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## Transferrin Aptamers Increase the *In Vivo* Blood-Brain Barrier Targeting of Protein Spherical Nucleic Acids

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### Abstract

The systemic delivery of exogenous proteins to cells within the brain and central nervous system (CNS) is challenging due to the selective impermeability of the blood-brain barrier (BBB). Herein, we hypothesized that protein delivery to the brain could be improved *via* functionalization with DNA aptamers designed to bind transferrin (TfR) receptors present on the endothelial cells that line the BBB. Using  $\beta$ -Galactosidase ( $\beta$ -Gal) as a model protein, we synthesized protein spherical nucleic acids (ProSNAs) comprised of  $\beta$ -Gal decorated with TfR aptamers (Transferrin-ProSNAs). The TfR aptamer motif significantly increases the accumulation of  $\beta$ -Gal in brain tissue *in vivo* following intravenous injection over both the native protein and ProSNAs containing nontargeting DNA sequences. Furthermore, the widespread distribution of  $\beta$ -Gal throughout the brain is only observed for Transferrin-ProSNAs. Together, this work shows that the SNA architecture can be used to selectively deliver protein cargo to the brain and CNS if the appropriate aptamer sequence is employed as the DNA shell. Moreover, this highlights the importance of DNA sequence design and provides a potential new avenue for designing highly targeted protein delivery systems by combining the power of DNA aptamers together with the SNA platform.

### Graphical Abstract

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Author Contributions

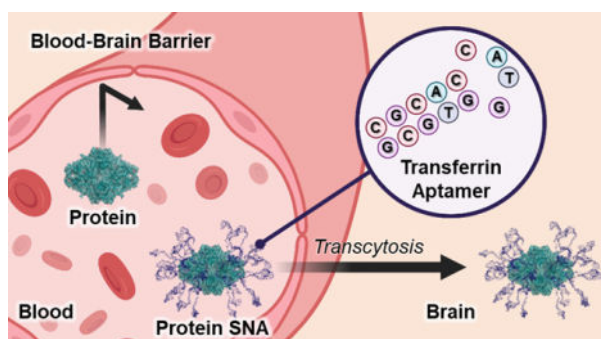
All authors contributed to this work.

Supporting Information.

The Supporting Information is available free of charge on the ACS Publications website at DOI \_\_\_\_\_:

Experimental procedures, oligonucleotide sequences, UV-Vis traces, SDS PAGE gel of purified ProSNAs, DLS, circular dichroism spectra of protein and ProSNA constructs, TEER measurements of *in vitro* BBB model, and fluorescence histology images.

Add financial interests here.



## Keywords

Spherical Nucleic Acids; Protein Delivery; Blood-Brain Barrier; Cellular Uptake

## INTRODUCTION

The targeted delivery of proteins to the brain and central nervous system (CNS) is inherently challenging<sup>1–4</sup> because of the many natural mechanisms and barriers that living systems have evolved to maintain homeostasis.<sup>5</sup> Firstly, unmodified proteins are rapidly degraded by serum proteases<sup>6</sup>, which inherently limits their effective delivery to target sites. Moreover, the blood-brain barrier (BBB) is a highly selective barrier that prevents circulating materials from reaching the CNS and is regulated by endothelial cells whose tight junctions form the impermeable walls of blood vessels<sup>7, 8</sup>. Finally, even if the issues with proteolysis and the BBB are overcome, the cell membrane itself prevents the passive movement of exogenous materials into cells within the brain. As such, strategies to improve the intracellular delivery of proteins to the brain and CNS are warranted.

The BBB is a highly regulated barrier, where endothelial cells control the influx of nutrients and molecules into the brain via both passive and active pathways. Lipid-soluble substances ( $\text{CO}_2$  and  $\text{O}_2$ ) freely cross the BBB<sup>9</sup>, while larger solutes gain access to the brain through active transport via receptors abundant in the endothelium<sup>10, 11</sup>. Accordingly, transport receptors, such as those for insulin<sup>12, 13</sup>, glucose<sup>14</sup>, or transferrin<sup>15–22</sup> (TfR), provide potential avenues for the transport of proteins across the BBB through chemical modifications with the respective ligands. Indeed, lysosomal enzymes modified with antibodies specific to both the insulin<sup>12, 13</sup> and TfR receptor<sup>16, 21, 22</sup> have shown significantly increased BBB penetration *in vivo*. Likewise, aptamers—short single-stranded DNA or RNA oligonucleotides that can assume secondary structures with high affinity and specificity to a target—have been evolved to transport oligonucleotides across the BBB via TfR receptors<sup>20</sup>.

Spherical nucleic acids (SNAs) comprise a class of nanomaterials where oligonucleotides are densely functionalized around a nanoparticle core in a highly oriented manner.<sup>23, 24</sup> This spherical arrangement imparts biological properties to the composite oligonucleotides that are distinct from their linear counterparts, including rapid cellular uptake through scavenger receptor-mediated endocytosis and enhanced stability to degrading enzymes.<sup>25–29</sup> Importantly, proteins that are normally cell impermeable and highly susceptible to protease

degradation are rendered permeable and protease-resistant when functionalized with oligonucleotides as protein spherical nucleic acids (ProSNAs).<sup>29–31</sup> Previous work has found that the DNA shell not only protects the protein, but also alters its biodistribution following systemic administration.<sup>29</sup>

Herein, we hypothesized that the intracellular delivery of proteins to cells within the CNS could be maximized using the ProSNA platform via covalent functionalization with DNA aptamers specific to the TfR receptor. DNA shell design is a critical parameter influencing the cellular uptake and biodistribution of SNAs<sup>29, 32–37</sup>, and we reasoned that a ProSNA's distribution could be specifically directed by introducing a targeting sequence. Therefore, we synthesized ProSNAs bearing TfR aptamers using  $\beta$ -Galactosidase ( $\beta$ -Gal) as a model protein (Transferrin-ProSNAs, Figure 1) and assessed their uptake in an *in vitro* BBB model and their accumulation in the brain following intravenous injection into healthy mice.

## RESULTS & DISCUSSION

Transferrin-ProSNAs Are Successfully Synthesized through the Functionalization of  $\beta$ -Galactosidase with Transferrin Aptamer DNA. The transferrin (TfR) aptamer sequence was selected based on prior research using SELEX to evolve and truncate aptamers, resulting in a 14-mer sequence with a strong binding affinity to the extracellular domain of the mouse transferrin receptor.<sup>38–40</sup> A scrambled DNA sequence, which differs from TfR by four bases in the loop region and has no receptor binding affinity, was synthesized as a control for aptamer-mediated uptake (Table S1.1). Protein SNAs bearing either TfR aptamers (Transferrin-ProSNAs) or the scrambled DNA sequence (Scr-ProSNAs) were synthesized using published protocols<sup>29, 30</sup>.  $\beta$ -Gal was chosen as a model system to assess brain delivery efficacy because it is a large (464 kDa) protein, easily amenable to DNA conjugation, and does not naturally traverse cell membranes. To track the distribution of  $\beta$ -Gal *in vivo*, an AlexaFluor647 (AF647) fluorophore tag was conjugated to surface-accessible cysteines on the protein. Following successful fluorophore labeling, the lysine residues were modified with azide-functionalized-PEG<sub>4</sub>, which was subsequently reacted with DBCO-dT-terminated aptamer DNA strands through a strained-promoted alkyne azide cycloaddition (Figure S1, Table S1.2)<sup>41</sup>. A thymidine DNA spacer (T4) was appended to the 3' end of each sequence to distance the aptamer from the protein surface and promote sequence folding. UV-vis spectroscopy showed that the extent of oligonucleotide loading were nearly identical (39 Transferrin DNA per protein and 37 Scr DNA per protein) regardless of sequence (Figure S2, Table S1.3). Covalent DNA conjugation to the protein core was confirmed using sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), which showed mobility shifts commensurate with the increased mass following DNA addition (Figure S3). Moreover, dynamic light scattering (DLS) showed an increase in hydrodynamic diameter (Figure S4) following DNA functionalization, confirming the successful synthesis of ProSNAs ( $\beta$ -Gal AF647:  $18 \pm 2$  nm, Transferrin-ProSNA:  $24 \pm 3$  nm). Finally, circular dichroism (CD) spectroscopy confirmed that the secondary structure of  $\beta$ -Gal was not altered following fluorophore and DNA conjugation (Figure S5).

### Transferrin-ProSNAs Penetrate Cell Monolayer Faster in an *In Vitro* BBB Model.

With Transferrin-ProSNAs in hand, we sought to determine the rate at which the aptamer is able to promote receptor-mediated transcytosis.<sup>11</sup> For these experiments, bEnd.3 mouse brain endothelial cells, known to highly express the TfR receptor<sup>15, 42</sup>, were cultured to form a monolayer and simulate the tight endothelial junctions that comprise the BBB. To confirm that the bEnd.3 cell barrier was not compromised by the treatment, transendothelial resistance (TEER) measurements of the monolayer was assessed immediately before beginning the experiment, as well as 24 h post-treatment. TEER values for the bEnd.3 cells did not significantly change over time in the presence of the protein (Figure S6) indicating a restrictive barrier and match the reported finding for a stable cell layer<sup>43</sup>. Once the monolayer was established, Transferrin-ProSNAs, Scr-ProSNAs, or  $\beta$ -Gal AF647 were applied to the apical side (Figure 2A). The fluorescence of the media on the basolateral side was measured over time using a plate reader fluorimeter as a proxy for protein transcytosis, as a means to determine the rates of transport. The fluorescence intensity of the media in wells treated with both ProSNA constructs increased steadily as a function of time and at a much faster rate than the AF647 modified protein or TMR dextran 70 kDa (Figure 2B). It is to be expected that the Scr-ProSNA would exhibit some degree of transcytosis capability, as it has been previously shown in a clinical trial that SNAs synthesized with gold cores cross the BBB and accumulate in glioblastoma brain tumors following systemic injection, presumably facilitated by scavenger-mediated movement.<sup>44, 45</sup>

From these analyses, the apparent permeability coefficient ( $P_{app}$ ) was calculated to precisely analyze the kinetics of monolayer transcytosis. This index is widely used as part of the screening process of the flux of a molecule across a barrier, to measure the integrity of the *in vitro* BBB model<sup>46, 47</sup>. A significant increase in fluorescence on the basolateral side was observed at 90 min in cells incubated with Transferrin-ProSNAs (Figure 2C). In addition, Transferrin-ProSNAs showed a higher  $P_{app}$  value ( $7.571 \cdot 10^{-6}$  cm/s) relative to both  $\beta$ -galactosidase ( $6.628 \cdot 10^{-6}$  cm/s) and Scr-ProSNAs ( $P_{app} = 6.941 \cdot 10^{-6}$  cm/s), presumably due to their increased affinity for the transferrin receptor. A table for all the calculated values is included in the SI (Table S2.1).

### Transferrin-ProSNAs Significantly Increase the *in vivo* Accumulation of Proteins in the Brain.

By employing TfR aptamers as the DNA shell, we hypothesized that Transferrin-ProSNAs can target the BBB. Therefore, to study the brain targeting efficiency of Transferrin-ProSNAs, CD-1 mice ( $n = 6$ ) were injected intravenously (IV) with either Transferrin-ProSNAs, Scr-ProSNAs, or native  $\beta$ -Gal at a dosage of 6.5 mg/kg with respect to protein concentration. At 1-h post-injection, the biodistribution of each construct was assessed using an *in vivo* imaging system (IVIS) to measure the fluorescence signal in each organ (Figure 3) following sacrifice and perfusion. Importantly, Transferrin-ProSNAs show a significant increase in brain accumulation ( $p = 0.0001$ , Figure 3A, C) as compared to the Scr-ProSNAs, which only differs in sequence by four bases. Moreover, the targeting by Transferrin-ProSNAs is specific to the brain, as there was no significant increase in accumulation in other filtration organs when comparing mice treated with Transferrin-ProSNAs to those

administered Scr-ProSNAs (Figure 3B, D). Together, this indicates that the DNA aptamers remain folded and recognize their target receptor in a living system.

### **Transferrin-ProSNAs Visualized Throughout the Brain via Confocal Microscopy.**

Encouraged by the increased brain accumulation of Transferrin-ProSNAs on a whole-organ level, we evaluated the tissue and cellular distribution of all materials using immunofluorescent histochemistry (IFC) of brain tissue slices. CD-1 mice ( $n = 3$ ) were injected intravenously with either Transferrin-ProSNAs, Scr-ProSNAs, or  $\beta$ -Gal AF647 at a dosage of 6.5 mg/kg with respect to protein concentration. At 1-h post-injection, animals were sacrificed and perfused, then brain tissues were sectioned and incubated with a DAPI nucleus stain. Strikingly, the AF647 signal originating from the protein conjugated fluorophore was present throughout the entirety of the brain and not sequestered in a single region. Intense, punctate red fluorescence signals present in the same z-plane as DAPI signals, indicating that Transferrin-ProSNAs are taken up by cells (Figure 4, Figure S7). Conversely, there was no detectable signal in brain tissue slices of animals administered the Scr-ProSNA (Figure S8). This is in agreement with the whole-organ-level data, where minimal brain accumulation was observed via IVIS for these two systems. Together, this shows that not only to Transferrin-ProSNAs increase the accumulation of protein in the brain *via* enhanced BBB penetration, but also enable the efficient cellular entry in cells present in the brain.

## **CONCLUSIONS**

Herein, we show that the delivery of proteins to the brain is enhanced through their transformation into ProSNAs comprised of DNA aptamers that bind the TfR receptor. This is significant, because although it has been previously shown that adding a DNA shell changes the biodistribution of proteins<sup>29</sup>, this is a demonstration of using the oligonucleotide sequence as a tissue targeting moiety. This is especially exciting when one considers the vast number of targets and disease sites that aptamers can be easily designed against<sup>48–50</sup>. Moreover, DNA aptamers have been previously successfully employed in ProSNAs as nucleic acid recognition elements to probe pH or glucose levels in living cells.<sup>51</sup> As such, it may be possible to design systems that simultaneously target disease sites (using a targeting aptamer) and probe intracellular conditions upon localization (using a recognition aptamer). Taken together, this work illustrates that DNA shell design can have a profound impact on the resulting distribution of SNAs and can be used to target hard-to-reach organs through intelligent sequence design.

## **MATERIALS AND METHODS**

### **Synthesis $\beta$ -Galactosidase Protein Spherical Nucleic Acids.**

Oligonucleotides (TfR is 5'-GCG TGG TAC CAC GCT TTT T DBCO-dT-3' and Scr is 5'-GCG TGT GCT CAC GCT TTT T DBCO-dT-3') were synthesized using standard phosphoramidite chemistry protocols using universal CPG solid supports and phosphoramidites, as well as coupling reagents purchased from Glen Research. Synthesized strands were purified by HPLC and analyzed using MALDI-MS. ProSNA synthesis was

performed based on literature precedence<sup>29, 30</sup>.  $\beta$ -Gal from an E. coli overproducer was first dissolved in PBS. Next, ten equivalents of AF647-C2-maleimide were added, and the reaction was shaken for 2 hours at 25 °C. Unconjugated AF647 was removed by repeated rounds of centrifugation using a 100 kDa MWCO filter, and the number of fluorophore modifications per protein was calculated using UV-Vis spectroscopy. Then 350 equivalents of NHS-PEG4-azide crosslinker were added to  $\beta$ -Gal-AF647, and the reaction was shaken overnight at 25°C. Unconjugated linker was removed by ten rounds of centrifugation using a 100 kDa filter, and the number of azide modifications was assessed by MALDI-MS. Finally, 350 equivalents of DBCO-dT terminated DNA strands were mixed with  $\beta$ -Gal-AF647-azide and allowed to incubate for 72 h at 25 °C with shaking. Unreacted DNA strands were removed by successive rounds of centrifugation in a 100 kDa filter until the filtrate did not have a detectable absorbance at 260 nm. The number of DNA strands per protein was calculated based on UV-Vis spectroscopy. Characterization of successful covalent conjugation was assessed by SDS PAGE gel.

### Transendothelial electrical resistance (TEER) measurements.

The transendothelial resistance of bEnd.3 cell monolayers was measured using a Millicell ERS-2 voltammeter equipped with a MERSSTX01 electrode after cells reached confluence in a transwell plate. Once TEER values reached that of standard monolayers with tight junctions, cells were treated and resistances measured again at 24 h post-treatment. The final TEER result is expressed  $(\Omega \times \text{cm}^2) = (\text{TEER total } (\Omega) - \text{TEER blank } (\Omega)) \times \text{cm}^2$ .

### In Vitro Blood-Brain Barrier Model.

6.5 mm Transwell<sup>®</sup>-COL collagen-coated 0.4  $\mu\text{m}$  Pore PTFE membrane inserts were placed in a 24-well plate and pre-incubated with full cell culture media overnight at 37°C. The next day, bEnd.3 endothelial cells were seeded on the apical side of the transwell insert and cultured at 37°C in 5%  $\text{CO}_2$ . After nine days, media was removed, fresh media was added to each well, and treatments added to the appropriate insert. The treatment consisted of 60 nM with respect to protein concentration (300  $\mu\text{L}$ ) of  $\beta$ -Gal-AF647, Transferrin-ProSNA, or Scr-ProSNA diluted in full cell culture media and added to the apical side. Cell culture medium (1000  $\mu\text{L}$ ) was added to the basolateral side of the insert. Cells were incubated at 37°C in 5%  $\text{CO}_2$  and 75  $\mu\text{L}$  of media was removed from the lower chambers at set timepoints and fluorescence measured via plate reader fluorimeter ( $\lambda_{\text{Ex}} = 640 \text{ nm}$ ,  $\lambda_{\text{Em}} = 681 \text{ nm}$ ) and transferred back to the basolateral side. Cumulative transport across the membrane was plotted as increase in fluorescence as a function of time. The slopes of the linear regions were used to calculate the permeability coefficients as previously described<sup>52</sup>. The following equation was used:

$$P_{\text{app}} = (dQ/dt)/(C_0 \times A)$$

Where  $dQ/dt$  is the transport rate, defined as the slope obtained from the regression of the transported amount,  $C_0$  is the initial concentration on the donor side and  $A$  is the surface area of the inserts (0.33  $\text{cm}^2$ ).

### **Ex Vivo Near-Infrared Fluorescence (NIRF) Imaging.**

Female CD1 mice (28–30 g) were administered a single injection via tail vein at a dose of 6.5 mg  $\beta$ -Gal / kg body weight. After 1 h, mice ( $n = 6$ ) were humanely euthanized by cardiac perfusion with PBS while anesthetized. Tissues were harvested and fixed. Organs were imaged using an IVIS system using 650 nm/700 nm excitation/emission filters, and data was quantified by measuring radiant efficiency with the Living Image software.

### **Brain Histology and Imaging.**

Female CD1 mice (28–30 g) were administered a single injection via tail vein at a dose of 6.5 mg  $\beta$ -Gal / kg body weight. 1 h post-injection, mice ( $n = 3$ ) were humanely euthanized by cardiac perfusion while anesthetized. Each mouse was first perfused with 20 mL of PBS, then 20 mL of 4% paraformaldehyde (PFA), and the brains were dissected and stored in 4% PFA with shaking at 4 °C for 2 h. Organs were equilibrated at 4°C in 15% sucrose then 30% sucrose until the organs sunk to the bottom of their respective container. Each organ was then embedded and cryo-sectioned in the sagittal plane at the midline at a 10  $\mu$ m thickness. Slides were stained with DAPI to visualize nuclei. A Leica TCS SP8 Confocal Microscope was used to image different regions of each slide at a 40X magnification.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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### **ABBREVIATIONS**

<b>CNS</b>	Central nervous system
<b>BBB</b>	Blood-brain barrier
<b>TfR</b>	transferrin

<b>β-Gal</b>	β-Galactosidase
<b>SNA</b>	Spherical nucleic acid
<b>ProSNA</b>	Protein spherical nucleic acid
<b>Scr</b>	scrambled
<b>AF647</b>	AlexaFluor647
<b>SDS-PAGE</b>	Sodium dodecyl sulfate polyacrylamide gel electrophoresis
<b>CD</b>	Circular dichroism
<b>IV</b>	Intravenous
<b>IVIS</b>	In vivo imaging system
<b>IFC</b>	Immunofluorescent histochemistry
<b>HPLC</b>	High-performance liquid chromatography
<b>MALDI-MS</b>	Matrix-assisted laser desorption ionization-time of flight mass spectroscopy
<b>PBS</b>	Phosphate-buffer saline
<b>MWCO</b>	Molecular weight cut off
<b>PFA</b>	paraformaldehyde

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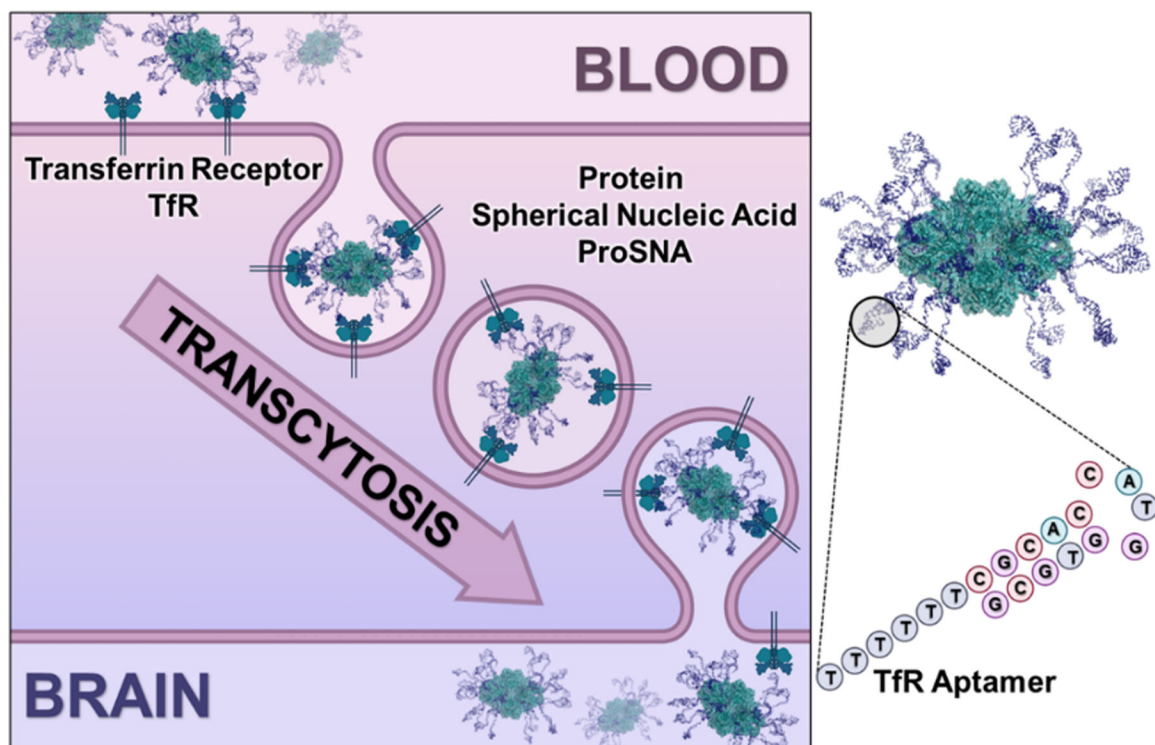
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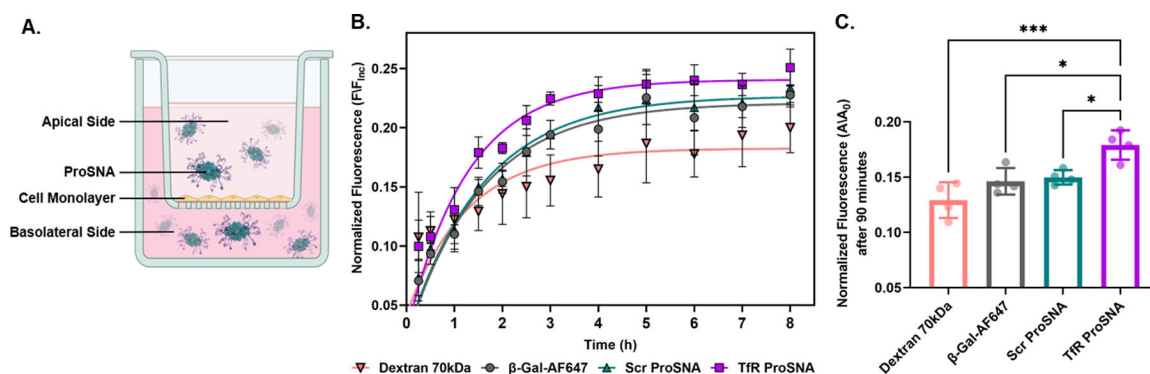
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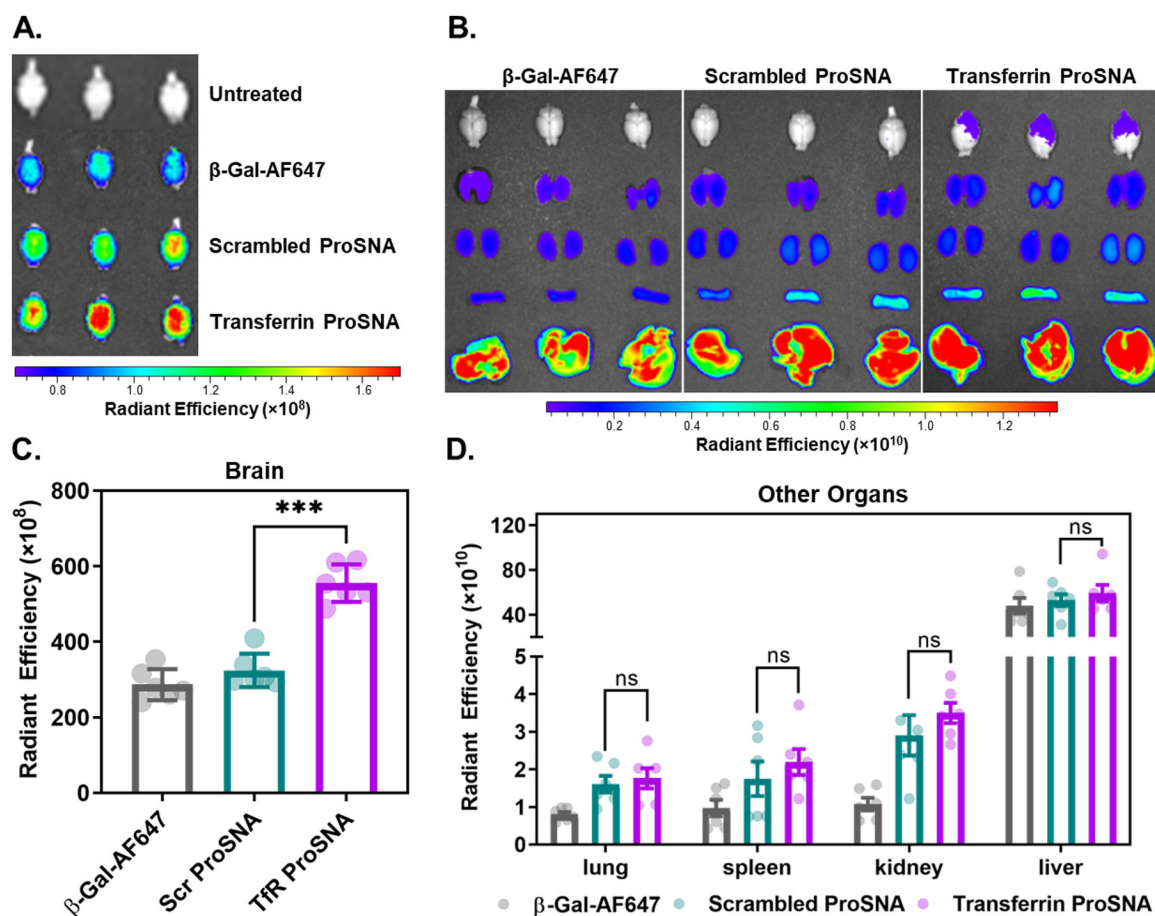
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**Figure 1. Receptor-Mediated Transcytosis of ProSNAs Across the Blood-Brain Barrier.** Cartoon representation of  $\beta$ -Gal ProSNAs being shuttled across an endothelial cell to the brain via the transferrin receptor. The surface of  $\beta$ -Gal is decorated with a shell of transferrin-specific aptamers covalently conjugated to surface lysine residues, resulting in a protein spherical nucleic acid (ProSNA). The representation was adapted from PDB ID 1PX3.

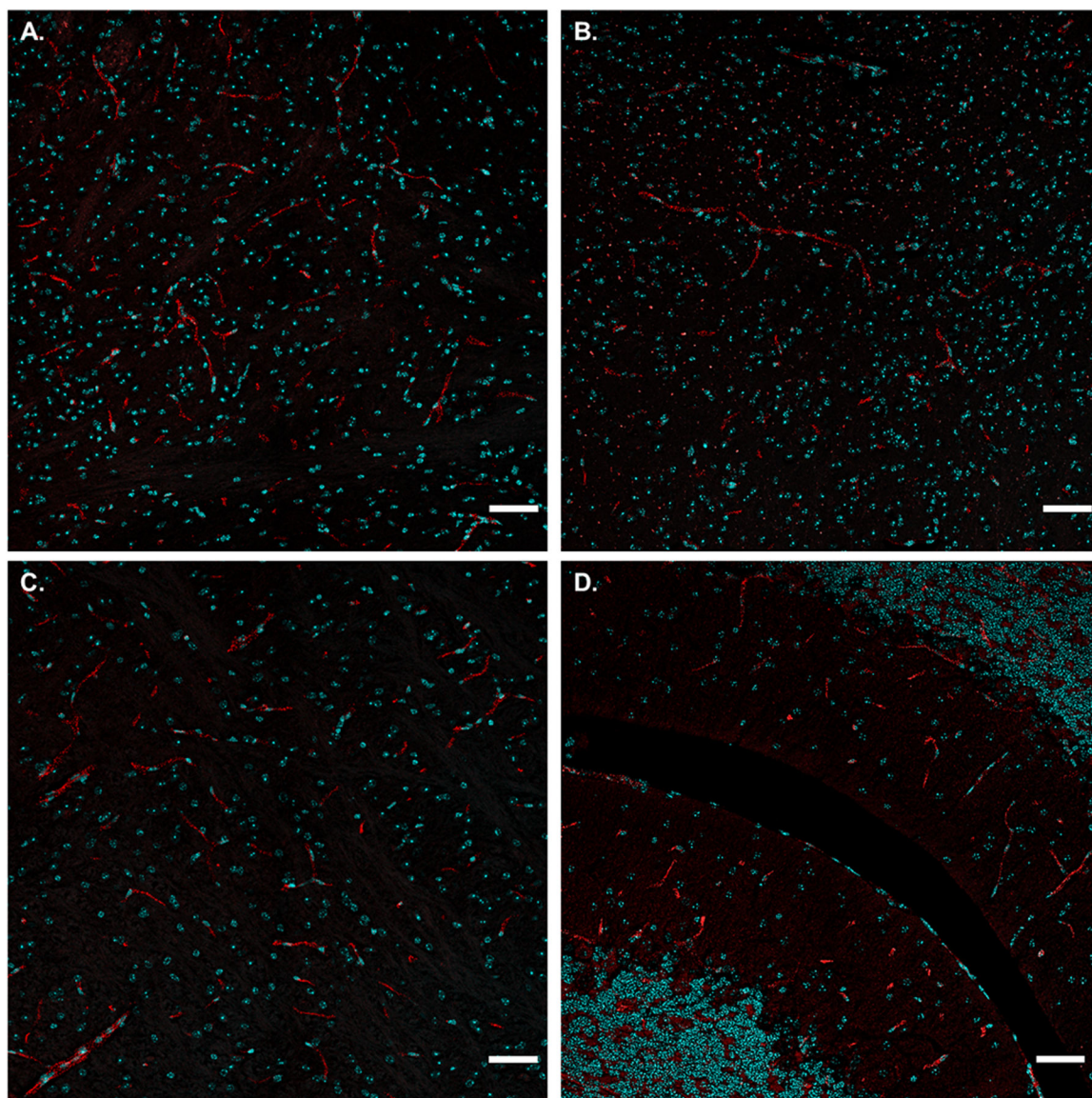


**Figure 2. *In vitro* BBB Analysis Demonstrates Specificity of Transferrin Aptamer for Receptor.** (A) Representation of ProSNAs crossing an *in vitro* BBB model. Cells were treated with fluorescently labeled proteins on the apical side of a transwell insert. (B) Normalized fluorescence readings of the basolateral side were taken periodically and plotted to measure the protein's transcytosis efficiency. Each point represents the mean of  $n=4$  measurements, the error bars show SD, and the line is a nonlinear regression fit graphed using GraphPad Prism. (C) Explicit fluorescence significance comparison after 90 minutes incubation of either Dextran 70 kDa,  $\beta$ -Gal-AF647, TfR ProSNA or Scr ProSNA. The statistical significance was analyzed using ordinary one-way ANOVA, followed by tukey multiple comparison. \*\*\*\*  $p$ -value < 0.0001, \*\*\*  $p$  < 0.001, \*\*  $p$  < 0.01, and \*  $p$  < 0.05.



**Figure 3. Biodistribution of  $\beta$ -Galactosidase Following Systemic Injection in Mice.**

After a 1-h treatment with 6.5 mg of  $\beta$ -Gal/kg mouse via a tail-vein injection; mice ( $n = 6$ ) were sacrificed, perfused, and their organs dissected and fixed. Using IVIS, the fluorescence from the protein's AF647 tag was measured in the (A) brain and (B) other filtration organs. A region of interest was drawn around each organ, and the fluorescence radiant efficiency were quantified (C-D). Each dot represents an individual mouse, the bars represent the mean, and the error bars show SD. A paired  $t$ -test was performed in GraphPad Prism. \*\*\*  $p < 0.0001$ ,  $ns =$  not significant



**Figure 4. Brain Distribution of Transferrin-ProSNAs.**

Representative confocal images of brain histology slides from mice 1-h after intravenous injection of 6.5 mg of  $\beta$ -Gal/kg mouse. Punctate red fluorescence signal (ProSNA) is present throughout the brain, in regions such as: (A) thalamus, (B) hypothalamus, (C) pons, and (D) cerebellum. Nuclei are stained with DAPI (cyan) to approximate the location within the cell. Scale bar = 50  $\mu$ m. Further imaging in other brain regions is represented in Figure S7.