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## DEVELOPMENT OF PHYLOGENETIC TREE BASED ON KIMURA'S METHOD



Pankaj Bhambri  
Franky Goyal  
**Development of  
Phylogenetic Tree based on  
Kimura's Method**  
Based on Un-weighted Pair Group method with  
Arithmetic Mean (UPGMA) and Neighbor Joining (NJ)  
Scoring Techniques



Condition: New. Publisher/Verlag: LAP Lambert Academic Publishing | Based on Un-weighted Pair Group method with Arithmetic Mean (UPGMA) and Neighbor Joining (NJ) Scoring Techniques | The research in bioinformatics has accumulated large amount of data. It is the study of Bio-molecules information. Bioinformatics offers different knowledge discovery concepts for molecular biology and has many practical applications. DNA sequence alignment is one of the applications of the bioinformatics. Multiple sequence alignment is used to align the biological sequences along a column...

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