



Nuclease resistance of DNA nanostructures

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Abstract | DNA nanotechnology has progressed from proof-of-concept demonstrations of structural design towards application-oriented research. As a natural material with excellent self-assembling properties, DNA is an indomitable choice for various biological applications, including biosensing, cell modulation, bioimaging and drug delivery. However, a major impediment to the use of DNA nanostructures in biological applications is their susceptibility to attack by nucleases present in the physiological environment. Although several DNA nanostructures show enhanced resistance to nuclease attack compared with duplexes and plasmid DNA, this may be inadequate for practical application. Recently, several strategies have been developed to increase the nuclease resistance of DNA nanostructures while retaining their functions, and the stability of various DNA nanostructures has been studied in biological fluids, such as serum, urine and cell lysates. This Review discusses the approaches used to modulate nuclease resistance in DNA nanostructures and provides an overview of the techniques employed to evaluate resistance to degradation and quantify stability.

It has been almost 40 years since Ned Seeman published his thoughts on using DNA as a building block for the assembly of nanostructures, a field now called DNA nanotechnology¹. Seeman's original vision was to use DNA as a framework to crystallize other hard-to-crystallize macromolecules. Early work on DNA-based construction mostly involved the creation of DNA junctions, motifs and devices^{2–4}, and, over the past four decades, multiple strategies have been developed to create diverse architectures using DNA⁵. The library of DNA nanostructures, ranging from the nanometre to the micrometre scale^{6–8}, now includes geometric patterns⁹, polyhedral objects¹⁰, nanoscale bunnies¹¹, a box with a lid¹², a cargo-moving assembly line¹³ and even a miniature Mona Lisa¹⁴. More recently, the focus has moved from proof-of-concept construction using DNA towards the applications of these structures. Being a natural material, DNA nanostructures have found the most use in biological applications^{15–17}. Indeed, nanocages built from DNA have been used in drug delivery¹⁸, reconfigurable DNA devices in biosensing¹⁹ and DNA structures conjugated to fluorophores and ligands for imaging²⁰ and modulating cellular behaviour²¹. Although each of these applications has specific prerequisites, one common requirement is that the DNA nanostructures should remain stable under physiological conditions and be able to withstand degradation by nucleases.

For practical application, DNA nanostructures need to meet certain biostability thresholds. For example, in

biosensing, DNA nanodevices must remain intact when mixed with samples (typically, biofluids such as blood, serum and urine), as unpredictable and sudden degradation can lead to false signals. In drug delivery, the biostability of DNA nanostructures is important to protect the drug from harsh physiological conditions until it reaches the destination site in the body. In cellular modulation and bioimaging, DNA nanostructures need to remain largely intact in the cellular or bodily environment to complete their functions (to deliver regulating proteins or accumulate at target tissues, respectively). A major reason for degradation of DNA nanostructures in these conditions is the presence of nucleases (enzymes that degrade nucleic acids). To address this issue, strategies have been developed to modulate the biostability of DNA nanostructures and to increase their suitability for biological applications.

In this Review, I discuss the inherent biostability of DNA nanostructures and the potential use of nanostructures with enhanced nuclease resistance in biological applications. In particular, I focus on the methods developed to tune the nuclease resistance of these structures, the techniques used to analyse their nuclease resistance and the comparative analysis of results from key studies. More general ideas of DNA nanotechnology are beyond the scope of this Review, and interested readers are directed to other reviews covering a range of topics, including biosensing^{22–24}, drug delivery^{25–27}, bioimaging^{28–30}, cellular programming^{31,32} and scaffolding^{33,34}.

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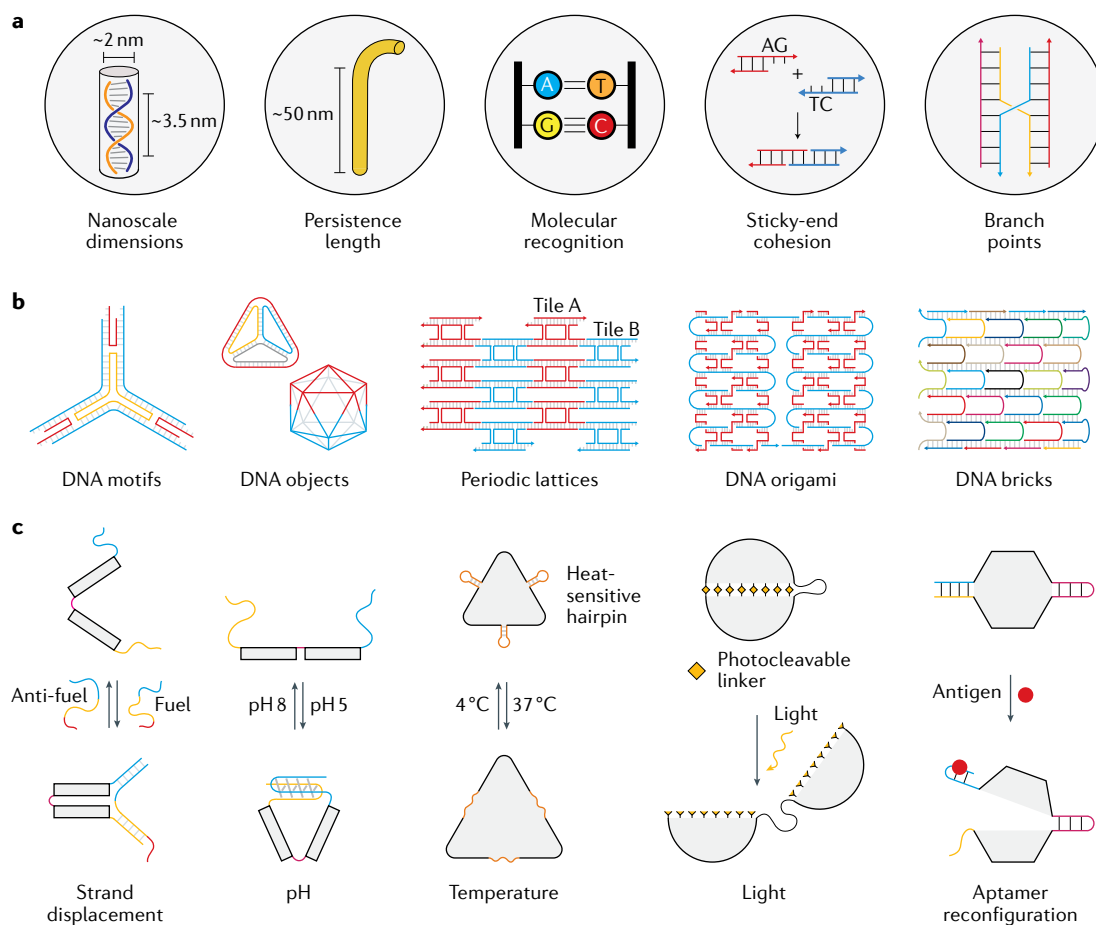


Fig. 1 | Concept, design and construction of DNA nanostructures. **a** | Properties of DNA that make it suitable for the bottom-up construction of nanostructures. **b** | Structural DNA nanotechnology involves the construction of DNA-based structures with different geometries, periodicities or spatial patterns, including motifs³⁶, objects such as polyhedra^{35,144}, periodic 2D lattices¹⁴⁵, DNA origami³⁷ and DNA bricks³⁸. **c** | Dynamic DNA nanotechnology involves the construction of DNA devices that operate in response to external stimuli, such as toehold-based strand displacement for nucleic acids¹⁰⁷, pH changes⁴³, temperature changes⁴⁵, light⁴⁶ (ultraviolet or near-infrared, for example) and antigen–aptamer interactions⁴¹.

Construction using DNA

DNA is well known as the carrier of genetic information, but is also a natural nanoscale material. The characteristics of DNA that make it a suitable building block for nanoscale architectures include its nanoscale dimensions, persistence length of ~50 nm (that is, its ‘stiffness’ for robust construction) and predictable Watson–Crick base pairing, which allows for programmed assembly (FIG. 1a). Furthermore, multidimensional assembly is possible by creating branched DNA structures that can be connected by short single-stranded extensions (sticky ends) to create larger assemblies. Building using DNA has led to two areas of DNA nanotechnology: structural and dynamic. Structural DNA nanotechnology mainly involves the design and construction of DNA-based structures with different geometries, periodicity or spatial patterns (FIG. 1b). Researchers have created various DNA nanostructures using tile-based assembly^{35,36}, DNA origami³⁷ and DNA brick strategies³⁸. In dynamic DNA nanotechnology, DNA structures are designed to reconfigure upon recognition of external moieties, such as nucleic acids^{39,40}, antigens^{12,41} or ligands^{40,42},

or in response to environmental changes, including changes in pH^{43,44}, temperature⁴⁵ or light^{46,47} (FIG. 1c). Such stimuli-responsive devices are useful in biosensing and as components for triggered release in drug-delivery carriers. Construction principles for DNA nanostructures, assembly strategies and the types of motifs and structures have been discussed elsewhere^{3,6,48,49}.

DNA nanotechnology applications

Developments in DNA nanotechnology have not always adhered to the principle of ‘form follows function’: in most cases, the function of a DNA nanostructure has followed its form. Demonstrations of the applications of DNA in different fields have increased exponentially in the past decade^{50,51}. In materials science, DNA can serve as a copolymer for creating integrated chips^{52,53}, pooled DNA libraries are used in data storage and molecular computation^{54,55}, and DNA-based moulds have been used for the assembly of metallized nanosheets and electrical nanowires^{56,57}. In chemistry, DNA has been used in directing polymer construction⁵⁸ and site-specific chemical reactions⁵⁹, and as scaffolds for the structure

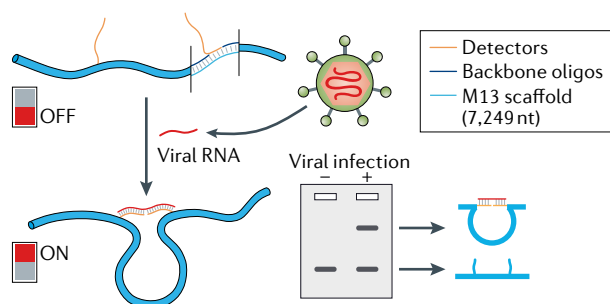
DNA origami

A technique that uses hundreds of short ‘staple’ strands to fold a long piece of single-stranded ‘scaffold’ DNA into desired shapes.

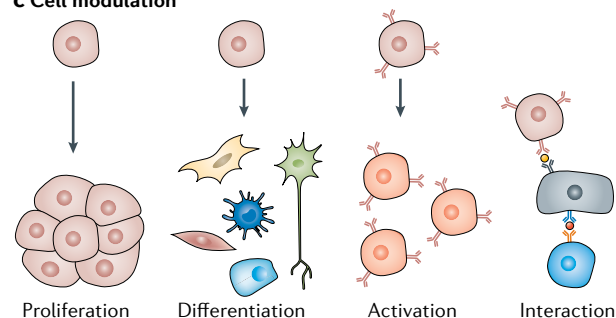
a Diagnostics



Example: viral RNA detection



c Cell modulation



Example: structurally tunable extracellular matrix

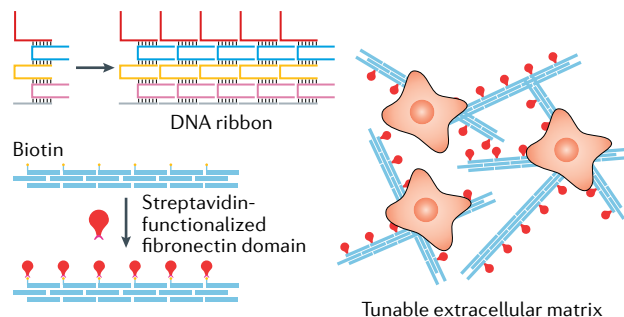
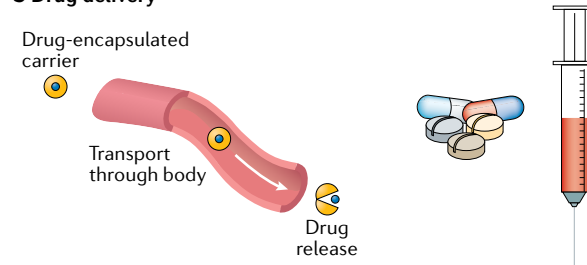
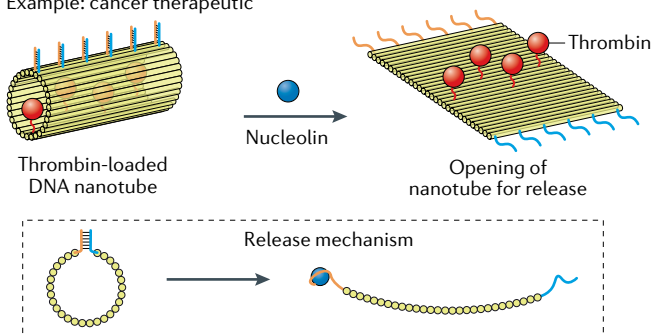


Fig. 2 | Biological applications of DNA nanostructures. **a** | DNA nanostructures can be used in diagnostics for disease detection and monitoring. The example shown is a DNA nanoswitch made from a long M13 scaffold strand and short complementary backbone oligonucleotides. When the single-stranded detectors of the nanoswitch bind to viral RNA, the nanoswitch reconfigures to a looped 'on' state that can be visualized on a gel⁶⁵. **b** | DNA nanostructures are used as carriers to deliver drugs to specific sites in the body or to cellular compartments. For example, a DNA origami nanotube loaded with thrombin (an anticoagulant) can be targeted towards tumours using a nucleolin-targeting aptamer. On reaching the tumour, the

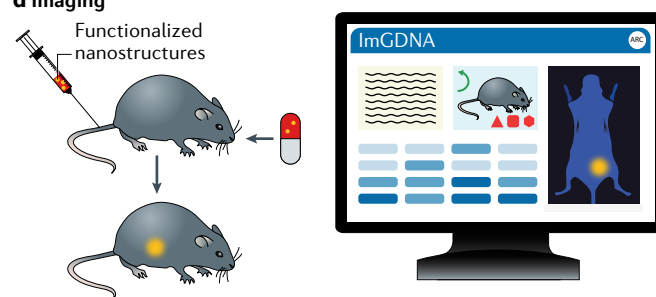
b Drug delivery



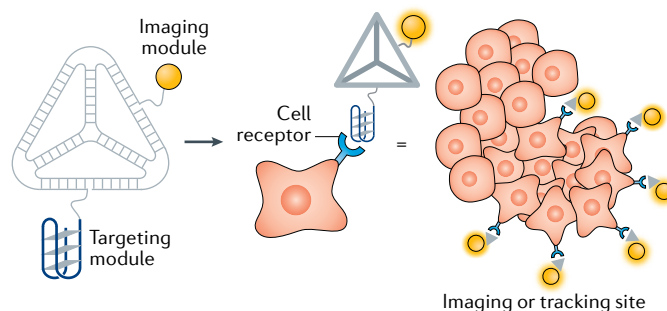
Example: cancer therapeutic



d Imaging



Example: imaging cancer cells



determination of macromolecules^{60–62}. However, most of the recent studies are in biological applications, some of which are briefly discussed below.

Biosensing. DNA-based biosensors are cost-effective and sensitive^{63,64}, and are being developed for use as point-of-care diagnostic tools (FIG. 2a). DNA-based sensing

strategies rely on the recognition event between the DNA nanostructure and a target analyte (such as in nucleic acid and protein sensing) or the conformational change of a DNA-based device (upon sensing, for example, pH changes or light)²². Advantages of DNA nanostructures for biosensing include the highly precise design based on base pairing, the large surface-to-volume ratios and the low-cost synthesis of functionalized DNA strands. So far, DNA nanostructures have been used to detect nucleic acids (including microRNAs¹⁹, viral RNAs⁶⁵ and pathogenic DNA⁶⁶), antigens⁶⁷, antibodies⁶⁸ and small molecules⁴⁰, as well as in sensing in vivo pH changes⁴³.

Drug delivery. In biomedicine, DNA nanostructures have potential utility as drug-delivery carriers (FIG. 2b). DNA provides precise control over the size and geometry of the nanostructures, generating homogeneous populations of drug carriers — a unique advantage over other self-assembling materials and nanoparticles⁴⁸. DNA strands can also be synthesized with functional moieties that are useful for imaging⁶⁹, site-specific targeting⁷⁰ and triggered delivery⁷¹. Furthermore, DNA nanostructures are biocompatible and exhibit quick renal clearance and minimal toxicity⁷². DNA nanostructures can be tailored to carry various drug cargos, including unmethylated cytosine–phosphate–guanine (CpG) sequences for immunotherapy⁷³, monoclonal antibodies as immune checkpoint inhibitors⁷⁴, small interfering RNA⁷⁵ and antisense oligonucleotides⁷⁶ for gene therapy, proteins for vaccine development⁷⁷ and metallic nanoparticles for photodynamic therapy⁷⁸. Functionalized DNA nano-carriers are, thus, useful in controlling the circulation time, drug-release rate and specificity to a particular target site, resulting in increased delivery efficiency and efficacy of the encapsulated drug^{25,26}.

Cell modulation. DNA nanostructures have also been used in modulating cell behaviour and activity⁷⁹ (FIG. 2c). For example, DNA tetrahedra promoted the proliferation and differentiation of stem cells⁸⁰, and peptide-functionalized DNA nanotubes were used in the differentiation of neural stem cells into neurons⁸¹. In immunotherapy, DNA nanostructures can activate macrophages to provide anti-inflammatory and anti-oxidative responses⁸². In other studies, DNA nanostructures have been used to influence cell migration, a factor important in wound healing and tumour cell metastasis²¹. The activation of specific cellular pathways by DNA nanostructures and their effects are currently under investigation⁸³, which may lead to the development of tools useful in regenerative medicine and gene therapy.

Bioimaging. DNA nanostructures tagged with different imaging modules, such as organic fluorophores, radioactive isotopes and quantum dots, are used for quantitative functional imaging in cells and live model organisms (FIG. 2d). The use of DNA polyhedra and many versions of DNA devices have been demonstrated in imaging tumour-related messenger RNAs^{84,85} and membrane proteins⁸⁶, as well as in tracking cell-entry pathways of drug-carrying DNA nanostructures⁶⁹. Functionalized

DNA nanostructures are also used as imaging standards to calibrate super-resolution microscopes, such as in direct stochastic optical reconstruction microscopy (dSTORM) and stimulated emission depletion (STED)⁸⁷. In addition, DNA-PAINT (DNA-based point accumulation for imaging in nanoscale topography) is being explored for imaging cellular architectures and the real-time trajectories of membrane proteins in live cells⁸⁸.

DNA nanostructure stability

As a molecule, DNA is highly stable, with a half-life of ~500 years⁸⁹. However, when assembled into a nanostructure, the stability of DNA can vary, depending on the environment. In each biological application described above, the stability of the assembled DNA nanostructure is a key parameter. To establish the viability of DNA nanostructures in different applications, the stability of these structures is often tested at elevated temperatures⁹⁰, in high-salt and low-salt conditions⁹¹, and in the presence of nucleases⁹², chaotropic⁹³ agents and crowding agents⁹⁴. In most cases, biological operations are tested at the physiological temperature of 37 °C. Depending on the application, stability at 37 °C for multiple hours might suffice. In terms of the salt and cation concentration, there needs to be a compromise between the conditions required for assembly versus those in the body. In addition, the stability and functionality of DNA structures are also dependent on where they are used, and can be affected by the local density, molecular flexibility and interactions with other molecules present in the sample, such as biological nucleic acids and serum opsonin proteins (which can lead to opsonization)⁹⁵.

A major challenge to the application of DNA-based nanostructures is their susceptibility to attack by a variety of nucleases present in body fluids such as blood, urine and saliva. Nucleases cleave various DNA and RNA substrates (BOX 1), and are involved in biological processes such as DNA repair, replication and recombination⁹⁶. Nuclease activity is also required for structural alterations of nucleic acids, such as in topoisomerization⁹⁷, site-specific recombination⁹⁸ and RNA splicing⁹⁹. Furthermore, nucleases form an integral part of the host immune system in degrading foreign nucleic acids¹⁰⁰ and have a role in programmed cell death¹⁰¹. Although nucleases have important biological functions, they also interfere with the function of DNA-based structures in biological applications, necessitating the study of DNA nanostructure biostability in vivo or in conditions that mimic body fluids. As substitutes for physiological conditions, the stability of DNA nanostructures is tested in vitro using cell and tissue culture media supplemented with mammalian serum (such as fetal bovine serum (FBS)), which contains various nucleases¹⁰².

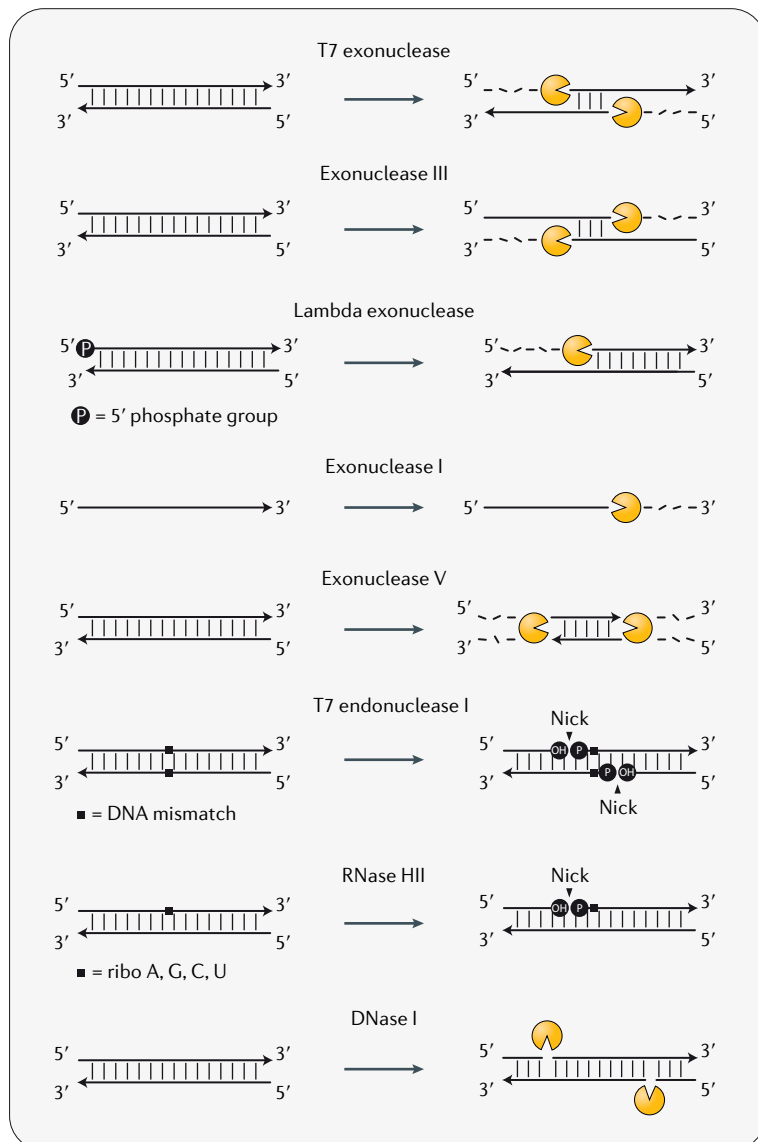
Although not all applications require DNA nanostructures to be nuclease resistant, it would be an advantage in most. Exceptions include applications such as biomolecular analysis, in which the molecules under observation are usually isolated and purified before their interactions are tested. Thus, even if they are typically present in a biofluid or tissue, the experiments do not occur in these conditions and are rarely affected by

Opsonization

Non-specific interactions of any foreign body, including DNA nanostructures, with plasma proteins, resulting in the formation of a protein corona.

Box 1 | Examples of nucleases and their activities

Nucleases are enzymes that cleave phosphodiester bonds between the sugar and phosphate moieties of DNA. Depending on their activity, nucleases are categorized as exonucleases (cleavage reactions occur at the terminus) or endonucleases (cleavage occurs within the DNA strand). Exonucleases can be further classified as 5'-end processing (such as T7 exonuclease) or 3'-end processing (such as exonuclease III) according to their polarity of consecutive cleavage, with some enzymes preferring 5'-phosphorylated double-stranded DNA (for example, lambda exonuclease). Exonucleases can also be single-strand-specific (for example, exonuclease I) or double-strand-specific (such as exonuclease V). Endonucleases display similar specific activities, with some catalysing cleavage at mismatch sites within double-stranded DNA (for example, T7 endonuclease I), at the site of a ribonucleotide (such as RNase HII) or non-specifically at random sites (such as DNase I). The activity of nucleases can also be sequence-specific or structure-specific.



Enzyme unit

The amount of enzyme that will catalyse or produce a specific amount of a substrate or product, respectively, under the specified conditions of an assay.

nucleases in the system. There are also some applications in which the native stability of DNA nanostructures is an advantage, or even susceptibility to nuclease degradation might be beneficial (for quick or spontaneous drug release, for example). Nevertheless, in many cases, the stability of DNA nanostructures needs to be enhanced before they are used. In this regard, several strategies

are being developed to create nuclease-resistant DNA nanostructures.

Strategies to modulate nuclease resistance

Strategies to modulate nuclease resistance can begin at the design stage, involve chemical modification or protective coating of the nanostructure, or occur at the point of use by solution treatment, each of which is discussed in more detail in the following sections. Unless otherwise mentioned, the studies discussed below were carried out at the physiological temperature of 37 °C. The structures tested for each strategy, the analysis technique and the resulting nuclease resistance are compiled in TABLE 1.

Design and inherent nuclease resistance. The structural design of DNA nanostructures influences the nuclease resistance. DNA tiles and motifs contain multiple strands hybridized together in a tightly woven fashion. Similarly, larger (and more complex) origami structures contain multiple helical domains packed into bundles (arranged in, for example, square or honeycomb lattices)^{37,103}. This close-packed structure in DNA origami and other multi-helix structures is more nuclease resistant than linear duplexes or plasmid DNA¹⁰⁴ (FIG. 3a). For example, 1 enzyme unit (U) of DNase I completely degraded 65 ng of duplex plasmid DNA (pET24b) in only 5 min, whereas it took ~60 min to degrade 2 ng of a 24-helix-bundle DNA origami structure¹⁰⁴. This result indicates the potential of multilayer DNA origami objects as encapsulation agents for use with intercalating drugs, for example. However, DNA nanostructures are required to be hollow to encapsulate cargos such as nanoparticles and proteins. Hollow wireframe structures, such as DNA tetrahedra¹⁰⁵, have also shown higher stability in 10% FBS (up to 42 h) than linear duplexes, possibly owing to their non-native geometries. The influence of molecular topology on nuclease resistance has also been noted¹⁰⁶: a reconfigurable DNA device based on DNA tweezers¹⁰⁷ was more nuclease resistant in the closed state than in the open state (~37 h versus 20 h in 70% human serum, respectively), possibly owing to the stacking of two helices side by side in the closed state¹⁰⁶ (FIG. 3b).

Although, in the above examples, the structures are fortuitously nuclease resistant, this feature can also be intentionally designed into DNA nanostructures. In DNA tetrahedra, placement of restriction sites at specific locations on the duplex edges affected the activity of the enzyme¹⁰⁵. Structures with the restriction site located near the vertex were protected against degradation by DdeI restriction enzyme, whereas those with the restriction site in the middle fully degraded (FIG. 3c). Another possible strategy to enhance the nuclease resistance is to increase the number of crossovers within the nanostructures. For example, a study⁹² on paranemic crossover (PX) DNA¹⁰⁸ revealed the exceptional nuclease resistance of the motif, attributed to the continuous crossovers within the structure (FIG. 3d). Compared with duplexes, PX DNA was up to ~2,800-fold more resistant to four different nucleases (DNase I, exonuclease V, T5 exonuclease and T7 exonuclease), as well as to human serum and urine. Motifs with fewer crossovers

Table 1 | Reported nuclease resistance of various DNA nanostructures

Strategy	Structure (size)	Test environment	Analysis technique	Stability metrics
Design				
Close-packed helices ¹⁰⁴	24-HB (100 nm)	1 U DNase I	AGE	Duplex plasmid DNA: degraded in 5 min Close-packed helices: degraded in 60 min
Topology ¹⁰⁶	Tweezers (~14 nm)	70% human serum	PAGE, FRET	Closed state: degraded in ~37 h Open state: degraded in ~20 h
Restriction-site location ¹⁰⁵	Ligated tetrahedron (7 nm)	Ddel restriction enzyme	PAGE	Middle of edge: fully degraded Near vertex: fully protected
Increased crossovers ⁹²	PX, DX and duplex DNA (~13 nm)	DNase I, exonuclease V, T5 and T7 exonucleases (for 1 h); 10% FBS, human serum and urine (for 24 h)	PAGE	PX: ~100% intact DX: 0–30% intact Duplex: almost fully degraded
Solution treatment				
FBS heat treatment ¹⁰⁹	Octahedron (50 nm)	Media + 10% FBS for 24 h	AGE	Without heat treatment: 0% intact Heated FBS: almost 100% intact
Nuclease inhibitors (actin) ¹⁰⁹	Octahedron (50 nm), nanotube (400 nm), nanorod (89 nm)	Media + 10% FBS for 24 h	AGE	Without actin: 0% intact With actin: ~100% intact
Minor-groove binders (DAPI) ¹¹⁰	Pentagonal bipyramid (~40 nm)	10% mouse serum	AGE	Native structures: stable for 3 h With groove binders: stable for 24 h
Ethylenediamine buffer ¹¹¹	Tetrahedron (14 nm)	0.5 U DNase I	PAGE	In TAE/Mg ²⁺ buffer: 0% intact In ethylenediamine buffer: ~100% intact
Chemical modifications				
Ligation ^{105,113}	Tetrahedron (7 nm), prism (7 nm)	10% FBS	PAGE	ssDNA: degraded in 0.8 h Ligated tetrahedron: degraded in 42 h
Crosslinking (click chemistry) ¹¹⁴	Nanotube (~30 nm)	Exonuclease I for 3 h	AGE	Native: fully degraded Crosslinked: partially degraded
Crosslinking (UV-induced T–T dimers) ¹¹⁵	Brick-like DNA origami (~70 nm)	0.4 U ml ^{−1} DNase I	AGE	Native: stable for 10 min Crosslinked: stable for 60 min
Hexanediol and hexaethylene glycol ^{113,116}	Triangular prism (7 nm) ¹¹³ , tetrahedron (7 nm) ¹¹⁶	Media + 10% FBS	PAGE	Lifetime of unmodified prism: 18 h Lifetime with hexanediol: 55 h
L-DNA ¹¹⁷	4-Arm junction (~5 nm), nanotube (30–70-nm width, ~μm long)	2 U μl ^{−1} (exonuclease I) or 20 U μl ^{−1} (exonuclease III) for 45 min	PAGE, AFM	Native junction: completely degraded L-DNA junction: almost fully intact
Unnatural base pairs ¹¹⁸	6-Arm junction (~14 nm)	T7 exonuclease for 12 h	PAGE	Native structure: completely degraded Modified: partially degraded
Protective coatings				
HSA–DNA dendrite conjugates ¹²⁰	Cube (~7 nm)	Media + 10% FBS for 48 h	PAGE	ssDNA: 33-min half-life Protected cube: up to 22-h half-life
Dendritic oligonucleotides ¹²¹	DNA brick (~50 nm)	100 U ml ^{−1} DNase I for 1 h	AGE	Native: fully degraded with 5 U ml ^{−1} Coated: 50% degraded with 100 U ml ^{−1}
PEGylated lipid bilayer ¹²²	Octahedron (76 nm)	20 U DNase I for 24 h	Fluorescence	Without envelope: ~30% intact With envelope: ~85% intact
Cationic polysaccharides ¹²³	Origami rod (350 nm), bottle (50 × 25 nm)	10 U ml ^{−1} DNase I	AGE, TEM	Native structure: stable for 1 h With protection: stable for 24 h
PEG–polylysine block copolymers ¹²⁶	Rectangle (~100 nm), 6-HB (~600 nm), truss (~20 × 200 nm)	1 μl of 16 U ml ^{−1} DNase I for 16 h	TEM, AFM	Native: 0% intact Protected: 100% intact
	24-HB (~100 nm)	1 μl of 2,000 U ml ^{−1} DNase I for 16 h		
Oligolysine–PEG copolymer ¹²⁴	Origami barrel (~60 nm)	Media + 10% FBS	AGE, TEM	Native structure: 5-min half-life With oligolysine: 50-min half-life

Table 1 (cont.) | Reported nuclease resistance of various DNA nanostructures

Strategy	Structure (size)	Test environment	Analysis technique	Stability metrics
Protective coatings (cont.)				
Crosslinking of oligolysine coating ¹²⁵	Origami barrel (60–90 nm)	1 U μL^{-1} DNase I	AGE, TEM	Without crosslinking: 16-min half-life With crosslinking: ~66-h half-life
BSA–dendron conjugates ¹²⁷	60-HB (20 \times 20 \times 33 nm)	10 U DNase I in 20 μL reaction for 1 h at RT	AGE	Native: <20% intact With protection: ~100% intact
Peptoids ¹²⁸	Octahedron (29 nm)	15 $\mu\text{g mL}^{-1}$ DNase I for 30 min	AGE, TEM	Native structure: completely degraded With protection: almost fully intact
Silica coating ¹³⁰	24-HB (~100 nm)	4 U mL^{-1} DNase I for 3 h	AGE, TEM	Native structure: completely degraded
	13-Helix ring (66 nm)	0.5 U mL^{-1} DNase I for 3 h		Silicified structures: almost fully intact

Unless mentioned otherwise, listed experiments were conducted at 37 °C. AFM, atomic force microscopy; AGE, agarose gel electrophoresis; BSA, bovine serum albumin; DAPI, 2-(4-amidinophenyl)-1H-indole-6-carboxamide; DX, double crossover; EDTA, ethylenediaminetetraacetic acid; FBS, fetal bovine serum; FRET, Förster resonance energy transfer; HSA, human serum albumin; n-HB, n-helix bundle; PAGE, polyacrylamide gel electrophoresis; PEG, polyethylene glycol; PX, paranemic crossover; RT, room temperature; ssDNA, single-stranded DNA; TAE, tris-acetate-EDTA; TEM, transmission electron microscopy; UV, ultraviolet.

Restriction enzyme

An enzyme that recognizes a specific, short nucleotide sequence and cuts the DNA only at that specific site, which is known as a restriction site or target sequence.

Paranemic crossover (PX) DNA

A four-stranded DNA motif that consists of two adjacent double-helical DNA domains connected by strand crossovers at every possible point between two side-by-side helices.

showed reduced stability, suggesting that the increased number of crossovers in PX DNA confers additional biostability.

Solution treatment and nuclease inhibitors. The effect of nucleases on the lifetime of DNA nanostructures can be minimized by using nuclease inhibitors or changing the environment in which the structures are assembled or used. One approach is to expose the samples to conditions that denature the nucleases. For example, DNA origami nanostructures (an octahedron, nanotube and nanorod) incubated in RPMI cell culture medium supplemented with 10% FBS degraded completely within 24 h (REF.¹⁰⁹). By contrast, heating the FBS to 75 °C for

10 min before addition to the cell culture medium prolonged the nanostructure lifetime in the medium (with almost no degradation), owing to the inactivation of nucleases in the serum (FIG. 4a). However, heating the FBS might also have a more general effect on serum proteins, which could affect cell growth and phenotype.

Nuclease inhibitors can also be used to reduce the effect of nucleases in solution. For example, the addition of a tenfold molar excess of actin, a known nuclease inhibitor, extended the lifetime of DNA nanostructures against DNase I (REF.¹⁰⁹) (FIG. 4b). In contrast to heat treatment of FBS, which is incompatible with *in vitro* tissue culture, inclusion of actin (up to 200 nM) had no observable effect on cell growth and viability. In other cases, compounds that restrict access to the minor groove of double-stranded DNA can competitively inhibit DNase I activity (FIG. 4c). In a study reported in a recent preprint article, the efficiency of different classes of minor-groove binders on reducing the nuclease degradation of wireframe DNA origami structures (a two-helix pentagonal bipyramid) was tested¹¹⁰. Diamine 2-(4-amidinophenyl)-1H-indole-6-carboxamide (DAPI) was the most potent stabilizer, increasing the protection of structures in 10% mouse serum to 24 h, compared with the 3-h stability of native structures.

The conditions in which the nanostructures are assembled also affect their biostability. DNA tetrahedra assembled in ethylenediamine buffer exhibited enhanced nuclease resistance compared with those assembled in one of the most commonly used buffers, tris-acetate-EDTA/ Mg^{2+} (TAE/ Mg^{2+} ; where EDTA is ethylenediaminetetraacetic acid)¹¹¹ (FIG. 4d). In a 10-min reaction with DNase I, DNA tetrahedra assembled in TAE buffer completely degraded, whereas almost no digestion occurred in the ethylenediamine buffer. The increased resistance in ethylenediamine is possibly due to the absence of metal ions needed for enzymatic activity. Although this strategy increases the lifetime of DNA nanostructures in the presence of nucleases, further studies are needed to validate the compatibility of this buffer for the assembly of different DNA nanostructures and the toxicity of such organic compounds *in vivo*. For some structures, the inherent assembly

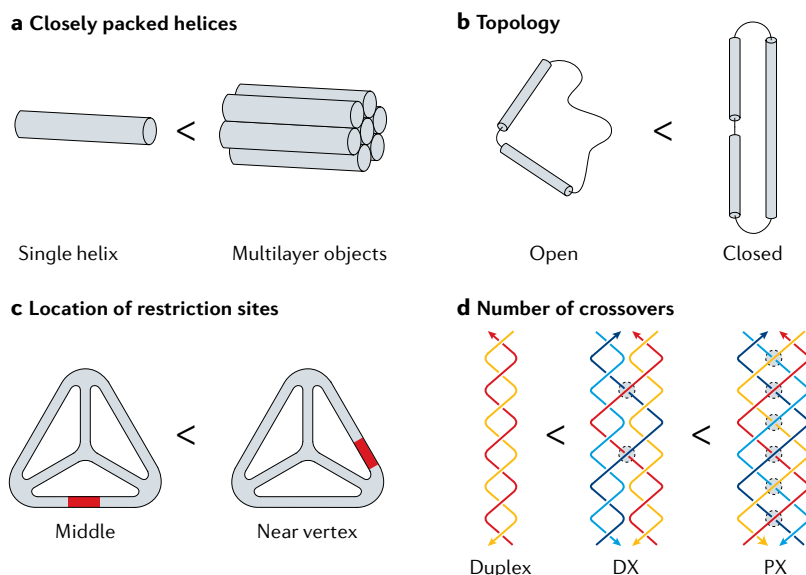


Fig. 3 | Nanostructure designs for enhanced nuclease resistance. **a** | Close-packed helices, as used in DNA origami or DNA bundles¹⁰⁴, are more nuclease resistant than linear duplexes or plasmid DNA. **b** | The biostability of nanostructures is also dependent on the topology, as seen in DNA tweezers¹⁰⁶, for which the closed state is more nuclease resistant than the open state. **c** | Changing the location of enzyme-specific sequences such as restriction sites¹⁰⁵ also influences the nuclease resistance. **d** | DNA motifs with a greater number of crossovers exhibit higher nuclease resistance³². For example, paranemic crossover (PX) DNA is more nuclease resistant than double crossover (DX) DNA.

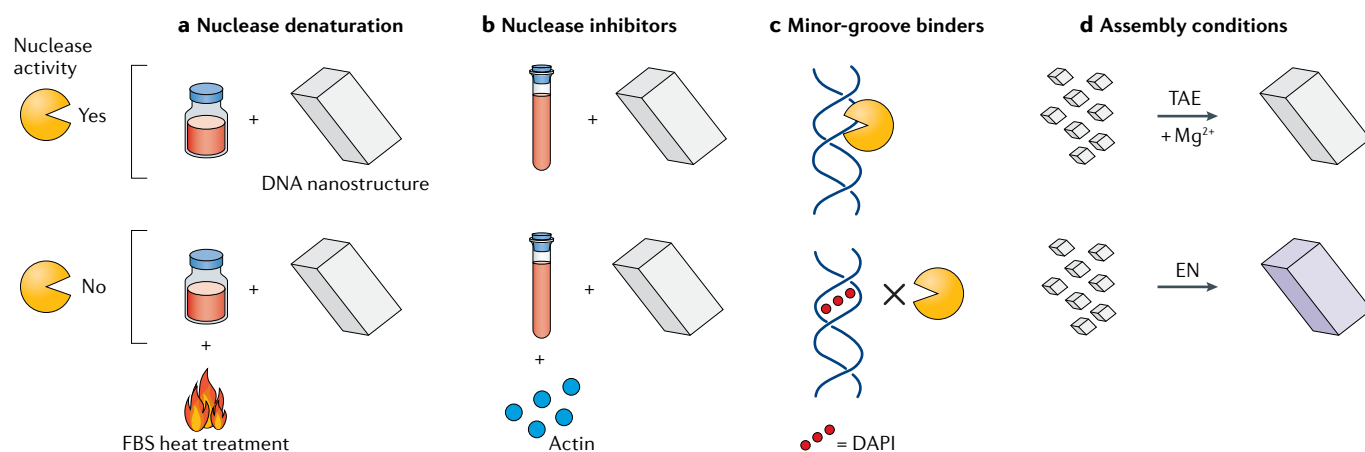


Fig. 4 | **Solution treatment to prevent nuclease degradation.** **a** | Heating serum samples denatures the nucleases, thus rendering them inactive¹⁰⁹. **b** | The addition of nuclease inhibitors, such as actin, to samples reduces nuclease activity, thus protecting DNA nanostructures from degradation¹⁰⁹. **c** | The addition of minor-groove binders such as DAPI (2-(4-amidinophenyl)-1H-indole-6-carboxamide) occludes nuclease binding to DNA nanostructures, thus minimizing degradation¹¹⁰. **d** | The assembly conditions can be varied to increase the biostability. For example, DNA nanostructures were minimally degraded when assembled in ethylenediamine (EN) buffer, but degraded fully when assembled in the commonly used buffer tris-acetate-EDTA (TAE) with Mg^{2+} (REF.¹¹¹). EDTA, ethylenediaminetetraacetic acid; FBS, fetal bovine serum.

units can be provided as fuel in the solution to heal the nanostructures as they are degraded by nucleases, thus reversing the degradation process. This approach was demonstrated using micrometre-scale DNA nanotubes assembled from double crossover (DX) tiles¹¹². The DNA nanotubes degraded within 24 h in 10% FBS-supplemented medium. Upon introducing the monomer DX tiles into the solution, these units incorporated into the damaged regions of the DNA nanotubes and, thus, repaired the nanotubes, prolonging their lifetime to 96 h.

Chemical modifications. Numerous studies have reported chemical modifications to DNA nanostructures to enhance nuclease resistance. These modifications strengthen the weaknesses in the nanostructure design, making DNA nanostructures less susceptible to nuclease digestion. For example, the presence of single-stranded segments and internal nicks in DNA nanostructures make them vulnerable to nuclease attacks¹⁰⁵. Once constructed, internal nicks can be eliminated by covalently linking the strand termini by ligation or crosslinking. Reducing the number of free termini by enzymatic ligation was effective in enhancing the nuclease resistance of DNA tetrahedra¹⁰⁵ and prisms¹¹³, increasing the mean lifetime of these structures to ~42 h and ~200 h, respectively, in 10% FBS (FIG. 5a). Crosslinking the helices within nanostructures by azide-alkyne click reactions has also helped DNA nanostructures to withstand nuclease degradation¹¹⁴ (FIG. 5b). Crosslinked DNA nanotubes had greater protection against exonuclease I compared with native structures. In another example, thymidines were positioned in close proximity within DNA origami nanostructures and crosslinked using ultraviolet light, leading to the site-specific formation of covalent bonds and cyclobutane pyrimidine dimers¹¹⁵ (FIG. 5c). The additional covalent bonds increased the lifetime of

the nanostructures from 10 min (for native structures) to 60 min in the presence of 0.4 U ml^{-1} DNase I.

In addition to post-assembly modifications, the inherent stability of nanostructures can be increased by modifying the DNA backbone and nucleobases of component strands. In one example, DNA strand termini were functionalized with hexaethylene glycol and hexanediol groups, and then used in building DNA prisms^{113,116} (FIG. 5d). These structures had a lifetime of 55–62 h in FBS-supplemented culture medium, whereas the unmodified native DNA nanostructures degraded in less than 18 h. Another route to enhancing nuclease resistance is by using modified DNA as the component strands for assembly. Two-dimensional DNA arrays and a DNA tetrahedron built using L-DNA (that is, regular right-handed DNA) counterparts¹¹⁷ (FIG. 5e). The L-DNA structures remained almost fully intact after treatment with exonuclease I and exonuclease III for 45 min, whereas the native structures fully degraded. In another study, six-arm junctions containing unnatural DNA base pairs (2-thiothymidine:A and 5-methyl-isocytidine:isoG) were only partially digested by T7 exonuclease after 12 h, whereas the unmodified junction completely degraded¹¹⁸ (FIG. 5f). In the context of chemical modifications, a less-explored option is the incorporation of 2'-5'-linked oligonucleotides into nucleic acid nanostructures to enhance resistance against ribonucleases¹¹⁹.

Protective coatings. Modifying the component strands also allows the nanostructures to be coated with a protective layer that resists nuclease activity. In one such strategy, the component strands of DNA cubes were modified with dendritic alkyl chains to form amphiphiles that bind to human serum albumin (HSA)¹²⁰ (FIG. 6a). In stability tests in Dulbecco's modified Eagle

L-DNA

Mirror-image form of naturally occurring DNA, forming a left-handed double helix, in contrast to the right-handed structure of 'regular' D-DNA.

Dendron

Monodisperse, wedge-shaped branched molecule with multiple terminal groups and a single reactive function at the focal point.

medium supplemented with 10% FBS, single-stranded DNA modified with a dendritic alkyl chain and conjugated to HSA had a half-life of 33 min, increasing to 10 h for a DNA cube with four such modified strands conjugated to HSA and to 22 h for a DNA cube with eight HSA-conjugated strands. A protective coating formed from dendritic oligonucleotides that feature at one end a three-pronged group attached to three oligonucleotides has also been demonstrated¹²¹ (FIG. 6b). The stem of the dendritic oligonucleotide can hybridize to single-stranded handles on DNA brick structures, forming a thick DNA coating. DNA bricks with this coating were only 50% digested with 100 U ml⁻¹ of DNase I in 1 h, whereas native structures were fully digested with only 5 U ml⁻¹ of the nuclease. In another example, a DNA octahedron was enveloped by a PEGylated lipid bilayer, which increased the percentage of intact structures from 30% to 85% when incubated in 20 U DNase I for 24 h (REF.¹²²) (FIG. 6c).

The high negative charge of DNA can also be exploited to create a polycationic coating through electrostatic interactions with DNA. For example, cationic polysaccharides such as chitosan and synthetic linear polyethyleneimine were used to coat DNA origami rods, yielding structures that were stable in 10 U ml⁻¹ of DNase I for 24 h (REF.¹²³) (FIG. 6d). Using a similar strategy, DNA origami barrels were coated with oligolysines to increase the lifetime of nanostructures in culture media to 50 min,

compared with 5 min for native structures under the same conditions¹²⁴. Furthermore, conjugating PEG to these oligolysines¹²⁴ (FIG. 6e) or crosslinking the oligolysine-PEG coating using glutaraldehyde¹²⁵ (FIG. 6f) further increased the nuclease resistance of these nanostructures to ~36 h in culture medium and ~66 h in 1 U μ l⁻¹ of DNase I, respectively. PEG-conjugated polylysines have also been used for coating DNA origami structures, forming DNA origami polyplex micelles¹²⁶. The electrostatic interactions between PEG-polylysine and DNA enabled the origami structure to remain intact for 16 h in the presence of DNase I and in FBS-supplemented RPMI medium, whereas native structures completely degraded. This protection strategy was also compatible with functionalized DNA origami structures, causing no loss of functional moieties when applied to six-helix bundles decorated with gold nanoparticles or streptavidin-coated quantum dots.

Peptides and proteins can also be used to coat DNA nanostructures. In one example, a protein corona was formed around DNA origami nanostructures to make them nuclease resistant¹²⁷. Protein-dendron conjugates were created by anchoring bovine serum albumin to a dendron through a cysteine-maleimide bond (FIG. 6g). The dendron part of the conjugate functions as a cationic binding domain that electrostatically attaches to the negatively charged DNA origami surface. Almost 100% of the protected DNA origami structures remained

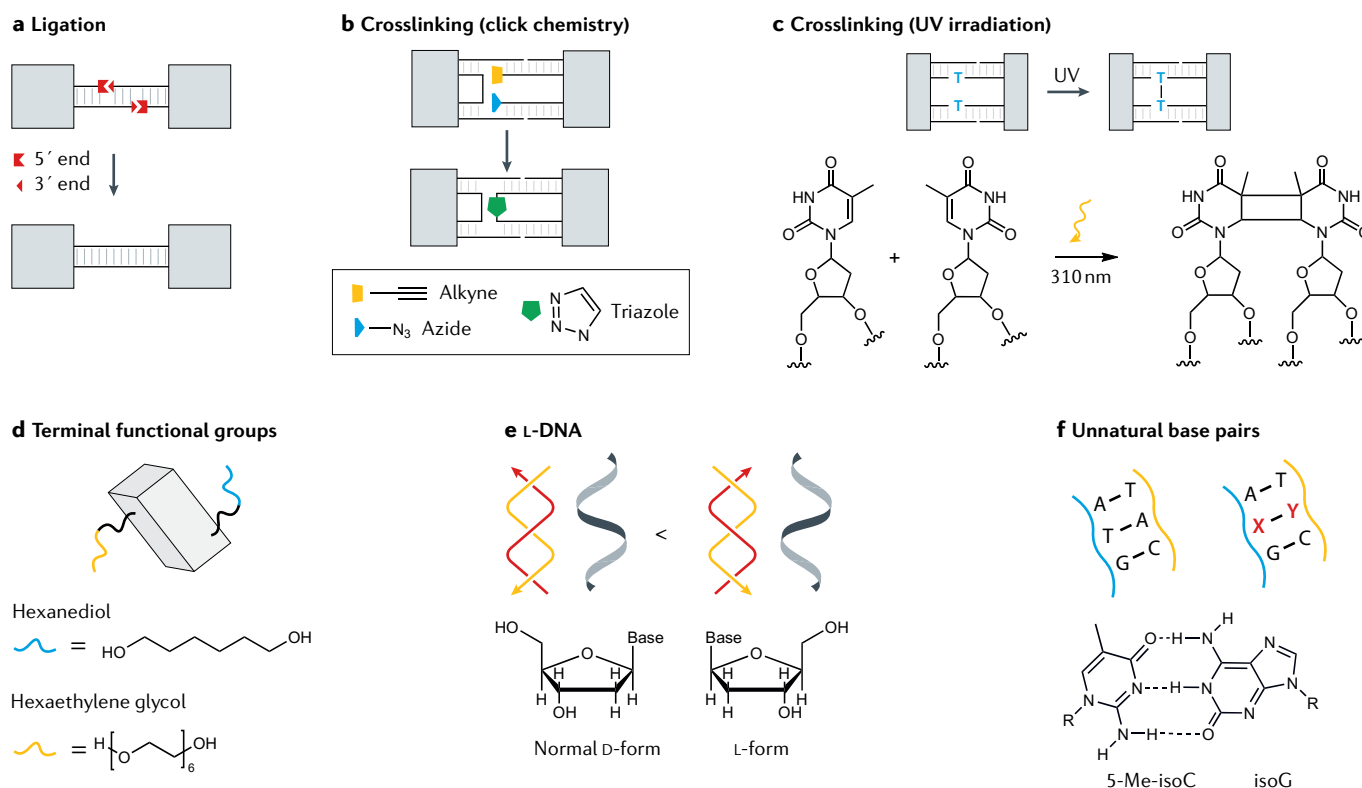
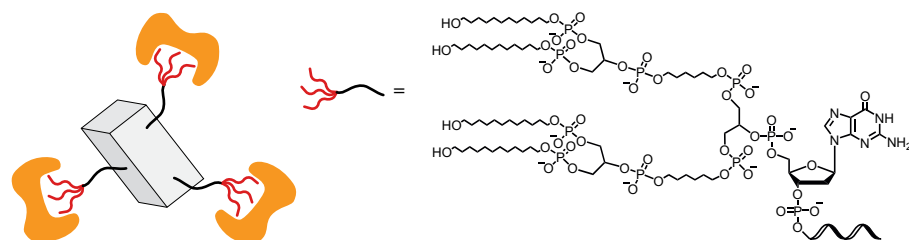
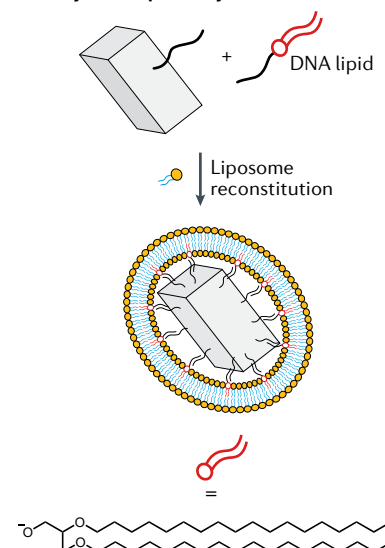


Fig. 5 | Chemical modifications to enhance nuclease resistance. Various chemical modifications have been demonstrated to increase the nuclease resistance of DNA structures. These approaches include ligation of strand termini¹⁰⁵ (part a); crosslinking of component strands by click chemistry¹¹⁴ (part b); crosslinking of strands through the formation of a thymidine dimer under ultraviolet (UV) irradiation¹¹⁵ (part c); the introduction of terminal functional groups, such as hexaethylene glycol and hexanediol¹¹³ (part d); the use of L-DNA¹¹⁷ (part e); and the use of unnatural base pairs (where 5-Me-isoC is 5-methyl-isocytidine; part f)¹¹⁸.

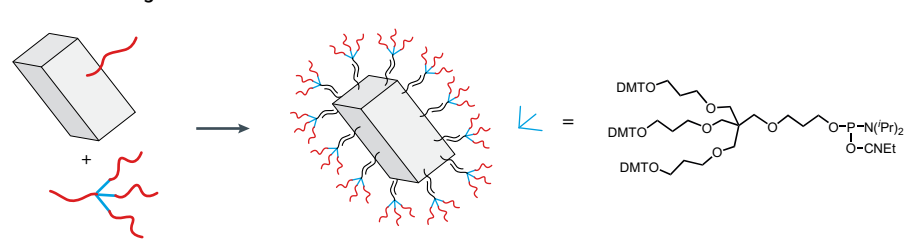
a HSA-DNA dendrite



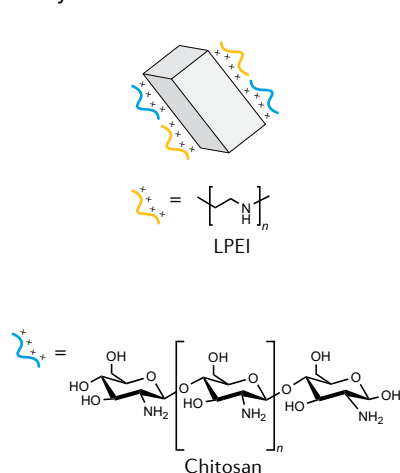
c PEGylated lipid bilayer



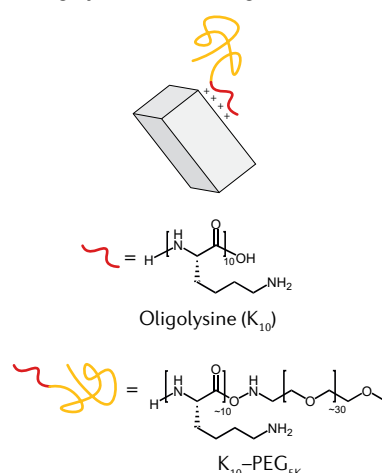
b Dendritic oligonucleotides



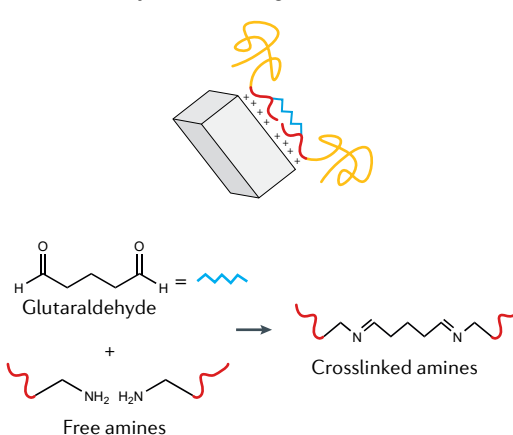
d Polycation shell



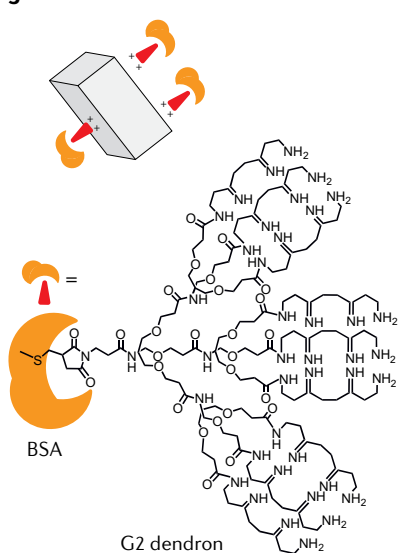
e Oligolysine-PEG coating



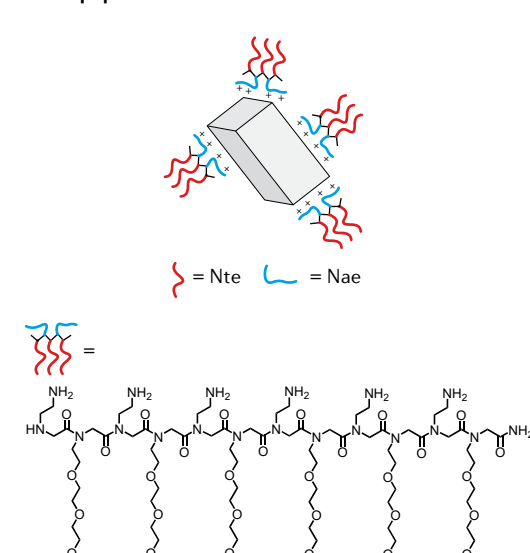
f Glutaraldehyde crosslinking



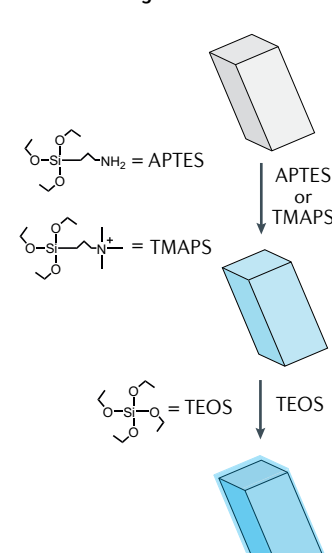
g BSA-dendron



h DNA peptoid



i Silica coating



◀ Fig. 6 | **Protective coatings to enhance nuclease resistance.** Various protective coatings have been developed to increase the nuclease resistance of DNA nanostructures.

These approaches include a coating formed through the complexation of human serum albumin (HSA) and DNA dendrites¹²⁰ (part **a**); coating nanostructures with dendritic DNA through hybridization to DNA handles¹²¹ (part **b**); a PEGylated (where PEG is polyethylene glycol) lipid bilayer protective envelope¹²² (part **c**); the charge-based accumulation of a polycationic shell¹²³ (part **d**); an oligolysine-PEG coating¹²⁴ (part **e**); a coating formed through glutaraldehyde crosslinking of oligolysines¹²⁵ (part **f**); a bovine serum albumin (BSA)-dendron block copolymer coating¹²⁷ (part **g**); a coating comprising DNA peptoids¹²⁸ (part **h**); and silica-based coatings¹³⁰ (part **i**). APTES, (3-aminopropyl)triethoxysilane; DMTO, dimethoxytrityloxy; LPEI, linear polyethyleneimine; Nae, *N*-(2-aminoethyl)glycine; Nte, *N*-(2-(2-methoxyethoxy)ethoxy)ethylglycine; TEOS, tetraethyl orthosilicate; TMAPS, *N*-trimethoxysilylpropyl-*N,N,N*-trimethylammonium chloride.

intact on exposure to 10 U of DNase I for 1 h at room temperature, whereas only 20% of the native structures remained intact. A peptoid-based coating has also been developed to protect DNA origami octahedral structures against nucleases¹²⁸ (FIG. 6h). Compared with peptides, peptoids provide enhanced protection against proteolytic cleavage¹²⁹. The peptoid sequences contained two types of monomer: positively charged *N*-(2-aminoethyl) glycine to facilitate electrostatic complexation with the DNA structure and neutral *N*-(2-(2-methoxyethoxy) ethoxy)ethylglycine for surface passivation. In the presence of DNase I, the bare octahedra completely degraded, whereas peptoid protection preserved the structures. To demonstrate the utility in biomedical applications, bovine serum albumin was encapsulated within the DNA octahedra, which protected the encapsulated protein from hydrolysis by trypsin. In another study, a solution-based method was used to coat 24-helix-bundle DNA origami nanostructures with a thin layer of silica¹³⁰ (FIG. 6i). This coating enabled the DNA origami structures to withstand degradation by DNase I.

Analysis techniques

The predominant techniques used to analyse the nuclease digestion of DNA nanostructures are polyacrylamide or agarose gel electrophoresis (PAGE and AGE, respectively), fluorescence, transmission electron microscopy (TEM) and atomic force microscopy (AFM). All these techniques are commonly used to characterize the formation of DNA nanostructures, but gel electrophoresis is the most often-used technique to quantify the reduction in intact structures on incubation with nucleases or body fluids (FIG. 7a). Typically, nanostructures are incubated for different times with nucleases or serum (or with different concentrations of these), and the incubated samples run on PAGE or AGE. The band corresponding to the structure is then quantified to obtain the level of degradation¹³¹. Fluorescence and Förster resonance energy transfer (FRET) techniques are also often used to quantify degradation of nanostructures^{106,122}. Upon degradation of DNA nanostructures conjugated to fluorescent dyes, the fluorophore and the quencher (or the FRET pair) separate, causing an increase in fluorescence levels (FIG. 7b). This increase can be monitored over time to study the degradation kinetics of DNA nanostructures.

Gel electrophoresis and fluorescence techniques provide information on the degradation levels of nanostructures but do not provide information on structural

changes. To obtain such structural information, TEM and AFM are used to provide before and after snapshots of DNA nanostructures incubated with body fluids or nucleases^{111,132} (FIG. 7c,d). Real-time AFM was recently used to monitor the degradation of different 2D DNA origami shapes when incubated with nucleases¹³³. This technique enables visualization of nanostructures as they degrade and is, thus, useful in determining the parameters for the design of improved DNA-based nanocarriers (for example, in determining whether nucleases affect vertices more than edges).

Overall, gel electrophoresis and fluorescence techniques are the easiest to adapt for most laboratories and provide timed degradation analysis. AFM and TEM require specific instrumentation and training, and are time-consuming, making them the least practical for monitoring the nuclease digestion of DNA nanostructures. The advances in real-time-AFM potentially address the issue of long experimental observation times but is still an expensive instrument not available in most laboratories. There are also challenges associated with the use of gel-based analysis techniques. For example, in the case of cationic block copolymers, polyplexed DNA origami need to be decomplexed (by dextran sulfate, for example) before they can be analysed on a gel (also applicable to TEM)¹²⁶. Moreover, in studies involving FBS or human serum, the serum proteins could cause a background signal in both AGE and TEM, thus necessitating purification of treated samples before analysis.

Mechanism of nuclease resistance

There is still a lack of understanding of the mechanisms of reported nuclease resistance for different structures or protection strategies. Depending on the type of nuclease used, different parameters, such as DNA sequence, backbone geometry, groove width, curvature and flexibility, can all contribute to nuclease resistance¹³⁴. For DNA nanostructures, the non-native geometries and sizes could have a role in inhibiting enzyme binding, thus decreasing the efficiency of specific or non-specific enzymatic cleavage¹⁰⁵. Close-packed helices and frequent crossovers in DNA nanostructures can also occlude enzyme binding, thus negating the effect of nucleases^{92,133}. Furthermore, nucleases such as DNase I cause distortion on binding to helices¹³⁵, and, thus, the increased stiffness of shorter duplex regions or multi-helical edges within DNA nanostructures might prevent the binding of such enzymes. In the case of different modifications and coatings, strong multivalent interactions of the envelope with the DNA could hinder nuclease adsorption and, thus, reduce the degradation. In DNA origami, the scaffold strand already contains restriction sites. In a study on 2D structures, such as a DNA triangle and rectangle, the position of the restriction site post-folding and the introduction of defects (missing staple strands) affected the overall activity of restriction endonuclease on the structures¹³⁶. In a follow-up computational study, global conformational fluctuations between metastable states of DNA origami structures were shown to affect the reactivity of restriction endonucleases on the structures¹³⁷. Simulations of a 2D DNA origami triangle with four or eight staple

Peptoid

Small protein-like polymer in which the side chains are attached to backbone nitrogen atoms, rather than to the α -carbons, as in peptides.

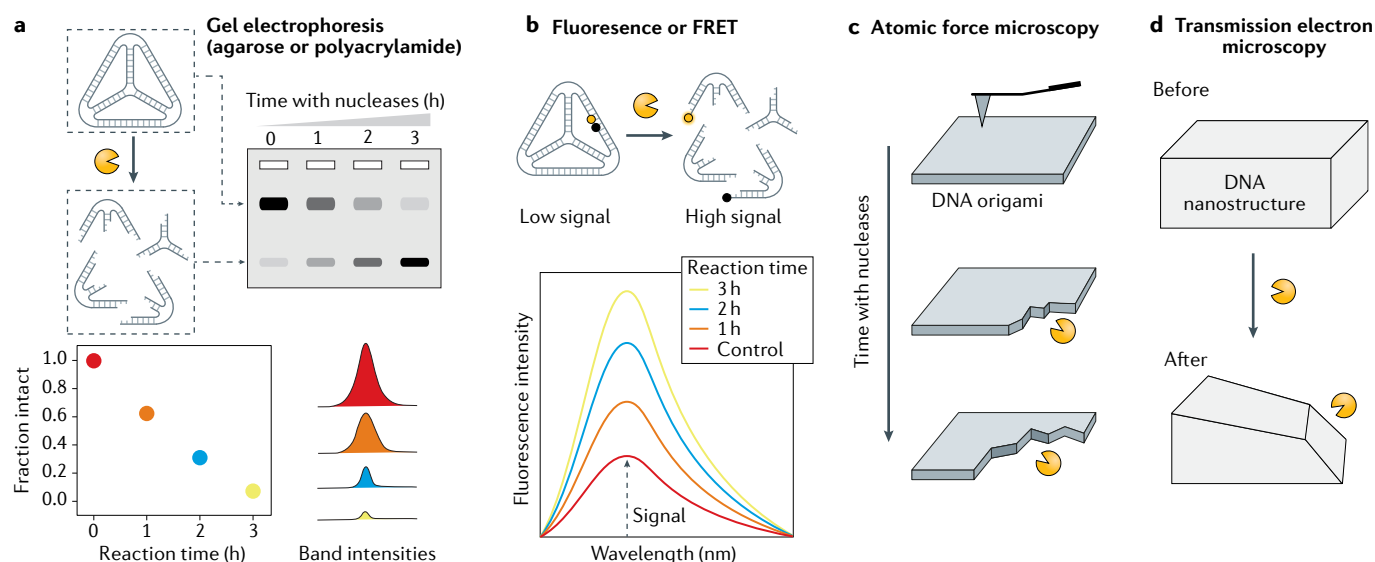


Fig. 7 | Techniques used to analyse nuclease degradation of DNA nanostructures. The main techniques used to analyse DNA nanostructures are polyacrylamide or agarose gel electrophoresis (part **a**), fluorescence or Förster resonance energy transfer (FRET; part **b**), atomic force microscopy (AFM; part **c**) and transmission electron microscopy (TEM; part **d**).

strands omitted from the structure, each resulting in a metastable assembly, revealed that the accessibility of the enzyme to the structure is dependent on the inter-conversion between these metastable states. This analysis showed that the reactivity of restriction enzymes is dependent on the steric overlap between the enzyme and the adjacent helices, thus, providing a route to designing nanostructures with higher or lower nuclease resistance.

A checklist for biostability studies

Design principles. Researchers have designed a large library of DNA nanostructures, each requiring specific assembly conditions. Given this requirement, one has to choose a strategy for modulating the nuclease resistance of a DNA nanostructure on the basis of the design of the underlying DNA assembly. For example, DNA prisms in which the strand termini contained hexaethylene glycol and hexanediol groups showed enhanced nuclease resistance^{113,116}, but the same strategy applied to a larger wireframe DNA origami pentagonal bipyramid did not further stabilize the structure in 10% FBS¹¹⁰. Although there is no general set of rules to make this choice, some studies have noted how the stability numbers reported could apply to similar structures. In DNA origami, most structures share a common scaffold strand (M13) and, thus, have similar GC content if completely double-stranded, eliminating sequence as a factor in stability variations. Therefore, the difference in stability arises from the design of the nanostructures, how closely the helical bundles are packed and the length of the staple strands¹⁰⁴. Even if the stability reported for an origami structure is not representative for objects built with similar specifications, these results could be useful in designing new objects with an expected robustness. In designing other structures, adapting the edges to be multi-helical domains might increase the persistence length of the wireframe structures and, thus, provide additional stability.

Types of nucleases and concentrations. The nuclease of choice in most studies is DNase I. However, some studies have tested the degradation of DNA nanostructures in the presence of other nucleases, showing that the level of degradation is dependent on the exonuclease or endonuclease used^{92,104,118}. In two studies, enzymes such as exonuclease VIII, lambda exonuclease, Mse I restriction endonuclease and T7 exonuclease did not have any effect on DNA origami objects, whereas DNase I, *Escherichia coli* exonuclease I, exonuclease T, T7 endonuclease and exonuclease III degraded them^{104,115}. Similarly, PX DNA showed different levels of resistance to DNase I, exonuclease V, T7 exonuclease and T5 exonuclease; however, PX DNA displayed much higher nuclease resistance than DX or duplex structures in all cases⁹². Thus, although generic trends exist, the activity of different nucleases could differ. The concentration of nucleases tested is also an important factor in determining the extent of degradation. Through calibration of the nuclease levels in FBS by comparing DNA origami nanostructures incubated in 1–20% FBS and different concentrations of DNase I, it was estimated that typical tissue culture conditions may contain between 256 and 1,024 U l⁻¹ equivalent of DNase I activity¹⁰⁹. These studies indicate that it is important to know the specific levels of nucleases in different biofluids and to test relevant amounts of nucleases in such biostability studies.

Reagents. Another factor to consider when testing the stability of DNA nanostructures is the type of serum used and the freeze–thaw protocols. The level of nuclease activity in different FBS lots and frozen aliquots has been observed to vary¹⁰⁹. In this report, the nuclease activity was highest after initial thawing of the FBS and was lost after a few weeks when FBS-supplemented medium was stored at 4 °C (REF.¹⁰⁹). Thus, it is important to follow the specific reagent-handling protocol across multiple experiments to validate the nuclease-resistance values

of different structures. Nuclease activity in body fluids also varies widely between species^{138,139}, and studies on DNA nanostructures have also found the stability of nanostructures to vary in sera from different species¹¹⁰. Future studies could work with human-derived solutions instead of animal sera to make the results more relevant for human applications.

Choice of protection strategy. The type of application will determine the level of biostability needed and which strategy to use for modulating nuclease resistance. In drug delivery, partial digestion of DNA nanostructures could trigger release of the drug cargo or attached functional moieties (such as fluorophores for tracking). However, slow or delayed degradation of nanostructures could be useful in the spontaneous release of drugs. The addition of nuclease inhibitors or the heat treatment of samples are potential options for biosensing applications in which the sample can be pre-processed before addition of the DNA sensor. However, use of DNA nanostructures in vivo requires strategies that obviate the need to alter the environment, as the addition of external factors might influence other biomolecular processes. Furthermore, the choice of strategy will also depend on the type of biofluid the structures will be in (blood, urine, saliva), and prior knowledge of the types of nucleases present in these fluids and their levels will be useful in biostability studies. Once the stabilization strategy is chosen, it is imperative to test the functionality of these structures after the protection process; chemical modifications or coatings should not affect the binding of sensing elements or targeting moieties to DNA nanostructures. For example, the peptoid coating of DNA origami structures did not affect the encapsulation of cargos such as proteins and nanoparticles within the nanostructure, suggesting potential use in drug delivery¹²⁸. In addition, polyplex micelles comprising cationic polysaccharides have been successfully used to stabilize plasmid DNA for gene therapy¹⁴⁰, indicating the potential use of similarly stabilized DNA nanostructures.

In vivo stability and immune response. The integrity of DNA nanostructures in vivo affects the immune response in cells or animals. When used as drug carriers, the immune response elicited by intact and degraded nanostructures might differ in some cases, whereas in others, it might be dependent on the total mass of DNA and not the design or integrity of the nanostructures¹⁰⁹. Thus, care has to be taken to test the intactness of DNA nanostructures in studies in which the specific immune

response is important. For example, the oligolysine-PEG block copolymer coating of DNA nanostructures had negligible effect on cell viability or enzyme kinetics, indicating minimal immune response in the cells^{125,141}. Studies to test in vivo stability and immune response might also require additional functionalities to track the nanostructure through the body or a cellular pathway, and can use newly developed techniques, such as a 'hydroporator', to deliver DNA nanostructures directly into cells for monitoring¹⁴².

Conclusion

As DNA nanotechnology moves towards real-life applications, enhancing the stability of nanostructures in biological environments is of increasing importance. The strategies discussed in this Review provide an overview of methods to modulate the nuclease resistance of DNA nanostructures, rendering them more useful in biological applications. There is already a large library of DNA nanostructures available, but future studies will need to focus on upgrading these structures to be application-ready.

The studies discussed herein have used different structures as examples in different environments (different culture media, with or without serum supplement, or in a combination of nucleases). From these studies, it is apparent that DNA nanostructures possess higher nuclease resistance than duplexes, and that all the strategies discussed can enhance this resistance. However, it is difficult to further compare results from these studies, as they report results using different metrics (lifetime, half-life, percentage intact) and use different nanostructures. To facilitate comparison, the DNA nanotechnology research community is encouraged to decide on a set of standardized reporting parameters for nuclease resistance and biostability. Such a system would benefit the generalization of design parameters and the choice of strategy for a specific structure and/or application. Future studies could also explore the effect of the shape and size of DNA nanostructures on nuclease resistance, reports on which are few.

DNA nanostructures have permeated many different fields and are closer to real-life applications than ever before. The ongoing development of stabilization strategies, construction methods, characterization tools and platforms that allow non-experts to design DNA nanostructures¹⁴³ is contributing to the rapid progress towards achieving the potential that DNA nanotechnology promises.

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