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**The co-assembly of lipids and other compounds has recently gained increasing interest. Here, we report the formation of stimuli-responsive lipid–DNA origami fibers through the electrostatic co-assembly of cationic lipids and 6-helix bundle (6HB) DNA origami. The photosensitive lipid degrades when exposed to UV-A light, which allows a photoinduced, controlled release of the 6HBs from the fibers. The presented complexation strategy may find uses in developing responsive nanomaterials e.g. for therapeutics.**

Lipid self-assembly is important for the formation of many cellular compartments in biological systems, but lipids can also co-assemble with other molecular entities into hierarchical structures with different structural morphologies.<sup>1</sup> Nucleic acids, for example, form highly ordered supramolecular assemblies (lipoplexes) when complexed with cationic lipids.<sup>2,3</sup> Lipoplexes have potential applications in nanomedicine<sup>3–5</sup> as well as in optoelectronics and synthetic chemistry.<sup>5</sup> We have recently demonstrated that lipids could also co-assemble with larger DNA origami nanostructures and that DNA origami may serve as nucleation templates for the growth of multilamellar lipid assemblies.<sup>6</sup> DNA origami are well-defined two- and three-dimensional (2D and 3D) DNA-based nanostructures prepared by folding a long-single-stranded DNA “scaffold” strand by a set of shorter oligonucleotides (“staples”).<sup>7–9</sup> Versatile and custom-shaped DNA origami can be utilized in, for example, drug delivery,<sup>10–12</sup> synthetic cells,<sup>13,14</sup> nanorobotics,<sup>15–17</sup> and bottom-up nanofabrication.<sup>18,19</sup> In addition, the phosphate groups in the DNA backbone give the DNA origami an overall negative charge, and therefore, DNA origami can direct formation of ordered

assemblies from positively charged compounds merely *via* electrostatic interactions.<sup>20–23</sup>

Currently, the research interests are increasingly focusing on functional nanomaterials that can respond to external stimuli.<sup>24</sup> Responsive nanomaterials that are able to release encapsulated compounds on demand would be advantageous especially for therapeutics.<sup>25</sup> In our previous work, we showed that DNA origami could be released from the multilamellar lipid assemblies by adding competitive polyanions.<sup>6</sup> However, for practical applications this disassembly approach is rather unfeasible and it would be advantageous to explore other stimuli-responsive disassembly methods. Light as a stimulus is non-invasive and straightforward to apply with high spatio-temporal control.<sup>26</sup> Therefore, we designed and synthesized a cationic photosensitive lipid, spermine-hydroxyethyl photolinker-oleic ester (**SpILO**, Fig. 1a), with the ability to co-assemble with DNA origami (Fig. 1b) into highly ordered lipid–DNA origami fibers through electrostatic and hydrophobic interactions (Fig. 1c). **SpILO** contains an *o*-nitrobenzyl group that undergoes photolytic degradation when exposed to UV-A light ( $\lambda \sim 350$  nm),<sup>27</sup> thus cleaving the oleic ester tail from the spermine head group (inset in Fig. 1a). The spermine group alone has low affinity to DNA, and by removing the lipid tail, the self-assembled multivalency of **SpILO** driven by hydrophobic interactions, is lost, and the **SpILO**-DNA origami fibers are disassembled.<sup>28–30</sup>

The photosensitive lipid **SpILO** was synthesized and characterized using standard methods (ESI,† Scheme S1 and Fig. S1, S2, S25–S35). Briefly, a photolabile *o*-nitrobenzyl linking group (**pll**) was functionalized with a *tert*-butyloxycarbonyl protected spermine (BOC-spermine, **1**) by an amide bond.<sup>27</sup> In a subsequent step, the BOC-spermine-**pll** (**2**) was coupled with the oleic acid in a Steglich esterification. Finally, the water-soluble **SpILO** was obtained by a deprotection of the BOC-protected spermine group by acidic hydrolysis. For the co-assembly with **SpILO**, we used a long, rod-shaped 6-helix bundle (6HB) DNA origami<sup>31</sup> as a model structure (Fig. 1b and ESI,† Fig. S3–S6). The negatively charged DNA origami interacts with the positively charged

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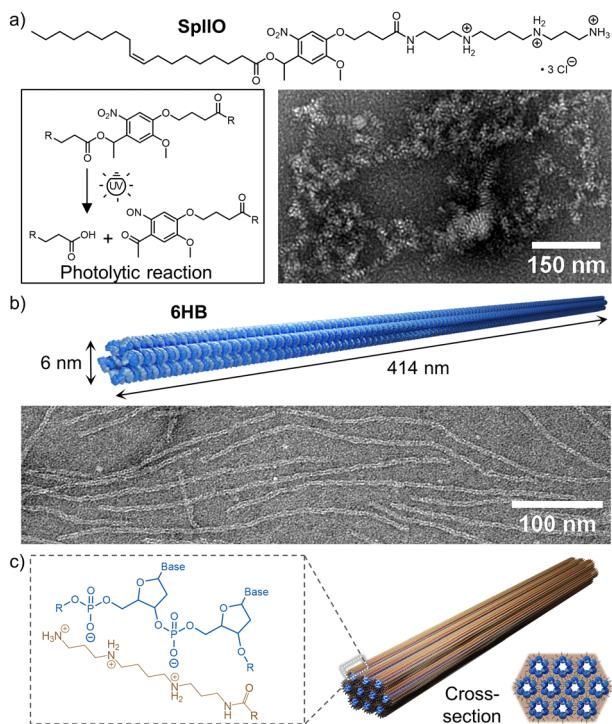
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**Fig. 1** The building blocks for the electrostatic co-assembly. (a) Chemical structure and transmission electron microscopy (TEM) image of the cationic lipid spermine-hydroxyethyl photolinker-oleic ester (**Spillo**,  $c = 0.34 \text{ mg mL}^{-1}$ ) in 1× FOB (1× TAE, 12.5 mM MgCl<sub>2</sub>). The inset shows the photolytic degradation of **Spillo**. (b) TEM image of the 6-helix bundle (6HB) DNA origami ( $c = 7.5 \text{ nM}$ ) in 1× FOB. (c) The assembly of highly ordered fibers is driven by hydrophobic interactions and electrostatic interactions between the negatively charged phosphate groups in the DNA backbone and the positively charged amines of **Spillo**. The TEM samples are negatively stained with 2% (w/v) uranyl formate.

amines in **Spillo**, which facilitates the electrostatic co-assembly (Fig. 1c).

**Spillo**-6HB complexes were formed by mixing the two compounds together in 6HB folding buffer (FOB; 1× Tris-acetate-EDTA (TAE) and 12.5 mM MgCl<sub>2</sub>) supplemented with different amounts of NaCl. Initially, the electrostatic co-assembly of **Spillo** and 6HBs was studied using an agarose gel electrophoretic mobility shift assay (EMSA, Fig. 2a). The binding of **Spillo** and the subsequent formation of complexes immobilize the 6HBs in the gel, which is observed as a gradual decrease in the intensity of the free 6HB band with increasing stoichiometric ratios,  $n_{\text{Spillo}}/n_{\text{6HB}}$ , between **Spillo** and 6HBs. High concentrations of free ions in the assembly buffer typically prevent assembly formation by screening the electrostatic interactions between the charged compounds.<sup>22,32</sup> However, in this case, the electrostatic co-assembly was not significantly affected by the NaCl concentration of the assembly buffer, and a  $n_{\text{Spillo}}/n_{\text{6HB}} \sim 1000$  was sufficient for complexation even at high NaCl concentrations (Fig. 2a and ESI,† Fig. S7). This further confirms that the co-assembly is driven not only by electrostatic but also hydrophobic interactions.

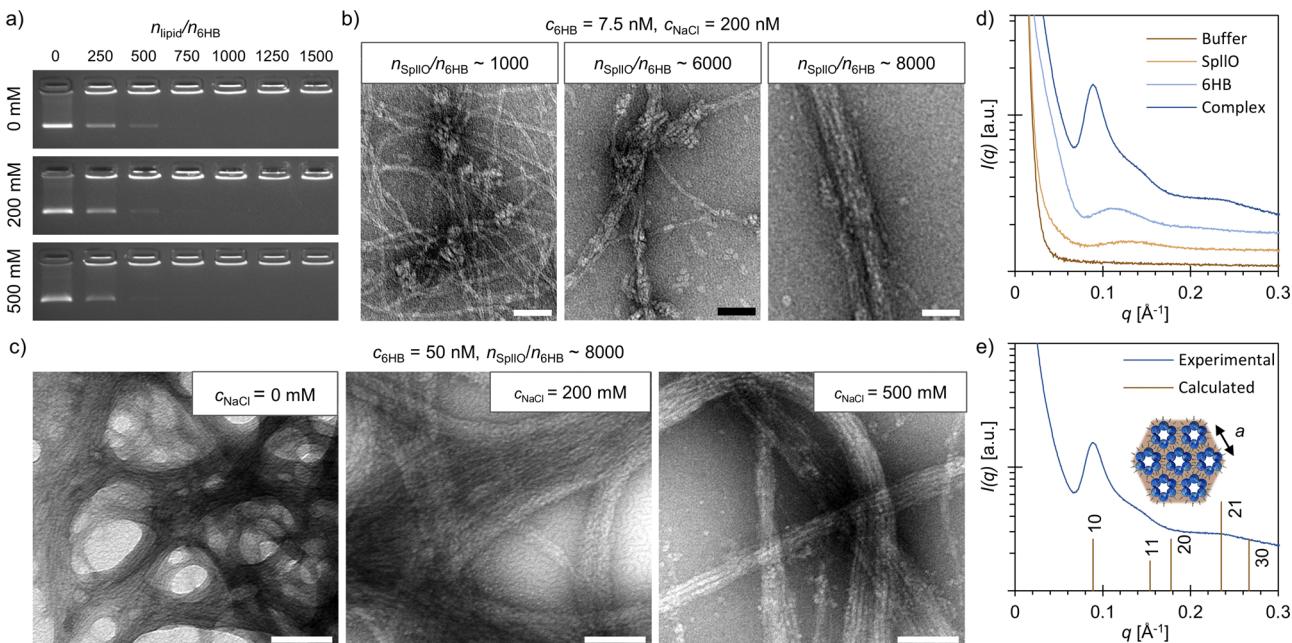
To gain additional insights into the assembly, we used transmission electron microscopy (TEM) to visualize the **Spillo**-6HB assemblies ( $c_{\text{6HB}} = 7.5 \text{ nM}$ ) formed in 1× FOB supplemented with 200 mM NaCl (Fig. 2b and ESI,† Fig. S8–S10). At  $n_{\text{Spillo}}/n_{\text{6HB}} \sim 1000$ , the 6HBs were, as also suggested by agarose gel EMSA, highly aggregated, although mostly uncoated. Small lipid-like structures were visible on the 6HB interfaces, which indicates that **Spillo** acts as a supramolecular “glue” bringing the structures together. By increasing the  $n_{\text{Spillo}}/n_{\text{6HB}}$  considerably, the amount of uncoated 6HBs decreased, and a  $n_{\text{Spillo}}/n_{\text{6HB}} \sim 8000$  was found to be required to entirely encapsulate the 6HBs with **Spillo**. Furthermore, the TEM images revealed that the structural morphology of the **Spillo**-6HB complexes changed from less ordered aggregates towards more fiber-like structures with increasing  $n_{\text{Spillo}}/n_{\text{6HB}}$ .

The initial co-assembly, shown in Fig. 2a and b, was carried out using relatively low 6HB concentrations ( $c_{\text{6HB}} = 7.5 \text{ nM}$ ), but for further characterization, we formed **Spillo**-6HB complexes at remarkably higher concentrations ( $c_{\text{6HB}} \geq 50 \text{ nM}$ , ESI,† Fig. S11–S16). As for the assemblies formed at low 6HB concentrations, a  $n_{\text{Spillo}}/n_{\text{6HB}} \sim 8000$  was found to be sufficient for the assembly (ESI,† Fig. S12–S14). At high concentrations ( $c_{\text{6HB}} = 50 \text{ nM}$ ), we also studied the effect of the NaCl concentration on the assembly formation more thoroughly. TEM shows that large, micrometer-sized fibers are formed for all three NaCl concentrations, but that the most well-aligned fibers are formed at  $c_{\text{NaCl}} = 200 \text{ mM}$  (Fig. 2c and ESI,† Fig. S11, S14, S15). In the absence of NaCl the fibers tend to aggregate, whereas at high NaCl concentrations ( $c_{\text{NaCl}} = 500 \text{ mM}$ ) the fibers are smaller and the 6HBs are more loosely packed. This is in line with the small-angle X-ray scattering (SAXS) data measured for **Spillo**-6HB assemblies formed at different NaCl concentrations, which indicates that 200 mM NaCl would be the optimum salt concentration for the assembly (ESI,† Fig. S17).

SAXS measurements were conducted on both the **Spillo**-6HB complexes and the individual components to confirm the observations from TEM (Fig. 2e and ESI,† Fig. S18). The **Spillo**-6HB complexes had a clearly different scattering profile than the individual building blocks. Additional analysis of the SAXS data for the **Spillo**-6HB complex further suggests that the complexes form a 2D hexagonal lattice. The peak positions found at  $q = 0.089, 0.148$  and  $0.235 \text{ \AA}^{-1}$  fit approximately with the reflections from [10], [11] and [21] planes for a 2D hexagonal lattice with a lattice constant of  $a = 7.1 \text{ nm}$ . The lattice constant is slightly larger than the 6HB diameter (6 nm), and therefore matches the dimensions of the building blocks. Previously, similar fiber-like structures have been reported when 6HBs were co-assembled with positively charged phthalocyanines.<sup>22</sup>

Plain DNA origami are typically susceptible to digestion by nucleases, such as DNase I,<sup>33</sup> but the structural stability against nucleases can be enhanced by applying different coatings or by encapsulating the structures.<sup>34,35</sup> Co-assembly with **Spillo** ( $c_{\text{6HB}} = 7.5 \text{ nM}$ ,  $n_{\text{Spillo}}/n_{\text{6HB}} \sim 8000$ ) was also observed to shield the 6HBs and slow down the DNase I digestion. The plain 6HBs were completely digested when exposed to DNase I at a concentration of 25 Kunitz units (KU)  $\text{mL}^{-1}$  for 1 h at RT, while the





**Fig. 2** Characterization of the complexes formed by the electrostatic co-assembly of **Spillo** and 6HBs. (a) Agarose gel electrophoretic mobility shift assay (EMSA) for a constant amount of 6HBs ( $c_{6\text{HB}} = 7.5 \text{ nM}$ ) complexed with increasing amount of **Spillo** in 1× FOB supplemented with different amounts of NaCl. The gels (2% (w/v)) were run at 95 V for 45 min. TEM images, samples negatively stained (2% (w/v) uranyl formate), of complexes formed at (b) different stoichiometric ratios  $n_{\text{Spillo}}/n_{6\text{HB}}$  ( $c_{6\text{HB}} = 7.5 \text{ nM}$ ,  $c_{\text{NaCl}} = 200 \text{ mM}$ ), scale bars 50 nm, (c) different NaCl concentrations ( $c_{6\text{HB}} = 50 \text{ nM}$ ,  $n_{\text{Spillo}}/n_{6\text{HB}} \sim 8000$ ), scale bars 100 nm. (d) Small-angle X-ray scattering (SAXS) data measured from the individual building blocks and a **Spillo**–6HB complex ( $n_{\text{Spillo}}/n_{6\text{HB}} \sim 8000$ ) in 1× FOB with 200 mM NaCl. (e) Experimental data and theoretical peak positions for a 2D hexagonal lattice with a lattice constant of  $a = 7.1 \text{ nm}$ .

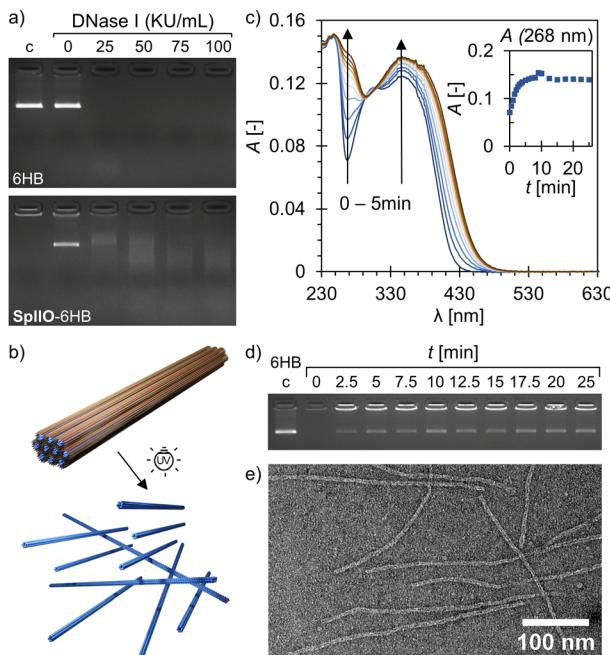
6HBs co-assembled with **Spillo** were only partially digested (Fig. 3a). The DNase I was inactivated with sodium dodecyl sulfate (SDS) before running the gel, which disassembled the **Spillo**–6HB complexes and released the 6HBs. The 6HBs embedded in the fibers formed at high concentrations ( $c_{6\text{HB}} = 50 \text{ nM}$ ) withstand DNase I at least to the same extent as the 6HBs co-assembled with **Spillo** at low concentrations (ESI,† Fig. S19).

The responsiveness of the assembly was demonstrated by exposing the **Spillo**–6HB complexes to UV-A light ( $\lambda \sim 350 \text{ nm}$ ). Irradiation with UV-A light cleaves the photosensitive *o*-nitrobenzyl group of **Spillo**, which facilitates a controlled, optically triggered release of the 6HBs from the assemblies (Fig. 3b). The photolytic degradation of plain **Spillo** was confirmed by exposing an aqueous **Spillo** solution to UV-A light and following the degradation over time using UV/Vis spectroscopy (Fig. 3c). UV-A irradiation changed the absorbance spectra significantly, and clear increases in the absorbance at 268 nm and 347 nm were observed, which is typical for the proposed photolytic degradation.<sup>27,36</sup> Moreover, after  $\sim 8 \text{ min}$  of irradiation, the **Spillo** degradation leveled off and reached a plateau. Further irradiation after that resulted in a decrease again of the absorbance at 347 nm (ESI,† Fig. S20). In addition, also TEM demonstrated that **Spillo** is degraded by UV-A light (ESI,† Fig. S21). The disassembly of the complexes ( $c_{6\text{HB}} = 7.5 \text{ nM}$ ,  $n_{\text{Spillo}}/n_{6\text{HB}} \sim 8000$ ) and the subsequent release of the 6HBs upon irradiation was studied using an agarose gel EMSA (Fig. 3d and ESI,† Fig. S22). A band with similar electrophoretic

mobility as the plain 6HB appeared in the gel after only 2.5 min of UV-A exposure, and maximum 6HB recovery of  $\sim 30\%$  was observed after  $\sim 10 \text{ min}$ . The co-assembly with **Spillo** entangles the 6HBs (as shown in Fig. 2b), and therefore some 6HBs will remain aggregated also after the UV-A exposure although the supramolecular multivalency effect is destroyed. DNA origami are known to maintain their structural integrity even upon high-dose UV-A irradiation,<sup>37</sup> and TEM further confirmed that intact 6HBs are released from the complexes (Fig. 3e and ESI,† Fig. S23). The 6HBs could also be recovered from the larger and more well-aligned fibers formed at high 6HB concentrations ( $c_{6\text{HB}} = 50 \text{ nM}$ ), but a longer irradiation time was required (ESI,† Fig. S24).

In conclusion, we have demonstrated a straightforward and scalable approach for constructing well-aligned and photo-responsive fibers by the co-assembly of a cationic photosensitive lipid **Spillo** and 6HB DNA origami. The lipid co-assembly provided protection for the 6HBs against DNase I digestion, and the **Spillo**–6HB complexes could be controllably disassembled and the 6HBs released through UV-A light exposure. Lipid nanoparticles are already widely used for mRNA delivery,<sup>38</sup> but stimuli-responsive lipid-based systems for the delivery of larger DNA origami could find intriguing uses in therapeutics. The high addressability of the DNA origami allows a variety of drugs, therapeutic proteins and antibody fragments to be precisely positioned onto the structure,<sup>39</sup> and genes folded within DNA origami have been successfully expressed in cells.<sup>40</sup>

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**Fig. 3** Functionality of the **Spillo**–6HB complexes. (a) Agarose gel EMSA of bare 6HBs (top) and **Spillo**–6HB complexes after incubation with DNase I for 1 h. The DNase I is deactivated by sodium dodecyl sulfate (SDS) before the gel run, and c is a control sample without added SDS and DNase I. (b) Schematic of the release of 6HB upon irradiation of the **Spillo**–6HB fibers with UV-A light. (c) UV/Vis spectra of **Spillo** ( $c = 0.1 \text{ mg mL}^{-1}$ ) after different UV-A irradiation times. The inset shows the absorbance at 268 nm as a function of irradiation time. (d) Agarose gel EMSA of **Spillo**–6HB complexes after UV-A exposure. (e) TEM image showing intact 6HBs released from **Spillo**–6HB complexes after 10 min of UV-A irradiation. The TEM sample is negatively stained with 2% (w/v) uranyl formate, and in all experiments the **Spillo**–6HB complexes are formed at  $c_{6\text{HB}} = 7.5 \text{ nM}$  and  $n_{\text{Spillo}}/n_{6\text{HB}} \sim 8000$ . The gels (2% (w/v)) were run at 95 V for 45 min.

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## Conflicts of interest

The authors declare no conflict of interest.

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