1. Record Nr. UNINA9910410043303321 Autore YU Zutao Titolo Artificial Assemblies with Cooperative DNA Recognition / / by Zutao YU Pubbl/distr/stampa Singapore:,: Springer Singapore:,: Imprint: Springer,, 2020 **ISBN** 981-15-4423-9 Edizione [1st ed. 2020.] Descrizione fisica 1 online resource (XV, 136 p. 136 illus., 119 illus. in color.) Collana Springer Theses, Recognizing Outstanding Ph.D. Research, , 2190-5053 Disciplina 660.6 Soggetti Biotechnology Bioorganic chemistry Nucleic acids Pharmaceutical chemistry Bioorganic Chemistry **Nucleic Acid Chemistry Medicinal Chemistry** Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia Note generali "Doctoral thesis accepted by Kyoto University, Kyoto, Japan." Nota di contenuto 1. Synthetic transcription factors (Syn-TFs): design, progress and perspectives -- 2. Pip-HoGu, an artificial assembly with cooperative DNA recognition capable of mimicking transcription factor pairs -- 3. Orthogonal PNA dimerization domains empower DNA binders with cooperativity and versatility mimicking that of the transcription factor pairs -- 4. Advanced DNA binding system mimicking the cooperative function of transcription factor pairs precisely recruits the epigenetic modifiers to the DNA repeat binding sites. Sommario/riassunto This book presents three types of synthetically cooperative DNA recognizing assemblies, in order to advance the development of programmable DNA-binding pyrrole-imidazole polyamides (PIPs). PIPs represent the best-characterized class of small molecule DNA binders that can be modified to bind with any predetermined DNA sequence and regulate gene expression patterns in a transgene-free and costeffective manner. PIPs are characterized by their small molecular size, high binding affinity, programmability, sequence selectivity, and

moderate cell permeability. In recent years, there have been numerous

novel studies on the applications of these biological tools; this research is thoroughly reviewed in the first chapter. There are several critical issues, however, that impede the further broad study of PIPs, which greatly concern the author. For instance, the short PIP version has an excessively hi^10 bp; this significantly decreases cell permeability. Moreover, the conventional binding strategy for PIP design cannot apply to flexible DNA binding—for example, the DNA-binding mode of a transcription factor pair. In this book, the author describes the development of three kinds of cooperative DNA-binding systems that help resolve the current highly problematic issues concerning PIPs. These three systems offer a range of significant advantages, such as favorable sequence selectivity, long recognition sequence, higher binding affinity, and a flexible gap distance. Released at a critical juncture in the application of PIPs, this book will greatly facilitate their use as therapeutic drugs in the treatment of cancer and hereditary diseases, and in regenerative medicine. .