field strengths represents the best way to directly measure the concentrations of native glucose in the brain of human volun-

teers. This is supported by the observation of resolved glucose resonances in the rat brain at 9.4 Tesla (27).

In summary, our experiments demonstrate that $^1$H-MRS at 4 Tesla is a powerful investigative tool with which to directly and noninvasively measure the concentration of glucose in cerebral tissue. Using this technique, we demonstrated that the infusion of insulin during constant hyperglycemia was without effect on brain glucose concentrations and that the kinetics of glucose transport calculated from data acquired in the presence of insulin are not different from those calculated from data acquired in the absence of insulin. In addition, we found that the kinetic constants calculated for both gray and white matter–rich regions in the presence of insulin are similar, indicating that the hormone has similar effects on glucose uptake/metabolism in both types of cerebral tissue. All together, this study demonstrated the power that MRS brings to the study of in vivo metabolism and illustrated how the method can be used to uniquely and directly address questions important in human physiology and disease.

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