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Ligands located within a cholesterol domain enhance gene delivery to the target tissue

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Abstract

Targeted gene delivery provides enormous potential for clinical treatment of many incurable diseases. Liposomes formulated with targeting ligands have been tested extensively both in vitro and *in vivo*, and many studies have strived to identify more efficacious ligands. However, the environment of the ligand within the delivery vehicle is generally not considered, and this study assesses the effect of ligand micoenvironment by utilizing a lipoplex possessing a cholesterol domain. Our recent work has shown that the presence of the targeting ligand within the cholesterol domain promotes more productive transfection in cultured cells. In the present study, lipoplexes having the identical lipid composition were formulated with different conjugates of the folate ligand such that the ligand was included in, or excluded from, the cholesterol domain. The effect of locating the ligand within the cholesterol domain was then tested in a xenograft tumor model in mice. Lipoplexes that included the ligand within the cholesterol domain showed significantly higher luciferase expression and plasmid accumulation in tumors as compared to lipoplexes in which the ligand was excluded from the domain. These results demonstrate that the microenvironment of the ligand can affect gene delivery to tumors, and show that ligand-mediated delivery can be enhanced by locating targeting ligands within a cholesterol domain.

Keywords

folate; nanodomain; targeted gene delivery in vivo; liposome; tumor targeting; cholesterol domain; serum stable; ligand microenvironment

Introduction

Delivery of DNA/siRNA into cells offers the potential to develop potent vaccines and novel therapeutics to cure many diseases that are currently difficult to treat with traditional therapies, e.g., hereditary diseases, cancer [1-4]. Cationic liposomes have been used extensively in gene delivery both in vitro and in vivo. However, it has been shown that cationic lipoplexes are taken up predominantly in the lung and liver [5-11]. Furthermore, it is known that lipoplexes interact with serum proteins and blood cells in the circulation, resulting in uptake by the cells of the mononuclear phagocyte system (MPS), classically known as the reticuloendothelial system [12-15]. When targeting a cancer therapeutic to

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tumors, uptake by these alternate tissues/cells limits accumulation in the tumor thereby reducing efficacy. In efforts to enhance uptake within tumors, researchers have utilized various targeting ligands (e.g., antibodies, peptides) binding specifically to receptors that are highly expressed in tumors [16-21]. Strategies for improving tumor targeting have largely focused on identifying ligands with greater affinities for the receptor on the tumor cell. However, it is clear that cell membranes are composed of different regions (e.g., "rafts") that possess specialized functions, and therefore the microenvironment of the ligand within the delivery vehicle may play a role in the ligand mediated drug delivery when it interacts with the membrane.

Another factor that is thought to play a large role in the ability of delivery vehicles to distribute to tumors is maintenance of a small particle size and prolonged circulation in the blood [23,25,26]. Nanoparticles that are endowed with these properties are believed to be "passively targeted" to tumors via the enhanced permeation and retention effect (EPR) [22-24]. Typically, PEGylated components are incorporated into particulate delivery systems because PEG is known to endow particles with greater stability in blood which increases deposition in tumors [25-27]. However, PEGylation is also known to have detrimental impacts on cellular uptake and intracellular trafficking that compromises the ultimate delivery efficiency [28-31]. Recent studies by Li et al. [22] on siRNA delivery concluded that PEGylation reduces delivery efficiency in cell culture by approximately 10 fold, and in vivo studies have also observed decreased tumor accumulation with PEGylated gene delivery systems [32].

As an alternative strategy to PEGylation, our previous work has demonstrated the remarkable serum stability of lipoplexes formulated with high cholesterol contents [33, 34]. Additional studies demonstrated that this formulation strategy extends circulation time and enhances tumor distribution to a greater extent than that observed with PEGylated lipoplexes [32]. Although the exact mechanism by which cholesterol imparts stability has yet to be fully elucidated, we proposed that cholesterol imparts sufficient rigidity such that serum protein adsorption/insertion is dramatically attenuated, thereby reducing the aggregation that accelerates clearance [33, 34]. It follows that the incorporation of targeting ligands into a cholesterol-stabilized lipoplex would circumvent the detrimental effects of PEGylation noted above, while taking advantage of the ability of targeting ligands to facilitate uptake by tumors.

Further work on the characterization of lipoplexes formulated with elevated cholesterol has shown that a region of pure cholesterol ("cholesterol domain") is formed at 69% (w/w) cholesterol; a concentration at which we observe a distinct increase in both serum stability and in vitro transfection [34]. Also, serum protein binding studies indicated that protein adsorption to these cholesterol domains is negligible, suggesting that these domains represent a distinctly different environment on the lipoplex that may offer advantages with regard to ligand-mediated drug delivery [34, 35]. Because domain formation is predominantly governed by hydrophobic interactions in the acyl chain region of the bilayer [36-38], the incorporation of folate-cholesterol into lipoplexes results in the ligand being able to partition into the cholesterol domain. In contrast, ligands anchored to the lipoplex via diacyl lipids (folate-DSPE) are excluded from the domain. Accordingly, the utilization of these different folate conjugates in lipoplexes allows us to assess the effect of the local environment (i.e., within or excluded from the cholesterol domain) on ligand-mediated delivery.

Our most recent work has incorporated these folate conjugates into lipoplexes possessing domains to assess their effect on *in vitro* transfection [35]. Enhanced transfection was observed when the folate ligand was conjugated to cholesterol and therefore able to partition

into the cholesterol domain. Conversely, conjugation of folate to a diacyl lipid (DSPE) that results in exclusion from the cholesterol domain caused a reduction in transfection rates as compared to lipoplexes lacking the folate ligand [35]. Curiously, studies assessing cellular uptake and internalization clearly showed that the enhanced transfection observed with formulations possessing folate within the cholesterol domain does not result from greater binding and/or uptake [35]. These results suggest that location of the targeting ligand within the cholesterol domain facilitates targeted gene delivery by altering intracellular trafficking such that uptake is more productive.

In the present study, we investigated the ability of cholesterol domains to promote ligandmediated gene delivery to tumors in vivo. As in our previous cell culture experiments mentioned above, lipoplexes were formulated with different conjugates of the folate ligand such that the ligand was either included in, or excluded from, the cholesterol domain. Transgene expression (luciferase) and DNA accumulation (determined by real-time PCR) were assessed in the tumor and other tissues 24 hrs after intravenous injection. Consistent with our cell culture studies, lipoplexes that located the ligand within the cholesterol domain (i.e., folate-cholesterol) showed significantly higher luciferase expression and plasmid accumulation in tumors as compared to lipoplexes in which the ligand was excluded from the domain (i.e.,folate-DSPE) or lipoplexes lacking the ligand. Our observation that the microenvironment of the ligand can have significant effects on ligand-mediated gene delivery suggests that vehicles possessing multiple microenvironments (domains) can offer advantages over nanoparticles possessing uniform surface properties.

Materials and Methods

Materials

Luciferase plasmid DNA without the CpG motif was a generous gift from Dr. Manfred Ogris in Ludwig-Maximilians-Universität München and was prepared with endotoxin level lower than 0.03 EU/μg by Genscript Inc. (Piscataway, NJ). N-(1-(2, 3-dioleoyloxy) propyl)- N, N, N-trimethylammonium chloride (DOTAP), 1,2-distearoyl-sn-glycero-3 phosphoethanolamine-N-[methoxy(polyethylene glycol)-2000] (PEG₂₀₀₀-DSPE) and 1,2distearoyl-sn-glycero-3-phosphoethanolamine-N-[folate(polyethylene-glycol)-2000] (DSPE-PEG₂₀₀₀-Folate) were purchased from Avanti Polar Lipids (Alabaster, AL). Synthetic cholesterol was purchased from Sigma-Aldrich (St. Louis, MO). Folate-PEG₂₀₀₀-Cholesterol was custom synthesized by GLS synthesis Inc (Worcester, MA) and $PEG₂₀₀₀$ -Cholesterol was synthesized by Dr. Michael Wempe as described by Zhao et al [39]. The identity of the PEG-cholesterol was confirmed by NMR, and the purity was estimated to be > 95%. The luciferase assay kit was obtained from Promega (Madison, WI). Fetal bovine serum (FBS) was purchased from Mediatech Inc. (Manassas, VA) and was filtered with 0.22-µm low protein binding cellulose acetate filter from Fisher Scientific (Pittsburgh, PA) before use. All chemicals were of reagent grade or higher quality.

Liposome and lipoplex preparation

DOTAP combined with cholesterol (wt/wt: 31/69) was mixed in chloroform. The lipid mixture was dried under a stream of nitrogen gas and placed under vacuum (100 mTorr) for 2 hours to remove residual chloroform, and dried lipids were subsequently resuspended in autoclaved, distilled water and sonicated. Cationic liposomes were prepared immediately before use as previously described [32]. Lipoplexes were prepared by mixing equal volumes of DNA (50 μ g/ml) and liposomes (0.625 mM), and incubated at room temperature for 15 min. To increase the amount of luciferase DNA delivered into the mice while keeping the injection volume low to avoid hydrodynamic effects, the lipoplexes were concentrated tenfold (to 250 μg DNA/mL) with Millipore ultrafree 100K MW centrifugal filters (Bedford,

MA). This method for concentrating the lipoplexes did not alter the particle size as detected by dynamic light scattering (data not shown). To prepare folate or PEG conjugated lipoplexes, folate-cholesterol, folate-DSPE, PEG-DSPE or PEG-cholesterol was mixed with DOTAP and cholesterol in chloroform following the procedures described above.

Animal use and tumor inoculation

All procedures were approved by the University of Colorado Denver Committee on Animal Research. KB cells (ATCC #CCL-17, 1×10^7 cells/mL) in a volume of 0.1 mL were subcutaneously injected into the right flanks of male athymic nude mice (National Cancer Institute, Bethesda, MD) 4-6 weeks old. Animals with tumors were treated with lipoplexes when tumor volume reached 100 mm^3 .

In vivo treatment protocols

Mice with tumors were administered lipoplexes via either intratumoral or intravenous routes. For intratumoral administration, lipoplexes with 5 μg plasmid DNA encoding luciferase were injected into the tumor directly. Mice were sacrificed 24 hrs after injection and tumors were collected for luciferase expression. For the intravenous route, animals with tumors were administered lipoplexes via a single i.v. bolus dose of 50 μg plasmid DNA encoding luciferase in the tail vein. Following treatment, mice in each formulation were sacrificed at 24 hrs. Tumor, liver, spleen, kidney, heart and lung were collected from each animal after sacrifice by carbon dioxide exposure, quickly frozen in liquid nitrogen, and stored at -80°C until analysis. Tissues were analyzed for luciferase expression using a luciferase assay kit (Promega) and plasmid DNA levels were quantified via real-time PCR.

Quantification of DNA in tissues

Total DNA from each tissue was extracted with a Qiagen DNeasy tissue kit (Qiagen, Valencia, CA) following the protocol provided by the manufacturer. Briefly, tissue samples are first lysed using Proteinase K and the lysate is loaded onto the DNeasy mini columns. During centrifugation, DNA is selectively bound to a silica-gel membrane. DNA is then washed to remove impurities and total DNA is eluted and quantified by A260/280 measurements in an Agilent UV/Vis spectrophotometer. The plasmid DNA encoding luciferase was then amplified and quantified via real-time quantitative PCR in an ABI GeneAmp 9700 Sequence Detection System (PE Biosystems, Foster City, CA). Extracted DNA was added to a SYBR Green PCR master mix (Qiagen) and 0.3 mM of primers (Integrated DNA Technologies, Inc, Coralville, IA). The sequences of primers are: 5'-TGA AGA GGT ATG CCC TGG TC-3' and 5'-CCA GCC TCA CAG ACA TCT CA-3'. In order to quantify the amount of DNA present in a reaction tube, standard curves were generated and the amounts of plasmid in the samples were determined by interpolation [32].

In vitro firefly luciferase assay

Extraction of luciferase from mouse tissues was carried out as described [24]. Frozen tissues were homogenized in 0.5 mL/organ of Cell Lysis buffer (Promega) except liver in 1 mL/ organ of Cell Lysis buffer. After thawing in 37°C water baths, the homogenates were centrifuged at $13,000 \times g$ for 10 min. Luciferase activity was assessed using 20 μ L supernatant with 100 μL Luciferase Assay Reagent (Promega).

Statistical analysis

A one-way analysis of variance (ANOVA) was used to determine statistical significance $(p<0.05)$ among the mean values for luciferase expression and DNA accumulation. A Bonferroni's multiple comparison test was used to determine statistical significance ($p<0.05$) between formulations.

Results

Optimization of folate-cholesterol concentration by intra-tumoral injection

In order to determine the optimal ligand concentration for tumor-targeted delivery, the folate ligand was incorporated into lipoplexes prepared with DOTAP/cholesterol (wt/wt: 31/69). Lipoplexes formulated with different levels of folate were then injected directly into xenograft tumors of KB cells in nude mice. Twenty-four hours after injection, mice were sacrificed and tumors were collected and analyzed for luciferase expression. The optimal transfection efficiency was observed at the concentration of 0.4 mol% folate-cholesterol, with a 7-fold increase as compared to the control lacking the folate ligand (Fig.1). Higher ligand concentrations (>0.4 mol %) did not further improve luciferase expression in tumors, instead transfection efficiency actually decreased dramatically at higher ligand concentrations. Therefore, further experiments employed lipoplexes prepared with 0.4 mol% folate ligand.

Targeted systemic gene delivery

To test delivery to the tumor, folate-targeted lipoplexes containing DOTAP/cholesterol (wt/ wt: 31/69) were prepared with two different conjugates of cholesterol (i.e. folate-cholesterol and folate-DSPE) in order to assess the effect of locating the targeting ligand within the cholesterol domain. Lipoplexes without ligands as well as PEGylated formulations prepared by inclusion of PEG-DSPE or PEG-cholesterol were included as controls. Systemic gene delivery into nude mice with tumors was carried out via a single i.v. bolus dose of 50 μ g plasmid DNA encoding luciferase in the tail vein. Twenty four hours post treatment, mice were sacrificed and organs (tumor, liver and lung) were collected for analysis. In tumors, lipoplexes in which the ligand was located within the domain (folate-cholesterol) showed significantly higher levels of luciferase expression (2-fold higher than the no ligand control), whereas formulations in which the ligand was excluded from the domain (folate-DSPE) exhibited formulations incorporating PEGylated components (i.e., PEG-DSPE) that were excluded from the domain also resulted in reduced expression. In this regard, PEGylation is well known to adversely affect the intracellular trafficking of lipoplexes, however attachment of the targeting ligand (folate) at the distal end of PEG-DSPE did not appear to help increase transfection. Interestingly, formulations in which PEG was located within the domain (PEG-cholesterol) did not show the negative effect on transfection observed with PEG-DSPE, with the luciferase expression being comparable to markedly reduced expression (Fig. 2A). In addition, the use of the no ligand control. In the liver, all the formulations exhibited comparable luciferase expression with the exception of lipoplexes containing PEG-cholesterol (Fig. 2B). In the lung, lipoplexes with folate-cholesterol showed levels of luciferase expression comparable to the no ligand control, and a significant difference was observed between formulations where the ligand was located within the domain (folate-cholesterol) as compared to formulations in which the ligand was excluded from the domain (folate-DSPE, Fig. 2C). As mentioned above, alteration of the ligand microenvironment (i.e., included in or excluded from the domain) affects transfection in agreement with studies in cell culture. However, it is interesting that altering the microenvironment of the PEG can also have a dramatic effect on transgene expression (Figs 2A-C). While this effect was observed in all the tissues tested, it was most striking in the lung where luciferase expression was 7-fold higher when PEG was present within the domain as compared to formulations where PEG was excluded from the domain (Fig. 2C).

In order to quantify the biodistribution of plasmid DNA, tissue samples were also analyzed by real-time PCR. As shown in Fig. 3A, the tumor accumulation of plasmid DNA was significantly higher (2.7-fold) with the formulation incorporating folate-cholesterol as compared to formulations with folate-DSPE or lacking a ligand. In contrast, DNA

accumulation in the liver was reduced by all formulations except those incorporating folatecholesterol (Fig. 3B). Significantly higher levels of DNA accumulation were observed in the liver with folate-cholesterol as compared to the no ligand control and PEG-cholesterol. Interestingly, formulations containing PEG-cholesterol delivered significantly higher levels of DNA to the lung (Fig. 3C), consistent with the increased luciferase expression in the lung observed with this formulation (Fig 3A). In other organs tested (spleen, kidney and heart), luciferase expression and DNA accumulation were reduced dramatically in the kidney for all the formulations (S1. B and E). Formulations including PEG-cholesterol showed higher luciferase expression than other formulations, despite its comparably low level of DNA accumulation in the spleen, kidney and heart (S1. D, E and F). In the spleen, formulations with folate-cholesterol showed significantly higher accumulation than that with PEGcholesterol, while its transgene expression was significantly lower.

Discussion

Tumor-targeted gene delivery utilizing lipid based systems holds tremendous promise for cancer gene therapy. However, due to the serum-induced aggregation upon intravenous injection, PEGylated components have been predominantly employed to sterically shield the delivery vehicles from blood components [28, 40, 41]. Our previous studies [33, 34] have shown that lipoplexes with high levels of cholesterol exhibited enhanced transfection in vitro and resistance to serum-induced aggregation, and thus could be utilized as a potential alternative to PEGylation. Furthermore, formulating lipoplexes with high levels of cholesterol (69% by weight) caused the formation of a cholesterol domain (determined by differential scanning calorimetry) that corresponds with a significant increase in transfection [34]. The presence of a domain creates a distinctly different microenvironment within the lipoplex characterized by a neutral charge. In contrast, the microenvironment outside of the cholesterol domain contains cationic lipids involved in DNA binding. It is possible that the non-ionic character contributes to the undetectable levels of serum protein that adsorbs to the cholesterol domain, which further distinguishes it from the rest of the lipoplex [35].

Incorporation of a targeting ligand that partitions into the cholesterol domain (folatecholesterol) was shown to enhance transfection in KB cells that overexpress folate receptor [35]. The present study utilized the same delivery system for targeted gene delivery in vivo, and our initial experiments were aimed at optimizing ligand concentration via intra-tumoral injection. Luciferase expression in the tumor reached the highest level in lipoplexes formulated with a ligand concentration of 0.4 mol%, while higher ligand concentrations (± 1 mol %) caused a decrease in transfection rate (Fig. 1). This result is entirely consistent with the *in vitro* result [35], as well as with our assertion that cell culture experiments employing 50% serum provide a more realistic assessment of delivery system performance upon intravenous administration, at least in terms of transfection.

The main goal of the present study was to test the *in vivo* effect of locating a ligand within the cholesterol domain, and we utilized a xenograft model where a subcutaneous human tumor cell line that over-expresses the folate receptor (KB cells) was used to produce tumors in athymic nude mice. Lipoplexes with the folate ligand located within the cholesterol domain showed significantly higher transgene expression after intravenous injection than that observed with the identical lipoplex formulation when the folate ligand was excluded from the domain (Fig. 2A). Similarly, accumulation of plasmid DNA in tumors was significantly higher when the ligand was located within the domain (Fig 3A). Because the identical lipoplex formulation was utilized with each ligand conjugate, it is highly unlikely that differences in serum stability and/or clearance can explain our findings. Also, the fact that the ligand microenvironment alters tumor accumulation calls into questioning the conclusion that accumulation in tumors is governed exclusively by the EPR effect [22-24].

Considering that smaller particle sizes should enhance the EPR effect, it would be expected that the formulation incorporating folate-DSPE would exhibit superior accumulation in the tumor due to its smaller diameter compared to that with folate-cholesterol (153 \pm 8.8 vs. 197.7 \pm 16.8 nm; [35]). In fact, plasmid DNA accumulation in the tumor was 2.7-fold *higher* with the larger particle wherein the folate ligand was located within the cholesterol domain (Fig. 3A). These data strongly suggest that factors other than EPR can affect accumulation at the tumor site. It should be pointed out that previous studies concluding that EPR governs tumor accumulation monitored localization at 4 hrs, whereas we monitored accumulation after 24 hrs. Considering the rapid loss of plasmid from tissues [32, 42]; Xu, L unpublished] (presumably due to extracellular and/or lyososomal degradation), our data showing greater tumor accumulation with formulations incorporating the folate ligand within the cholesterol domain may reflect greater plasmid persistence within the target tissue. Considering our cell culture experiments showing that this delivery system endowed with folate-cholesterol increases transfection despite reduced uptake (as compared to folate-DSPE) [43], we suggest that the greater tumor accumulation is due to more productive intracellular trafficking after the initial deposition.

Furthermore, these in vivo results suggest a correlation of tumor accumulation with transgene expression (R^2 =0.74, Fig. 4A). A better correlation is seen in the lung (R^2 =0.96), but there is a clear lack of correlation in the liver $(R^2=0.02)$, suggesting that liver accumulation involves less productive uptake of lipoplexes than that in tumor or lung. More specifically, formulations incorporating the ligand within the cholesterol domain resulted in relatively high levels of accumulation in the liver despite low transfection rates (compare Figs. 2B and 3B). The ability of folate to stimulate specific uptake by Kupffer cells that also express folate receptors in the liver [43] may contribute to these observations.

It is well-known that cationic lipid-based nucleic acid formulations accumulate predominantly in lung and liver after intravenous injection [44]. Although the present study shows that incorporation of the folate ligand into a cholesterol domain increases transgene expression and accumulation in the tumor, this formulation still predominantly accumulates in the lung. Exclusion of the folate ligand from the cholesterol domain resulted in decreased transfection in the tumor as well as reduced accumulation in both the lung and liver, but transgene expression and accumulation remained predominantly in the lung.

The incorporation of PEG-DSPE in lipid-based formulations has been widely used to improve the circulation lifetime and reduce the liver and lung accumulation after intravenous injection, and our results with formulations employing PEG-DSPE are consistent with previous studies. However, these formulations also exhibited reduced transfection in the tumor in agreement with studies showing that even relatively low levels of PEGylation (≤ 0.5 mol %) reduce both transgene expression and DNA accumulation [28, 45]. Surprisingly, location of the PEGylated components within the domain (i.e., PEGcholesterol) exhibited a very different effect, and showed higher luciferase expression in tumor, liver and lung as compared to the non-PEGylated control. In addition, lipoplexes formulated with PEG-cholesterol delivered significantly more DNA to the lung when compared to the non-PEGylated control; just the opposite to that observed with PEGylation by PEG-DSPE. These results are very consistent with our observations in cell culture studies with KB cells, and the ester bond linking PEG to cholesterol may allow PEG to be more readily released via hydrolysis than PEG-DSPE [35]. Similarly, the cholesterol anchor may allow more rapid dissociation of the PEG moiety from the lipoplex as compared to PEG-DSPE [28, 35]. However, the release of PEG from formulations containing PEG-cholesterol cannot explain the enhanced transfection observed with this formulation. In addition, PEGylation with PEG-DSPE caused a decreased accumulation in tumor, liver, and lung, indicating that the steric stabilization provided by PEGylation cannot explain our

observations with PEG-cholesterol. Our studies in cell culture allowed us to determine that internalization was not increased by PEG-cholesterol, even though transfection was significantly enhanced [46]. Furthermore, PEG-cholesterol *reduced* transfection rates when incorporated into lipoplexes lacking a cholesterol domain, suggesting that the presence of a hydrophilic moiety within the cholesterol domain may trigger more productive trafficking that is ultimately responsible for the observed increase in transfection.

In conclusion, our results demonstrate that the microenvironment of the ligand can affect targeted delivery. More specifically, incorporation of the folate ligand within a cholesterol domain significantly increases transfection rates in tumors. The enhanced transfection correlated with an approximately 2.7-fold greater accumulation in the tumor, in contrast to previous studies concluding that accumulation is controlled exclusively by the EPR effect [22-24]. The observation that transfection correlates with accumulation in tumors even in formulations lacking a ligand (i.e., PEG-cholesterol and PEG-DSPE) suggests that the enhanced gene delivery was achieved by increasing plasmid levels within the tumor cells due to more productive intracellular trafficking. In addition, the high liver accumulation observed when the folate ligand is present in a cholesterol domain (folate-cholesterol vs. folate DSPE) suggests that the domain facilitates ligand interaction with folate receptors in the liver. We suggest that the ability of the high cholesterol formulations incorporating folate-cholesterol to enhance accumulation in vivo is due to the location of the ligand within a cholesterol domain that does not adsorb serum proteins and thereby reduces protein fouling [34, 35]. The in vivo results presented here are also consistent with our previous cell culture studies indicating that the presence of other hydrophilic moieties (e.g., PEG) within the cholesterol domain may trigger more productive intracellular trafficking that contributes to the enhanced transfection [35, 46]. We speculate that this improved trafficking may result in more plasmid molecules avoiding intracellular degradation in the endosomal/lysosomal pathway after uptake via the folate receptor. Future studies will be needed to tease out the relative contributions of increased uptake and more productive intracellular trafficking, as well as to assess the applicability of these findings to targeting with other ligands.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.

Luciferase expression in tumors 24 hours after intra-tumoral injection. The data represent the mean + one standard error of four mice in each group. Asterisk indicates a significant difference (p<0.05) as compared to the no ligand control.

Figure 2.

Luciferase expression in tumors (A), livers (B) and lungs (C) 24 hours after intravenous injection. A) Asterisk indicates a significant difference $(p<0.05)$ as compared to the no ligand control and those with folate-DSPE or PEG-DSPE. B) Asterisk indicates a significant difference $(p<0.05)$ as compared to the no ligand control. C) Asterisk indicates a significant difference (p<0.05) for comparisons between folate-cholesterol and folate-DSPE and between PEG-cholesterol and PEG-DSPE. The data represent the mean + one standard error of 8-12 mice in each group. NL: no ligand; F-chol: folate-cholesterol; F-DSPE: folate-DSPE; PEG-chol: PEG-cholesterol.

Figure 3.

DNA accumulation in tumors (A), livers (B) and lungs (C) 24 hours after intravenous injection. The data represent the mean + one standard error of 8-12 mice in each group. NL: no ligand; F-chol: folate-cholesterol; F-DSPE: folate-DSPE; PEG-chol: PEG-cholesterol. A) Asterisk indicates a significant difference $(p<0.05)$ as compared to the no ligand control and the formulation with folate-DSPE. B) Asterisk indicates a significant difference ($p<0.05$) as compared to the no ligand control and the formulation with PEG-cholesterol. C) Asterisk indicates a significant difference $(p<0.05)$ as compared to formulations with PEG-DSPE or without ligand.

Figure 4.

Correlation of accumulation and transfection in tumor, liver, and lung. Luciferase expression from each formulation was correlated with its corresponding DNA accumulation in tumor (A), liver (B) and lung (C). The correlation coefficient (R^2) was determined by linear regression as shown in the figure. The data represent the mean of 8-12 mice in each formulation.