

Structural diversity of photoswitchable sphingolipids for optodynamic control of lipid microdomains

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ABSTRACT Sphingolipids are a structurally diverse class of lipids predominantly found in the plasma membrane of eukaryotic cells. These lipids can laterally segregate with other rigid lipids and cholesterol into liquid-ordered domains that act as organizing centers within biomembranes. Owing the vital role of sphingolipids for lipid segregation, controlling their lateral organization is of utmost significance. Hence, we made use of the light-induced *trans-cis* isomerization of azobenzene-modified acyl chains to develop a set of photoswitchable sphingolipids with different headgroups (hydroxyl, galactosyl, phosphocholine) and backbones (sphingosine, phytosphingosine, tetrahydropyran-blocked sphingosine) that are able to shuttle between liquid-ordered and liquid-disordered regions of model membranes upon irradiation with UV-A ($\lambda = 365$ nm) and blue ($\lambda = 470$ nm) light, respectively. Using combined high-speed atomic force microscopy, fluorescence microscopy, and force spectroscopy, we investigated how these active sphingolipids laterally remodel supported bilayers upon photoisomerization, notably in terms of domain area changes, height mismatch, line tension, and membrane piercing. Hereby, we show that the sphingosine-based (Azo- β -Gal-Cer, Azo-SM, Azo-Cer) and phytosphingosine-based (Azo- α -Gal-PhCer, Azo-PhCer) photoswitchable lipids promote a reduction in liquid-ordered microdomain area when in the UV-adapted *cis*-isoform. In contrast, azo-sphingolipids having tetrahydropyran groups that block H-bonding at the sphingosine backbone (lipids named Azo-THP-SM, Azo-THP-Cer) induce an increase in the liquid-ordered domain area when in *cis*, accompanied by a major rise in height mismatch and line tension. These changes were fully reversible upon blue light-triggered isomerization of the various lipids back to *trans*, pinpointing the role of interfacial interactions for the formation of stable liquid-ordered domains.

SIGNIFICANCE Sphingolipids are predominantly found at the plasma membrane of cells and are recurrently associated to segregated liquid-ordered domains with important cellular functions (e.g., signaling). Controlling the lateral organization of lipids within microdomains is thus of utmost significance. In this work, we took advantage of the light-triggerable *trans-cis* isomerization of azobenzene-modified acyl chains and designed novel photoswitchable sphingolipids with varying headgroups and sphingoid backbones. By combining atomic force and fluorescence microscopies, we were able to evaluate the ability of these photoswitchable lipids to reversibly alter the properties and amount of lipid domains on phase-separated model membranes by light, having identified the importance of sphingoid base functionality for a differential optical control of liquid-ordered properties by photoswitchable sphingolipids.

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INTRODUCTION

Sphingolipids are a major component of eukaryotic (notably mammalian) membranes and play a crucial role in signaling inside cells (1–3). Members of this lipid class, like ceramide (Cer) or sphingomyelin (SM), have a backbone formed from sphingosine and have been widely studied in terms of their biophysical properties, behavior on membrane models, and affinity to other lipids (4–11). Another important, but less studied, class of sphingolipids is phytosphingolipids, which



are abundant in plants and fungi (12–14). These lipids have phytosphingosine as a sphingoid base (2), a backbone with increased polarity compared with sphingosine. Sphingolipids are also largely localized in the plasma membranes of eukaryotic cells when compared with inner organelle membranes (3). Here, they are usually linked to so-called lipid rafts (15), liquid-ordered (L_o) phase domains composed of rigid lipid species (sphingolipids and saturated phospholipids) and sterols, thought to be a means by which cells organize or segregate important proteins within the membrane (16–18). In addition, sphingolipids have been shown to be distributed asymmetrically in the plasma membrane, being highly enriched on the outer leaflet (19).

From a molecular point of view, sphingolipids can form stable H-bond and hydrophobic interactions with other sphingolipids (e.g., SM) and cholesterol (Chol) (2,20). While *in vivo* such lipid interactions are assumed to be transient and lead to the formation of nanoscopic domains (21,22), these can, e.g., be observed *in vitro* on supported membrane model systems, as segregated rigid L_o phase microdomains embedded in a fluid bulk liquid-disordered (L_d) membrane phase (23–27). The presence of hydroxyl (–OH) groups on the backbone of sphingolipids are particularly relevant for the formation of L_o domains, as interfacial H-bonding markedly stabilizes the interactions among sphingolipids and Chol (20,28). In fact, the central role of the 3-OH moiety on the sphingosine backbone of sphingolipids, like Cer or SM, has been thoroughly scrutinized (13,29–35). Likewise, the presence of a second 4-OH hydroxyl group on phytosphingosine-based lipids, like PhCer, further strengthens H-bonding and domain thermostability (13). In contrast, hindering H-bonding by adding a methyl, ethyl, or tetrahydropyranyl (THP) group at the 3-OH hydroxyl severely affected the molecular packing and ability of those blocked sphingolipids to interact with Chol (29,36). Indeed, functionalization of the 3-OH of SM by THP greatly decreases gel-phase stability (lowering the melting temperature by 10°C), impedes tight contacts with Chol, and increases the rate of sterol desorption from vesicles containing this blocked SM analog (29).

As lipid segregation plays a crucial physiological role in biomembranes, new tools to investigate and control membrane phase properties are urgently needed. In that regard, strategies based on photopharmacology (37,38), which take advantage of the high spatiotemporal precision of light, are particularly appealing. In 2016, we reported that photo-switchable Cers (ACes), which have an azobenzene photo-switch incorporated in the lipid fatty acid chain, enable optical control of lipid raft-mimicking microdomains within synthetic membranes (39). While recent advancements demonstrated the potential of azobenzene-modified photo-switchable lipids for controlling/altering membrane properties without significant physiological perturbations (39–47), the structural diversity of these photoactive molecules is nonetheless still fairly limited. This stands in stark contrast

to the impressive diversity of sphingolipids found in nature. Indeed, diverse sphingolipids can serve as docking sites for various toxins (e.g., Shiga toxin binds to Gb3 (48) or cholera toxin B binds to GM1 (49)) or can even modulate the uptake of viruses by the cells (e.g., SV40 requires GM1 as receptor (50) or HIV-1 gp120 surface protein binds to GalCer on epithelial cells (51)). Hence, controlling their lateral organization within membranes is of vital importance. An expanded palette of photoswitchable sphingolipids could therefore offer new photoresponsive *N*-acyl azobenzene sphingolipids with more complex headgroup functionalities and other types of sphingoid backbones.

In our present work, we then aimed to develop a new set of photoswitchable sphingolipids with increased functionalization and incorporate these various photoswitchable lipids (photolipids) into L_d - L_o phase-separated supported model membranes. Our main goal was to investigate the influence of these modifications on the reversible remodeling of membrane microdomains upon photoswitching, as well as on fundamental mechanical properties of lipid bilayers. To this end, we performed atomic force microscopy (AFM) combined with fluorescence confocal microscopy, following the generated changes in domain area, height, and line tension.

MATERIALS AND METHODS

Synthesis of *N*-acyl azobenzene-modified sphingolipids

A protocol for the synthesis and analysis of all photolipids can be found in the [supporting material](#). Also, a schematic overview of all synthesis can be found on page 28 of the [supporting material](#).

Membrane model systems

Throughout this work, small unilamellar vesicles (SUVs) and supported lipid bilayers (SLBs) were used as lipid membrane model systems. These were primarily composed by *N*-stearoyl-*D*-erythro-sphingosylphosphorylcholine (C18-SM, or simply SM), 1,2-dioleoyl-*sn*-glycero-3-phosphocholine (DOPC), and Chol, which were purchased from Avanti Polar Lipids (Alabaster, AL, USA), into which different photolipids, notably Azo-Cer, Azo-PhCer, Azo-THP-Cer, Azo- α -Gal-PhCer, Azo- β -Gal-Cer, Azo-SM, or Azo-THP-SM, were mixed. Also, native *N*-stearoyl-*D*-erythro-sphingosine (C18-Cer), *D*-galactosyl- β -1,1'-*N*-stearoyl-*D*-erythro-sphingosine (C18- β -Gal-Cer), and *N*-stearoyl-4-hydroxysphinganine (C18-PhCer) were used as controls. Unless otherwise stated, the typical lipid composition was DOPC:Chol:SM:photolipid with a 10:6.7:5:5 mol ratio. For fluorescence detection, lipid mixtures were also doped with 0.1 mol % Atto655-DOPE, purchased from ATTO Technology (Siegen, Germany).

First, all lipid stock concentrations were determined in duplicate on a Mettler-Toledo UMX2 ultrabalance (Greifensee, Switzerland) with an accuracy of ± 0.1 μ g. SUVs were then obtained through bath sonication of multilamellar vesicles. Briefly, the desired lipid mixtures dissolved in cholofom:methanol (7:3) were added to a glass vial, and the solvent was then evaporated using N_2 flow, followed by vacuum drying in a desiccator. Lipids were rehydrated by adding HEPES buffer (10 mM HEPES, 150 mM NaCl [pH 7.4]), reaching a final lipid concentration of 10 mM, and then vigorously vortexed forming a suspension of multilamellar vesicles. These

were then diluted to 1 mM with HEPES buffer and sonicated in an ultrasonic bath for 10–20 min until the suspension became clear, giving rise to SUVs.

SLBs were prepared by deposition and fusion of SUVs on top of freshly cleaved mica glued on a borosilicate coverglass, as described elsewhere (39). Shortly, SUV suspensions (at 1 mM lipid concentration) were deposited in the presence of 2 mM CaCl₂ on freshly cleaved mica. The samples were then incubated for 20 min at 65°C, rinsed with HEPES buffer, and allowed to slowly cool down to room temperature for at least 1 h.

UV-visible (UV-vis) spectral and size characterization of vesicles with azo-sphingolipids

The mean number-normalized diameters and monodispersity of the formed SUVs were checked via dynamic light scattering using a Zetasizer Nano ZSP (Malvern Instruments, Worcestershire, UK), with an incident wavelength of 633 nm and a backscatter detection angle of 173°. Measurements were performed at 25°C and a total lipid concentration of 100 μM using disposable polystyrene cuvettes and a thermal equilibration time of 2 min.

UV-vis spectra of the various azo-sphingolipids embedded within SUVs were collected with Hellma SUPRASIL precision quartz cuvettes (10 mm light path) on a Jasco V-650 spectrophotometer (Tokyo, Japan) before and after illumination with UV-A or blue lights. More precisely, SUV suspensions at 150 μM lipid concentration, composed of DOPC:Chol:SM:photolipid (10:6.7:7:3 mol ratio), were here utilized.

Because of the high polydispersity of the SUVs (Fig. S1), we also examined the UV-vis spectra of extruded large unilamellar vesicles with a DOPC:Chol:SM:photolipid composition (10:6.7:5:5 mol ratio). These samples were then solubilized with 0.1% (v/v) Triton X-100 to eliminate the contribution of scattering and to more easily detect a possible shift in the main azobenzene peak. Extrusion was performed on an Avestin Lipofast extruder (Ottawa, Canada) by passing the multilamellar vesicle suspensions 21 times through polycarbonate membrane filters with 100 nm pore size.

Laser scanning confocal fluorescence microscopy

Fluorescence microscopy was performed on a Zeiss LSM510 Meta laser scanning microscope (Jena, Germany) using a water immersion objective (C-Apochromat, 40 × 1.2 W UV-VIS-IR). Samples were excited with the 633 nm line of a He-Ne laser for Atto655 excitation. Images were typically recorded with 1 Airy unit pinhole and 512 × 512 pixel resolution. Image analysis was performed using Fiji software (<http://fiji.sc/Fiji>).

Segmentation methods

In order to quantify the lipid domain (L_o phase), the confocal data were processed by a custom-made MATLAB script for batch processing. The algorithm performs basic segmentation operations based on thresholding (Otsu's method), morphological erosion, and dilation operations. The output is an image in Portable Network Graphic format of the positive mask of the dark regions corresponding to the lipid domains and a text file containing the calculated area ratio of the domains for each image.

AFM and force spectroscopy

AFM was performed on a JPK Instruments Nanowizard Ultra (Berlin, Germany) mounted on the Zeiss LSM510 Meta laser scanning confocal microscope (Jena, Germany). High-speed and normal-speed AFM, both in AC mode, were done with USC-F0.3-k0.3 ultrashort cantilevers from Nanoworld (Neuchâtel, Switzerland) with typical stiffness of 0.3 N/m. The cantilever oscillation was tuned to a frequency of 100–150 kHz, and the amplitude was kept below 10 nm. Images were acquired with a typical

256 × 256 pixel resolution. Scan rate was set to 12.5–150 Hz for high-speed AFM (i.e., line acquisition taking 6.7–80 ms and full frames 1.7–20.5 s) and to 2–6 Hz for normal-speed AFM. All measurements were performed at room temperature. The force applied on the sample was minimized by continuously adjusting the set point and gain during imaging. Height, error, deflection, and phase-shift signals were recorded, and images were line fitted as required. Data were analyzed using JPK data processing software v.6.0.55 (JPK Instruments) and Gwyddion v.2.49 (Czech Metrology Institute, Jihlava, Czechia).

Line tension (γ) was determined as previously reported (52) using the equation by Cohen and co-workers (53):

$$\gamma = \frac{\sqrt{B_s K_s B_r K_s}}{\sqrt{B_r K_s} + \sqrt{B_s K_s}} \cdot \frac{\delta^2}{h_0^2} - \frac{1}{2} \cdot \frac{(J_s B_s - J_r B_r)^2}{\sqrt{B_r K_s} + \sqrt{B_s K_s}}, \quad (1)$$

with δ being the L_q-L_o height mismatch; h the monolayer thickness with $h_0 = (h_r + h_s)/2$; B the elastic splay modulus; K the tilt modulus; and J the spontaneous curvature of the monolayer. Herein, the subscripts r and s refer to the L_o (rigid) and L_q (soft) membrane phases, respectively. For calculating the effective heights, we used the height mismatches obtained for the various samples and considered a thickness of the L_q bilayer of 3.9 nm, as measured in Fig. S7. Finally, as described by Cohen and co-workers (53) and García-Sáez et al. (52), and since the values of the elastic moduli of the model here employed are unknown, we assumed the case of flexible/soft “raft” domains with $B_r = B_s = 10 k_B T$, $K_r = K_s = 40$ mN/m, and $J_r = J_s = 0$.

Force spectroscopy measurements were performed using uncoated silicon cantilevers CSC38 from MikroMasch (Tallinn, Estonia), with a spring constant of 0.12 N/m, as previously described (54,55). Shortly, sensitivity and spring constant calibration were done via the thermal noise method. The total z-piezo displacement was then set to 300 nm, the indenting approach speed was set to 800 nm/s, the retraction speed was 200 nm/s, and the maximal setpoint was set to 5–7 nN. Force measurements were carried out at different points of the lipid bilayers. An average of 200 and 900 approach force curves per illumination state were, respectively, acquired for DOPC:Chol:Azo-THP-SM (10:6.7:10 mol ratio) and DOPC:Chol:Azo-Cer (10:6.7:10 mol ratio) SLBs. The collected curves were then baseline subtracted (offset and tilt) and corrected for cantilever bending using the JPK data processing software v.6.0.55 (JPK Instruments). Identification of the breakthrough events (and respective force values) was done using the step fit, offset, and statistical quantification functions on the JPK data processing software. Finally, the retrieved yield forces were plotted as histograms using OriginPro2015 (OriginLab, Northampton, MA, USA).

Compound switching on SUVs and SLBs

Photoswitching of the photolipid compounds was achieved using a CoolLED pE-2 LED light source (Andover, UK) for illumination at $\lambda = 365$ and 470 nm. The light source was typically operated for ~ 20 s at 80% power. For the UV-vis spectroscopic experiments with SUVs inside cuvettes, the light beam was guided by a fiber-optic cable directly to the cuvette top. For microscopic experiments, the light beam was guided by an optical fiber directly through the objective of the LSM510 Meta microscope via a collimator at the backport.

Statistical analysis

Statistical analysis of the collected AFM and confocal microscopy data was performed by one-way analysis of variance followed by Bonferroni comparison tests (95% confidence interval), using SigmaPlot 12.3 (Systat Software, Chicago, IL, USA). Tables showing the data information (mean, standard deviation, and sample size), as well as the recovered t statistics and p values can be found in the [supporting material](#).

RESULTS AND DISCUSSION

Synthesis of photoswitchable sphingolipids

Inspired by the structural design of our simpler azo-Cers (ACes) (39) and more complex α -galactosyl-phytoceramides (40), we introduced five new azobenzene-modified sphingolipids, namely Azo-PhCer, Azo-THP-Cer, Azo- β -Gal-Cer, Azo-SM, and Azo-THP-SM. These photolipids featured (Fig. 1 A) 1) a FAAzo-4 fatty acid (37) at the *N*-acyl chain (equivalent to a $\Delta 9$ unsaturation when in the *cis*-isoform) and 2) sphingoid backbones based on naturally

occurring sphingosine and phytosphingosine or hydroxyl-blocked sphingosine with a THP protecting group, as well as 3) lipid headgroups presenting either a free -OH, galactosyl, or phosphocholine moiety. For our comparative studies, Azo-Cer (previously named ACe-1) (39) and Azo- α -Gal-PhCer (40) (previously named GalACer-4), having the same FAAzo-4 moiety able to undergo *trans-cis* photoisomerization (Fig. 1 B), were also assessed.

The synthesis and characterization of Azo-Cer and Azo- α -Gal-PhCer were reported elsewhere (39,40). Azo-PhCer was prepared analogously to Azo-Cer by the coupling of

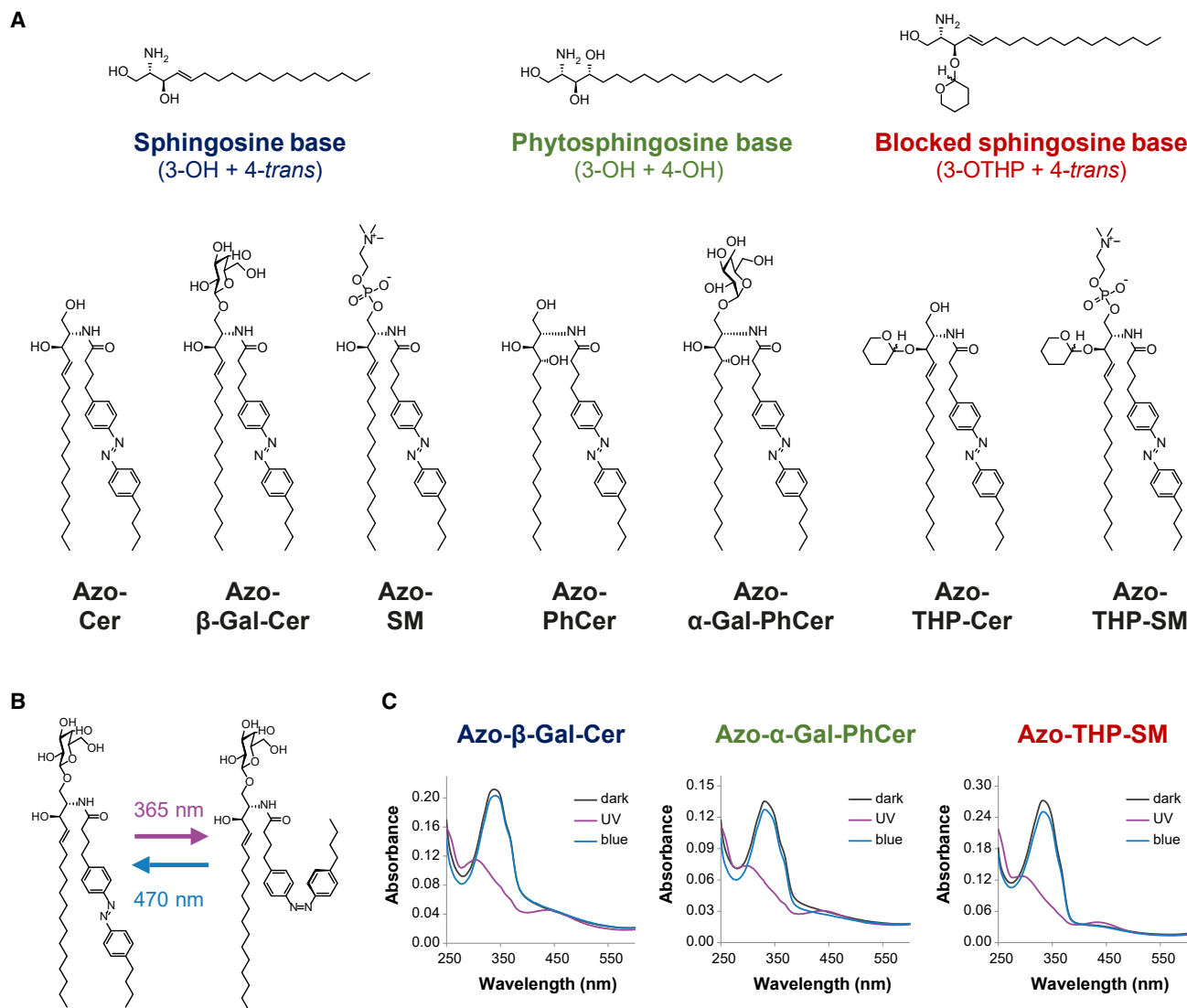


FIGURE 1 Structural and spectral properties of photoswitchable azo-sphingolipids. (A) Chemical structures of the *N*-acyl azobenzene-modified (FAAzo-4 fatty acid) sphingolipids here tested, subdivided according to their sphingoid backbone: Azo-Cer, Azo- β -Gal-Cer, and Azo-SM with a sphingosine base and Azo-PhCer and Azo- α -Gal-PhCer with a phytosphingosine base, as well as Azo-THP-Cer and Azo-THP-SM displaying a 3-OH-blocked sphingosine base with a THP protecting group. (B) Schematics of light-induced *trans-cis* isomerization for an azo-sphingolipid, notably Azo- β -Gal-Cer. Application of UV-A light ($\lambda = 365$ nm) leads to the formation of a *cis*-photolipid, while illumination with blue light ($\lambda = 470$ nm) leads to the formation of a *trans*-photolipid. (C) UV-vis absorbance spectra of photoswitchable sphingolipids (notably Azo- β -Gal-Cer, Azo- α -Gal-PhCer, and Azo-THP-SM) incorporated in SUVs at the dark-adapted state (black curves), as well as after the sequential shining of UV-A (purple curves) and blue light (blue curves). To see this figure in color, go online.

phytoceramide with FAAzo-4 using HBTU as a coupling agent (see [supporting material](#)). Additional protecting group manipulations yielded Azo-THP-Cer.

For the synthesis of Azo- β -Gal-Cer, we used a benzoyl protected alcohol and the azide as protecting groups (56) (see [supporting material](#)). Azides do not coordinate to the primary alcohol, and thereby the nucleophilicity of the sphingosine is greatly enhanced. Glycosylation of azidosphingosine with trichloroacetimidate yielded protected glycoside in 92% yield and excellent β -selectivity. Staudinger reduction using PBU_3 and subsequent amide coupling with FAAzo-4 (37) using 1-ethyl-3-(3-dimethylaminopropyl) carbodiimide, followed by global deprotection gave Azo- β -Gal-Cer (see [supporting material](#)).

The SM derivatives were prepared from Azo-THP-Cer, which was phosphorylated using 2-cyanoethyl- N,N,N',N' -tetraisopropylphosphorodiamidite and 1H-tetrazole, followed by reaction with choline tosylate (see [supporting material](#)). An oxidation directly yielded Azo-THP-SM. Finally, deprotection under acidic conditions gave the unprotected Azo-SM (see [supporting material](#)).

Light responsiveness of membrane-embedded azo-sphingolipids

Next, we incorporated our newly synthesized **Azo- β -Gal-Cer**, **Azo-SM**, **Azo-PhCer**, **Azo-THP-SM**, and **Azo-THP-Cer** photolipids, as well as **Azo-Cer** and **Azo- α -Gal-PhCer**, into L_d - L_o phase-separated mixtures containing DOPC, Chol, and SM (18:0-SM) and formed SUVs as previously described (39,57). Unless otherwise stated, a molar ratio of 10:6.7:5:5 (DOPC:Chol:SM:photolipid) was typically chosen.

We started with dynamic light scattering to determine the typical size and monodispersity of the sonicated SUVs. As shown in [Fig. S1](#), the average number-normalized diameters were between 40 and 60 nm. Then, we collected UV-vis spectra of those various SUV suspensions and characterized the photodynamic properties of the different azobenzene-modified sphingolipids within a membrane environment. As seen in [Figs. 1 C](#) and [S2](#) (see [supporting material](#)), the azo-sphingolipids displayed an absorbance maximum, λ_{max} , at ~ 340 nm when in the dark-adapted state prior to irradiation with UV-A or blue light. This peak corresponds to the $\pi \rightarrow \pi^*$ transition and is characteristic for the *trans*-azobenzene isoform. First illumination with UV-A light ($\lambda = 365$ nm) led to the reduction of the abovementioned absorbance peak and to the appearance of a new λ_{max} at ~ 450 nm. This new peak corresponds to the $n \rightarrow \pi^*$ transition and is characteristic for the *cis*-azobenzene isoform. Subsequent irradiation with blue light ($\lambda = 470$ nm) led then to a back isomerization of the *N*-acyl azobenzene moieties into the *trans*-isoform, as confirmed by the disappearance of the absorbance peak at ~ 450 nm and the reemergence of the absorbance peak at ~ 340 nm.

Our results indicate that the tested photoswitchable sphingolipids are in the *trans*-configuration for the dark- and blue light-adapted states and are mostly in *cis*-configuration for the UV light-adapted state. To infer if significant spectral shifts occur due to headgroup functionality and sphingoid polarity, we performed additional experiments examining the UV-vis spectra of extruded large unilamellar vesicles with a DOPC:Chol:SM:photolipid composition (10:6.7:5:5 mol ratio), followed by solubilization with 0.1% (v/v) Triton X-100 to eliminate the contribution of scattering. When we normalize the spectral maxima ([Fig. S2 E](#)), we find a stronger spectral red shift of ~ 7 nm for the blocked THP photolipids (maximum 338 nm) compared with the azo-sphingolipids and azo-phytosphingolipids (maximum 331 nm). Thus, it appears that the presence of the THP moiety only slightly affects the spectral azobenzene properties of our photolipids.

Subsequently, we deposited SUVs composed of quaternary DOPC:Chol:SM:photolipid mixtures doped with 0.1 mol % Atto655-DOPE (dye labeling the L_d regions) on top of freshly cleaved mica to form SLBs via vesicle fusion. By collecting fluorescence confocal images, we first evaluated membrane integrity and the presence of phase separation for samples having sphingosine- (Azo-Cer, Azo- β -GalCer, Azo-SM), phytosphingosine- (Azo-PhCer, Azo- α -GalPhCer), and blocked THP-sphingosine-based (Azo-THP-SM) lipids. As seen in [Figs. S3](#) and [S4](#), all tested SLBs displayed L_d - L_o phase separation at the dark-adapted state, with μm -sized rigid L_o domains (*dark areas in fluorescence images*) segregated within a fluid L_d matrix (*red areas in fluorescence images*).

The photoresponsiveness and ability of the azo-sphingolipids to then remodel/reorganize phase-separated SLBs were further assessed directly after irradiation with UV-A light ($\lambda = 365$ nm). For membranes lacking a photolipid (control sample with a DOPC:Chol:SM composition at a mol ratio 10:6.7:10), no light-induced remodeling was observed ([Fig. S3 A](#)). In contrast, for SLBs containing azo-sphingolipids, stark lipid rearrangement dependent on the amount of photolipid present was reported. Here, lipid bilayers with the highest amount of azo-sphingolipid tested (18.7 mol %; DOPC:Chol:SM:photolipid with mol ratio 10:6.7:5:5) showed strong reorganization of the L_o domains with admixing of fluorescently marked L_d lipids and blurring of the domain boundaries directly after UV-A irradiation ([Fig. S3, B–G](#)). Intermediate amounts of photolipid (11.2 mol %; DOPC:Chol:SM:photolipid with mol ratio 10:6.7:7:3) led to a lower domain remodeling activity. Finally, for SLBs with the lowest amount of azo-sphingolipid tested (3.7 mol %; DOPC:Chol:SM:photolipid with mol ratio 10:6.7:9:1), only a minor admixing of L_d - L_o domains was observed ([Fig. S4](#)). Thus, the higher the concentration of azo-sphingolipids on the membranes, the stronger the reorganization of the phase-separated domains directly after UV-triggered photoisomerization of the *N*-azobenzene acyl

chain of lipids into *cis*. Specifically, the observed mixing of the L_d marker from the highly fluorescent L_d matrix regions to the nonfluorescent L_o domain regions is evidence that lipid content is exchanged between the two domains immediately after UV illumination. To be precise, this phenomenon hints that the membrane gets “fluidified” once the photolipids are switched, as L_d -localizing lipids appear to “populate” (i.e., diffuse into) the more rigid L_o domains.

Remodeling of membrane domains by sphingosine-based azo-sphingolipids

After the initial characterization and screening mentioned above, we systematically investigated the light-induced remodeling of phase-separated supported membranes with 18.7 mol % photolipid (DOPC:Chol:SM:photolipid, 10:6.7:5:5 mol ratio) by first performing high-speed AFM to gain more insight into the activity of the photolipids. AFM enables us to capture minor dynamic changes in membrane architecture very accurately due to its exquisite sub-nm resolution. Herein, we started by analyzing bilayers containing the sphingosine-based photolipids Azo-Cer, Azo- β -GalCer, and Azo-SM.

At the dark-adapted state, those various membranes displayed a height mismatch between the L_d and L_o regions of 1.1–1.7 nm, similar to what we reported before for phase-separated bilayers with Azo-Cer (ACes) lipids (39). Application of UV-A light ($\lambda = 365$ nm) to bilayers having Azo-Cer, Azo- β -GalCer, or Azo-SM led then to the generation of an L_d phase, with no major alteration of the L_d - L_o domain height mismatch (i.e., minor increase of ~ 0.1 nm).

As seen from single images in Figs. 2 A and S5 B and Videos S1, S2, S3, S4, S10, S11, and S12, the UV-induced isomerization of the *N*-acyl chains from a straight *trans*-form into a kinked *cis*-form promoted an apparent “fluidization” of the phase-separated membranes, with an overall robust decrease of the total L_o area for all three azo-sphingolipids (Fig. S6 C). Directly after irradiation with UV-A light, small nanoscopic L_d “lakes” were formed within the more rigid, thicker L_o domains, consistent with the influx of L_d markers into the L_o regions observed in Fig. S3. Because the height of the L_o domains is largely unaffected by the isomerization of these photolipids in *cis*, the nanoscopic L_d lakes detected by AFM may be the reservoirs into which the fluorescent L_d markers distribute immediately after photoisomerization. These membrane nanopockets with diameters below the diffraction limit would be poorly resolved in fluorescence microscopy, which explains the fluorescently smeared appearance of the L_o regions when detected under the confocal microscope (Figs. S3 and S4).

The number of fluid L_d lakes then rapidly dropped in order to reduce surface tension. While the majority of the smaller nanoscopic L_d lakes seem to vanish toward the outer fluid L_d matrix, a few larger fluid L_d lakes remained trapped

inside rigid L_o domains, appearing to grow primarily via domain fusion and/or Ostwald ripening (58). Finally, if all the high-speed AFM area values (Figs. 2, A and C, and S5 B; $n = 7$) for this class of photolipids are grouped, an average L_o area decrease of $\sim 25\%$ (area $L_o(\text{UV})/\text{area } L_o(\text{dark}) = 0.75 \pm 0.14$) can be estimated, as depicted in Fig. S6.

After a few minutes of equilibration, subsequent application of a brief pulse of blue light ($\lambda = 470$ nm) to those phase-separated membranes reversed the effect, with an L_o phase being generated. Here, back isomerization of the *N*-acyl chains from a kinked *cis*-form into a straight *trans*-form, stimulated by blue light, promoted a rigidification of the membranes (Figs. 2 A and S5 B), with an increase of the total L_o area (Fig. S5 C) back to its original equilibrium dark-adapted value (area $L_o(\text{blue})/\text{area } L_o(\text{dark}) = 1.01 \pm 0.18$; from grouped high-speed AFM area values in Fig. S6). More specifically, upon irradiation with blue light, small rigid L_o “islands” were firstly formed within the fluid L_d matrix (Videos S1, S2, S3, S4, S10, S11, S12, and S13). These taller L_o islands then vanished, as preexisting L_o domains grew primarily via Ostwald ripening and domain fusion. Moreover, L_o domains displayed height values like the ones reported before for the initial dark-adapted state. Interestingly, those changes could be repeated over multiple cycles without dissipation effects, with the amount of L_d - L_o phase separation alternating between two defined steady states (or area levels) (Fig. 2, C and D).

Besides changes in L_d - L_o phase separation, we also observed sporadic generation of holes on our supported bilayers after the blue light-triggered conversion of the azo-sphingolipids’ *N*-acyl chains from *cis* to *trans* (Fig. S7 A). The presence of holes allowed us to recover the total membrane thickness, which was ~ 5.2 nm (L_d thickness ~ 3.9 nm; Fig. S7, B and C), in agreement with previously reported values for membranes of comparable lipid composition (59–61).

In summary, the tested azo-sphingolipids with a sphingosine backbone display a similar photoswitching profile independently of the type of headgroup. These lipids are able to increase the amount of L_d phase on phase-separated membranes upon conversion to the *cis*-isoform after UV-A light irradiation and increase the amount of L_o phase upon conversion to the *trans*-isoform after irradiation with blue light.

Remodeling of membrane domains by phytosphingosine-based azo-sphingolipids

Next, we recapitulated the same high-speed AFM procedures on membranes with photoswitchable phytosphingosine-based sphingolipids displaying two hydroxyl groups (3-OH + 4-OH) on the phytosphingosine backbone. More precisely, we investigated the ability of these photolipids to interfere with the L_d - L_o phase separation on SLBs

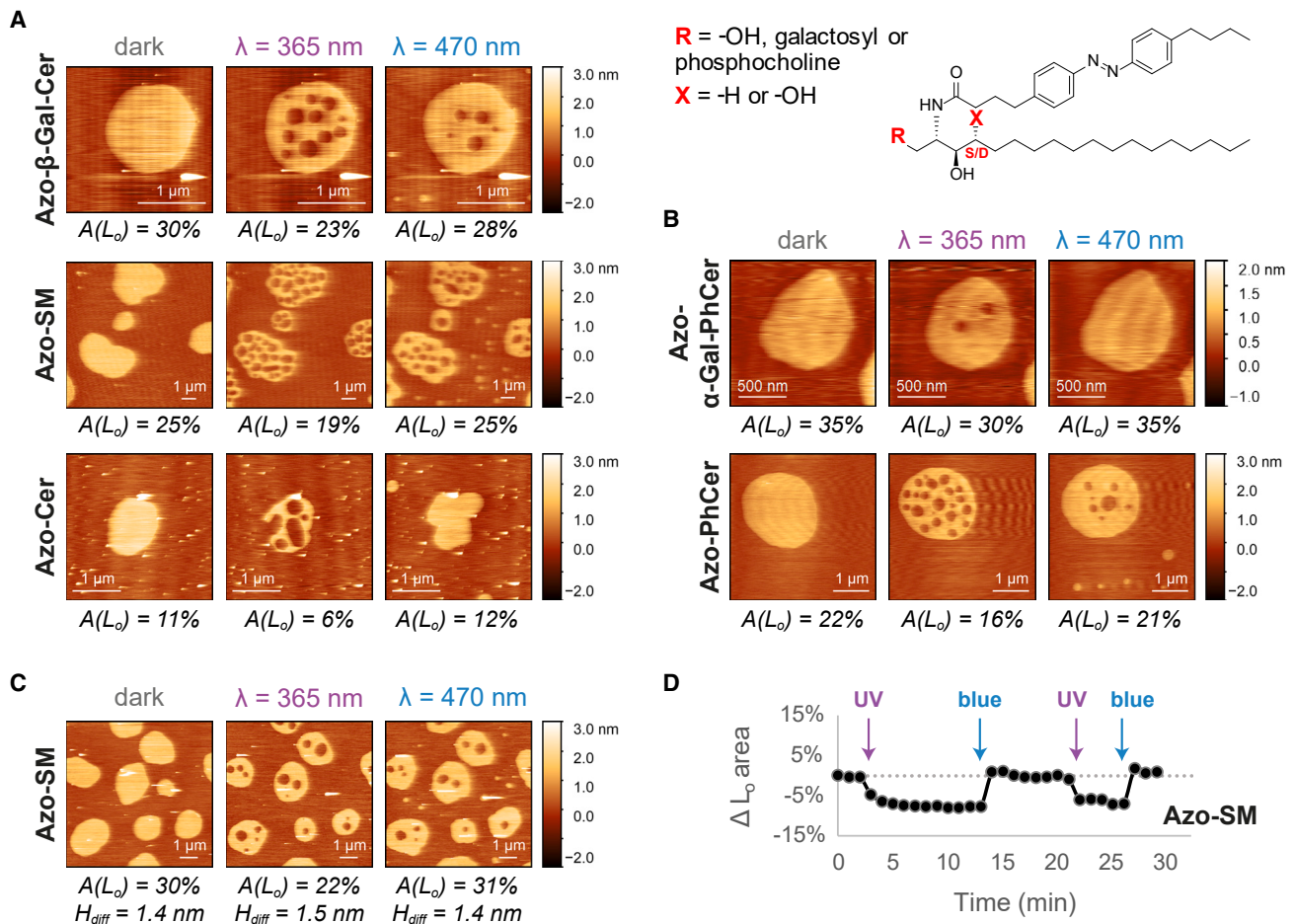


FIGURE 2 Lateral remodeling of phase-separated membranes containing nonblocked azo-sphingolipids upon light trigger analyzed by high-speed AFM. (A and B) Changes in the area of L_o domains (of depicted snapshots) before/after illumination with UV-A ($\lambda = 365 \text{ nm}$) and blue ($\lambda = 470 \text{ nm}$) lights on DOPC:Chol:SM:photolipid (10:6.7:5:5 mol ratio) SLBs having (A) Azo- β -Gal-Cer (see Video S1), Azo-SM (see Video S2), or Azo-Cer (see Video S3) with a sphingosine backbone ($X = -\text{H}$) and varying headgroup functionality ($R = \text{galactosyl}$, phosphocholine, or $-\text{OH}$, respectively) or (B) Azo- α -Gal-PhCer (see Video S5) or Azo-PhCer (see Video S6) with a phytosphingosine backbone ($X = -\text{OH}$) and varying headgroup functionality ($R = \text{galactosyl}$ or $-\text{OH}$, respectively). (C and D) Reversible lateral remodeling of a phase-separated SLB containing Azo-SM (DOPC:Chol:SM:Azo-SM; 10:6.7:5:5 mol ratio) upon UV-A/blue light irradiation, as seen in Video S4. (C) AFM images of the SLB at the dark-, UV-, and blue light-adapted states, displaying the area occupied by the L_o phase and the L_d - L_o height mismatches. (D) Relative variation of total L_o area of the SLB over time, shown in Video S4, upon shining short pulses (marked with arrows) of UV-A and blue light. To see this figure in color, go online.

when compared with sphingosine-based lipids having only one hydroxyl (3-OH) on their backbone.

In the dark-adapted state, phase-separated DOPC:Chol:SM:photolipid SLBs with Azo-PhCer and Azo- α -Gal-PhCer exhibited a domain height mismatch of 1.2–1.8 nm (Figs. 2 B and S5 C), very close to the L_d - L_o height differences here reported for membranes with sphingosine analogs (Figs. 2 A and S5 B). Upon photoactivation, the L_d - L_o phase-separated SLBs containing either Azo-PhCer (with a hydroxyl headgroup) or Azo- α -Gal-PhCer (with a bulkier galactosyl headgroup) behaved in a similar way to membranes with sphingosine-based azo-sphingolipids by exhibiting, at the end, an identical phenotype of membrane remodeling (Fig. S6 C).

As seen in Figs. 2 B and S5 C and Videos S6, S7, S13, S14, and S15, after irradiation with UV-A light ($\lambda =$

365 nm), L_d lakes initially appeared inside existing L_o domains, and the total L_d phase membrane area lowered by $\sim 23\%$ (area $L_o(\text{UV})/\text{area } L_o(\text{dark}) = 0.78 \pm 0.18$, from grouped high-speed AFM area values in Fig. S6, extracted from Figs. 2 B and S5 C; $n = 5$), while the domain height mismatch did not change majorly. Then, after irradiation with blue light ($\lambda = 470 \text{ nm}$), L_o islands initially formed inside the L_d regions, and the total L_o phase membrane area subsequently increased to the initial equilibrium dark-adapted values (area $L_o(\text{blue})/\text{area } L_o(\text{dark}) = 1.03 \pm 0.20$, from grouped high-speed AFM area values in Fig. S6).

Our results confirm that the bulkiness of the neutral headgroup does not play a role in the membrane remodeling ability of our photoswitchable phytosphingolipids. Moreover, the increased backbone polarity of the phytosphingosine backbone does not seem to affect the way

azo-phytosphingolipids engage in interactions with their neighboring lipids when compared with azo-sphingolipids. Hereby, we conclude that the tested Azo-Cer, Azo- β -GalCer, Azo-SM, Azo- α -Gal-PhCer, and Azo-PhCer establish stable interactions with other sphingolipids (such as SM) and sterols (such as Chol) inside L_o domains when their azobenzene acyl chain is in the *trans*-isoform (imitating a “straight” saturated acyl chain) and with unsaturated phosphatidylcholines (e.g., DOPC) inside L_d regions when the azobenzene is in the *cis*-isoform (mimicking a “bent” unsaturated chain).

It is worth noting that the normalized extent of L_o domain area changes produced by *trans-cis* isomerization of the *N*-acyl azobenzene chain in these two classes of photolipids (Fig. S6 C) appears to be robust and independent of domain size, scanned area dimensions, and different L_o phase surfaces scanned locally (Figs. 2, 3, and S5). A direct comparison of all AFM time lapses (Videos S1, S2, S3, S4, S5, S6, S7, S8, S9, S10, S11, S12, S13, S14, S15, S16, and S17) also shows that it is difficult to assign whether the contribution of Ostwald ripening or domain fusion to the restructuring of L_d lakes is different for each photolipid type. This was particularly evident in membranes containing Azo- α -Gal-PhCer and Azo-Cer (with completely different backbones and headgroups), where some of the bilayers appeared to show a more rapid remodeling and domain fusion-based mecha-

nism of the L_d lakes upon UV irradiation, while in other bilayers/areas, the remodeling/expulsion of the L_d lakes trapped inside L_o domains appeared to occur slowly by Ostwald ripening (for Azo-Cer, see Videos S3 and S12; for Azo- α -Gal-PhCer, see Videos S5, S13, and S14).

Remodeling of membrane domains by 3-OH-blocked azo-sphingolipids

In order to infer the exact role of H-bonding and sphingoid base polarity for the mode of action of azo-sphingolipids, we used high-speed AFM to further investigate the photo-switching and lateral membrane remodeling activities of the azo-sphingolipids Azo-THP-Cer and Azo-THP-SM. These lipids have the 3-OH group on their sphingosine backbone protected with a bulky THP moiety. Notably, the final protecting step resulted in an inseparable mixture of diastereomers at the THP linkage (see Fig. 3), which was used in all experiments.

In the dark-adapted *trans*-form, DOPC:Chol:SM:photolipid bilayers containing 3-OH-blocked Azo-THP-SM or Azo-THP-Cer (Figs. 3 and S5 D; Videos S7, S8, S9, S16, and S17) had L_o domains with irregular borders and lower heights (~ 0.6 – 0.9 nm) when compared with SLBs with non-blocked counterparts (Fig. 2). Such a noticeable effect on the global architecture of L_o domains is evidence that

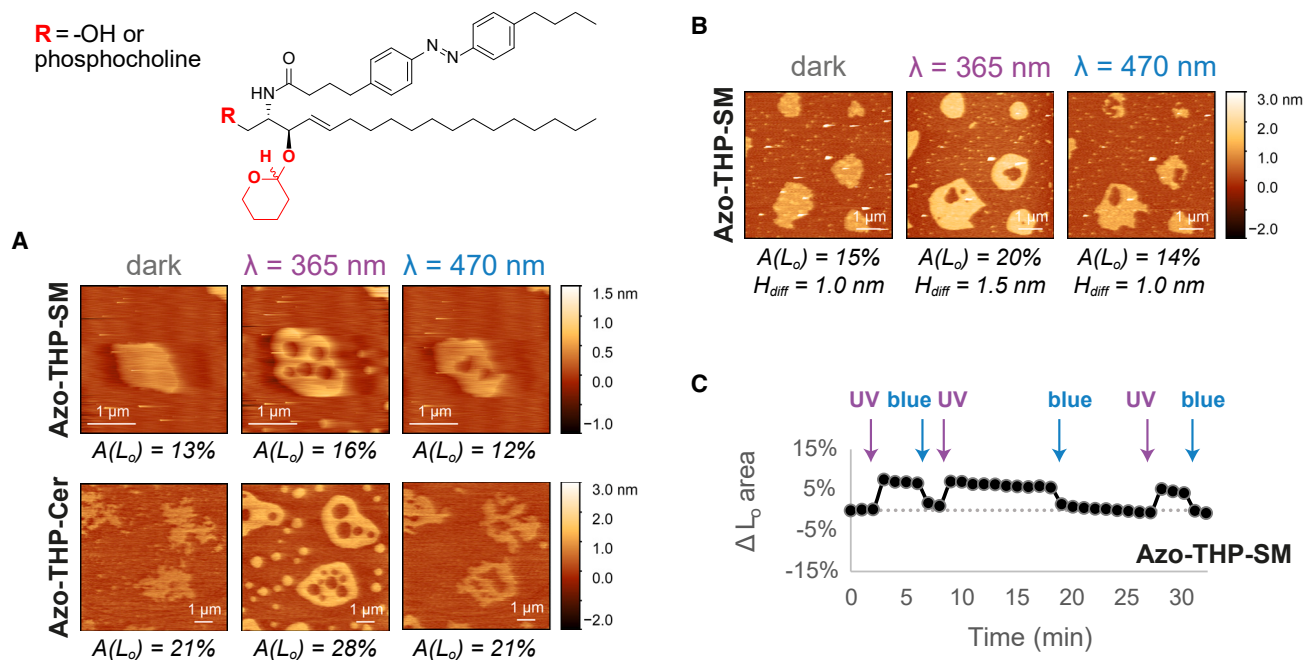


FIGURE 3 Lateral remodeling of phase-separated membranes containing 3-OH-blocked azo-sphingolipids upon light trigger analyzed by high-speed AFM. (A) Changes in the area of L_o domains (of depicted snapshots) before/after illumination with UV-A ($\lambda = 365$ nm) and blue ($\lambda = 470$ nm) lights on DOPC:Chol:SM:photolipid (10:6.7:5:5 mol ratio) SLBs having Azo-THP-SM (see Video S7) or Azo-THP-Cer (see Video S8) with a 3-OH-blocked (THP-protected) sphingosine backbone and varying headgroup functionality (R = phosphocholine or -OH, respectively). (B and C) Reversible lateral remodeling of a phase-separated SLB with Azo-THP-SM (DOPC:Chol:SM:Azo-THP-SM; 10:6.7:5:5 mol ratio) upon UV-A/blue light irradiation, as seen in Video S9. (B) AFM images of the SLB at the dark-, UV-, and blue light-adapted states, displaying the area occupied by the L_o phase and the L_d - L_o height mismatches. (C) Relative variation of total L_o area of the SLB over time, shown in Video S9, upon shining short pulses (marked with arrows) of UV-A and blue light. To see this figure in color, go online.

3-OH-blocked lipids are able to reduce the molecular packing within the L_o phase, as previously reported for 3-OH-blocked stearyl-SM (29,62).

When the membranes with THP-protected photolipids were irradiated with UV-A light ($\lambda = 365$ nm) and the lipids converted to the *cis*-isoform, the total area of L_o phase markedly increased $\sim 23\%$ (area $L_o(\text{UV})/\text{area } L_o(\text{dark}) = 1.23 \pm 0.21$, from grouped high-speed AFM area values in Fig. S6, extracted from Figs. 3 and S5 D; $n = 5$), with L_o domains getting larger, rounder, and noticeably higher (~ 1.1 – 1.4 nm) (Fig. 3, A and B). Quite strikingly, directly after exposure to UV-A light, L_d lakes inside preexisting L_o domains, as well L_o islands within the L_d matrix were transiently formed. The preexisting L_o domains then grew in total area, mainly via Ostwald ripening as L_o islands disappeared, whereas only a few larger L_d lakes appeared at the end trapped inside the enlarged L_o domains.

Subsequent illumination with blue light ($\lambda = 470$ nm) led to an overall decrease of both total L_d - L_o height mismatch and L_o area back to the initial dark-adapted state values (area $L_o(\text{blue})/\text{area } L_o(\text{dark}) = 0.98 \pm 0.20$, Fig. S6), as the 3-OH-blocked photolipids isomerized back to the *trans*-isoform. Interestingly, no noticeable formation of large L_d lakes or L_o islands was observed here. The L_o domains rapidly shrank with their domain borders becoming irregular (less rounded) and more unstable, as large rapid fluctuations were visible (Videos S7, S8, S9, S16, and S17). This clearly indicates that the THP-protected azo-sphingolipids, when in the *trans*-isoform, severely affect line tension of the phase-separated domains. Finally, the reported L_o domain height and area changes within the phase-separated bilayers could be repeated over multiple illumination cycles, as seen in Fig. 3 C and Videos S7, S8, S9, S16, and S17.

To sum up, photoswitchable sphingolipids having their 3-OH sphingoid moiety blocked with a THP group promote a clearly distinct light-induced reorganization of L_d - L_o phase-separated membranes when compared with nonblocked counterparts. These blocked lipids are able to significantly increase the percentual amount and height of the L_o phase upon UV-triggered isomerization to the *cis*-isoform and to decrease both these parameters upon blue light-triggered isomerization to the *trans*-isoform. Moreover, the formation of L_d lakes and L_o islands only after the application of UV light (and not blue light) suggests an altered distribution profile of these photolipids within the L_d and L_o phases when compared with nonblocked photolipids.

Phase-separation area changes by photoswitchable sphingolipids

After demonstrating that the various nonblocked versus hydroxyl-blocked photoswitchable sphingolipids reorganize phase-separated membranes differently, we set out to quantitatively compare the extent by which these lipids alter the

total distribution of phase separation, as well as other structural membrane parameters.

To begin, since AFM only allows us to follow a limited number of L_o domains simultaneously, we acquired additional large field-of-view fluorescence confocal images (Figs. S8, S9 A, and 4 A) to obtain better statistics for determining ensemble area values independent of domain size and number. We analyzed changes in L_o total area before/after UV/blue irradiation on DOPC:Chol:SM control SLBs ($n = 5$ images) and, most importantly, DOPC:Chol:SM:photolipid SLBs containing 18.7 mol % Azo-Cer, Azo- β -GalCer, Azo-SM, Azo-PhCer, Azo- α -GalPhCer, Azo-THP-Cer, or Azo-THP-SM (number of images per photolipid, $n = 5$ – 8) doped with 0.1 mol % Atto655-DOPE for fluorescent detection of the L_d phase.

Usage of fluorescence allowed us to easily generate binary masks (Fig. S8), from which L_o phase areas could be straightforwardly estimated. Instead of collecting the confocal images immediately after photoisomerization (Fig. S3), which would have been smeared due to the short-lived nanoscopic L_d lakes, we let the SLBs equilibrate here for 20 min after brief irradiation with UV/blue light. In this way, we were able to reduce the number of hard-to-resolve L_d lakes and L_o islands and obtained a clearer macroscopic L_d - L_o phase separation (Figs. 4 A and S9 A).

All average absolute L_o area values recovered for the various SLBs are shown in Fig. S9 B. These values are particularly important as a way to assess the extent of membrane phase-separation perturbation caused by the presence of *trans*-photolipids prior to photoactivation. Briefly, control membranes without photolipids (DOPC:Chol:SM 10:6.7:10 mol ratio) displayed a total surface coverage by L_o domains of 45%. A similar average value of $45\% \pm 4\%$ was obtained for membranes with 18.7 mol % nonblocked photolipids in *trans*. Only the SLBs with THP-protected *trans*-photolipids showed significantly lower total L_o domain coverage, about $29\% \pm 3\%$ ($p < 0.001$), when compared with the control membranes without photolipids. This large reduction in total L_o area for membranes with hydroxyl-protected azo-sphingolipids clearly supports our AFM observation that THP-blocked *trans*-photolipids affect L_o phase properties.

To facilitate data comparison for the photoswitchable conversion of phase separation by our photolipids, the recovered L_o areas were normalized before/after UV/blue light illumination by the average L_o area for the different individual SLBs at the dark-adapted state (Fig. 4 C). This allowed us to analyze the relative changes in the total L_o area (for *cis*- and *trans*-photolipids) and compare it with the behavior of the DOPC:Chol:SM control lipid samples without photolipids (Fig. 4 C). Relevant statistical comparisons were then performed using one-way analysis of variance followed by Bonferroni comparison *t*-tests (95% confidence interval), and the respective *t* statistics and *p* values are listed as supporting material in Table S3.

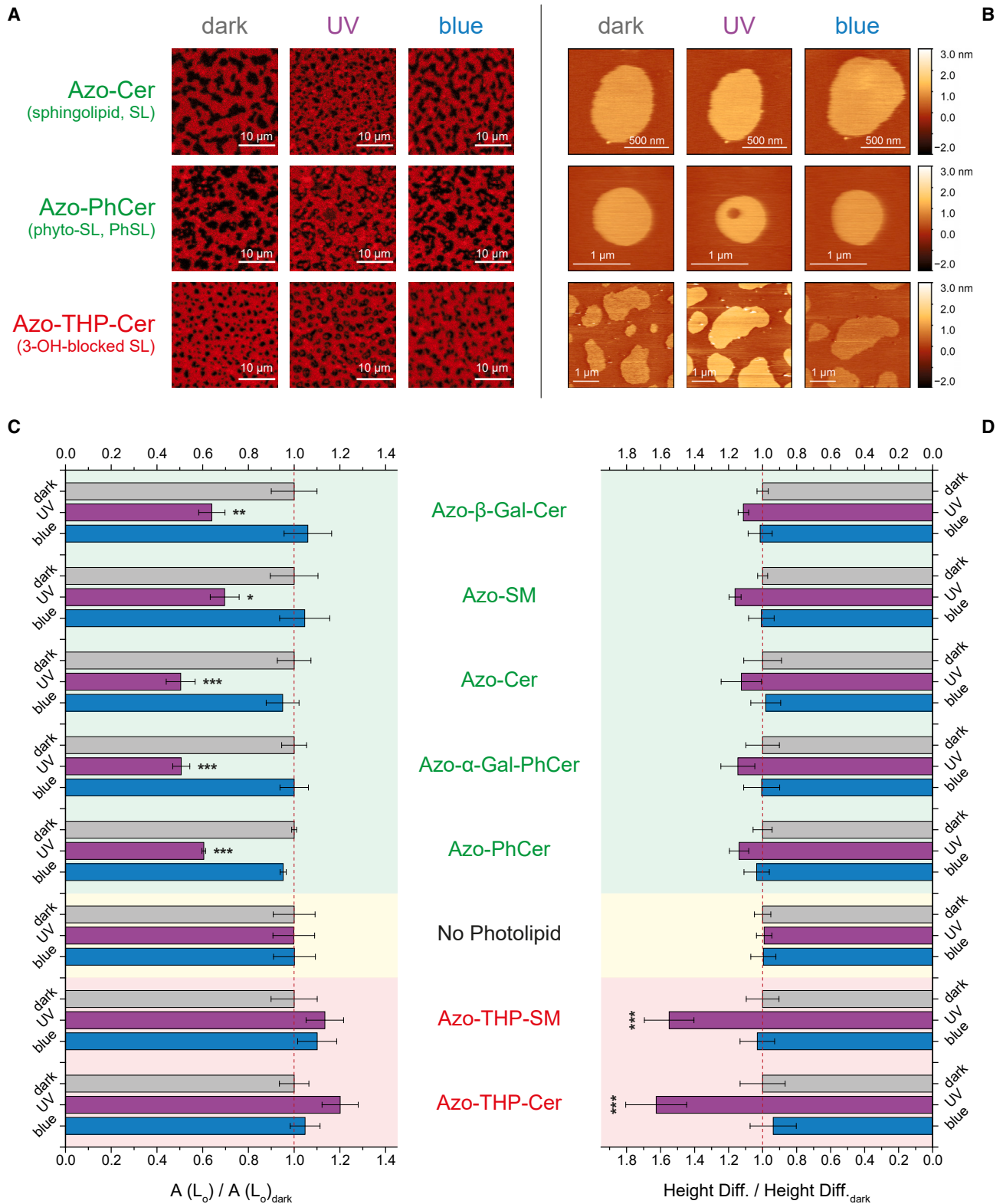


FIGURE 4 Normalized changes in the total L_o phase area and L_d-L_o height difference on phase-separated SLBs having different types of azo-sphingolipids upon application of UV-A ($\lambda = 365$ nm) and blue ($\lambda = 470$ nm) lights. Fluorescence confocal (A) and AFM (B) images of DOPC:Chol:SM:photolipid (10:6.7:5:5 mol ratio) SLBs with sphingosine-based Azo-Cer, phytosphingosine-based Azo-PhCer, or 3-OH-blocked sphingosine-based Azo-THP-Cer, all having the same $-OH$ headgroup but distinct sphingoid backbone. Normalized L_o areas (C) and L_d-L_o height differences (D), respectively, recovered

(legend continued on next page)

As seen in Figs. 4 C and S9 B, the total L_o area for phase-separated SLBs having sphingosine- or phytosphingosine-based photolipids significantly decreased (with p values between $p < 0.05$ and $p < 0.001$) on average by 41% (area $L_o(\text{UV})/\text{area } L_o(\text{dark}) = 0.59 \pm 0.04$) after UV-A illumination (when the lipids are in *cis*) and augmented back to the original dark-adapted state (area $L_o(\text{blue})/\text{area } L_o(\text{dark}) = 1.00 \pm 0.02$) upon blue light irradiation (when the lipids are in *trans*). For SLBs having Azo-THP-SM or Azo-THP-Cer, on the contrary, the total area of the L_o phase did not statistically change ($p > 0.05$) when compared with control membranes without a photolipid. Here, only a slight increase by 17% (area $L_o(\text{UV})/\text{area } L_o(\text{dark}) = 1.17 \pm 0.03$) after illumination with UV-A light is seen, while the recorded amount of L_o phase decreased back to the original dark-adapted state value (area $L_o(\text{blue})/\text{area } L_o(\text{dark}) = 1.07 \pm 0.03$) after the application of a blue light pulse.

Interestingly, if we compare the changes in L_o area after azo-sphingolipid photoisomerization determined by fluorescence (Figs. 4 C and S9) versus high-speed AFM data (Fig. S6), the area changes for confocal microscopy seem to be skewed toward detecting higher amounts of L_d phase after UV irradiation. This skew may be a direct consequence of the limited pixel resolution of conventional laser-scanning confocal microscopy for detecting nanoscale L_o domains when compared with AFM. Despite this instrumental bias, similar trends in membrane domain area variations were detected with both fluorescence confocal and AFM techniques.

These experiments corroborate that sphingosine- and phytosphingosine-based photolipids rely on the same principles for reshuffling membrane phase-separated domains, whereas the THP-protected counterparts, owing to their distinct physicochemical properties, follow a markedly different mechanism. Finally, although there is no statistical difference between the various sphingosine- and phytosphingosine-based *cis*-photolipids ($p = 1$), the obtained t statistics for their comparison with the control DOPC:Chol:SM suggest a possible hierarchy. Here, Azo-Cer and the two azo-phytosphingolipids (Azo- α -Gal-PhCer and Azo-PhCer), the latter being able to establish stronger H-bonding interactions, appeared to possess the strongest “remodeling activity” (with $p < 0.001$) (see Table S3 for more details).

Domain height mismatch changes by photoswitchable sphingolipids

Our results so far clearly point out that blocking the interfacial hydroxyl on the sphingoid backbone has a marked effect

on the molecular organization of individual lipids and on the global architecture of L_o domains. Thus, to quantitatively ascertain how photoswitchable sphingolipids affect the structure and physicochemical properties of L_o domains within phase-separated membranes, we collected zoomed-in and high-resolution low-speed AFM images ($n = 5-9$) of individual L_o domains on DOPC:Chol:SM:photolipid SLBs prior to and 20 min after brief illumination with UV-A/blue light (to let domains equilibrate and reduce the amount of L_d lakes and L_o islands), as depicted in Figs. 4 B, S10, and S11 A. This acquisition mode allows us to follow the membrane contour with an increased signal/noise ratio and therefore determine more accurately the height differences between the L_o domains and the surrounding L_d matrix (Fig. S10). This is an important parameter, as it relates to the hydrophobic mismatch between the saturated (“nonbent” acyl chains) lipids in L_o domains with increased chain order (63) and the unsaturated (bent acyl chains) lipids in less packed L_d regions.

Altogether, the average height difference between L_o and L_d regions (at the dark-adapted state for the various SLBs having either sphingosine- or phytosphingosine-based photolipids) was 1.3 ± 0.1 nm. This value corresponds to the mean of all average L_d - L_o height differences (\pm standard error) obtained for membranes containing Azo-Cer, Azo- β -GalCer, Azo-SM, Azo-PhCer, or Azo- α -GalPhCer (Fig. S11 B) and was very close to the less precise values previously reported using high-speed AFM. Owing to the exquisite z -resolution of AFM, we also identified that the L_o domains of SLBs having azo-sphingolipids with smaller headgroups (e.g., Azo-Cer and Azo-PhCer) were slightly less elevated (1 ± 0.1 nm) than the L_o domains of SLBs having azo-sphingolipids with larger headgroups (e.g., Azo-SM, Azo- β -GalCer, and Azo- α -GalPhCer). The later-displayed L_d - L_o height mismatches (1.4 ± 0.1 nm) are closer to the values recovered (1.8 ± 0.1 nm) for control ternary mixtures without a photolipid. Hence, our observations corroborate a preferred localization of the *trans*-azo-sphingolipids inside L_o domains, as these lipids could then engage hydrophobic packing and stable H-bonding interactions with SM and Chol, altering slightly the height of L_o domains due to the different headgroup size and N -acyl chain length (e.g., C18:0 acyl chain: 21.2 Å vs. FAAzo-4: 17.9 Å, retrieved from Chem3D, PerkinElmer, Boston, MA, USA).

Interestingly, upon applying UV-A light to SLBs having these nonblocked photolipids, the L_d - L_o height mismatch increased on average by 14% (1.4 ± 0.1 nm; Fig. 4 D). Although this variation is not statistically relevant, it is in line with the exclusion of *cis*-azo-sphingolipids from the

from fluorescence confocal and AFM data for phase-separated SLBs having either azo-(phyto)sphingolipids with free 3-OH (marked in green), no azo-sphingolipid (controls with SM, marked in yellow), or THP-protected azo-sphingolipids with the 3-OH blocked (marked in red). Error bars correspond to standard error of the mean ($n = 5-8$ confocal images each). Statistical analysis: *cis*-photolipids versus no photolipid (*** $p < 0.001$, ** $p < 0.005$, * $p < 0.05$). See Figs. S9 and S11 for details. To see this figure in color, go online.

L_o phase and SM being then the predominant sphingolipid molecule inside those domains. Subsequent irradiation with blue light led to the decrease of the domain height by 13% back to the original values reported for the dark-adapted state (Fig. 4 D), corroborating a repartitioning of *trans*-azo-sphingolipids back to the L_o phase.

For membranes having hydroxyl-blocked photolipids, the height of L_o domains at the dark-adapted state was lower, and the domain boundaries more irregular, when compared with membranes with nonblocked counterparts. Indeed, the L_d - L_o height mismatch observed for SLBs with Azo-THP-SM or Azo-THP-Cer was below 1 nm (0.7 ± 0.1 nm; Fig. S11 B), similar to the values observed using high-speed AFM and nearly 0.5 and 1.0 nm lower than the height mismatch found for SLBs with Azo-SM or control membranes lacking azo-sphingolipids, respectively. The lower L_o height observed for membranes with 3-OTHP-lipids indeed corroborates that these lipids establish altered interactions with other colipids in the L_o phase when in the *trans*-isoform but, most importantly, validates that blocking H-bonding severely alters the molecular packing, as well as the line tension within L_o domains.

This destabilization effect can be overcome once the hydroxyl-blocked azo-sphingolipid is converted to its *cis*-isoform upon illumination with UV-A light. Indeed, after applying UV-A to the phase-separated SLBs with Azo-THP-SM, the L_d - L_o height mismatch significantly increased ($p < 0.001$) by 55% (Fig. 4 D) to average values above 1 nm (1.1 ± 0.1 nm). This elevation in height suggests that Azo-THP-SM and Azo-THP-Cer lipids are expelled from the L_o phase when in the *cis*-isoform, leaving the L_o domains mainly composed by SM and Chol. Without the interference of these THP-protected lipids, SM and Chol molecules can then establish more stable H-bonding and tighter hydrophobic chain packing interactions, giving rise to taller, rounder, and larger L_o domains. In opposition, irradiation of the phase-separated SLBs with blue light leads to a marked reduction of the L_o domain height (Fig. 4 D) back to the initial dark-adapted state value (0.7 ± 0.1 nm). As the 3-OH-blocked azo-sphingolipids isomerize back to their *trans*-isoform, these lipids could then reestablish hydrophobic chain packing interactions with the other L_o -localizing lipids, destabilizing the existing H-bonding interactions between SM and Chol.

Domain line tension changes by photoswitchable sphingolipids

A parameter closely linked to the domain height mismatch is line tension, which can be perceived as the interfacial energy arising at the boundaries of coexisting phases and is an important driving force for membrane shape transformation (e.g., budding (4,64,65) and fusion (66)). In order to estimate the approximate values of line tension for the various L_d - L_o membranes with distinct blocked and nonblocked photoswitchable sphingolipids, based on the height

mismatch values measured using low-speed AFM, we used the theoretical model for flexible domains implemented by Cohen and co-workers (Eq. 1). This model describes a quadratic dependence of the line tension with the phase height mismatch (53) and was previously used to estimate line tension on phase-separated membranes with similar lipid compositions (52).

Overall, as seen in Fig. S11 C, line tension values ranged from 1.9 to 4.3 pN for L_d - L_o phase-separated SLBs with sphingosine- and phytosphingosine-based lipids in the *trans*-isoform. When the lipids are in the *cis*-isoform and partition to the L_d phase instead, a statistically nonsignificant yet small increase in line tension by 24% (Fig. S12) was recovered, with the values being very close to the ones gauged for phase-separated SLBs lacking azo-sphingolipids (5.4 ± 0.6 pN).

In contrast, L_d - L_o bilayers with THP-protected azo-sphingolipids (such as Azo-THP-SM and Azo-THP-Cer) in the *trans*-isoform possess noticeably reduced line tension values (~ 1.2 pN): 2.2- to 4.5-fold lower than the domain line tension measured for SLBs with nonblocked photolipid counterparts (Fig. S11 C). Next, when the hydroxyl-blocked lipids are in their *cis*-isoform and locate in the L_d phase, the domain line tension greatly increases ($p < 0.001$) by $\sim 120\%$ (Fig. S12), reaching values close to the ones reported for nonblocked counterparts (2.6 pN). Hence, THP-protected photolipids, when in *trans*, greatly reduce the line tension of L_o domains in opposition to the azo-sphingolipids with free interfacial hydroxyls, appearing to possess additional line-active (lineactant) properties.

Line-active molecules are known to concentrate at the boundaries of membrane phases (67–70), reducing the hydrophobic mismatch and line tension around phase-separated domains (e.g., L_o versus L_d). Herein, hybrid lipids such as palmitoyl-oleyl-phosphatidylcholine, possessing both a saturated and unsaturated fatty acid chain, are of particular relevance. When added to ternary mixtures made of lipids with two saturated tails (e.g., DPPC or DSPC), two unsaturated tails (DOPC), and a sterol (Chol), palmitoyl-oleyl-phosphatidylcholine was shown to interact with the L_o phase and disturb the chain ordering of L_o domains in a so-called “partitioning and loosening” mechanism (71). Such an effect will then promote the reduction of line tension and formation of nanoscopic domains, hence lowering the differences in physical properties between the L_o and L_d phases (71–75). Interestingly, Azo-THP-SM and Azo-THP-Cer also appear to reduce line tension and promote the formation of nanoscopic domains in a similar way, sharing moreover key structural similarities with hybrid lipids: 1) the *trans*-azobenzene *N*-acyl chains mimic saturated fatty acids prone to localize within L_o domains, and 2) the bulky THP moiety at the sphingosine base appears to interfere with the molecular packing of lipids and therefore be susceptible to preferentially localize in the less packed L_d phase.

To sum up, the hybrid chain properties in addition to the blockage of H-bonding could be possible explanations for the observed perturbation of the L_o domain boundaries and reduction of domain height by THP-protected azo-sphingolipids in the *trans*-isoform. It is plausible that the blocked photolipids in the *trans*-configuration are not exclusively localized in the L_o phase but rather behave as a hybrid lipid able to interact with both the L_d and L_o phases.

Here, the blocked *trans*-photolipids partitioned inside the L_o phase could loosen the chain packing of the L_o domains, reduce line tension, and be responsible for the formation of L_d lakes within the L_o phase when converted into *cis*. Moreover, since the nominal height mismatch (Fig. S11 B) and total area (Fig. S9 B) of the L_o phase are significantly reduced ($p < 0.001$) on membranes with hydroxyl-blocked photolipids when compared with membranes with nonblocked photolipid equivalents (i.e., *trans*-Azo-SM and *trans*-Azo-Cer), we can also speculate that a small fraction of the destabilized “rigid” colipids (i.e., C18-SM and Chol) might be localized in the L_d phase as well. This would further substantiate the mechanism of L_o island formation after UV irradiation, as such a pool of rigid colipids in the L_d phase could serve as nuclei for the formation of L_o islands once the fraction of L_d -localizing THP photolipids is isomerized into *cis* and no longer destabilizes this pool of rigid colipids.

Changes in the mechanics of homogeneous membranes promoted by the photoisomerization of azo-sphingolipids

Despite the differences in membrane domain remodeling by THP-protected versus nonprotected azo-sphingolipids, we also observed the generation of occasional membrane holes when isomerizing 3-OH-blocked photolipids from the bent *cis*-isoform to the straight *trans*-isoform, as depicted in Fig. S13 for phase-separated membranes with Azo-THP-SM. This hole formation clearly indicates that SLBs containing THP-protected azo-sphingolipids, similarly to membranes with nonblocked counterparts (Fig. S7 A), globally expand after UV-A illumination and subsequently compress upon illumination with blue light due to the bending and unbending on the *N*-acyl chains, respectively. Hence, photoswitching of azo-sphingolipids not only influences lipid phase separation (as discussed so far) but irrefutably affects basic mechanical properties of the membrane, such as packing and stiffness/fluidity.

To evaluate whether photoswitchable sphingolipids indeed interfere with global packing of membranes, irrespective of phase separation, we sought to perform additional analyses on a nonphase-separated system. In this context, we explored the possibility of using ternary compositions that form only a single L_o phase. It has been reported in the literature that certain sphingolipid-containing ternary mixtures such as DOPC:GalCer:Chol (37.5:37.5:25 mol %)

can exhibit a single L_o phase in the presence of significant amounts of Chol (76), whereas other mixtures with the same ratio, e.g., DOPC:SM:Chol (37.5:37.5:25 mol %), can exhibit L_d - L_o phase separation (77). Some information is already available on ternary mixtures containing photolipids from previous work with colleagues, notably for the phosphatidylcholine Azo-PC. Specifically, for equimolar DPhPC:Azo-PC membrane mixtures, Urban et al. reported L_d - L_o phase separation in the presence of 10–35 mol % Chol and a single L_o phase with Chol above 35% (45). In addition, the authors also reported a single phase on giant vesicles for DOPC:Azo-PC with 20 mol % Chol (78).

With this in mind, we examined bilayers containing either Azo-Cer (as nonblocked azo-sphingolipid) or Azo-THP-SM (as 3-OH-blocked azo-sphingolipids) in a DOPC:Chol:photolipid composition with 10:6.7:10 mol ratio. As shown in Fig. S14 (using fluorescence confocal microscopy) and Fig. S15 (using AFM), these membrane compositions are indeed homogenous and do not exhibit domains either before or after irradiation with UV-A/blue light.

Next, we performed additional AFM-based force spectroscopy measurements, in particular nanoindentation experiments (54,55,79–84), on these ternary mixed homogenous SLBs having Azo-Cer (Fig. S16 A) or Azo-THP-SM (Fig. S16 D). In short, when indenting such membranes with an AFM tip, a typical jump (or discontinuity) corresponding to the force required to pierce (or break through) the SLB can be easily identified within the collected force-displacement curves (as seen in Fig. S16, B and E). The extent of such breakthrough forces is then directly linked to the mechanical properties of the membrane: lower forces are expected when the membranes are more fluid (or less compact) and higher forces when these are stiffer (or more compact). Thus, upon recording a set of force curves prior and after illumination with UV-A ($\lambda = 365$ nm) and blue ($\lambda = 470$ nm) lights, we evaluated the breakthrough events and displayed the recovered forces needed to pierce the membranes (Fig. S16, C and F) as histograms normalized by the average breakthrough force obtained for the dark-adapted state. Nonnormalized values are depicted in Fig. S17.

For homogenous DOPC:Chol:Azo-Cer SLBs, an average breakthrough force of 2.8 ± 0.3 nN ($n = 900$ curves) was recorded in the dark-adapted state (Fig. S17 A). After illumination with UV light, and consequent conversion of Azo-Cer to its bent *cis*-isoform, the force required for piercing the membrane reduced by $\sim 30\%$ (Fig. S16 C) to 1.9 ± 0.2 nN (Fig. S17 B). Irradiation with blue light, on the contrary, promoted an increase of the breakthrough force back to its original average value (2.8 ± 0.3 nN; Fig. S17 C), as Azo-Cer would back isomerize to its *trans*-isoform. Thus, for nonblocked azo-sphingolipids, we confirmed that the *cis*-isoform expands/fluidifies the membrane, while the *trans*-isoform compacts/stiffens it. In this context, our force data clearly back up a previous study (44) based on

fluorescence recovery after photobleaching, which showed that lateral lipid diffusion on a membrane made of Azo-PC (phosphatidylcholine analog with a FAAzo-4 acyl chain) was higher when the photolipid was in its *cis*-isoform and slower when in the *trans*-isoform.

Interestingly, the measured average breakthrough force required to pierce a bilayer with Azo-THP-SM was significantly higher than for bilayers containing Azo-Cer (Fig. S16, B and E). Although we cannot exclude a contribution of the THP moiety at the interface, the measured increase is mainly due to the larger phosphocholine headgroup of Azo-THP-SM compared with the smaller hydroxyl headgroup of Azo-Cer. Indeed, Garcia-Manyes et al. (81) had previously found a clear correlation of membrane piercing forces for lipids with the same acyl chains as a function of headgroup size. In that seminal work, the authors found that DPPA with the smallest headgroup had the lowest breakthrough force, followed by DPPE, DPPC, and DPPS.

When we instead analyze the normalized changes in breakthrough force for nonphase-separated DOPC:Chol:Azo-THP-SM bilayers as a function of photoisomerization, then similar trends as for Azo-Cer were recorded. More precisely, the piercing force needed to break through those membranes ($n = 200$ curves) also reduced by $\sim 25\%$ (Fig. S16 F), from 5 ± 0.4 (Fig. S17 D) to 3.8 ± 0.4 nN (Fig. S17 E), upon irradiation with UV light and formation of *cis*-Azo-THP-SM, in agreement with a global expansion or fluidification of the membrane. Subsequently, the membrane breakthrough force also reverted back close to the original value (4.7 ± 0.6 nN; Fig. S17 F) upon illumination with blue light and formation of *trans*-Azo-THP-SM, in agreement with a global compaction or rigidification of those membranes.

Therefore, based on this force spectroscopy outcome for homogenous membranes, we can argue that the opposite changes in L_o area for phase-separated SLBs containing either 3-OH-blocked or nonblocked azo-sphingolipids (observed throughout the previous manuscript sections) are mainly due to different types of interactions these photolipids engage in via their sphingoid backbone with neighboring lipids within L_o domains and are not directly linked to the structural properties of the *N*-acyl photoswitch per se.

Comparison of the behavior of photolipids in direct relation to their natural sphingolipid counterparts

In this article, we have shown that photoswitchable sphingolipids with an azobenzene *N*-acyl chain are able to change the L_d - L_o ratio on phase-separated membranes and to affect packing on homogeneous membranes in a photoisomerization-dependent manner. However, a fundamental question remains as to how similar and biocompatible such photolipids are with respect to their natural counterparts.

Photolipids appear to maintain phase separation in the presence of native counterparts, as recently shown by Socrer et al. for photo-Gb3 (85) and previously by us for Azo-Cer (39). For example, in our previous work (39), we have shown that Azo-Cer not only maintains L_d - L_o phase separation but is also able to sustain a liquid-gel phase when mixed with DOPC and SM in the absence of Chol. From previous results with Azo-PC (45), it also appears that photolipids are, in principle, capable of forming phase separation themselves. However, the exact ratios required for these processes may differ from those of the native 18:0 or 16:0 counterparts and depend on the nature of the colipids used.

Concerning native sphingolipids, several studies in the literature suggest that small differences in headgroup, acyl chain, or sphingoid base are sufficient to significantly alter membrane properties and lateral organization. For example, it is well known that Cers have a strong ability to induce gel-fluid phase separation on membranes, as reported elsewhere (86,87). Particularly important is the presence of significant amounts of Chol, which is known to liquefy such rigid gel phases. As previously mentioned in changes in the mechanics of homogeneous membranes promoted by the photoisomerization of azo-sphingolipids, no gel phases were found for equimolar DOPC:SM (77) or equimolar DOPC:GalCer (76) mixtures with 25 mol % Chol. For mixtures containing Cer, Castro et al. (88) also reported that increasing amounts of Chol liquefied the gel phases.

Since the presence or absence of certain phases depends then on the ratiometric combination and the nature of the sphingolipids and colipids (i.e., unsaturated phospholipid and Chol), we still decided to investigate the phase-separation properties of natural C18 lipids in membranes under the same ratiometric combinations of colipids that we tested here for the photolipids. Therefore, we collected AFM and fluorescence images of quaternary DOPC:Chol:SM:C18-sphingolipid mixtures and finally compared them with those of quaternary DOPC:Chol:SM:photolipid (all bilayers with a molar ratio of 10:6.7:5:5 and 0.1 mol % Atto6550DOPE). Such a comparison will ultimately give us a clearer picture on how biocompatible our photolipids are in relation to their natural counterparts.

As summarized in Fig. S18, most membranes with photolipids ended up showing equivalent phase-behavior properties when compared with membranes having their natural sphingolipid counterparts. Indeed, DOPC:Chol:SM membranes with 18.7 mol % azo- β -GalCer and azo-PhCer exhibited similar total L_o area (in *trans*) as well as similar L_d - L_o height deviation as membranes with 18.7 mol % 18:0- β -GalCer and 18:0-PhCer (native counterparts). Membranes with 18.7 mol % Azo-SM showed a similar total L_o area (in *trans*) yet a slightly smaller (~ 0.4 nm) but significant ($p < 0.001$) deviation in the L_d - L_o height mismatch compared with membranes with 18:0-SM.

A clearly different behavior was observed only for membranes with 18.7 mol % Azo-Cer, which did not exhibit any gel phase compared with their native 18:0-Cer counterpart. While L_d - L_o phase separation was observed for bilayers with 18:0-SM, 18:0- β -GalCer, and 18:0-PhCer, a gel phase (visible by AFM and fluorescence microscopy) was observed for membranes with 18:0-Cer instead. AFM analysis also shows that lipid bilayers with 18:0-Cer form a third L_o phase with an intermediate height (Fig. S18 B), which is consistent with previous results (10). However, apart from these differences, no significant variations can be observed when comparing the values of the recorded L_d - L_o height for both SLBs with azo-Cer or 18:0-Cer, as both membranes show a similar L_d - L_o mismatch height of 1.1 nm (Table S2).

Based on these last results, photolipids do not seem to behave dramatically differently from their native counterparts, at least in terms of their ability to maintain L_d - L_o phase separation and preserve their respective height deviation. This is very important information confirming that azobenzene-modified lipids are indeed biomimetic, or even bioequivalent, molecules and can therefore be considered a suitable option for manipulating native membrane properties.

CONCLUSIONS

In this work, we evaluated physicochemical foundations for the membrane remodeling ability by a family of photo-switchable sphingolipids, deciphering the relative contributions of the lipid headgroup and sphingoid backbone. We synthesized new types of *N*-acyl azobenzene sphingolipids with varying headgroup and sphingoid base functionalities. Then, we studied, with the help of atomic force and fluorescence microscopies, the propensity of these photolipids to alter membrane properties and laterally remodel L_d - L_o phase-separated supported membranes. Overall, we demonstrated that the headgroup type (simple hydroxyl versus more complex galactosyl or phosphocholine) does not interfere with the photoswitching ability of the various azo-sphingolipids within L_d - L_o lipid mixtures. Owing to the photodynamic reversibility of the azobenzene *N*-acyl chain, we further highlighted that *trans*-photolipids (i.e., dark-adapted and blue light-illuminated states) predominantly localize within preexisting L_o domains and compact membranes, while *cis*-photolipids (i.e., UV-A-illuminated state) preferentially locate within the more fluid L_d membrane regions and expand membranes.

Importantly, our results provide clear evidence that the nature of the sphingoid backbone, and its ability to engage stable H-bonding interactions with other colipids, plays a fundamental role in the way photoswitchable sphingolipids remodel L_d - L_o phase-separated membranes and change the amount, size, and height of L_o domains. Sphingosine- and phytosphingosine-based lipids, with their free interfacial 3-OH and 4-OH hydroxyls, do not significantly alter

the height of L_o domains when in the *trans*-isoform. In contrast, THP-protected lipids, with their interfacial 3-OH blocked, greatly interfere with the molecular packing and line tension of L_o domains, markedly reducing the overall L_o height mismatch. Whereas nonblocked azo-sphingolipids will promote a decrease of the total L_o phase area upon UV trigger, THP-protected azo-sphingolipids will increase the total L_o area, as well as induce a marked rise in L_o domain height after illumination with UV light.

Taken together, the structural diversity of the photo-switchable sphingolipids presented here, as well as an exquisite understanding of how these lipids alter important membrane properties, may offer new strategies for controlling the structure of biological lipid bilayers and the localization of membrane-interacting proteins. Thus, by further expanding the headgroup repertoire of photolipids, we may soon be in the position to target the fate of biologically relevant proteins on membranes using light as trigger. Such an endeavor would not only open up new, exciting avenues for optodynamic applications in the fields of synthetic biology, structural biology, or biophysics but would also offer novel perspectives toward the development of innovative photoresponsive drugs and pharmacological therapies.

SUPPORTING MATERIAL

Supporting material can be found online at <https://doi.org/10.1016/j.bpj.2023.02.029>.

AUTHOR CONTRIBUTIONS

H.G.F. and S.M.L. conducted the biophysical experiments and analyzed data. N.H. synthesized and purified most photolipids, in cooperation with N.W., H.T.-R., and J.A.F. P.S. and D.T. provided funding. D.T. supervised the synthesis of photolipids and H.G.F. their biophysical characterization. H.G.F., N.H., and S.M.L. prepared the first manuscript draft. All authors contributed to revision of the manuscript.

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DECLARATION OF INTERESTS

The authors declare no competing interests.

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