

R documentation

of ‘/Users/stevenmartell/Documents/CURRENT’ etc.

November 11, 2010

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| Riscam-package | <i>This package contains supporting functions and routines for the stock</i> |
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Description

Provides R support for Integrated Statistical Catch Age Model (iscam), includes routines for reading ADMB output and custom graphics and table routines for summarizing model results.

Details

| | |
|-----------|------------|
| Package: | Riscam |
| Type: | Package |
| Version: | 1.0 |
| Date: | 2010-11-10 |
| License: | GPL |
| LazyLoad: | yes |

~~ An overview of how to use the package, including the most important ~~ functions ~~

Author(s)

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See Also

~~ Optional links to other man pages, e.g. ~~ <pkg> ~~

A

*Pacific hake results***Description**

Output form the hake assessment that is provided in the iscam users guide

Usage

```
data (A)
```

Format

The format is: List of 42 \$ f : num -379 \$ nlvec : num [1:6, 1:2] -121.6 30.4 -407.5 55.2 11- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:2] "V1" "V2" \$ ro : num 3.21 \$ rbar : num 0.845 \$ bo : num 3.75 \$ kappa : num 8.8 \$ m : num 0.228 \$ tau : num 1.35 \$ sig : num 0.56 \$ age_tau2 : num [1:2] 0.387 0.681 \$ ngear : num 2 \$ yr : num [1:33] 1977 1978 1979 1980 1981 ... \$ yrs : num [1:34] 1977 1978 1979 1980 1981 ... \$ iyr : num [1:13] 1977 1980 1983 1986 1989 ... \$ age : num [1:15] 1 2 3 4 5 6 7 8 9 10 ... \$ la : num [1:15] 14.2 24.6 32.1 37.5 41.5 ... \$ wa : num [1:15] 0.0144 0.0743 0.1652 0.2646 0.3574 ... \$ fa : num [1:15] 1.32e-05 1.16e-03 3.58e-02 2.19e-01 3.53e-01 ... \$ log_sel : num [1:66, 1:16] 1 1 1 1 1 1 1 1 1 1- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:16] "V1" "V2" "V3" "V4" ... \$ vax : num [1:15] 0 0 0 0 0 0 0 0 0 0 ... \$ obs_ct : num [1:2, 1:33] 0.133 0 0.104 0 0.137- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:33] "V1" "V2" "V3" "V4" ... \$ ct : num [1:2, 1:33] 1.33e-01 3.55e-31 1.04e-01 3.31e-31 1.37e-01- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:33] "V1" "V2" "V3" "V4" ... \$ ft : num [1:2, 1:33] 0.326 0 0.32 NA 0.36- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:33] "V1" "V2" "V3" "V4" ... \$ bt : num [1:34] 1.15 1.06 1.09 1.09 1.34 ... \$ sbt : num [1:34] 1.028 0.926 0.804 0.715 0.935 ... \$ rectype : num 1 \$ rt : num [1:32] 4.346 0.262 0.77 17.201 0.19 ... \$ log_rt(syr,nyr): num [1:33] -1.123 1.469 -1.339 -0.262 2.845 ... \$ delta : num [1:32] 1.471 -1.303 -0.175 2.977 -1.626 ... \$ it : num [1:13] 1.92 2.12 1.65 2.86 1.24 ... \$ pit : num [1:13] 0.977 0.825 1.504 2.684 3.227 ... \$ epsilon : num [1:13] 0.6728 0.9408 0.0911 0.0624 -0.9579 ... \$ F : num [1:33, 1:15] 0.10063 0.01509 0.00483 0.00547 0.0261- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:15] "V1" "V2" "V3" "V4" ... \$ a_sage : num [1:2] 2 2 \$ a_nage : num [1:2] 15 15 \$ A : num [1:46, 1:16] 1977 1978 1979 1980 1981- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:16] "V1" "V2" "V3" "V4" ... \$ Ahat : num [1:46, 1:16] 1977 1978 1979 1980 1981- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:16] "V1" "V2" "V3" "V4" ... \$ N : num [1:34, 1:15] 0.325 4.346 0.262 0.77 17.201- attr(*, "dimnames")=List of 2\$: NULL\$: chr [1:15] "V1" "V2" "V3" "V4" ... \$ fmsy : num 0.322 \$ msy : num 0.206 \$ bmsy : num 1.48 \$ fit :List of 10 ..\$ npar : int 174 ..\$ nlogl : num -379 ..\$ maxgrad : num 5.41e-05 ..\$ npar : num 175 ..\$ logDetHess: num 888 ..\$ names : chr [1:175] "theta[1]" "theta[2]" "theta[3]" "theta[4]"\$ est : num [1:175] 1.167 0.688 -1.478 -0.168 0.294\$ std : num [1:175] 0.3258 0.2142 0.0489 0.1187 0.0435\$ cor : num [1:175, 1:175] 1 -0.238 0.126 0.116 -0.215\$ cov : num [1:175, 1:175] 0.10614 -0.01659 0.00201 0.00448 -0.00304 ...

Examples

```
data(A)
## maybe str(A) ; plot(A) ...
```

| | |
|-----------|-------------------|
| read.admb | <i>Data input</i> |
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Description

Reads the parameter files, correlation file, and report file outputs from an ADMB program.

Usage

```
read.admb(ifile)
read.fit(ifile)
read.rep(fn)
read.psv(fn, nsamples=10000)
```

Arguments

| | |
|----------|---|
| ifile | the file name (without the extension) of the ADMB program |
| fn | the file name with the extension, use for read.rep & read.psv |
| nsamples | the number of posterior samples to read in from the binary *.psv file |

Details

The function read.admb reads the contents of the ADMB output parameter file (*.par) and the correlation file (*.cor) as well as the report file (*.rep). The function read.rep, just reads the report file only.

The read.psv function reads the binary output from the -mcsave command line option in ADMB. The default number of samples (rows) is 10,000. Note that this function is also called from read.admb if, and only if, the *.psv file exists.

Author(s)

S. Martell, Anders Nielsen, and the original contribution came from George Watters

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.

## The function is currently defined as
function(ifile)
{
  ret=read.fit(ifile)

  fn=paste(ifile, '.rep', sep='')
  A=read.rep(fn)
  A$fit=ret

  pfn=paste(ifile, '.psv', sep='')
  if(file.exists(pfn))
  A$mc=read.psv(pfn)
```

```
return(A)  
}
```

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