

Postdoctoral Position in Phylogenetics at the University of Alaska Fairbanks

Preliminary applications for a post-doctoral position in Mathematical Phylogenetics are being accepted now. Under an NIH grant joint with Stanford University and University of New Mexico, we seek a candidate with a background in mathematics, statistics or quantitative biology to join our group in Alaska. Applicants should be interested in working on mathematical, computational, and statistical aspects of gene-tree/species-tree problems.

The position will provide opportunities to visit and collaborate with the research groups of biologist N. Rosenberg (Stanford) and statistician J. Degnan (University of New Mexico) for interdisciplinary training. However, the PIs (E. Allman and J. Rhodes) in Alaska are mathematicians, and the successful applicant should have background and interests appropriate to join their research program (e.g., applied algebra, combinatorics, population genetics, and/or phylogenetics).

A start date of January 2017 is preferred, with six months of funding available during the academic year 2016-2017, and nine-months for each of the two subsequent academic years.

Please e-mail e.allman@alaska.edu and j.rhodes@alaska.edu to express interest by February 28, 2016. The e-mail should include a brief explanation of your interest in the position, an outline of educational background and training in relevant subject areas, and a CV. A full application process will follow.