

Supplementary Information

Assembly of dsDNA nanocircles into dimeric and oligomeric aggregates

Damian Ackermann, Goran Rasched, Sandeep Verma, Thorsten L. Schmid, Alexander Heckel, and Michael Famulok*

Instrumentation. Automated solid-phase oligonucleotide synthesis was performed on an ABI 394 DNA-Synthesizer (*Applied Biosystems*). Mass spectra of the oligonucleotides were performed on a Bruker APEX IV Fourier-Transform Ion-Cyclotron-Resonance (FT-ICR). Polyacrylamide gel electrophoresis (PAGE) was performed on a Mini Protean 3 system from *BioRad* and for the electroelution of the DNA architectures a 422 Electroeluter from *BioRad* was used. Instruments for measuring atomic force microscopy (AFM) see below.

Buffer solutions. TEAAc: 100 mM triethylammonium acetate in H₂O; DNA storage buffer: 50 mM NaCl, 10 mM Tris·HCl in H₂O, pH 7.5; TAE-Mg buffer: 40 mM Tris, 20 mM acetic acid, 2.5 mM EDTA, 12.5 mM MgCl₂, pH 7.5.

Synthesis of anthracene modified oligonucleotide. The anthracene modified oligonucleotide was prepared by standard phosphoramidite chemistry (0.2 μmol scale) and deprotected and cleaved from solid support by treating it with 1 mL aq. NH₃-solution (33%, DNA-grade) at 55°C for 5 h. The oligonucleotide was purified by RP-HPLC (Agilent 1100 HPLC System, Nucleosil 100-5 C18 column, CS-Chromatographie Service GmbH, Germany) Flow rate: 1.0 mL/min; buffer A: 0.1 M TEAAc in H₂O; buffer B: MeCN; gradient: 0 → 30% B in 20 min. The oligonucleotide was characterized by ESI-MS: calculated 7104.4, found 7127.7 [M+Na].

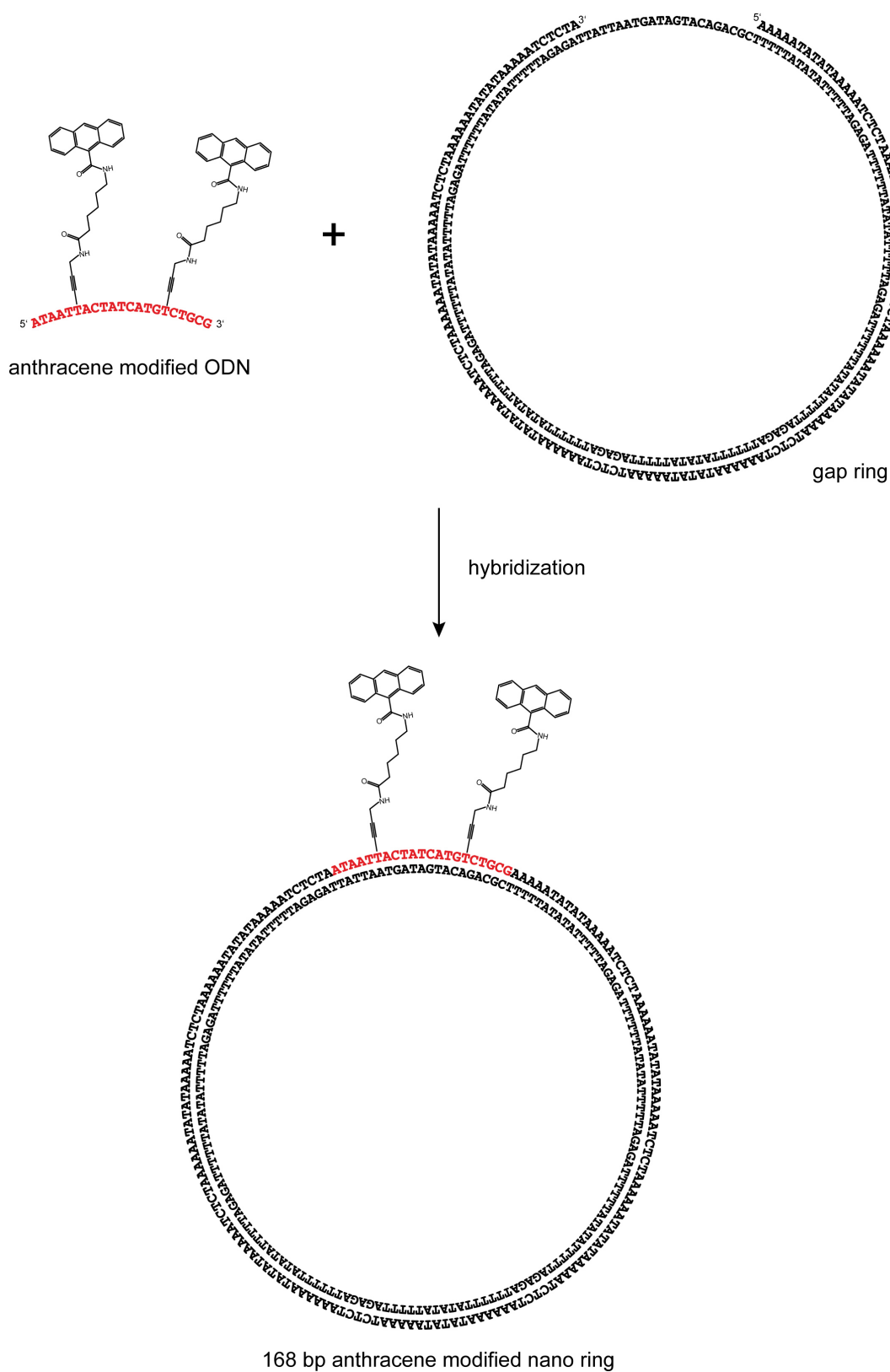
Synthesis of gap containing DNA nanocircles. All ssDNA were ordered as 5'-phosphate oligonucleotides from *Metabion*. For the ligation of the subunits the following optimised protocol was used: 4 μM of each precursors ssDNA, 3'000 U/ml T4 DNA ligase (*New England Biolabs*) and 1x ligation buffer (*New England Biolabs*) in a total volume of 200 μl. Prior to the addition of the ligase (400 u) the solution was annealed to 65°C and then allowed to cool down to 15°C during 3h. Ligation was performed at 15°C during 14 h. Complete consumption of the precursor DNA sequences was verified by native PAGE (7% acrylamide, 200 V, 40 min). As a reference a fully double stranded DNA nanoring was simultaneously run on the gel, indicating the correct band for the gap-containing DNA nanoring. The product band was excised from the gel and subjected to electroelution at 200V for 2h. After collecting the product solution the DNA nanocircles were isolated by EtOH precipitation and the pellet dissolved in 100 μl 1x DNA buffer.

Oligonucleotides for intercalator modified DNA nanocircle

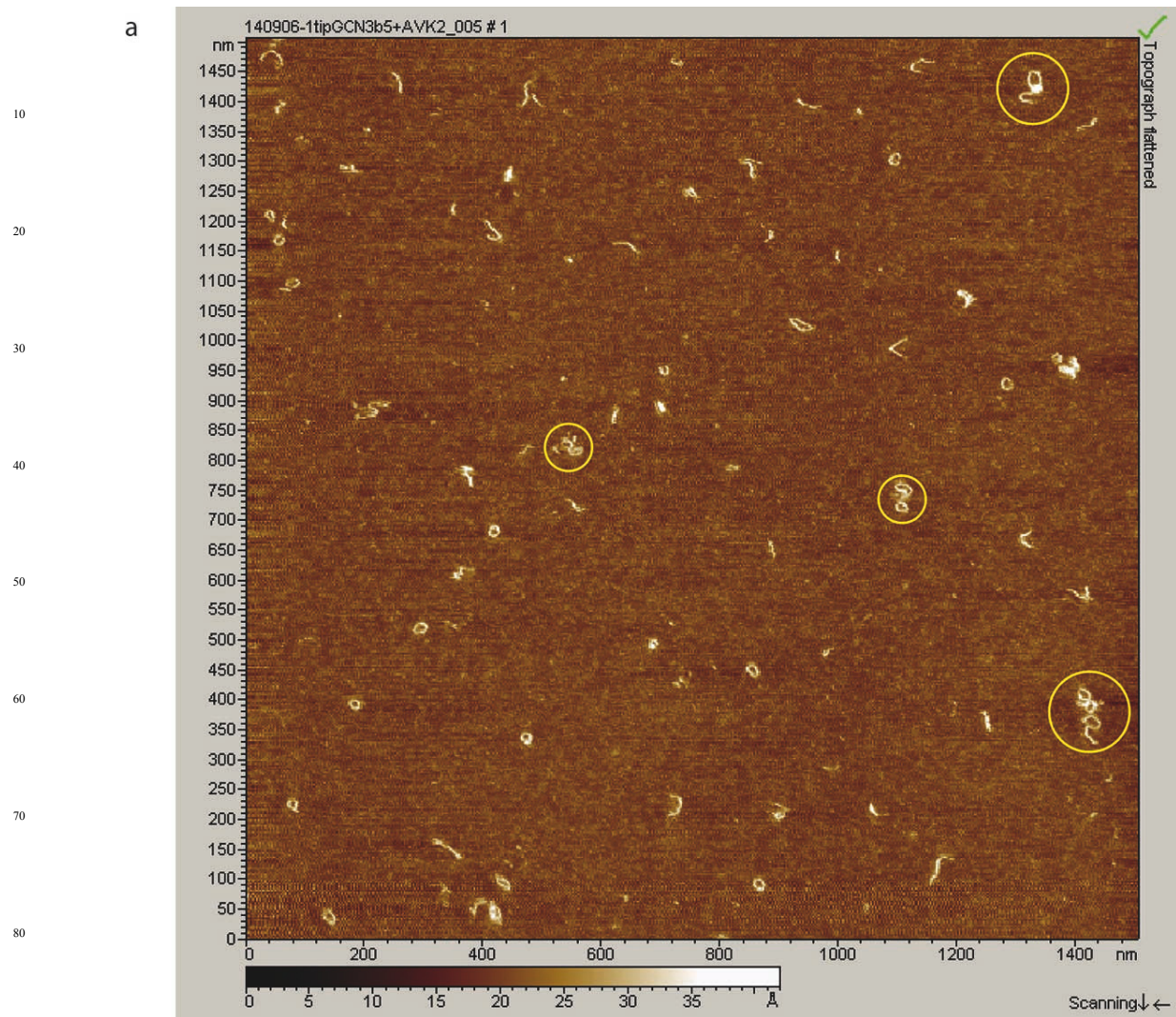
name	sequence
Alpha-f	5' -TCTCTAAAAAATATATAAAAAATCTCTAAAAAATATATAAAAAATCTCTAAAAAATAT-3'
Alpha-r	3' -TTTTTTATATATTTTTAGAGATTTTTTATATATTTTTAGAGATTTTTTATATATTT-5'

Water was added to reach a volume of 100 μ L and the sample was incubated at RT for 20 min. Before scanning another 300 μ L (10 mM) Tris-HCl-buffer were added to the liquid cell. The images were processed with WSxM 3.0 beta 9.0 software (Nanotech Electronica, Spain).

5 **AFM imaging methode II** (used for dumbbell DNA architectures). AFM images were taken on a Veeco (NY, USA) Dimension 3100 AFM with a Nanoscope IIIa controller in tapping mode. Imaging was performed both, in buffer or in air. 30 μ l of a solution of 0.1 mg polyornithin (P3655, Sigma Aldrich) in 1 ml TAE-Mg buffer, the same was used for AFM imaging, was applied on freshly cleaved mica (muscovite grade, Plano, Wetzlar, Germany) and incubated for ca. 2 minutes. The surface was
10 rinsed with milli-Q water and dried with a stream of nitrogen. For imaging, the respective samples were diluted with TAE-Mg buffer to a final concentration of between 2-5 nM. 4 μ l of this solution were applied on the pretreated mica surface and were incubated for a minute. For dry imaging the TAE-Mg buffer was briefly washed off with a few ml milli-Q water and immediately dried with a stream of nitrogen. For liquid mode imaging the sample was not washed and dried. 30 μ l of TAE-Mg
15 buffer were added both to the sample and on the probe holder. Liquid mode: A Veeco fluid cell (DTFML-DD) was used with Veeco DNP-S tips (0.12-0.58 N/m) in tapping mode. The amplitude set-point was set to 300 mV, typical resonance frequencies of the cantilevers were between 19 and 20 kHz. Scanning frequencies were between 1-2 Hz. Dry mode: ACT probes (25-75 N/m) from APP Nano (Santa Clara, CA, USA) were used for imaging in air. Typical resonance frequencies of the can-
20 tilevers were between 300 and 400 kHz. The amplitude set-point was set to 0.3 V, scanning frequencies were between 0.5 to 2 Hz. Prior to imaging the all AFM probes were irradiated with a UV hand lamp for 2-12h from a distance of 1-2 cm. AFM raw data was processed with the Nanoscope(R) 5.31 software (Veeco).

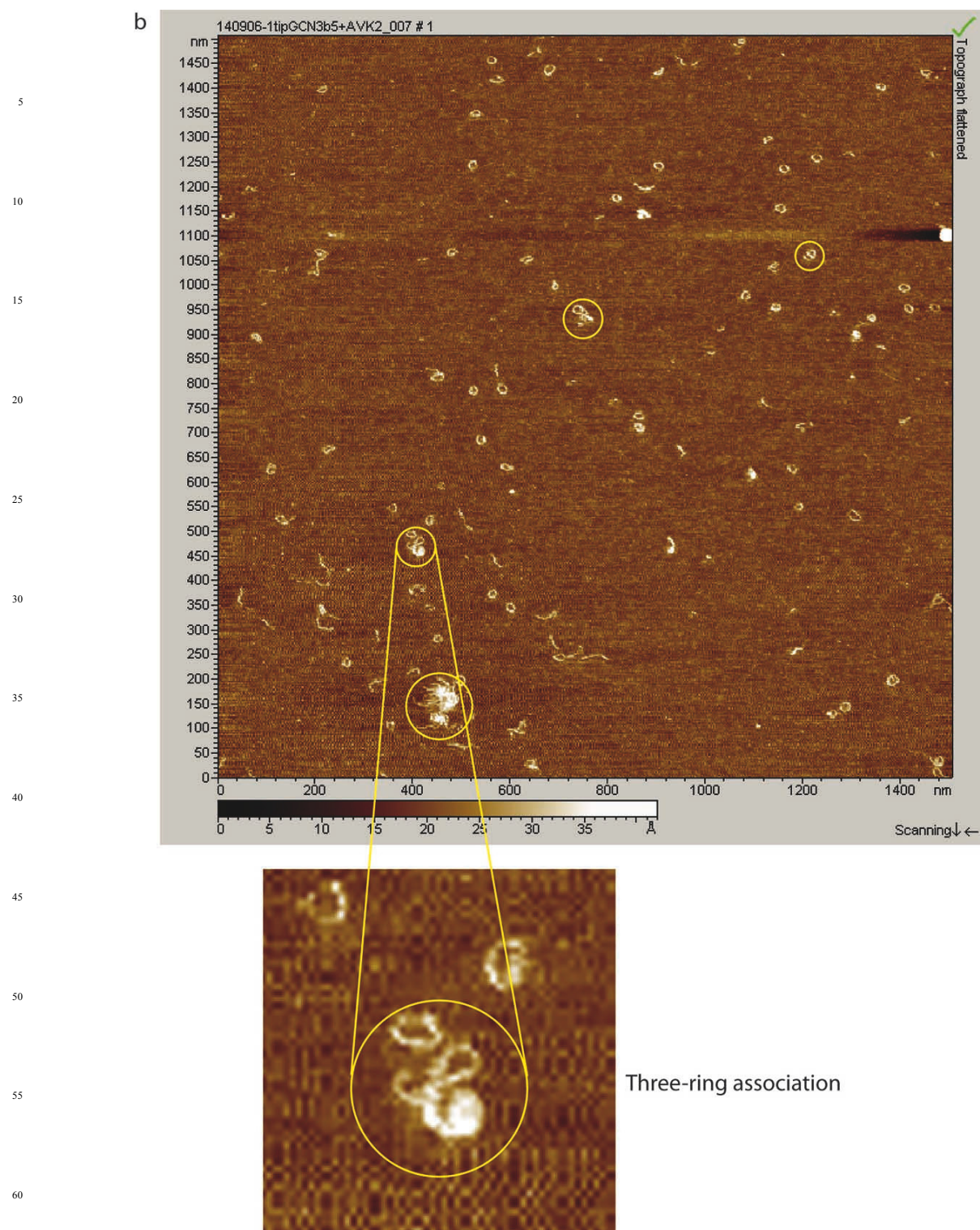


Supplementary Figure S1. Structure of the anthracene modified DNA nano ring. The modified 21-mer DNA sequence is hybridized (but not ligated) to the complementary ss-sequence of the gap ring.

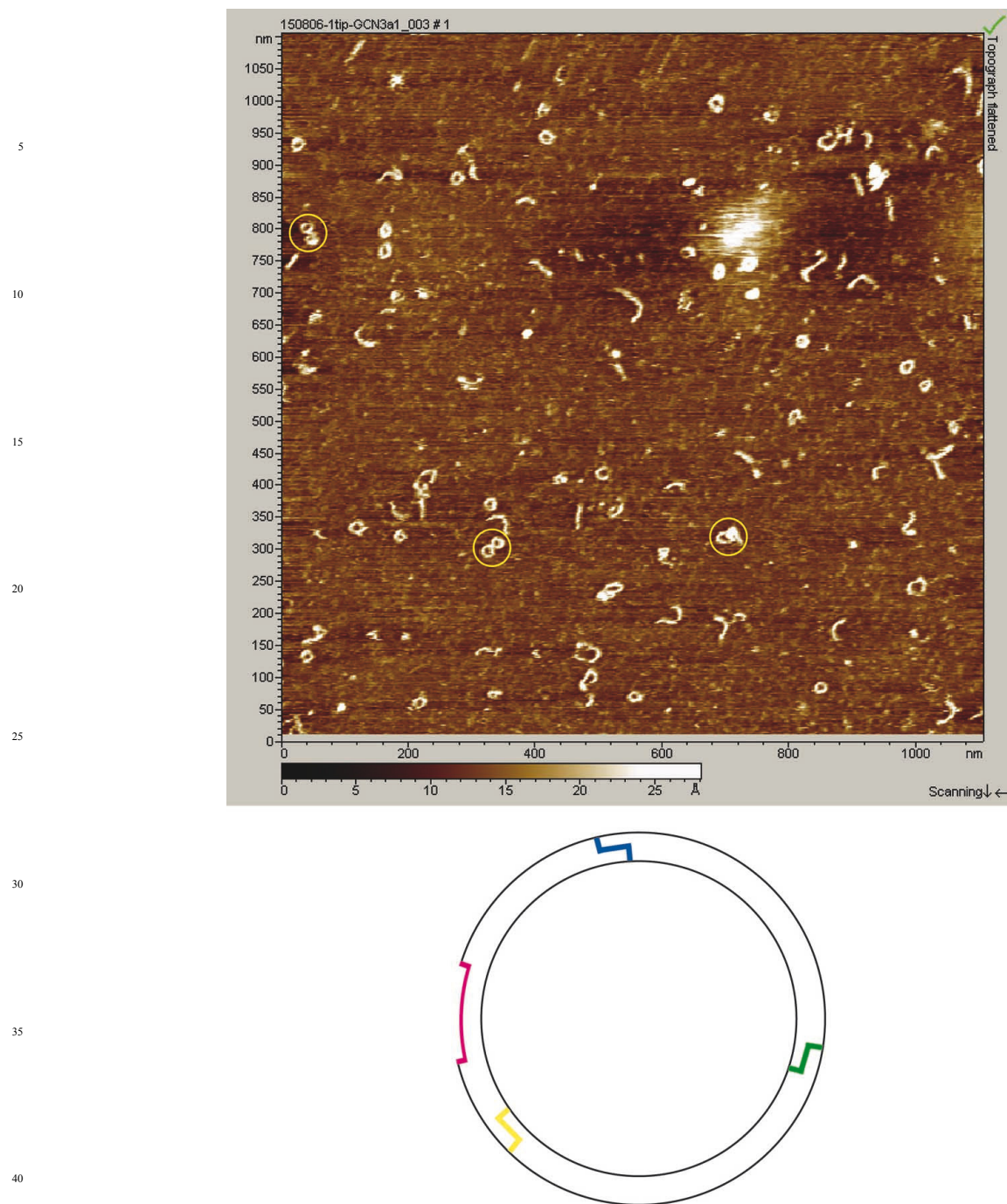


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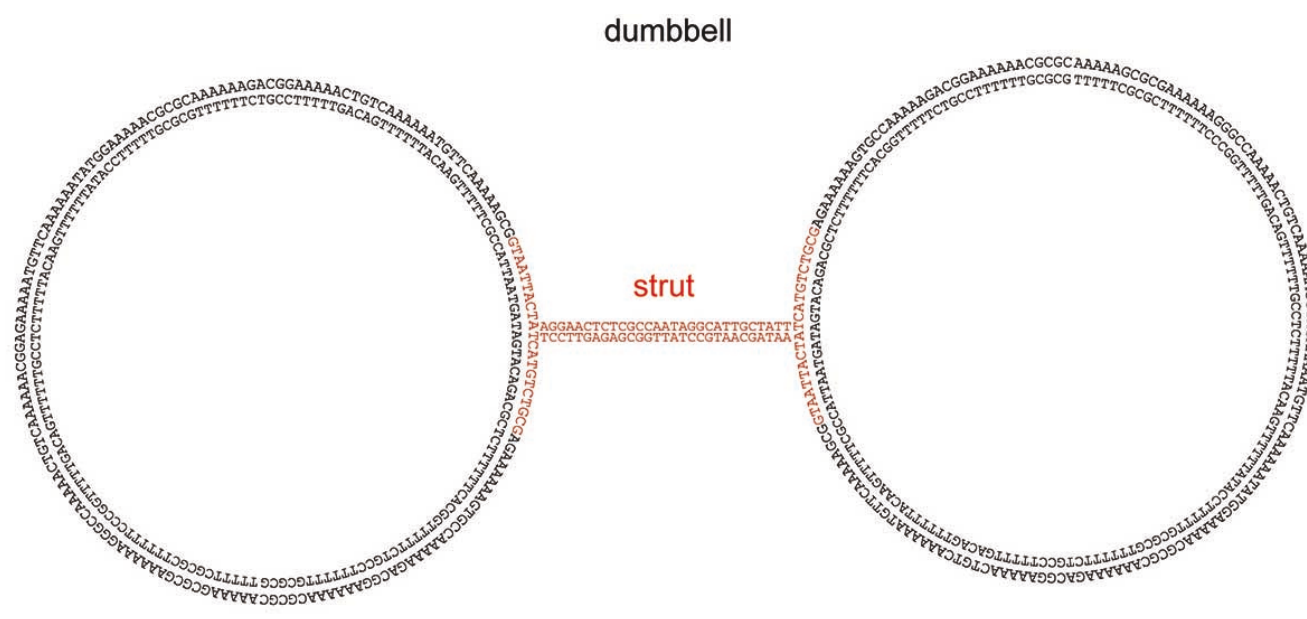
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65 **Supplementary Figure S2.** AFM scan of additional dsDNA minicircles in the presence of the anthracene-modified ODN. a) AFM scan displaying double- and triple ring aggregates. b) AFM scan displaying double and triple aggregates as well as highly oligomeric clusters.



Supplementary Figure S3. AFM scan of ss-gap containing dsDNA minicircles (bottom panel) in the absence of the anthracene-modified ODN. The AFM scan (top panel) displays few double ring aggregates.



Supplementary Figure S4. Sequences of the gap-ring and strut ODNs used in this study.