

Table of Contents

- Introduction G. Williams
- Nucleic Acid and Protein Sequence Databases R. Cotton
- Phenotype, Mutation, and Genetic Linkage Databases and Their Links to Sequence Databases R. Guigo
- DNA Composition, Codon Usage, and Exon Prediction W. Taylor
- Properties of Amino Acids in Sequences M. Gribskov
- Sequence Comparison J. Epplen
- Simple Repetitive Sequences in DNA Databanks. P. Bucher
- Gene Feature Identification D. Higgins
- Multiple Sequence Alignment K. Triman
- RNAs C. Ponting and D.J. Blake
- Predicting the Evolution, Structure, and Function of Proteins from Sequence Information D. Jones
- Structure Databases and Structure Alignments M. Gribskov
- Bourne, Smith, Integrated Data Resource for Protein Kinases R. Baldock
- Gene Expression Databases P.D. Karp
- Pathways and Development