

DNA Circuits

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Deoxyribonucleic acid (DNA), a genetic material, encodes all living information and living characteristics, e.g., in cell, DNA signaling circuits control the transcription activities of specific genes. In recent years, various DNA circuits have been developed to implement a wide range of signaling and for regulating gene network functions. In particular, a synthetic DNA circuit, with a programmable design and easy construction, has become a crucial method through which to simulate and regulate DNA signaling networks. Importantly, the construction of a hierarchical DNA circuit provides a useful tool for regulating gene networks and for processing molecular information. Moreover, via their robust and modular properties, DNA circuits can amplify weak signals and establish programmable cascade systems, which are particularly suitable for the applications of biosensing and detecting. Furthermore, a biological enzyme can also be used to provide diverse circuit regulation elements.

synthetic DNA circuit

DNA strand displacement

DNA self-assembly

DNA networks

1. Introduction

Recently, various artificial DNA circuits have been established and widely applied to many fields such as medical diagnosis [1][2][3], molecular detection, and information processing [4][5][6][7][8][9][10]. Particularly, synthetic DNA circuits, designed and constructed in vitro, perform an important role in effectively controlling the gene networks in cell [11][12][13]. Synthetic DNA circuits have been demonstrated as possessing superiority in simulating and regulating DNA signaling, due to the properties of programmability and easy operation [14][15][16][17][18]. More importantly, synthetic DNA circuits have the potential to promote complex biological information processes and provide a new way to achieve gene analysis and molecular information processing [19][20][21].

Using predesigned specific base pair recognition, synthetic DNA circuits can modulate complex gene networks to implement diverse biofunctions. Recently, a variety of bioengineering and biocomputing functions have been regulated by varying the architectures and integrations of DNA circuits, such as their signal simulation [5][22], the molecular switch, catalytic cycle, cascade amplification [23][24][25][26][27][28][29][30], and logic gates [31][32][33][34][35][36][37]. In fact, most of the DNA circuits are implemented and regulated for the DNA strand displacement, whereby the longer DNA strand is able to hybridize with the complementary strand to displace the shorter one [38][39][40]. Through a DNA strand displacement reaction (SDR), a synthetic DNA circuit can be used to precisely regulate complex gene networks and molecular biosystems, e.g., DNA neural network systems that are constructed to implement pattern recognition [41].

2. The Research Studies on DNA Circuits

2.1. Cascading DNA Circuits In Vitro

The cascade effect is a signal transmission phenomenon, moving from the upstream to the downstream unit within an integrated system [42]. In a cascading DNA circuit, the DNA signal can be transmitted to a downstream reaction along a long-range signaling path. During the cascading reaction process, the free energy of the whole system gradually reduces toward a more stable energy state [43]. As a result, a cascading DNA circuit is a precisely designable information processing system that is able to produce the expected signal. Therefore, cascading DNA circuits are usually used to build continuous multiple-layer reaction systems [44], with the capacity to transmit molecular information and perform a computing function.

Interestingly, one of the typical applications of a cascading DNA circuit is to construct a DNA computing system with continuous passing information. Imitating the modularization of electronic circuits, Seelig et al. [45] designed a cascading DNA circuit using ssDNA as the input/output signal through DNA strand displacement. The process is illustrated in **Figure 1a**.

Recently, Qian et al. [46] designed a molecule calculator with the function of computing the square root of binary data by designing a DNA cascade circuit based on the basic neural network model (**Figure 1b**). Afterwards, based on the principle of the construction of another Neuronal connection pattern, Cherry and Qian [41] designed a winner-take-all neural network realizing the digital pattern recognition function using a DNA circuit (**Figure 1c**).

Since the DNA neural network circuit has been demonstrated to perform complex logic operations at the gene manipulation level, the DNA circuit designed according to its model is also designed for a rapid and accurate diagnosis of cancer [47]. **Figure 1d** illustrates a DNA neural network circuit that uses miRNA in clinical serum samples to detect cancer. In this system, different kinds of miRNA from the clinical serum samples are first identified as input signals by the DNA neural network circuit, and are then sequentially implemented by DNA multiplication, summation and subtraction to achieve a rapid and accurate cancer diagnosis.

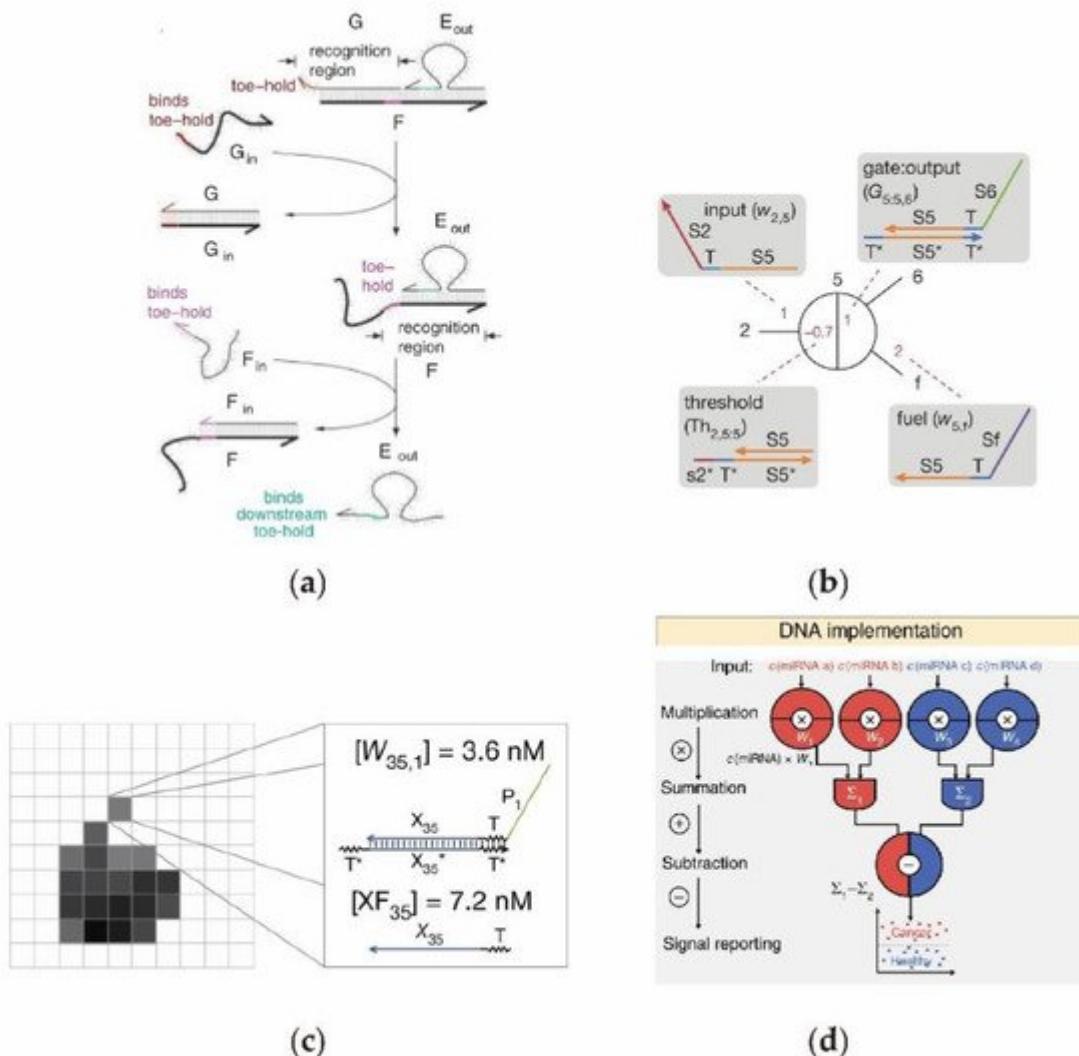


Figure 1. Cascade circuit to perform neural networks. **(a)** Schematic diagram of the DNA cascade circuit based on creating toehold sites. **(b)** Structure of DNA circuit neurons with weights and thresholds comprised of DNA strands, where multi-input and multi-output functions are performed. **(c)** Schematic diagram of the DNA neural network circuit to implement handwritten character recognition system. **(d)** The neural network computing model designed using the DNA cascade circuit to classify and identify Cancer.

2.2. Enzyme-Free Catalytic DNA Circuit

In natural gene networks, protein enzymes are typically used as catalysts, while DNA seldom performs a catalyzing function. This means that the input DNA signals are continuously consumed in a normal noncatalytic DNA circuit reaction, thus leading to a limited reaction efficiency. In synthetic DNA circuit, by introducing a recyclable DNA as a catalyst the reaction efficiency is greatly improved, and only a small amount of input DNA is required to release a large amount of output DNA [48]. Therefore, a catalytic DNA circuit can be used to construct a more efficient artificial nucleic acid system without the use of a bioenzyme.

In 2006, Seelig et al. [49] first proposed a DNA circuit with high catalytic efficiency based on DNA strand replacement by establishing a catalytic cycling unit. **Figure 2a** illustrates the operation of this enzyme-free catalytic

circuit based on the principle of DNA displacement.

Since the linear DNA structure is more stable and easily transformed, Zhang et al. [50] designed an enzyme-free DNA circuit with a high catalytic performance, using single-stranded DNA as a catalyst. As shown in **Figure 2b**, the DNA catalyst was used repeatedly in the catalytic cycle without being consumed, and a large amount of outputDNA was produced along with the generation of waste and intermediate products.

Zhang et al. [23] introduced an inhibitor DNA and activator DNA into the catalytic system. As shown in **Figure 2c**, in contrast to a simple linear ssDNA catalyst, a loop DNA catalyst can be used to perform the catalytic function, where the catalytic process can be dynamically switched according to the specific inhibitors and activators. By combining the enzyme-free catalytic circuit with DNA tile technology, David Yu Zhang et.al realized the rapid and large-scale assembly of micro-nanotubes [51]. The linear catalytic circuit in **Figure 2d** continuously generates the single-stranded DNA required to assemble large-scale nanotubes, thereby achieving the function of efficiently increasing the size of nanotubes within a short period.

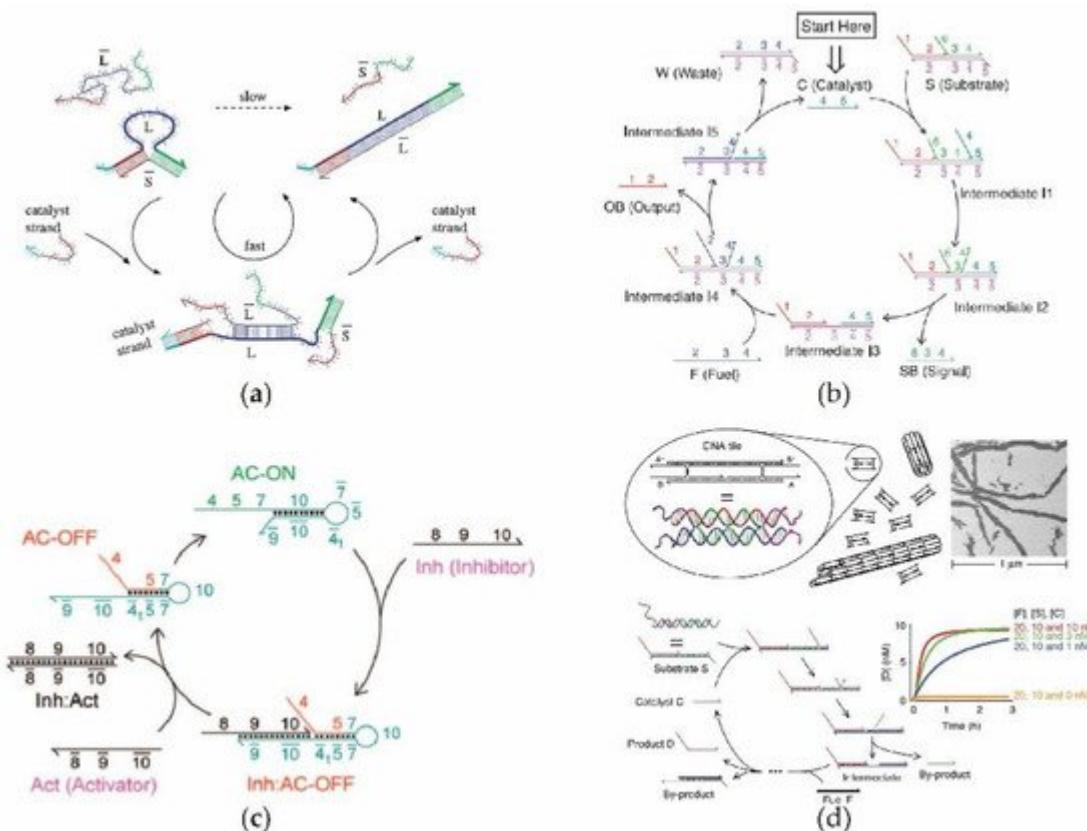


Figure 2. Schematic diagram of catalytic DNA circuit **(a)** The DNA catalytic circuit in which the catalyst can be used repeatedly in each reaction cycle. **(b)** Catalytic DNA circuit using single-stranded DNA as a catalyst, where DNA intermediates, output products, and signals are produced during each cycle with sufficient fuel DNA. **(c)** Catalytic DNA circuit using neck ring DNA as catalyst. During the catalytic process, the structures of neck ring DNA are dynamically changed. **(d)** Large-scale assembly of DNA tile using DNA enzyme-free catalytic circuit.

2.3. DNAzyme-Based DNA Circuit

DNAzyme is a specific single-stranded DNA fragment with a catalytic cutting function, high catalytic activity, and specific recognition capacity. One of its most important roles is to cleave the RNA site via esterification. By taking advantage of this property, the DNA circuit can directly employ DNAzyme to accurately control its circuit structure and signal transmission mode [52][53][54]. On the other hand, the structure and activity of DNAzyme can be precisely regulated, thus providing the stronger controllability for DNA circuits [27][55][56][57][58][59][60].

Utilizing a metal-ion ion dependent DNAzyme, Moshe et al. [61] constructed a DNA circuit that was regulated by uranium dioxide and magnesium. As shown in **Figure 3a**, only once both DNAzymes were activated simultaneously could the target DNA sequence be released.

Moreover, by combining gold nanoparticles with DNAzyme, Bi et al. [62] designed a DNAzyme logic circuit characterized by colorimetric and UV/vis detection. As illustrated in **Figure 3b**, circle DNA with different RNA modification sites can be cut into several DNA fragments by the DNAzyme, resulting in the aggregation of gold nanoparticles.

The potential of the upstream output signal being successfully transmitted to the downstream as an input signal without a signal leakage in the process of transmission remains a major challenge in DNAzyme cascade circuits. Unlike the traditional DNAzyme with fixed structures, Elbaz et al. [63] constructed a DNA circuit with a controllable DNAzyme by splicing the structure into a hybrid structure (**Figure 3c**).

The signal catalytic function of DNAzyme provides the advantage reusability in the process of RNA cleavage, however, it also causes the DNAzyme-based circuit to become uncontrollable, which greatly limits the time domain characteristics of the circuits. Compared with the previous unchangeable DNAzyme, Harding et al. [64] designed a DNA circuit using the DNAzyme with a dynamic switching ability (**Figure 3d**). Therefore, the activity of the DNAzyme can be well regulated using an additional input DNA signal.

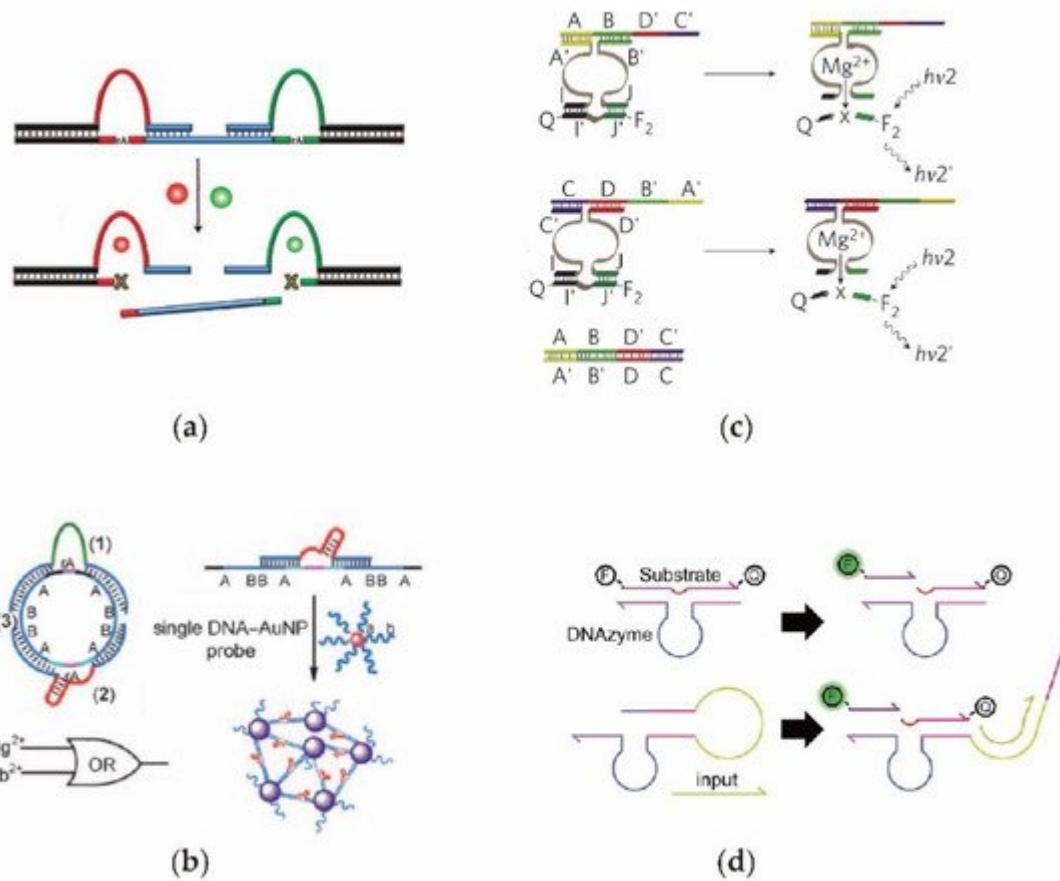


Figure 3. Schematic diagrams of DNAzyme-based synthetic DNA circuit. **(a)** DNAzyme DNA circuit able to recognize metal ions. **(b)** DNAzyme DNA circuit to realize controllable self-assembly of nanoparticles to construct logic gates. **(c)** Editable DNA circuit constructed by the combinations of variable DNA scaffolds. **(d)** Controllable DNAzyme switches with repeated open/off control.

2.4. Protein Enzyme-Assisted DNA Circuit

A biological enzyme is a type of biomolecule that can specifically recognize the substrate to perform its catalytic function [65]. According to their catalytic reaction properties, biological enzymes can be generally divided into categories such as oxidoreductase, transferases, and hydrolases. Due to its high specificity, strong catalytic ability, and fast reaction speed, the enzyme-assisted DNA circuit is suitable for building a more intelligent biological computing system. Recently, biological enzymes have been used to build functional DNA circuits [66][67][68].

Through their simulation of ecosystems, Fuji et al. [5] designed an enzyme-assisted DNA circuit with oscillating and competitive functions (Figure 4a). In the system, both the predator P and prey N were dynamically generated and degraded in the presence of the polymerase, nicking enzyme, and exonuclease.

In 2017, inspired by signaling networks in living cells, Lenny et al. [69] designed an enzyme-driven DNA circuit that can dynamically regulate the signal strength during a reaction (Figure 4b). The researchers applied this toolbox to various DNA circuits to realize the bionic electronic functions by using DNA as the information carrier.

Song et al. [70] established a logic DNA circuit composed of simple ssDNA and polymerase-triggered DNA strand displacement (Figure 4c). Significantly, since the DNA circuit consists of only single-stranded DNA, the problems of signal leakage and signal reset are resolved well. Through the combination of the enzyme- and entropy-driven DNA catalytic reaction, Zhang et al. [71] constructed a dual-catalytic recyclable DNA circuit (Figure 4d).

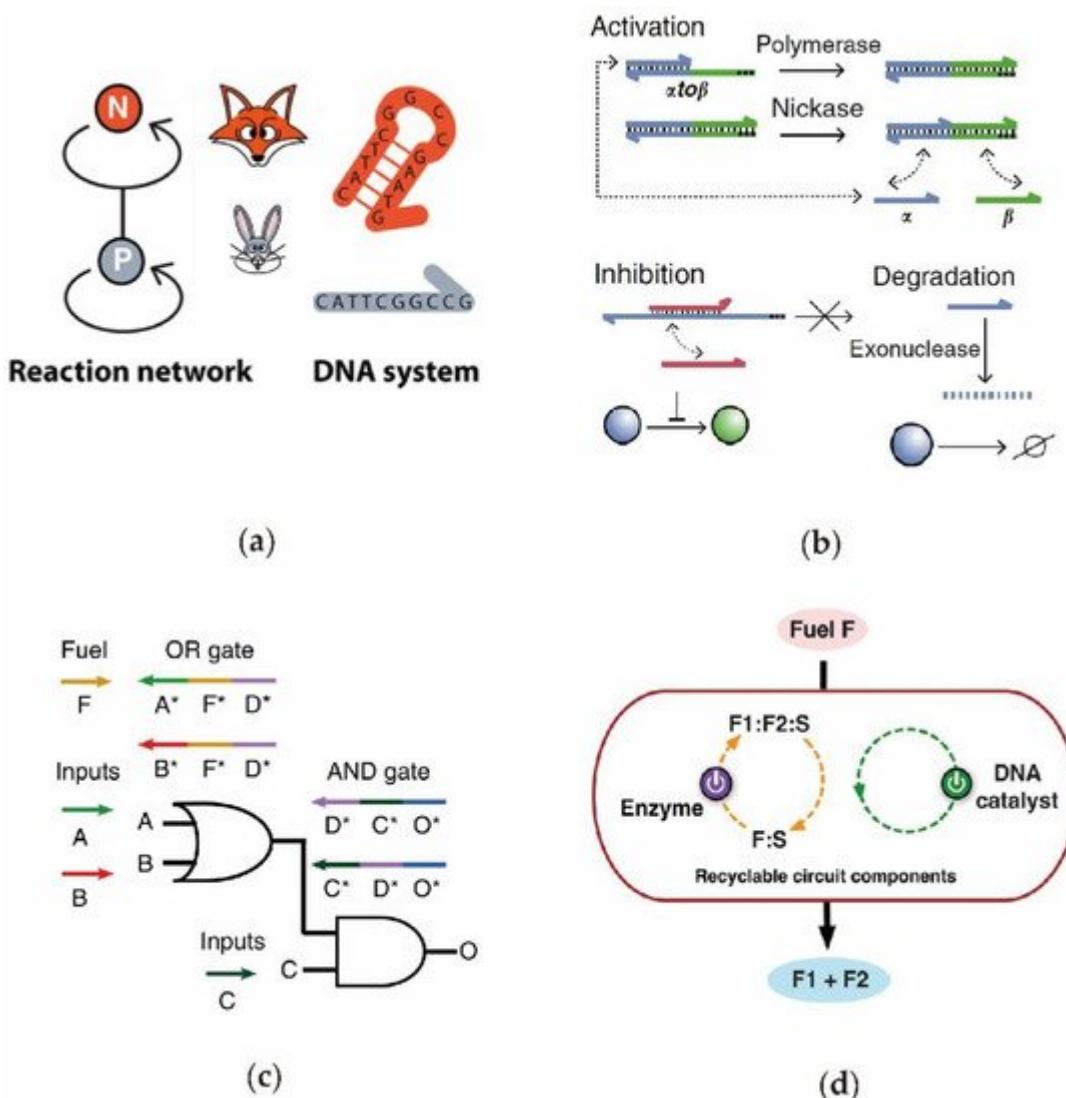


Figure 4. Designs of biological enzyme-involved DNA circuits. (a) Schematic diagram of the DNA logic gates using strand displacement polymerase. (b) Highly efficient dual-catalytic recyclable DNA circuit driven by nicking enzyme and DNA catalyst. (c) Schematics for the oscillating DNA circuit based on protein enzyme. (d) The enzyme-toolbox-directed DNA circuit with activation, inhibition, and subtraction functions.

2.5. The DNA Circuits on Origami Surface

A DNA origami assembly was developed by Rothemund in 2006, in which hundreds of short DNA fold with a M13mp18 scaffold to construct various nanometer shapes, such as squares, circles, and triangles. As the surface of DNA origami can modify DNA in a practicable way, DNA circuits can be constructed onto the origami surface to achieve vector-based logic operations and robot movements [72][73][74][75][76][77]. Such an on-surface DNA circuit

operation is a new method of gene regulation, exhibiting the properties of specific spatial designs and modular constructions.

As shown in **Figure 5a**, by utilizing synthetic molecular motors that move autonomously along liner tracks, Shelley et al. [78] designed a programmed network system on the surface of a rectangular DNA origami. After moving, the majority of motors can follow the correct path to the predesigned target aim. This programmable on-surface DNA circuit motor has the potential to build complex and controllable information transmission systems. In 2014, to investigate the kinetic and efficiency of on-surface DNA circuit, Teichmann et al. [79] installed DNA circuit units of signal sender and receiver on an origami surface (**Figure 5b**). In the on-surface circuit, the sender unit was situated in the middle (red color point) of the origami platform and the receiver units were distributed evenly on the plane around the sender. Interestingly, researchers can regulate the signal transmission by adjusting the distributions of gate units, so as to optimize the efficiency and accuracy of the on-surface DNA circuits. In addition, the precise arrangement of on-surface circuit is of great significance for DNA-based memories and information processing. In 2017, Winfree developed a DNA robot with the ability to carry cargo in sequence on the origami surface by designing a randomly moving DNA circuit [80] (**Figure 5c**). Gourab et al. [42] constructed a surface logic DNA circuit whereby the signal transmission path can be programmed to realize the directional logic operation as shown in **Figure 5d**.

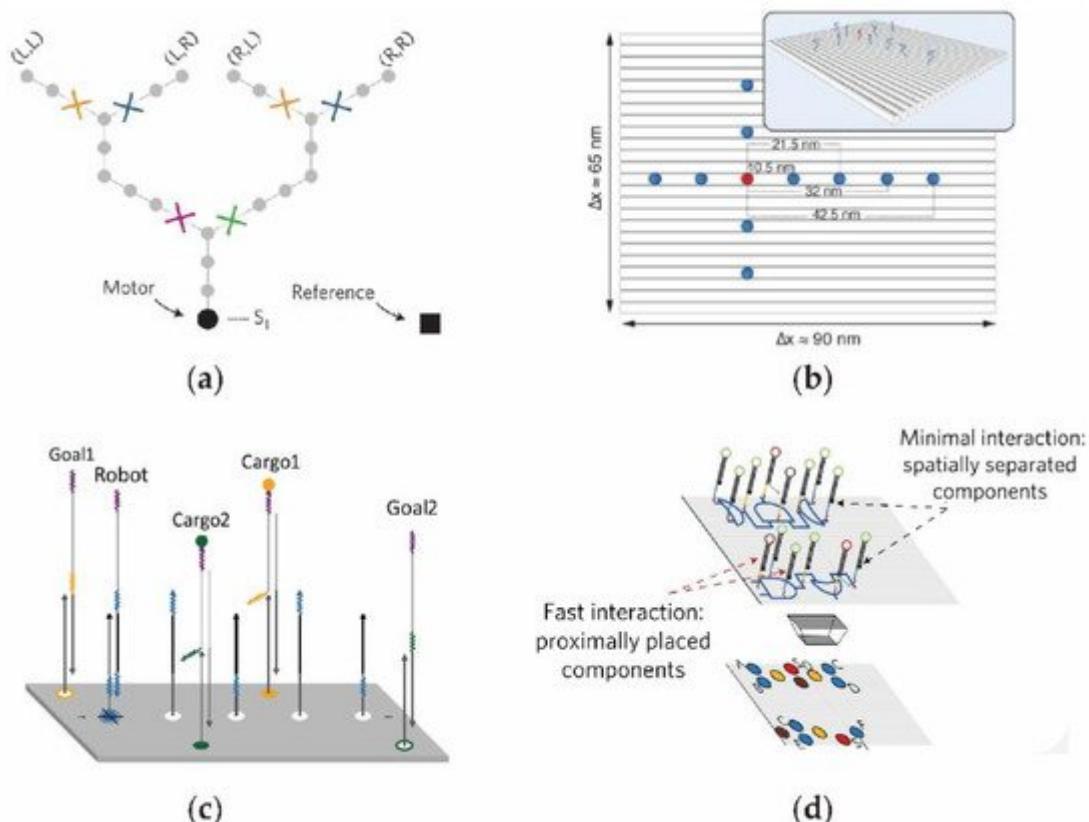


Figure 5. Design of DNA circuits on origami surface. (a) DNA walking motor that can follow the path on the DNA origami surface. (b) On-surface DNA circuit that can establish signal transmissions with addressable positions. (c) DNA transport cargos on origami surface. (d) On-surface DNA logic gates.

2.6. The DNA Circuits Combined with Nanoparticles

Compared to other macroscopic materials, nanoparticles possess the advantages of well-controlled sizes and shapes [81][82][83], visible spectroscopy characteristics [84][85][86][87], and multiple chemical modifications. Therefore, nanoparticles have been widely used in multiple disciplines, e.g., in medicine, bioengineering, and molecular signal detection [88][89][90][91][92]. Recently, combined with a DNA circuit, nanoparticles have been widely applied to perform the functions of biosensing, molecular signal detection, and DNA computing [93][94][95][96].

3. Summary

Although DNA circuits have engendered remarkable progresses in recent years, there are still some critical aspects that need to be improved upon to broaden the related application fields. The most important aspect is the signal leakage and distortion that exists in various complex DNA circuits [41][46]. The primary reason for this phenomenon is that some redundant base complementary pairings easily emerged during the circuit operations. At present, the design methods implemented to program DNA sequences are mainly based on the combinations of a manual design and simple software. Therefore, it is still difficult to completely eliminate these interference signals using pre-established methods, and a more automatic and universal sequence design platform is eagerly desired in the future. In addition, another important consideration to account for is the method by which to ensure the DNA circuit is reusable, thereby promoting its practical applications. Compared with the repeated electronic circuits, DNA circuits cannot be reused after completing the first complete operation [70][47][97], which inhibits the subsequent development of more advanced intelligent DNA circuits. Specifically, in large computing DNA circuits, it is important to establish a reusable circuit mechanism, to meet requirements of frequently adjusted circuit parameters and to test the structural stability of the system. In fact, many studies have been undertaken to improve the repeatability of the DNA circuits. More recently, Zhang et al. reported a nicking-assisted recycling strategy for implement reusable DNA circuits to achieve a repeatability of over 20 times during one reaction cycle [71].

References

1. Du, Y.C.; Cui, Y.X.; Li, X.Y.; Sun, G.Y.; Zhang, Y.P.; Tang, A.N.; Kim, K.; Kong, D.M. Terminal Deoxynucleotidyl Transferase and T7 Exonuclease-Aided Amplification Strategy for Ultrasensitive Detection of Uracil-DNA Glycosylase. *Anal. Chem.* 2018, 90, 8629–8634.
2. Shen, H.; Wang, J.; Liu, H.; Li, Z.; Jiang, F.; Wang, F.B.; Yuan, Q. Rapid and Selective Detection of Pathogenic Bacteria in Bloodstream Infections with Aptamer-Based Recognition. *ACS Appl. Mater. Interfaces* 2016, 8, 19371–19378.
3. Zhang, Q.; Jiang, Q.; Li, N.; Dai, L.; Liu, Q.; Song, L.; Wang, J.; Li, Y.; Tian, J.; Ding, B.; et al. DNA Origami as an In Vivo Drug Delivery Vehicle for Cancer Therapy. *ACS Nano* 2014, 8, 6633–6643.
4. Organick, L.; Ang, S.D.; Chen, Y.J.; Lopez, R.; Yekhanin, S.; Makarychev, K.; Racz, M.Z.; Kamath, G.; Gopalan, P.; Nguyen, B.; et al. Random access in large-scale DNA data storage. *Nat.*

Biotechnol. 2018, 36, 242–248.

5. Fujii, T.; Rondelez, Y. Predator-Prey Molecular Ecosystems. *Acs Nano* 2013, 7, 27–34.

6. Guo, M.Y.; Chang, W.L.; Ho, M.C.; Lu, J.; Cao, J.N. Is optimal solution of every NP-complete or NP-hard problem determined from its characteristic for DNA-based computing. *Biosystems* 2005, 80, 71–82.

7. Zhou, C.; Duan, X.Y.; Liu, N. A plasmonic nanorod that walks on DNA origami. *Nat. Commun.* 2015, 6, 6.

8. Kim, J.; White, K.S.; Winfree, E. Construction of an in vitro bistable circuit from synthetic transcriptional switches. *Mol. Syst. Biol.* 2006, 2, 68.

9. Xie, Z.; Liu, S.J.; Bleris, L.; Benenson, Y. Logic integration of mRNA signals by an RNAi-based molecular computer. *Nucleic Acids Res.* 2010, 38, 2692–2701.

10. Wang, K.; Ren, J.; Fan, D.; Liu, Y.; Wang, E. Integration of graphene oxide and DNA as a universal platform for multiple arithmetic logic units. *Chem. Commun.* 2014, 50, 14390–14393.

11. Zhu, J.B.; Zhang, L.B.; Dong, S.J.; Wang, E.K. Four-Way Junction-Driven DNA Strand Displacement and Its Application in Building Majority Logic Circuit. *Acs Nano* 2013, 7, 10211–10217.

12. Fern, J.; Schulman, R. Design and Characterization of DNA Strand-Displacement Circuits in Serum-Supplemented Cell Medium. *ACS Synth. Biol.* 2017, 6, 1774–1783.

13. Wei, Q.; Huang, J.; Li, J.; Wang, J.; Yang, X.; Liu, J.; Wang, K. A DNA nanowire based localized catalytic hairpin assembly reaction for microRNA imaging in live cells. *Chem. Sci.* 2018, 9, 7802–7808.

14. Prokup, A.; Hemphill, J.; Deiters, A. DNA Computation: A Photochemically Controlled AND Gate. *J. Am. Chem. Soc.* 2012, 134, 3810–3815.

15. Green, A.A.; Kim, J.; Ma, D.; Silver, P.A.; Collins, J.J.; Yin, P. Complex cellular logic computation using ribocomputing devices. *Nature* 2017, 548, 117–121.

16. Rinaudo, K.; Bleris, L.; Maddamsetti, R.; Subramanian, S.; Weiss, R.; Benenson, Y. A universal RNAi-based logic evaluator that operates in mammalian cells. *Nat. Biotechnol.* 2007, 25, 795–801.

17. Hemphill, J.; Deiters, A. DNA computation in mammalian cells: microRNA logic operations. *J. Am. Chem. Soc.* 2013, 135, 10512–10518.

18. Zong, Y.; Zhang, H.M.; Lyu, C.; Ji, X.; Hou, J.; Guo, X.; Ouyang, Q.; Lou, C. Insulated transcriptional elements enable precise design of genetic circuits. *Nat. Commun.* 2017, 8, 52.

19. Phillips, A.; Cardelli, L. A programming language for composable DNA circuits. *J. R. Soc. Interface* 2009, 6 (Suppl. S4), 18.
20. Chen, K.; Kong, J.; Zhu, J.; Ermann, N.; Predki, P.; Keyser, U.F. Digital Data Storage Using DNA Nanostructures and Solid-State Nanopores. *Nano Lett.* 2019, 19, 1210–1215.
21. Lopez, R.; Chen, Y.J.; Dumas Ang, S.; Yekhanin, S.; Makarychev, K.; Racz, M.Z.; Seelig, G.; Strauss, K.; Ceze, L. DNA assembly for nanopore data storage readout. *Nat. Commun.* 2019, 10, 2933.
22. Fern, J.; Scalise, D.; Cangialosi, A.; Howie, D.; Potters, L.; Schulman, R. DNA Strand-Displacement Timer Circuits. *ACS Synth. Biol.* 2017, 6, 190–193.
23. Zhang, D.Y.; Winfree, E. Dynamic allosteric control of noncovalent DNA catalysis reactions. *J. Am. Chem. Soc.* 2008, 130, 13921–13926.
24. Graugnard, E.; Kellis, D.L.; Bui, H.; Barnes, S.; Kuang, W.; Lee, J.; Hughes, W.L.; Knowlton, W.B.; Yurke, B. DNA-Controlled Excitonic Switches. *Nano Lett.* 2012, 12, 2117–2122.
25. Chen, X.; Briggs, N.; McLain, J.R.; Ellington, A.D. Stacking nonenzymatic circuits for high signal gain. *Proc. Natl. Acad. Sci. USA* 2013, 110, 5386–5391.
26. Wang, F.; Elbaz, J.; Willner, I. Enzyme-free amplified detection of DNA by an autonomous ligation DNAzyme machinery. *J. Am. Chem. Soc.* 2012, 134, 5504–5507.
27. Wang, S.; Yue, L.; Shpilt, Z.; Cecconello, A.; Kahn, J.S.; Lehn, J.M.; Willner, I. Controlling the Catalytic Functions of DNAzymes within Constitutional Dynamic Networks of DNA Nanostructures. *J. Am. Chem. Soc.* 2017, 139, 9662–9671.
28. Tian, Y.; Mao, C. Molecular gears: A pair of DNA circles continuously rolls against each other. *J. Am. Chem. Soc.* 2004, 126, 11410–11411.
29. Wang, F.; Elbaz, J.; Orbach, R.; Magen, N.; Willner, I. Amplified analysis of DNA by the autonomous assembly of polymers consisting of DNAzyme wires. *J. Am. Chem. Soc.* 2011, 133, 17149–17151.
30. Wang, F.; Elbaz, J.; Teller, C.; Willner, I. Amplified detection of DNA through an autocatalytic and catabolic DNAzyme-mediated process. *Angew. Chem.* 2011, 50, 295–299.
31. Zhong, W.; Tang, W.; Tan, Y.; Fan, J.; Huang, Q.; Zhou, D.; Hong, W.; Liu, Y. A DNA arithmetic logic unit for implementing data backtracking operations. *Chem. Commun.* 2019, 55, 842–845.
32. Shlyahovsky, B.; Li, Y.; Lioubashevski, O.; Elbaz, J.; Willner, I. Logic Gates and Antisense DNA Devices Operating on a Translator Nucleic Acid Scaffold. *ACS Nano* 2009, 3, 1831–1843.
33. Lake, A.; Shang, S.; Kolpashchikov, D.M. Molecular Logic Gates Connected through DNA Four-Way Junctions. *Angew. Chem.* 2010, 49, 4459–4462.

34. Li, W.; Zhang, F.; Yan, H.; Liu, Y. DNA based arithmetic function: A half adder based on DNA strand displacement. *Nanoscale* 2016, 8, 3775–3784.

35. Lin, H.Y.; Chen, J.Z.; Li, H.Y.; Yang, C.N. A simple three-input DNA-based system works as a full-subtractor. *Sci. Rep.* 2015, 5, 10686.

36. Wu, C.; Wang, K.; Fan, D.; Zhou, C.; Liu, Y.; Wang, E. Enzyme-free and DNA-based multiplexer and demultiplexer. *Chem. Commun.* 2015, 51, 15940–15943.

37. Zhu, J.; Zhang, L.; Li, T.; Dong, S.; Wang, E. Enzyme-Free Unlabeled DNA Logic Circuits Based on Toehold-Mediated Strand Displacement and Split G-Quadruplex Enhanced Fluorescence. *Adv. Mater.* 2013, 25, 2440–2444.

38. Chen, X. Expanding the rule set of DNA circuitry with associative toehold activation. *J. Am. Chem. Soc.* 2012, 134, 263–271.

39. Zhang, D.Y.; Winfree, E. Control of DNA Strand Displacement Kinetics Using Toehold Exchange. *J. Am. Chem. Soc.* 2009, 131, 17303–17314.

40. Xing, Y.; Yang, Z.; Liu, D. A Responsive Hidden Toehold To Enable Controllable DNA Strand Displacement Reactions. *Angew. Chem.* 2011, 50, 11934–11936.

41. Cherry, K.M.; Qian, L. Scaling up molecular pattern recognition with DNA-based winner-take-all neural networks. *Nature* 2018, 559, 370–376.

42. Chatterjee, G.; Dalchau, N.; Muscat, R.A.; Phillips, A.; Seelig, G. A spatially localized architecture for fast and modular DNA computing. *Nat. Nanotechnol.* 2017, 12, 920–927.

43. Srinivas, N.; Ouldridge, T.E.; Sulc, P.; Schaeffer, J.M.; Yurke, B.; Louis, A.A.; Doye, J.P.; Winfree, E. On the biophysics and kinetics of toehold-mediated DNA strand displacement. *Nucleic Acids Res.* 2013, 41, 10641–10658.

44. Frezza, B.M.; Cockroft, S.L.; Ghadiri, M.R. Modular multi-level circuits from immobilized DNA-Based logic gates. *J. Am. Chem. Soc.* 2007, 129, 14875–14879.

45. Seelig, G.; Soloveichik, D.; Zhang, D.Y.; Winfree, E. Enzyme-free nucleic acid logic circuits. *Science* 2006, 314, 1585–1588.

46. Qian, L.; Winfree, E. Scaling Up Digital Circuit Computation with DNA Strand Displacement Cascades. *Science* 2011, 332, 1196–1201.

47. Zhang, C.; Zhao, Y.; Xu, X.; Xu, R.; Li, H.; Teng, X.; Du, Y.; Miao, Y.; Lin, H.C.; Han, D. Cancer diagnosis with DNA molecular computation. *Nat. Nanotechnol.* 2020, 15, 709–715.

48. Green, S.J.; Lubrich, D.; Turberfield, A.J. DNA hairpins: Fuel for autonomous DNA devices. *Biophys. J.* 2006, 91, 2966–2975.

49. Seelig, G.; Yurke, B.; Winfree, E. Catalyzed relaxation of a metastable DNA fuel. *J. Am. Chem. Soc.* 2006, 128, 12211–12220.

50. Zhang, D.Y.; Turberfield, A.J.; Yurke, B.; Winfree, E. Engineering entropy-driven reactions and networks catalyzed by DNA. *Science* 2007, 318, 1121–1125.

51. Zhang, D.Y.; Hariadi, R.F.; Choi, H.M.; Winfree, E. Integrating DNA strand-displacement circuitry with DNA tile self-assembly. *Nat. Commun.* 2013, 4, 1965.

52. Elbaz, J.; Moshe, M.; Shlyahovsky, B.; Willner, I. Cooperative multicomponent self-assembly of nucleic acid structures for the activation of DNAzyme cascades: A paradigm for DNA sensors and aptasensors. *Chemistry* 2009, 15, 3411–3418.

53. Orbach, R.; Mostinski, L.; Wang, F.; Willner, I. Nucleic acid driven DNA machineries synthesizing Mg²⁺-dependent DNAzymes: An interplay between DNA sensing and logic-gate operations. *Chemistry* 2012, 18, 14689–14694.

54. Huang, P.J.; Lin, J.; Cao, J.; Vazin, M.; Liu, J. Ultrasensitive DNAzyme beacon for lanthanides and metal speciation. *Anal. Chem.* 2014, 86, 1816–1821.

55. Wu, N.; Willner, I. DNAzyme-Controlled Cleavage of Dimer and Trimer Origami Tiles. *Nano Lett.* 2016, 16, 2867–2872.

56. Endo, M.; Takeuchi, Y.; Suzuki, Y.; Emura, T.; Hidaka, K.; Wang, F.; Willner, I.; Sugiyama, H. Single-Molecule Visualization of the Activity of a Zn(2+)-Dependent DNAzyme. *Angew. Chem.* 2015, 54, 10550–10554.

57. Aleman-Garcia, M.A.; Orbach, R.; Willner, I. Ion-responsive hemin-G-quadruplexes for switchable DNAzyme and enzyme functions. *Chem.* 2014, 20, 5619–5624.

58. Wang, F.; Orbach, R.; Willner, I. Detection of metal ions (Cu²⁺, Hg²⁺) and cocaine by using ligation DNAzyme machinery. *Chem.* 2012, 18, 16030–16036.

59. Yang, J.; Wu, R.; Li, Y.; Wang, Z.; Pan, L.; Zhang, Q.; Lu, Z.; Zhang, C. Entropy-driven DNA logic circuits regulated by DNAzyme. *Nucleic Acids Res.* 2018, 46, 8532–8541.

60. Orbach, R.; Willner, B.; Willner, I. Catalytic nucleic acids (DNAzymes) as functional units for logic gates and computing circuits: From basic principles to practical applications. *Chem. Commun.* 2015, 51, 4144–4160.

61. Moshe, M.; Elbaz, J.; Willner, I. Sensing of UO₂²⁺ and Design of Logic Gates by the Application of Supramolecular Constructs of Ion-Dependent DNAzymes. *Nano Lett.* 2009, 9, 1196–1200.

62. Bi, S.; Yan, Y.; Hao, S.; Zhang, S. Colorimetric logic gates based on supramolecular DNAzyme structures. *Angew. Chem.* 2010, 49, 4438–4442.

63. Elbaz, J.; Lioubashevski, O.; Wang, F.; Remacle, F.; Levine, R.D.; Willner, I. DNA computing circuits using libraries of DNAzyme subunits. *Nat. Nanotechnol.* 2010, 5, 417–422.

64. Harding, B.I.; Pollak, N.M.; Stefanovic, D.; Macdonald, J. Repeated Reuse of Deoxyribozyme-Based Logic Gates. *Nano Lett.* 2019, 19, 7655–7661.

65. Enghiad, B.; Zhao, H.M. Programmable DNA-Guided Artificial Restriction Enzymes. *ACS Synth. Biol.* 2017, 6, 752–757.

66. Yang, X.L.; Tang, Y.A.; Mason, S.D.; Chen, J.B.; Li, F. Enzyme-Powered Three-Dimensional DNA Nanomachine for DNA Walking, Payload Release, and Biosensing. *Acs Nano* 2016, 10, 2324–2330.

67. Wang, F.; Zahid, O.K.; Swain, B.E.; Parsonage, D.; Hollis, T.; Harvey, S.; Perrino, F.W.; Kohli, R.M.; Taylor, E.W.; Hall, A.R. Solid-State Nanopore Analysis of Diverse DNA Base Modifications Using a Modular Enzymatic Labeling Process. *Nano Lett.* 2017, 17, 7110–7116.

68. Esadze, A.; Rodriguez, G.; Weiser, B.P.; Cole, P.A.; Stivers, J.T. Measurement of nanoscale DNA translocation by uracil DNA glycosylase in human cells. *Nucleic Acids Res.* 2017, 45, 12413–12424.

69. Meijer, L.; Joesaar, A.; Steur, E.; Engelen, W.; van Santen, R.A.; Merkx, M.; de Greef, T. Hierarchical control of enzymatic actuators using DNA-based switchable memories. *Nat. Commun.* 2017, 8, 11.

70. Song, T.; Eshra, A.; Shah, S.; Bui, H.; Fu, D.; Yang, M.; Mokhtar, R.; Reif, J. Fast and compact DNA logic circuits based on single-stranded gates using strand-displacing polymerase. *Nat. Nanotechnol.* 2019, 14, 1075–1081.

71. Zhang, C.; Wang, Z.; Liu, Y.; Yang, J.; Zhang, X.; Li, Y.; Pan, L.; Ke, Y.; Yan, H. Nicking-Assisted Reactant Recycle To Implement Entropy-Driven DNA Circuit. *J. Am. Chem. Soc.* 2019, 141, 17189–17197.

72. Johnson, J.A.; Dehankar, A.; Winter, J.O.; Castro, C.E. Reciprocal Control of Hierarchical DNA Origami-Nanoparticle Assemblies. *Nano Lett.* 2019, 19, 8469–8475.

73. Wang, J.; Yue, L.; Li, Z.; Zhang, J.; Tian, H.; Willner, I. Active generation of nanoholes in DNA origami scaffolds for programmed catalysis in nanocavities. *Nat. Commun.* 2019, 10, 4963.

74. Tikhomirov, G.; Petersen, P.; Qian, L. Triangular DNA Origami Tilings. *J. Am. Chem. Soc.* 2018, 140, 17361–17364.

75. Tian, Y.; Wang, T.; Liu, W.; Xin, H.L.; Li, H.; Ke, Y.; Shih, W.M.; Gang, O. Prescribed nanoparticle cluster architectures and low-dimensional arrays built using octahedral DNA origami frames. *Nat. Nanotechnol.* 2015, 10, 637–644.

76. Castro, C.E.; Kilchherr, F.; Kim, D.N.; Shiao, E.L.; Wauer, T.; Wortmann, P.; Bathe, M.; Dietz, H. A primer to scaffolded DNA origami. *Nat. Methods* 2011, 8, 221–229.

77. Chao, J.; Wang, J.; Wang, F.; Ouyang, X.; Kopperger, E.; Liu, H.; Li, Q.; Shi, J.; Wang, L.; Hu, J.; et al. Solving mazes with single-molecule DNA navigators. *Nat. Mater.* 2019, 18, 273–279.

78. Wickham, S.F.; Bath, J.; Katsuda, Y.; Endo, M.; Hidaka, K.; Sugiyama, H.; Turberfield, A.J. A DNA-based molecular motor that can navigate a network of tracks. *Nat. Nanotechnol.* 2012, 7, 169–173.

79. Teichmann, M.; Kopperger, E.; Simmel, F.C. Robustness of Localized DNA Strand Displacement Cascades. *ACS Nano* 2014, 8, 8487–8496.

80. Thubagere, A.J.; Li, W.; Johnson, R.F.; Chen, Z.; Doroudi, S.; Lee, Y.L.; Izatt, G.; Wittman, S.; Srinivas, N.; Woods, D.; et al. A cargo-sorting DNA robot. *Science* 2017, 357, 9.

81. Chen, G.; Gibson, K.J.; Liu, D.; Rees, H.C.; Lee, J.H.; Xia, W.; Lin, R.; Xin, H.L.; Gang, O.; Weizmann, Y. Regioselective surface encoding of nanoparticles for programmable self-assembly. *Nat. Mater.* 2019, 18, 169–174.

82. Lan, X.; Lu, X.; Shen, C.; Ke, Y.; Ni, W.; Wang, Q. Au nanorod helical superstructures with designed chirality. *J. Am. Chem. Soc.* 2015, 137, 457–462.

83. Lan, X.; Liu, T.; Wang, Z.; Govorov, A.O.; Yan, H.; Liu, Y. DNA-Guided Plasmonic Helix with Switchable Chirality. *J. Am. Chem. Soc.* 2018, 140, 11763–11770.

84. Thacker, V.V.; Herrmann, L.O.; Sigle, D.O.; Zhang, T.; Liedl, T.; Baumberg, J.J.; Keyser, U.F. DNA origami based assembly of gold nanoparticle dimers for surface-enhanced Raman scattering. *Nat. Commun.* 2014, 5.

85. Kühler, P.; Roller, E.M.; Schreiber, R.; Liedl, T.; Lohmüller, T.; Feldmann, J. Plasmonic DNA-Origami Nanoantennas for Surface-Enhanced Raman Spectroscopy. *Nano Lett.* 2014, 14, 2914–2919.

86. Xin, L.; Lu, M.; Both, S.; Pfeiffer, M.; Urban, M.J.; Zhou, C. Watching a Single Fluorophore Molecule Walk into a Plasmonic Hotspot. *Acs Photonics* 2019, 6, 985–993.

87. Urban, M.J.; Zhou, C.; Duan, X.; Liu, N. Optically Resolving the Dynamic Walking of a Plasmonic Walker Couple. *Nano Lett.* 2015, 15, 8392–8396.

88. Kyriazi, M.E.; Giust, D.; El-Sagheer, A.H.; Lackie, P.M.; Muskens, O.L.; Brown, T.; Kanaras, A.G. Multiplexed mRNA Sensing and Combinatorial-Targeted Drug Delivery Using DNA-Gold Nanoparticle Dimers. *Acs Nano* 2018, 12, 3333–3340.

89. Chai, H.; Miao, P. Bipedal DNA Walker Based Electrochemical Genosensing Strategy. *Anal. Chem.* 2019, 91, 4953–4957.

90. Chen, Y.; Corn, R.M. DNAzyme footprinting: Detecting protein-aptamer complexation on surfaces by blocking DNAzyme cleavage activity. *J. Am. Chem. Soc.* 2013, 135, 2072–2075.

91. Jiang, X.; Wang, H.; Chai, Y.; Li, H.; Shi, W.; Yuan, R. DNA Cascade Reaction with High-Efficiency Target Conversion for Ultrasensitive Electrochemiluminescence microRNA Detection. *Anal. Chem.* 2019, 91, 10258–10265.

92. Zhou, J.; Lai, W.; Zhuang, J.; Tang, J.; Tang, D. Nanogold-functionalized DNAzyme concatamers with redox-active intercalators for quadruple signal amplification of electrochemical immunoassay. *ACS Appl. Mater. Interfaces* 2013, 5, 2773–2781.

93. Guo, X.; Li, F.; Bai, L.; Yu, W.; Zhang, X.; Zhu, Y.; Yang, D. Gene Circuit Compartment on Nanointerface Facilitating Cascade Gene Expression. *J. Am. Chem. Soc.* 2019, 141, 19171–19177.

94. Yang, Y.; Huang, J.; Yang, X.; He, X.; Quan, K.; Xie, N.; Ou, M.; Wang, K. Gold Nanoparticle Based Hairpin-Locked-DNAzyme Probe for Amplified miRNA Imaging in Living Cells. *Anal. Chem.* 2017, 89, 5850–5856.

95. Liang, C.P.; Ma, P.Q.; Liu, H.; Guo, X.; Yin, B.C.; Ye, B.C. Rational Engineering of a Dynamic, Entropy-Driven DNA Nanomachine for Intracellular MicroRNA Imaging. *Angew. Chem.* 2017, 56, 9077–9081.

96. Gines, G.; Zadorin, A.S.; Galas, J.C.; Fujii, T.; Estevez-Torres, A.; Rondelez, Y. Microscopic agents programmed by DNA circuits. *Nat. Nanotechnol.* 2017, 12, 351–359.

97. Yao, G.; Li, J.; Li, Q.; Chen, X.; Liu, X.; Wang, F.; Qu, Z.; Ge, Z.; Narayanan, R.P.; Williams, D.; et al. Programming nanoparticle valence bonds with single-stranded DNA encoders. *Nat. Mater.* 2019, 19, 781–788.

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