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By Andreas Gogol-D

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In the context of genomics, annotation is the process of marking the genes and other biological features in a DNA sequence. This process needs to be automated because

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Dynamic visual data mining 23 Nucleotide sequence analysis The nucleotide sequence analysis category consists of the following six modules (Figure 1):

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1. Biotechniques. 1999 Jul;27(1):76-8. Biological sequence analysis using regular expressions. Horton RM. Attotron Biosensor Corporation, Carson City, NV, USA

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Biological Sequence Analysis: Probabilistic -

Biological Sequence Analysis: Probabilistic Models of Proteins and Nucleic Acids, Richard Durbin, Sean R. Eddy, Anders Krogh, Graeme J. Mitchison

Biological Sequence Analysis using the SeqAn C++ -

Before the SeqAn project, there was clearly a lack of available implementations in sequence analysis, even for standard tasks. Implementations of needed algorithmic

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HMMER - Official Site -

HMMER is used for searching sequence databases for homologs of protein sequences, and for making protein sequence alignments. It implements methods using

Sequence analysis - Wikipedia, the free -

In bioinformatics, sequence analysis is the process of subjecting a DNA, RNA or peptide sequence to any of a wide range of analytical methods to understand its