

1. Record Nr.	UNINA9910254300903321
Autore	Yamagishi Michel Eduardo Beleza
Titolo	Mathematical Grammar of Biology // by Michel Eduardo Beleza Yamagishi
Pubbl/distr/stampa	Cham : , : Springer International Publishing : , : Imprint : Springer, , 2017
ISBN	3-319-62689-2
Edizione	[1st ed. 2017.]
Descrizione fisica	1 online resource (XII, 82 p. 19 illus., 17 illus. in color.)
Collana	SpringerBriefs in Mathematics, , 2191-8198
Disciplina	576.58 577.88
Soggetti	Biomathematics Bioinformatics Biology—Philosophy Genetics and Population Dynamics Computational Biology/Bioinformatics Philosophy of Biology
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di contenuto	Chapter 01- Introduction -- Chapter 02- Modeling Human Nucleotide Frequencies -- Chapter 03- Expanding the Grammar of Biology -- Chapter 04 - "In God We Trust; All Others, Bring Data."- References.
Sommario/riassunto	This seminal, multidisciplinary book shows how mathematics can be used to study the first principles of DNA. Most importantly, it enriches the so-called "Chargaff's grammar of biology" by providing the conceptual theoretical framework necessary to generalize Chargaff's rules. Starting with a simple example of DNA mathematical modeling where human nucleotide frequencies are associated to the Fibonacci sequence and the Golden Ratio through an optimization problem, its breakthrough is showing that the reverse, complement and reverse-complement operators defined over oligonucleotides induce a natural set partition of DNA words of fixed-size. These equivalence classes, when organized into a matrix form, reveal hidden patterns within the DNA sequence of every living organism. Intended for undergraduate and graduate students both in mathematics and in life sciences, it is

also a valuable resource for researchers interested in studying invariant genomic properties.

---