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Statistical Science 2003, Vol. 18, No. 2, 241-255 C)
Institute Of Mathematical Statistics, 2003
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The Use Of The Bootstrap In The Area Of Systematic
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Results Of Multiple Sequence Alignment That Is Called As Bootstrap Replicate. Each Bootstrap Replicate Is In The Form Of Phylogenetic Tree Contains Same Number Of Species [10]. The Branches Repeated In Maximum Bootstrap Replicates Have High Confidence. This Method Has Been Very Successful Among Alignment Based Methods [11]. Jan 1th, 2024

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