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This book is organized into three parts. In part I (Chapters 1-7), we introduce the field of bioinformatics. Topics include access to sequence (and other) data, pairwise sequence alignment, BLAST and other methods of searching databases, multiple sequence alignment, and phylogeny. In part II (Chapters 8-14), we discuss functional genomics, moving from DNA (including next-generation sequencing) to RNA (including RNAseq) to protein.

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