Transportan-derived cell-penetrating peptide delivers siRNA to inhibit replication of influenza virus in vivo

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Introduction: In this study, we report on the development of an effective delivery system for siRNAs; a novel cell-penetrating peptide (CPP), T9(dR), obtained from transportan (TP), was used for in vivo and in vitro testing.

Methods: In this study, toxicity of T9(dR) and TP and efficient delivery of siRNA were tested in 293T, MDCK, RAW, and A549 cells. Furthermore, T9(dR)- and TP-delivered siRNAs against nucleoprotein (NP) gene segment of influenza virus (siNP) were studied in both cell lines and mice.

Results: Gel retardation showed that T9(dR) effectively condensed siRNA into nanoparticles sized between 350 and 550 nm when the mole ratio of T9(dR) to siRNA was ≥4:1. In vitro studies demonstrated that T9(dR) successfully delivered siRNA with low cellular toxicity into several cell lines. It was also observed that T9(dR)-delivered siRNAs inhibited replication of influenza virus more efficiently as compared to that delivered by TP into the MDCK and A549 cells. It was also noticed that when given a combined tail vein injection of siNP and T9(dR) or TP, all mice in the 50 nmol siNP group infected with PR8 influenza virus survived and showed weight recovery at 2 weeks post-infection.

Conclusion: This study indicates that T9(dR) is a promising siRNA delivery tool with potential application for nucleotide drug delivery.

Keywords: cell-penetrating peptide, CPP, siRNA, inhibition, influenza virus, IV, transportan, nucleoprotein, NP

Introduction

Influenza A virus infection is a leading cause of almost 200,000 hospitalization and 25,000 annual deaths in the United States.1,2 In addition, influenza A virus is capable of causing pandemics, making the discovery of novel therapeutics against influenza A imperative.3 Although vaccination is an effective strategy to prevent influenza virus infection, less coverage of potent vaccination and vaccinal mismatch of seasonal influenza viruses demand serious consideration. Anti-influenza drugs are efficient in treating influenza virus infection, yet they fail to provide complete protection due to frequent emergence of drug-resistant influenza viruses.4 Furthermore, anti-influenza drug must be taken within 48 hours after the onset of symptoms to obtain a clinically successful rapid response.5,6 Nucleoprotein (NP) of influenza virus is a structural protein which encapsidates the negative-strand viral RNA. Among eight gene segments of type A influenza virus, NP gene is highly conserved.7 Therefore, NP may be targeted for the development of genetic therapy for influenza virus infection.

RNAi is a sequence-specific biological process in which short (21–26 nucleotides), double-stranded RNA molecules inhibit the expression of target genes through a series
of steps after the formation of RNAi-induced silencing complex. In the past decades, RNAi technique was also introduced into antivirus research. An array of siRNAs have been designed to interfere with the expression of gene segments of influenza viruses and have been shown to inhibit the replication of influenza virus in mammalian cells and mouse model. However, with negative charge, it is hard for siRNA to access intracellular active site. To address this issue, a plethora of delivery platforms have been developed. Current delivery tools for siRNA are cationic lipid and virus-based vectors, but the latter has been facing problems when used in clinical trials. Cell-penetrating peptides (CPPs), consisting of a class of highly effective short peptides, are able to penetrate cell membranes and transport different components into cells with high efficiency. When CPPs non-covalently bind to oligonucleotides and form non-covalent nanoparticles, they retain the translocation properties. These short, synthetic peptide-based vectors were considered as biocompatible and economical candidates for the delivery of therapeutic oligonucleotides. Binding potent agents to CPPs has been suggested to improve their entry into cells and bioavailability. Among the published CPPs, transportan (TP), a 27 amino acid chimeric peptide which is a combination of N terminal fragment of neuropeptide galanin and membrane interacting wasp venom peptide mastoparan, is an effective cargo delivery tool with low cytotoxicity. Several results have shown that analogs and derivatives of TP can transport small proteins and DNA and RNA oligomers into cells via direct membrane interactions (transient pore formation), however, inefficiently. Arginine-rich cell-penetrating peptide is another kind of delivery implemented for small molecules. Based on the above findings, we designed and synthesized a CPP, T9(dR), with a length of 36 amino acids consisting of TP and nona-D-arginine residues (9(dR)). In vitro studies showed that T9(dR) induced efficient transportation of siRNA into a variety of cell lines with low toxicity and high efficiency. This occurred when the cell lines were treated with a complex of T9(dR) and siRNA. In addition, in vitro and in vivo experiments demonstrated that T9(dR) delivered siNP into epithelial cells in respiratory tract and inhibited replication of influenza virus more efficiently than TP. These results showed that T9(dR) is an interesting vector for siRNA delivery which shows an intriguing perspective for therapy of influenza virus infection.

**Materials and methods**

**Cell culture, influenza virus, and mice**

Madin-Darby Canine Kidney (MDCK) epithelial cells (CCL-34; American Type Culture Collection [ATCC], Manassas, VA, USA) were cultured in modified Eagle’s medium (HyClone, Logan, UT, USA) with 10% fetal calf serum, 1 mM sodium pyruvate, 1× nonessential amino acids, 5 μg/mL gentamycin sulfate, and 4 mM L-glutamine (Thermo Fisher Scientific, Waltham, MA, USA). A549 cells (CCL-185; ATCC), HEK 293T cells (CRL-11268; ATCC), and RAW cells (CRL-2278; ATCC) were cultured in DMEM supplemented with 10% FBS and penicillin (100 U/mL)/streptomycin (100 μg/mL). PR8 influenza virus (H1N1) was produced and rescued using the influenza viral plasmid system consisting of pHW-PB2, pHW-PB1, pHW-PA, pHW-NA, pHW-NS (gifted by Dr Webster), propagated in 10-day-old embryonated eggs (Baisaisi Laboratories, Jinan, People’s Republic of China), titered with MDCK cells, aliquoted, and stored at −80°C for use. LD$_{50}$ of influenza viruses in mice was determined using the Reed–Muench method. BALB/c mice were purchased from experimental animal center of Shandong Province and housed under specific pathogen-free conditions in the LARC facilities at Qingdao Agricultural University.

**Peptide and siRNA**

Published reports stated that TP and 9(dR) could deliver functional siRNA into different cell lines. In our study, we designed a chimeric peptide T9(dR) by linking TP and 9(dR). The sequence was GWTLNSAGYLLGKINLKALAALA and synthesized by GenScript (Nanjing, People’s Republic of China). Unconjugated siRNA against GFP gene (siGFP) and cy3 (an indicator for uptake of siRNA) conjugated siGFP: sense: 5′-AAGCGUACCUGAGUUCAdTdT-3′, anti-sense: 5′-UGAACCUCAGGAUCGCUdTdT-3′, 5′-GGAUCUUAAUUCUCUGAGdTdT-3′ and guide strand: 5′-CUCCGAAGAAAAAGAUCCdTdT-3′ were purchased from GenScript.

**Gel retardation assay and nanoparticle size measurement**

A mixture of siRNA and T9(dR) or TP in PBS (pH 7.4) was kept at room temperature for 15 minutes. T9(dR) siRNA binding capability was evaluated by electrophoresis through 2% agarose in TAE buffer (40 mM Tris–HCl, 1 v/v% acetic acid, and 1 mM EDTA) at 100 V for 20 minutes. The siRNA bands were stained with ethidium bromide and imaged using a VersaDoc Imaging System (Bio-Rad Laboratories, Inc., Hercules, CA, USA). To test the stability of a complex consisting of CPP and siRNA, 100 nmol siRNA and 400 nmol T9(dR) or 1.6 μmol TP were dissolved in 50 μL PBS (pH 7.4), respectively, mixed gently, and kept in room temperature for 15, 30,
and 60 minutes. The size of the complexes was determined at 25°C and 37°C using the Zetasizer Nano Zeta potential analyzer (Malvern Instruments, Malvern, UK) at 15, 30, and 60 minutes after mixing CPP and siRNA together.

Assessment of CPP cytotoxicity
The MTT assay, a colorimetric method, with NAD(P)H-dependent cellular oxidoreductase enzymes being involved, for measuring the activity of mitochondrial succinate dehydrogenase in living cells, was used. In brief, HEK 293T cells were seeded in 96-well plates at a density of 2×10^4 cells/well in 0.2 mL of medium. After 16 hours, CPPs were kept at 25°C for 15 minutes. Then CPPs with different concentrations were added into wells. After 24 hours of incubation, 20 µL MTT solution was added into each well to reach the final concentration of 10 mg/mL. After additional incubation for 2 hours, the medium was removed followed by the addition of 100 µL dimethylsulfoxide, and absorbance was read in a microplate reader (BioTek, Winooski, VT, USA) at 590 nm wavelength.

Fluorescence and flow cytometry
About 2×10^4 A549 cells or 3×10^4 HEK 293T cells were seeded in an eight-well chamber and incubated at 37°C with 5% CO₂ overnight. Subsequently, the cells were treated with a mixture of 4 nmol T9(dR) and 1 nmol cy3 conjugated siGFP or 16 nmol TP and 1 nmol cy3-conjugated siGFP (incubated at room temperature for 15 minutes). At 24 hours post-transfection, the cells in the chamber were fixed in 4% paraformaldehyde for 30 minutes, washed thrice with Na₃PO₄/PBS, and mounted with DAPI (Southern Biotech, Birmingham, AL, USA) and observed on microscope (Nikon eclipse, Chiyoda, Japan). About 5×10⁵ A549 cells, 8×10⁵ HEK 293T cells, 8×10⁵ RAW cells, and 5×10⁵ MDCK cells were seeded in six-well plates, respectively. On the next day, the cells were treated with 40 nmol T9(dR) and 10 nmol cy3 complex conjugated siGFP or 160 nmol TP and 10 nmol cy3 complex conjugated siGFP (incubated at room temperature for 15 minutes). At 24 hours post-transfection, the cells were infected with PR8 H1N1 influenza viruses in 50 µL PBS. After being challenged with influenza virus, mice were observed daily and recorded for weight and clinical symptoms.

Statistical analysis
Comparisons between treated groups were performed by using a nonparametric one-way ANOVA with the Tukey multiple comparison test and Fisher’s exact, and the final data were analyzed by using the log-rank test. The analyses were performed by using GraphPad Prism version 5.0 for Windows (GraphPad Software, Inc., La Jolla, CA, USA). P-values <0.05 were considered to be significant.

Ethical statements
In this study, animal experimental protocol was approved by the Institutional Animal Care and Use Committee at Qingdao Agricultural University. All animal experimental tests were carried out in accordance with the 2016 standards of laboratory animal in China and other related regulations in Animal Welfare Act.

Results
Non-covalent nanoparticle formation
In this study, T9(dR), GWTLNSAGYLLGKINLKA LAALAKKILdRdRdRdRdRdRdR, and TP were synthesized. T9(dR) or TP and siRNA formed non-covalent nanoparticle based on the phenomenon that positively charged peptides could condense negatively charged siRNA predominantly through electrostatic and hydrophobic interactions.
interactions and form nanoparticles.34,35 To evaluate the avidity of T9(dR) or TP with siRNA, gel retardation assays were performed. Results in Figure 1A showed that when the mole ratio of T9(dR) and siRNA against siGFP was ≥4:1, T9(dR) efficiently condensed siRNA and formed complexes, but the effective ratio of TP to siGFP is ≥16:1. Therefore, T9(dR) and siRNA were used in the ratio of 4:1 and TP and siRNA was used in ratio of 16:1 in follow-up experiments. To further assess the association rate of T9(dR) or TP, siGFP and CPP were mixed in PBS and kept at 25°C or 37°C for 15, 30, and 60 minutes, and then the sample sizes were measured on Zetasizer Nano at 25°C or 37°C. Results in Figure 1B showed that nanoparticles formed by T9(dR) and siGFP are around 350 nm at 15 minutes and nanoparticles formed by TP and siGFP are around 450 nm (P<0.05). In addition, the size became larger as the time extended, and the dissociation of the nanoparticles at 37°C is faster than that at 25°C.

Lower toxicity of T9(dR) than TP
Currently, most of the commercial siRNA transfection reagents are cationic lipids, which are with unbalanced toxicity to cells.36,37 We made a comparative analysis of the toxicity of T9(dR) and TP by using MTT assay. Results in Figure 2
showed when 293T cells in six-well plate were treated with 10 µmol T9(dR) or TP, the cell viability was not substantially affected. Based on viability–regression curves, the CPP lethal doses sufficient to kill 50% T9(dR)- or TP-treated cells were 85 and 90 µmol, respectively. These data could be a guide to further test T9(dR) or TP.

Higher delivery efficiency of T9(dR) than TP
To validate the efficient delivery by T9(dR) of siRNA, transportation experiments were conducted on different cell lines with T9(dR) or TP and cy3-conjugated siRNA complex. Uptake of nanoparticles of T9(dR) and siRNA by cells were indicated by cy3-positive cells. As expected, results in Figures 3 and 4 showed that T9(dR) efficiently transported siRNA into HEK 293T, A549, MDCK, and RAW cells. To further evaluate the delivery efficiency of T9(dR), the delivery efficiencies of T9(dR) and TP were compared in different cell lines. Results in Figure 4 showed that T9(dR) had higher delivery efficiency than TP in all cell lines employed.

siNP delivered by T9(dR) inhibited influenza virus replication in vitro
To confirm siRNA delivered by T9(dR) function properly, complex of siNP (20 nmol) and T9(dR) or TP was incubated in MDCK cells or A459 cells. Subsequently, cells were infected with influenza virus of MOI = 0.01. At 24 hours post-infection, the supernatants were harvested. Virus titer in supernatant were titrated. In Figure 5, the results indicated that siNP delivered by T9(dR) or TP inhibited the replication of PR8 influenza virus, but siNP delivered by T9(dR) has higher efficiency than TP. The results suggest that functional siRNA was transported into A549 and MDCK cells.

Figure 3 T9(dR) or TP transported siRNA into (A) 293T and (B) A549 cell lines. After T9(dR) or TP and cy3-conjugated siGFP were incubated for 15 minutes, the complex was transfected into cells seeded in eight-well chamber. At 24 hours post-transfection, cells were fixed, mounted with DAPI, and observed under a microscope (20×).

Abbreviation: TP, transportan.
siRNA delivered by T9(dR) showed potent inhibition on influenza virus replication in mice

To test whether T9(dR) delivered siRNAs inhibit influenza virus replication in vivo, 6-week old BALB/c mice were intravenously injected with complex of T9(dR) or TP with siGFP, or siNP. At 6 hours post-infection, the mice were infected with 100× MLD<sub>50</sub> PR8 H1N1 influenza virus. After infection, clinical symptoms and weight in mice were recorded every day. Mice treated with PBS or siGFP were

Figure 4 The delivery efficiency of T9(dR) with siRNA in A549, 293T, RAW, and MDCK cells (A–D). After T9(dR) and cy3-conjugated siGFP were incubated for 15 minutes, the complex was transfected into cells seeded in six-well plate. At 24 hours post-transfection, cells were run in a flow cytometer to detect cy3-positive cells. **P-values <0.05 were considered to be significant.

Abbreviation: TP, transportan.

Figure 5 T9(dR) delivered functional siRNA into (A) MDCK and (B) A549 cell lines. MDCK and A549 cells were treated with siNP and infected with influenza virus of MOI = 0.01. At 24 hours post-infection, viral titers in supernatant were titrated by standard plaque assay. **P-values <0.05 were considered to be significant.

Abbreviations: MOI, multiplicity of infection; NP, nucleoprotein; TP, transportan.
observed with poor growth, which included ruffled fur, and reduced activity with decreased food and water intake. Data in Figure 6 showed that mice administrated with PBS or siGFP experienced weight loss and death. Although the group injected with 50 nmol siNP expressed just weight loss ($P$-value <0.005), a complete recovery was observed on day 14 post-infection.

**Discussion**

Infection by human influenza A viruses incurs a huge burden to public health worldwide with an estimated 3–5 million hospitalization and 250,000–500,000 deaths every year. In the United States, around 200,000 infections with severe illness and 20,000 deaths annually were due to influenza A virus. Although influenza vaccines and anti-influenza drugs provide protection, their efficacies are inconsistent. Vaccination is an effective way to prevent infection by influenza viruses. Reportedly, there was 50% mismatch between the seasonal influenza viruses and seasonal influenza vaccines from 1997 to 2005. In addition, the poor coverage of vaccination makes the prophylaxis of influenza ineffective. The mutation of neuraminidase (NA) also created a lot of oseltamivir-resistant influenza strains; hence, reinforcing development of new prophylactic and therapeutic tools is necessary according to the surveillance results of annual influenza epidemic.

siRNA, as a potent tool to inhibit expression of target gene, has been investigated intensively in therapeutics. The siRNAs against conserved regions of the influenza gene segments are powerful promising tools to inhibit replication of influenza virus through degradation of target mRNA. However, naked siRNA was rapidly destroyed by serum nuclease. Due to the large size and anionic charge, the major impediment for therapeutic use of siRNAs is the lack of effective platform to deliver siRNA to desired cell types in vivo. Most of the current siRNA delivery implements are chemical regents and often exhibit unbalanced toxicity in both in vitro and in vivo trials. CPPs demonstrate a greater efficiency for the delivery of siRNAs. So far, many CPPs are identified from variable sources. Usually, the CPPs are 5–30 amino acids in length. However, how the length and structure affect the transportation ability of CPPs is still unknown. siRNAs binding to CPP form nanoparticles and confer resistance to nuclease degradation; however, the nanoparticles are driven by electrostatic and hydrophobic interaction creating instability in liquid environment. Thus, it is crucial to enhance the biological stability of the nanoparticles in biological fluids for a system with better delivery efficiency. Several methods were employed to increase the stability in vivo. In this study, we developed T9(dR) for effective delivery of siRNA in vivo. Initially, results of gel retardation showed that T9(dR) and siRNA bind properly

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**Figure 6** siNP transported by T9(dR) inhibited the replication of PR8 influenza virus in vivo. BALB/c mice were divided into seven groups, intravenously injected PBS, siGFP, siNP with T9(dR) or TP, and infected with influenza. Mouse weight was recorded (A) and survival was calculated (B). $P$-values <0.05 were considered to be significant.

**Abbreviations:** NP, nucleoprotein; TP, transportan.
We treated mice with complexes formed by T9(dR) or TP and siNP before infection with PR8 H1N1 influenza virus. The results showed that siNP functioned well in vivo. It suggested that complexes formed by T9(dR) and siNP were resistant to digestion in serum; however, we did not know whether treatment before infection influence the initiation of infection by influenza viruses. When 10 nmol siNP gene was delivered into mouse lung by T9(dR) or TP, we observed partial protection. When 50 nmol siNP was delivered into mouse lung, delivery efficiency of T9(dR) was higher than that observed for TP. These results suggested that T9(dR) would be a promising functional siRNA delivery tool. In this study, only siRNA against the NP gene segment was employed. In future investigation, mixture of siRNA against the NP, M, and polymerase gene segments will be tested to improve the inhibition on influenza viruses. Perhaps, the cocktail siRNAs will enhance the protection against influenza viruses.

**Conclusion**

In this study, we designed and synthesized a TP-derived CPP, T9(dR). Results of in vitro and in vivo trials showed T9(dR) delivered siRNA with higher efficiency as compared to delivery by TP. This study indicated that T9(dR) would prove to be a promising nucleotide delivery tool with potential application for nucleotide drug and vaccine.

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**Disclosure**

All authors report no conflicts of interest in this work.

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