

Brief Description:

Checks pedigrees for Mendelian Errors and, when errors are found, systematically jackknives every typed pedigree member to determine if eliminating this member will remove all Mendelian Errors from the pedigree.

=====

Installation procedures

Installation procedures in this file are for setting up a 'local library' for a user. If a large group of users plan to use the same library, then install on a system-wide 'library' location designated by a system administrator.

=====

R Installation

The mend.err package is not available for distribution through any R websites. To install, download the README.mend.err file with mend.err_x.y.z.tar.gz and execute the following command with the local "INSTALL LIBRARY DIR PATH" correctly specified.

```
R CMD INSTALL -l /INSTALL LIBRARY DIR PATH/ mend.err_x.y.z.tar.gz
```

Additional details on R may be found on the R project website (<http://www.r-project.org>).

=====

User Documentation:

Please find specific instructions in the help files for the mend.err functions, for example by typing help(mend.err) at the R prompt.