

Package ‘hierfstat’

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Title Estimation and tests of hierarchical F-statistics

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Description This R package allows the estimation of hierarchical F-statistics from haploid or diploid genetic data with any numbers of levels in the hierarchy, following the algorithm of Yang (Evolution, 1998, 52(4):950-956). Functions are also given to test via randomisations the significance of each F and variance components, using the likelihood-ratio statistics G -see Goudet et al (Genetics, 1996, 144(4): 1933-1940)

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URL <http://www.r-project.org>, <http://www.unil.ch/popgen/software/hierfstat.htm>

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hierfstat *General information on the hierfstat package*

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