

**DNA-binding Cell-penetrating Peptide-based  
TRAIL Over-expression in Adipose Tissue-derived  
Mesenchymal Stem Cells Inhibits  
Glioma U251MG Growth**

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Yonsei University  
Department of Medicine

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Directed by Professor Kum Whang

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This certifies that the dissertation  
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## ABBREVIATIONS

**TNF** Tumor necrosis factor

**TRAIL** Tumor necrosis factor (TNF)-related apoptosis-inducing ligand

**MSCs** Mesenchymal stem cells

**ASCs** Adipose tissue-derived mesenchymal stem cells

**ASC-TRAIL** ASCs overexpressing TRAIL

**CPP** Cell-permeation peptide

**PBS** Phosphate-buffered saline

**DMEM** Dulbecco's Modified Eagle's Medium

**FBS** Fetal bovine serum

**CM** Conditioned medium

**SDS-PAGE** Sulfate-polyacrylamide gel electrophoresis

**PE** Phycoerythrin

## ABSTRACT

# DNA-binding Cell-penetrating Peptide-based TRAIL Over-expression in Adipose Tissue-derived Mesenchymal Stem Cells Inhibits Glioma U251MG Growth

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*Background/Aim:* Genetic manipulation of stem cells using non-viral vectors is still limited due to low transfection efficiency. We investigated whether the DNA-binding cell-permeation peptides (CPP) can enhance the transfection efficiency of non-viral vectors in adipose tissue-derived mesenchymal stem cells (ASCs) and whether ASCs over-expressing TRAIL through CPP can inhibit the growth of glioma U251MG cells *in vitro* and *in vivo*. *Materials and Methods:* ASCs were genetically engineered to over-express TRAIL by using CPP, pCMV3-TRAIL and lipid-based transfection reagents (X-tremeGENE). *Results:* The transfection efficiency of ASCs increased by approximately 7% using CPP; 53.9% of ASCs were transfected and TRAIL expression in ASCs increased by approximately 3 times compared to X-tremeGENE alone. ASCs over-expressing TRAIL using CPP inhibited growth of glioma U251MG

cells both *in vitro* and in the U251MG xenograft model. *Conclusion:* CPP can be used as an enhancer for genetically manipulating ASCs and tumor treatment.

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Keywords: Adipose tissue-derived mesenchymal stem cells, tumor necrosis factor-related apoptosis-inducing ligand, genetic engineering, glioma, cell permeation peptide.

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## **I. INTRODUCTION**

### ***1.1 Research background and aim***

As a pro-apoptotic gene, tumor necrosis factor (TNF)-related apoptosis-inducing ligand (TRAIL) induces tumor- or infected cell-specific apoptosis through interaction with two agonistic TRAIL receptors, TRAIL-R1/DR4 and TRAIL-R2/DR5 (Pan et al., 1997; MacFarlen et al., 1997). Although TRAIL is effective in tumor regression, including colorectal cancer (Marini et al., 2006), glioblastoma (Nagane et al., 2000), and NSCLC (Voortman et al., 2007), the short half-life of TRAIL in plasma due to rapid clearance by the kidneys shows poor bioavailability (Kelly et al., 2001).

Mesenchymal stem cells (MSCs) have the tumor tropism and can act as a double-edged sword that promotes or suppresses the tumor growth (Rhee et al, 2015; Liang et al, 2021). Owing to the tumor tropism of MSCs, they have been used as a vehicle to deliver anticancer drugs, including cancer therapy

genes (e.g. suicide genes, cytokines and pro-apoptotic genes) for cancer treatment (Pawitan et al., 2020). Moreover, MSCs as a vehicle increase expression of carried genes locally in tumor microenvironments. MSCs expressing TRAIL have been applied to several preclinical models of sarcoma (Guiho et al., 2018), lung cancer metastasis (Loebinger et al., 2009), renal cancer (Grisendi et al., 2010), colorectal cancer (Luetzkendorf et al., 2010), lymphomas (Yan et al., 2013), and breast cancer (Reagan et al., 2012).

Viral or non-viral vector systems are used for the genetic engineering of MSCs (Marofi et al., 2017). Retrovirus, lentivirus, adenovirus, and adeno-associated viruses have been extensively used as viral vectors for gene delivery into MSCs (Marofi et al., 2017; Nowakowski et al., 2013; Oggu et al., 2017). Retroviral or lentiviral vectors, which are integrating viral vectors, can be used to correct genetic pathology or to continuously express target genes throughout the patient's lifespan, but adenovirus can be used for non-inherited diseases or diseases requiring the transient expression of therapeutic genes (Park et al., 2003). Viral vectors have high gene transfer efficiency, but have problems related to potential immunogenicity and insertional mutagenesis (Oggu et al., 2017; Dewey et al., 1999). Single or combinations of cationic lipids, surfactants, peptides, polysaccharides, nanoparticles (gold, magnetic iron), and synthetic polymers have been used for genetic manipulation when using non-viral vectors (Foldvari et al., 2016; Peer et al., 2007). Non-viral vectors are preferable for expressing the therapeutic gene for a short period, but transfection efficiency using such vectors is lower than that of viral vectors (Chira et al., 2015).

Previously, we reported that adipose tissue-derived mesenchymal stem cells (ASCs) cultured at high density, express TRAIL and inhibit the growth of several cancer cell lines, including Huh7, H460, MCF-7, and MDA-MB-231 (Ryu et al., 2014; Jung et al., 2019; Byun et al., 2020). However, the tumor-suppressive capability was significant in an H460 xenograft model (Jung et al., 2019).

Therefore, we investigated whether the DNA-binding cell permeation peptide (CPP) can enhance the transfection efficiency of non-viral vectors in ASCs and whether ASCs overexpressing TRAIL (ASC-TRAIL) through CPP can inhibit the growth of glioma U251MG in vitro and in vivo.

## II. MATERIAL AND METHODS

### *2.1. Cell culture*

This study was approved by the Institutional Review Board of Yonsei University Wonju College of Medicine (CR319048). Human adipose tissues were obtained from three healthy donors (24–38 years of age) who provided written informed consent through elective liposuction procedures under anesthesia at the Wonju Severance Christian Hospital (Wonju, Korea). The adipose tissue-derived mesenchymal stem cells (ASCs) were isolated using a modified protocol as described by Zuk *et al.* (Zuk *et al.*, 2001). Briefly, after removing the contaminated blood cells and local anesthetics with phosphate-buffered saline (PBS) washing, the adipose tissues were digested with 0.075% type IA collagenase (Sigma-Aldrich, St. Louis, MO, USA) in PBS to obtain mononuclear cells. The mononuclear cells ( $5 \times 10^6$ ) were seeded in 100 mm culture dishes with low-glucose Dulbecco's Modified Eagle's Medium (DMEM; Gibco, Rockville, MD, USA) containing 10% fetal bovine serum (FBS; Gibco) and penicillin/streptomycin. After 2 days, the non-adherent cells were removed by changing the media. The ASCs at passage 3 to 5 were used for this experiment. The human glioma cell line U251MG was purchased from the Korean Cell Line Bank (Seoul, Korea). The U251MG cells were maintained in DMEM (Gibco) supplemented with 10% FBS and penicillin/streptomycin. The U251MG and ASCs were indirectly co-cultured using a Transwell plate.

### *2.2. Synthesis of CPP and analysis of its binding to the supercoiled plasmid*

The DNA-binding CPP was ordered from Peptron, Inc. (Daejeon, Korea). Briefly, CPP peptides were synthesized by Fmoc solid-phase peptide synthesis using ASP48S (Peptron) and purified by the reverse phase Prominence HPLC (Shimadzu, Kyoto, Japan) using a Vydac Everest C18 column (250

mm × 22 mm, 10 μm, HiChrom, Berkshire, UK). Elution was carried out with a water-acetonitrile linear-gradient (10–75% (v/v) of acetonitrile) containing 0.1% (v/v) trifluoroacetic acid. Molecular weights of the purified peptides were confirmed using LC/MS (Shimadzu). To confirm the binding affinity of CPP with supercoiled plasmids, 1 μg of pCMV3-GFP (SinoBiological, Wayne, PA, USA) and CPP (0–10 μg) were incubated in 10 μl of DNA binding buffer (50 mM Tris-HCl, pH 7.5, 10 mM MgCl<sub>2</sub>, 250 mM KCl) for 15 min at room temperature (20 to 25°C). The complex of pCMV3-GFP with CPP was loaded in 1% agarose gel, separated by electrophoresis in TAE buffer (40 mM Tris-HCl, 20 mM acetic acid, 1 mM EDTA), and visualized using a ChemiDoc XRS+ system (Bio-Rad, Hercules, CA, USA).

### ***2.3. ASC Transfection***

The X-tremeGENE HP DNA transfection reagents (Roche, Basel, Switzerland) were used to transfect ASCs with pCMV3-TRAIL (SinoBiological). Briefly, ASCs were seeded in a 100 mm culture dish at a density of  $5.2 \times 10^5$  cells and cultured for 24 h. About 10 μg of pCMV3-TRAIL were diluted with 1 ml of serum-free DMEM and incubated with 20 μl of X-tremeGENE HP reagents with or without 5 μg of CPP for 20 min at room temperature (20 to 25°C). Then, the transfection complex was added dropwise to the cells. The cells were incubated for 3 days to express TRAIL and the conditioned medium (CM) was recovered by centrifugation and then stored at -80°C until further analysis. The transfection efficiency with X-tremeGENE HP reagents with or without CPP was determined by using pCMV3-GFP (SinoBiological) and the GFP-positive cells were evaluated using a flow cytometer.

#### **2.4. MTT assay**

The U251MG cells were plated at  $2 \times 10^4$  cells/cm<sup>2</sup> in 96-well plates. After 24 h, the U251MG cells were treated with CM obtained from ASC and ASC-TRAIL cultures for an additional 48 h, and then 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT, Sigma) dissolved in phosphate-buffered saline (PBS) was added to each well (final concentration, 5 mg/ml) and incubated at 37°C for 2 h at room temperature (20-25°C). The MTT formazan was dissolved in 100  $\mu$ l DMSO and incubated for a further 15 min with shaking before the optical density of each well was read at 570 nm using a microplate reader (Molecular Devices, San Jose, CA, USA).

#### **2.5. Immunoblotting**

The cells were lysed in sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) sample buffer (62.5 mM Tris-HCl, pH 6.8; 1% SDS; 10% glycerol; and 5%  $\beta$ -mercaptoethanol). Protein samples were boiled for 5 min, subjected to SDS-PAGE, and transferred to an immobilon polyvinylidene difluoride membrane (Millipore, Billerica, MA, USA). The membrane was blocked with 5% skim milk in Tris-HCl buffered saline containing 0.05% Tween 20 and then incubated with primary antibodies against caspase 3, cleaved caspase 3, and PARP (1:2000, Cell Signaling Technology, Danvers, MA, USA), TRAIL (1:1000, R&D Systems, Minneapolis, MN, USA), and GAPDH (1:1000, Santa Cruz Biotechnology, Dallas, TX, USA). This was followed by treatment with peroxidase-conjugated secondary antibodies (1:2000, Santa Cruz Biotechnology). The membrane was treated with EZ-Western Lumi Pico or Femto (DOGEN, Seoul, Korea) and visualized using a ChemiDoc XRS+ system (Bio-Rad).

## **2.6. Apoptosis assay**

The phycoerythrin (PE)-Annexin-V apoptosis detection kit I (BD Biosciences, San Jose, CA, USA) was used according to the manufacturer's instructions. The cells were harvested, washed twice with cold PBS, and resuspended in binding buffer. Cells were stained with PE-Annexin-V and 7-aminoactinomycin D (7-AAD) for 15 min at room temperature (20-25°C) in the dark. The cells were then analyzed without washing on a flow cytometer (BD FACSAria III) within 1 h.

## **2.7. Animal studies**

All the animal experiments were performed according to institutional guidelines and approved by the Institutional Animal Care and Use Committee of Yonsei University Mirae Campus at Wonju (YWCI-202012-N-006). Five-week-old athymic nude mice were purchased from Central Lab Animal Inc. (Seoul, Korea). Before transplantation, the U251MG or ASCs were washed with PBS thrice, once with Hank's Balanced Salt Solution (HBSS; Sigma-Aldrich), and then resuspended in HBSS for transplantation into the nude mice. The U251MG ( $2 \times 10^6$  cells/mice) and ASCs ( $2 \times 10^6$  cells/mice) were mixed just prior to co-injection. The cells were suspended in 100  $\mu$ l HBSS and subcutaneously injected into the flanks of nude mice. The sham groups were injected only with U251MG ( $2 \times 10^6$  cells/mice). Three weeks later, mice were euthanized in a CO<sub>2</sub> chamber; thereafter, the tumors were excised and photographed, and the tumor weights were measured using an electronic balance.

## **2.8. Statistical analysis**

Data are presented as the mean  $\pm$  standard deviation (SD) or standard error (SE) of the mean. To compare the group means, the Student's *t*-test and one-way analysis of variance were performed, followed by Scheffe's test. A *P* value < 0.05 was considered significant.

## III. Results

### ***3.1. Binding affinity of CPP for pCMV3-GFP***

The DNA-binding CPP peptides are composed of 24 amino acids with a cell-penetrating domain, a linker, and a DNA-binding domain (Figure 1A). To confirm that the synthesized CPP bind to the pCMV3-GFP plasmid, 1  $\mu\text{g}$  of pCMV3-GFP was incubated with different doses of CPP (0-10  $\mu\text{g}$ ) for 15 min. The supercoiled pCMV3-GFP plasmid (SC, 1  $\mu\text{g}$ ) almost disappeared because it bound with 1  $\mu\text{g}$  of CPP for 15 min and then retarded in the well (Figure 1B). To determine the time required for the optimal binding of CPP and pCMV3-GFP, 1  $\mu\text{g}$  of pCMV3 and 0.5  $\mu\text{g}$  CPP were incubated for 0–2 h. About 50% of the supercoiled pCMV3-GFP disappeared after 15 min and SC was rarely observed after 30 min (Figure 1C). Therefore, it was confirmed that 1  $\mu\text{g}$  of pCMV3-GFP could bind to 1  $\mu\text{g}$  of synthesized DNA-binding CPP within 15 min, suggesting that the synthesized CPP has proper DNA-binding capability.

### ***3.2. Enhancement of transfection efficiency of ASCs with plasmid vectors by DNA-binding CPP***

Since the transfection efficiency using a non-viral vector is reported to be low in MSCs (0–40%) (Abdul et al., 2014), we investigated whether CPP can increase the transfection efficiency of non-viral vectors in ASCs. First, the expression levels of TRAIL were determined in ASCs, which were transfected with pCMV3-TRAIL, X-tremeGENE HP transfection reagents, and/or CPP. In the presence of CPP, ASCs over-expressed TRAIL (ASC-TRAIL) and the amount of TRAIL expression increased by more than 2.2 times; and when 1  $\mu\text{g}$  of CPP per 2  $\mu\text{g}$  of pCMV3-TRAIL was used, the expression of TRAIL increased the most by 3 times (Figure 2A). In ASC-TRAIL, TRAIL secreted into the culture medium was also increased in the presence of CPP, and when 1  $\mu\text{g}$  of CPP was used, secreted TRAIL was increased by about 37.6% and the amount of secreted TRAIL was  $754.21 \pm 15.35$  pg/ml compared

to the control at  $548.51 \pm 12.22$  pg/ml (Figure 2B). In addition, flow cytometry analysis indicated that the transfection efficiency of the pCMV3-GFP in the presence of CPP was increased by about 7%, from  $53.88 \pm 5.01\%$  to  $46.71 \pm 3.19\%$  in the case of using only X-tremeGENE (Figure 2C). Therefore, when CPP is used together with X-tremeGENE HP and a non-viral vector, it is possible to further increase gene expression by increasing the transfection efficiency of ASCs.

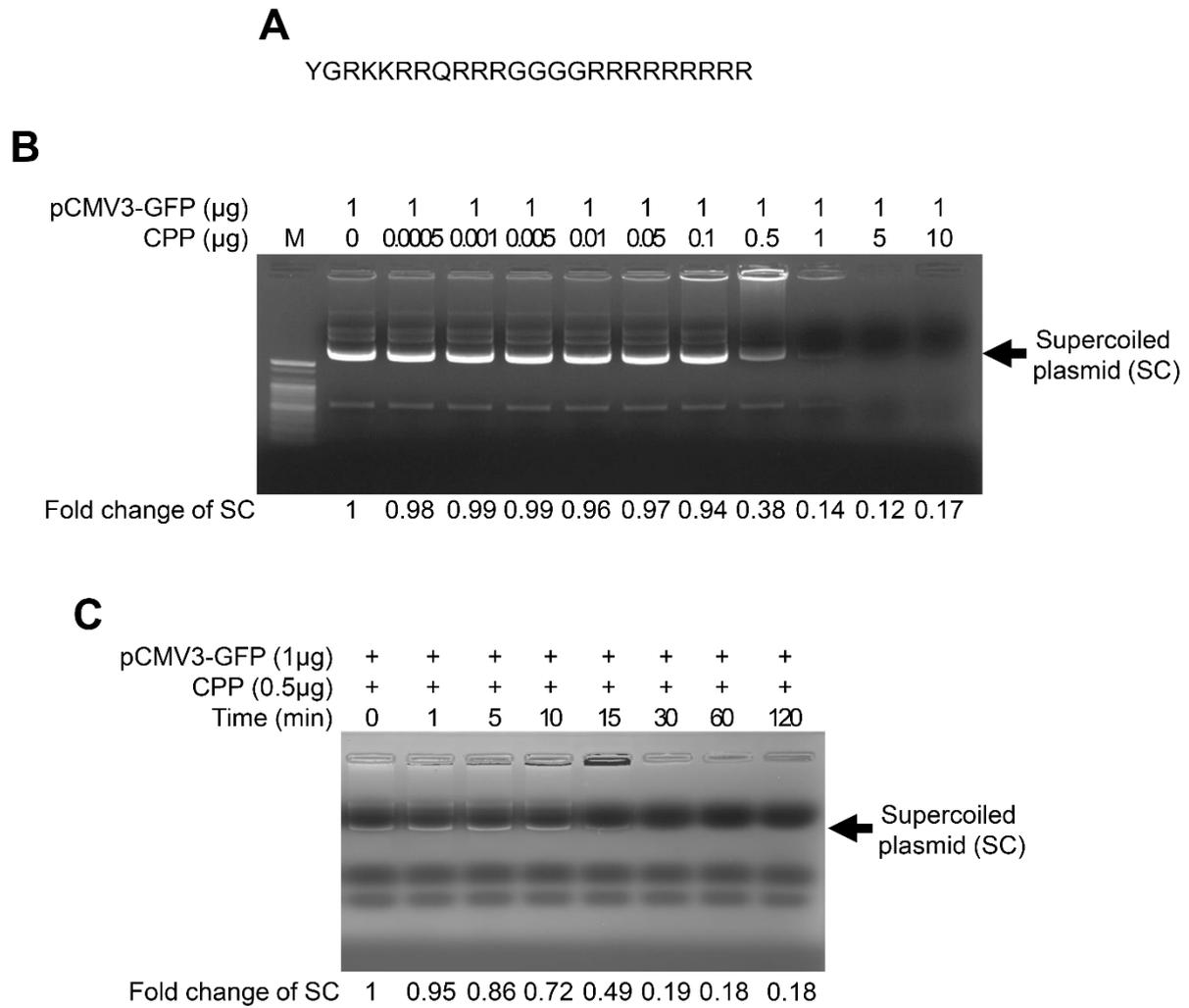
### ***3.3. Cytotoxicity of U251MG by ASC-TRAIL***

To determine whether ASC-TRAIL exhibit cytotoxicity against U251MG glioma cells, these cells were incubated for 2 days with conditioned medium (CM) obtained from ASC or ASC-TRAIL. CM of ASC and ASC-TRAIL induced approximately  $5.78 \pm 0.03\%$  and  $29.27 \pm 0.03\%$  cytotoxicity in U251MG cells, respectively (Figure 3A). In addition, CM of ASCs and ASC-TRAIL induced cleavage of caspase 3 and PARP in U251MG cells, and the levels of cleaved caspase 3 and PARP were higher in U251MG cells treated with CM from ASC-TRAIL than those of cells treated with CM from ASCs (Figure 3B). Next, we analyzed the death of U251MG cells following indirect co-culture with ASC or ASC-TRAIL cells. In indirect co-culture, ASCs reduced apoptosis of U251MG cells, while ASC-TRAIL increased the rate of death in U251MG cells (Figure 3C). These results suggest that ASC-TRAIL can exhibit anti-cancer effects by inducing apoptosis of U251MG cells.

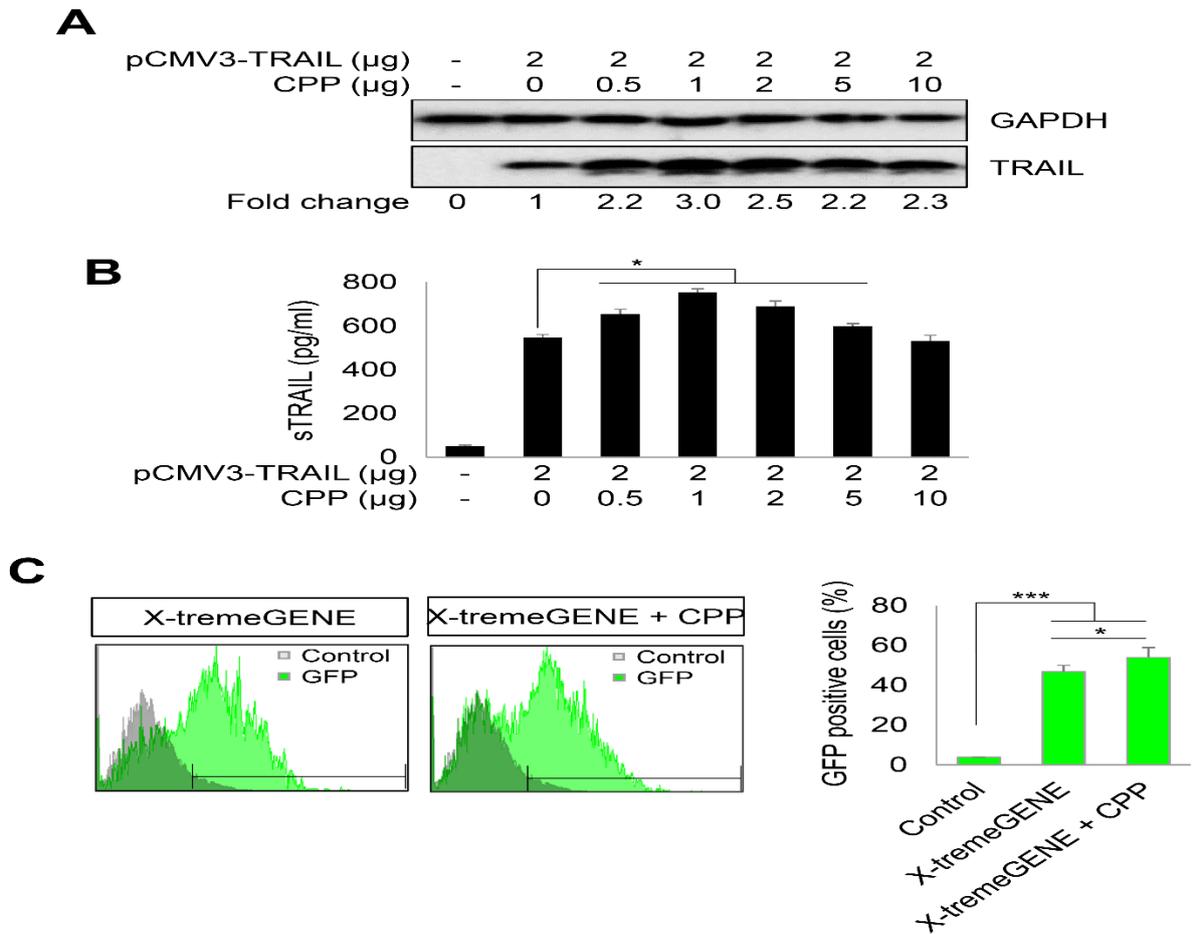
### ***3.4. Anti-tumor effects of ASC-TRAIL in xenograft animal model***

Although ASCs cultured at high density are able to express TRAIL and induce cell death in H460 lung cancer cells *in vitro*, insignificant inhibition of tumor growth was observed in a xenograft animal model (Jung et al., 2018). To investigate whether ASC-TRAIL transfected using CPP show antitumor activities, nude mice (n=5, each) were injected with U251MG only, U251MG + ASC, or U251MG + ASC-TRAIL subcutaneously. After cell transplantation, the increase in tumor size was very slow in both U251MG

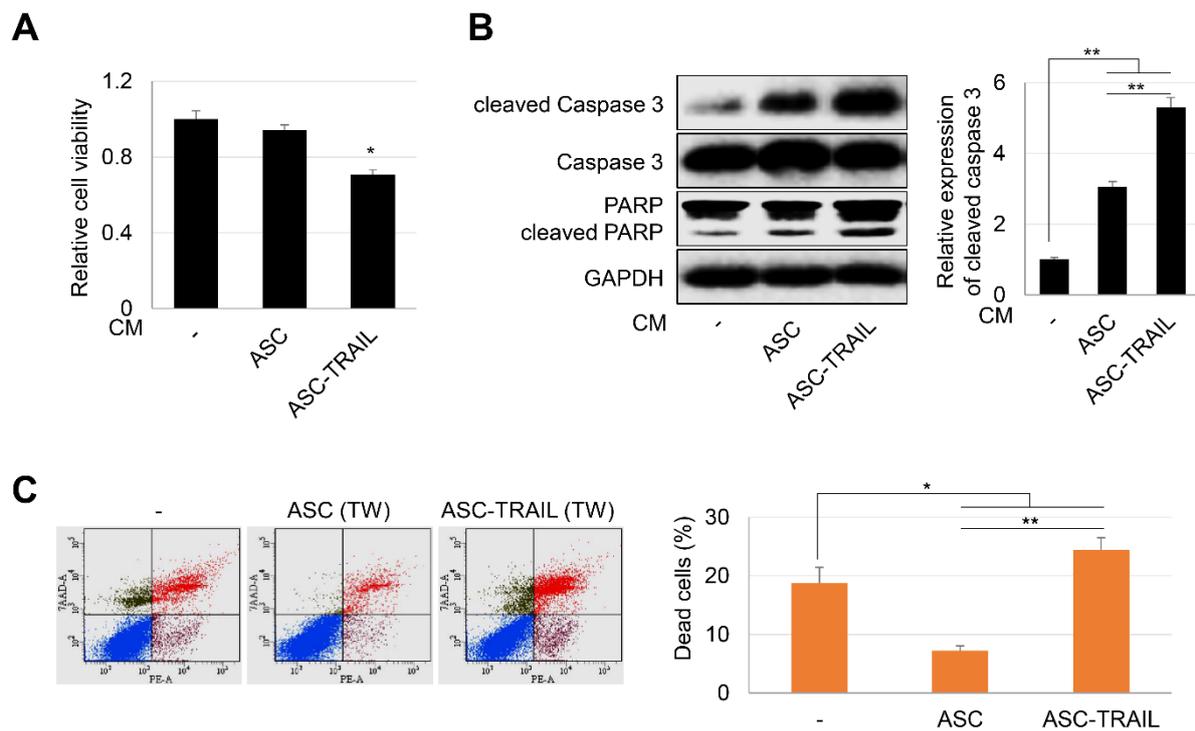
only and U251MG + ASC-TRAIL groups, compared to the U251MG + ASC group. On the 21st day after transplantation, the nude mice were euthanized, and the tumor weight was measured (Figure 4). The weight in the U251MG and U251MG + ASC groups was  $0.17 \pm 0.04$  gram and  $0.60 \pm 0.09$  gram; the latter was about 353% heavier than the former. However, the weight in the U251MG + ASC-TRAIL group was  $0.03 \pm 0.03$  grams, which was 15% of that in the sham control (Figure 4B). Therefore, the naive ASCs can promote tumor growth, whereas ASC-TRAIL can inhibit tumor growth in the U251MG xenograft model, suggesting the use of ASC-TRAIL in tumor treatment.



**Figure 1. The binding affinity of cell-permeation peptides (CPP) for pCMV3-GFP.** (A) Amino acid sequences of CPP. (B and C) Binding affinity of CPP and pCMV3-GFP. One  $\mu\text{g}$  of pCMV3-GFP was incubated with CPP (0–10  $\mu\text{g}$ ) for 15 min in 10  $\mu\text{l}$  of DNA binding buffer. The binding affinity was judged by the degree of reduction in a supercoiled plasmid (SC). One microgram pCMV3-GFP bound to 1  $\mu\text{g}$  of CPP by about 86% affinity within 15 min (B). The time required for optimal binding was determined as the time that SC decreased by 50% after incubating 1  $\mu\text{g}$  of pCMV3-GFP with 0.5  $\mu\text{g}$  of CPP (C).

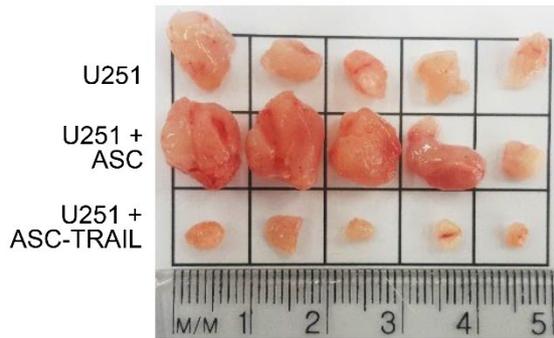


**Figure 2. TRAIL expression in adipose tissue-derived mesenchymal stem cells (ASCs) and transfection efficiency in the presence of cell-permeation peptides (CPP).** (A) TRAIL expression in ASC-TRAIL. ASCs were transfected with pCMV3-TRAIL and CPP for 3 days and then TRAIL expression was detected by immunoblotting. (B) TRAIL secretion levels of ASC-TRAIL. The ASC-TRAIL was transfected in the presence of different doses of CPP (0–10 μg) for 3 days and then soluble TRAIL was analyzed in the conditioned medium (CM) by ELISA. The data are expressed as the mean ± SD from three independent experiments. \* $p \leq 0.05$ . (C) The transfection efficiency in ASCs treated with the pCMV3-GFP and/or CPP. The rate of green fluorescence protein (GFP)-positive cells was analyzed by flow cytometry. The data are expressed as the mean ± SD from three independent experiments. \* $p \leq 0.05$ ; \*\*\* $p \leq 0.001$

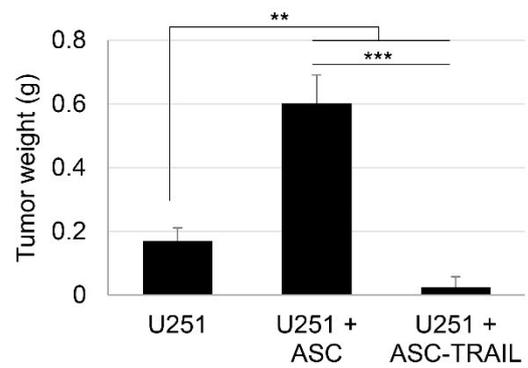


**Figure 3. Death of U251MG cells by adipose tissue-derived mesenchymal stem cell (ASC)-TRAIL and its conditioned medium (CM).** (A) Viability of U251MG cells exposed to CM obtained from ASC and ASC-TRAIL. U251MG cells were cultured with CM obtained from ASC and ASC-TRAIL for 2 days and then cell viability was determined by the MTT assay. Data are shown as the mean  $\pm$  SD from four independent experiments. \* $p \leq 0.05$ . (B) Caspase 3 activation by CM obtained from ASC and ASC-TRAIL. U251MG cells were treated with CM from ASC and ASC-TRAIL for 2 days and cleaved caspase 3 levels in U251MG cells were detected by immunoblotting. Data are shown as the mean  $\pm$  SD from three independent experiments. \*\* $p \leq 0.01$ . (C) Death of U251MG cells indirectly co-cultured with ASC and ASC-TRAIL. After indirect co-culture for 2 days, U251MG cells were stained with PE-annexin-V and 7-AAD to detect the dead cell populations. The dead cell population was the sum of annexin-V-positive/7-AAD-negative, annexin-V-positive/7-AAD-positive, and annexin-V-negative/7-AAD-positive cells. Data are shown as the mean  $\pm$  SD from three independent experiments. \* $p \leq 0.05$ ; \*\* $p \leq 0.01$ .

**A**



**B**



**Figure 4. Anti-tumor effects of TRAIL-overexpressing adipose tissue-derived mesenchymal stem cells (ASCs) in xenograft animal models.** (A) Morphology of tumor mass. At 3 weeks after injection of U251MG and/or ASCs, mice (n=5) were euthanized in a CO<sub>2</sub> chamber and the tumor mass was separated and photographed. (B) Tumor weight at 3 weeks after injection with ASCs. Tumor weight was measured using an electronic balance. Data are shown as the mean ± SE. \*\* $p \leq 0.01$ ; \*\*\* $p \leq 0.001$  (n=5).

## IV. DISCUSSION

Due to the low efficiency (0-40%) of the prevalent transfection methods using non-viral vectors in MSCs, viral vectors (high transfection efficiency, >80%) are widely used to genetically engineer MSCs in preclinical and clinical studies. However, although viral vectors have high gene transfer efficiency and enable stable expression, they have potential immunogenicity and insertional mutagenesis problems (Oggu et al., 2017; Dewwy et al., 1999). For the genetic modification of MSCs using non-viral vectors, electroporation, cationic liposomes, and polymers have shown transfection efficiency of about 40%, 2-35%, and 25-40%, respectively, which is lower than that of viral vectors (>80%). Human ASCs could be transfected with high efficiency (> 80%) by using linear polyethyleneimine and enhancers, which are mixture of histone deacetylase 6 inhibitor and fusogenic lipids (Ho et al., 2020). Moreover, solid gold nanoparticles and the pentapeptide Ku70 have been shown to enhance cellular uptake and expression of the target gene with transfection efficiency of more than 80% (Flanagan et al., 2011).

We observed that with CPP as an enhancer, the transfection efficiency of X-tremeGENE increased up to 53.9%, TRAIL expression increased by 3 folds, and ASC-TRAIL could inhibit tumor growth in vitro and in vitro. Transfection efficiency of our system was about 25% lower than that of Ho and Murpski's method (Ho et al., 2020; Muroski et al., 2014). However, compared to X-tremeGENE alone, CPP with X-tremeGENE increased TRAIL expression in ASCs by more than 3 times and TRAIL secretion by about 37.6%.

The genetic engineering methods of MSCs using a non-viral vector must not only have high transfection efficiency but also not cause toxicity to MSCs. Electroporation, a method of relatively high MSC transfection efficiency (40%) of non-viral vectors, greatly reduces the viability of MSCs (<50% viable) (Flanagan et al., 2011). Compared to cancer cell lines, MSCs proliferate very slowly. Therefore, the transfection method should not affect MSC viability. We confirmed that when CPP was used together with lipofectamine 2000 (Invitrogen, Eugene, OR, USA), PolyMag magnetofection reagent (OZ Biosciences, Marseille, France), and CombiMag transfection reagent (OZ Biosciences), the

expression of the target gene could be increased dramatically. More importantly, when CPP was used together with above transfection reagents, cytotoxicity in MSCs was rarely observed (Data not shown). In U251MG xenograft model, ASC-TRAIL reduced tumor mass, but ASC increased tumor mass significantly.

In conclusion, CPP can be used as an enhancer for non-viral genetic manipulation of ASCs that can be applied to clinical studies for cancer treatment.

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## VI. ABSTRACT OF KOREAN

DNA결합 세포투과펩타이드 기반 TRAIL 과발현 지방조직유래 중간엽 줄기세포의

신경교종 U251MG 성장 억제

신재식

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*연구 배경 및 목적:* 비바이러스 벡터를 이용한 줄기 세포의 유전적 조작은 낮은 형질 감염 효율로 인해 여전히 제한적이다. 이번 연구에서 DNA 결합 세포투과펩타이드(CPP)가 지방조직유래 중간엽 줄기세포(ASC)에서 비바이러스 벡터의 형질감염 효율을 향상시킬 수 있는지, 그리고 세포투과펩타이드(CPP)를 통해 TRAIL을 과발현하는 지방조직유래 중간엽 줄기세포(ASC)가 세포 실험 및 동물 실험에서의 신경교종 U251MG의 성장을 억제할 수 있는지 연구하였다. *재료 및 방법:* 지방조직유래 중간엽 줄기세포(ASC)는 CPP, PCMV3-TRAIL 및 지질기반 형질감염시약(X-tremeGENE)을 사용하여 TRAIL을 과발현하도록 유전적으로 조작하였다. *결과:* 지방조직유래 중간엽 줄기세포(ASC)의 형질감염 효율은 세포투과펩타이드(CPP)를 사용하여 약 7% 증가하였다. 지방조직유래 중간엽 줄기세포(ASC)의 53.9%가 형질감염 되었고 지방조직유래 중간엽 줄기세포(ASC)에서 TRAIL 발현이 지질 기반 형질감염 물질(X-tremeGENE) 단독 사용에 비해 최대 3배 증가하였다. 세포투과펩타이드(CPP)를 사용하여 TRAIL을 과발현하는 지방조직유래 중간엽 줄기세포(ASC)

는 세포 실험 및 U251MG 이종 이식 모델 모두에서 신경교종 U251MG의 성장을 억제하였다. 결론: 세포투과펩타이드(CPP)는 지방조직유래 중간엽 줄기세포(ASC)를 유전적으로 조작하여 종양을 치료하기 위한 증폭자로 사용할 수 있을 것으로 사료된다.

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핵심 되는 말: 지방조직유래 중간엽 줄기세포, 종양괴사인자 관련 세포사멸 유도 리간드, 유전공학, 신경교종, 세포투과펩타이드.