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Synthesis and study of the complex formation of a cationic alkyl chain bola amino alcohol with DNA: *In vitro* transfection efficiency

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Abstract

Recent studies point out that bolaamphiphiles can be used in non-viral gene therapy. Due to their bipolar character, they may span a membrane and thus stabilize or destabilize it, which could be relevant for DNA transfer across a biological membrane. Since there are only very few studies on bolaamphiphile application in DNA transfection, it is difficult to assess whether they will bring additional advantages to the class of non-viral vectors. A bolaamphiphile with a hydrophobic chain of 22 carbon atoms with trimethylammonium and hydroxyl groups at each end was synthesised (22-hydroxydocos-1-yl-N,N,N-trimethylammonium bromide). It has been shown that this bolaamphiphile can transfect the green fluorescent protein plasmid in cells if mixed with a cationic helper lipid. It can compete with standard cationic lipids in terms of transfection, but is at the same

time less toxic. The potential of this class of molecules in gene delivery results from the fact that they will confer high stability to the DNA vector.

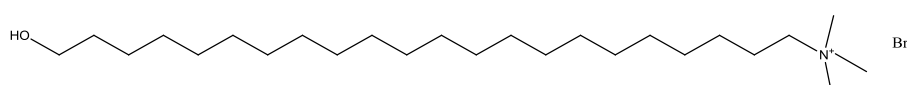
Introduction

In recent years, the research interest in non-viral gene transfection has gradually increased due to issues associated with the safety of viral vectors such as immune response and inflammation [1], activation of oncogenes and hindrance of the expression of tumor suppressor genes [2-4]. Beside these drawbacks, viral gene transfection is also expensive and has the additional limitation of the gene size: only small nucleic acid sequences can be transported by the modified viruses [5,6]. Non-viral vectors, which are polymer or lipid based, are safer, but have rather low transfection efficiency and lower stability. Thus, the research and development effort for non viral vectors is very much focused on the improvement of their transfection efficiency and stability. Bolaamphiphiles have two hydrophilic head groups linked by a hydrophobic spacer that can have one, two or three alkyl chains. These surface-active molecules can self-assemble in water, and form monolayer lipid membranes which have a higher packing density and are less permeable than lipid bilayer membranes [7,8]. Mixed with fatty acids and trimethylammonium amphiphiles, single-chain bolaamphiphiles increase the stability and encapsulation efficiency of vesicles [9]. The bolaamphiphiles have their origin in the lipids of archaeobacteria. Their membrane is formed by tetraether isoprenoid based bolaamphiphiles, rendering them resistant to high temperature and low pH [10]. The prefix “bola” is associated to the shape of an old South American missile weapon that consists of two balls connected by a cord. There are only few molecules in nature with these properties, thus most bolaamphiphiles are synthesized chemically [11,12]. Bolaamphiphiles can be assembled into distinct structures, that could be used as gene carriers [13]. Yoshimura *et al.* investigated anchored and cationic bolaamphiphiles with steroids, lithocholic acid and cholestane as the hydrophobic part of the molecule [14]. This approach was based on DC-cholesterol, which is an established transfection agent used in cationic liposomes [15,16]. Unsymmetric cationic

bolaamphiphiles similar to archaeobacterial lipids were prepared by the group of Benvegnu [17]. These compounds contain a C22 or C32 chain with a sugar moiety on one side and a quaternized glycinamide on the opposite side. The bolaamphiphile can be intercalated in lipid bilayer vesicles if the size of its hydrophobic part is close to twice of the size of the hydrophobic part of the lipid molecules [18]. Bolaamphiphiles can also acquire an U-shape to better blend with the bilayer [19-21].

Réthoré *et al.* proposed diglycerol tetraether lipids with two quaternary glycine amid moieties at both ends, also called “archaeosomes”, as gene vectors [22]. A more complex molecule containing a folic acid component connected via a polyethylene glycol chain with such an “archaeosome” molecule was synthesized and characterized with regard to gene transfection [23]. The folate residue was introduced for targeting, since folate receptors are overexpressed by human tumor cells. These bola lipid derived transfecting agents exhibit also an enhanced membrane stability. Based on the synthesis and characterization of new bola compounds [8,24] and their application as gene carrier [25,26] this study was focused on the complexation behavior of simple long chain amino alcohol membranes with DNA. We report the synthesis of a new type of bola lipid, study its aggregation, transfection and cytotoxicity in combination with DNA.

The structure of compound **BA** is shown in Figure 1. A permanent positive charge was introduced by the use of a quaternary trimethylammonio group. The other end contains the hydroxyl function as polar group. The synthesized molecule can self-assemble in water and form structures with crystal shape. The molecule was mixed with the cationic lipid DOTAP (Figure 1) to form stable vesicles with small sizes due to the two polar heads and the long carbon chain present in the bolaamphiphile [13].



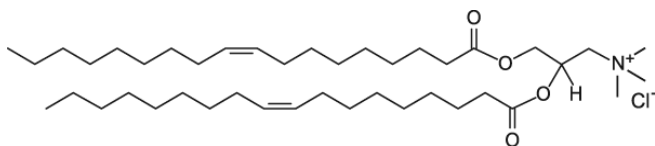


Figure 1: Chemical structures of 22-Hydroxydocos-1-yl-N,N,N-trimethylammonium bromide (**BA**, top) and 1,2-dioleoyl-3-trimethylammonium-propane (**DOTAP**, bottom).

Materials and methods

Chemicals

DOTAP (1,2-dioleoyl-3-trimethylammonium-propane, chloride salt, MW 698.542) and Rhod PE (1,2-dipalmitoyl-sn-glycero-3-phosphoethanolamine-N-(lissamine rhodamine B sulfonyl), ammonium salt, MW 1249.65) were purchased from Avanti Polar Lipids. Chloroform was acquired from Merck. All the other chemicals were supplied by Sigma-Aldrich. The solvents for the Grignard-reaction and for chromatography were dried and distilled before use. The green fluorescent protein (GFP) plasmid (pEGFP-N1, BD Biosciences Clontech) encoding for green fluorescent protein, hereafter designed as pDNA, was amplified and isolated with a PlasmidPrep Midi Flow kit (GE Healthcare) and the concentration and purity of pDNA were assessed using a NanoDrop 1000 (Thermo Scientific) apparatus. The deionized water was purified with a Milli-Q apparatus with the specific resistance of 18.2 MΩ·cm. All solutions, except those containing DNA, were filtered through a 200 nm porous membrane.

Synthesis

2-(Docos-21-en-1-yloxy)tetrahydro-2H-pyran (1): A Grignard-reagent was prepared from 0.11 mol (2.67 g) magnesium in 150 ml dry THF under inert conditions by dropping 0.09 mol (20.96 g) 1-bromoundec-1-ene into the slurry of magnesium turnings at such a rate that the temperature was always between 40 °C and 50 °C. Then the mixture was stirred for further 3 h at 50 °C. After cooling to room temperature, the Grignard reagent was decanted under inert conditions from the magnesium residue. The magnesium deposit was washed with 30 ml dry THF which was

combined with the Grignard solution. The combined solutions were cooled down to -5 °C under argon. Then, 0.059 mol (19.85 g) of 2-(11-bromoundec-1-yloxy)tetrahydro-2H-pyran dissolved in 20 ml dry THF were added at once followed by 20 ml of a 0.1 M solution of lithium tetrachlorocuprate in THF. The mixture was stirred for 3 hours at -5 to 0 °C. The mixture was poured into 200 ml ice-cold 5% ammonia chloride solution, and the product was extracted three times with 100 ml ether. The combined ethereal solutions were washed with brine and dried over sodium sulfate. After evaporation of the solvent, the crude compound **1** was purified using column chromatography with heptane/ether and ammonia (0.1 ml/100ml eluent) using gradient technique. Yield: 19.49 g (81%), white waxy solid, mp: 33 °C. ESI-MS: m/z 409.3 (M+H), 432.4 (M+NaO). Calcd. for C₂₇H₅₂O₂: C, 79.35; H, 12.82. Found: C, 79.44; H, 12.71. ¹H-NMR (CDCl₃, 400 MHz): δ = 1.23-1.38 (m, 36H, [chain]), 1.45-1.88 (m, 6H, [-CH₂-CH₂-O-, ring]), 1.99-2.04 (m, 2H, [-CH₂-CH=]), 3.32-3.40 (m, 1H, [-HCH-O-]), 3.45-3.51 (m, 1H, [-HCH-O-]), 3.68-3.75 (m, 1H, [-HCH-O-]), 3.83-3.89 (m, 1H, [-HCH-O-]), 4.55-4.57 (t, 1H, [-O-CH-O-]), 4.95 (q, 2H, [=CH₂]), 5.75-5.83 (m, 1H, [-CH=]) ppm.

22-(Tetrahydro-2H-pyran-2-yloxy)docosanol (2): To a solution of 0.04 mol (16.32 g) of compound **1** in 120 ml dry THF 80 ml of a 0.5 M 9-BBN solution in THF were added dropwise for 20 min via a syringe under stirring. The mixture was stirred for further 6 h at room temperature. At 20 °C, 28 ml ethanol, 8.48 ml NaOH (6M) and 17 ml H₂O₂ (30 %) were dropped successively into the mixture, where the temperature was raised to 50 °C. The solution was stirred further for 1h at 50 °C. After cooling down to room temperature, K₂CO₃ was added, and the organic phase was separated. The organic layer was then dried over K₂CO₃ and evaporated. The crude product was purified by column chromatography using CHCl₃/ether and TEA (0.1 ml/100 ml eluent) and the gradient technique. Yield: 15.72 g (92 %), white waxy solid, mp: 55-56 °C. ESI-MS: m/z 427.3 (M+H), 449.1 (M+Na). Calcd. for C₂₇H₅₄O₃: C, 76.00; H, 12.76. Found: C, 75.81; H, 12.52. ¹H-NMR (CDCl₃, 400 MHz): δ = 1.21-1.40 (m, 38H, [chain]), 1.44-1.82 (m, 8H, [-CH₂-CH₂-O-]),

3.34-3.40 (m, 1H, [-HCH-O-]), 3.47-3.52 (m, 1H, [-HCH-O-]), 3.64 (t, 2H, [-CH₂OH]), 3.68-3.74 (m, 1H, [-HCH-O-]), 3.82-3.889 (m, 1H, [-HCH-O-]), 4.56 (t, 1H, [-OCH-O]) ppm.

2-(22-Bromdocosan-1-yloxy)tetrahydro-2H-pyran (3): 0.03 mol (12.82 g) of the alcohol **2** were dissolved in 60 ml dry chloroform. 0.036 mol (5.0 ml) TEA and 20 mg DMAP were added to the solution. After the mixture was cooled down to 0 °C, a cooled solution of 0.036 mol (2.73 ml) methansulfonic acid chloride was added at such a rate that the temperature remained at 0 °C. The mixture was stirred for another hour at that temperature and for further 3 h at room temperature. 0.012 mol (1.6 ml) TEA in 20 ml water were dropped into the solution. The organic layer was separated, washed with 20 ml water and dried with sodium sulfate. After evaporation of the solvent, the residue was dried over P₂O₅ in vacuum. The crude methane sulfonic acid ester was dissolved in 40 ml dry acetone. After addition of 0.0875 mol (7.6 g) lithium bromide, the mixture was stirred for 3 h under reflux. The mixture was cooled down, the solvent was removed to half of the total volume and the residue was poured into 100 ml cold water. The mixture was extracted two times with 100 ml ether and the collected organic layers were washed with 100 ml water. After drying the solution over sodium sulphate, the solvent was evaporated and the residue was purified using column chromatography with heptane/ether-gradient and 0.1 ml TEA/100 ml eluent. Yield: 12,42 g (84,6 %), white pasty solid, EI-MS (m/z): 489 [M]⁺. ¹H-NMR (CHCl₃, 400 MHz): δ= 1.18-1.36 (m, 36H, [chain]), 1.35-1.6 (m, 6H, [-CH₂-CH₂-CH₂-CH₂,ring]), 1.66-1.86 (m, 4H, [-CH₂-CH₂-O-CH], [Br-CH₂-CH₂-]), 3.36-3.42 (m, 2H, [-CH₂-CH₂-O-CH]), 3.49-2.53 (m, 2H, [Br-CH₂-CH₂-]), 3.66-3.76 (m, 2H, [-CH-O-CH₂-], ring), 4.47-4.51 (m, 1H, [O-CH-O]) ppm.

22-(Tetrahydro-2H-pyran-2yloxy)docos-1-yl-N,N,N-trimethylammonium bromide (4): 5 mmol (2.44 g) of compound **3** were dissolved in 20 ml chloroform/acetonitrile (1:1). 50 mmol of an alcoholic solution of trimethyl amine (33 %) was added, and the mixture was heated to 50 °C in a closed vessel for 10 h. Afterwards, the mixture was kept for several days at room temperature

(dc control). The solvent and the excess of amine were evaporated and the crude substance was purified using mplc or by column chromatography with chloroform/methanol/water gradient technique. Yield: 2.01 g (73.4 %), white solid, mp: 160-162 °C. ESI-MS (m/z): 468.7 [M-Br]⁺. HR-MS (m/z) calcd.: 468.4781[M-Br]⁺. Found: 468.4759. Elemental analysis calcd. for C₃₀H₆₂BrNO₂: C, 65.64; H, 11.39; N, 2.55; Br, 14.56. Found: C, 65.6; H, 11.34; N, 2.46; Br, 14.8. ¹H-NMR (CDCl₃, 400MHz): δ = 1.18-1.34 (m, 34H, [-(CH₂)₁₈-]), 1.48-1.81 (m, 10H, [-(CH₂)₃-CH-O-CH₂-CH₂-],[-(CH₂-CH₂-N(CH₃)₃)]), 3.33-3.39 (m, 1H, [-CH-O-CH₂-]), 3.45 (s, 9H, [-N(CH₃)₃]), 3.48-3.55 (m, 1H, [-CH-O-CH₂-] and 2H [-CH₂-N(CH₃)₃]), 3.68-3.73 (m, 1H, [-CH-O-CH₂-]), 3.82-3.85 (m, 1H, [-CH-O-CH₂-]), 4.54-4.56 (m, 1H, [-CH₂-O-CH-]) ppm. ¹³C-NMR (CDCl₃, 100 MHz): δ = 19.7-29.74 [chain], 30.78 [-O-CH-CH₂-], 53.32 [-N(CH₃)₃], 62.3 [-CH₂-O-CH], 66.99 [CH-O-CH₂], 67.65 [-CH₂-N(CH₃)₃], 98.79 [-O-CH-O-] ppm.

22-Hydroxydocos-1-yl-N,N,N-trimethylammonium bromide (BA): 1 mmol (0.54 g) of compound **4** and 10 mg p-toluensulfonic acid were dissolved in 50 ml methanol. The mixture was heated for 3 h under reflux. The solvent was then removed to half of the volume and 50 ml water were added. The mixture was extracted three times with 50 ml chloroform. The collected organic phases were dried (sodium sulphate), and the solvent was removed in vacuum. The crude compound **BA** was purified using column chromatography or mplc with chloroform/methanol/ammonia and gradient-technique. Yield: 0.27 g (58 %), white solid, mp: 154-157 °C. ESI-MS (m/z): 384.6 [M-Br]⁺. HR-MS (m/z) calcd.: 384.4205 [M-Br]⁺. Found: 384.4195. ¹H-NMR (CDCl₃, 400 MHz): δ = 0.97-1.16 (m, 36H, [-(CH₂)₁₈-]), 1.29-1.32 (m, 2H, [HO-CH₂-CH₂-]), 1.50-1.56 (m, 2H, [-CH₂-CH₂-N(CH₃)₃]), 2.92 (s, 9H, [-N(CH₃)₃]), 3.07-3.12 (m, 3H, [HO-CH₂-], [-CH₂-N(CH₃)₃]), 3.31-3.35 (t, J=6.6 Hz, 2H, [HO-CH₂-]) ppm.

Purification. The compounds were purified using column chromatography and mplc. The ¹H and ¹³C NMR-spectra were recorded with a Varian Gemini at 400 and 100 MHz, respectively. The

solvent was used as internal standard. The EI-mass spectra were obtained with an AMD 402 (70eV) (Fa. AMD Intecta GmbH, Harpstedt). ESI-MS was performed using a Finnigan MAT 710C. For the high resolution mass spectra a Q-TOF 2 (Waters/Micromass, Manchester, UK) was used.

Preparation of Liposomes

Liposomes were prepared by the thin lipid film hydration method. Briefly, the appropriate lipid mixture (**BA**:DOTAP) at different molar ratios (1:1, 1:5 and 1:10) was dissolved in chloroform, which was then evaporated under N₂ flow. The lipid film was hydrated with 1 mL of 10 mM Hepes buffer at pH 7.4 (ionic strength 0.00325 M). The suspension was vortexed for 15 minutes, sonicated for 15 min at 60 °C and extruded at 60 °C through Whatman filters of decreasing pore sizes (down to 100 nm pore size membranes) using a Lipex pressure extruder apparatus (Northern Lipids Inc.). The total lipid concentration was 1 mM.

Dynamic light scattering and zeta potential

The hydrodynamic diameter and the zeta potential of the liposomes and lipoplexes were measured at room temperature with a Malvern Zetasizer Nano series instrument by, respectively, dynamic light scattering (DLS) and laser Doppler velocimetry. The results were averaged from three measurements for each sample and were obtained for at least three independent procedures of sample preparation. The final concentration of liposomes was 0.1 mg/mL, and the concentration of pDNA was 0.01 mg/mL.

Morphology studies

Transmission Electron Microscopy (TEM) and Confocal Laser Scanning Microscopy (CLSM) were used for the characterization of the structures formed by the bolaamphiphile **BA** alone. For the TEM (Jeol JEM-1400, JEOL) analysis, a drop of the suspension was placed on copper grids (Formvar/carbon on 400 mesh – Agar) and stained with 2% (w/v) uranyl acetate. For CLSM

imaging (Leica TCS SP5 II, Leica Microsystems), the bolaamphiphile solution was previously mixed with 0.2% (w/w) of Rhod PE lipid (fluorescent marker) in chloroform, which was evaporated and the film was hydrated, in order to mix properly the two lipids in solution.

In vitro transfection studies

The gene transfer and transgene expression after cellular transfection was monitored using pDNA encoded with GFP. The HEK 293 cell line was used in this study and was grown in Minimum Essential Medium (Lonza), supplemented with 10% (v/v) heat inactivated fetal bovine serum (FBS – Gibco), 2 mM L-Glutamine (Lonza), 100 U Penicillin/Streptomycin (Gibco) and 1% MEM non-essential amino acid solution (Sigma). Cells were cultured in a humidified 5% CO₂ atmosphere at 37 °C. For transfection studies cells were seeded at approximately 70% confluence in 12-well tissue culture plates 24 h before the experiments. Before transfection, growth medium was removed and cells were washed twice with PBS, pH 7.4. Liposome-pDNA complexes were prepared by adding equal volumes of pDNA solution at 0.01 mg/mL to the liposome suspension. The lipid:pDNA ratio was 10:1 (w:w), corresponding to a N:P ratio of 5 for BA:DOTAP liposomes (charge ratio between the cationic groups (amine) of the liposome and the anionic groups (phosphate) of the pDNA) [27-29]. DNA solution was added dropwise to the liposomes, and the mixture was kept under magnetic stirring for at least 15 minutes. Cells were then incubated with 200 µL of lipoplexes prepared with liposomes and pDNA in OPTI-MEM (Gibco). The final concentration of pDNA was 2 µg/well. The negative control corresponded to cells treated with liposomes without pDNA. After 5 h, the medium was replaced by fresh growth medium, and transfection was assessed after 48 h incubation time by flow cytometry (FACS Canto II, BD Biosciences). Twenty thousand events were measured for each sample. Experimental data were statistically analyzed applying one-way analysis of variance (ANOVA) following the Tukey test approach. Microscopy images of transfected cells were obtained using the CLSM.

Results and Discussion

Synthesis

For the preparation of the 22-hydroxy-docos-1-yl-N,N,N-trimethylammonio bromide (**BA**) the copper-catalyzed Grignard coupling of suitable building blocks was used [24,30]. Thus, the commercially available 11-bromoundec-1-ene was transformed into the corresponding Grignard reagent using THF as solvent with respect to the following coupling step. After separation of the excess magnesium the Grignard-reagent was converted into the 2-(docos-21-en-1-yloxy)tetrahydro-2H-pyran (**1**) by reaction with 2-(11-bromoundecane-1-yloxy)tetrahydro-2H-pyran with 0.1 M solution of dilithium tetrachlorocuprate below 0 °C. The terminal double bond was transformed into the primary hydroxyl group per hydroboration reaction with 9-BBN followed by common oxidation with hydrogen peroxide. Starting from **2** the introduction of the bromine atom was realized in two steps. First of all the hydroxyl function was converted to the methane sulfonic acid ester using the corresponding acid chloride. The sulfonic ester moiety was then replaced by bromine with dry lithium bromide in acetone. For the introduction of the trimethylammonio group an ethanolic solution of trimethylamine and compound **3** were heated in a closed vessel until the quaternation was complete. The final compound was delivered under mild conditions with pyridiniumtosylate in methanol under reflux.

Bolaamphiphile self-assembled structure

Attempts of extruding samples containing only compound **BA** were not successful, even at high temperatures (80 °C). The characterization of the structure by CLSM and TEM analysis shows that **BA** forms microstructures, some of which present a crystal shape (Figure 2).

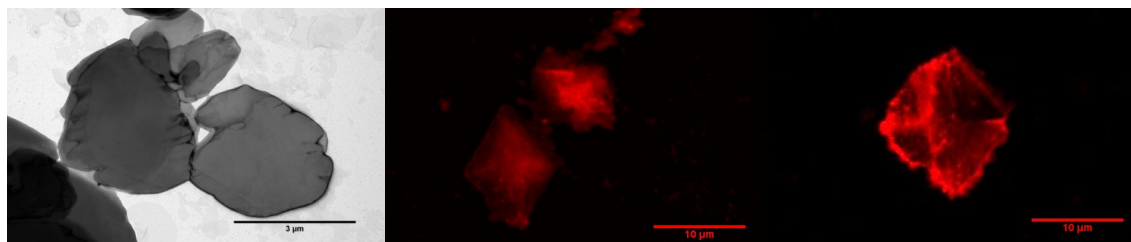


Figure 2: Self-assembled bolaamphiphile **BA** structures observed by TEM (left) and CLSM (right). Scale bar is, respectively, 3 and 10 μm .

Crystalline structures are expected, if the size of the two polar groups is not too different. For instance, they could assemble into a bilayer, where the OH groups will be oriented towards the inner space and the N- groups to the outer space (Figure 3.a). They may also form a mixed orientation where an OH group and a N- group will be in the inner space, as represented in Figure 3.b [7].

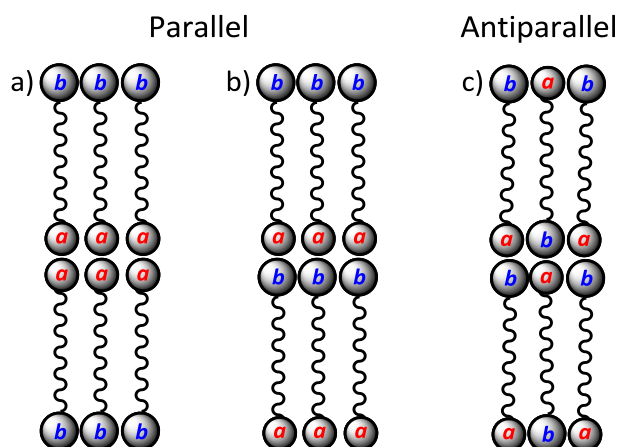


Figure 3: Schematic representation of possible arrangements of asymmetric bolaamphiphiles in crystals: a) parallel a,a; b) parallel a,b; c) antiparallel a,b and b,a (adapted from [7]).

Previous works report that these amphiphiles self-assemble into small nanoparticles (50-100 nm) [31,29], suggesting the formation of an asymmetric monolayer, which was confirmed by Langmuir monolayer studies and Grazing incidence X-Ray diffraction [19,32-34]. However, these structures are influenced by the number of carbons in the hydrophobic part of the molecule as well as the forces present in the hydrophilic part. The hydrophobic tail of compound **BA** has no double bonds, which would allow the formation of monolayer vesicles, where the OH-end group would be

directed to the inside of the vesicle and the N group to the outside. Due to its long hydrocarbon chain, **BA** can arrange itself in a U-shape, reported in the literature for similar molecules [35,36]. In this case, a bilayer would be formed and its structure could be more rigid, generating bigger particles that cannot be extruded. Mixing the compound **BA** with other lipids at appropriate ratios resulted in the formation of large unilamellar vesicles (LUV), where the bolaamphiphile **BA** was intercalated in the lipid bilayer. **BA** was mixed with a well-known cationic lipid commonly used in gene delivery [37], DOTAP, at **BA**:DOTAP molar ratios of 1:1, 1:5 and 1:10. For the ratio 1:1, the size of the structures was smaller than that obtained for **BA** alone (Figure 4a), but extrusion of the sample was still not possible. At the molar ratios of 1:5 and 1:10, **BA**:DOTAP vesicles were extruded, and it was possible to obtain large unilamellar vesicles using filters with a pore size of 100 nm. Vesicles prepared with **BA**:DOTAP (1:5 mol:mol) were used for *in vitro* cell culture studies, and the results were compared with those obtained for DOTAP liposomes. In the literature, mixtures of bolaamphiphiles with other lipids, such as cholesterol [38] and DOPE [29], have been reported mainly to stabilize the liposomes within the complexes formed with pDNA. Since the bolaamphiphile **BA** does not self-assemble into small vesicles, we propose the use of this molecule as a helper lipid to increase the stability and the transfection efficacy of lipid bilayer vesicles. **BA** was thus included into DOTAP vesicles. The zeta potential of **BA**:DOTAP (1:5 mol:mol) vesicles is slightly lower than that of DOTAP vesicles (Figure 4b), which indicates a different distribution of the positive charges at the surface of the vesicles.

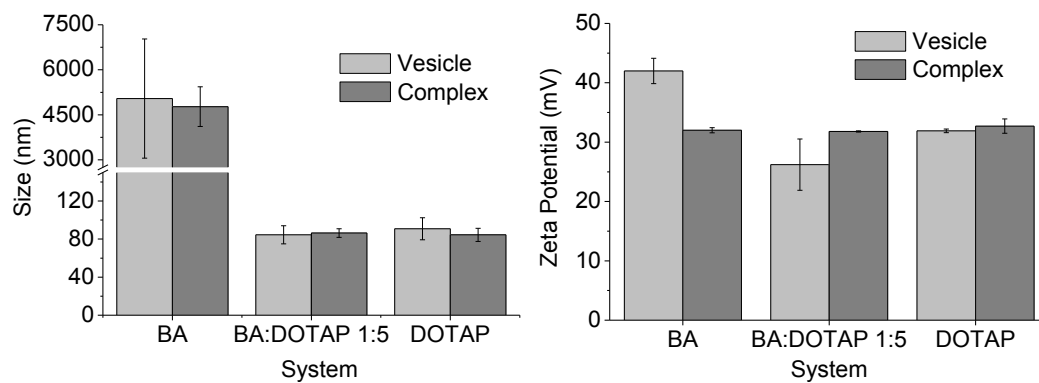


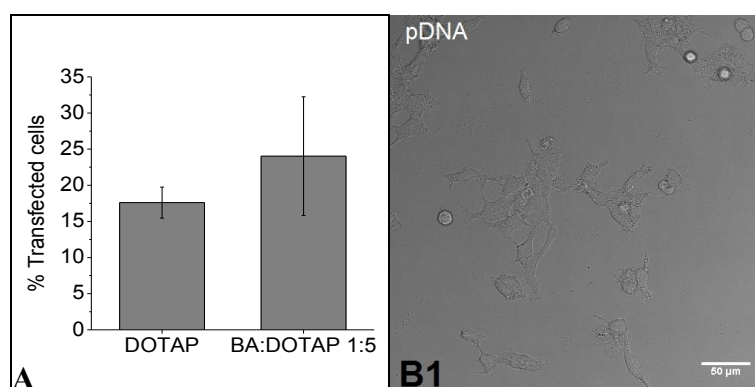
Figure 4: (a) Hydrodynamic diameter of vesicles of BA, DOTAP and their mixture at a ratio of 1:5 and (b) zeta potential values before and after the formation of a complex with DNA.

The complex formation with DNA was obtained at the ratio of lipid:pDNA 10:1 (w:w). The zeta potential values were about 32 mV for all the pDNA:vesicle complexes (lipoplexes) (Figure 4).

The size measurements of the lipoplexes reveal values similar to those obtained for the liposomes alone, however, the pDNA has an effect on the population distribution, as it seems to reduce the polydispersity index (PDI) of the sample. The parameters measured here indicate that all lipoplexes are stable and have a positive charge, which is a desirable condition for cell transfections studies.

Cell transfection studies

The transfection efficiency of the lipoplexes was evaluated using the HEK 293 cell line, a human embryonic kidney derived cell line. Cells were transfected with plasmid DNA (pDNA) encoding green fluorescent protein (GFP). Flow cytometry analysis revealed that GFP expression was induced similarly by DOTAP/pDNA and BA:DOTAP/pDNA (Figure 5A). Naked pDNA did not transfect cells (Figure 5B1).



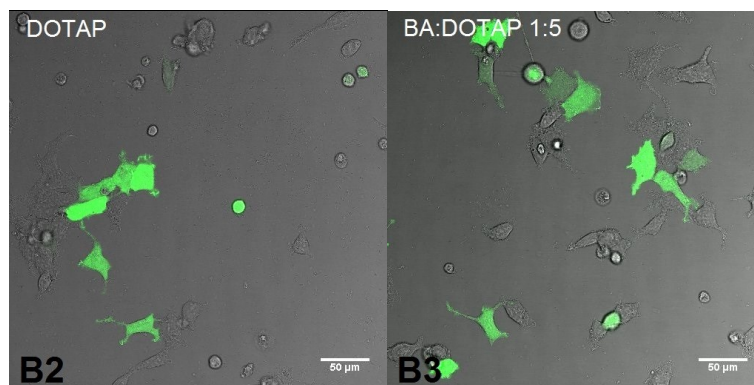


Figure 5: Transfection of the HEK cell line with green fluorescent protein (GFP) plasmid (pDNA) at a 10:1 L/D ratio: (A) cytometry analysis of transfection levels for DOTAP and BA:DOTAP 1:5 systems. Results are expressed as mean \pm S.D. (n=3). (B) Illustrative confocal images of GFP expression when transfected with naked pDNA (B1), DOTAP (B2) and BA:DOTAP 1:5 (B3). The scale bar is 50 μ m.

DOTAP has been proven to be an efficient and widely used system to complex, deliver and transfect DNA into, for example, human hematopoietic stem cells [39] and cervical cancer cells [40]. Combined with other lipids, such as cholesterol, DOTAP showed to be efficient in transporting genetic material *in vivo* [41-44]. Our results demonstrate that BA mixed with DOTAP at the molar ratio of 1:5 has the potential of being used as a helper lipid in vectors for pDNA delivery, evidenced by the positive results obtained with respect to the DOTAP system. In all cell trials, the BA:DOTAP system achieved higher values than pure DOTAP, although according to the statistical analysis, they are not significantly different at the 0.05 level, indicating that both systems have a similar ability to transfect cells. The cytometry technique and confocal analysis, however, showed that DOTAP lipoplexes cause more cellular death than BA:DOTAP lipoplexes, indicating that the latter system is less cytotoxic.

Conclusions

The properties associated to bolaamphiphiles are very promising with respect to application as delivery systems and as bio sensors [45]. Concerning the delivery of drugs/ DNA, the formation of nanoparticles, vesicles or small structures is desirable. As shown here, not all bolaamphiphiles can assemble as vesicles or other structures in the nanometer size range. This might be related to

the presence of a single alkyl chain in the bola-molecule. In a lipid membrane, the bolaamphiphile can extend completely across the membrane or form a U-shape conformation in a single membrane leaflet, depending on the length and flexibility of the hydrophobic chain. The bolaamphiphile considered here can in principle adopt both conformations: Either fully extended in a DOTAP bilayer with a thickness of approximately 3.7 nm [46] (the monomolecular layer of long alkyl monofunctional silanes (docosyl C22) has an average thicknesses of 3.5 nm [47]) or U-shape, as single-chain bolaamphiphiles form U-shapes at the air–water interface [35]. The mixed system BA:DOTAP formed stable complexes with DNA and the transfection efficiencies were similar or better than those of the pure DOTAP system. An important aspect was also the decreased toxicity of the BA:DOTAP-DNA complexes. We thus propose vesicles that combine a bolaamphiphile with cationic lipids to form stable and positively charged liposomes capable of compacting efficiently genetic material by electrostatic interactions.

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