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## EphA2 targeting peptide tethered bio-reducible poly(cystamine bisacrylamide - diamino hexane) for the delivery of therapeutic pCMV-RAE-1 $\gamma$ to pancreatic islets

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### Abstract

The pathogenesis of type-1 diabetes is complicated, and a clear, single mechanism has yet to be identified. Reports have indicated that the activating receptor NKG2D plays an important role in the development of disease. Exploiting a natural phenomenon observed in tumors, plasmid DNA encoding for a soluble ligand to NKG2D (sRAE-1 $\gamma$ ) was isolated and engineered into a plasmid expression system. A polymeric gene delivery system was developed to deliver the soluble RAE-1 plasmid locally to the pancreatic islets for the prevention of type-1 diabetes. The bio-reducible cationic polymer poly(cystamine bisacrylamide – diamino hexane) (p(CBA-DAH)) was modified with poly(ethylene glycol) (PEG) and the targeting peptide CHVLWSTRC, known to target the EphA2 and EphA4 receptors. The PEG serves to improve stability and tissue selectivity, while the peptide will target EphA2 and A4, overexpressed in the pancreatic microvasculature. The targeting polymer Eph-PEG-p(CBA-DAH) shows selective uptake by the target cell line, indicative of the targeting properties that will be seen in systemic administration. Using the delivery system, the therapeutic plasmid can be delivered to the pancreas, reduce interactions between the beta-cells and infiltrating NKG2D positive lymphocytes, and effectively protect beta-cells from autoimmune destruction and prevent type 1 diabetes.

### Keywords

Copolymer; Gene Therapy; Drug Delivery; Biodegradation

### Introduction

Type 1 diabetes is an autoimmune disease in which the beta-cells in the pancreatic islets are destroyed by autoreactive immune cells [1, 2]. Specifically, CD4<sup>+</sup> and CD8<sup>+</sup> T-cells are considered essential in the development of type 1 diabetes [3–5]. Additionally, the interaction of co-stimulatory receptors on T-cells and their interaction with the ligands on beta-cells is thought to be critical for the autoimmune response. The activating receptor

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NKG2D is thought to be directly involved in the progression of type-1 diabetes [6–9]. Retinoic acid early inducible gene-1 (RAE-1) is a tightly regulated high affinity ligand for the NKG2D receptor in mice [10, 11]. The expression of RAE-1 on pancreatic islet beta-cells indicates a close relationship between this receptor and the pathogenesis of type-1 diabetes in the non obese diabetic (NOD) mouse model [12].

It has been shown that the systemic blockage of the NKG2D receptor with monoclonal antibodies can prevent the occurrence of type-1 diabetes in the NOD mouse [13]. However, systemic blockage of the NKG2D receptor is not clinically applicable, as NKG2D has important functions in immunosurveillance, including detection of cancerous and virally infected cells [14–17]. Similarly, by blocking the NKG2D receptor specifically around the pancreas, the beta cells can be protected from cell-mediated cytotoxicity. Specifically, delivering a gene coding for soluble RAE-1 locally to the pancreas will result in the binding and internalization of the NKG2D receptor of infiltrating lymphocytes. This will allow the pancreas to undergo “immune evasion,” and will halt the progression of type-1 diabetes.

The main purpose for designing a targeting gene carrier is to deliver therapeutic genes specifically to a target tissue or target cells, thereby minimizing side effects associated with systemic therapeutics [18]. By using a bioreducible polymer, pDNA will be released upon cell entry due to the difference in reduction potential [19, 20]. Additionally, the degradation of the polymer will reduce cytotoxicity and allow for clearance in vivo [21]. The targeting gene carrier consists of three basic components: a bioreducible cationic backbone, a PEG linker, and a targeting peptide. The cationic polymer is essential in complexing with DNA to form polyplex, providing protection from endonucleases, and supplying a mechanism for endonuclease escape [22]. By using a bioreducible polymer, pDNA will be released upon cell entry due to the difference in reduction potential [19, 20]. Additionally, the degradation of the polymer will reduce cytotoxicity and allow for clearance in vivo [21]. The addition of a PEG linker will not only act to connect the targeting ligand to the backbone polymer, but will also serve to stabilize polyplex, shield surface charge, and increase circulation time in vivo [23]. The targeting peptide NH<sub>2</sub>-GGGCHVLWSTRC (Eph) will be added to the distal end of the PEG chain. This peptide has been shown to target the receptors EphA2 and A4, which are overexpressed in the pancreatic islet microvasculature [24, 25].

Here, we report the design of an effective polymeric gene delivery system to deliver a plasmid coding for soluble RAE-1 $\gamma$  to the pancreatic islet microvasculature for the prevention of type-1 diabetes.

## Materials and Methods

### Materials

N-Boc-1,6-diaminohexane, N-Boc-DAH) was purchased from Alfa Aesar (Ward Hill, MA). Hyperbranched polyethylenimine (bPEI, M<sub>w</sub> 25 kDa), trifluoroacetic acid (TFA), triisobutylsilane (TIS), tithiothreitol (DTT), and 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyltetrazolium bromide (MTT) were purchased from Sigma-Aldrich (St. Louis, MO). N,N'-cystaminebisacrylamide (CBA) was purchased from PolySciences, Inc. (Warrington, PA). Heterobifunctional N-hydroxysuccinimide-poly(ethylene glycol)-maleimide (NHS-PEG-Mal, M<sub>w</sub> 2000) was purchased from JenKem Technology (Allen, TX). The targeting peptide Eph (NH<sub>2</sub>-GGGCHVLWSTRC) was synthesized at the University of Utah Core Peptide Synthesis facility. SYBR Green Nucleic acid stain was purchased from Invitrogen (Carlsbad, CA). Luciferase assay system with reporter lysis buffer was purchased from Promega (Madison, WI). DMEM, phosphate buffered saline, penicillin/streptomycin, Trypsin-like enzyme (TrypLE), Trypan Blue, YOYO-1 nucleic acid dye, and AlexaFluor 647 carboxylic acid, succinimidyl ester were purchased from Invitrogen (Carlsbad, CA).

Fetal bovine serum (FBS) was purchased from HyClone (Logan, UT). Luciferase Assay System and Reporter Lysis Buffer were purchased from Promega (Madison, WI). Superscript III First Strand Synthesis kit, Platinum Taq polymerase, pcDNA3.1/V5-His TOPO TA vector, Purelink PCR purification kit, NuPage Bis-Tris gels, nitrocellulose membranes, and anti-V5-AP antibody were purchased from Invitrogen (Carlsbad, CA). Alkaline Phosphatase development kit was purchased from BioRad (Hercules, CA). PCR primers were purchased from Integrated DNA Technologies (IDT, San Diego, CA). RNeasy Kit and Maxi-prep plasmid purification kit were purchased from Qiagen (Valencia, CA). KpnI and NotI restriction enzymes were obtained from New England Biolabs (Ipswich, MA). All materials and solvents were used as received without further purification.

### **Synthesis of p(CBA-DAH)**

The polymer poly(CBA-DAH) was synthesized as previously reported [26].

### **Conjugation of NHS-PEG-Mal to p(CBA-DAH)**

The conjugation of NHS-PEG-Mal to p(CBA-DAH) was adapted from manufacturer's protocols. The NHS ester in heterobifunctional NHS-PEG-Mal was reacted with the primary amine in p(CBA-DAH). NHS-PEG-Mal (PEG  $M_w$  2000) was dissolved in PBS (pH 7.4), and was added to the p(CBA-DAH) solution drop wise at a 1.5:1 molar ratio and reacted in the dark overnight while stirring. The polymer was collected and purified with dialysis (MWCO 3500) and lyophilized. Conjugation was confirmed with  $^1\text{H-NMR}$  (400 MHz,  $\text{D}_2\text{O}$ ).

### **Thiolation of Eph Peptide**

The thiolation of the Eph peptide was carried out according to the manufacturers protocol. Eph peptide was dissolved in PBS, pH 8.0 + 2.0 mM EDTA. Traut's reagent was dissolved and added with stirring to the peptide solution in a 2:1 molar ratio for one hour. The peptide was then purified with a D-Salt polyacrylamide desalting column (Pierce, Rockford, IL), MWCO 1.8 kDa. Elution was monitored with a nanoDrop ND-1000 spectrophotometer at 280 nm wavelength and characterized via MALDI-ToF.

### **Conjugation of Eph-SH to PEG-p(CBA-DAH)**

The conjugation of Eph-SH to PEG-p(CBA-DAH) was completed in two steps. First, the conjugation of heterobifunctional NHS-PEG-Mal to p(CBA-DAH) was completed as described above. Prior to dialysis and lyophilization, thiolated peptide (Eph-SH) is added to the reaction. The thiolated peptide is dissolved in PBS, pH 7.4, and added drop wise with stirring. The peptide is added at a 1.5 fold molar excess over the amount of NHS-PEG-Mal to ensure adequate conjugation. Conjugation was confirmed with  $^1\text{H-NMR}$  (400 MHz,  $\text{D}_2\text{O}$ ).

### **Characterization of p(CBA-DAH)**

The approximate molecular weight of p(CBA-DAH) was determined by size-exclusion chromatography (SEC). The AKTA FPLC system (Pharmacia) was used equipped with a Superose 12 column. Molecular weight was calculated using the Wyatt Optilab T-Rex and the Minidawn Treos, detecting refractive index and light scattering, respectively. The samples were run in a 0.3 M sodium acetate buffer (pH 5.0).

### **Agarose Gel Electrophoresis**

Polyplex was formed in a HEPES buffer solution (20 mM HEPES, pH 7.4, 5% glucose) at weight:weight ratios ranging from 0.5 to 40. Samples were mixed via pipetting, vortexed at a low speed for 10 seconds, and incubated at room temperature for 30 minutes. Agarose gels (1% w/v) were prepared in Tris-acetate-EDTA (TAE) buffer with SYBR Green Nucleic

Acid Gel stain. The samples were electrophoresed at 100V for 30 minutes. DNA was then visualized with a UV illuminator and analyzed using Gel Documentation Systems.

### Polyplex Size Measurements

Dynamic light scattering (DLS) was used to determine the hydrodynamic radius of polyplexes [27]. Polyplexes were prepared by vortexing 60  $\mu\text{g}$  pDNA in 500  $\mu\text{L}$  HEPES buffer solution with an equal volume of polymer solution to yield the desired w:w ratios.

### DNase Protection Assay

The ability of each polymer to protect plasmid DNA from enzymatic degradation was determined using a modified version of a previously reported method [28].

### Cell lines

The cell lines MS1s, NIH 3T3s, and 293T that were used for experiments were obtained from ATCC (catalog numbers CRL-2279, CRL-1658, and CRL-11268, Manassas, VA). These cells were cultured in DMEM supplemented with 10% FBS and 1% penicillin/streptomycin in a humidified incubator at 37°C with 5% CO<sub>2</sub>. All cell lines used for experiments were between passages 3–9.

### Cellular Uptake Assay

Cellular uptake assay was performed with some modifications as previously described [29]. MS1 cells, NIH 3T3 cells, and 293 cells were seeded at a density of  $2 \times 10^5$  cells/well in a 6-well plate in growth medium and grown to 70–80% confluence prior to transfection. Medium of each well was exchanged for 1 mL fresh serum-free medium, and cells were treated with polyplex solutions (2  $\mu\text{g}$  of YOYO-1 labelled pDNA per well) at 10:1 w:w ratios for 4 h at 37 °C, control was given no treatment. For Eph competition assay, 0.1 mM free Eph peptide was added to the media just prior to transfection. Following incubation, cells were washed two times with ice-cold D-PBS, incubated with 0.4 % trypan blue solution for 5 min, washed with DPBS, and collected via trypsinization. Uptake was examined with the BD FACScan analyzer at a minimum of  $1 \times 10^4$  cells gated per sample. Analysis was performed by using Becton Dickinson CellQuest software.

### Luciferase Transfection and Expression

Cells were plated at an initial density of  $4.0 \times 10^4$  cells/well in 24 well plates. 24 hours following plating, luciferase (pCMV-Luc) was complexed with p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) at indicated w:w ratios in HEPES buffer and incubated for 30 minutes. Media in each well was replaced with 0.5 mL fresh, serum-free media. Polyplex was added to each well (0.5  $\mu\text{g}$  DNA/well), and the cells were incubated for 4 hours at 37°C, 5% CO<sub>2</sub>. Medium was then replaced with 0.5 mL complete medium, and cells were incubated for an additional 44 hours. Cells were washed once with PBS, and 200  $\mu\text{L}$  Reporter Lysis Buffer was added (Promega). Cells were then taken through one freeze-thaw cycle, scraped and cell lysate was collected. Experiments were performed in quadruplicate and analyzed via ANOVA.

### Cytotoxicity Assay

MTT assay was performed. In parallel with the luciferase assay, cells were plated at  $2 \times 10^4$  cells/well in 24-well plates. 24 hours following plating, cells were transfected as described for Luciferase transfection. After 20 hours, 20  $\mu\text{L}$  MTT solution (5mg/mL) was added. After 4 hours, media was removed and 200  $\mu\text{L}$  DMSO was added. 150  $\mu\text{L}$  of the solution was transferred to a 96-well plate and absorbance was measured at 570 nm. Treated cells are relative to untreated control, were performed in quadruplicate, and analyzed via ANOVA.

### Isolation of sRAE-1 $\gamma$

Total RNA was collected from Min6 cells using a RNeasy mini kit according to manufacturers recommendation (Qiagen). A cDNA library was built via RT-PCR with the Superscript III first strand synthesis system (Invitrogen) and oligodT primers. To isolate the gene for soluble RAE-1 $\gamma$ , primers were designed with PrimerQuest (Integrated DNA Technologies, Inc) using a reported sequence for RAE-1 $\gamma$  (NCBI sequence ID: NM\_009018) and screened on BLAST for cross-reactivity. The gene for soluble RAE-1 $\gamma$  was isolated using PCR amplification using gene specific primers with sequences as follows:

Forward: 5' – ATGGCCAAGGCAGCAGTGA – 3'

Reverse: 5' – AGAAGTAGAGTGGCTGGGA – 3'

The expected PCR product was 693 bp long.

### Insertion of sRAE-1 $\gamma$ into Expression Cassette

The isolated gene was inserted into the multiple cloning site of pcDNA3.1/V5-His TOPO TA vector according to manufacturer's instructions. Verification of gene insertion was confirmed by restriction enzyme digestion followed by gel electrophoresis. The orientation and open reading frame (ORF) integrity was confirmed via sequencing

### Determine Expression Profile of pCMV-sRAE-1 $\gamma$

The expression profile of pCMV-sRAE-1 $\gamma$  was characterized with verification of both transcription and translation. MS1 cells were transfected with pCMV-RAE-1 $\gamma$ /PEI polyplexes. The growth medium was collected at 24 and 48 hours following transfection, centrifuged at 14,000  $\times$  g for 2 minutes and stored at  $-20^{\circ}\text{C}$ . Protein was separated via SDS-PAGE then transferred to a nitrocellulose membrane. Protein was detected with anti-V5-AP antibody and colorimetric detection.

## Results

### Synthesis, Conjugation, and Characterization of p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH)

The bio-reducible poly(disulfide amine) poly(cystaminebisacrylamide diamino-hexane) p(CBA-DAH) was synthesized by a Michael-type addition reaction between the monomers N,N'-cystaminebisacrylamide (CBA) and N-Boc-1,6-diamino-hexane (N-Boc-DAH). By using a monomer with one N-boc protected nitrogen, a linear, comb-like polymer with a CBA backbone and DAH pendant chains result. This polymer contains one reducible disulfide bond, one tertiary amine, and one pendant primary amine per repeating unit. SEC data indicates the MW is 6kDa.

$^1\text{H-NMR}$  studies were used to verify the synthesis of p(CBA-DAH) and the conjugation of PEG-p(CBA-DAH) and Eph-PEG-p(CBA-DAH). Synthesis of p(CBA-DAH) was verified by comparing peaks to those previously published [26]. The appearance of a peak for PEG at  $\delta$  3.55 ppm indicates the successful conjugation of PEG. By calculating the ratio of the integral of the peak for PEG with one of the known peaks of p(CBA-DAH) (and normalizing to the relative number of protons in each peak per repeat unit and molecular weight), the ratio of PEG:p(CBA-DAH) was identified as 1.35:1 PEG:p(CBA-DAH).

For Eph-PEG-p(CBA-DAH), the thiolated targeting peptide Eph-SH was conjugated to PEGylated p(CBA-DAH). Again, by measuring the integral of the peaks produced by the methyl groups in the valine, leucine, and threonine (5 methyl groups, 15 protons), and

comparing it to the integral of the peak produced for PEG, the relative number of PEG chains modified with the Eph peptide can be identified. After calculations, it was found that roughly 63% of the PEG chains had been modified with an Eph targeting peptide (Figure 1).

### Electrophoretic Mobility Shift Assay

The ability of each of the three polymers to form polyplexes with plasmid DNA was investigated with an electrophoretic mobility shift assay. In Figure 2(A), the complexation ability of each polymer is examined. As can be seen, p(CBA-DAH) is capable of complexing pDNA at w:w ratios as low as 0.5:1. PEG-p(CBA-DAH) can form stable polyplex at a w:w ratio of 3:1, and Eph-PEG-p(CBA-DAH) can form polyplex at w:w ratios of 5:1. In Figure 2(B), the ability of the polyplexes to release their DNA cargo is shown by treating formed polyplex with 2.5 mM DTT for 30 minutes. Clearly, p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) are all able to release their plasmid DNA at w:w ratios up to 40:1. These results indicate that p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) will be able to unpackage the pDNA cargo upon entry into the reductive intracellular environment, enhancing gene expression [30, 31].

### Polyplex Size Measurement

As can be seen in Figure 3, the particles have effective diameters of less than 250 nm at w:w ratios of 5:1 and above, reaching 200 nm around w:w ratios of 8:1. This is the ideal particle size for efficient cellular uptake via endocytosis and for circulation in vivo [32]. The PEG-p(CBA-DAH) particles remain small and well formed upto w:w ratios of 40:1, however Eph-PEG-p(CBA-DAH) shows an increased particle size above w:w ratio of 10:1. For this reason, w:w ratios up to 10:1 were used for experiments with Eph-PEG-p(CBA-DAH).

### DNase Protection

The ability of the three synthesized polymers to protect pDNA from enzymatic degradation was determined with a DNase protection assay. As can be seen in Figure 4, p(CBA-DAH) is capable of protecting plasmid DNA from enzymatic degradation up to two hours, while p(CBA-DAH) can protect up to roughly 90 min and PEG-p(CBA-DAH) is able to protect up to 30 min.

### Cellular uptake of p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH)

The uptake of PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) was analyzed in MS1s, NIH 3T3s, and 293Ts. MS1s, a pancreatic islet microvasculature cell line, is representative of the target cell and expresses EphA2 and EphA4 receptor. NIH 3T3 and 293T cells do not express EphA2 or EphA4, and are used as controls. As can be seen in Figure 5, the addition of PEG shields the polyplex and prevents non-specific uptake by all cell lines, as can be seen by the low fluorescence in cells treated with PEG-p(CBA-DAH). The recovery of uptake was observed in the MS1 cell line (the target cell line) by the addition of Eph targeting peptide, but not in NIH 3T3 or 293T cells. Furthermore, the functionality of Eph peptide targeting was verified by reducing the uptake in MS1 cells with a free competition assay. The addition of free peptide had no significant effect on the uptake of Eph-PEG-p(CBA-DAH) in NIH 3T3 cells and 293T cells. These results verify that the Eph-PEG-p(CBA-DAH) polymer can specifically target MS1 cells, the pancreatic islet microvasculature endothelial cells.

This result has implications for the ability of Eph-PEG-p(CBA-DAH) to deliver therapeutic genes specifically to the targeted tissue of the pancreatic islet microvasculature.

## Luciferase Expression

Luciferase expression is indicative of uptake, endosomal escape, polyplex unpacking, nuclear entry, transcription, and translation of a delivered gene. As can be seen in Figure 6, there is recovery of delivery with the addition of the Eph targeting peptide to PEG-p(CBA-DAH). However, there is variability in the Luciferase assay, and the results were not significant. The overall trend showed an increase in luciferase expression with delivery with the targeting peptide, which we assume was recovered through receptor-mediated uptake. As with the cellular uptake information, these results indicate that the targeting polymer Eph-PEG-p(CBA-DAH) is capable of delivering plasmid DNA specifically to target cells.

## Cytotoxicity

The addition of PEG is well known to reduce toxicity of cationic polymers [33]. Additionally, the toxicity of bioreducible polymers such as p(CBA-DAH) has been shown to be markedly reduced compared to bPEI and PLL. Thus, it was expected that the polymers p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) would all show very little to no cytotoxicity. As can be seen in Figure 7, none of these three polymers show any statistically significant cytotoxicity up to a w:w ratio of 40:1.

## Construction of sRAE-1 Plasmid

Cloning of the sRAE-1 into the plasmid was also confirmed with restriction enzyme digestion. Digestion with NotI and KpnI produced a 770 bp fragment (sRAE-1), and a 5.4 kb fragment (vector), confirming the successful insertion of sRAE-1. The digestion of pCMV-RAE-1 $\gamma$  is shown in Figure 8. Lane 1 is a 1 Kb plus marker (Invitrogen), lane 2 shows the pcDNA3.1/v5-His vector alone, and lane 3 shows the complete constructed plasmid pCMV-RAE-1 $\gamma$  digested with KpnI and NotI.

## Characterization of pCMV-RAE-1 $\gamma$

Expression and secretion of sRAE-1 $\gamma$  were evaluated following transfection of pCMV-RAE-1 $\gamma$  into pancreatic islet endothelial cells (MS1s). Transcription was verified using RT-PCR, and translation and secretion were verified using western blot. For RT-PCR, total RNA was collected from cells 24 hours following transfection. Reverse transcription was performed with both oligo dT primers and gene specific primers, and product was amplified using forward and reverse primers designed for initial isolation. As can be seen in Figure 9, sRAE-1 $\gamma$  is expressed in transfected cells (lanes 3 and 5), but not in untreated controls (lanes 2 and 4), verifying transcription of exogenous sRAE-1 $\gamma$ .

For western blot, cells were transfected with pCMV-RAE-1 $\gamma$  and samples of supernatant were collected at various time points. The total protein content was then assayed (BCA protein assay kit, Pierce) and equal amounts of protein were loaded and separated via SDS-PAGE. Following separation, protein was transferred to a nitrocellulose membrane, sRAE-1 $\gamma$  was stained with an anti-V5-AP antibody, and the blot was developed with an AP detection kit (BioRad). There is a time dependent secretion of soluble RAE-1 $\gamma$ . Lane 1 shows the protein ladder, with bands at 40 and 50 kDa as shown in Figure 10. In lanes 2–5, samples collected at 24 hours can be seen, while in lanes 6–9, samples collected at 48 hours are seen.

## Discussion

The use of p(CBA-DAH) as a polymeric gene carrier is attractive for several reasons. The inclusion of reducible disulfide bonds allows for the rapid triggered degradation upon entering the cytoplasmic compartment [34, 36]. The presence of primary and tertiary amines in the reducible polymer allow for the self-assembly of nucleic acid/polymer polyplexes

[38]. Further, the presence of primary amines allows for straightforward modification of the polymer with PEG, and helps maintain solubility of the polymer [26]. Also, the polymer is synthesized from commercially available monomers, allowing for a straightforward synthesis.

The size of polyplexes plays a vital role in biodistribution for in vivo delivery [39–41]. Accumulation in the liver and spleen is prevalent with particle sizes that are too big (above 250 nm), so the ability to form polyplexes with diameters below that cutoff is important in designing a system for targeted in vivo delivery. All three polymers were able to reach sub-250 nm particle sizes at optimal w:w ratios, as can be seen in Figure 3.

The ability to condense DNA is essential for successful delivery to target tissues. The formation of dense polyplex between polymers and pDNA allows for the protection of the DNA from enzymatic degradation as well as packaging to facilitate cellular uptake [41, 42]. All three polymers were able to successfully condense pDNA at low w:w ratios, as was seen in the electrophoretic mobility shift assay in Figure 2 a). Furthermore, the ability to unpackage the pDNA following cellular uptake and endosomal escape enhances transfection efficiency [35–38]. The triggered release of plasmid DNA upon incubation in a reductive environment mimics the polyplex behavior in the cytoplasmic compartment. It was seen in Figure 2 b) that treatment of formed polyplexes with DTT can cause the triggered release of plasmid DNA, indicating that these polymers have an environment-responsive element for the triggered release following endosomal escape [29–45].

The ability to protect from enzymatic degradation is also essential for these polymers to be used for systemic delivery [46]. As was seen in Figure 5, the targeting polymer Eph-PEG-p(CBA-DAH) was able to protect plasmid DNA from DNase degradation up to 120 minutes (Figure 4). For p(CBA-DAH), some DNA degradation could be seen at 120 minutes. PEG-p(CBA-DAH) was only able to protect from Dnase degradation for roughly 30 minutes, however this result was not unexpected as the addition of PEG is known to interfere with condensation ability of cationic polymers [23].

The targeting ability of Eph-PEG-p(CBA-DAH) to the islet microvasculature endothelial cells was examined with transfection experiments with EphA2-expressing MS1 cells and EphA2-negative NIH 3T3 and 293T cells. It was expected that the addition of PEG to the backbone polymer would reduce cellular uptake of polyplex, as PEG has been previously reported to shield the cationic charge of polyplex and prevent non-specific interactions with cells [47–49]. This reduction in uptake was then expected to be recovered in the target cells, specifically MS1, the pancreatic islet microvasculature endothelial cell that expresses EphA2, with the addition of the targeting peptide, as has been shown with antibody fragments and other targeting peptides [50–52]. The uptake ability was not expected to recover in non-targeted cell lines, namely NIH 3T3 and 293T. The Eph-PEG-p(CBA-DAH) polymer was able to deliver plasmid DNA to MS1 cells with about 5-fold higher uptake efficiency than PEG-p(CBA-DAH) control in MS1 cells (Figure 5). In NIH 3T3 cells and 293T cells, however, there was no significant difference in uptake efficiency between Eph-PEG-p(CBA-DAH) and PEG-p(CBA-DAH). These results suggest that the Eph-PEG-p(CBA-DAH) polymer has specificity toward EphA2 expressing cells. The enhanced cellular uptake was verified to be due to specific interaction between the targeting peptide and the EphA2 receptor with a competition experiment carried out in the presence of free targeting peptide (0.1 mM) during transfection with Eph-PEG-p(CBA-DAH). The uptake efficiency of Eph-PEG-p(CBA-DAH) was significantly reduced to levels similar to that of PEG-p(CBA-DAH) complexes (Figure 5). In transfection experiments expressing luciferase, although there is a trend of increase in expression in the cells treated with Eph-PEG-p(CBA-



DAH) compared to PEG-p(CBA-DAH), the results were not significant due to the variability in the experiment (Figure 6).

It is thought that this targeting polymer will have significant advantages when delivered in vivo. It is expected to have lower transfection efficiency with the addition of PEG to a backbone polymer, as most of the cellular uptake in unmodified cationic polymers is due to non-specific electrostatic interactions between polyplex and the anionic cell membrane [53].

The construction of an expression plasmid for a constitutively expressed soluble ligand to the NKG2D receptor, once successfully delivered locally to the pancreatic islets, will allow for the pancreatic islet beta cells to undergo immune evasion and prevent NKG2D mediated cytotoxicity to the beta cells. The therapeutic potential of the blockade of the NKG2D receptor for the prevention of type 1 diabetes remains largely unexplored, mainly due to the potential for side effects including the reduction in tumor surveillance, DNA damage detection, and viral infection [15, 54]. The systemic blockade of the NKG2D receptor is not an ideal therapy, yet remains the only in vivo tested therapeutic to prevent type 1 diabetes by receptor blockade in the form of systemic anti-NKG2D IgG delivery [13]. The construction of a therapeutic plasmid to deliver the gene that will encode for a protein to be synthesized locally around the pancreatic islets capable of blocking the NKG2D receptor of pancreatic infiltrating immunocytes will allow for the benefit of the blocking therapy while reducing the side effects expected from a systemic blockade.

The RAE-1 $\gamma$  expressing plasmid was designed for high constitutive expression with elements to aid in detection and purification [55]. The CMV promoter, in addition to being a strong, constitutively active promoter in many tissues, also shows relatively low transcription in hepatocytes [56]. The inclusion of the V5 epitope will enhance detection ability, both in vitro with western blot and in vivo with immunohistochemistry [57].

## Conclusions

The synthesis of p(CBA-DAH) and modification to PEG-p(CBA-DAH) and Eph-PEG-p(CBA-DAH) were successfully completed. The polymers are all able to form polyplexes with pDNA and release that pDNA rapidly in a reductive environment due to the reduction of disulfide bonds in the backbone. All three polymers are capable of forming nano-sized particles when complexed with pDNA. Both p(CBA-DAH) and Eph-PEG-p(CBA-DAH) show the ability to protect pDNA from enzymatic degradation. It was shown that the addition of PEG to the backbone polymer reduced non-specific cellular uptake and transfection, and that the addition of a targeting peptide Eph could recover the cellular uptake and transfection efficiency selectively in a specific cell line. Additionally, the construction and characterization of expression of pCMV-RAE-1 $\gamma$  was successfully completed. The therapeutic gene was inserted in the expression plasmid, as was verified by sequencing and restriction enzyme digestion. The expression capabilities of the pCMV-RAE-1 $\gamma$  plasmid were verified with RT-PCR and western blot. The therapeutic protein srRAE-1 $\gamma$  was secreted, as was also verified by western blot. Therefore, the combination therapy of the polymer Eph-PEG-p(CBA-DAH) and the therapeutic pCMV-RAE-1 $\gamma$  is a viable therapeutic strategy for the prevention of type 1 diabetes.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

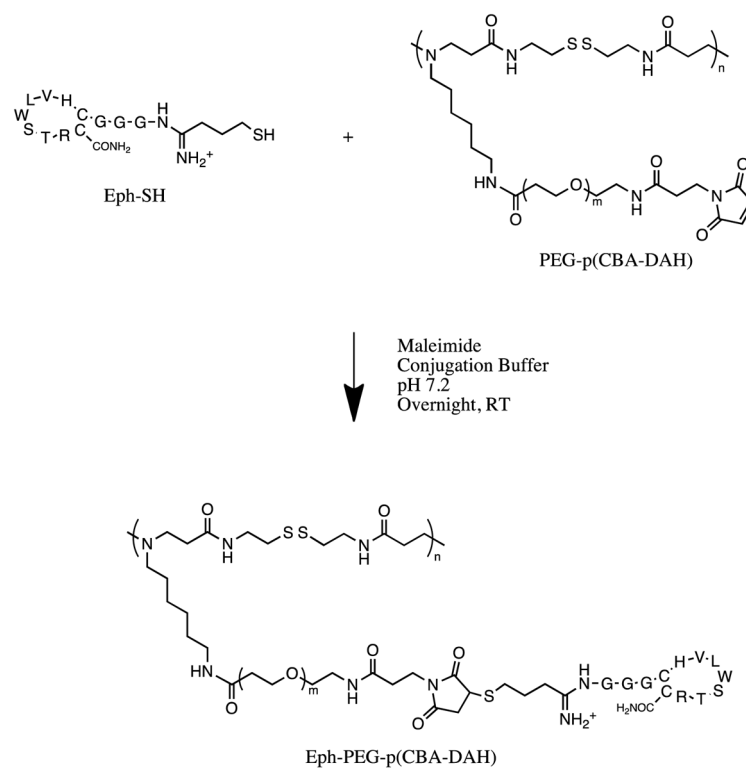
Funding by NIH Grants DK085075.

## References

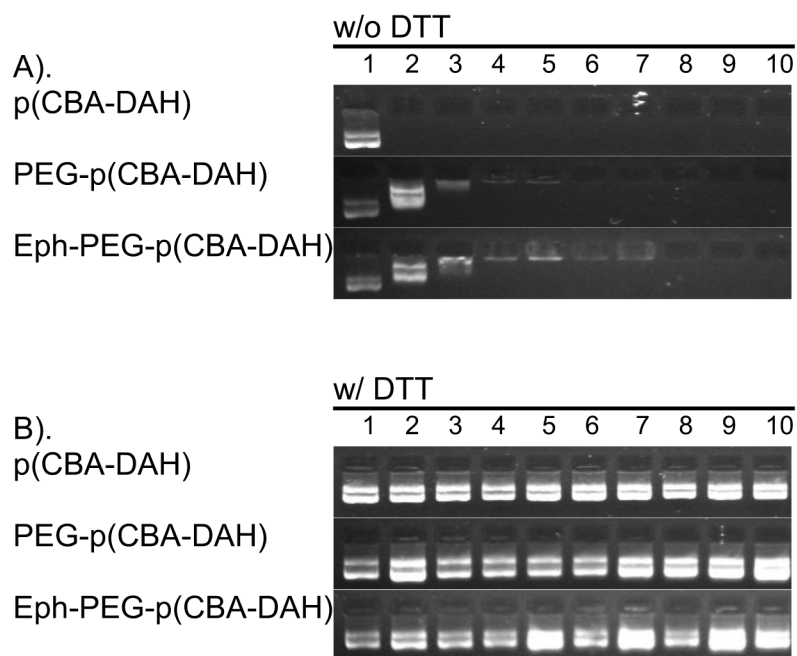
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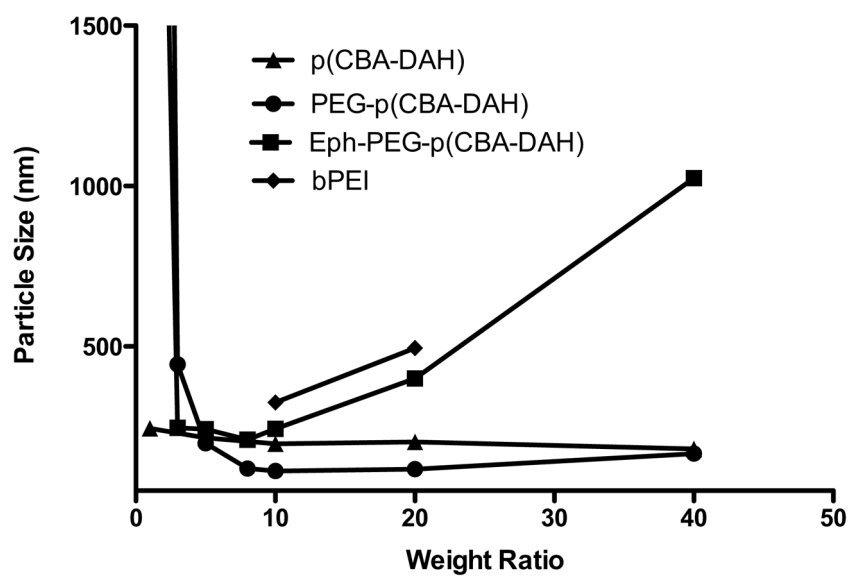
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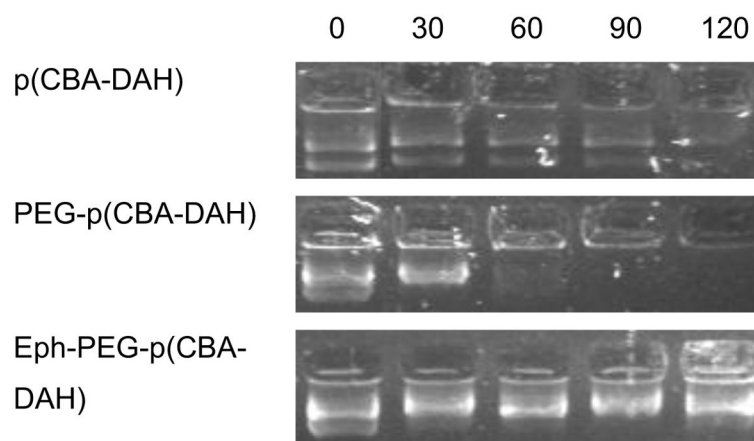
**Figure 1.**  
Conjugation scheme for Eph-PEG-p(CBA-DAH)



**Figure 2.** Electrophoretic mobility shift assay of p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) with pDNA polyplex (A) without DTT and (B) with DTT (2.5 mM for 30m). Lane 1: naked pDNA; Lanes 2–10: polymer/pDNA polyplexes at N:P ratios of 0.5:1, 1:1, 3:1, 5:1, 10:1, 15:1, 20:1, 30:1, 40:1.

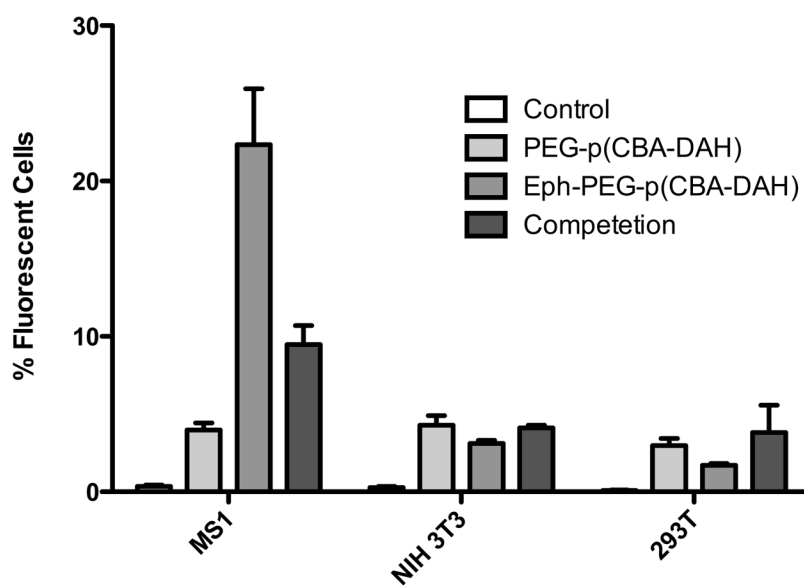


**Figure 3.** Average particle size of polymer/pDNA polyplexes at w:w ratios from 1:1 to 40:1. bPEI/pDNA polyplex sizes were determined at N:P ratios of 10:1 and 20:1. The large size of Eph-PEG-p(CBA-DAH) at w:w ratios above 10 is due to particle aggregation.

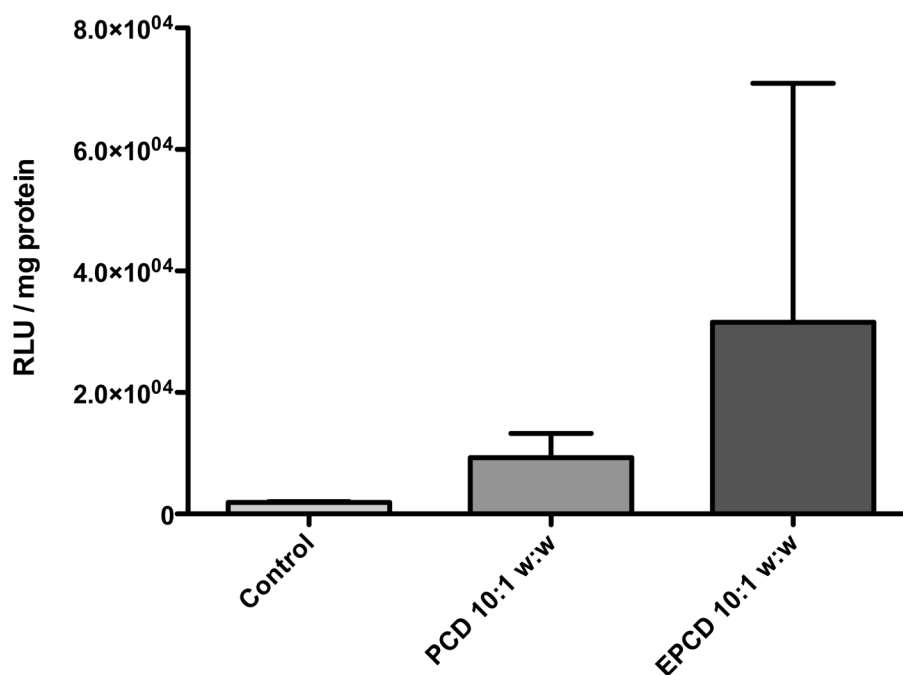


**Figure 4.** DNase protecting ability of p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) at 0, 30, 60, 90, and 120 minutes. W:W ratios of 10:1 were used.

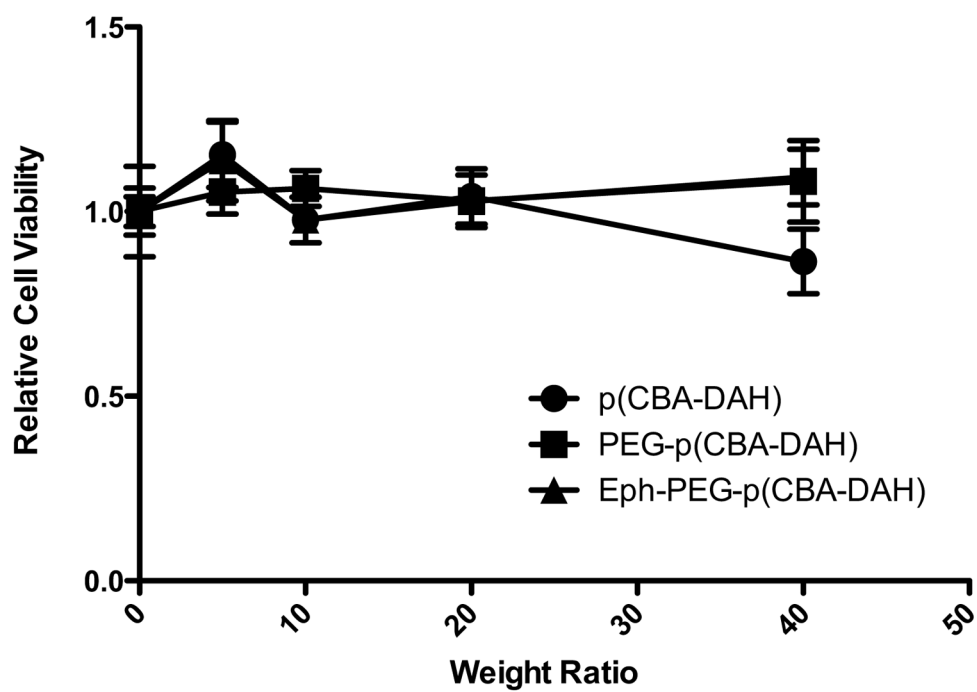




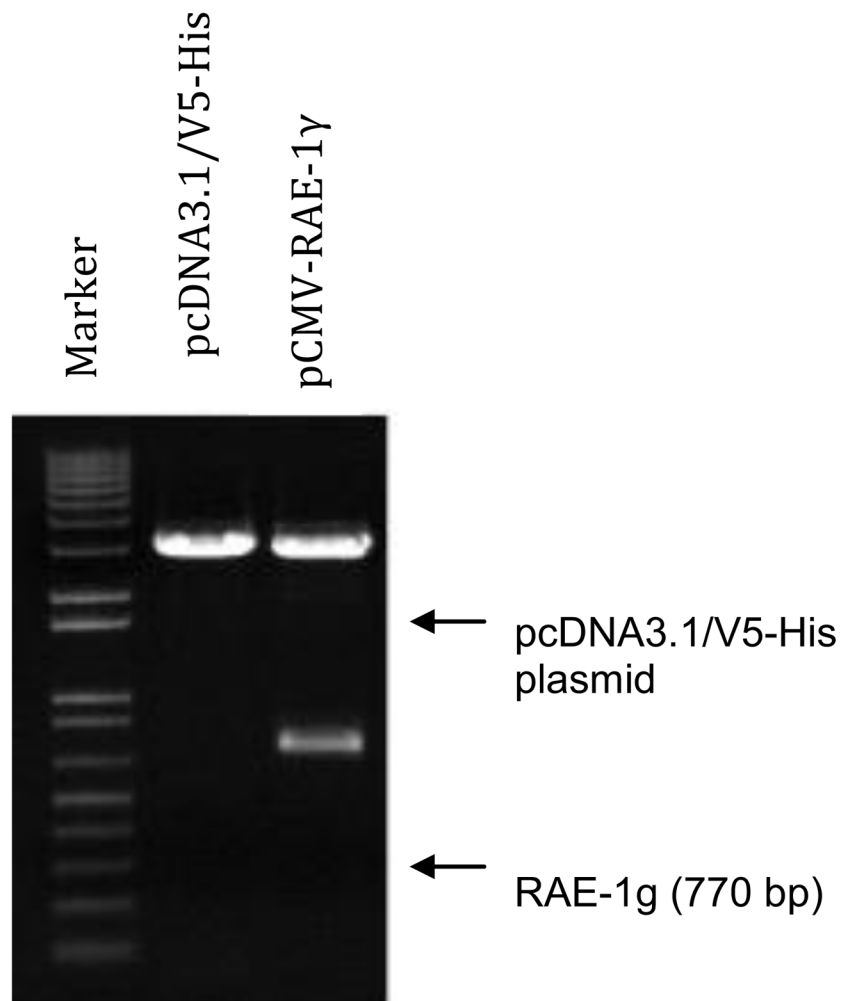
**Figure 5.** Cellular uptake of polyplexes of pDNA with PEG-p(CBA-DAH) (PCD), and Eph-PEG-p(CBA-DAH) (EPCD). The effect of addition of free Eph peptide (0.1 mM) was examined in an uptake competition assay (EPCD Comp).



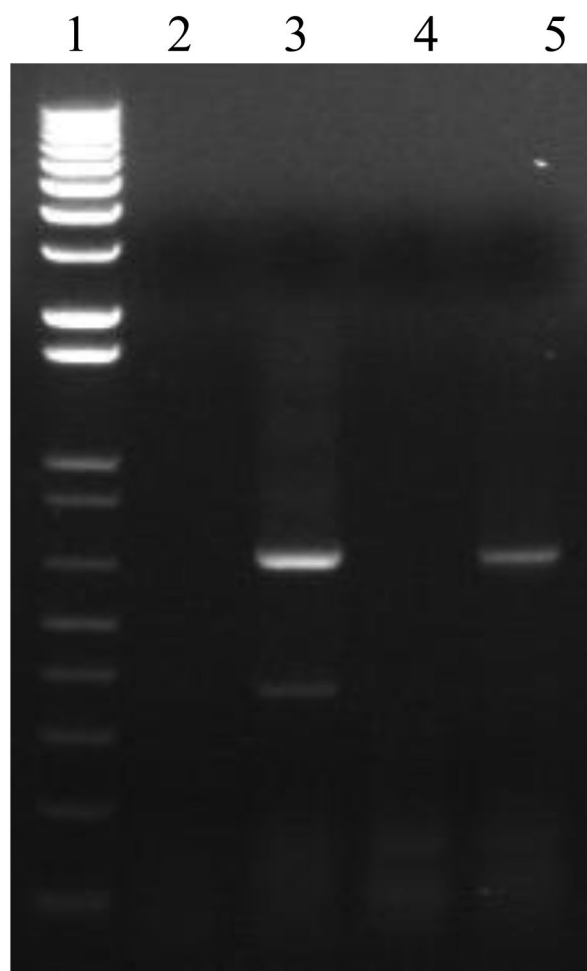
**Figure 6.** Luciferase expression in MS1 cells transfected with pCMV-Luc delivered with PEG-p(CBA-DAH) (PCD), and Eph-PEG-p(CBA-DAH) (EPCD). Polyplexes were formed at a 10:1 w:w ratio.



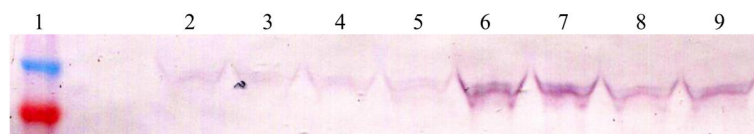
**Figure 7.** Cytotoxicity of p(CBA-DAH), PEG-p(CBA-DAH), and Eph-PEG-p(CBA-DAH) in MS1 cells at various w:w ratios.



**Figure 8.** Digestion of pCMV-RAE-1 $\gamma$  with KpnI and NotI.



**Figure 9.** RT-PCR verifies transcription of sRAE-1  $\gamma$ . Lane 1: 1 kb+ size marker; lane 2: control nontransfected – oligo dT primer; lane 3: pCMV-RAE-1 $\gamma$  transfected – oligo dT primer; lane 4: control non-transfected – gene specific primers; lane 5: pCMV-RAE-1 $\gamma$  – gene specific primers.



**Figure 10.** Western blot verifies translation and secretion of sRAE-1 $\gamma$  following transfection in MS1 cells. Lane 1: Novex Sharp prestained ladder (40 and 50 kDa bands shown). Lanes 2–5, 24h. Lanes 6–9, 48h.