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Designer lipids for drug delivery: from heads to tails

Aditya G. Kohli^{1,2}, Paul H. Kierstead³, Vincent J. Venditto², Colin L. Walsh^{1,2}, and Francis C. Szoka^{1,2}

¹The UC-Berkeley-UCSF Graduate Program in Bioengineering, University of California Berkeley, Berkeley, CA 94720

²Department of Bioengineering, Therapeutic Sciences and Pharmaceutical Chemistry, University of California San Francisco, San Francisco, CA 94143

³Department of Chemistry, University of California, Berkeley, CA 94720

Abstract

For four decades, liposomes composed of both naturally occurring and synthetic lipids have been investigated as delivery vehicles for low molecular weight and macromolecular drugs. These studies paved the way for the clinical and commercial success of a number of liposomal drugs, each of which required a tailored formulation; one liposome size does not fit all drugs! Instead, the physicochemical properties of the liposome must be matched to the pharmacology of the drug. An extensive biophysical literature demonstrates that varying lipid composition can influence the size, membrane stability, *in vivo* interactions, and drug release properties of a liposome. In this review we focus on recently described synthetic lipid headgroups, linkers and hydrophobic domains that can provide control over the intermolecular forces, phase preference, and macroscopic behavior of liposomes. These synthetic lipids further our understanding of lipid biophysics, promote targeted drug delivery, and improve liposome stability. We further highlight the immune reactivity of novel synthetic headgroups as a key design consideration. For instance it was originally thought that synthetic PEGylated lipids were immunologically inert; however, it's been observed that under certain conditions PEGylated lipids induce humoral immunity. Such immune activation may be a limitation to the use of other engineered lipid headgroups for drug delivery. In addition to the potential immunogenicity of engineered lipids, future investigations on liposome drugs *in vivo* should pay particular attention to the location and dynamics of payload release.

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Corresponding Author: Francis C. Szoka, szoka@cgl.ucsf.edu, School of Pharmacy, University of California, 513 Parnassus Ave., Box 0912, San, Francisco, CA, 94143, USA, 415-476-3895.

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Conflict of Interest

F.C.S. declares a conflict of interest due to his involvement in a liposome company. The other authors declare no conflict of interest.

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Keywords

Biophysics; Immunology; Liposomes; Payload Release; Targeting

1. Introduction

Lipid vesicles, or liposomes, are micro- or nano-structures formed from a bilayer of lipid surrounding an aqueous core. In the past 30 years they have been widely used to modify the pharmacokinetics, biodistribution, and cellular trafficking of drugs, nucleic acids, and proteins. Liposomal therapeutics have had preclinical and commercial success with more than 46,000 publications, 850 patents, and 13 clinically approved liposome agents with greater than \$750 million in revenue in 2011. These drugs continue to advance through the clinic, and the results from a number of pivotal phase III trials, including those from Merrimack Pharmaceuticals (NCT01494506) and Celator Pharmaceuticals (NCT01696084), will be available in the next 18 months.

Liposomes can be tailored to deliver a range of cargo using a diverse toolbox of lipids with well-characterized biophysical behavior. Lipids in this toolbox can be naturally occurring or rationally designed using a variety of hydrophilic headgroups, linkers, and hydrophobic moieties. Selecting the appropriate combination of lipids and the method of assembly provides control over liposome macrostructures, biophysical characteristics, and subsequent *in vivo* behavior.

At the most fundamental level, the properties of a liposome depend upon the subtle physicochemical interactions among the various lipid species in its composition. A wealth of research has focused on the design, synthesis and characterization of naturally occurring and synthetic lipids. Individual lipids can be combined to form a myriad of superstructures including bilayers, and bilayer properties can be tuned to modulate drug release and membrane stability (Figure 1A,B). In a simplified bilayer model acyl chain length dictates bilayer thickness and phase transition temperature (T_m), acyl chain saturation controls bilayer fluidity, and headgroup interactions impact inter- and intra-lipid molecular forces (Figure 1B). Liposome behavior can be adjusted by incorporating synthetic lipids such as lipid prodrugs, fusogenic lipids and functionalizable lipids into the bilayer (Figure 1C). As a result, there have been 50 years of synthetic efforts to develop novel lipids with properties that improve delivery while maintaining low cytotoxicity and immunogenicity. A number of databases classify lipids by structure [1], organize information related lipid T_m and phase preferences into phase diagrams [2], or detail methods for liposome characterization (cyberlipid.org; lipidmaps.org). This abundance of information provides accessible resources to guide the development of lipids for drug delivery.

As a starting point, nature has provided a variety of lipids that have evolved to satisfy diverse structural and functional purposes. Phospholipids with neutral, zwitterionic, or anionic headgroups, such as: phosphatidylcholine (PC), sphingomyelin, and phosphatidylethanolamine (PE), are the primary components of cell membranes and are essential for membrane stability and intracellular trafficking. Glycerides are neutral lipids that serve as energy sources and signaling molecules in mammalian cells. Naturally

occurring anionic lipids, including phosphatidylglycerol, phosphatidylinositol, cardiolipin, phosphatidic acid, and phosphatidylserine are also found in mammalian cell membranes, and play a critical role in cellular signaling, lipid-protein interactions, and membrane trafficking [3–7]. These naturally occurring lipids are components of FDA approved therapeutics such as Doxil®, AmBisome®, and DepoCyt® [8]. Half a century of characterization of the physicochemical properties of these lipids allows the lipid engineer to build from a wealth of structure-function relationships to design systems with control over stability and payload release.

1.1 Synthetic lipids for drug delivery

There are three key steps in liposomal drug delivery that can be improved with synthetic lipids: 1) extended circulation of the liposome after intravenous administration, 2) directed lipid headgroup interactions and cell targeting and 3) controlled payload release (Figure 2, 3). Synthetic lipids can be formulated in liposomes alongside naturally occurring lipids to serve these structural or functional roles.

After administration, liposomes circulate in the bloodstream and accumulate in tumors by the enhanced permeability and retention (EPR) effect. Increasing the circulation half-life of the liposomes allows a higher fraction of the dose to transit to the tumor and increases the probability that liposomes will extravasate into the tumor parenchyma. Functionalizing lipid headgroups with polymers, proteins, or peptides, can extend liposome circulation time by reducing liposome adhesion to mononuclear phagocyte system (MPS) cells and preventing destabilizing interactions with serum proteins (Figure 2).

The next step in delivery involves delivering payload to specific cell types by attachment of targeting ligands. While cell targeting is not a requisite liposome characteristic, a number of next-generation delivery systems look to take advantage of targeting particular cell types. To that end, engineering lipids with chemistries that allow for facile attachment of proteins, sugars, or other targeting moieties are of particular interest. In addition to targeting, lipids with specific and programmable interactions in the headgroup can direct macrostructure formation and membrane biophysics. Such synthetic lipids can be used to probe the intra- and inter-molecular forces governing superstructure formation.

In the final delivery step, liposomes must release their cargo in the appropriate tissue or cellular compartment over the desired timescale. There are a number of ways to control contents release from liposomes: 1) change the liposome formulation, 2) make the cargo more hydrophilic such that it loads into the aqueous compartment, and 3) use remote loading to cause the cargo to accumulate inside the liposome [9]. These techniques are the subject of a number of excellent reviews and will not be discussed in this article [10]. An additional method to control liposome contents release relies on including triggerable lipids that are sensitive to environmental stimuli (pH, shear stress, oxidative environment, etc.) in the liposome formulation. Such lipids allow for burst delivery in the appropriate cellular compartment and have been of particular interest in siRNA delivery, where engineered pH-dependent fusogenic lipids allow for delivery to the cytosol. These lipids have also been extensively reviewed [11,12]. Herein we focus on a number of alternatives to these triggerable systems to control liposome functionalization and payload release, including

lipid prodrugs, lipids with inverted headgroup architectures, and lipids with covalently attached hydrophobic structural or drug moieties (Figure 2).

While naturally occurring lipids are the workhorses of liposomal systems, clinical advances would not have been possible without the development of a number of synthetic lipids. In this review we focus on synthetic lipids developed over the past 10 years to probe certain aspects of liposome behavior or to improve liposome *in vivo* stability and therapeutic activity. We concentrate on lipids for drug delivery and highlight selected successes and failures of such lipids in animal studies or in the clinic.

2. Lipids that extend circulation

2.1 Polymer headgroup lipids

Conventional liposomes (CL) are cleared from circulation by the phagocytic cells of the MPS and preferentially accumulate in the spleen and liver [13]. Sterically stabilized liposomes (SL) are modified on the surface with a hydrophilic polymer that extends from the surface in a brush or mushroom configuration. This steric barrier around the liposome decreases protein adsorption to the lipid membrane, conceals surface charge, reduces liposome adhesion to cell surfaces, and, as a consequence of these factors, extends circulation time. This prolonged circulation improves the exposure of rapidly eliminated drugs by extending payload release and drug exposure and improving biodistribution to the target site. This is especially true for passive targeting strategies that depend on the EPR effect [14], in which liposomes preferentially escape the poorly organized tumor vasculature and accumulate near the tumor blood vessels. Steric stabilization further reduces the fraction of the drug that distributes to the liver, spleen, and bone marrow.

The first SLs incorporated glycolipids such as GM1 ganglioside, cerebroside sulfate, or phosphatidylinositol [15,16]. A significant breakthrough in the design of SLs came in the late 1980's with the attachment of poly(ethyleneglycol) (PEG) to the liposome surface. PEG increased circulation half-life of proteins and other biomaterials, and it was discovered that similar effects were achievable by including PEGylated lipids into liposome formulations [17,18]. Doxil®, a liposomal doxorubicin formulation incorporating PEG-2000-1,2-distearoyl-*sn*-glycero-3-phosphoethanolamine (PEG₂₀₀₀-DSPE) is the only FDA approved PEGylated liposomal therapeutic [19,20]; its development was recently reviewed [21]. By forming a steric barrier around the liposome, PEG-modified lipids embedded in a lipid bilayer decrease interactions with serum opsonins, cellular ligands/receptors and other pre-existing serum factors [22] while reducing adhesion to other membrane surfaces [23].

In the early work on PEGylated proteins it was assumed that PEG was immunologically inert [17]. However, it is now recognized that PEG can induce an immune response under certain conditions. Importantly, PEGylated drugs and liposomes can initiate the accelerated blood clearance (ABC) effect [24]. In this effect, PEGylated materials induce an anti-PEG IgM response upon the first injection [26]. Subsequent injections are then labeled for removal and are rapidly cleared from circulation and accumulate largely in the liver and spleen [24,25]. This effect reduces the widespread utility of PEG.

Other synthetic polymer-modified lipids also increase circulation half-life by providing a hydrophilic steric coat, and may serve as alternatives to PEG (Table 1, Figure 4). HPMa (poly[N-(2-hydroxypropyl) methacrylamide]) [27–29], PVP (poly(vinylpyrrolidone)) [30,31], PMOX (poly(2-methyl-2-oxazoline)) [32,33], PAcM (poly(N-acryloyl morpholine)) [34,35], PAA (poly(acrylamide)) [36], PG (poly(glycerol)) [37,38], PVA (poly(vinylalcohol)) [39–41], pNIPAM (poly(n-isopropylacrylamide)) [42], and pAAs (poly(amino acids)) [43–46] all increase the circulation half-life of liposomes *in vivo*. However, inconsistent experimental procedures make it difficult to directly compare these polymers. While only PMOX coated liposomes demonstrated similar circulation times to PEG [33], a number of favorable properties could make these alternative polymers a better choice for certain drug delivery applications [44]. In particular, polymers that avoid the ABC effect and have a lower viscosity than PEG may be advantageous for the therapeutic delivery [44]. Future studies should focus on developing, characterizing, and comparing these polymers.

2.2 Alternatives to polymer headgroups for extending circulation

Polymer coatings for liposomes are limited by their high intrinsic viscosity and their induction of the IgM-mediated ABC effect. They can also hinder liposome uptake into diseased cells [50] and fail to stop absorption of serum proteins that promote clearance such as IgG [51]. Approaches that avoid these limitations have been promising in preclinical studies. Masking nanoparticles with markers of “self” is also emerging as an alternative approach to extending liposome circulation. Red blood cell surface proteins have been investigated as nanoparticle coatings that allow for evasion of the RES. Hu and colleagues coated PLGA nanoparticles by extruding them along with disrupted erythrocyte membranes [52]. These nanoparticles, coated in an erythrocyte membrane, had extended circulation compared to nanoparticles coated with PEG or unmodified nanoparticles in mice. In a more defined approach, Rodriguez and colleagues computationally designed a 21 amino acid peptide mimetic of CD47, a marker of self on erythrocytes that impedes phagocytosis by signaling through phagocyte receptor signal regulatory protein- α (SIRP- α) [51]. Nanoparticles opsonized with IgG were coated with peptide or with a PEG brush. While PEG had no mitigating impact on macrophage uptake of the opsonized particles *in vitro* and *in vivo*, the peptide prolonged nanoparticle circulation four fold. Further, nanoparticles coated with hCD47 showed a longer circulation time than those coated with PEG. To our knowledge, this peptide has yet to be tested in a liposomal system. A number of additional factors, including shape, surface chemistry, and mechanical properties are under investigation to improve the *in vivo* properties of nanoparticles [53]. Going forward, it will be important to perform controlled studies comparing these alternatives to PEG and other polymers for extended circulation *in vivo*. Such studies should place a heavy emphasis on understanding the immunogenicity of these systems.

3. Immunological considerations in lipid headgroup design

Humoral immune responses to natural and synthetic lipids occur under a myriad of conditions in various animals [55], and immune responses to drug delivery systems have been recently reviewed by Jiskoot and colleagues [54]. B cells secrete antibodies when the B

cell receptor both possesses an affinity for a specific epitope and is activated (Figure 5). Recognition of antigen by a resting or naïve B cell occurs at the cell surface by membrane bound IgM or IgD. Subsequent B cell activation and antibody induction can occur through different mechanisms involving either a T cell dependent (T-D) response or T cell independent (T-I) response. The pathway for antibody generation during a T-D response requires protein to be processed and loaded in the major histocompatibility complex class II (MHC II) by antigen presenting cells (APCs) for T cell activation and release of cytokines for B cell activation. Activated B cells then undergo class switching to generate excreted forms of IgM, IgG, IgE or IgA depending on the cytokine stimulus from the T cell. T-D responses arising from liposomal protein and peptide antigens in the context of vaccine design have been reviewed [55] and are not the focus of this discussion. Rather, we will examine antibody responses to various non-protein molecules presented on liposomes to identify potential issues to consider in the design of novel lipids for liposomal drug delivery.

In the absence of protein and consequent T cell activation, B cells orchestrate a humoral immune response in the T-I pathway to generate IgM and IgG antibodies. Both antibody isotypes are responsible for complement activation, which may result in adverse response to non-protein antigens displayed on liposomes. There are two types of T-I responses: *T-I type 1* responses depend on toll-like receptor (TLR) agonists to activate B cells, and *T-I type 2* responses depend on multivalent ligand interactions to activate B cell receptors for activation.

In general, most naturally occurring lipids do not exhibit a T-I response. Exceptions to this generalization include the glycosphingolipids (e.g. ceramide) and acidic phospholipids (e.g. cardiolipin), which are observed in certain autoimmune disorders, such as anti-phospholipid syndrome. This syndrome results in severe blood clots leading to stroke, heart attack and miscarriage [56]. Some synthetic headgroup modified lipids have been shown to induce immune responses including polymer and hapten conjugates. As discussed, polymer modifications, including PEG [25], induces an IgM response leading to the advanced blood clearance by the MPS. This IgM response is thought to be caused by B cell activation through receptor ligation by the polymer repeat units (Figure 5B) [57]. While the ABC effect has only been observed to date with PEG, this phenomenon must be considered when developing other liposome-polymer conjugates.

Haptens do not naturally induce immune responses unless conjugated to a carrier such as a protein or liposome (Figure 5A,C). While the immune response is generated against the hapten presented on the carrier, the response persists toward the unconjugated hapten. The first immune response from a synthetic hapten-lipid conjugate was reported by Uemura and colleagues in 1974 with the attachment of 2,4-dinitrophenol or fluorescein isothiocyanate (FITC) to the headgroup of PE lipids [58]. Since then, a number of haptened liposomes have been investigated [59,60] and a number of biophysical studies have identified the role of membrane fluidity [61], linker length between the lipid and hapten [62], and hapten concentration (Table 2, *left*) [63] on immunogenicity. While antibody responses to PC lipids are rare, immune responses were observed when the PC headgroup was appended to PE lipids using a six-carbon spacer [64]. Importantly, memory IgM was also observed in response to haptened liposomes, suggesting that multiple injections would also lead to an

advanced blood clearance of the haptened formulations [65]. Several haptened liposomes have been evaluated, but the Super Hapten Database reports more than 7000 known haptens [66]. Given the vast number of known haptens and their chemical diversity, we believe that most synthetic lipid headgroups will induce an immune response.

Immunomodulatory molecules may also be incorporated in liposomes to induce specific immune responses. Adjuvants cause a T-I type 1 response through activation of the TLR signaling pathway. Monophosphoryl lipid A (MPLA), a component of endotoxin, has been the most widely studied adjuvant incorporated in liposomes and provides B cell activation as an agonist for TLR 4 [67,68]. Immune responses to haptens in the presence of other adjuvants, including nucleic acid CpG motifs [60] and Freund's Adjuvant [58], have also been investigated (Figure 5D; Table 2, *right*). Incorporation of MPLA in liposomes induces antibody responses toward the liposome in the form of MPLA specific antibodies as well as anti-phospholipid antibodies including anti-phosphocholine, anti-sphingomyelin [67] and anti-squalene antibodies [68]. Antibody responses toward the liposome may destabilize the membrane through complement dependent damage [69]. Lipid specific antibodies directed toward DMPG, MPLA, cholesterol and phosphatidyl inositol-4-phosphate (PIP) were induced in human subjects after immunization with a candidate vaccine to *Plasmodium falciparum* [70]. Although antibodies toward liposomal lipids were observed, the authors suggest that these formulations appear to have acceptable safety records and that such vaccines would be well tolerated [70]. Longitudinal studies, however, must be performed to determine the long-term effects of such antibodies in serum. Other lipid adjuvants including the lipid core peptide (LCP) system [71] or Pam3Cys [72] have also been incorporated into liposomal formulations to induce peptide specific immune responses. In each of these reports, however, the adjuvant is included in the formulations to achieve an immune response toward a specific antigen. These data indicate that it is critical to prepare endotoxin free liposomal formulations to avoid the immunogenicity of drug delivery systems [73].

3.1 Inducing immune tolerance with liposomes

Liposome encapsulated cytotoxic agents have been widely used in drug delivery applications [74], but their activity has also been exploited to induce immune tolerance. Depleting a specific population of B cells using targeted cytotoxic liposomes leads to immune tolerance against a specific epitope. Liposomes encapsulating cytotoxic agents can be targeted to specific B cell receptors. Upon interaction with the B cell membrane, the liposome is endocytosed and the payload is released, thereby ablating the specific B cell population. A number of examples of this approach exist in the literature, and Doxil® is a particularly relevant example. While PEGylated liposomes induce an ABC effect, this effect is not seen in Doxil®, as B cells with a receptor that recognizes PEG are deleted by the cytotoxic activity of doxorubicin. Deletion of these B cells leads to immune tolerance to PEG and prevents the generation of IgM and the ABC effect [75–80]. This tolerance effect has also been observed with protein agents. Liposomes displaying bovine serum albumin (BSA) and containing methotrexate were found to depress the anti-BSA plaque-forming cells in mice [81,82]. In a similar fashion, tolerance to ovalbumin was achieved in the presence of doxorubicin [83,84], which was reexamined again 16 years later and published as a novel strategy [85]. The exciting concept of tolerance induction using cytotoxic agents

is a field of research that is largely unexplored, but may be exploited for autoimmune disorders and allergic immune responses.

In summary, it is critical to be aware of the potential immune responses that may be induced by synthetically engineered lipids and novel formulations. The potential for immunoreactivity using synthetic lipids and lipid conjugates and the clearance of liposomal formulations through immune dependent mechanisms, including B cell responses, serum antibodies and complement activation, suggest that immune responses to novel lipids must be evaluated. The effect of immunoreactivity and clearance may also have implications on the pharmacokinetics, biodistribution, safety, and efficacy of liposomal formulations.

4. Lipids for directed headgroup interactions

Lipid headgroups can act as points for liposome functionalization for cargo, polymer coatings, or targeting ligands. These functionalization points often are covalent coupling moieties that allow for the attachment of ligands that alter pharmacokinetics or biodistribution, such as cell or tissue targeting groups, membrane-active peptides, or polymer coatings. In addition, headgroups contribute to lipid-lipid interactions, and influence the biophysical characteristics and macromolecular behavior of a liposome. Modifications to the headgroup allow for the control of specific molecular interactions such as H-bonding, pi-stacking, and electrostatics.

4.1 Lipids with a nucleic acid headgroup

Nucleolipids are lipids with a nucleic acid headgroup, and are unique because they can interact with nucleic acids via hydrogen bonding, pi stacking, and electrostatics. They are an excellent model system to study the hierarchy of molecular forces between lipid headgroups and how those forces contribute to superstructure formation. Further, these headgroups are biologically relevant, as they complex with single-stranded nucleic acids via base pairing and are useful in nucleic acid and drug delivery [88] and as lipid prodrugs [6]. Advancements in nucleolipid structures and their applications in transfection have recently been reviewed [86,89,90]. Herein we focus on engineered lipids with a monomeric nucleoside or nucleotide headgroup with a particular emphasis on understanding the forces governing headgroup interactions with nucleic acids.

The deoxyribose sugar is the scaffold for nucleolipid construction (Figure 6A). In general, a nucleobase (A, T, C, G or U) is conjugated to the 1' position by a beta-glycosidic linkage. This base allows for selective interactions with other nucleolipids, single-stranded nucleic acids, or drugs via electrostatics, hydrogen bonding, and pi-stacking. The 2' and 3' positions of the ribose are generally functionalized to hydrophobic domains via an ester or ether linker. Finally, anionic, cationic, zwitterionic and non-ionic groups have been conjugated to the 5' position in order to alter the behavior and distinct self-assembly properties of these lipids [89].

A few studies isolate the role of H-bonding and electrostatics in nucleolipid headgroup interactions with nucleic acids by engineering lipids that lack the functional groups required for such interactions. Ceballos and colleagues synthesized nucleolipids with 3-nitropyrrrole,

5-nitroindole or 4-nitroimidazole headgroups, universal bases known to form complementary base pairs with all four natural bases [91,92]. The headgroups are further functionalized with a quaternized amine at the 5' position and two oleyl chains to allow for lipid self-assembly (Figure 6B–D). These bases lack hydrogen-bonding capabilities, and are limited to interactions via pi-stacking interactions and electrostatics. These forces were sufficient to complex siRNA and knock down glyceraldehyde-3-phosphate dehydrogenase (GAPDH) protein levels in a number of cell lines. However, the hierarchies between pi-stacking and electrostatic interactions remain unclear in these lipids. To our knowledge, this is the first example of protein knockdown using nucleolipids transfecting siRNA. Interestingly, the authors increased the nucleolipid affinity for siRNA by altering the stereochemistry of the base at the 1' position. This increase in affinity correlates with an increase in transfection efficiency, and this relationship was corroborated by density functional theory modeling, a quantum mechanical method to investigate the electronic structure of a molecule [92]. Defining such quantitative structure-function relationships is useful in designing amphiphilic molecules [93,94].

Since hydrogen-bonding and pi-stacking interactions are weaker than electrostatic interactions via cationic groups with DNA, it is difficult to decouple their role in directing nucleolipid-DNA complexes [95]. To understand the function of these forces, Banchelli *et al.* synthesized nucleolipids with a formal negative charge [96]. Two octanoyl chains were conjugated to a nucleobase at the 5' position via an anionic phosphate linkage and an adenosine nucleotide was linked to the 1' position (Figure 6E). The hydrogen bonding and pi-stacking interactions of these nucleolipid overcame the repulsive force between the negative charge in the headgroup and the negative charge on nucleic acids in order to complex polyuridylic acid [96]. In a similar vein, Khiati and co-workers synthesized anionic nucleotide-lipids to decouple cationic interactions from H-bonding and pi-stacking (Figure 6F) [97]. Again, the forces of H-bonding and pi-stacking dominated the anionic repulsive forces between the nucleolipid and nucleic acid to complex and transfect eGFP into HEK cells *in vitro* [97]. These studies show that electrostatic interactions are not necessary to complex and transfect nucleic acids *in vitro*, as H-bonding and pi-stacking forces can be sufficient.

The selective intermolecular binding properties of nucleolipids have been incorporated into other rationally designed lipids. In a series of papers, Ma and colleagues designed synthetic multivalent hydrogen-bonding lipids with a melamine or cyanuric acid headgroup that replace the sugar-phosphate backbone of DNA with a phospholipid [98,99]. These lipids formed large unilamellar vesicles and demonstrated complementary base pairing interactions in membrane mixing and surface plasmon resonance studies (Figure 6G,H). Polidori and colleagues extensively investigated the role of lipid shape on macromolecular structure formation of synthetic lipids with a H-bonding tris(hydroxymethyl) aminomethane (tris) moiety linked with an aminoglycerol group [100]. Increasing the length of the hydrophobic tails increased the T_m of the lipids and altered the length of the tubules they formed, their stability, and their ability to stably entrap carboxyfluorescein (CF). Substitution of an ester linkage for a carbamate linkage allowed for the formation of unstable vesicles rather than stable tubes, highlighting the lipid linkage as an important parameter in lipid design and

behavior. Godeau and coworkers synthesized H-bonding glycosyl-nucleoside lipids as low-molecular-weight hydrogelators for nucleic acid delivery [101]. Lipids were first conjugated to a nucleobase using click chemistry and subsequently clicked with glycosyl groups (Figure 6I) to achieve reversible nano-fibers, hollow nanotubes, and hydrogels capable of transfecting cultured cells in serum [102]. This chemistry allows for a universal scaffold for facile lipid functionalization and engineering. Two lipid parameters dictated the formation of gels in solution: 1) H-bonding between headgroups and 2) saturation of acyl chains. H-bonding headgroups (30 fold) and saturated acyl chains (25 fold) dramatically lowered the concentration necessary for gelation [101].

While many publications have sought to simplify the synthesis, characterize the biophysics, and assess the *in vitro* behavior of nucleolipids and H-bonding lipids, little is published on their *in vivo* activity. Special attention, however, must be paid to the immunogenicity of such lipids. As previously discussed, nucleolipid headgroups may act as haptens, especially when delivering known TLR agonists such as double-stranded DNA or single-stranded RNA [103]. Such studies should be prioritized in order to understand the potential of these lipid systems for drug delivery.

4.2 Headgroups for attachment of targeting ligands

In addition to engineered lipid headgroups that rely on intrinsic molecular interactions such as H-bonding and pi-stacking, a myriad of synthetic lipid headgroups have been developed that contain specific chemistries that permit controlled covalent attachment of targeting ligands or functional groups. In particular, lipids with maleimide, avidin, ether, ester, thiol, carboxylic acid, and hydrazine moieties in the headgroup have been extensively reviewed [104,105]. The maleimide lipid is most commonly used as a functionalization point of lipids because of the reactivity of the group with a free thiol; the maleimide is typically separated from the lipid headgroup by a spacer that reduces steric hindrance in the coupling reaction. This is particularly useful, as a thiol containing ligand, such as a single chain antibody fragment, can be coupled to preformed liposomes containing a small mole fraction of maleimide lipid [106].

These covalent attachment approaches are broadly applicable but can require complex chemistry and can partially inactivate the proteins [107]. Further, the number and location of attachment sites on the ligand dictate the orientation of ligand attachment. For example, a maleimide group can form a thioether bond with any solvent exposed cysteine on a protein.

Several attempts to engineer lipid headgroups for facile non-covalent attachment have been pursued with limited success. Protein A, which interacts specifically with the Fc region of IgG, can be adsorbed to the surface of polymer nanoparticles. Addition of IgG to the nanoparticles results in uniform orientation of the antibodies and nearly 100% attachment efficiency. However, this system is limited by its stability *in vivo*, as the adsorbed protein is displaced by proteins in serum [108].

A number of groups have linked His-tagged proteins to liposomes using nickel chelating moieties [109–113]. His-tags are popular motifs that can be easily engineered into proteins and act as handles for protein purification or binding by nickel chelation. This interaction is

reversible by stripping chelated nickel from NTA [114]. In an effort to develop a general attachment approach for liposomes, van Broekhoven and Altin [115,116] engineered trivalent nitrilotriacetic acid (tris-NTA) lipids with nanomolar affinities for polyhistidine tagged (His-tag) proteins. Our group provided an alternative synthetic route to the tris-NTA lipid [117,118]. Increasing the valency of the chelating moiety in the lipid headgroup increased the affinity for His-tagged proteins (Figure 7). Yeast cytosine deaminase (yCD) and monomeric Katushka (mKate), a far red-fluorescent protein, maintained their activity while attached to the liposomes via the tris-NTA lipid. These proteins were stably attached to the liposome in fetal calf serum and mouse plasma. However, liposome attachment via tris-NTA lipids did not enhance the circulation time of proteins *in vivo*, likely due to competing interactions from plasma proteins and other histidine motifs. While the *in vivo* applications of these chelating lipids are limited, they have shown utility in subcutaneous vaccine delivery [119] and they can effectively be used to simplify rapid screening of lipid-protein conjugates *in vitro* to identify binding and internalizing antibodies for use in targeted drug delivery [120].

5. Lipids that direct membrane biophysics and payload release

Ideally, liposomes are designed to be stable in circulation until they reach the target site. Upon interaction with the appropriate compartment in target cells, the payload should be rapidly released, particularly if cytotoxicity is the objective. Hydrophilic liposome payload can be encapsulated in the aqueous core of the liposome or substituted for polar liposome headgroups in the bilayer as a lipid prodrug. In both scenarios, destabilization of the membrane and release of the therapeutic are critical steps in delivery.

As such, a keen understanding of membrane biophysics and stability is integral to tuning drug release. The T_m , phase, and composition of the bilayer drive its stability. At the T_m , the bilayer undergoes a gel to liquid phase transition causing lipid-packing defects that increase bilayer permeability. Lipid components can dictate the T_m and other bilayer physical properties. Generally, lipids with long saturated acyl chains have higher T_m than those with shorter or unsaturated chains. The inter- and intramolecular interactions of lipid headgroups further control T_m and membrane permeability. Regardless of the lipid composition, cholesterol can be included in bilayers to eliminate the phase transition and encourage a stable gel-like phase. In addition to intrinsic liposome parameters, microenvironmental factors such as redox state, pH, temperature, and enzyme activity have a profound impact on stability and can be used to trigger drug release [121,122].

5.1 Covalent attachment of hydrophobic moieties

Sterols are important components of natural membranes that play a critical role in regulating membrane fluidity. Cholesterol is the most common sterol in mammalian membranes and is the preferred sterol in several FDA approved liposome therapies including DaunoXome®, Myocet®, Depocyt®, Marqibo®, and Doxil® [123]. While cholesterol or cholesterol esters do not form bilayer structures on their own, their amphipathic nature allows for their inclusion in liposome bilayers. Incorporation of cholesterol into liposomes at 30 mol % eliminates the phase transition of diacylphospholipids [124,125], reduces membrane permeability [126,127], and forces the bilayer into a stable gel-like state. Because of their

stability, liposomes of this composition are widely used in the formulation of chemotherapeutic drugs [128]. Below 30 mol %, cholesterol does not pack uniformly in diacylphospholipids and fails to completely eliminate their phase transition [122]. Above 50 mol %, cholesterol phase separates in the membrane and can form crystals [129]. As such, changes in cholesterol composition can have profound effects on liposome formation, membrane stability, and permeability. *In vivo*, free cholesterol can rapidly transfer from liposomes into biomembranes and lipoproteins [130–132]. This constant flux of cholesterol destabilizes the liposomes and promotes contents release while in circulation.

In order to prevent such transfer of cholesterol, our group synthesized a family of sterol-modified phospholipids (SMLs) by covalently attaching cholesterol to the glycerol backbone of phosphatidylcholine (Figure 8). An early example of such a cholesterol containing phospholipid was used to investigate the role of cholesterol binding on ATPase activity in a lipid micelles but had not been demonstrated to form liposomes [133]. We showed that SMLs readily formed liposomes by themselves or when mixed with diacyl PC lipids. By anchoring cholesterol in the membrane, we reduced cholesterol transfer and increased *in vivo* liposome stability. SMLs are easily synthesized, commercially available, and recapitulate the biophysical properties of liposomes incorporating cholesterol [134–136]. Compared to liposomes formulated with free cholesterol, SMLs eliminate the phase transition of diacylphospholipids [135], exhibit similar permeability to entrapped hydrophilic molecules [135,136] and demonstrate similar membrane fluidity [136]. SMLs maintain these properties while preventing cholesterol transfer from the bilayer [135].

In vivo, SML liposomes encapsulating doxorubicin demonstrate comparable efficacy to Doxil® in a C-26 colon carcinoma model [135]. Interestingly, we found that at acyl chain lengths of C₁₆ and C₁₈, SML liposomes are more stable in circulation than liposomes containing free cholesterol. Further, we found that these SML liposomes had improved uptake into and slower clearance from the liver and spleen compared to traditional liposomes [136]. These studies highlight the stability of SML systems *in vivo* and their potential utility as drug carrier systems.

In addition to cholesterol, hydrophobic moieties for therapy and diagnostics can be covalently coupled to lipids to prevent loss in circulation. Liposomal formulations of porphyrins, hydrophobic and photosensitive agents with applications in photodynamic therapy, have been used to improve their solubility. These nanoparticle systems are limited by the amount of porphyrin that can be included in the formulation (15 mol %) and by transfer of porphyrin out of the bilayer *in vivo* [137,138]. Anchoring of pyropheophorbide, a chlorophyll-derived porphyrin analogue, to lyso PC prevented bilayer transfer of the porphyrin and enhanced its self-quenching in liposomes [137]. Lipid systems composed of these engineered lipids along with PEG-DSPE and cholesterol were termed “porphysomes”. These nanoparticles were safe at high doses in mice, demonstrated favorable pharmacokinetics, and could be loaded with hydrophilic payload. After accumulation in tumors, these theranostic particles could be imaged for diagnostic purposes or irradiated for photothermal therapy [137]. This interesting technology has been extended to applications in triggerable systems and acoustic imaging [139,140].

5.2 Inverse zwitterlipids (IZ)

In addition to engineering the lipid hydrophobic domain, modifications to the lipid headgroup can influence membrane stability and permeability. The influence of lipid headgroup properties such as hydrogen bonding capability, size and charge on superstructure formation and molecular interactions has been rigorously examined. For example, it is well known that the relative size of the lipid headgroup to the hydrophobic tails determines the phase preference of the lipid: lipids with a small headgroup (i.e. PE) prefer an unstable hexagonal phase while those with larger headgroups (i.e. PC) prefer a stable bilayer phase [10]. However, a number of aspects of lipid headgroup architecture have not been well characterized. In particular, modifications to the zwitterionic PC headgroup have been sparsely studied despite their pivotal role in biological membranes.

The influence of the position and choice of charged moieties in zwitterionic headgroups on lipid behavior is not well understood. These headgroups are oriented with an anion proximal to, and a cation distal to, the membrane interface. In the case of the PC headgroup, the anionic phosphate group is anchored at the membrane interface while the cationic amine extends into the aqueous space (Figure 9). In a series of recent studies, our group has systematically examined modifications to the PC headgroup. We set out to understand the role of 1) the relative location of charged moieties in the headgroup and 2) the type of charged moiety on lipid behavior. To that end, we synthesized a library of lipids with an inverted architecture: the cation is proximal to the membrane while the anion extends into the aqueous space. This family includes phosphates (CP), ethyl phosphates (CPe), carboxylates (AQ), sulfonates (SB) and sulfates (CS) as the anionic moiety (Figure 9). AQ, SB, CP and CPe lipids form liposomes when formulated with cholesterol. This new class of inverse zwitterionic (IZ) lipids has distinct biophysical properties from traditional PC lipids including elevated T_m and limited interactions with divalent cations. Most importantly, these studies have highlighted the pivotal role of the charge at the bilayer interface in drug permeability and ionic interactions of liposomes. While single chain surfactants with an inverted headgroup architecture have been studied, these studies extend the understanding of headgroup charge inversion in lipid systems [141].

The charge at the bilayer interface dictates the ion interactions of the liposome membrane. Liposomes composed of AQ and SB lipids interact with anions according to the Hofmeister series (Figure 10A), a classification of ions according to their ability to salt in or salt out proteins. Ions with high charge densities (F^- , Cl^-) are strongly hydrated, while ions with low charge densities (I^- , ClO_4^-) are weakly hydrated. Anions with low charge density interact with the bilayer of AQ and SB membranes more than those with high charge density. Further, all anions interact with AQ and SB membranes to a greater extent than PC membranes (Figure 10A). We believe that these interactions are directed by the positive charge at the membrane interface.

While the position of the positive charge enhances interactions with anions, it reduces interactions with cations. Divalent cations, especially Ca^{2+} , are of particular interest because they are ubiquitous in biological systems and can interact with lipid headgroups to cause membrane destabilization via aggregation, fusion, or alteration of surface charge [142]. CP, CPe, AQ and SB liposomes do not aggregate in the presence of physiological levels of Ca^{2+} ,

and IZ liposomes interact with Ca^{2+} less than PC liposomes [143–145]. Reduced divalent cation interactions make IZ liposomes interesting candidates to investigate as *in vivo* delivery systems. In addition to ions found in biological environments, the charge at the membrane interface can influence the permeation of entrapped liposome cargo. CPE liposomes are more permeable to negatively charged contents and relatively less permeable to neutral contents than PC liposomes [144]. In this way, IZ lipids may be useful to tune release properties of certain entrapped liposome drugs.

The inverted headgroup architectures also alter inter- and intra- headgroup interactions. The T_m is a measure of the intramolecular forces driving hydrophilic interactions between lipid headgroups and hydrophobic and van der Waals interactions between lipid tails. Stronger interactions in these two lipid regions lead to elevated T_m . PE lipids have strong headgroup interactions because of their H-bonding capability, and as such, have higher T_m than non H-bonding PC lipids. Similarly, lipids with long saturated acyl chains have stronger hydrophobic interactions and thus higher T_m than those with shorter or unsaturated chains. IZ lipids have very high T_m characteristic of PE lipids rather than PC lipids (Figure 10B). To our knowledge, CS lipids have the highest T_m of any lipid, natural or synthetic (Figure 10B). Interestingly, these T_m are dependent on the ionic character of the surrounding solution. For SB, CS and certain chain lengths of AQ, anion interactions with the membrane according to the Hofmeister series shift the transition from PE-like to PC-like. This shift is likely driven by inter- or intra- headgroup interactions that are disrupted when anions interact with the bilayer surface.

IZ lipids have potential as triggerable *in vivo* delivery systems. AQ liposomes have similar pharmacokinetics and biodistribution to PC liposomes, but have dramatically different biophysical properties that may be exploited for thermo-responsive systems [145]. Further, CP and CS lipids may be useful in enzyme-triggered systems. The CP lipid headgroup can be cleaved by alkaline phosphatase, yielding a lipid with a net positive charge that may destabilize the membrane and release trapped cargo (unpublished data). Such triggerable systems may be useful in drug delivery to organs enriched in phosphatase or sulfatase activity.

5.3 Triggerable lipids

As previously described, liposomes are designed to optimize stability and circulation time in order to drive a high dose of payload to the target site. However, this stability in circulation must be balanced with the release of payload in diseased tissue. Incomplete release of drug at the site of action can limit the clinical success of therapeutics, as evidenced by the failure of liposomal cisplatin in the clinic [146]. To improve release, researchers have developed a number of triggerable liposome systems. The trigger can be an external cue (heat, light, ultrasound) or intrinsic to the disease site (pH, redox environment, enzymes). These systems have been thoroughly reviewed [147,148], but have been disappointing in practice [147]. This is especially true for systems dependent on external triggers, since only primary tumors, and not nascent metastasis, can be targeted via such cues. ThermoDox® (Celsion Corporation) is a prime example of the failure of these systems in the clinic. This heat sensitive liposome formulation incorporates lysolipid that promotes liposome degradation at

mildly elevated temperatures [149] and is administered alongside radiofrequency ablation or ultrasound. However, incorporation of lysolipids can destabilize the liposome in circulation [150] and may lead to contents release before the liposome reaches the tumor site.

ThermoDox® recently failed to meet its primary endpoint in a Phase III study in patients with hepatocellular carcinoma (Celsion Corporation press release January 31, 2013).

5.4 Lipid Prodrugs

Incorporation of small molecule drugs or prodrugs into self-assembling, amphiphilic molecules can improve their biodistribution, pharmacokinetics, safety profile, trafficking and stability [3,6]. In particular, lipid prodrugs can improve the characteristics of small molecule drugs that are not well suited for traditional liposomal encapsulation and delivery due to their molecular properties. In principal, a hydrophilic drug is conjugated to a hydrophobic lipid tail such that it is substituted for a polar lipid headgroup; this amphiphilic lipid prodrug can then be incorporated into a liposome. The long history of these prodrugs have been reviewed [6]. However, a number of recent clinical advances and setbacks suggest that the clinical on the development of these drugs will be challenging. In this section we review recent developments in lipid prodrugs of doxorubicin (dox), gemcitabine, cytarabine (Ara-C), mitomycin C (MMC), and paclitaxel. We focus our discussion on clinical and *in vivo* preclinical developments.

Dox, a topoisomerase II inhibitor, is the chemotherapeutic most often studied in nanoparticle formulations because of the thorough understanding of its pharmacology and the success of Doxil® in the clinic. Systemic doses of dox lead to myelosuppression, gastrointestinal toxicity and cardiotoxicity. These issues make doxorubicin a prime candidate for nanoparticle formulation, and a number of dox prodrugs have had preclinical success. The most promising dox derivative is not a lipid prodrug, but an albumin binding dox derivative: Aldoxorubicin® (CytRX) (Figure 11). Binding of doxorubicin to albumin improves the pharmacokinetics and efficacy of the free drug and has shown promise in early clinical trials [151]. A number of lipid prodrugs of dox are in preclinical testing. A docosahexanoic acid (DHA) dox conjugate proved more efficacious than free dox in L1210 leukemia and B16 melanoma models [152]. Further, Duhem and colleagues conjugated dox to tocopherol succinate via an amide linkage which self assembled into 250 nm macrostructures when stabilized with PEG lipid [153]. These nanoparticles have improved efficacy over free dox in CT26 tumors *in vivo*. While these dox derivatives are promising, future experiments should be carried out to benchmark these dox-prodrug systems to Doxil®.

MMC is an alkylating chemotherapeutic agent limited by toxicities such as leukopenia, thrombocytopenia, and mucous membrane toxicity [154]. A number of lipid prodrugs have been developed to improve the therapeutic index of MMC. In particular, Gabizon and colleagues have developed an MMC prodrug in which the drug is conjugated to a 1,2-distearoyl glycerol lipid via a cleavable dithiobenzyl linker (Figure 11) [155]. In this system, the MMC acts as the polar lipid headgroup and the prodrug is easily incorporated into liposome membranes. This prodrug liposome system has demonstrated impressive efficacy in a range of preclinical models [155,156], and is now advancing in a phase I dose escalation study (Promitil®, LipoMedix, NCT01705002).

Nucleoside analog chemotherapeutics, including gemcitabine and Ara-C, are a major class of chemotherapeutics. Small structural differences have a profound impact on the activity of these molecules: Ara-C is primarily used to treat hematological tumors while gemcitabine, which is structurally similar except for two fluorine atoms, is used against solid tumors. Both drugs are limited by short circulation times and can be rapidly deactivated by deamination reactions *in vivo*. As such, a wealth of research has been focused on lipid prodrug modifications of these compounds to prolong circulation of the active form. Gemcitabine lipid prodrugs have shown promise in preclinical studies and are comprehensively reviewed [157]. Modification of Ara-C at the 5' position with an elaidic acid chain (Elacytarabine®, Aqualis ASA) circumvented drug resistance *in vitro* and resulted in an improvement in *in vivo* anti-tumor efficacy as compared to unmodified Ara-C (Figure 11) [158,159]. However, Elacytarabine® demonstrated no advantage in efficacy over the control arm in two phase III studies in acute myeloid leukemia and pancreatic cancer (Aqualis ASA press release November 12, 2012 and April 1, 2013). While the lipid prodrugs improved pharmacokinetic and biodistribution of Ara-C, these changes did not translate into a survival advantage in patients. Another prodrug, CP-4126, is a 5' elaidic ester of gemcitabine that has advanced into the clinic (Figure 11) [160]. Orally administered CP-4126 was poorly absorbed and subject to metabolism before systemic exposure [160]. As such, future work with this compound will focus on intravenous administration. These studies highlight the difficulties and unknowns in moving lipidated prodrugs into the clinic.

Similar to the nucleoside analog prodrugs, paclitaxel lipid prodrugs have had disappointing clinical outcomes. Paclitaxel is highly hydrophobic and its encapsulation in liposomes is problematic. Modification of paclitaxel at the 2' hydroxyl with DHA (Taxoprexin®, Protarga Inc.) improved anti-cancer efficacy by extending circulation [161]. However, these improvements in pharmacokinetics did not translate into improvements in efficacy, as Taxoprexin® had only modest activity in gastric and esophageal adenocarcinoma [162]. Further, a phase III trial in metastatic melanoma showed no survival advantage of Taxoprexin® versus dacarbazine [163]. Squalenoylation of paclitaxel and gemcitabine may represent a new class of prodrugs for clinical evaluation [164–166] but their promise must be viewed in the light of the challenges experienced by the other lipid prodrugs reviewed above.

6. Conclusions

The large variation of lipid structures outlined in this review can be combined to form liposomes with multifunctional capabilities including increased serum stability, extended circulation, ligand targeting, optimized drug loading and triggered release. These lipids have been vital tools for probing lipid membrane biophysics, especially for understanding the role of inter- and intra-molecular interactions on liposome superstructure formation and behavior. Such studies have led to a wealth of well-characterized lipid structure function relationships that are useful to those looking to synthesize novel molecules. For this reason, systemic manipulation of lipid headgroups, linkers and hydrophobic domains to further our understanding of lipid biophysics remains a worthwhile and productive area of research.

While a number of novel lipids have been synthesized and published for applications in drug delivery, their pharmacokinetics, pharmacodynamics and immunogenicity are rarely thoroughly characterized [74]. Looking forward, such studies should be prioritized in order to realize the translational potential of these systems. Special attention should be paid to commonly overlooked areas such as lipid immunogenicity, and extreme care should be taken to ensure that liposome preparations administered *in vivo* are endotoxin free to avoid confounding immunological factors.

The setbacks of Lipoplatin®, Thermodox®, Elacytarabine®, and Taxoprexin® in the clinic provide insight into the critical parameters in drug delivery to solid tumors. While in circulation, liposome payload should be stably encapsulated in order to maximize accumulation in the tumor. However, once the liposome extravasates from the vasculature into the tumor, it must quickly release its payload uniformly throughout the tumor. The complex process of payload release then involves three parameters: 1) stability in circulation, 2) distribution in the tumor, and 3) payload release over the appropriate timescale. Future work should focus on the balance between these parameters in order to optimize formulations for drug delivery. As has been written elsewhere [167], the incorporation of engineered lipids into a delivery system must take into account the higher costs, complexity of manufacturing, and complicated intellectual property of the multi-component systems. In order to justify the added costs of these systems, liposomes, like other nanomedicines, must offer significant clinical advantages in both safety and efficacy.

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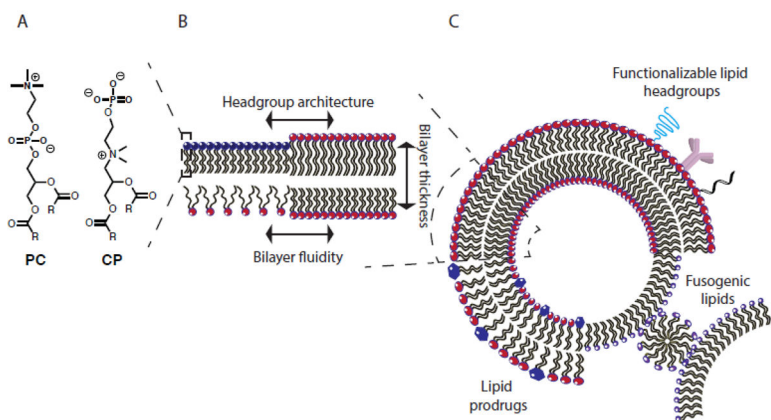


Figure 1. Modulating liposome behavior

Liposome behavior can be controlled across a number of length scales: (A) engineering of individual lipids (B) modification of bilayer biophysics and (C) inclusion of lipids that direct macroscopic liposome behavior and interactions.

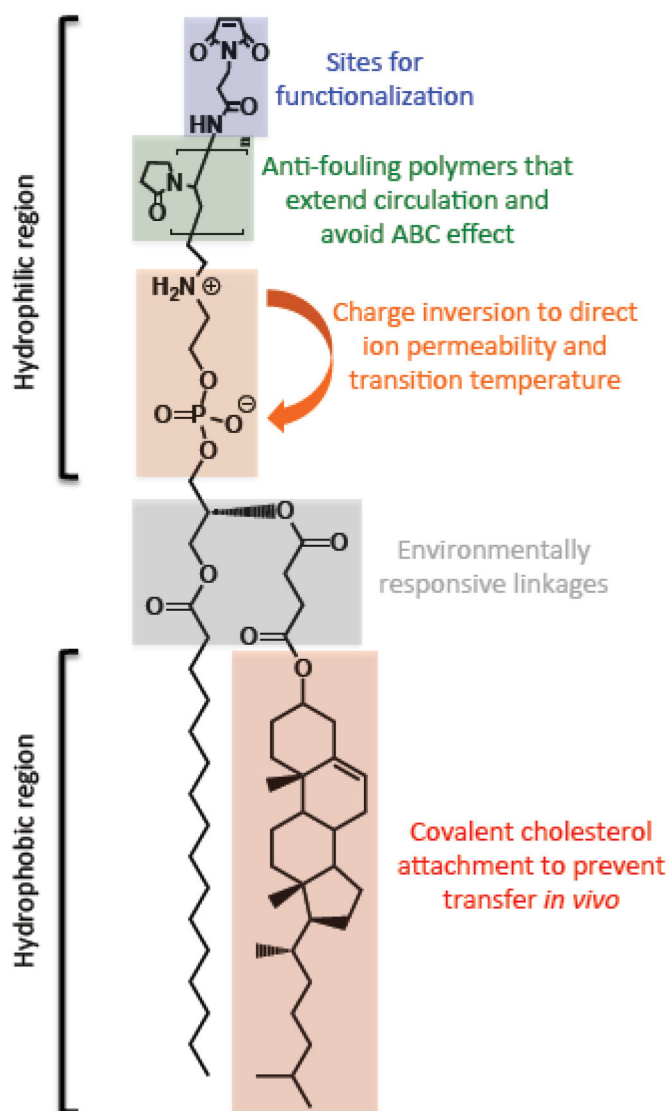


Figure 2. Overview of lipid engineering

Potential modifications to a single lipid are highlighted along with their potential functional consequences.

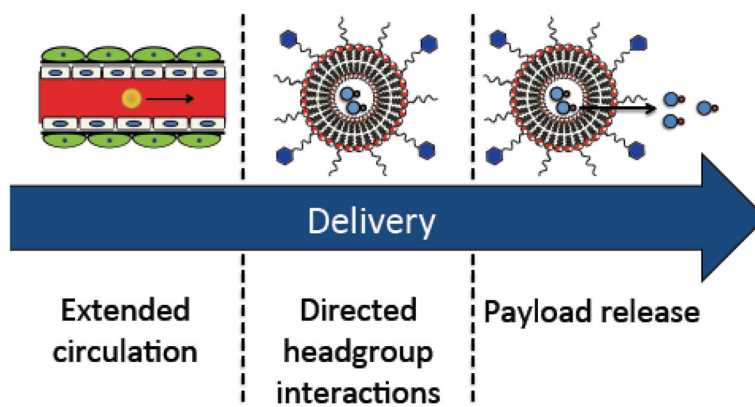


Figure 3. Three key steps in liposome drug delivery

After intravenous administration, liposomes circulate in the bloodstream and accumulate at the disease site. Directing liposome headgroup interactions allows for targeting to the appropriate cellular or tissue compartment. Finally, membrane thickness, fluidity and interfacial charge orientation control payload release.

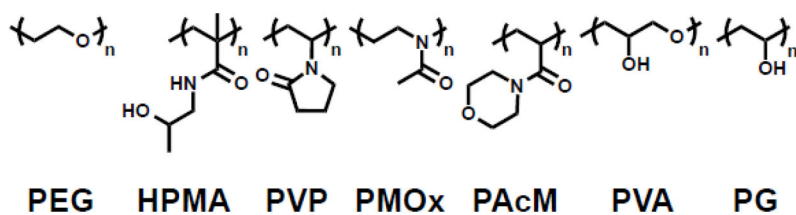


Figure 4.
Polymer headgroup lipids for extended circulation of liposomes.

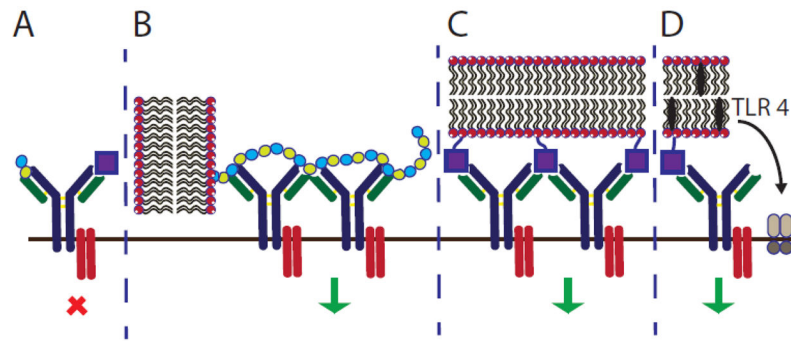


Figure 5. Pathways to B cell activation

(A) Short polymers or monomeric haptens fail to activate B cells. (B) Long polymers, such as PEG, or (C) multivalent haptens on liposomes can activate B cells by cross-linking B cell receptors. (D) Inclusion of TLR4 agonists alongside haptens can also activate B cells.

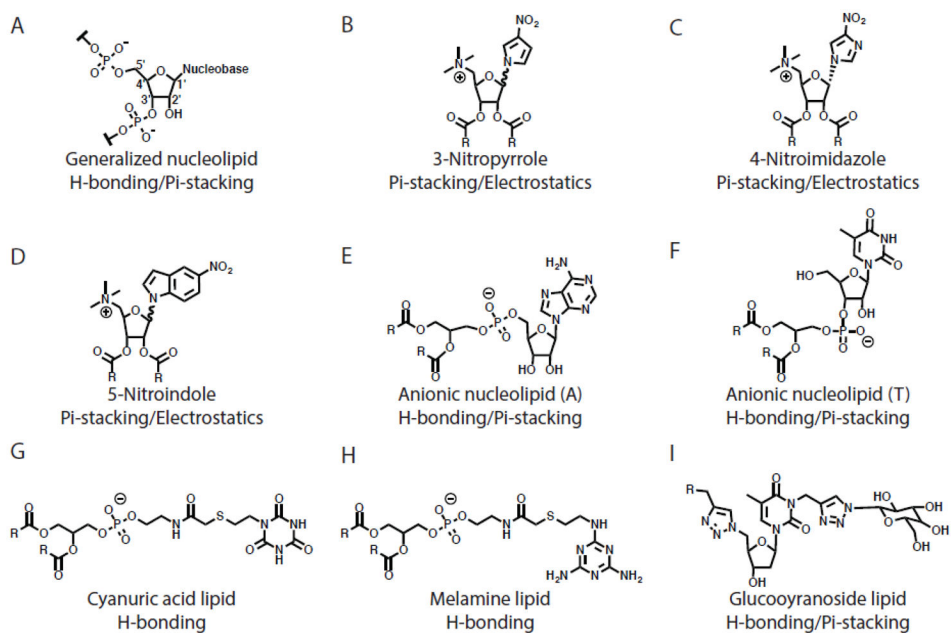


Figure 6. Nucleolipid headgroup structures

Predicted forces governing interactions with nucleic acids are shown.

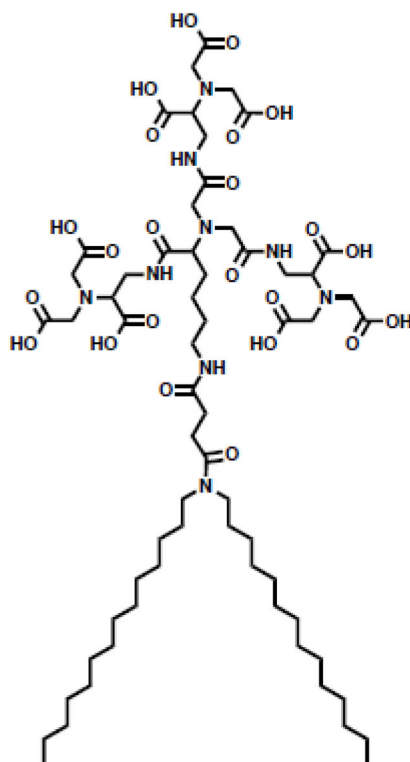


Figure 7.
Structure of Tris-NTA lipid.

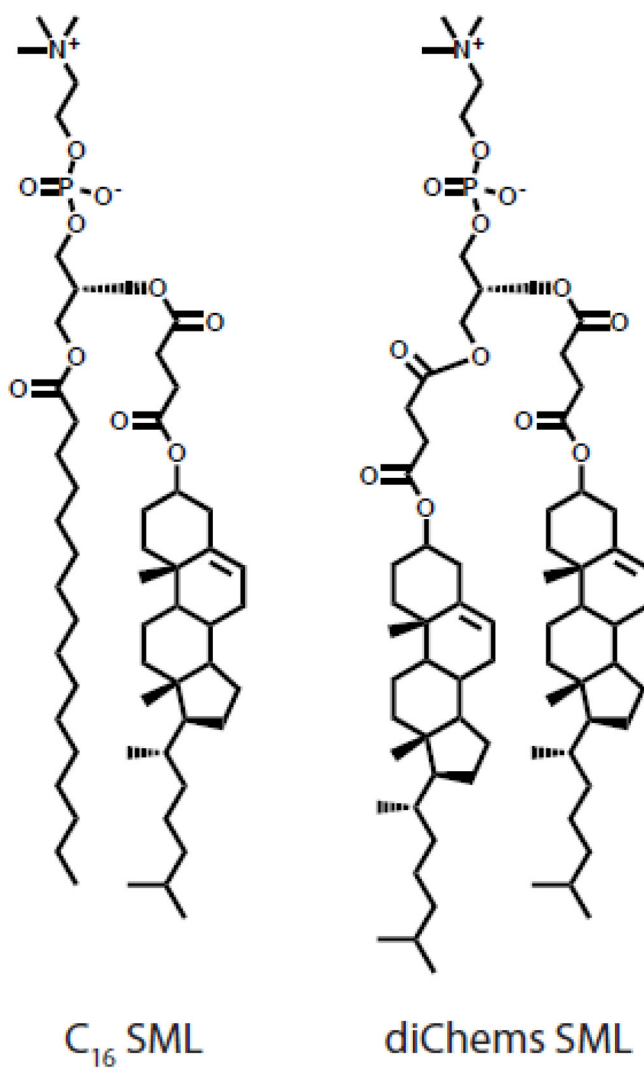


Figure 8.
Structure of sterol modified lipids (SMLs).

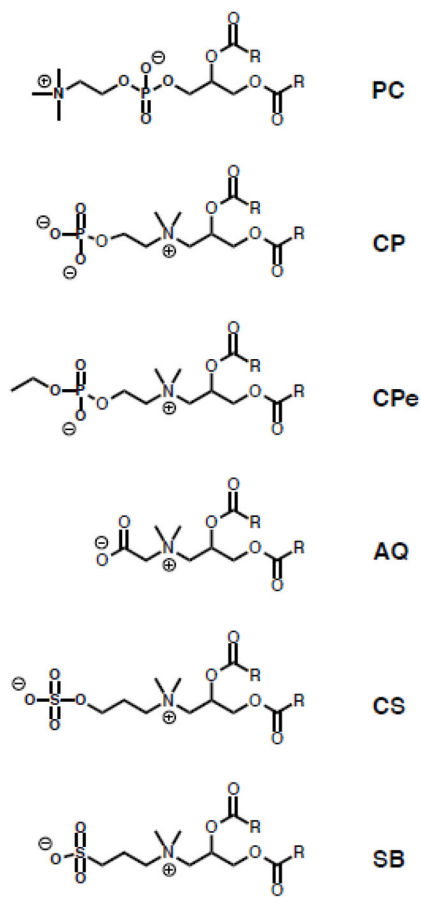


Figure 9. Structures of lipids with inverted headgroup architecture
Structures of inverse zwitterlipids (IZ) lipids.

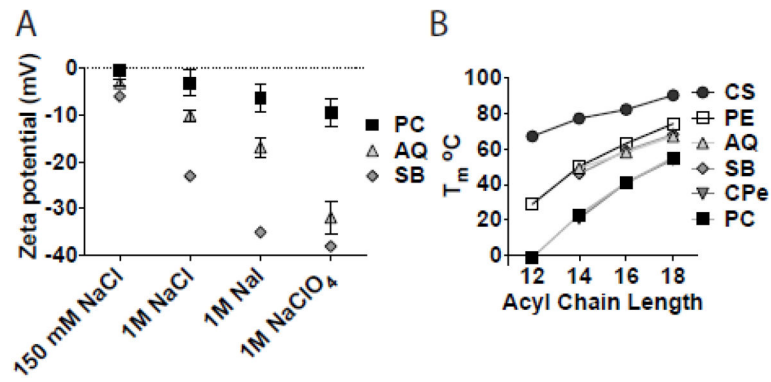


Figure 10. Biophysical properties of IZ lipids

(A) IZ lipids (AQ, SB) preferentially interact with ions according to the Hofmeister series.

(B) IZ lipids have elevated T_m compared to PC lipids.

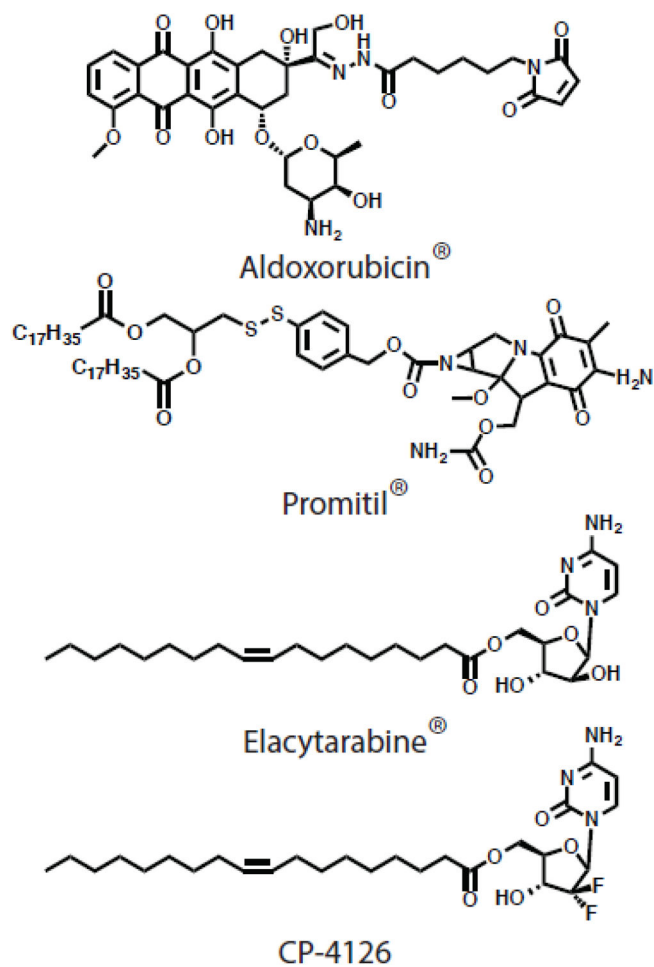


Figure 11. Prodrug structures

Selected structures of prodrugs that have advanced into the clinic.


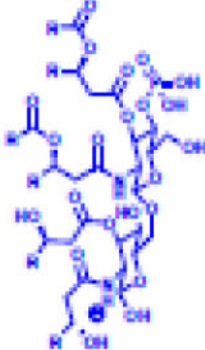
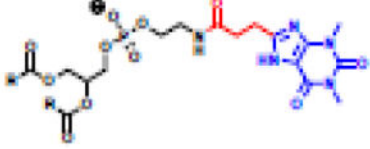


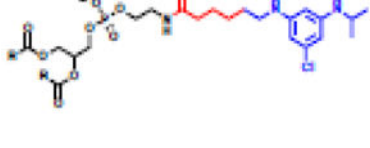
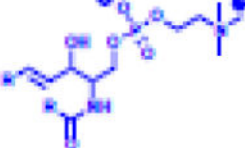
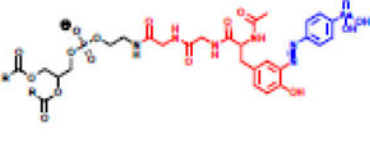
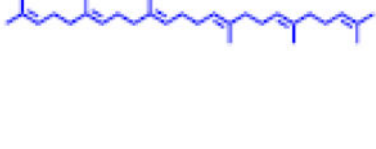
Table 1

Summary of published data on polymer modified long circulating liposomes.

Polymer	M _w	Lipid anchor	Liposome characteristics	Model	Circulation half-life	Ref.
PVP	6–15 kDa	PE/Palmityl	3 and 7 mol % polymer 155–190 nm	Mice	0.7–2.2 h	[34,36]
PMOX	2–5 kDa	DSPE	5 mol % polymer 90–112 nm	Mice/Rats	17.8 h	[33,47]
PACM	6 kDa	PE	3 and 7 mol % polymer 155–180 nm	Mice	1.5–2.8 h	[34]
HPMA	3–4 kDa	Oleic acid	0.3 and 3 mol % polymer 150–200 nm	Mice	0.8–2.3 h	[48]
PVA	6–20 kDa	(C ₁₆ H ₃₃ -S-)	0.33–8.5 mol % polymer 100–300 nm	Rats	NA	[39–41]
PG	0.2–3 kDa	DPPG/DSPE	2–15 mol % polymer 100–150 nm	Mice/Rats	8.8–22.1 h	[37,38]
pNIPAM-MAA	29 kDa	DODA	0.3 w/w	Rats	7.7–9.8 h	[42]
pAAs	3–4 kDa	PE/SA/DOD/DODA	1–15 mol % polymer 105–370 nm	Rats	2.8 h	[43,44] [45,46]
PEG	2 kDa	DSPE	5–7 mol % 100 nm	Mice	>20 h	[49]

Table 2

(Left) Selected examples of hapten conjugated lipids that induce immune responses in the absence of adjuvants. (Right) Selected examples of hapten conjugated lipids that induce immune responses in the presence of adjuvants. In all structures, the R group represents a non-specific lipid moiety.

<i>-Adjuvant</i>		<i>+Adjuvant</i>	
PEG	[25,75–80]	MPLA, MPLA	[67]
			
Theophylline	[59]		
			
TEMPO	[61]	phosphatidylcholine MPLA	[67]
			
Atrazine	[63]	Sphingomyelin, MPLA	[67]
			
Azobenzene arsonate	[65]	Squalene, MPLA	[68]
			
Dinitrobenzene	[65]	Biotin, CpG	[60]
