

Multifunctionalized polyethyleneimine-based nanocarriers for gene and chemotherapeutic drug combination therapy through one-step assembly strategy

Dandan Jiang^{1,*}
Mingfang Wang^{1,*}
Tianqi Wang¹
Bo Zhang¹
Chunxi Liu²
Na Zhang¹

¹Department of Pharmaceutics, Key Laboratory of Chemical Biology (Ministry of Education), School of Pharmaceutical Sciences, Shandong University, Jinan, China;
²Pharmaceutical Department, Qilu Hospital of Shandong University, Jinan, China

*These authors contributed equally to this work

Abstract: Gene therapy combined with chemotherapy to achieve synergistic therapeutic effects has been a hot topic in recent years. In this project, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene (*TRAIL*) and doxorubicin (Dox)-coloaded multifunctional nanocarrier was constructed based on the theory of circulation, accumulation, internalization, and release. Briefly, polyethyleneimine (PEI) was selected as skeleton material to synthesize PEI-polyethylene glycol (PEG)-TAT (PPT). Dox was conjugated to PEI using C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone (C6-SANH), and a pH-sensitive Dox-PEI (DP) conjugate was obtained. Then, intracellular cationic pH-sensitive cellular assistant PPT and DP were mixed to condense *TRAIL*, and *TRAIL*-Dox coloaded PPT/DP/*TRAIL* (PDT) nanocarriers were obtained by one-step assembly. *TRAIL* was completely condensed by DP or PPT when mass ratios (DP/PPT to *TRAIL*) were up to 100:64, which indicated that DP and PPT could be mixed at any ratio for *TRAIL* condensation. The intracellular uptake rate of PDT was enhanced ($P < 0.05$) when the contents of PPT in PPT+DP increased from 0 to 30%. Free Dox and *TRAIL*-loaded nanocarriers (PPT/C6-SANH-PEI/*TRAIL* [PCT]) were selected as controls to verify the synergistic antitumor effects of PDT. Compared with free *TRAIL*, *TRAIL*-protein expression was upregulated by PDT and PCT on Western blotting assays. The in vitro cytotoxicity of PDT was significantly enhanced compared to free Dox and PCT ($P < 0.01$). Furthermore, murine PDT nanocarriers showed higher in vivo antitumor ability than both the Dox group ($P < 0.05$) and the murine PCT group ($P < 0.05$). These results indicated that the *TRAIL* + Dox synergistic antitumor effect could be achieved by PDT, which paves the way to gene-drug combination therapy for cancer.

Keywords: multifunctional, gene therapy, chemotherapy, *TRAIL*, one-step assembly strategy, CAIR theory

Introduction

Mutation and inactivation of tumor-suppressor genes are important pathogenesises during the development of tumors.^{1,2} Advances in understanding and manipulating genes have set the stage to prevent or treat cancer by altering human genetic material.^{3,4} Even though gene therapy has been increasingly applied in clinical trials for cancer treatment over the past decade, its clinical applications have not been growing as expected.^{5,6} Till now, only two cancer-gene therapies have been approved in China, and none has been approved by the US Food and Drug Administration. The limited clinical outcome of gene therapy might be due to tumor heterogeneity and drug resistance. Combination therapies using multiple therapeutic strategies can synergistically

Correspondence: Na Zhang
Department of Pharmaceutics, Key Laboratory of Chemical Biology (Ministry of Education), School of Pharmaceutical Sciences, Shandong University, 44 West Culture Road, Jinan, Shandong 250012, China
Tel +86 531 8838 2015
Fax +86 531 8838 2548
Email zhangnancy9@sdu.edu.cn

elevate antitumor effects with lowering doses of each agent, hence reducing side effects.⁷ Recently, the combination of chemotherapy and gene therapy has attracted a lot of attention with enhanced therapeutic effects of drugs and transfection efficiency of genes.^{1,8}

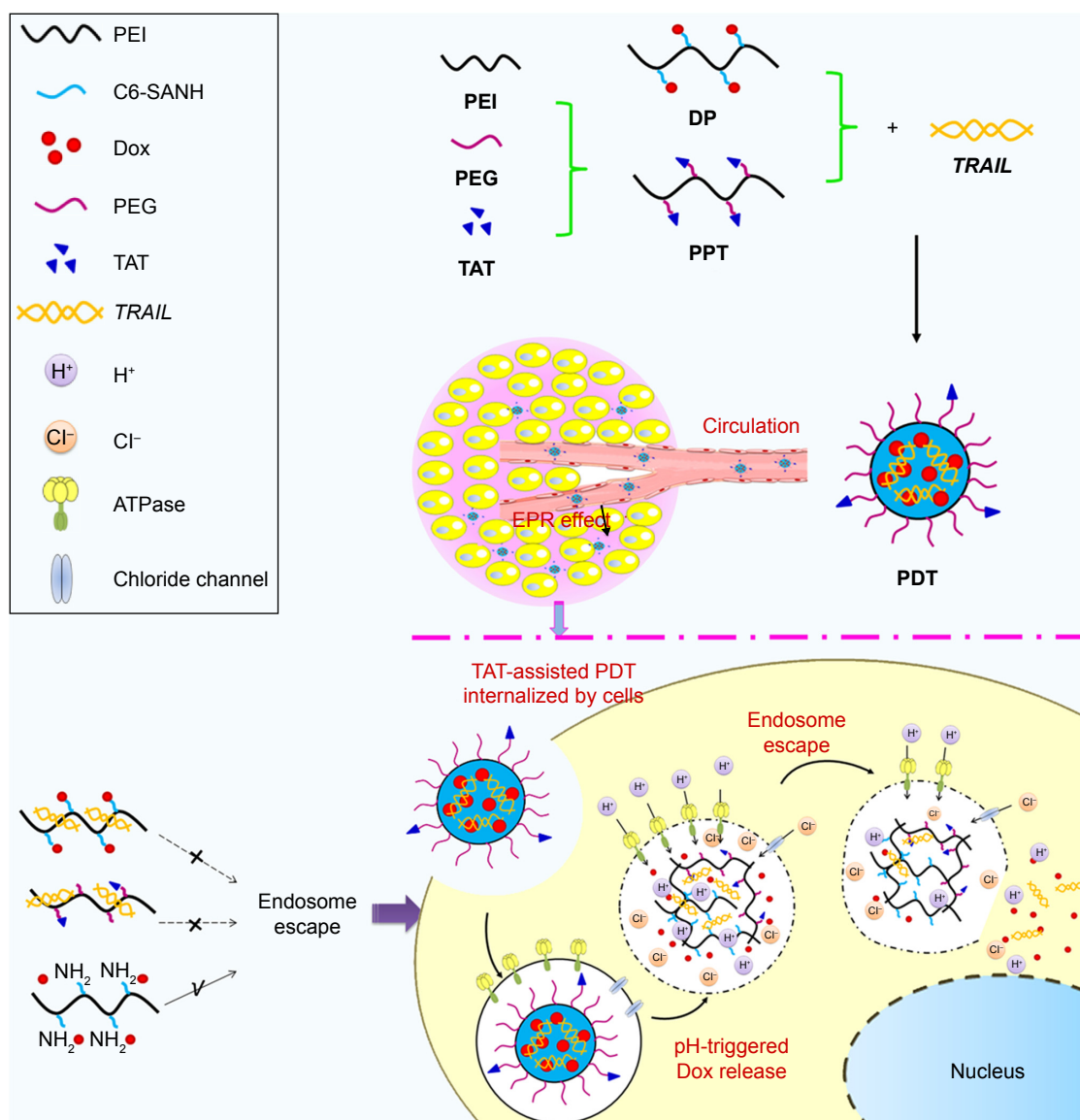
The human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene (*TRAIL*) is in the TNF superfamily of related proteins and can induce apoptosis in cancer cells instead of most normal cells.⁹ *TRAIL* has shown impressive anticancer activity in preclinical models, and this promise has promoted the application of *TRAIL* in clinical trials.^{10,11} To enhance *TRAIL*-induced apoptotic potency, combination therapies of *TRAIL* and chemotherapeutics have been being studied in recent years.^{12,13} Doxorubicin (Dox), which has been used clinically to treat several kinds of tumors in clinical settings, was selected as the model chemotherapeutic in this work.¹⁴ There are several advantages to the Dox and *TRAIL* combination. *TRAIL*-induced apoptosis requires clustering of DR5 within ceramide-enriched membrane platforms.¹⁰ DR5 clustering can facilitate recruitment of the adaptor protein FADD and pro-caspase 8 and transactivation of caspase 8, leading to induction of apoptosis.^{15,16} Triggered release of ceramide in plasma membrane or addition of exogenous ceramide can both mediate the formation of ceramide-enriched membrane platforms and enhance apoptosis in cancer cells.¹⁷ The potential of Dox to promote ceramide production has been identified by Vitovski et al.¹⁸ Dox enhances *TRAIL*-induced apoptosis in cancer cells via ceramide-enriched membrane platforms. Dox has been identified to have the greatest potential for sensitization of tumor cells to *TRAIL* through upregulation and activation of DR4 and DR5.¹⁹ Combining short-term *TRAIL* treatment with Dox prevents the development of *TRAIL*-resistant cells.^{20,21}

The rate-limiting step in successful gene–drug combination therapy is the preparation of highly efficient nanocarriers.^{22,23} Nanocarrier-based gene–drug combination therapy can be classified into three groups: general chemotherapy combined with gene-carrying nanocarrier, use of separate nanocarriers for chemotherapy and gene therapy, and codelivery of chemical and gene therapy within a single nanocarrier.²² Among these three strategies, gene–drug combination therapy using a single nanocarrier is highly effective, due to its ability to affect multiple disease pathways in the same tumor cell.²³ For example, Wang et al²⁵ prepared a hyaluronic acid-decorated polyethylenimine (PEI)–poly(lactide-*co*-glycolic acid) nanoparticle system for targeted codelivery of Dox and miR5423p. The nanoparticles increased cytotoxicity

in MDA-MB231 cells and further promoted triple-negative breast cancer-cell apoptosis via activating p53 and inhibiting surviving expression. Xu et al²⁶ prepared double-walled microspheres consisting of a poly(lactide-*co*-glycolic acid) core surrounded by a polylactic acid shell to achieve the codelivery of “chi-p53” and Dox, resulting in a combined antitumor effect.

This codelivery of gene and chemotherapeutic drug was achieved by developing complex nanocarriers with various materials. Nowadays, one of the main challenges for preparation of gene–drug coloaded nanocarriers is how to make the preparation process more simplified and controllable.²⁶ In this project, one-step assembly nanocarriers were designed by conjugating a chemotherapeutic drug to condensed gene material. For gene delivery, PEI has attracted much attention for its high transfection efficiency and multifunctional modification ability.^{27,28} For example, Yang et al³⁰ prepared nanohydrogels comprising carboxymethylcellulose complexed with cationic branched PEI (bPEI) as gene-delivery vehicles. The gene-delivery efficacy showed that the carboxymethylcellulose nanogel complexed with bPEI showed high uptake and gene-transfection ability. Therefore, PEI, which can both achieve gene condensation and carriage of chemotherapeutic drugs, was selected as the skeleton material in this project.

A gene–drug coloaded nanocarrier delivering its cargo efficiently to the cytoplasm of cancer cells in solid tumors must go through a cascade of four steps: long circulation in the blood (C), accumulation in the tumor via the enhanced permeability and retention effect (A), internalization by tumor cells (I), and intracellular drug release (R), collectively referred to as the CAIR theory. As such, a nanocarrier achieves high delivery and therapeutic efficacy only if it efficiently meets all conditions of the CAIR theory.³⁰ Therefore, to develop PEI-based nanocarriers in view of the CAIR theory, multiple elements should be modified in the skeleton material (Scheme 1): in blood circulation, polyethylene glycolation (PEGylation) can be used to make nanocarriers able to escape extracellular nucleases, evade the reticuloendothelial system, and avoid aspecific interaction with blood cells;^{31–34} to increase accumulation at tumor sites, suitable particle sizes are necessary³⁰ (nanocarriers around 100 nm have longer blood-circulation times and better tumor accumulation than smaller particles); and cell-penetrating peptides, which can improve translocation across the plasma membrane and increase intracellular drug concentration, are often used to increase internalization of payloads by tumor cells.³² TAT peptide, a classical cationic cell-penetrating peptide,



Scheme 1 Preparation and efficient delivery process of TRAIL and Dox by PDT based on CAIR theory.

Abbreviations: Dox, doxorubicin; PDT, PEI-PEG-TAT/Dox-PEI/TRAIL; CAIR, circulation, accumulation, internalization, and release; PEI, polyethyleneimine; SANH, succinimidyl 6-hydrazinonicotinate acetone hydrazide; PEG, polyethylene glycol; ATPase, adenosine triphosphatase; DP, Dox-PEI; PPT, PEI-PEG-TAT; EPR, enhanced permeability and retention.

is known to facilitate and enhance cellular internalization efficiently in a large variety of electrostatically or covalently bound cargoes in a nontoxic fashion.^{34,35} For example, Koren et al³⁷ prepared pH-sensitive PEGylated liposomes modified with TAT peptide, which showed increased cytotoxicity with exposed TAT (lower pH values) than shielded TAT (higher pH values). Peng et al³⁸ synthesized TAT peptide conjugated novel cationic metal nanoparticles as highly efficient carriers for gene delivery to stem cells. Therefore, the TAT peptide was selected to improve gene and drug translocation across the plasma membrane in this project. Finally, effective intracellular gene and drug release can be achieved by pH-sensitive

chemical bonds^{38,39} and endosomal escape materials.^{40,41} To achieve the pH-responsive release of the chemotherapeutic drug Dox, Dox-PEI (DP) conjugates were prepared through hydrazone bonds based on our group's previous work.⁴³ The pH-buffering ability of PEI can be used to achieve endosomal escape of cargoes through its "proton sponge" effect. As shown in Scheme 1, there are only free amino groups on PEI, and endosomal escape is achieved. The programmable release of Dox and gene via an endosomal pH-triggered mechanism has been verified in previous work.⁴² In short, Dox can be released from a Dox-gene co-loaded nanocarrier rapidly, while the gene is protected from degradation under

acidic endosomal conditions. Furthermore, due to the proton-sponge effect of free PEI, the gene and Dox are released into the cytoplasm.

In this project, PEI-based intracellular assistant material (PEI-PEG-TAT [PPT]) and pH-sensitive material (DP) were synthesized and characterized by ^1H nuclear magnetic resonance (NMR) spectroscopy. Then, PPT and DP of ideal properties were used to condense *TRAIL* to form Dox-*TRAIL* coloaded PPT-DP-*TRAIL* nanocarriers (PDT). Morphology, particle-size distribution, and ζ -potential of PDT were investigated. Cell-uptake testing was performed in HEPG2 and SKOV3 cells to verify the intracellular uptake-assistance ability of PPT. The expression of the *TRAIL* protein of PDT was investigated by Western blot assay. Finally, the synergistic antitumor effect of *TRAIL*-Dox coloaded PDT was confirmed by in vitro cytotoxicity testing and in vivo antitumor evaluation.

Materials and methods

Materials

Branched PEI (molecular weight [MW] 25 kDa) was purchased from Sigma-Aldrich (St Louis, MO, USA). C6-S-HyNic (C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone [C6-SANH]) was purchased from Solulink (San Diego, CA, USA). Dox was purchased from Meilun Biology Technology (Dalian, China). Bifunctional PEG derivatives with a maleimide and *N*-hydroxysuccinimide ester group (Mal-PEG-NHS; MW 2 kDa) were purchased from Kaizheng Biotech Development (Beijing, China). TAT with a cysteine on the C-terminal (sequence YGRKKRRQRRRC) was purchased from and synthesized by Shanghai Apeptide. MTT was purchased from Sigma-Aldrich. GoldView was purchased from Saibaisheng Biological Engineering (Beijing, China). Lipofectamine 2000 was purchased from Thermo Fisher Scientific (Waltham, MA, USA). All other reagents were of commercial special grade and used without further purification. Human *TRAIL* under an hEF1 α promoter and EGFP under a CMV promoter plasmid were purchased from Yingrun Biotechnology (Changsha, China). Murine *TRAIL* under a CMV promoter plasmid was purchased from Vigene Biosciences (Rockville, MD, USA).

Cell culture

HEPG2 and SKOV3 cells, both purchased from the Chinese Academy of Sciences (Shanghai, China), were kindly provided by the Institute of Immunopharmacology and Immunotherapy of Shandong University (Jinan, China). They were both cultured in DMEM supplemented with 10% FBS at

37°C with 5% CO_2 . The sensitivity of *TRAIL* on HEPG2 and SKOV3 cells has been identified before.⁴⁴⁻⁴⁷

Synthesis of multifunctional materials

Intracellular pH-sensitive cationic DP was synthesized and characterized by ^1H NMR spectroscopy (Avance DPX-300; Bruker, Billerica, MA, USA) as per our previous work.²⁹ C6-SANH-PEI was the intermediate product in the DP-synthesis process, which was also characterized by ^1H NMR spectroscopy in our previous work. Synthesis of the cellular internalization-assistant material PPT is depicted in Figure 1. First, PEI-PEG was synthesized through amide reaction between the NHS ester group of PEG and the amino groups of PEI. Briefly, PEI and NHS-PEG-Mal (2 kDa) were dissolved in dimethyl sulfoxide 20 mL in a 50 mL glass flask under stirring. Then, 20 μL TEA was added to the reaction solution and stirred for 48 hours at room temperature. The product obtained was purified by dialysis against distilled water (MW cutoff [MWCO] 8,000–14,000 Da) for 48 hours and lyophilized. The chemical structure of the PEI-PEG was confirmed by ^1H NMR spectroscopy.

Then, PPT was synthesized. Briefly, TAT and PEI-PEG were dissolved in 1 mM EDTA containing PBS buffer (20 mL) and reacted for 48 hours in a nitrogen atmosphere. TAT was coupled to the end of PEI-PEG via the reaction between the Mal residues of the PEG and sulfhydryl groups of TAT. The target product was purified by dialysis against distilled water (MWCO 8,000–14,000 Da) and then lyophilized. The chemical structure of PPT was also confirmed by ^1H NMR.

Agarose gel-electrophoresis assay

The condensation abilities of DP-*TRAIL* (DT), C6-SANH-PEI/*TRAIL* (CT), and PPT-*TRAIL* were analyzed with 0.8% agarose gel. To obtain specified weight ratios of multifunctional materials, *TRAIL* was added to different concentrations of DP, C6-SANH-PEI, and PPT solutions at equal volumes under vortexing (Vortex 5; Kylin-Bell Lab Instruments, Jiangsu, China). The mixed solutions were incubated at room temperature for 30 minutes to form DP-*TRAIL*, CT, and PPT-*TRAIL* complexes.

Briefly, the concentration of *TRAIL* was fixed at 80 $\mu\text{g/mL}$, and DT, CT, and PPT-*TRAIL* solutions were all prepared at different mass ratios (25:128, 25:64, 25:32, 25:16, 25:8, 25:4, and 25:2). The obtained DT, CT, and PPT-*TRAIL* complexes with different mass ratios were mixed with an appropriate amount of 6 $\times$ loading buffer and electrophoresed on a 0.8% (w:v) agarose gel containing an appropriate amount of GoldView (Tris-acetate-EDTA buffer, 90 V,

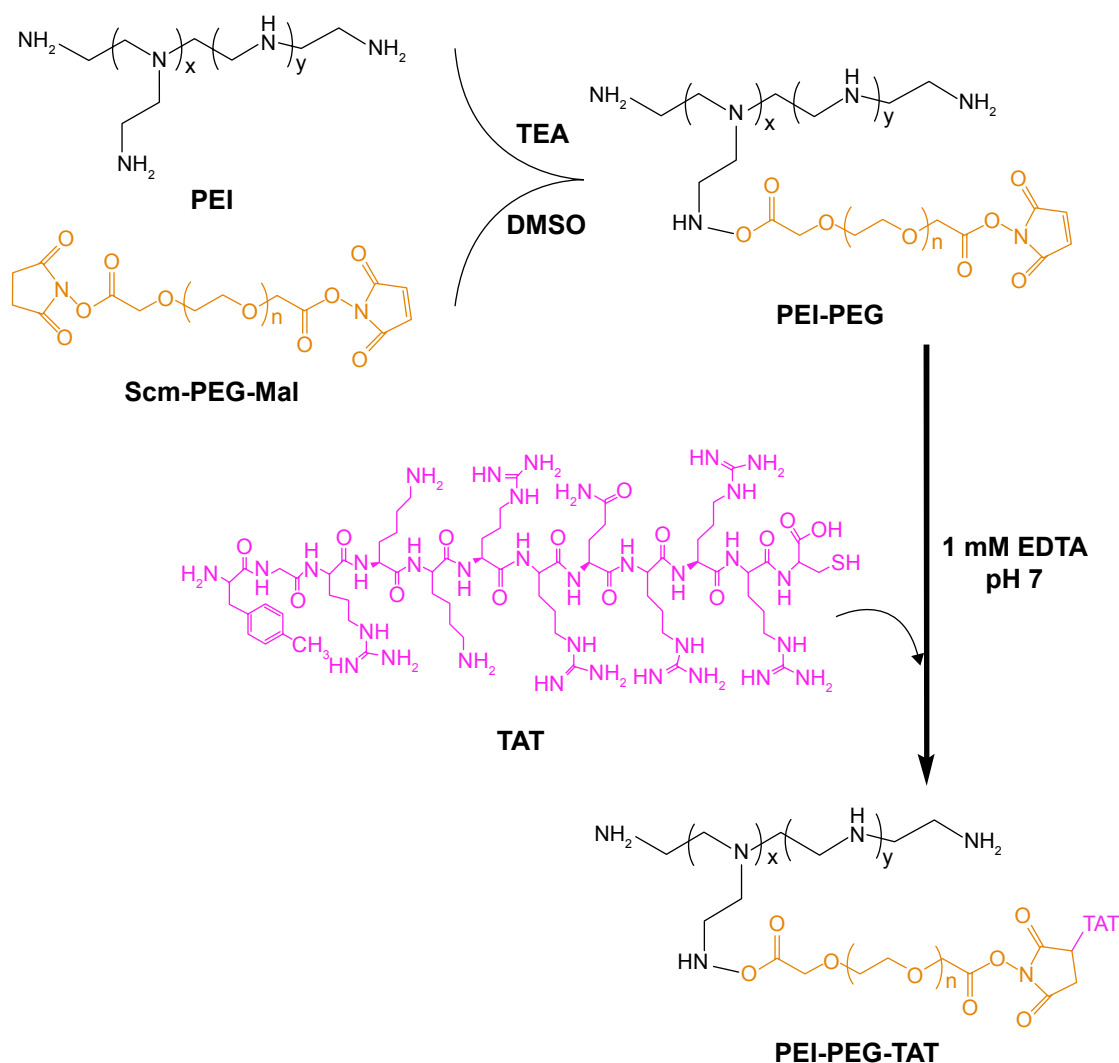


Figure 1 Synthesis route of PEI-PEG-TAT.

Abbreviations: PEI, polyethylenimine; PEG, polyethylene glycol; TEA, triethylamine; DMSO, dimethyl sulfoxide; Scm, succinimidyl; Mal, maleimide.

20 minutes). The electrophoretic mobility of DT, CT, and PPT-*TRAIL* was visualized by an ultraviolet transilluminator and a digital imaging system (IS-2200; Alpha Innotech, San Leandro, CA, USA).

In vitro transfection test of DT

Since DP and PPT have similar condense ability, the first step of prescription optimization was the determination of mass ratio between DP and *TRAIL*. DT particle sizes with mass ratios of 25:16, 25:8, 25:4, and 25:2 were measured with a nanoparticle analyzer (DelsaNano S; Beckman Coulter, Brea, CA, USA). The transfection efficiency of DT with suitable particle sizes was further evaluated to determine the optimal preparation.

In the in vitro transfection test, pEGFP-N1 plasmid (pDNA), which contains the gene encoding EGFP, expressed

from a CMV promoter, was used as the reporter gene. Briefly, HEPG2 and SKOV3 cells were seeded in 24-well culture plates at a density of 5×10^4 cells/well. Culture media were removed after cells had reached 80% confluence, and cells were treated with fresh media (without FBS) containing DP-pDNA with different mass ratios (2 μ g pDNA/well) for 4 hours at 37°C in an incubator. Then, the culture media containing DP-pDNA were removed and washed twice with cold PBS and all the cells cultured for another 48 hours with fresh media. At the end of the incubation, cells were washed with PBS three times and cell nuclei labeled by incubating the samples with Hoechst 33342 for 10 minutes. Finally, the cells were washed with PBS three times and 1 mL PBS added to each cell. Dox and GFP expression in cells was observed using inverted fluorescence microscopy (BX40; Olympus, Tokyo, Japan). Fluorescent images were obtained

with a DP70 digital imaging system (Olympus), and the results were analyzed by Olysia BioReport imaging software (version 3.2; Olympus).

Optimization of PPT content in PPT+DP by intracellular uptake test

The second step of prescription optimization was the determination of PPT content in PPT+DP. The optimal mass ratio of DT was fixed at 25:4 according to the in vitro transfection test. The mass ratio of PPT+DP:*TRAIL* was also fixed at 25:4 to keep consistent condensation ability. To select the optimal PPT content in the mixed materials of PPT+DP, PPT+DP mixtures with different PPT content (0, 1%, 5%, 10%, 20%, 30%, 40%, and 50%) were used to condense *TRAIL* (80 µg/mL). Particle sizes of PDT with different PPT contents were characterized with the DelsaNano S.

The internalization abilities of PDT with different PPT contents in HEPG2 and SKOV3 cells were visualized and quantified by fluorescence microscopy (BX40) and flow cytometry, respectively. Briefly, HEPG2 and SKOV3 cells were seeded in 24-well culture plates at a density of 5×10^4 cells per well. Culture media were removed after cells had reached 80% confluence, and cells were treated with fresh media containing PDT with different PPT contents for 4 hours at 37°C in an incubator. Then, culture media containing the tested PDT with different PPT contents were removed and washed twice with cold PBS. Cells were incubated with Hoechst 33342 and washed twice with cold PBS 10 minutes later. Then, cellular uptake of PDT was observed using inverted fluorescence microscopy with the DP70 digital imaging system and the results analyzed with Olysia BioReport. After that, all cells were harvested and washed three times with cold PBS. The fluorescence intensity of the cells was measured using flow cytometry equipped with a 488 nm argon laser for excitation. For each sample, 10,000 events were collected and fluorescence detected. The percentage of positive events was calculated from the events within the gate divided by the total number of events, excluding cell debris. Finally, the transfection efficiency of PDT with optimal PPT content was further evaluated by the method in the “In vitro transfection test of DT” section.

Characterization of DT, PDT, PCT, and PCD

To establish the synergistic effect of *TRAIL* and Dox on PDT, DT, PPT-CT (PCT), and PPT/C6-SANH-PEI/pDNA (PCD) were prepared and used as controls in the subsequent

experiments. Particle sizes and ζ -potential of DT, PDT, PCT, and PCD were analyzed with the DelsaNano S. The morphology of DT, PDT, PCT, and PCD was examined by transmission electron microscopy (JEM-1200EX; JEOL, Tokyo, Japan).

Western blotting assay

The successful expression of the *TRAIL* protein of PDT was the precondition for the exertion of *TRAIL*–Dox combined/synergistic effects, and expression of the *TRAIL* protein of PDT was determined by Western blotting assay. Briefly, HEPG2 cells were seeded in six-well plates at a density of 4×10^5 cells/well. Culture media were removed after cells had reached 80% confluence, and cells were treated with fresh media containing PDT, PCT, PCD, and Dox for 4 hours at 37°C in an incubator, with the Lipofectamine 2000/*TRAIL* as positive control and PCD as negative control (equivalent to 5 µg of *TRAIL* or 1.75 µg of Dox each well). Then, media were removed and supplemented with DMEM with 10% FBS. After incubation for another 48 hours, media were discarded and washed twice with cold PBS. Then, the protein was extracted from the cells in each group. The sodium dodecyl sulfate polyacrylamide-gel electrophoresis was applied to separate the specified proteins, which were transferred to polyvinylidene fluoride membranes. After that, membranes were blocked by 5% skimmed milk and incubated with the primary antibodies anti-*TRAIL* (Bioss, Beijing, China) and anti- β -actin (Sigma-Aldrich) overnight at 4°C. HRP-conjugated antibodies were incubated with the membranes after washing. An enhanced chemiluminescence kit was used to visualize the bands in darkness, and bands were quantified using the IS-2200 and AlphaEaseFC 4.0 (Alpha Innotech).

In vitro cytotoxicity assays

To evaluate the synergistic effects of Dox and *TRAIL* in PDT, the MTT assay was used to evaluate the in vitro antitumor effects of PDT in HEPG2 and SKOV3 cells. Briefly, HEPG2 and A549 cells were seeded in 96-well plates at a density of 8,000 cells per well and incubated for 24 hours at 37°C. Then, different concentrations of Dox, PCT, PDT, and PCD (equivalent Dox concentrations were 0.01, 0.05, 0.1, 0.5, 1, 2, 4, 6, 8, and 10 µg/mL and *TRAIL* concentrations 0.029, 0.14, 0.29, 1.4, 2.9, 5.8, 11.6, 17.4, 23.2, and 29 µg/mL) were added to five wells each and incubated for 48 hours. Following that, 20 µL of MTT reagents (5 mg/mL) was added to each well and the cells incubated at 37°C for another 4 hours. The media were removed, and formazan crystals

formed by the living cells were dissolved in 200 μ L dimethyl sulfoxide per well. Absorbance at 570 nm of the solution was measured on a microplate reader (model 680; Bio-Rad, Laboratories, Hercules, CA, USA), and all experiments were repeated three times.

In vivo antitumor evaluation

The in vivo synergistic antitumor effect of PDT was proved by using murine *TRAIL* and evaluating the synergistic antitumor effect of murine *TRAIL*–Dox coloaded PPT–DP–murine *TRAIL* nanocarriers (mPDT) on H22 tumor-bearing BALB/c mice. Briefly, H22 tumor-bearing BALB/c female mice were prepared by subcutaneous injection at the right axillary space with 0.1 mL cell suspension containing 10^6 H22 tumor cells. Ten days later, when tumor volumes were approximately 100 mm³, mice were randomly assigned to four treatment groups (n=3 for each group). Then, mice were treated every 2 days with normal saline (NS; intravenous [IV]), Dox solution (Dox 1 mg/kg, IV), mPCT (mouse *TRAIL* 1 mg/kg, IV) and mPDT (Dox 1 mg/kg, mouse *TRAIL* 1 mg/kg, IV). Body weights and tumor volumes were recorded every other day. Tumor volumes were measured with a Vernier caliper every other day using the formula (length $\times$ width²)/2, where length is the longest dimension and width the widest dimension. At day 12, the mice were killed and tumors from each group surgically excised, rinsed with NS, wiped, and photographed. Then, tumor sections (5 mm) were cut from 10% neutral buffered formalin-fixed and paraffin-embedded tissue blocks. Apoptotic cell death in tumor tissue was evaluated histologically by TUNEL staining (in situ cell-death-detection kit; Hoffman-La Roche, Basel, Switzerland) and imaged by fluorescence microscopy (BX4).

Hemolysis assessment

Whole blood from rabbits was collected in heparinized tubes and centrifuged at 3,000 rpm for 4 minutes at 4°C. Then, hemocyte suspensions were washed with NS (5–10 mL) three times and diluted to 2% erythrocyte suspension. The erythrocyte suspension was always freshly prepared and used within 24 hours after collection of DT, PDT, PCT, and PCD (0.1 mL) with the same concentration of in vivo administration were tested. NS and deionized water were used as negative (no hemolysis) and positive (100% hemolysis) controls, respectively. Specimens were incubated at 37°C $\pm$ 0.5°C for 3 hours and centrifuged at 3,000 rpm for 10 minutes before collecting the supernatant. The absorbance value of the hemoglobin released from the erythrocyte cells

was measured at 576 nm. All trials were performed three times. The hemolysis rate was calculated thus:

$$HR = \frac{(A_e - A_n)}{(A_p - A_n)}$$

where A_e , A_n , and A_p are the absorbance value of the experimental group, the negative-control group, and the positive-control group, respectively.

Statistical analysis

All experiments were repeated a minimum of three times and measured at least in triplicate. Results are reported as means $\pm$ SD. Statistical significance was analyzed using Student's *t*-test. Differences between the experimental groups were considered significant when $P < 0.05$.

Results and discussion

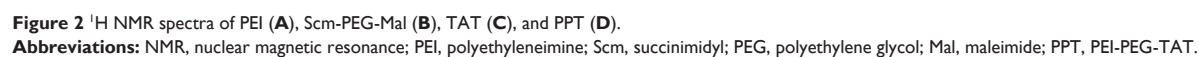
Synthesis of multifunctional materials

Successful synthesis of functionalized materials paved the way to construct desirable nanocarriers. DP and C6-SANH-PEI were synthesized and characterized by ¹H NMR in our previous work.³⁴ In addition, the intracellular pH-sensitivity of DP was confirmed in 0.1 M PBS buffer at pH 7.4 and 5. Dox was released from DP completely after 96 hours at pH 5 and only 27.84% of Dox was released from DP at pH 7.4.

¹H NMR spectra of PEI, succinimidyl (Scm)–PEG–Mal, TAT, and PPT are shown in Figure 2. The peaks (Figure 2D) at 2.2–3.2 ppm and 3.2–4 ppm were related to PEI and SCM-PEG–Mal, respectively, which indicated that SCM-PEG–Mal was successfully conjugated to PEI. Moreover, the successful conjugation of TAT to PEI–PEG was also confirmed by the chemical shifts belonging to TAT ($-\text{CH}_2-$, 1.6–1.9 ppm; aromatic protons of TAT, 6.8–7.3 ppm) in Figure 2D. These results indicated that the cellular internalization-assistant material PPT was successfully synthesized. The integration ratio of PEI (3.2–4 ppm), PEG (2.2–3.2 ppm), and TAT (6.8–7.3 ppm) indicated that the molar ratio among PEI, PEG, and TAT was approximately 1:95:2. The modification ratio of PEG:PEI was enough for long circulation of PDT in blood.

Agarose gel-electrophoresis assay

The condensation ability of DP, C6-SANH-PEI, and PPT was analyzed by 0.8% agarose gel. As shown in Figure 3, *TRAIL* was completely condensed by DP, C6-SANH-PEI, and PPT when mass ratios (*TRAIL*) were up to 100:64, which indicated the condensation abilities of DP, C6-SANH-PEI, and PPT to



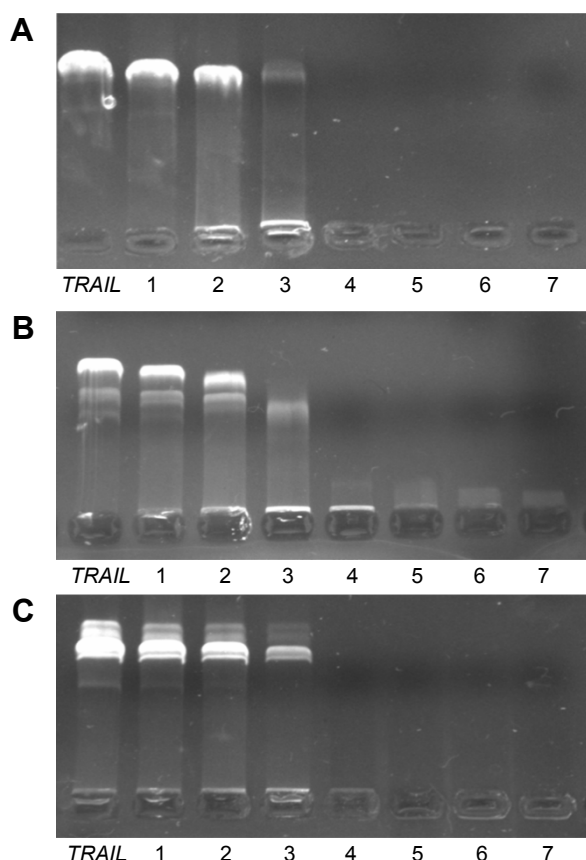


Figure 3 TRAIL-retardation assays for DP-TRAIL (A), C6-SANH-PEI/TRAIL (B), and PPT-TRAIL (C).

Abbreviations: DP, doxorubicin-PEI; SANH, succinimidyl 6-hydrazinonicotinate acetone hydrazine; PEI, polyethyleneimine; PPT, PEI-polyethylene glycol-TAT; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

condense TRAIL were similar. Therefore, DP, C6-SANH-PEI, and PPT could be mixed at any ratio to condense TRAIL.

In vitro transfection test of DT

As shown in Table 1, particle sizes of DT were 184.9 ± 5 nm, 85.8 ± 3.9 nm, 81 ± 3.5 nm, and 103.5 ± 2.1 nm when mass ratios of DT were 100:64, 100:32, 100:16, and 100:8, respectively. In view of the toxicity of PEI and the multimodal particle-size distribution of DT (100:8), DP:pDNA with mass ratios of 25:16–25:4 were used to perform the in vitro transfection test.

As shown in Figure 4, cells were incubated with Hoechst 33342 at 37°C for 1 hour to label cell nuclei with blue.

Table 1 Particle size of DT prepared under different mass ratios of DP:TRAIL

DP/TRAIL	100:64	100:32	100:16	100:8
Size, nm	184.9 ± 5	85.8 ± 3.9	81 ± 3.5	103.5 ± 2.1
PDI	0.247 ± 0.016	0.386 ± 0.074	0.35 ± 0.037	0.381 ± 0.036
Modality	Unimodal	Unimodal	Unimodal	Multimodal

Note: Data presented as mean $\pm$ standard deviation.

Abbreviations: DT, doxorubicin-polyethyleneimine-TRAIL; DP, doxorubicin-polyethyleneimine; PDI, polydispersity index; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

The transfection efficiency of DP-pDNA enhanced with mass ratios of DP:pDNA increased from 100:64 to 100:16 in both HEPG2 and SKOV3 cells. Meanwhile, the intensity of Dox was also enhanced with mass ratios of DP/pDNA increased in both HEPG2 and SKOV3 cells, which might have been due to the increased concentration of DP in DP-pDNA. When Dox and pDNA were merged with the nucleus, the white color (merge of red, green, and blue) meant both Dox and transcribed pDNA were in the nucleus, which means both were released from the DP-pDNA after 48-hour incubation.

Optimization of PPT in PPT+DP by cell-uptake test

Particle size is an important factor that affects the in vivo distribution of nanocarriers. With mass ratio of PPT+DP:TRAIL fixed at 25:4, PDT with different PPT contents was analyzed with the DelsaNano S. As shown in Figure 5, PDT particle sizes were about 80 nm, with PPT content of 0%–30%, and there were no significant differences. When the PPT contents in PPT+DP were 40% and 50%, particle sizes were around 100 nm. To confirm the optimal PPT contents in PDT, further investigation would be needed.

HEPG2 and SKOV3 cells were chosen to determine the cellular uptake of PDT with different PPT contents and select the optimal PPT contents in PDT. As shown in Figure 6 (HEPG2 cells) and Figure 7 (SKOV3 cells), cellular uptake of PDT was enhanced, with PPT contents in PDT increased from 0% to 30% ($P < 0.05$). When the PPT contents in PDT increased from 40% to 50%, cellular uptake dropped compared to PDT with 30% PPT contents ($P < 0.05$ vs 30% PPT contents). This result could have been due to the relatively large particles of PDT. For this reason, PDT with 30% PPT content was selected as the optimal formulation.

As shown in Figure 8, inverted fluorescence microscopy was consistent with cell-uptake analysis through flow cytometry: the intensity of Dox enhanced with PPT contents in PDT increased from 0 to 30%. When PPT contents in PDT increased from 40% to 50%, the intensity of Dox dropped compared to PDT with 30% PPT contents. As shown in Figure 9, the results showed that pDNA was able to be transcribed well with DP-PPT-pDNA.

Characterization of DP, PDT, PCT, and PCD

The therapeutic effect of the nanocarriers was significantly affected by the physicochemical characteristics. Particle sizes of DT, PDT, PCT, and PCD were analyzed. As shown in Table 2, DT, PDT, PCT, and PCD had similar average sizes, and average particle size was about 80 nm. The ζ -potentials of DT,

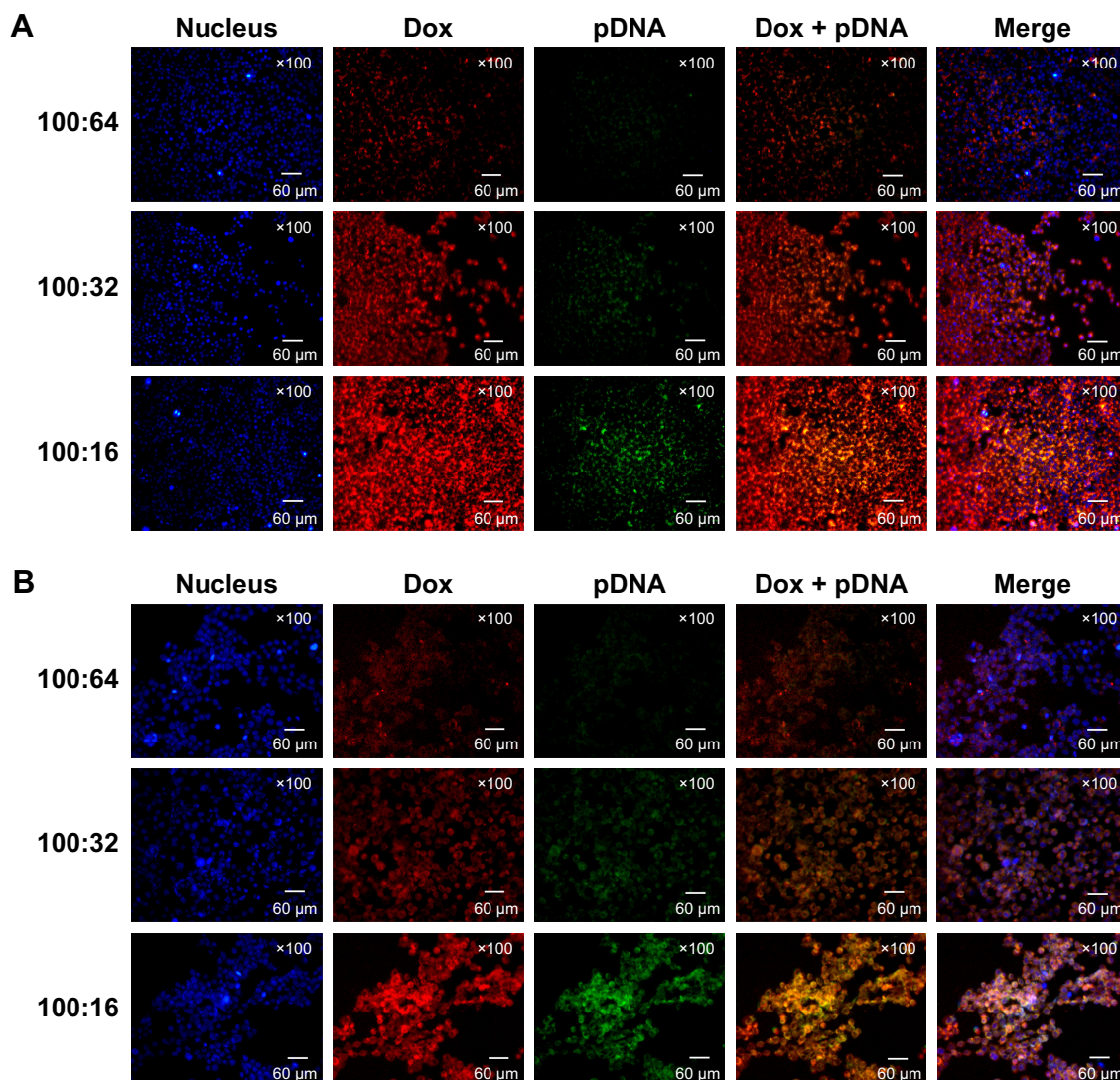


Figure 4 Fluorescence micrography of DP-pDNA complexes in HEPG2 cells (A) and SKOV3 cells (B) following 48-hour incubation at 37°C.

Notes: Green, EGFP transfection; merge column, merged images of nuclei, Dox, and pDNA; yellow, colocalization of Dox and EGFP; white, colocalization of nuclei, Dox, and pDNA.

Abbreviations: Dox, doxorubicin; DP, doxorubicin–polyethyleneimine.

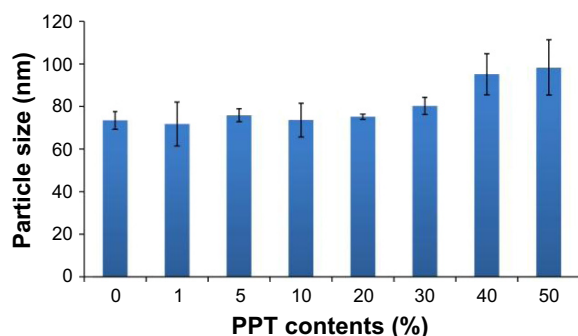


Figure 5 Particle sizes of PDT with different PPT content in mixture materials (PPT+DP).

Abbreviations: PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine–TRAIL; PPT, polyethyleneimine–polyethylene glycol–TAT; DP, doxorubicin–polyethyleneimine; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

PDT, PCT, and PCD were 12.3 ± 1.971 mV, 12.6 ± 0.289 mV, 11 ± 2.11 mV, and 9.33 ± 1.44 mV, respectively. DT, PDT, PCT, and PCD were spherical under optimal conditions, as shown in Figure 10. Therefore, the subsequent evaluations (in vitro Western blot assay and in vitro cytotoxicity assay) would not have been affected by the similar physicochemical characteristics of the nanocarriers.

Western blot assay of TRAIL

The Western blot assay was used to evaluate TRAIL-protein expression after HEPG2 cells had been transfected by PDT, which was the precondition of drug–gene synergistic effects. As shown in Figure 11, TRAIL expression in HEPG2 cells

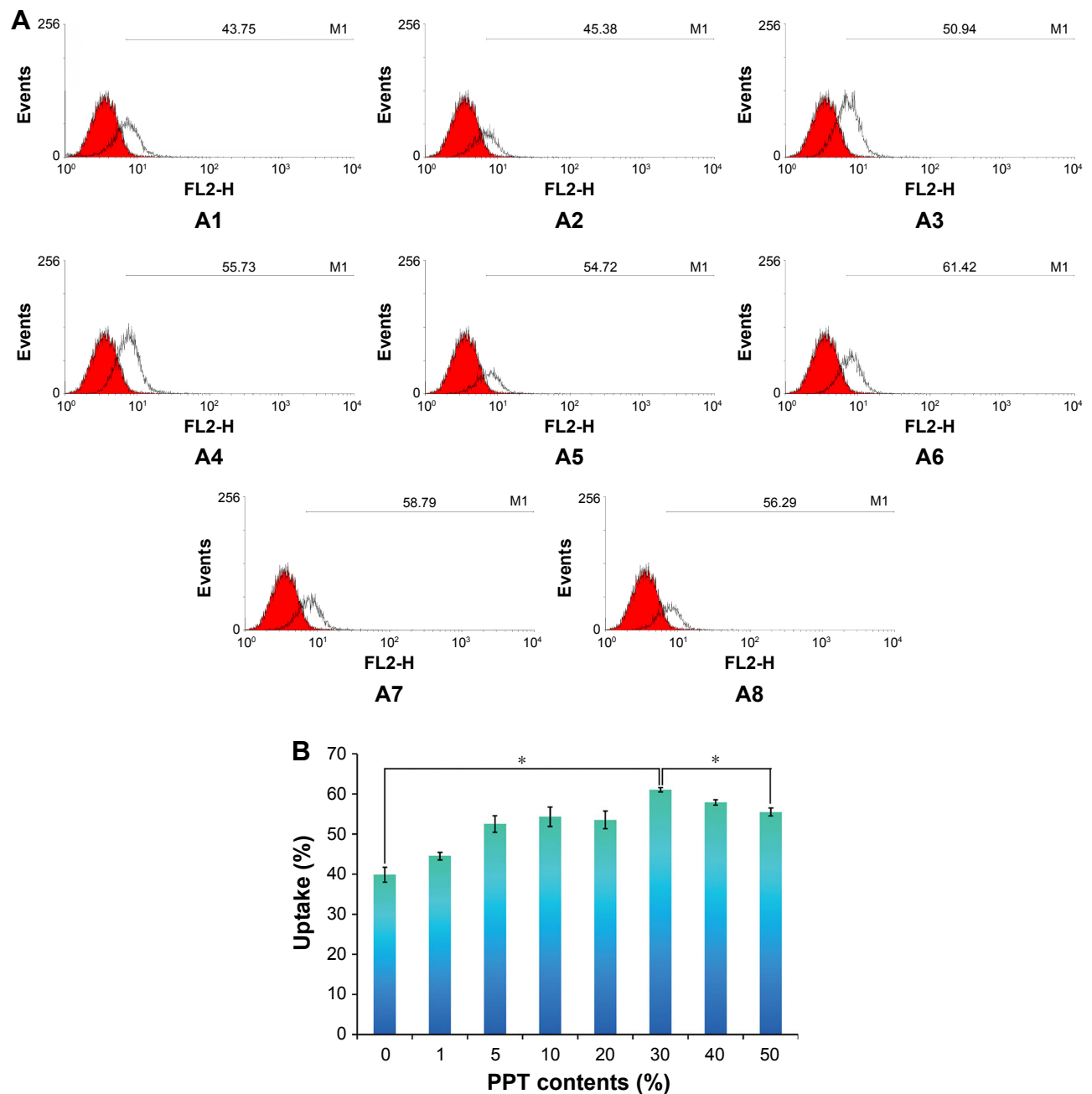


Figure 6 Representative histogram illustrating cellular uptake (**A**) and data analysis of the cellular uptake rates (%) (**B**) according to the results of flow cytometric analysis on HEPG2 cells.

Notes: PDT with different PPT contents of mixture materials (PPT+DP) in HEPG2 cells (* $P < 0.05$ vs 0% PPT contents). PPT content in DP+PPT was 0 (**A1**), 1% (**A2**), 5% (**A3**), 10% (**A4**), 20% (**A5**), 30% (**A6**), 40% (**A7**), and 50% (**A8**).

Abbreviations: PDT, polyethyleneimine-polyethylene glycol-TAT/doxorubicin-polyethyleneimine/TRAIL; PPT, polyethyleneimine-polyethylene glycol-TAT; DP, doxorubicin-polyethyleneimine; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

after treatment with PDT, PCT, PCD, and Dox was evaluated by Western blot assay. Among the four groups, PCD represented PPT/C6-SANH-PEI/pDNA and was a blank nanocarrier without Dox/TRAIL loaded. There was no TRAIL transcription in HEPG2 cells incubated with PCD. PCD was selected as the negative control and Lipofectamine 2000/TRAIL selected as positive control. Compared with PCD, TRAIL-protein expression was upregulated by PDT

and PCT compared with PCD, while TRAIL expression in HEPG2 cells was not upregulated by free Dox or TRAIL. These results indicated that the PDT can be successfully transfected in HEPG2 cells and express TRAIL protein. In this case, the internalized PDT would effectively exert gene-therapy effects. However, the TRAIL expression of Lipofectamine 2000/TRAIL in HEPG2 cells was higher than PDT. For this reason, more efforts should be made

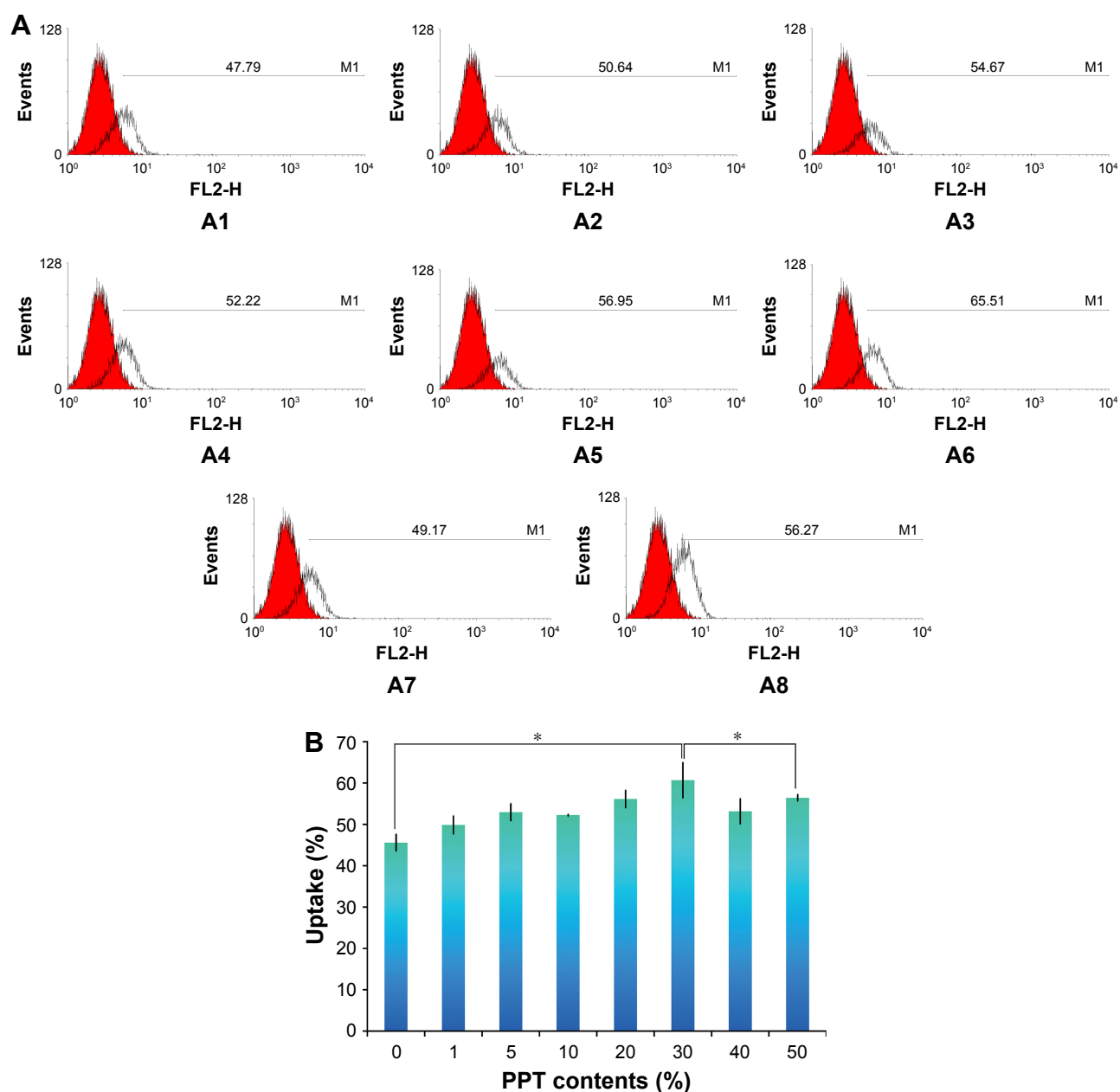


Figure 7 Representative histogram illustrating cellular uptake (**A**) and data analysis of the cellular uptake rates (%) (**B**) according to the results of flow cytometric analysis on SKOV3 cells.

Notes: Cellular uptake of PDT with different PPT contents of mixture materials (PPT+DP) in SKOV3 cells (* $P < 0.05$ vs 0% PPT contents). PPT content in (DP+PPT) was 0 (**A1**), 1% (**A2**), 5% (**A3**), 10% (**A4**), 20% (**A5**), 30% (**A6**), 40% (**A7**), and 50% (**A8**).

Abbreviations: PDT, polyethyleneimine-polyethylene glycol-TAT/doxorubicin-polyethyleneimine/TRAIL; PPT, polyethyleneimine-polyethylene glycol-TAT; DP, doxorubicin-polyethyleneimine; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

to enhance the transfection efficiency of PDT in future research.

In vitro cytotoxicity assay

The in vitro cytotoxicity assay was used to evaluate the in vitro synergistic antitumor effects of PDT. As shown in Figure 12 and Table 3, the antitumor effect of PDT was enhanced compared to free Dox and PCT in both HEPG2 and SKOV3 cells ($P < 0.05$). These results indicated that

enhanced antitumor effects were achieved by PDT. After Dox and TRAIL had been delivered to tumor cells, the cells would be killed by Dox and the expression of TRAIL. Therefore, synergistic antitumor effects of PDT were achieved. Furthermore, after HEPG2 and SKOV3 cells had been incubated with PCD for 48 hours, cell viability was above 80% at different concentrations. This result indicated that the blank nanocarriers PCD did not have obvious cytotoxicity in HEPG2 or SKOV3 cells.

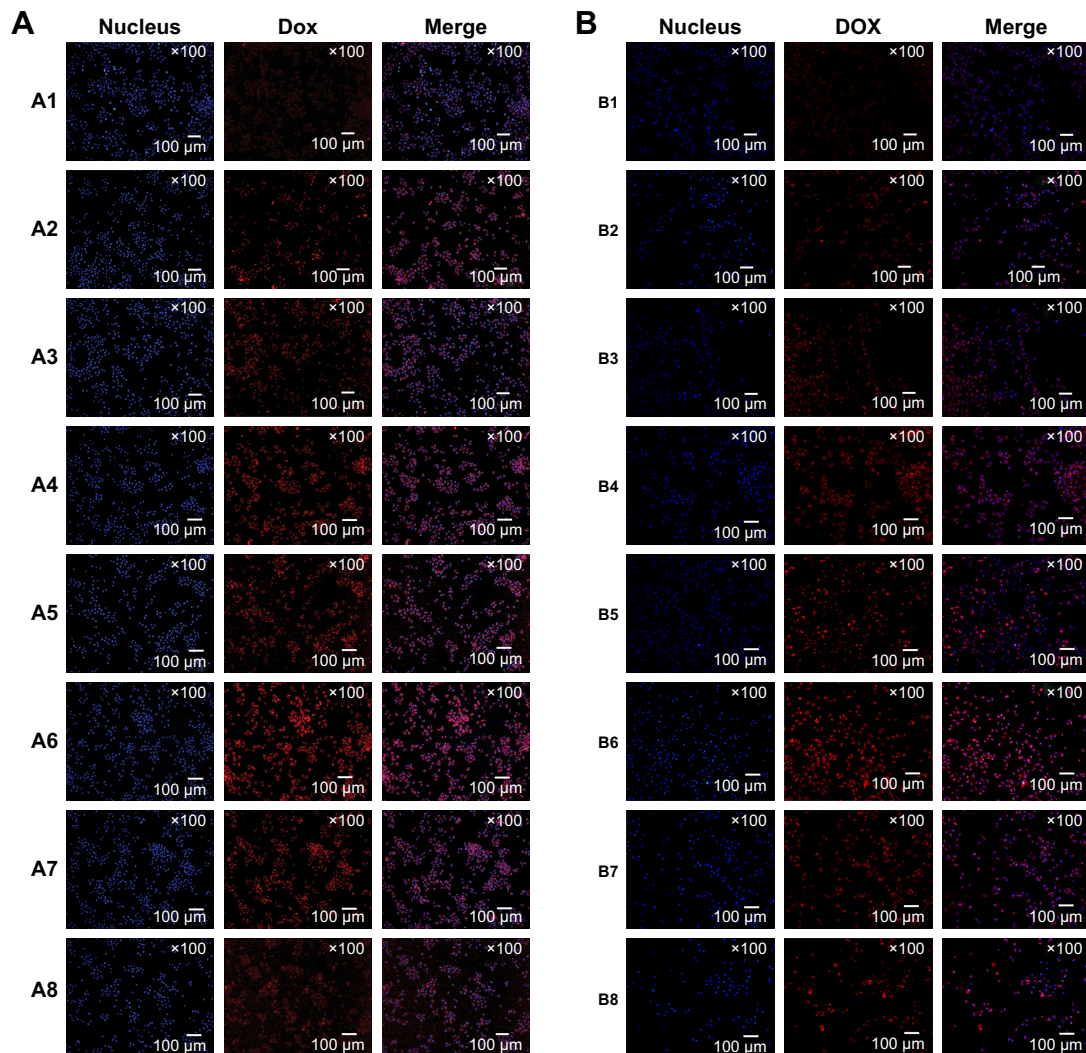


Figure 8 Cellular uptake observed by fluorescence microscopy of PDT with different PPT contents of mixture materials (PPT+DP).

Notes: HEPG2 cells (**A**) and SKOV3 cells (**B**) (* $P < 0.05$ vs PPT contents). PPT content in (DP+PPT) was 0 (**A1**, **B1**), 1% (**A2**, **B2**), 5% (**A3**, **B3**), 10% (**A4**, **B4**), 20% (**A5**, **B5**), 30% (**A6**, **B6**), 40% (**A7**, **B7**), and 50% (**A8**, **B8**).

Abbreviations: PDT, polyethyleneimine-polyethylene glycol-TAT/doxorubicin-polyethyleneimine/TRAIL; PPT, polyethyleneimine-polyethylene glycol-TAT; DP, doxorubicin-polyethyleneimine; Dox, doxorubicin; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

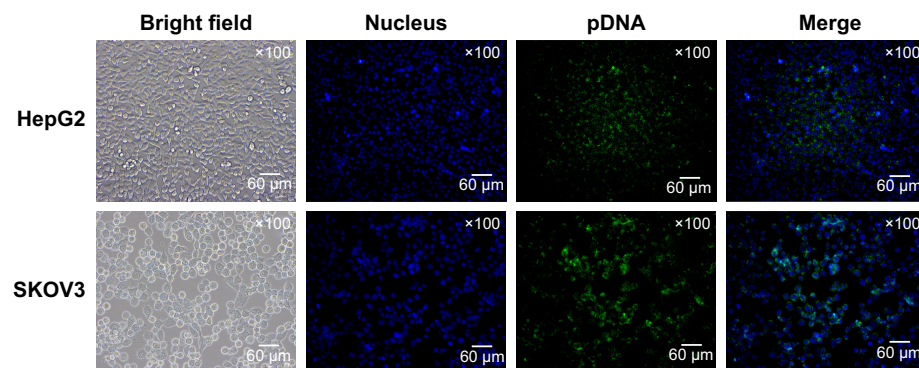


Figure 9 Fluorescent micrographs of DP-PPT-pDNA complexes in HEPG2 and SKOV3 cells following 48-hour incubation at 37°C.

Abbreviations: DP, doxorubicin-polyethyleneimine; PPT, polyethyleneimine-polyethylene glycol-TAT.

Table 2 Particle size and PDI of DT, PDT, PCT, and PCD

Nanoparticles	DT	PDT	PCT	PCD
Size, nm	73.52±3.97	80.29±3.66	87.7±5.49	84.38±6.23
PDI	0.235±0.033	0.210±0.013	0.252±0.020	0.173±0.043
ζ-Potential, mV	12.3±1.971	12.6±0.289	11.0±2.11	9.33±1.44

Note: Data presented as mean ± standard deviation.

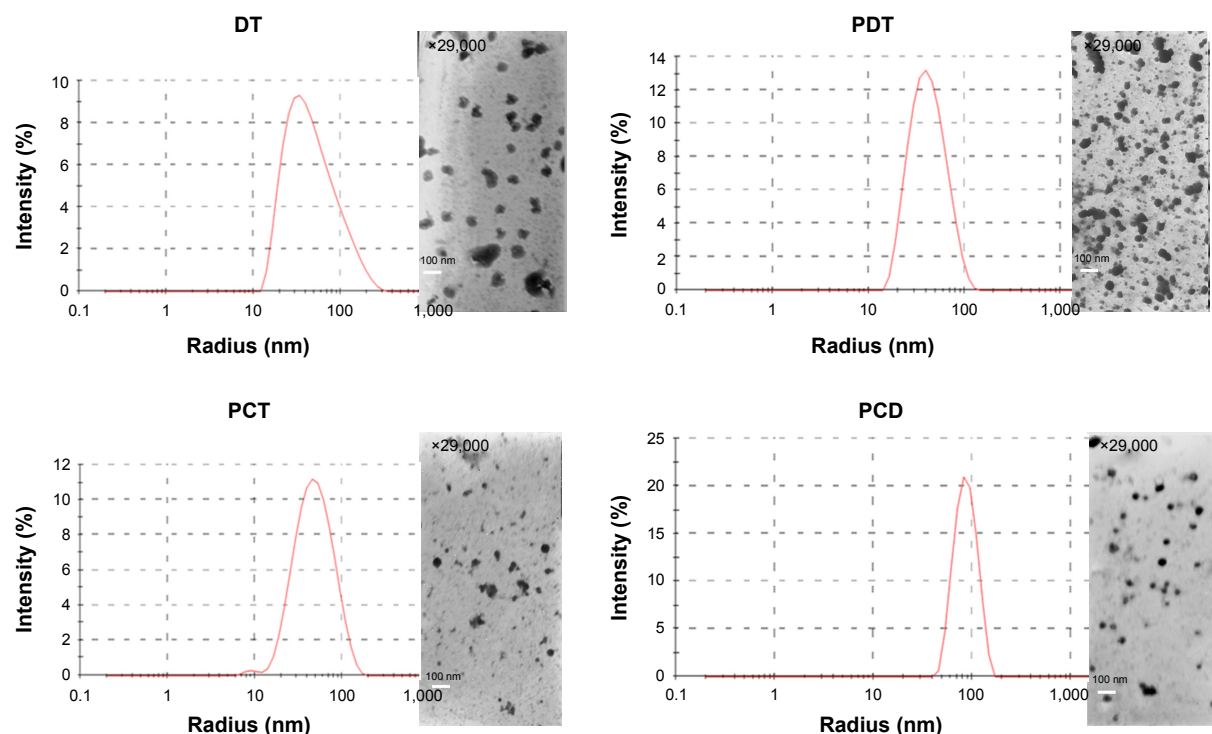
Abbreviations: PDI, polydispersity index; DT, doxorubicin–polyethyleneimine–TRAIL; PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine/TRAIL; PCT, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/TRAIL; PCD, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/DNA; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

In vivo antitumor evaluation

The antitumor efficacy of mPDT was assessed on H22 tumor-bearing BALB/c mice. Changes in body weight and tumor volumes in the mice are shown in Figure 13. After 12 days of treatment, the average tumor volume of mice treated with NS reached about 550 mm³ (Figure 13B), which was larger than that for the Dox ($P<0.05$), mPCT ($P<0.05$), and mPDT groups ($P<0.01$). For the mPDT group, average tumor volume was about 200 mm³, which was smaller than both the Dox group ($P<0.05$) and the mPCT group ($P<0.05$). Moreover, Figure 13C shows a direct visual representation of the tumor-suppression effect, in accordance with Figure 13B. These results indicated that mPDT had a stronger tumor-inhibition effect than Dox and mPCT, and a synergistic antitumor effect of Dox and TRAIL was achieved by mPDT.

Body-weight variations were monitored to monitor the adverse effects of various formulations. As shown in Figure 13A, no obvious body-weight loss induced by Dox, mPCT, or mPDT was observed in comparison with the NS group ($P>0.05$), which indicated that body-weight loss caused by systemic toxicities was not revealed in this test.

We next investigated the effect of treatments on apoptosis in vivo by TUNEL staining of paraffin-embedded sections of the H22-xenografted tumors. As shown in Figure 13D, there were greater degrees of apoptosis in the mPDT, Dox, and mPCT groups compared with the NS group. Either Dox or mPCT caused a modest increase in the number of TUNEL-positive cells (brown color) compared with the NS group. However, mPDT dramatically increased the number of

**Figure 10** Size distribution and transmission electron microscopy.

Abbreviations: DT, doxorubicin–polyethyleneimine–TRAIL; PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine/TRAIL; PCT, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/TRAIL; PCD, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/DNA; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

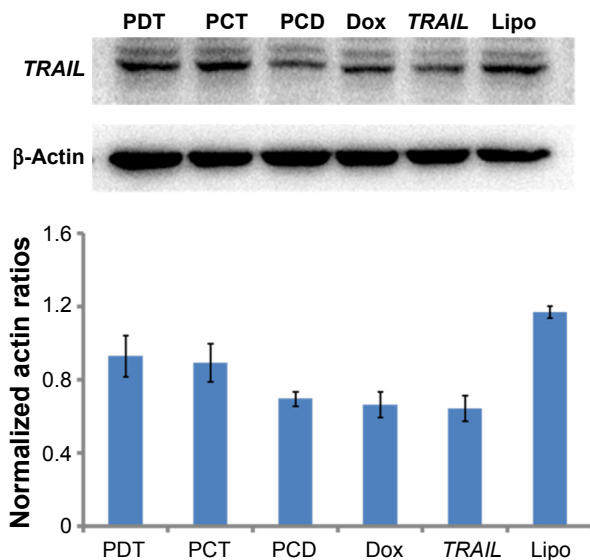


Figure 11 Western blot analysis of TRAIL expression in HEPG2 cells after treatment with PDT, PCT, PCD, Dox, TRAIL, and Lipo for 48 hours.

Abbreviations: PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine/TRAIL; PCT, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/TRAIL; PCD, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/DNA; Dox, doxorubicin; Lipo, Lipofectamine 2000; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

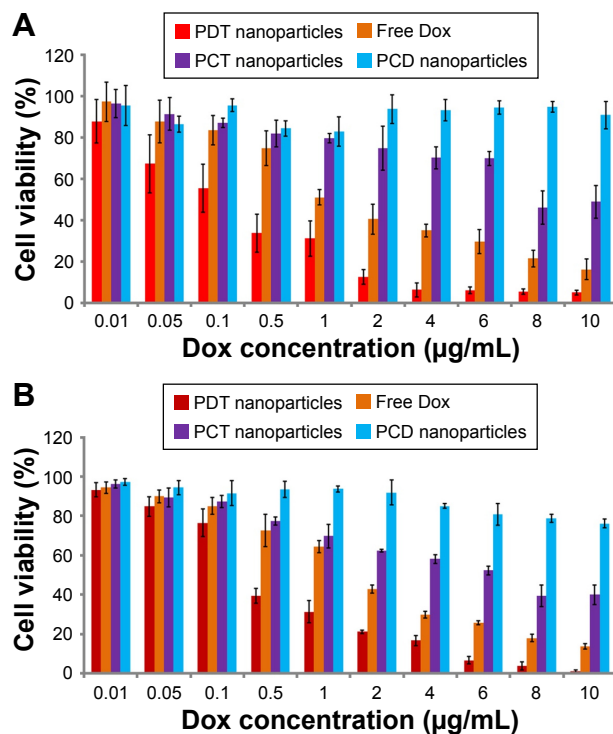


Figure 12 In vitro antitumor effects of free Dox, PCT, PDT and PCD in HEPG2 (A) and SKOV3 (B) cells.

Abbreviations: Dox, doxorubicin; PCT, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/TRAIL; PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine/TRAIL; PCD, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/DNA; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

Table 3 IC₅₀ of HEPG2 and SKOV3 cells incubated with free Dox, PCT, PDT, and PCD at 48 hours

Cell line	Free Dox	PCT	PDT	PCD
HEPG2 (μg/mL)	3.21±0.19**	9.72±4.28*	0.20±0.12	–
SKOV3 (μg/mL)	1.20±0.15**	5.00±0.83**	0.29±0.05	–

Notes: * $P < 0.05$ vs PDT; ** $P < 0.01$ vs PDT. Data presented as mean $\pm$ standard deviation.

Abbreviations: Dox, doxorubicin; PCT, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/TRAIL; PDT, polyethyleneimine–polyethylene glycol–TAT/doxorubicin–polyethyleneimine/TRAIL; PCD, polyethyleneimine–polyethylene glycol–TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone–polyethyleneimine/DNA; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

TUNEL-positive cells compared with the Dox and mPCT groups. These results indicated a significant increase in cell death in the tumors in the mPDT group compared with the Dox and mPCT groups. Taken together, these data further identify that a synergistic antitumor effect of Dox and TRAIL can be achieved by mPDT.

Hemolysis assessment

To determine whether the gene–drug codelivery nanocarriers were safe for IV injection, hemolytic properties were evaluated. No hemolysis or aggregation was observed for DT, PDT, PCT, or PDT after centrifugation. As shown in Table 4, hemolysis rates of DT, PDT, PCT, and PDT were less than 5%. Generally, a hemolysis percentage less than 5% is regarded as nontoxic and safe, so the hemolytic activities of DT, PDT, PCT, and PDT were negligible ($<5\%$). These results indicated that the prepared DT, PDT, PCT, and PDT had good hemocompatibility during the preliminary safety evaluation, and were suitable for IV administration.

Conclusion

In conclusion, Dox–TRAIL coloaded PDT based on one-step assembly was developed to achieve synergistic antitumor effects. The intracellular cationic pH-sensitive cellular assistant PPT and DP, with similar consistency condensation ability, were mixed along to condense the TRAIL by one-step assembly, resulting in the PDT obtained. The cellular uptake of PDT was enhanced with increased PPT contents in PDT ($P < 0.05$), which verified the intracellular uptake-assistance ability of PPT. TRAIL-protein expression was upregulated by PDT when compared with free TRAIL in Western blot assays. Furthermore, the in vitro antitumor effect of PDT was significantly enhanced compared to free Dox and PCT in both HEPG2 and SKOV3 cells ($P < 0.01$). mPDT had stronger tumor-inhibition effects than Dox ($P < 0.05$) and mPCT ($P < 0.05$) in vivo, which showed that the synergistic

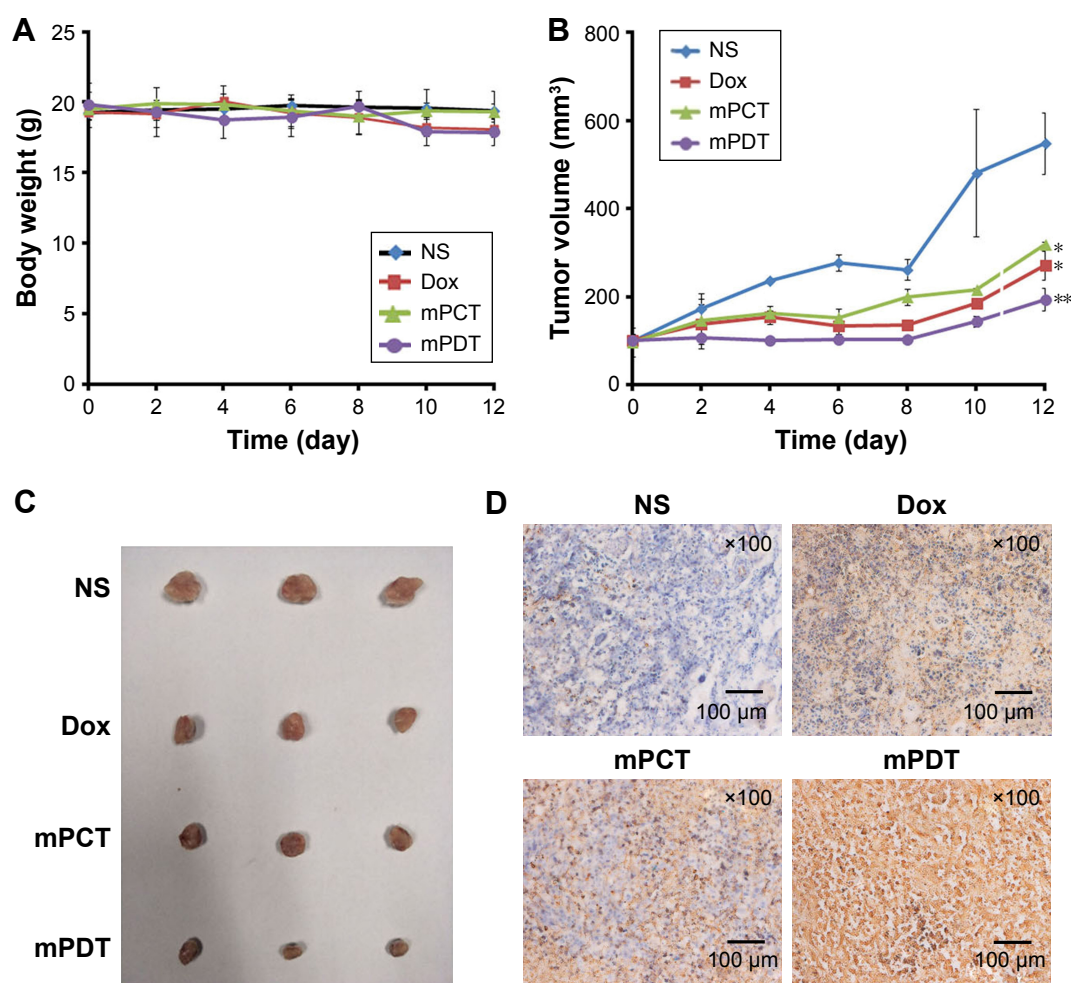


Figure 13 Results of different samples on H22 tumor-bearing mice in vivo antitumor evaluation.

Notes: Body-weight change in each treatment group (A); variation in tumor volume (B); tumors excised from each treatment group on day 12 (C); detection of cell death in xenografted H22 tumors carried out by TUNEL staining (D). Data given as means $\pm$ SD (n=3). * $P < 0.05$, ** $P < 0.01$ compared with the NS group.

Abbreviations: NS, normal saline; Dox, doxorubicin; mPCT, murine polyethyleneimine-polyethylene glycol-TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone-polyethyleneimine/TRAIL; mPDT, murine polyethyleneimine-polyethylene glycol-TAT/doxorubicin-polyethyleneimine/TRAIL; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

antitumor effect of Dox and TRAIL was achieved by mPDT. Therefore, the in vivo synergistic antitumor effect of PDT was identified indirectly. It can thus be said that Dox-TRAIL coloaded PDT holds promising potential as a multifunctional drug-gene codelivery nanocarrier for combination tumor therapy.

Table 4 Hemolysis rates of DT, PDT, PCT and PCD

Nanoparticles	DT	PDT	PCT	PCD
Hemolysis rate (%)	1.73	0	0	0

Abbreviations: DT, doxorubicin-polyethyleneimine-TRAIL; PDT, polyethyleneimine-polyethylene glycol-TAT/doxorubicin-polyethyleneimine/TRAIL; PCT, polyethyleneimine-polyethylene glycol-TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone-polyethyleneimine/TRAIL; PCD, polyethyleneimine-polyethylene glycol-TAT/C6-succinimidyl 6-hydrazinonicotinate acetone hydrazone-polyethyleneimine/DNA; TRAIL, the human tumor necrosis factor-related apoptosis-inducing ligand-encoding plasmid gene.

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Disclosure

The authors report no conflicts of interest in this work.

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