



## Proceedings Of The 5th Asia-Pacific Bioinformatics Conference

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**Summary** : Free proceedings of the 5th asia-pacific bioinformatics conference pdf download - high-throughput sequencing and functional genomics technologies have given us the human genome sequence as well as those of other experimentally medically and agriculturally important species and have enabled large-scale genotyping and gene expression profiling of human populations databases containing large numbers of sequences polymorphisms structures and gene expression profiles of normal and diseased tissues are being rapidly generated for human and model organisms bioinformatics is thus rapidly growing in importance in the annotation of genomic sequences the understanding of the interplay among and between genes and proteins the analysis of genetic variability of species the identification of pharmacological targets and the inference of evolutionary origins mechanisms and relationships this proceedings volume contains an up-to-date exchange of knowledge ideas and solutions to conceptual and practical issues of bioinformatics by researchers professionals and industrial practitioners at the 5th asia-pacific bioinformatics conference held in hong kong in january 2007 contents exploring genomes of distantly related mammals j a marshall graves subtle motif discovery for detection of dna regulatory sites m comin l parida using formal concept analysis for microarray data comparison v choi et al computing the quartet distance between evolutionary trees of bounded degree m stissing et al a randomized algorithm for comparing sets of phylogenetic trees s-j sul t l williams exact and heuristic approaches for identifying disease-associated snp motifs g huang et al the distance between randomly constructed genomes w xu semi-supervised pattern learning for extracting relations from bioscience texts s ding et al fast structural similarity search based on topology string matching s-h park et al and other papers readership academics researchers and graduate students in bioinformatics and computer science keywords bioinformatics computational biology systems biology statistical modeling comparative genomics evolutionary biology data mining structural bioinformatics statistical genetics

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