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Alexei J. Drummond; Remco R. Bouckaert**

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BEAST. BEAST (Bayesian Evolutionary Analysis Sampling Trees) is a program for evolutionary inference of molecular sequences designed by Andrew Rambaut and Alexei

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1. BMC Evol Biol. 2007 Nov 8;7:214. BEAST: Bayesian evolutionary analysis by sampling trees. Drummond AJ(1), Rambaut A. Author information: (1

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