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by: kal renganathan sharma abstract: a state-of-the-art textbook on bioinformatics covering the latest 21st-century technology.

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the hidden markov model ... anywhere there may be a model producing a sequence of observations. in bioinformatics, it has been used in sequence alignment, ...

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multiple sequence alignment zgene finding . what hmms do? a hmm is a statistical model for sequences of discrete symbols. ... hidden markov models in bioinformatics

APPLICATION OF HIDDEN MARKOV MODELS IN BIOINFORMATICS

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application of hidden markov models in bioinformatics ... random model: $1-v$ $1-v$ v 1 ... multiple sequence alignment: ...

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