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Debevec, E., R. B. Gates, M. Masuda, J. Pella, J. Reynolds, and L. W. Seeb. 2000. SPAM (Version 3.2) Statistics program for analyzing mixtures. Journal of Heredity 91(6):509-510.

**Abstract:** Overview: This article is an overview of SPAM (Statistics program for Analyzing Mixtures), a software tool for computing the maximum likelihood estimates for a mixed population of stock analysis. SPAM has two basic modes of operation, estimation mode and simulation mode. Results can be output in an estimation file or a simulation file, both of which can be edited, analyzed and viewed in a Windows environment. SPAM is available free of chafe from the Alaska Department of Fish and Game Gene Conservation Laboratory.

Keywords: None.

URL: http://jhered.oxfordjournals.org/content/91/6/509.full.pdf+html

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