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SOLVING SOME SEQUENCE PROBLEMS ON RUN-LENGTH ENCODED STRINGS

Condition: New. Publisher/Verlag: VDM Verlag Dr. Müller | Longest Common Subsequences, Edit Distances, and Squares | Measuring the similarity or difference between two strings is a fundamental problem to many applications. In bioinformatics, one has to predict the structures of RNA and proteins, to classify the functions of molecules, to infer the phylogeny of organisms, and to search entries in huge sequence databases. While processing electronic documents, one needs fast and flexible indexing techniques to perform searches. For this purpose,...

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