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Condition: New. Publisher/Verlag: VDM Verlag Dr. Müller | Basics, Concepts and Applications | Pattern matching plays a central role in different aspects of biological sequence analysis, and has been used in applications as diverse as short-gun sequencing, multiple sequence alignments, gene finding, analysis of repetition structures, searching for unique oligonucleotides, prediction of protein function and structure, sequence homology search, finding DNA-binding protein motifs, etc. Suffix trees and suffix arrays are primary data structures used in rapid pattern matching. And the problem of suffix sorting is a fundamental problem in constructing suffix...

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- Authored by Nan, Fei
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