AD

Award Number: W81XWH-FFËEGJI

 $\label{eq:tilde} TITLE: T`|aai' & aai' & a$

PRINCIPAL INVESTIGATOR: ÖLDØD * ÁY ^ã

CONTRACTING ORGANIZATION: Y ậlặæ (ÁT æ)• @ÁÜ 3&^ÁV} ãç^¦•ã: P[˘•đ] Ê√ÝÁÅ Ï €€Í Á

REPORT DATE: June 20FG

TYPE OF REPORT: Øð æ

PREPARED FOR: U.S. Army Medical Research and Materiel Command Fort Detrick, Maryland 21702-5012

DISTRIBUTION STATEMENT: Approved for public release; distribution unlimited

The views, opinions and/or findings contained in this report are those of the author(s) and should not be construed as an official Department of the Army position, policy or decision unless so designated by other documentation.

6					Form Approved	
Public reporting burden for this	collection of information is es	stimated to average 1 hour per re-	sponse, including the time for rev	iewing instructions, sea	arching existing data sources, gathering and maintaining the	
data needed, and completing a this burden to Department of D 4302. Respondents should be valid OMB control number. PI	and reviewing this collection or befense, Washington Headqua aware that notwithstanding a EASE DO NOT RETURN YO	f information. Send comments re arters Services, Directorate for Int ny other provision of Iaw, no pers UR FORM TO THE ABOVE ADI	garding this burden estimate or a formation Operations and Reports on shall be subject to any penalty DRESS.	iny other aspect of this s (0704-0188), 1215 Je y for failing to comply w	collection of information, including suggestions for reducing fferson Davis Highway, Suite 1204, Arlington, VA 22202- ith a collection of information if it does not display a currently	
1. REPORT DATE (DL	D-MM-YYYY)	2. REPORT TYPE		3.	DATES COVERED (From - To)	
4. TITLE AND SUBTIT	LE	Filidi		5a	CONTRACT NUMBER	
Multifunctional Viru	 s-Nanoshell Assei	mbly for Targeted H	vperthermia and Vir	al Gene		
Therapy for Breast	Cancer	nory for rangetourn		51	D. GRANT NUMBER	
				V	V81XWH-11-1-0297	
				50	:. PROGRAM ELEMENT NUMBER	
6. AUTHOR(S)				50	I. PROJECT NUMBER	
Dr. Fang Wei						
				56	. TASK NUMBER	
E-Mail: fwei@rice.edu					. WORK UNIT NUMBER	
7. PERFORMING ORC	SANIZATION NAME(S) AND ADDRESS(ES)		8.	PERFORMING ORGANIZATION REPORT	
William Marsh Rice	University				NUMBER	
Houston, TX 77005						
9. SPONSORING / MONITORING AGENCY NAME(S) AND ADDRESS(ES)				10	. SPONSOR/MONITOR'S ACRONYM(S)	
Fort Detrick, Marv	and 21702-5012					
				11	. SPONSOR/MONITOR'S REPORT	
					NUMBER(S)	
12. DISTRIBUTION / AVAILABILITY STATEMENT						
Approved for Public Release; Distribution Unlimited						
14. ABSTRACT						
vve aimed to develop a virus-Au NS assembly to specifically target breast cancer cells, and to use localized heat of NIR						
radiation of Au NS to destroy breast cancer cells in synergy with gene therapy. We proposed to develop Virus-hanoshell						
successfully completed majority of tasks 1 and 2 of our Statement of Work. Specifically, we have designed and synthesized a						
linker molecule with dual functionalities to be able to covalently attach $\Delta\Delta$ / to $\Delta\mu$ NS surface. Au NS was successfully						
functionalized with the linker molecules. AAV cansid surface was modified with a benzaldehvde functional group. Using higher						
titer AAV8, we were able to confirm and quantify the successful conjugation. Recombinant AAV vectors encoding a gene under						
regulation of a heat-responsive promoter were generated. By combining breast cancer targeting, gene therapy, and						
hyperthermia, the virus-NS assemblies have the potential to greatly reduce the recurrence of cancer, reduce side effects,						
increase patient survival rates, and improve the quality of life of breast cancer patients.						
15. SUBJECT TERMS						
n typermennia, Gen	e ulerapy, virus, A					
16. SECURITY CLASSIFICATION OF:			17. LIMITATION	18. NUMBER	19a. NAME OF RESPONSIBLE PERSON	
			OF ABSTRACT	OF PAGES	USAMRMC	
a. REPORT	b. ABSTRACT	c. THIS PAGE		_	19b. TELEPHONE NUMBER (include area	
	U		UU	9		
L		1	1	1	Standard Form 298 (Rev. 8-98)	

Standard Form 298 (Rev. 8-9 Prescribed by ANSI Std. Z39.18

Table of Contents

Page

Introduction	4
Body	4
Key Research Accomplishments	8
Reportable Outcomes	9
Conclusion	9
References	9

INTRODUCTION

Combination of hyperthermia with gene therapy may be a promising approach for breast cancer treatment. Previous work by others has shown that synergistic effects of hyperthermia-induced gene therapy increase the effects of hyperthermia at both the cellular and systemic levels. Moreover, viral gene delivery vectors delivering cargo driven by various heat-inducible promoters show positive anti-tumor effects with the combination of hyperthermia and gene therapy. We proposed to develop virus-nanoshell assemblies by attaching adeno-associated virus (AAV) to gold nanoshells (Au NS) through chemical bonds. AAV is currently one of the most promising and extensively investigated viral vectors for gene therapy. Au NSs are nanoparticles consisting of a dielectric core surrounded by a thin layer of Au. The unique property of Au NS is the tunability of its plasma resonance. Au NS can accumulate heat upon irradiation with NIR light, which is very useful for biomedical applications because tissues are transparent to NIR. Using NIR irradiation, the Au NS can be induced to generate localized heat in tissues. The generated heat will then activate the expression of delivered suicide genes that are under the regulation of heat-inducible promoters. Synergistic effects of hyperthermia and delivered genes should increase the specific and effective killing of breast cancer cells.

BODY

TASK #1: Generate virus NS assemblies

Au NS was synthesized via the following methods. SiO_2 was functionalized with 3aminopropyltrimethoxysilane (APTMS). The Au colloid was synthesized following Duff method. The functionalized SiO_2 was mixed with Au colloid to form a "seed" with approximately a 30% surface coverage of gold nanoparticles on the SiO_2 surface. The seed solution was mixed with HAuCl₄, and formaldehyde was added as a reducing agent to form a complete gold layer on the SiO_2 . TEM was used to confirm the size and completeness of the Au shell formation on the nanoparticles. UV-VIS was used to evaluate the plasmon resonance. Shown in Figure 1 (a)



insert is the TEM image of the Au NS. The size of the Au NS is about 164 nm in diameter. Shown in Figure 1 (a) is the UV-VIS spectrum of a batch of complete Au NS; the peak at 783 nm is the plasmon resonance of the Au NS. The narrowness of the peak indicates it has a complete Au layer on the surface. The photothermal capability of Au NS was tested by using 600 μ l Au NS solutions with OD (optical density) of 1.4 and absorbance of 783 nm. The solution was heated with 806 nm laser of 4W power and compared with 600 μ l water under the same experimental condition (Figure 1 (b)). In about 5 minutes, the Au NS solution was almost boiling but the temperature of the water only sample increased a few degrees.

Linker molecule polyethylene glycol (PEG) disulfide hydrazine was designed to use the disulfide or thiol group to form Au-S bonds to covalently attach the molecule on Au NS surface. Hydrazine group was used to form hydrazone bond with aldehyde functional group on adeno-associated virus (AAV). Hydrazone bond is stable in physiological condition and labile in acidic condition.

PEG disulfide hydrazine was synthesized using the scheme in Figure 2. First, commercially available PEG1102 was oxidized with iodine in methanol. The reaction mixture was purified with dialysis. Figure 3 shows the MALDI-MS of the starting material PEG1102 and oxidized PEG1102SSPEG1102. The expected mass of starting material is about 3000 Dalton and 6000 Dalton for the disulfide compound. The average mass obtained for the starting is 3121.6 Dalton and 6161.5 Dalton for the product.





Next, the PEG 1102 disulfide with ter-Butyloxycarbonyl (BOC) protection group was reacted with trifluoroacetic acid (TFA) in dichloromethane to remove the BOC group. The successful deprotection was confirmed by proton NMR as shown in Figure 4. The blue spectrum is the starting material (PEG1102), the red spectrum is disulfide PEG with BOC protection group (PEG1102-S-S-1102PEG). Both of them show an intense peak at 1.5 ppm, which is the characteristic peak of proton on t-butyl group. The green spectrum is the deprotected (PEG product disulfide



hydrazine); it showed a complete removal of the peak at 1.5 ppm, which suggested the successful deprotection of BOC group and formation of PEG disulfide hydrazine.

Next, 900 µl of 1mM deprotected disulfide PEG hydrazine was added into 5ml Au NS solution (about

1e9 particles/ml), protected from light, and rocked overnight at room temperature. The functionalized Au NS was purified by centrifugation with 500g for 5 minutes for three times. The synthesis scheme is shown in Figure 5.

To confirm the successful functionalization of Au NS, two tests were performed. First, we tested the stability of the fuctionalized Au NS by mixing Au NS solution with 100 mM NaCl and compared it with a negative control (Au NS only) and a positive control (Au NS with 100 mM NaCl). UV-Vis spectra for the three samples were collected every 10 minutes for 4 hours as shown in Figure 6. The positive control of Au NS with salt showed the absorbance of the Au NS decreased in the four hour period and the NSs were destroyed. In



contrast, the PEG protected NS remained stable over time. It is noteworthy that the absorbance peak of the NS had a slight red shift from 788 nm to 791 nm functionalization, after which is another indication of our successful chemical conjugation of Au NS with PEG hydrazine. Another performed was test we Dvnamic usina Liaht

Scattering (DLS) to confirm the hydrodynamic size of the functionalized Au NS. Au NS has a size of 164 nm; PEG-Au NS is 191 nm, which is 27 nm increases in diameter. It is about the size of the PEG chain, but it also depends on

the conformation of the PEG chain on the NS surface.

Our previous study showed that there are over 1,000 lysine residues on the surface of AAV. Among them, 300 are solvent accessible. By using NHS chemistry, the amine groups of lysine residues on AAV capsid can react with sulfo-S-4FB in PBS buffer to form amide bonds and covalently attach 4FB to the AAV capsid surface as shown in Figure 7. The product (AAV-4FB) has a terminal benzaldehyde group which later on will react with the linker molecule to form hydrazone bonds. The successful conjugation can be confirmed by reacting AAV-4FB with 2hydraxinopyridine as shown in Figure











Figure 8. (a) Colorimetric reaction used to quantify number of 4FB linkers on a AAV capsid. (b) Determination of molar substitution ratio (MSR) of 4FB on AAV capsid by UV Spectra.

8(a). 2-hydraxinopyridine reacts with AAV-4FB to form the compound which absorbs at 350 nm. The molar substitution ratio (MSR) can be estimated by using a colorimetric reaction and calculated to be 108 lysines on



AAV2 L (large loading).



Figure 10. TEM image of (a) AAV8 and (b) AAV8-4FB. Images were negatively stained with uranyl formate.

each capsid as shown in Figure 8 (b).

Since our lab mainly works on serotype AAV2, we had difficulty obtaining higher titer virus which prevented our ability to quantify amount of functional groups on AAV capsid. Toward the end of the project we were able to obtained higher titer AAV8 from our collaborator, which enabled us to quantify the MSR and continue the planned studies.

We performed western blot of AAV8, conjugated AAV8-4FB and compared them with AAV2 as shown in Figure 9. We found that after conjugation, the capsid protein VP2 and VP3

did not change. The conjugation AAV8-4FB has lower concentration of virus due to reaction and dialysis; the signals are not as strong as AAV8. We also used TEM to confirm the integrity of the virus capsid after conjugation as shown in Figure 10. We found that after conjugation, the size of the virus somewhat increased, but the capsid is still assembled.

Several attempts to synthesize AAV NS assembly under acidic conditions were not successful. We found Au NS is not stable under pH 5.0. Some studies have shown that the hydrazone formation reaction can also occur under basic conditions. We tested this by using two different buffers, sodium acetate solution (pH 7.2) and potassium carbonate solution (pH 10.0). 0.5 mM 2-hydraxinopyridine in 100mM sodium acetate or potassium carbonate solutions was made and 1mg/ml Sulfo-S-4FB was added. Then the solutions were incubated at 37 degrees for 30 minutes. Each sample was done in duplicate. The UV spectra were obtained as shown in Figure 11. Me₂S buffer was used as a positive control - it shows an obvious peak at 350 nm which is the indication of hydrazone bond formation. In NaAc buffer at pH 7.2, even though the peak shifted to 342 nm, it is still a large peak. In K_2CO_3 , the hydrazone bond formation was not successful. It appears if we need to have hydrazone bond formation in more basic conditions, NaAc would be a good choice.

TASK #2. Engineer AAV to target breast cancer cells

2c. Generate AAV vectors encoding heat-responsive promoters (HT-Prom) & therapeutic genes for heat-responsive gene therapy.

HSP70Bp Extraction from pGL4-HSP70B-Luc

Routine PCR was performed with two different primer sets (Table 1) in order to establish Ncol and BamHI restriction sites at either end of the HSP70Bp for an oligonucleotide annealing strategy (Figure 12). The resulting two PCR reaction mixes were treated with DPN1, combined, and annealed to construct the sticky- ended HSP70Bp using a set temperature of 95C for 4 minutes followed by temperature ramp decrease to 25°C over about 1

hour on a heat block. Annealing was carried out in TNE buffer (10 mM Tris, pH 7.5-8.0, 50 mM NaCl, 1 mM EDTA). The desired annealed product contained 5' sticky ends: a BamHI site on the upstream 5' end and an Ncol site on the downstream 3' end of the promoter, which includes a Kozak sequence directly upstream of the Ncol site, a sequence necessary for efficient translation in eukaryotic cells.¹ All products from the annealing procedure were also phosphorylated on the 5' ends using T4 Polynucleotide Kinase (New England Biolabs) to ensure phosphorylated sticky ends. Sticky end verification was performed by ligating the insert to itself in a concatemer test (Figure 13). The resulting HSP70Bp insert were ligated to the pAAV-GFP backbone vector.

Restriction Digest of pAAV-CMVp-GFP

The pAAV-GFP vector backbone was prepared by performing a double digest with Ncol and BamHI on pAAV-CMVp-GFP to remove the CMVp along with the Kozak sequence while retaining GFP. The digest also





Figure 12 HSP70 Bp Subcloning stragegy overview (pre-annealing)



7

1kb DNA Ladder HSP70Bp

Figure 13 Concatemer Test of HSP70Bp.

2000

1000-

500

removed the beta globin promoter and human beta globin fragments which were considered unnecessary for successful gene expression of GFP. The vector backbone was gel purified and extracted. The expected backbone size was ~ 6 kb along with three smaller fragments as a result of internal Ncol and BamHI restriction sites in CMVp and Beta globin promoter, respectively (Figures 14 and 15). The large ~6000 kb fragment with the other 3 smaller fragments denotes a successful digestion where largest band was extracted for purification and ligation with the HSP70Bp insert. The resulting vector was treated with calf intestinal phosphatase (New England Biolabs) to prevent re-ligation of the vector backbone.

Construction of pAAV-HSP70Bp-GFP

The resulting pAAV-GFP vector backbone and the HSP70Bp insert were ligated together using a standard ligation reaction with T4 DNA ligase in ligase buffer. A 1:6 and



1:0 (positive control) vector-to-insert ratio were used prepare the construct for transformation. The resulting ligation reactions were transformed into DH5α E.coli cells using heat shock (42C for 45 sec).

Colony Screening

Eight colonies were selected and two were sequenced. The sequencing overlap results for the desired plasmid are shown in Figure 16. The overlapping sequence matched perfectly for the region from the beginning of the HSP70B promoter, at the BamHI site, until midway into the GFP where a constructed sequencing primer attached. Base pair substitutions and insertions were detected upstream of the HSP70B promoter, but should not affect the overall effectiveness of the heat shock response.



KEY RESEARCH ACCOMPLISHMENTS

- Designed and synthesized linker molecule for the AAV NS assembly, purified the product and confirmed the product by MALDI-MS and ¹HNMR.
- Functionalized Au NS with linker molecule confirmed the product with stability test and DLS measurement.
- Conjugated 4FB molecules to AAV capsid surface through amide bonds. Conjugation was confirmed and quantified by UV-VIS spectroscopy. Western blot using B1 antibody and TEM confirmed the integrity of the conjugation.
- · Tested the experimental condition for functionalized Au NS and AAV hydrazone formation. Sodium acetate

solution of pH 7.2 would be a good choice.

Designed and constructed recombinant AAV encoding GFP driven by heat-inducible promoter. Colony
screening shows successful insertion of HSP promoter into the construct.

REPORTABLE OUTCOMES

• "Gene Therapy with Hyperthermia for Targeted Breast Cancer Treatment Using Virus-Nanoshell Assembly" oral presentation at 28th Southern Biomedical Engineering Conference, Houston, Texas, May 6, 2012.

CONCLUSION

We have successfully completed majority of tasks 1 and 2 of our Statement of Work. Specifically, we have designed and synthesized a linker molecule with dual functionalities to be able to covalently attach AAV to Au NS surface. Au NS was successfully functionalized with the linker molecules. AAV capsid surface was modified with a benzaldehyde functional group. Due to difficulties with low virus titers, we had to switch from AAV2 to AAV8 towards the latter part of our project. Using higher titer AAV8, we were able to confirm and quantify the successful conjugation. Recombinant AAV vectors encoding a gene under regulation of a heat-responsive promoter were generated. By combining breast cancer targeting, gene therapy, and hyperthermia, the virus-NS assemblies have the potential to greatly reduce the recurrence of cancer, reduce side effects, increase patient survival rates, and improve the quality of life of breast cancer patients.

REFERENCES

1. Kozak, M. "Point mutations define a sequence flanking the AUG initiator codon that modulates translation by eukaryotic ribosomes." *Cell* 44. 2 (1986): 283-292.