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Redesign Nature: Self-assembly of Nucleic Acids in Material Science, Molecular Computing, and Synthetic Biology

Nucleic acids DNA and RNA are capable of self-assembling into secondary structures following Watson-Crick base-pairing rules. Commonly, they are found in cells as double helices further condensed into chromatin (DNA), as protein-associated complexes (ribosomes), or as functional structures for translating genes into proteins (tRNA). The self-assembly process was used in recent years to design structures at the nanoscale, while strand-displacement and dynamic motifs allowed the reconfiguration of such nanostructures in a mechanical-like fashion.^{1,2} These architectures were decorated with metal nanoparticles, proteins, and fluorophore-quencher pairs to generate: Optically active chiroplasmonic nanostructures,^{3,4} programmed catalytic functions, and finely regulated dynamic nano-devices. The programmability of DNA was also used to generate dynamic junctions in hybrid nanomaterials comprising carbon nanotubes.⁵ Recently, hybrid RNA- DNA triplexes were used to design logic gates for molecular computing,⁶ and control transcription in a synthetic biology system.⁷

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Martedì 4 giugno 2024, Ore 14:00, aula A

Ospite: Prof.ssa Alessandra Gianoncelli