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The fasting polypeptide FGF21 can enter brain from blood

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Abstract

FGF21 recently has been proposed as a missing link in the biology of fasting, raising the question of whether it directly reaches the brain. We used multiple time-regression analysis to quantify the influx rate of this polypeptide across the blood-brain barrier (BBB), size-exclusion chromatography to examine degradation, capillary depletion to differentiate entry into brain parenchyma from retention in the microvasculature, and measurement of efflux rate to determine a possible confounding effect on measurement of entry. FGF21 was 94 % intact in serum and 75 % in brain 10 min after intravenous bolus delivery. Its influx rate was $0.23 \pm 0.12 \mu\text{l/g-min}$, nearly 4-times faster than that of the vascular marker albumin. At 10 min, about 0.5 % of the administered FGF21 was present in a gram of brain tissue. Of this, 70 % reached the parenchyma of the brain. Co-injection of excess FGF21 failed to inhibit the influx, showing a lack of saturation. Efflux, which occurred at the same rate as the bulk reabsorption of cerebrospinal fluid, also was not saturable. In summary, FGF21 shows significant, non-saturable, unidirectional influx across the BBB.

Keywords

FGF21; blood-brain barrier; metabolism

1. Introduction

The fibroblast growth factor (FGF) polypeptides were initially named by their ability to stimulate fibroblast proliferation. FGF21 activates glucose uptake of adipocytes, reduces plasma glucose and triglyceride concentrations in *ob/ob* and *db/db* mice, protects against diet-induced obesity when overexpressed in transgenic mice, and improves lipoprotein profiles in diabetic rhesus monkeys [10;11;19].

FGF21 recently was described as a missing link in the biology of fasting, important in regulating the adjustment to food deprivation [1;2;18]. The brain senses fasting with neurobehavioral changes, leading to various adaptations such as torpor, which is involved in the conservation of energy. Reitman posed the unanswered question of whether FGF21 reaches the brain directly [18].

Although it is now recognized that many peptides and polypeptides cross the blood-brain barrier (BBB), permeability varies with the peptide. Several peptides/polypeptides that have a molecular weight similar to that of FGF21, or even smaller, do not cross the BBB any faster

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than albumin. These include obestatin [14], adiponectin [14], orexin B [3], melanin-concentrating hormone [5], interleukin-10 [8], transforming growth factor- β [6], glial cell line-derived neurotrophic factor [7], and platelet-derived growth factors AA and BB [4]. The pharmacokinetics of the interactions of FGF21 with the BBB could have direct therapeutic implications for this recently discovered peptide. Therefore, we quantified the influx of intact FGF21 into the brain and its possible efflux after intracerebroventricular administration.

2. Materials and Methods

2.1. Animals and labeling

Male CD1 mice 5–6 weeks of age, used in accordance with a protocol approved by the Institutional Animal Care and Use Committee, were anesthetized by intraperitoneal injection of urethane. Recombinant human His-tagged full-length FGF21, containing 195 amino acids with a molecular weight of 20.996 kDa, was purchased from Phoenix Pharmaceuticals (Burlingame, CA). Bovine serum albumin (BSA) and the chloramine-T reagent used for radioactive labeling (radiolabeling) were purchased from Sigma (St. Louis, MO). Chloramine-T was used to radiolabel FGF21 with ^{125}I (Amersham, Piscataway, NJ), the reaction being stopped 1 min later with sodium metabisulfite. This material was purified on a Sephadex G-10 column. The specific activity of the ^{125}I -FGF21 was about 44 Ci/g.

2.2. Degradation

The supernatants from brain homogenates of 6 mice obtained 0, 10, and 20 min after injection of ^{125}I -FGF21 were pooled (2/sample) and run on size-exclusion chromatography. The time point 0 represents the processing control involving addition of ^{125}I -FGF21 to homogenate and blood to determine the extent of degradation occurring after collection. Each mouse in the experimental groups received an intravenous (iv) injection of 3 μCi of ^{125}I -FGF21 in 100 μl injected at time 0. At 10 and 20 min, arterial blood and brain were taken and processed on ice. Brain tissue was homogenized in 1 ml of lactated Ringer's solution containing 1% bovine serum albumin (LR/BSA) and Complete Protease Inhibitor Cocktail (Sigma). For size-exclusion chromatography, 100 μl of lyophilized brain supernatant was added to a 9 \times 6 cm column of polyacrylamide P10. Fractions of 100 μl were collected, as described previously [16].

2.3. Quantification of blood-to-brain influx

Multiple time-regression analysis was used as previously detailed [9;14]. ^{125}I -FGF21 and ^{131}I -albumin (about 30,000 cpm/ μl for each mouse) were delivered in the same injectate of 100 μl of freshly made LR/BSA. The injection was made into the isolated left jugular vein of anesthetized mice, each mouse representing a specific time point. At various times after iv injection, blood was collected by transection of the right carotid artery and the mice were decapitated immediately afterwards. Since degradation assays showed that substantial degradation had occurred at 20 min in the brain (see Results section), only the mice studied within 10 min after iv injection were used to calculate the influx rate from blood to the brain.

To test whether there was saturation of FGF21 permeation across the BBB, two groups of mice were studied simultaneously. The control group received only radioisotopes ($n = 7/\text{group}$), whereas the inhibition group received co-injection of 2 $\mu\text{g}/\text{mouse}$ of unlabeled FGF21 in addition to ^{125}I -FGF21 and ^{131}I -albumin ($n = 8/\text{group}$). At the end of the study, radioactivity in brain and serum was measured, and the brain/serum ratio of ^{125}I -FGF21 in each g of brain was measured. The serum values were also used in the calculation of exposure time, a theoretical steady-state value with correction for the decrease in blood concentration of ^{125}I -FGF21 after iv bolus injection. This represents the theoretical value correlated with each experimental time t as if the blood concentration of ^{125}I -FGF21 were constant from time 0 to

time t . It is obtained by division of the integral of serum radioactivity over time by the serum radioactivity at each experimental time [9]. The influx rate and volume of distribution of ^{125}I -FGF21 is determined from the linear regression correlation between the tissue/serum ratio of radioactivity and exposure time, as described previously [16]. The unidirectional influx rate K_i is the slope of the linear regression line, and the initial volume of distribution V_i is the intercept. Differences in the regression lines between groups were compared by the least square method with the GraphPad program.

2.4. Compartmental distribution by capillary depletion

Two groups of mice ($n = 4/\text{group}$) were used to determine whether the injected radioactive FGF21 actually reached brain tissue rather than remaining trapped in the cerebral vasculature. One group of mice was perfused intracardially to remove loosely adherent radiotracers in the brain vasculature whereas the other group did not receive intracardial perfusion for vascular washout. In both groups, each mouse received ^{125}I -FGF21 and ^{131}I -albumin in 100 μl of LR/BSA buffer by bolus injection into the left jugular vein. Ten min later, blood was collected by transection of the abdominal aorta. Either with or without vascular washout afterwards, the mouse was decapitated to obtain the cerebral cortex. The cortex was homogenized in 0.7 ml of capillary buffer and mixed thoroughly with 1.7 ml of 26 % dextran (about 67 kD). This mixture was centrifuged (5400 g, 15 min, 4 °C) with a swing bucket rotor to obtain the parenchymal and capillary fractions. After measurement with a dual-channel γ -counter, the ratios of tissue/serum radioactivity for ^{125}I -FGF21 were calculated. Group means are shown with their standard errors. Significant differences among the groups were determined by ANOVA followed by Tukey's post-hoc comparison test.

2.5. Quantification of brain-to-blood efflux

A slow rate of influx might reflect increased efflux. Therefore, one group of mice ($n = 9$) received only ^{125}I -FGF21 and ^{131}I -albumin (40,000 and 30,000 cpm, respectively) while a second group received 100-fold excess unlabeled FGF21 (40 ng/mouse) in addition to these two radioactively labeled compounds. A perfusion pump with a 5 μl Hamilton syringe connected to a PE10 cannula was used to deliver these solutions into the right lateral ventricle 1.0 mm lateral and 0.2 mm posterior to the bregma and 2.2 mm below the skull. The perfusion rate was 12 $\mu\text{l}/\text{min}$ and the duration of delivery was 5 sec; the cannula remained in place for another 5 sec before being slowly withdrawn. The mice were decapitated 2, 5, 10, 15, 20, 30, 45, and 60 min later. The control time 0 was determined after euthanasia of the mice by an overdose of urethane before injection, and the mice were decapitated immediately. The half-time disappearance of ^{125}I -FGF21 and ^{131}I -albumin was determined from the linear regression correlation between the brain radioactivity and time. Statistical significance between groups was determined with the Prism GraphPad program.

3. Results

3.1. Degradation of ^{125}I -FGF21

Size-exclusion chromatography confirmed that ^{125}I -FGF21 was mainly intact in both serum and brain 10 min after injection. After correction for degradation during ex-vivo processing determined from the controls, almost 100 % of the radioactivity in serum and 88 % in brain represented intact ^{125}I -FGF21. However, 20 min after iv delivery of ^{125}I -FGF21 there was substantial degradation in brain (41 % intact) and serum (83 % intact). These results are shown in Table 1 and figures 1A and 1B.

3.2. Blood-to-brain influx of ^{125}I -FGF21

The K_i of ^{125}I -FGF21 was $0.23 \pm 0.12 \mu\text{l/g}\cdot\text{min}$ and the initial volume of distribution was $13.04 \pm 1.19 \mu\text{l/g}$. This indicates a brain uptake of about 0.5 %/g brain at 10 min. With 1.36 μCi injected at a specific activity of 44 Ci/g, this shows that about 179 pg of ^{125}I -FGF21-1 entered one g of brain after a single bolus injection at this time. After addition of 2 μg /mouse of unlabeled FGF21 (about 65-fold excess), the resulting K_i ($0.36 \pm 0.15 \mu\text{l/g}\cdot\text{min}$) was not significantly different, indicating that the influx of FGF21 into brain was not inhibited at this dose (Fig. 2).

3.3. Uptake of ^{125}I -FGF21 by muscle and liver

Two peripheral organs, muscle and liver, were used as controls for the uptake of FGF21 into brain, an organ which is protected by the BBB consisting of less permeable microvessels reinforced by tight junctions. In the gluteus major muscle, ^{125}I -FGF21 had an influx rate 10-times higher than that in the brain. This is also significantly higher than that of albumin. Excess unlabeled FGF21 had no effect on the muscle uptake of either substance (Fig. 3A). Although FGF21 is produced by the liver, the influx of ^{125}I -FGF21 from blood to liver was even higher than that to the muscle, but this also did not show saturability in the presence of excess unlabeled FGF21 (Fig. 3B).

3.4. Compartmental distribution ^{125}I -FGF21 in the brain

Ten min after iv injection of ^{125}I -FGF21, $5.87 \pm 0.56 \mu\text{l/g}$ was present in the cortex. Of this total radioactivity, about 70 % ($4.14 \pm 0.56 \mu\text{l/g}$) reached the brain parenchyma. This was significantly higher than that retained in the capillaries ($0.40 \pm 0.06 \mu\text{l/g}$) or remaining loosely bound in the vessel lumen ($1.78 \pm 0.70 \mu\text{l/g}$). This reversible binding accounted for about 30 % of the total apparent brain uptake (Fig. 4).

3.5. Brain-to-blood efflux of ^{125}I -FGF21

The efflux rate of ^{125}I -FGF21 from brain was similar to that ^{131}I -albumin, their half-time disappearances being 17.6 and 17.4 min, respectively. These values were not significantly altered by inclusion of excess unlabeled FGF21 in the injectate (Fig. 5).

4. Discussion

FGF21 was originally isolated from mouse embryos. Northern blotting analysis showed that it had the highest expression in liver and lower levels in the thymus, but was undetectable in several other peripheral tissues. Human and mouse FGF21 have a high degree of homology (75 % of the amino acids) [13]. This justified the use of His-tagged human FGF21 in our mouse studies. FGF21 belongs to the FGF19 subfamily of metabolic regulators that also includes FGF15 and FGF23. They bind to the isoforms of FGFR1–4 that have intrinsic tyrosine kinase activity and regulate glucose and lipid homeostasis.

Overexpression (transgenic mice) and subcutaneous treatment of genetically obese and diabetic animals (ob/ob and db/db mice, Zucker diabetic fatty rats) can induce a reduction of blood glucose and triglyceride [10]. FGF21 and other members of the FGF19 family of ligands have a low level of binding to heparin/heparan sulfate. This enables them to circulate as hormones rather than being trapped by the extracellular matrix, but this also raises the necessity of cofactors for receptor activation, such as the transmembrane protein beta-Klotho [12]. Since FGF23 is produced by the brain, predominantly in the ventrolateral thalamus [20], it is not known whether FGF21, which is elevated by fasting, can cross the BBB to act on the FGF receptors in the brain. The permeability of the BBB to FGF21 can be best quantified by

multiple-time regression analysis [9]. This revealed slow but significant blood-to-brain penetration.

The results showed that intact FGF21 crossed the BBB at a rate comparable to that of polypeptides of similar size [15;17]. Size-gel chromatography showed that it crossed in intact form, and capillary depletion showed that it mainly entered the brain parenchyma rather than being trapped in the microvasculature that composes the BBB. We also ruled out the presence of an efflux transport system.

There was no saturation of the influx of FGF21 into liver or muscle. In doses of 125 or 750 µg/kg/d with once daily subcutaneous injection for a week, FGF21 administration does not cause mitogenicity or hypoglycemia, indicating that the influx to other organs does not induce metabolic dysfunction at large doses in rodents [10]. A similar lack of systemic toxicity for FGF21 has also been shown in studies in monkeys [11].

Thus, intact FGF21 crossed the BBB by simple diffusion. There was no saturable influx or efflux transport system. The small amount of FGF21 entering the CNS (about 0.5 %/g of brain tissue) could be physiologically important in fasting. To the question raised by the editorial asking whether FGF21 in the periphery can reach the brain directly [18], the answer is a definite yes.

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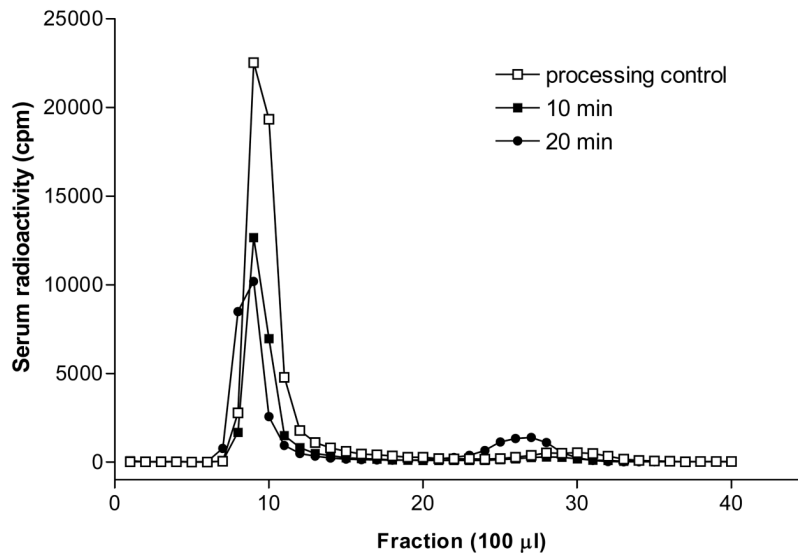


Fig. 1A

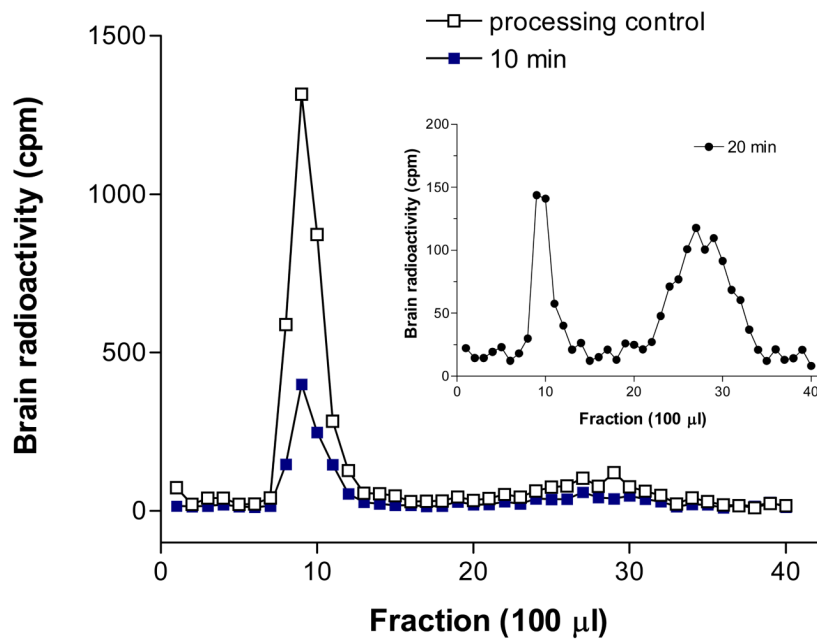


Fig. 1B

Fig. 1. Intact ^{125}I -FGF21 found by polyacrylamide size exclusion chromatography in samples of serum (1A) and brain (1B) 10 min after iv delivery. More degradation occurred at 20 min.

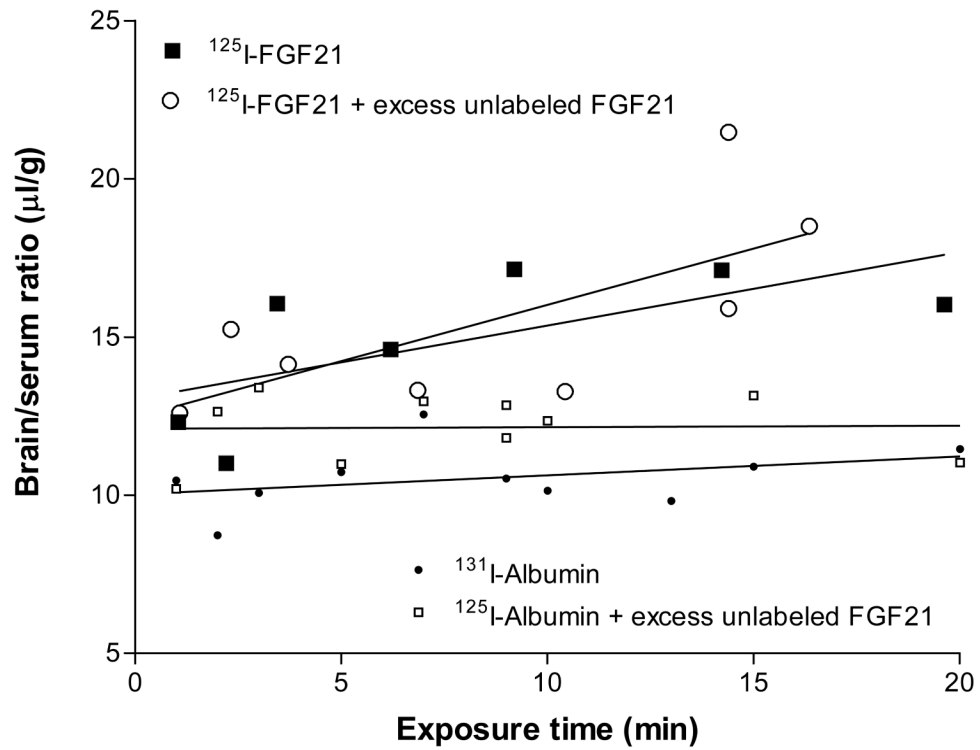


Fig. 2. The blood-to-brain influx rate of $^{125}\text{I-FGF21}$ after iv bolus injection was about 4-times higher than that of the vascular control, $^{131}\text{I-albumin}$. Excess unlabeled FGF21 did not affect the influx of either $^{125}\text{I-FGF21}$ or $^{131}\text{I-albumin}$.

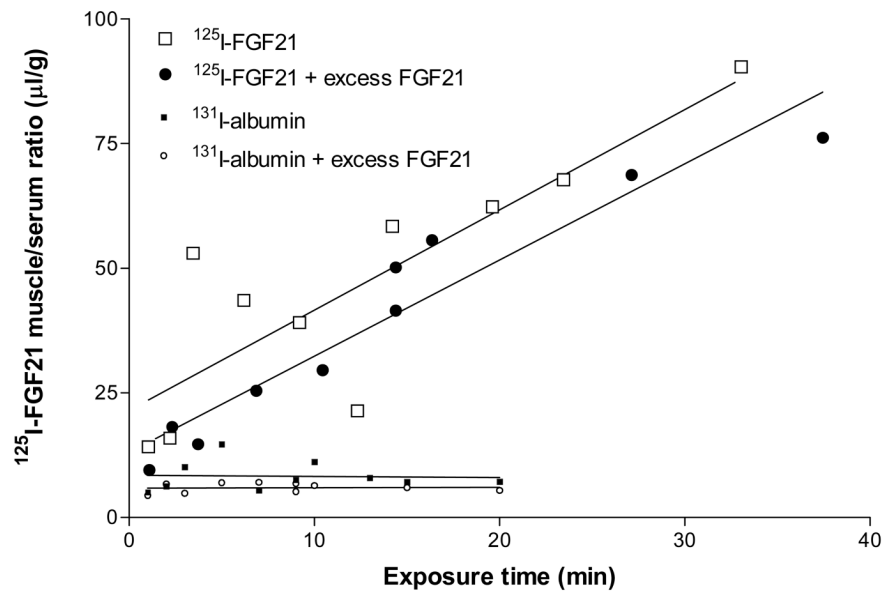


Fig. 3A

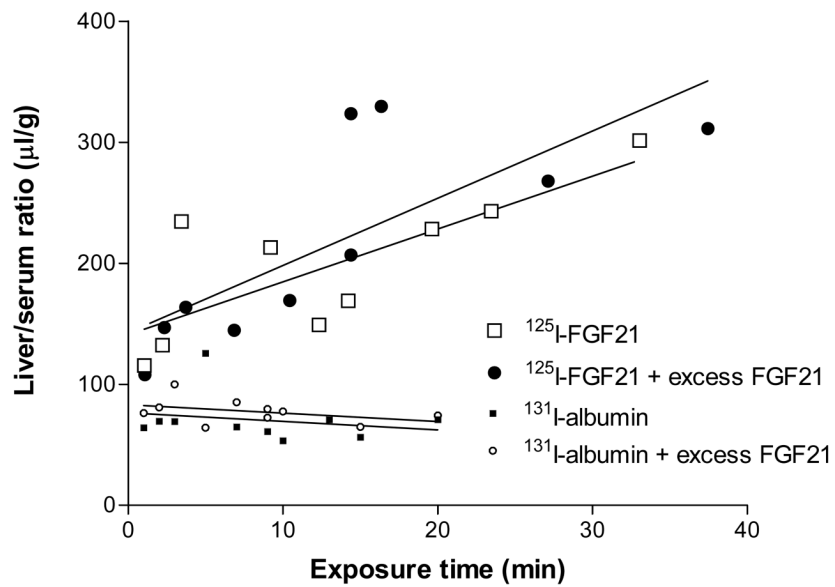


Fig. 3B

Fig. 3. The influx of $^{125}\text{I-FGF21}$ into muscle (3A) and liver (3B) was significantly higher than into brain and was also higher than that of $^{131}\text{I-albumin}$. There was no significant change caused by excess unlabeled FGF21. The influx rate of $^{125}\text{I-FGF21}$ into the liver was at least twice as fast as that into muscle.

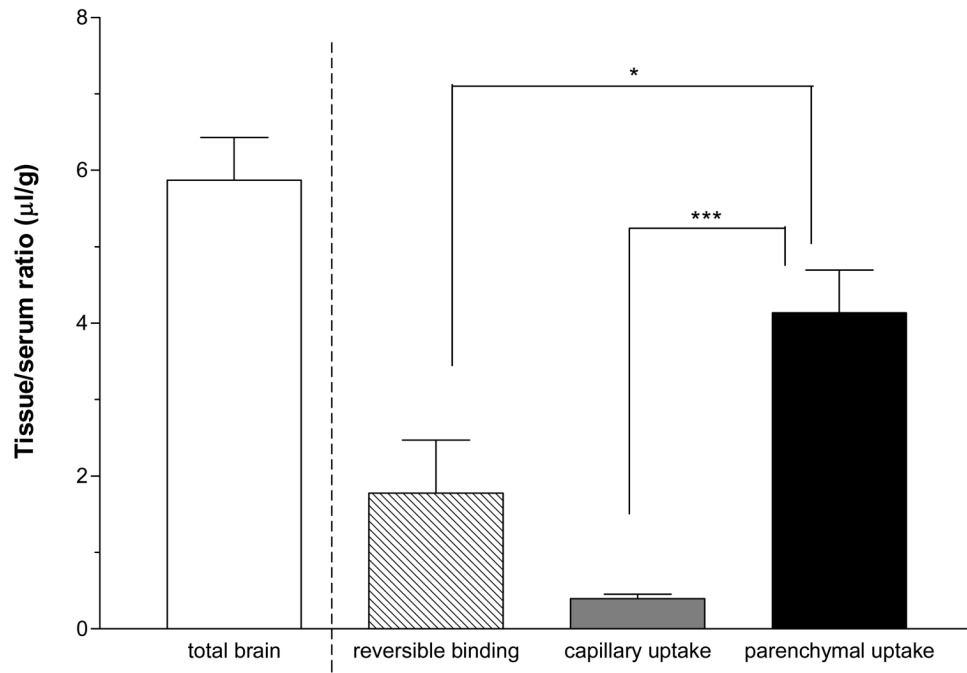


Fig. 4. Capillary depletion showed that most (70 %) of the ^{125}I -FGF21 in the total brain had reached the parenchyma 10 min after iv delivery. *: $p < 0.05$; **: $p < 0.01$.

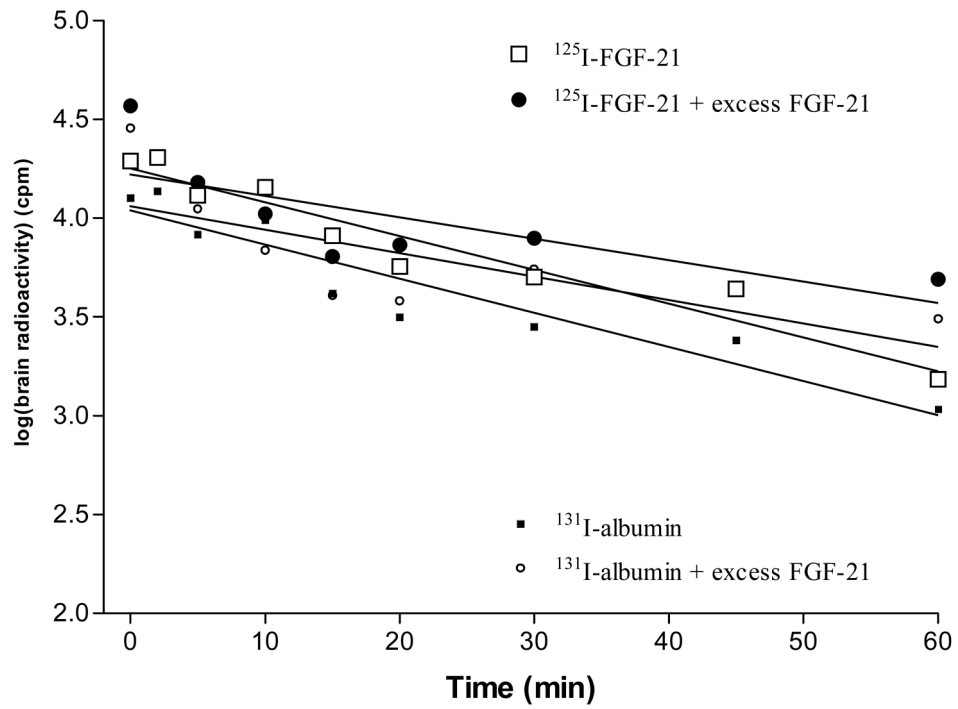


Fig. 5. The efflux of ^{125}I -FGF21 from brain was similar to that of ^{131}I -albumin. Excess unlabeled FGF21-1 showed no significant effect on the slow efflux of either substance.

Table 1¹²⁵I-FGF21 degradation assay after intravenous injection shown by size-exclusion chromatography

	Serum	Serum	Brain	Brain
	Raw data	corrected	Raw data	corrected
Control (0 min)	94.2%	100%	86.3%	100%
10 min	94.1%	99.9%	75.8%	87.8%
20 min	78.1%	82.9%	35.2%	40.7%