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A brain glioma gene delivery strategy by angiopep-2 and TAT-modified magnetic lipid-polymer hybrid nanoparticles

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Owing to the existence of the blood–brain barrier (BBB), most treatments cannot achieve significant effects on gliomas. In this study, synergistic multitarget Ang-TAT-Fe₃O₄-pDNA-(ss)373 lipid-polymer hybrid nanoparticles (LPNPs) were designed to penetrate the BBB and deliver therapeutic genes to glioma cells. The basic material of the nanoparticles was PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀, and is called (ss)373 herein. (ss)373 NPs, Fe₃O₄ magnetic nanoparticles (MNPs), DOTAP, and DSPE-PEG-MAL formed the basic structure of LPNPs by self-assembly. The Fe₃O₄ MNPs were wrapped in (ss)373 NPs to implement magnetic targeting. Then, the Angiopep-2 peptide (Ang) and transactivator of transcription (TAT) were coupled with DSPE-PEG-MAL. Both can enhance BBB penetration and tumor targeting. Finally, the pDNA was compressed on DOTAP to form the complete gene delivery system. The results indicated that the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs were 302.33 nm in size. In addition, their zeta potential was 4.66 mV, and they had good biocompatibility. The optimal nanoparticles/pDNA ratio was 5 : 1, as shown by gel retardation assay. In this characterization, compared with other LPNPs, the modified single Ang or without the addition of the Fe₃O₄ MNPs, the penetration efficiency of the BBB model formed by hCMEC/D3 cells, and the transfection efficiency of C6 cells using pEGFP-C1 as the reporter gene were significantly improved with Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs in the magnetic field.

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Introduction

Glioma is a kind of malignant tumor derived from the glial tissue of the central nervous system, which grows invasively and is highly aggressive.¹ The treatment of glioma is a comprehensive treatment based on surgical resection; however, the recurrence and metastasis after surgery remain difficult and intractable for clinical treatment.² Currently, there is an urgent need for further treatment to eliminate the residual tumor cells, and achieve a better therapeutic effect in the long term. Gene therapy has been considered a therapeutic method with high potential and promise for glioma treatment. The two important factors for implementing gene therapy successfully are penetrating the blood–brain barrier (BBB) efficiently, and achieving gene transfection of the glioma cells effectively.³ In order to achieve these two aims, we designed and synthesized the Ang-TAT-Fe₃O₄-DOTAP-(ss)373 lipid-polymer hybrid nanoparticle (LPNPs) as the gene delivery system.

The gene delivery system includes a viral vector and non-viral vector.⁴ Various viral vectors have been proven to be able to deliver genes into the central nervous system cells with large capacity and high efficiency. However, their application is limited owing to their biological specificity, such as immunogenicity and toxicity. By contrast, non-viral vectors with low immunogenicity and relative safety have attracted extensive attention.^{5,6} However, the disadvantages of non-viral vectors are poorer brain targeting and lower transfection efficiency for therapy of central nervous system diseases.⁷

To improve brain targeting, the penetration efficiency of the BBB should be improved first. The BBB mainly comprises the brain capillary endothelial cells and other different types of cells, like pericytes and astrocytes, that also contribute to the formation of the BBB.^{8,9} The main function of the BBB is the separation of the blood circulation system from the brain tissue, prevention of harmful substances from entering the brain, and providing a relatively stable internal environment for the brain tissue to maintain its normal function.¹⁰ While the BBB restricts harmful substances from entering the brain, it is also a barrier for gene delivery.¹¹ There is currently no safe and efficient method to circumvent the BBB, and work directly in the brain.¹² Several studies have shown that cell-penetrating peptides (CPPs), protein engineering, and magnetic targeting can effectively penetrate the BBB.^{13–15}

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Receptor-mediated endocytosis plays an important role in BBB penetration. Receptor-mediated transporters are highly expressed in brain capillary endothelial cells, including the insulin receptor, transferrin receptor protein 1, and low-density lipoprotein receptor-related protein 1.¹⁶ They can be used as targets for active brain-targeted gene therapy. Angiopep-2 peptide (Ang), an active target molecule with high affinity for lipoprotein receptor-related protein 1, has been proven to significantly improve the ability of nano-carriers to penetrate the BBB. It exhibits a much higher BBB transcytosis efficacy than transferrin and apotinin.^{17–19} CPPs can penetrate the plasma membrane and the nucleus of almost all live cells.^{20,21} As a kind of CPP, the transactivator of transcription (TAT) peptide has shown fast endosomal or lysosomal escape and quick nuclear localization, and can easily penetrate through biofilm barriers like the BBB.^{22,23} TAT and Ang can bind to DSPE-PEG₂₀₀₀-MAL with the cysteine structure through the MAL molecule group. Magnetic nanoparticles (MNPs) are increasingly used in brain-targeted therapy. In particular, Fe₃O₄ MNPs exhibit promising tumor targeting ability under magnetic fields,^{24–26} and exert a brilliant magnetic resonance imaging performance in a noninvasive manner.²⁷

Previous studies, which focused more on single or double target molecules, did not focus on multiple brain-targeted gene delivery systems like in this study.^{28,29} Considering the introduction of additional target molecules, Fe₃O₄ MNPs in particular, the size and instability of the nanoparticles will increase.²⁵ The aim of this study was to improve the brain targeting of the gene delivery system, while maintaining the stability of the carrier and the particle size being within the safe range.

Another problem to be solved is the improvement of the transfection efficiency of glioma cells. Therefore, PCL₃₇₅₀-SS-PEG₇₅₀₀-SS-PCL₃₇₅₀, which is abbreviated as (ss)373, was chosen as the basic material, and was used to form the basic structure of the LPNPs by self-assembly with DOTAP and DSPE-PEG-MAL. Considering that the low loading rate of pDNA is one of the reasons for the low transfection efficiency, it is necessary to increase the pDNA loading rate as much as possible. (ss)373 NPs have amphiphilicity, which indicates the hydrophobic core constituted by polycaprolactone (PCL) and hydrophilic shell constituted by polyethylene glycol (PEG). The hydrophobic core can carry Fe₃O₄ MNPs, and the hydrophilic shell can connect with DOTAP and DSPE-PEG-MAL efficiently. DOTAP with the positive charge can compress pDNA with the negative charge, and this may increase the loading rate of pDNA.³⁰

Another reason for the low transfection efficiency is that the vector cannot completely decompose and release enough pDNA in glioma cells. The reduction in the sensitivity of LPNPs may solve this problem. The concentration of glutathione in the cytosol, mitochondria, and cell nucleus is 100 to 1000 times higher than that in the extracellular fluids and circulation,³¹ and with a higher concentration in glioma cells.³² The high concentration of glutathione in glioma cells can cause a reductive cleavage of the disulfide-linked PEG shell in (ss)373 NPs, and then decompose the whole LPNPs associated with the rapid release of the encapsulated contents.³³ The dissociation of LPNPs in glioma cells can completely release the pDNA, which

may further increase the transfection efficiency of the glioma cells.

In this study, we focused on the construction and application of multiple brain-targeted Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs. Compared with other single or double molecule brain-targeted gene delivery systems, the multiple brain-targeted magnetic LPNPs may have better ability to penetrate the BBB and transfect glioma cells.

Experimental section

Materials

P_y-ss-PEG₇₅₀₀-ss-P_y was purchased from Biomatrik Inc. (Zhejiang, China). ϵ -Hexanolipid was purchased from Sigma-Aldrich Co. (St Louis, MO, USA). 1,2-Dioleoyl-3-trimethylammonium-propane (DOTAP) and 1,2-distearoyl-*sn*-glycero-3-phosphoethanolamine-*N* [maleimide (poly(ethylene glycol)-2000)] (DSPE-PEG₂₀₀₀-MAL) were purchased from Nanocs Biological Technology (Beijing, China). Oleic acid-modified Fe₃O₄ MNPs were synthesized by Xi'an Ruixi Biological Technology Co., Ltd. (Xi'an, China). Cell-penetrating peptide (TAT) with terminal cysteine (98%, YGRKKRRQRRRC) and Angiopep-2 peptide (Ang) with terminal cysteine (95%, TFFYGGSRGKRNNFKTEEYC) were synthesized by China Peptides Co., Ltd. (Shanghai, China). The plasmid DNA (pEGFP-C1) was provided by Hailing Zhang, a professor from the Institute of Medical Biology Chinese Academy of Medical Sciences. Tryptone and yeast extract were purchased from the OXOID Biological Technology Company (Shanghai, China). The Genopure Plasmid Maxi Kit was purchased from Shanghai Roche Pharmaceuticals Limited (Shanghai, China). 6 \times DNA loading buffer and 50 \times TAE were purchased from Solarbio Life Sciences Company (Beijing, China). The cy5-labeling kit for pDNA was obtained from Mirus Bio (Madison, WI, USA).

Preparation of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs

Preparation of PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀[(ss)373]. P_y-ss-PEG₇₅₀₀-ss-P_y (2 g) was weighed and dissolved fully in 4 mL dimethyl sulfoxide, and then 50 μ L mercaptoethanol and 40 μ L acetic acid were added. Magnetic stirring was performed on the resulting solution at 35 °C for 24 h. Next, the resulting solution was transferred to a dialysis bag (MWCO 8000–10 000) and placed in distilled water for 24 h. After freeze-drying, HO-ss-PEG₇₅₀₀-ss-OH was obtained as a pure white powder product.

HO-ss-PEG₇₅₀₀-ss-OH (1 g) and purified ϵ -hexanolipid monomer (1 g) were added to a polystyrene tube. The ϵ -hexanolipid monomer was completely mixed with HO-ss-PEG₇₅₀₀-ss-OH in the oven at 50–60 °C. A drop of the catalyst stannous caprylate was added to the reaction system, then vacuumed for 5 min, and filled with nitrogen for 5 min. This process was repeated thrice. Finally, the polymerization tube was sealed under the protection of nitrogen and reacted for 24 h in an oil bath at 100 °C. At the end of the reaction, the resulting solution was dissolved in 5–10 mL dichloromethane, and then poured into 250 mL anhydrous ether. After extraction and precipitation, the pure white product was purified thrice and dried in vacuum



at 350 °C. Finally, the pure white granular product, PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀, was obtained, and the product was characterized by ¹H NMR and Gel Permeation Chromatography (GPC) spectrum. PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀ is abbreviated to (ss)373 according to the first digit of the molecular weight of each group and the special disulfide bond structure linking each molecular group.

Preparation of Fe₃O₄-DOTAP-(ss)373 LPNPs. According to the appropriate proportions from our previous experiment³⁴ and the pre-experiment, 20 mg (ss)373 NPs, 1.02 mg DOTAP, 0.8 mg DSPE-PEG₂₀₀₀-MAL, and 100 μL oleic acid-modified Fe₃O₄ MNPs were dissolved in dichloromethane, and were fully mixed in an eggplant flask (5 mL). Then, the eggplant flask was installed on a rotary evaporation apparatus (Eyela N-1001, Rikakikai Co., Ltd., Tokyo, Japan) and the bottom of the flask was in contact with a water bath device, which was heated to 37 °C, rotated, and evaporated at 120 rpm and 580 kPa for 15–20 min. After the liquid in the eggplant flask evaporated completely, the rotary evaporation apparatus was attached to a vacuum system for drying to continue for 1 h. Next, the eggplant flask was removed to a dryer overnight. Sterile PBS (Gibco, Carlsbad, CA, USA) was added to the eggplant flask, and the resulting solution was hydrated at 65 °C for 5 h. Under the conditions of an ice bath, ultrasound was performed with a 5 mm probe for 10 min. Then, the reaction products were dialyzed with 500 mL PBS for 2 h, and the dialysate was changed thrice during dialysis. Eventually, (ss)373 NPs, DOTAP, DSPE-PEG₂₀₀₀-MAL, and Fe₃O₄ MNPs formed the LPNPs (DOTAP-Fe₃O₄-(ss)373 LPNPs) by self-assembly. Other control groups followed the same process, except for the addition of the oleic acid-modified Fe₃O₄ MNPs, and the DOTAP-(ss)373 LPNPs were finally prepared.

Preparation of Ang-TAT-Fe₃O₄-DOTAP-(ss)373 LPNPs. According to the previous study, the appropriate addition mole ratio of Ang : TAT is 1–2 : 1.²⁸ Ang-cys (0.6 mg) and TAT-cys (0.4 mg) (mole ratio = 1 : 1) were dissolved and mixed well in PBS containing the prepared DOTAP-Fe₃O₄-(ss)373 LPNPs, and the resulting solution was allowed to react completely at 4 °C for 12 h. Then, the reaction products were dialyzed with PBS for 2 h, and the dialysate was changed thrice during dialysis. Eventually, the Ang-TAT-Fe₃O₄-DOTAP-(ss)373 LPNPs were prepared.

In the other control groups, the prepared DOTAP-(ss)373 LPNPs were modified with either Ang or both Ang and TAT. By adding only 0.6 mg Ang-cys, 0.6 mg Ang-cys and 0.4 mg TAT-cys, and neither of them, the Ang-DOTAP-(ss)373 LPNPs, Ang-TAT-DOTAP-(ss)373 LPNPs, and DOTAP-(ss)373 LPNPs were obtained, respectively. Subsequently, the process was the same as that followed for the preparation of the Ang-TAT-Fe₃O₄-DOTAP-(ss)373 LPNPs.

Preparation and characterization of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs. Ang-TAT-Fe₃O₄-DOTAP-(ss)373 LPNPs were gently mixed with pDNA in different N/P ratios, including 2.5 : 1, 5 : 1, 10 : 1, and 20 : 1, and were fully complexed by shaking at 20 °C for 30 min.³⁵ Then, the carrier was dialyzed with 500 mL PBS for 2 h, and the dialysate was changed thrice during dialysis. Eventually, the different N/P ratios of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs were prepared. The prepared

systems were directly analyzed qualitatively by agarose-gel electrophoresis (Tanon 1600/1600R, Ewell Bio-Technology Co., Ltd., Guangzhou, China) to determine the optimal N/P ratio. The N/P ratio = 5 : 1 was chosen for subsequent experiments.

We used two fluorescent strategies to detect the position of pDNA in the LPNPs and cells. One was using Cy5 to label pDNA, which directly carried fluorescent molecules, and the labeling process was consistent with the description available in the literature.³⁶ Another was to use pEGFP-C1 as the reporter gene and GFP as the fluorescent tracer to measure the gene transfection efficiency of the C6 cells. Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs were prepared using Cy5-labeled pDNA and pEGFP-C1 with the N/P ratio of 5 : 1, which was also suitable for the control groups, including pDNA-(ss)373 LPNPs, Ang-pDNA-(ss)373 LPNPs, and Ang-TAT-pDNA-(ss)373 LPNPs.

The mean diameters and zeta potential of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs and control groups were measured using Zetasizer (Nano-ZS, Malvern instruments Ltd., Malvern, UK). The storage stability of the different modified LPNPs was determined by measuring the size change at 4 °C for 7 days. The work stability of the different modified LPNPs was determined by measuring the size change at 37 °C with the addition of ECM (Gibco, Carlsbad, CA, USA) supplemented with 10% FBS (Gibco, Carlsbad, CA, USA) for 48 h. The particle morphology was detected by AFM (Multimode 8, Veeco Instruments, Santa Barbara, CA, USA). A CLSM (LSM 710, Carl Zeiss Meditec AG, Gena, Germany) was used to observe the location of the Cy5-labeled pDNA in the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs.

Cell culture

Human brain microvascular endothelial (hCMEC/D3) cells (Thermo Fisher Scientific Company, Waltham, MA, USA) were cultured in ECM supplemented with 10% FBS. C6 Glioma (C6) cells (ACTT, Zhongyuan Ltd, Beijing, China) were cultured in DMEM/F12 medium (Gibco, Carlsbad, CA, USA) containing 10% FBS. All cells were cultured at 37 °C.

The cytotoxicity was evaluated by MTS assay.³⁷ Briefly, hCMEC/D3 and C6 cells were separately seeded in two 96-well plates at a density of 5 × 10³ cells per well, and incubated at 37 °C for 24 h. Then, Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs were diluted with PBS (pH 7.4) to 50 μg mL⁻¹, 100 μg mL⁻¹, 150 μg mL⁻¹, 200 μg mL⁻¹, 300 μg mL⁻¹, and 400 μg mL⁻¹. Each concentration of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was added to six wells of each 96-well plate. Two additional groups were set as the control. One was a blank control group that only had the ECM and incubated cells without LPNPs; the other was the negative control group that only had the ECM and the LPNPs without incubated cells. After processing, the two 96-well plates were incubated at 37 °C for 24 h. Then, the MTS reagent (20 μL) was added to each well and incubated at 37 °C for 2 h in the dark. The plate was read at 490 nm using a microplate reader (Bio-Rad, Hercules, CA, USA) to identify the cytotoxicity of the different modified LPNPs. In the pre-experiment, we determined that the appropriate concentration of pDNA was 0.4 μg mL⁻¹ (12-well plate). The corresponding concentration of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was calculated as



approximately $150 \mu\text{g mL}^{-1}$, and this concentration was used for the entire study.

Preparation of monolayer cell BBB model *in vitro* and measurement of penetration rate

A Transwell plate and hCMEC/D3 cells were used to establish the monolayer cell BBB model *in vitro*. The second or third generation passage hCMEC/D3 cells were placed at a density of 4×10^5 per well in the upper chambers of a Transwell 3460 (Corning, USA) plate. ECM medium was added to the lower chambers of Transwell plate. The medium was changed every day, and the Transwell plate was placed in an incubator at 37°C . The transendothelial electric resistance (TEER) value of the BBB model was detected using Millicell ERS (MERS00001, Millipore, MA, USA). When the TEER value reached above $100 \Omega \text{ cm}^2$ and was stabilized, the actin-tracker Green and DAPI were used for staining in the well. The formation of tight connections between the hCMEC/D3 cells was observed by CLSM to determine whether the BBB model was successfully constructed. We also detected the TEER value of the Ang-TAT- Fe_3O_4 -pDNA-(ss)373 LPNPs at the concentration of $150 \mu\text{g mL}^{-1}$, and the TEER value of the other different modified LPNPs in the control groups before and after penetrating the BBB under a magnetic field. The aim was to identify whether the carriers themselves would destroy the BBB structure. With the concentration of Ang-TAT- Fe_3O_4 -pDNA-(ss)373 LPNPs at $150 \mu\text{g mL}^{-1}$, the BBB

penetration rates of the different modified LPNPs were determined by observing the fluorescence intensity of the Cy5-labeled pDNA using a fluorescence spectrophotometer (FL 6500, PerkinElmer, MA, USA).

Preparation of the glioma model *in vitro* and measurement of the transfection rate of C6 cells after BBB penetration

C6 cells were seeded at a density of 4×10^5 cells per well in the lower chamber of a Transwell plate (70–80% confluence). The hCMEC/D3 cells, which had reached the criteria of the BBB model mentioned above, were placed in the upper chamber. To the upper chamber of the Transwell plate, a $2 \mu\text{g}$ per well mixture was added that contained $150 \mu\text{g mL}^{-1}$ Ang-TAT- Fe_3O_4 -pDNA-(ss)373 LPNPs and DMEM/F12 medium. The magnet was placed at the bottom of the plate, and the cells were cultivated for 24 h. After processing, the expression of GFP was observed in the different groups, which had penetrated the BBB model to transfect the C6 cells using CLSM, and the transfection rates of the different modified LPNPs were analyzed using flow cytometry (BD FACS Calibur, BD Biosciences, CA, USA).

Statistical analysis

All data are presented as the mean \pm standard deviation (SD). The differences among the different modified LPNPs were analysed by one-way analysis of variance (ANOVA) using SPSS

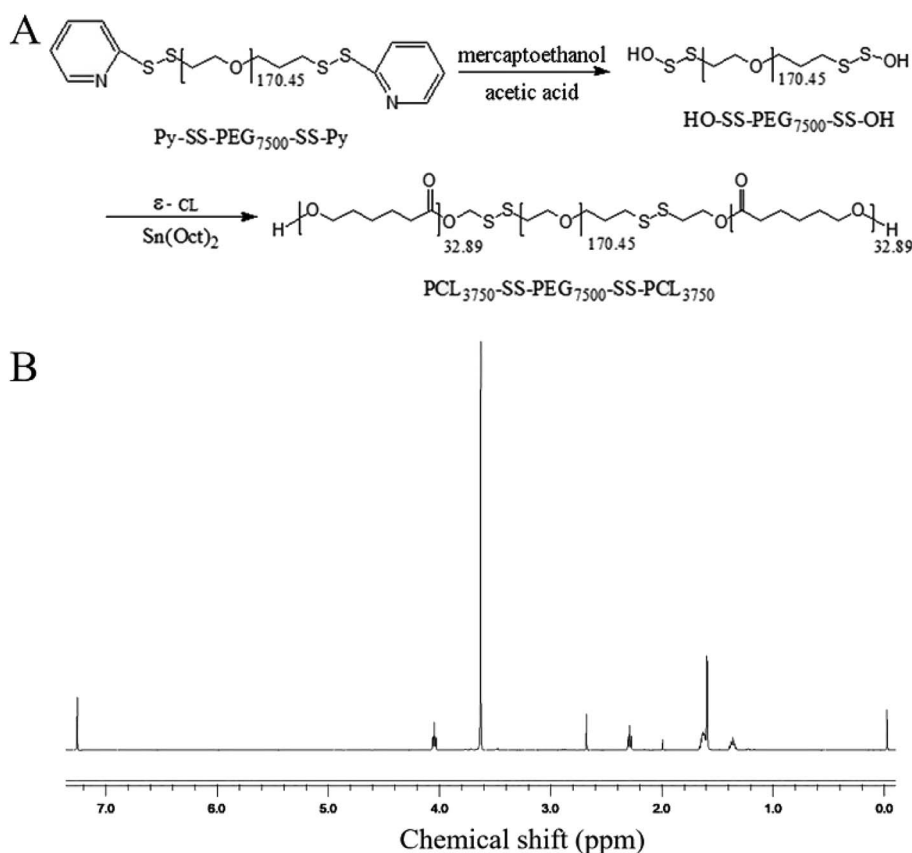


Fig. 1 (A) Synthesis of PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀; (B) ¹H NMR spectrum of PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀.



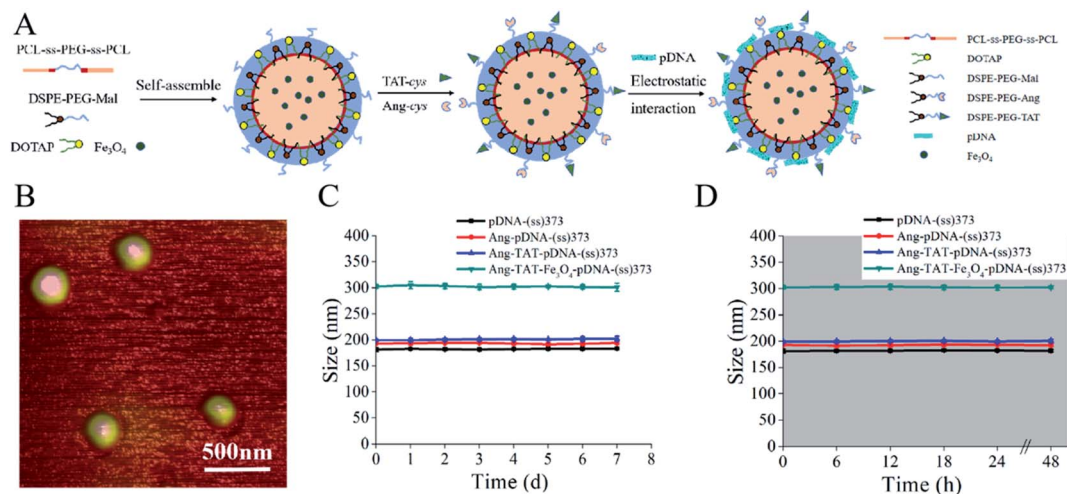


Fig. 2 (A) Schematic diagram of the preparation of Ang-TAT-Fe₃O₄-(ss)373 lipid-polymer hybrid nanoparticles (LPNPs). (B) Atomic force microscope image of Ang-TAT-Fe₃O₄-(ss)373 LPNPs. (C) Stability of LPNPs with different modifications stored at 4 °C dissolved in phosphate-buffered saline. (D) Stability of LPNPs with different modifications stored at 37 °C dissolved in endothelial cell medium with 10% fetal bovine serum.

23.0. *P*-value <0.05 was considered to be significant, and *P*-value <0.01 was considered as highly significant.

Results

The preparation and characterization of PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀ [(ss)373]

As shown in Fig. 1A, the reduction-sensitive material disulfide-bridged triblock copolymer poly(ϵ -caprolactone)₃₇₅₀-ss-poly(ethylene glycol)₇₅₀₀-ss-poly(ϵ -caprolactone)₃₇₅₀ (PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀) was synthesized by ring-opening polymerization of ϵ -CL initiated by HO-ss-PEG₇₅₀₀-ss-OH using Sn(Oct)₂ as a catalyst. The product was characterized by ¹H NMR. The ¹H NMR spectrum was recorded, and is shown in Fig. 1B. The peak at 3.63 ppm was assigned to the methylene protons of -OCH₂CH₂- in the PEG units. Furthermore, the peaks at 1.40, 1.64, 2.30 and 4.04 ppm were assigned to the methylene protons of -CH₂CH₂CH₂-, -CH₂*CH₂CH₂*, -COCH₂*, and -OCH₂*- in the PCL units, respectively. In addition, the number average molecular weight (*M_n*) based on ¹H NMR was 13 442. Besides, the Gel Permeation Chromatography (GPC) spectrum reported that the *M_n* was 19 844, and the weight average molecular weight (*M_w*) was 30 116. These results indicated that the copolymer was synthesized successfully.

Structure and characterization of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs

A schematic diagram of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs is shown in Fig. 2A. PCL₃₇₅₀-ss-PEG₇₅₀₀-ss-PCL₃₇₅₀ was successfully synthesized, and is named (ss)373 NPs herein. DOTAP, DSPE-PEG₂₀₀₀-MAL, and (ss)373 NPs formed the basic structure of LPNPs by self-assembly, which had a hydrophobic core and hydrophilic shell. The PCL groups of (ss)373 NPs constructed the hydrophobic core, and the Fe₃O₄ MNPs were wrapped in the hydrophobic core to achieve the effect for magnetic targeting. The PEG groups of the (ss)373 NPs formed the hydrophilic shell, which was separated from the hydrophobic core by disulfide bonds. Then, TAT-cys and Ang-cys were coupled with the MAL molecule group of DSPE-PEG₂₀₀₀-MAL. Finally, pDNA was compressed on the positively charged DOTAP to form a complete gene delivery system.

The morphology of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was characterized using atomic force microscopy (AFM). Fig. 2B indicates that the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs showed a well-formed spherical shape. As shown in Table 1, the average particle size of the (ss)373 NPs, pDNA-(ss)373 LPNPs, Ang-pDNA-(ss)373 LPNPs, and Ang-TAT-pDNA-(ss)373 LPNPs was 84.55 nm, 181.00 nm, 193.00 nm, and 199.33 nm, respectively.

Table 1 Size distribution (nm) and zeta potential (mV) of LPNPs with different modifications. Data represent mean \pm SD, *n* = 4

Formulation	Size (nm)	Zeta potential (mV)	Polydispersity index (PDI)
(ss)373	84.55 \pm 0.27	-11.73 \pm 0.37	0.28 \pm 0.03
pDNA-(ss)373	181.00 \pm 3.74	6.00 \pm 0.10	0.28 \pm 0.02
Ang-pDNA-(ss)373	193.00 \pm 0.82	8.36 \pm 0.22	0.27 \pm 0.03
Ang-TAT-pDNA-(ss)373	199.33 \pm 1.70	9.07 \pm 0.24	0.23 \pm 0.02
Ang-TAT-Fe ₃ O ₄ -pDNA-(ss)373	302.33 \pm 3.68	4.66 \pm 0.15	0.13 \pm 0.02



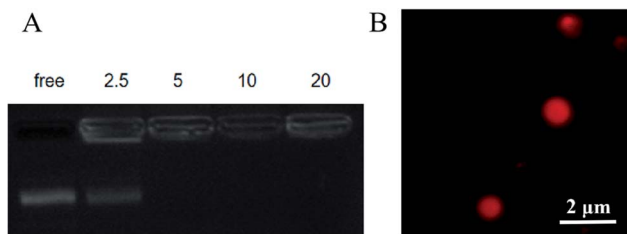


Fig. 3 (A) Gel retardation assay of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 lipid-polymer hybrid nanoparticles (LPNPs) at different N/P ratios. (B) Laser scanning confocal microscopy image of Ang-TAT-Fe₃O₄-(Cy5) pDNA-(ss)373 LPNPs.

The PDI values of those formulations all demonstrated a fairly homogeneous distribution. The average particle size of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs increased to 302.33 nm, demonstrating the successful encapsulation of the Fe₃O₄ MNPs. The average zeta potential of (ss)373 NPs, pDNA-(ss)373 LPNPs, Ang-pDNA-(ss)373 LPNPs, and Ang-TAT-pDNA-(ss)373 LPNPs was -11.73 mV, 6.00 mV, 8.36 mV, and 9.07 mV, respectively. Owing to the influence of Fe₃O₄, the average zeta potential of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was decreased to 4.66 mV. Fig. 2C shows that the different modified LPNPs were able to maintain the size for up to 7 days at almost a constant level at 4 °C in phosphate-buffered saline (PBS). Fig. 2D shows that the different modified LPNPs were able to maintain stability for at least 2 days at 37 °C in endothelial cell medium (ECM) supplemented with 10% fetal bovine serum (FBS).

Determination of the optimal nanoparticles/pDNA (N/P) ratio

Based on the results of the gel retardation assay, when the N/P ratio is >5 : 1, the entrapped pDNA can remain in the wells (Fig. 3A). Considering the improvement in the encapsulation rate, the N/P ratio = 5 : 1 was chosen for application in the entire study. As shown in Fig. 3B, as observed by laser scanning confocal microscopy (CLSM), the Cy5-labeled pDNA was connected tightly with the carrier, and still maintained a stable spherical shape.

Cytotoxicity of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs *in vitro*

The cytotoxicity of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs at a series of concentrations was evaluated in hCMEC/D3 and C6 cells using MTS reagent kits. As shown in Fig. 4, although the concentration of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs reached 400 μg mL⁻¹, the cell viability of both hCMEC/D3 and C6 cells was more than 95% at 2 h and 24 h, and there was no significant difference between the two time periods. According to the results of the pre-experiment, the appropriate concentration of pDNA was 0.4 μg mL⁻¹ (12-well plate) and the corresponding concentration of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was calculated as approximately 150 μg mL⁻¹, which indicated that there is no apparent cytotoxicity of the gene delivery system toward C6 and hCMEC/D3 cells. The Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs at the concentration of 150 μg mL⁻¹ were used in all experiments in this study.

BBB penetration rate of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs

Fig. 5A shows the *in vitro* monolayer BBB model formed by hCMEC/D3 cells to identify the BBB penetration efficiency of different modified LPNPs. The trans-endothelial electrical resistance (TEER) value of the BBB model was detected to ascertain whether the hCMEC/D3 cells had formed the BBB structure successfully. As shown in Fig. 5B, the TEER value in this study tended to be stable on the fourth day and exceeded 130 Ω cm². Moreover, as shown in Fig. 5C, the CLSM images of hCMEC/D3 cells stained with actin-tracker Green and DAPI further confirmed the formation of tight junctions. After different modified LPNPs had penetrated the BBB model, its TEER value was also detected. As shown in Fig. 5D, there was no significant change, which indicates that the different modified LPNPs had almost no damage on the integrity of the BBB model. Fig. 5E shows the BBB penetration rate of different modified LPNPs analyzed semi-quantitatively at 2 h, 6 h, 12 h, and 24 h with the magnet under the Transwell plate. The concentration of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was 150 μg mL⁻¹, and the pDNA was labeled by the Cy5 molecule in this experiment. The BBB penetration rate of pDNA-(ss)373 LPNPs, Ang-pDNA-

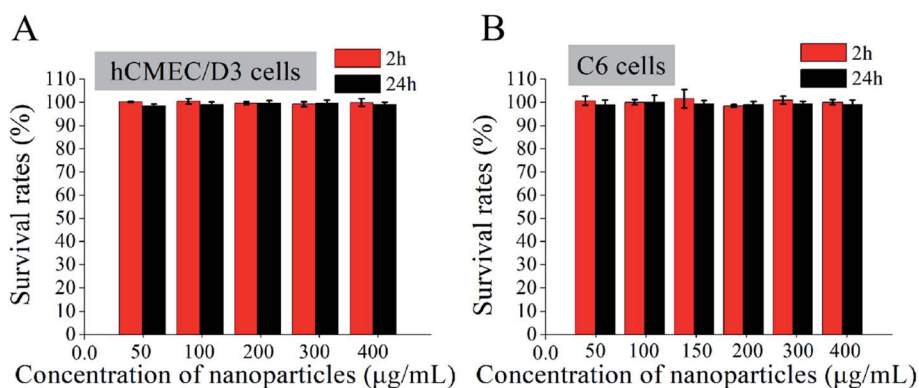


Fig. 4 (A) Survival rates of hCMEC/D3 cells cultured with different concentrations of Ang-TAT-Fe₃O₄-pDNA-(ss)373 lipid-polymer hybrid nanoparticles (LPNPs) at 2 h/24 h. (B) Survival rates of C6 cells cultured with different concentrations of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs at 2 h/24 h. All data are represented as the mean ± SD (n = 5).



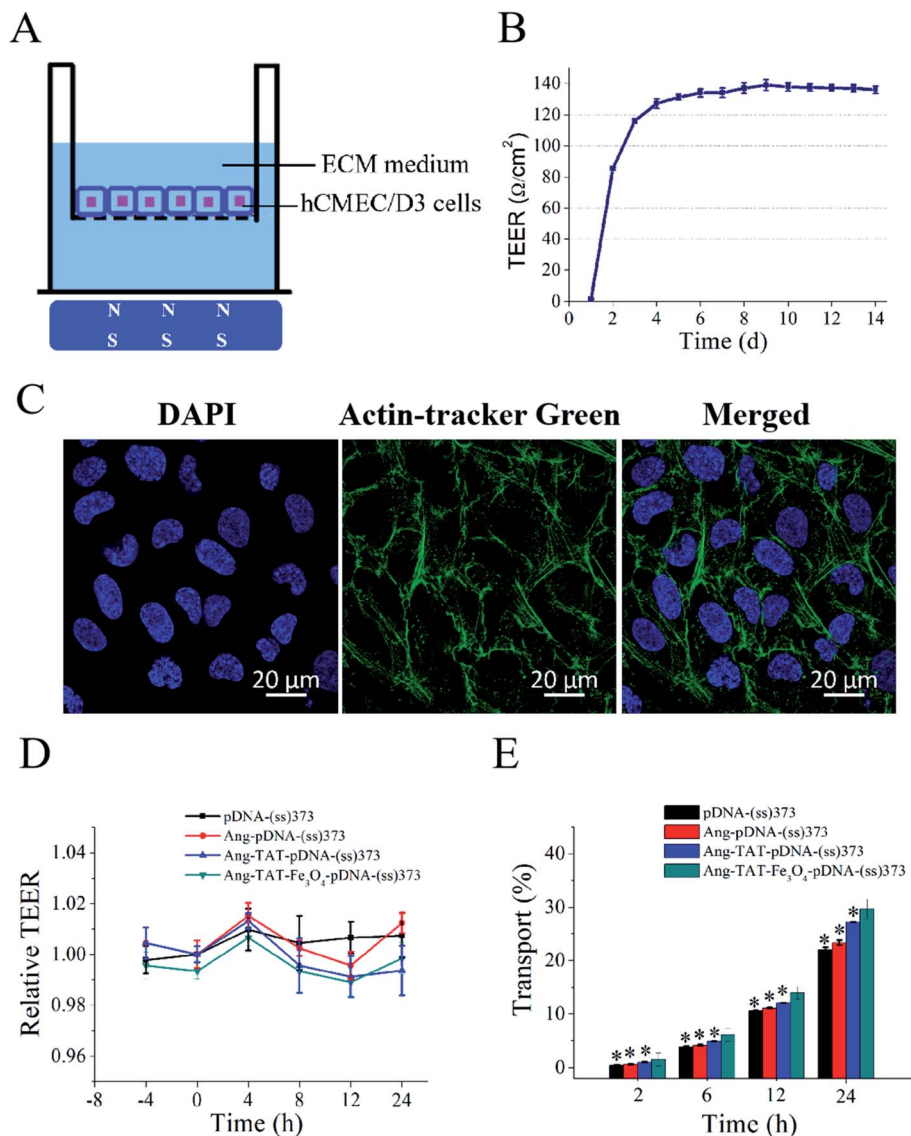


Fig. 5 (A) Schematic diagram of the BBB model constructed by the monolayer hCMEC/D3 cells *in vitro* [the model was placed on a magnetic field]. (B) Trend of trans-endothelial electrical resistance (TEER) values of the constructed BBB model with time. (C) Laser scanning confocal microscopy images of the actin-tracker Green- and DAPI-stained BBB model (scale bar = 20 μm). (D) TEER value changes in the BBB model after treatment with lipid-polymer hybrid nanoparticles (LPNPs) with different modifications. (E) Mean relative transport rate of LPNPs with different modifications in the BBB model. [Magnet placed under the BBB model.] All data are represented as mean \pm SD ($n = 3$); * $P < 0.05$ versus Ang-TAT-Fe₃O₄-pDNA-(ss)373.

(ss)373 LPNPs, Ang-TAT-pDNA-(ss)373 LPNPs, and Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was 0.44%, 0.63%, 1.02%, and 1.55%, respectively, at 2 h. The accumulated BBB penetration rate of pDNA-(ss)373 LPNPs, Ang-pDNA-(ss)373 LPNPs, Ang-TAT-pDNA-(ss)373 LPNPs, and Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was 22.00%, 23.35%, 27.22%, and 29.69% respectively, in a total of 24 h. Compared with the others, the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs showed a significant difference in penetrating the BBB ($P < 0.05$), and the penetration rate increased with prolonged incubation.

In vitro experiment for transfection of C6 cells through BBB

Fig. 6A shows the multiple targeting model constructed using hCMEC/D3 and C6 cells, which was designed to detect the

transfection of C6 cells through the BBB *in vitro*. The C6 cells were seeded in the lower chamber of Transwell plates. The hCMEC/D3 cells, which had reached the criteria of the BBB model mentioned above, were placed in the upper chamber. The magnet was placed under the Transwell plate to exert a magnetic targeting effect. Using pEGFP-C1 as the reporter gene, the green fluorescent protein (GFP) was expressed after successfully transfecting the C6 cells. GFP was used as a tracer to detect the transfection of C6 cells. The CLSM images in Fig. 6B show the transfection of C6 cells with the different modified LPNPs after 24 h. The concentration of Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was 150 $\mu\text{g mL}^{-1}$. There was almost no GFP expression in the free pEGFP-C1 group. With the addition of target molecules, the GFP expression increased



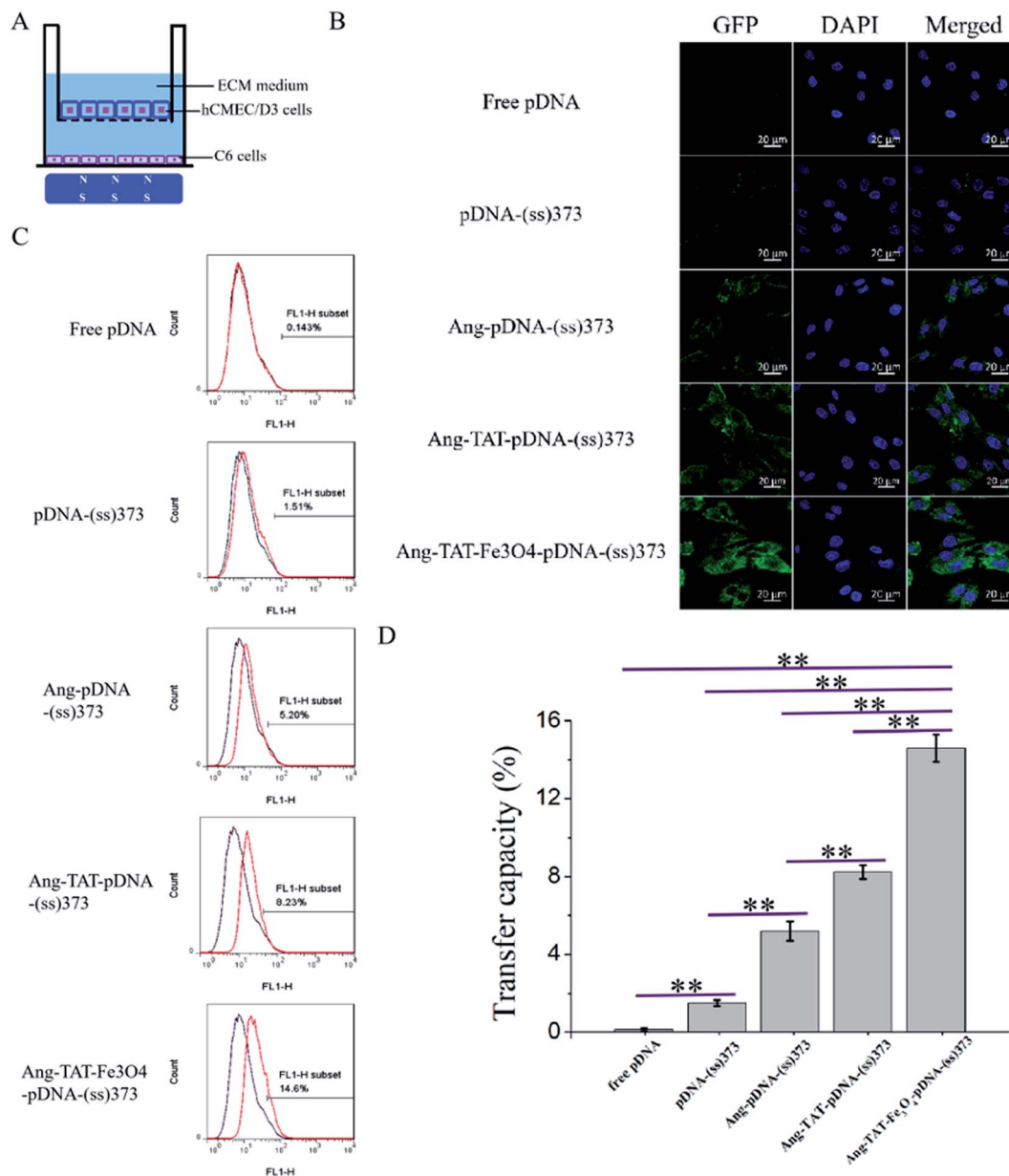


Fig. 6 (A) Schematic diagram of the multiple target model constructed using monolayer hCMEC/D3 and C6 cells [the model was placed on the magnetic field]. (B) Fluorescent images of free pDNA and lipid-polymer hybrid nanoparticles (LPNPs) with different modifications to transfect the C6 cells through the multiple target model under laser scanning confocal microscopy after 24 h (scale bar = 20 μm). (C) Ability of free pEGFP-C1 and LPNPs with different modifications to transfect C6 cells through the BBB model measured by flow cytometer after 24 h. (D) Transfer capacity of free pDNA and LPNPs with different modifications to transfect C6 cells through the BBB model. ** $P < 0.001$. a: Free pDNA; b: pDNA-(ss)373; c: Ang-pDNA-(ss)373; d: Ang-TAT-pDNA-(ss)373; e: Ang-TAT-Fe₃O₄-pDNA-(ss)373.

gradually. The maximum GFP expression was achieved in the pEGFP-C1 loaded with the Ang-TAT-Fe₃O₄-(ss)373 LPNPs group. The flow cytometry results are shown in Fig. 6C. The transfection rate of C6 cells with free pDNA, pDNA-(ss)373 LPNPs, Ang-pDNA-(ss)373 LPNPs, Ang-TAT-pDNA-(ss)373 LPNPs, and Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was 0.143%, 1.51%, 5.20%, 8.23%, and 14.6% at 24 h, respectively. Fig. 6D shows the processed data, which illustrates that compared with free pEGFP-C1, the transfection rate of pEGFP-C1 loaded with the different modified LPNPs was improved gradually. The transfection efficiency of the C6 cells with Ang-TAT-Fe₃O₄-pDNA-(ss)

373 LPNPs was the highest, reaching 14.6% with significant differences from that of other LPNPs at all detected time points ($P < 0.001$). With the addition of target molecules, the transfection efficiency of C6 treated with different modified LPNPs was significantly different ($P < 0.001$).

Discussion

Currently, it is urgent to improve the therapeutic effect of glioma, and gene therapy may be a potential and promising therapeutic method. Two key points to achieve glioma gene



therapy are to increase the BBB penetration efficiency and the transfection efficiency of glioma cells. We designed and synthesized Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs in this study to partially solve the two issues. The Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs showed greater efficiency of BBB penetration and glioma cells transfection *in vitro*. In this study, three methods were used to improve the targeting of the gene to the brain, namely, the introduction of Ang, TAT, and Fe₃O₄. Previous studies have introduced either or both Ang and TAT, which achieved good effects in BBB penetration.^{19,29,38} When the ratio of Ang : TAT is 1–2 : 1, relatively high targeting and low cytotoxicity can be obtained.²⁸ Magnetic targeting is also a popular brain targeting method.^{39–41} Our research integrated the above three targeting methods to greatly improve the penetration efficiency of the BBB under an external magnetic field.

However, with the introduction of Fe₃O₄, the size of the LPNPs also expanded. The mean diameter of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs increased to 302.33 nm. In this study, the result of the cytotoxicity assay showed that this particle size did not cause any apparent damage to the hCMEC/D3 and C6 cells. Moreover, excellent stability plays an important role in the long-term storage and normal exertion of functions. The Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs with this size were able to sustain at least 48 h without aggregation and decomposition under working conditions, as well as under 4 °C storage conditions. This result was similar to that reported by the other studies with a particle size greater than 300 nm, which was still within a safe range and did not cause damage to the brain cells.^{42,43} Therefore, Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs with a diameter of approximately 302.33 nm might safely and effectively penetrate the BBB, and enter glioma cells to achieve gene transfection *in vitro*.

The average zeta potential of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs was around 4.66 mV, which made them easier to approach the negatively charged membrane of the brain capillary endothelial cells.^{44,45} The reason the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs possessed a positive zeta potential was that DOTAP and DSPE-PEG₂₀₀₀-MAL are cationic liposomes, especially DOTAP, which can contribute to the whole lipid formulation to present a positive zeta potential.^{46,47} The positively charged DOTAP also allows it to connect tightly with the negative charge of pDNA.⁴⁸ Both TAT and Ang-2 have a positive charge. When they were used as a modification, the zeta potential increased positively.^{49,50} The zeta potential of Fe₃O₄ MNPs will change with different pH values. In this study, the pH value of the PBS buffer was 7.45, which was used to measure the zeta potential. At this pH value, the Fe₃O₄ MNPs show negative zeta potential and can neutralize parts of the positive charges.⁵¹

Preparing the BBB model successfully is essential for identifying the BBB penetration rate. In this study, the BBB model was established using a monolayer of hCMEC/D3 cells. The formation of the BBB structure was determined by two methods: by measuring the TEER value of the BBB model and observing the stained hCMEC/D3 cells with CLSM. It is generally considered that BBB in mammalian systems show high TEER values, well above 1000 Ω cm². However, it is difficult to achieve the TEER value above 1000 Ω cm² *in vitro*, especially in cell lines compared with that in primary culture. The

monolayers that are formed by hCMEC/D3 cells develop only a low to medium level TEER (around 30–50 Ω cm²) under static culture conditions, according to various reports.⁵² The TEER value in this study exceeded 130 Ω cm² and maintained stability. Previous studies have shown that the BBB structure is formed under this TEER value.^{53–55} It has also been reported that a higher TEER value can be obtained by co-culturing with other cells, such as astrocytes, which can increase the TEER value to approximately 40 Ω cm².⁵⁶ The successful release of pDNA by the gene delivery system in glioma cells is very important to improve gene transfection. In order to improve the gene transfection efficiency, the reduction-sensitive material (ss)373 NPs with disulfide bonds was selected. Previous studies have proven that the dissociation of carriers containing disulfide bonds increases in glioma cells.^{31,57} Disulfide bonds were introduced based on multi-targeting, so that the gene delivery system can decompose relatively specifically in C6 cells with a high concentration of glutathione. Through analyzing the flow cytometry results and CLSM images with GFP as the tracer, the introduction of disulfide bonds that can improve the transfection efficiency of C6 cells was confirmed. According to the results of this study, the mechanism of LPNPs penetrating the BBB should be receptor-mediated endocytosis. However, we did not perform the relevant experiments for further confirmation. In this experiment, the types of cells were relatively simple; only the most typically used hCMEC/D3 cells and C6 glioma cells were selected, and the results were not compared with those of other cell lines. In the follow-up studies, we will verify the effectiveness of the Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs-loaded therapeutic gene for glioma treatment through animal experiments, and further clarify the mechanism of BBB penetration.

Conclusions

To summarize, gene therapy is currently considered a promising approach for glioma treatment. The Ang-TAT-Fe₃O₄-pDNA-(ss)373 LPNPs designed and synthesized in this study are a multi-targeting gene delivery system with several advantages, such as high BBB penetration efficiency, low toxicity, stability, and high transfection efficiency. These advantages may provide new opportunities for the therapeutic gene to penetrate the BBB to treat brain diseases.

Author contributions

X. J. and L. Z. conceived and designed the experiments. L. Q., Y. Q., and D. Z. performed the experiments. L. Q. contributed to the data analysis. Writing, editing, and reviewing were performed by Y. W., L. L., W. X., L. L., and D. B. All authors have read and agreed to the published version of the manuscript.

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List of abbreviations

AFM	Atomic force microscopy
Ang	Angiopep-2 peptide
BBB	Blood-brain barrier
CLSM	Laser scanning confocal microscopy
CPPs	Cell-penetrating peptides
DOTAP	1,2-Dioleoyl-3-trimethylammonium-propane
DSPE-PEG ₂₀₀₀ -MAL	1,2-Distearoyl- <i>sn</i> -glycero-3-phosphoethanolamine- <i>N</i> [maleimide (poly(ethylene glycol)-2000)]
ECM	Supplemented with 10% fetal bovine serum
FBS	Endothelial cell medium
GFP	Green fluorescent protein
GPC	Gel permeation chromatography
LPNPs	Lipid-polymer hybrid nanoparticles
MNPs	Magnetic nanoparticles
PBS	Phosphate-buffered saline
PCL	Polycaprolactone
PCL ₃₇₅₀ -ss-PEG ₇₅₀₀ -ss-PCL ₃₇₅₀	Poly(ϵ -caprolactone) ₃₇₅₀ -ss-poly(ethylene glycol) ₇₅₀₀ -ss-poly(ϵ -caprolactone) ₃₇₅₀
PEG	Polyethylene glycol
(ss) ₃₇₃	PCL ₃₇₅₀ -ss-PEG ₇₅₀₀ -ss-PCL ₃₇₅₀
TAT	Transactivator of transcription
TEER	Trans-endothelial electrical resistance

Conflicts of interest

There are no conflicts to declare.

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References

- 1 L. Zhang, L. Pang, S. Zhu, J. Ma, R. Li, Y. Liu, L. Zhu, X. Zhuang, W. Zhi, X. Yu, L. Du, H. Zuo and Y. Jin, *Int. J. Pharm.*, 2020, **583**, 119384.
- 2 S. L. Hervey-Jumper and M. S. Berger, *Curr. Treat. Options Neurol.*, 2014, **16**, 284.
- 3 H. Yao, K. Wang, Y. Wang, S. Wang, J. Li, J. Lou, L. Ye, X. Yan, W. Lu and R. Huang, *Biomaterials*, 2015, **37**, 345–352.
- 4 S. A. Pena, R. Iyengar, R. S. Eshraghi, N. Bencie, J. Mittal, A. Aljohani, R. Mittal and A. A. Eshraghi, *J. Drug Target.*, 2020, **28**, 111–128.
- 5 B. Caffery, J. S. Lee and A. A. Alexander-Bryant, *Nanomaterials*, 2019, **9**, 105.
- 6 A. Wahane, A. Waghmode, A. Kapphahn, K. Dhuri, A. Gupta and R. Bahal, *Molecules*, 2020, **25**, 2886.
- 7 D. L. Puhl, A. R. D'Amato and R. J. Gilbert, *Brain Res. Bull.*, 2019, **150**, 216–230.
- 8 E. Neuwelt, N. J. Abbott, L. Abrey, W. A. Banks, B. Blakley, T. Davis, B. Engelhardt, P. Grammas, M. Nedergaard, J. Nutt, W. Pardridge, G. A. Rosenberg, Q. Smith and L. R. Drewes, *Lancet Neurol.*, 2008, **7**, 84–96.
- 9 A. Alexander, M. Agrawal, A. Uddin, S. Siddique, A. M. Shehata, M. A. Shaker, S. Ata Ur Rahman, M. I. M. Abdul and M. A. Shaker, *Int. J. Nanomed.*, 2019, **14**, 5895–5909.
- 10 N. J. Abbott, A. A. Patabendige, D. E. Dolman, S. R. Yusof and D. J. Begley, *Neurobiol. Dis.*, 2010, **37**, 13–25.
- 11 M. M. Patel and B. M. Patel, *CNS Drugs*, 2017, **31**, 109–133.
- 12 K. Nagpal, S. K. Singh and D. N. Mishra, *Expert Opin. Drug Deliv.*, 2013, **10**, 927–955.
- 13 Y. Song, D. Du, L. Li, J. Xu, P. Dutta and Y. Lin, *ACS Appl. Mater. Interfaces*, 2017, **9**, 20410–20416.
- 14 L. I. Goulatis and E. V. Shusta, *Curr. Opin. Struct. Biol.*, 2017, **45**, 109–115.
- 15 Y. Cui, Q. Xu, P. K. Chow, D. Wang and C. H. Wang, *Biomaterials*, 2013, **34**, 8511–8520.
- 16 S. Ohtsuki, C. Ikeda, Y. Uchida, Y. Sakamoto, F. Miller, F. Glacial, X. Decleves, J. M. Scherrmann, P. O. Couraud, Y. Kubo, M. Tachikawa and T. Terasaki, *Mol. Pharm.*, 2013, **10**, 289–296.
- 17 T. Tian, J. Li, C. Xie, Y. Sun, H. Lei, X. Liu, J. Xia, J. Shi, L. Wang, W. Lu and C. Fan, *ACS Appl. Mater. Interfaces*, 2018, **10**, 3414–3420.
- 18 Y. Bertrand, J. C. Currie, M. Demeule, A. Regina, C. Che, A. Abulrob, D. Fatehi, H. Sartelet, R. Gabathuler, J. P. Castaigne, D. Stanimirovic and R. Beliveau, *J. Cell Mol. Med.*, 2010, **14**, 2827–2839.
- 19 H. Xin, X. Sha, X. Jiang, L. Chen, K. Law, J. Gu, Y. Chen, X. Wang and X. Fang, *Biomaterials*, 2012, **33**, 1673–1681.
- 20 A. Ur Rahman, S. Khan and M. Khan, *J. Pharm. Pharmacol.*, 2020, **72**, 519–530.
- 21 M. Kristensen, D. Birch and H. Morck Nielsen, *Int. J. Mol. Sci.*, 2016, **17**, 185.
- 22 D. Zhang, J. Wang and D. Xu, *J. Controlled Release*, 2016, **229**, 130–139.
- 23 N. Todorova, C. Chiappini, M. Mager, B. Simona, I. I. Patel, M. M. Stevens and I. Yarovsky, *Nano Lett.*, 2014, **14**, 5229–5237.
- 24 J. Cai, Y. Q. Miao, B. Z. Yu, P. Ma, L. Li and H. M. Fan, *Langmuir*, 2017, **33**, 1662–1669.
- 25 L. Shen, B. Li and Y. Qiao, *Materials*, 2018, **11**, 324.



- 26 C. H. Su, C. Y. Tsai, B. Tomanek, W. Y. Chen and F. Y. Cheng, *Nanoscale*, 2016, **8**, 7866–7870.
- 27 N. Zhang, Y. Wang, C. Zhang, Y. Fan, D. Li, X. Cao, J. Xia, X. Shi and R. Guo, *Theranostics*, 2020, **10**, 2791–2802.
- 28 Y. Zhu, Y. Jiang, F. Meng, C. Deng, R. Cheng, J. Zhang, J. Feijen and Z. Zhong, *J. Controlled Release*, 2018, **278**, 1–8.
- 29 W. Han, G. Yin, X. Pu, X. Chen, X. Liao and Z. Huang, *J. Biomater. Sci., Polym. Ed.*, 2017, **28**, 1695–1712.
- 30 B. B. M. Garcia, O. Mertins, E. R. D. Silva, P. D. Mathews and S. W. Han, *Colloids Surf. B Biointerfaces*, 2020, **193**, 111131.
- 31 R. Cheng, F. Feng, F. Meng, C. Deng, J. Feijen and Z. Zhong, *J. Controlled Release*, 2011, **152**, 2–12.
- 32 Y. Su, Y. Hu, Y. Du, X. Huang, J. He, J. You, H. Yuan and F. Hu, *Mol. Pharm.*, 2015, **12**, 1193–1202.
- 33 H. Y. Wen, H. Q. Dong, W. J. Xie, Y. Y. Li, K. Wang, G. M. Pauletti and D. L. Shi, *Chem. Commun.*, 2011, **47**, 3550–3552.
- 34 L. H. Zhang, C. Y. Hu, S. J. Wu, Z. Chen, F. Fan, Y. Qin and D. W. Zhu, *J. Controlled Release*, 2017, **259**, E84–E85.
- 35 Q. X. Chen, K. Osada, Z. S. Ge, S. Uchida, T. A. Tockary, A. Dirisala, A. Matsui, K. Toh, K. M. Takeda, X. Y. Liu, T. Nomoto, T. Ishii, M. Oba, Y. Matsumoto and K. Kataoka, *Biomaterials*, 2017, **113**, 253–265.
- 36 M. Z. Cao, Y. Gao, N. S. Qiu, Y. Q. Shen and P. H. Shen, *J. Drug Deliv. Sci. Technol.*, 2020, **56**, 1773–2247.
- 37 A. Rizzo, S. Donzelli, V. Girenti, A. Sacconi, C. Vasco, A. Salmaggi, G. Blandino, M. Maschio and E. Ciusani, *J. Exp. Clin. Canc. Res.*, 2017, **36**, 76.
- 38 S. Huang, J. Li, L. Han, S. Liu, H. Ma, R. Huang and C. Jiang, *Biomaterials*, 2011, **32**, 6832–6838.
- 39 L. L. Israel, A. Galstyan, E. Holler and J. Y. Ljubimova, *J. Controlled Release*, 2020, **320**, 45–62.
- 40 A. Skouras, K. Papadia, S. Mourtas, P. Klepetsanis and S. G. Antimisiaris, *Eur. J. Pharm. Sci.*, 2018, **123**, 162–172.
- 41 F. D'Agata, F. A. Ruffinatti, S. Boschi, I. Stura, I. Rainero, O. Abollino, R. Cavalli and C. Guiot, *Molecules*, 2017, **23**, 9.
- 42 B. Ramaswamy, S. D. Kulkarni, P. S. Villar, R. S. Smith, C. Eberly, R. C. Araneda, D. A. Depireux and B. Shapiro, *Nanomedicine*, 2015, **11**, 1821–1829.
- 43 H. Ding, V. Sagar, M. Agudelo, S. Pilakka-Kanthi, V. S. Atluri, A. Raymond, T. Samikkannu and M. P. Nair, *Nanotechnology*, 2014, **25**, 055101.
- 44 X. Wang, Y. Zhao, S. Dong, R. J. Lee, D. Yang, H. Zhang and L. Teng, *Molecules*, 2019, **24**, 3540.
- 45 Y. C. Kuo and I. C. Chen, *J. Phys. Chem. B*, 2007, **111**, 11228–11236.
- 46 B. Dos Santos Rodrigues, S. Lakkadwala, T. Kanekiyo and J. Singh, *Int. J. Nanomed.*, 2019, **14**, 6497–6517.
- 47 B. D. S. Rodrigues, T. Kanekiyo and J. Singh, *Mol. Pharm.*, 2020, **17**, 2054–2063.
- 48 G. Caracciolo and H. Amenitsch, *Eur. Biophys. J.*, 2012, **41**, 815–829.
- 49 A. Borrelli, A. L. Tornesello, M. L. Tornesello and F. M. Buonaguro, *Molecules*, 2018, **23**, 295.
- 50 M. Demeule, A. Regina, C. Che, J. Poirier, T. Nguyen, R. Gabathuler, J. P. Castaigne and R. Beliveau, *J. Pharmacol. Exp. Ther.*, 2008, **324**, 1064–1072.
- 51 I. Antal, O. Strbak, I. Khmara, M. Koneracka, M. Kubovcikova, V. Zavisova, M. Kmetova, E. Baranovicova and D. Dobrota, *Nanomaterials*, 2020, **10**, 394.
- 52 B. Weksler, I. A. Romero and P. O. Couraud, *Fluids Barriers CNS*, 2013, **10**, 16.
- 53 Y. Mi, Y. Mao, H. Cheng, G. Ke, M. Liu, C. Fang and Q. Wang, *Fitoterapia*, 2020, **140**, 104447.
- 54 L. Chen, W. Liu, P. Wang, Y. Xue, Q. Su, C. Zeng and X. Shang, *Brain Res.*, 2015, **1606**, 44–53.
- 55 W. Liu, P. Wang, C. Shang, L. Chen, H. Cai, J. Ma, Y. Yao, X. Shang and Y. Xue, *Brain Res.*, 2014, **1573**, 17–26.
- 56 B. P. Daniels, L. Cruz-Orengo, T. J. Pasieka, P. O. Couraud, I. A. Romero, B. Weksler, J. A. Cooper, T. L. Doering and R. S. Klein, *J. Neurosci. Methods*, 2013, **212**, 173–179.
- 57 L. Lu, X. Zhao, T. Fu, K. Li, Y. He, Z. Luo, L. Dai, R. Zeng and K. Cai, *Biomaterials*, 2020, **230**, 119666.

