

Package ‘PBSadmb’

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admb

*Start the PBS ADMB GUI***Description**

Start up the PBS GUI for running ADMB.

Usage

```
admb(prefix="", wdf="admbWin.txt", optfile="ADopts.txt")
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
wdf	string name of the <i>window description file</i> that creates the GUI.
optfile	string name of options file (usually in user's working directory).

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#)

appendLog

*Append Data to Log File***Description**

Append summary information or output to a previously created log file.

Usage

```
appendLog(prefix, lines)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
lines	data to append to 'prefix'.log).

Value

No explicit value returned. Appends data into a log file 'prefix'.log.

Note

A wrapper function that can be called from a GUI exists as `.win.appendLog`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[startLog](#), [editADfile](#)

checkADopts

Check ADMB Options for Link Integrity

Description

Check that .ADopts has all required components and that links point to actual files on the hard drive.

Usage

```
checkADopts(opts=getOptions(.PBSadmb), check=c("admbpath", "gccpath", "editor"),
            warn=TRUE, popup=FALSE)
```

Arguments

opts	ADMB options values.
check	components of .ADopts to check.
warn	logical: if TRUE, print the results of the check to the R console.
popup	logical: if TRUE, display program location problems in a popup GUI.

Value

Boolean value where TRUE indicates all programs were located in the specified directories and FALSE if at least one program cannot be found. The returned Boolean scalar has two attributes:
 warn - named list of test results, and
 message - named vector of test results.

Note

A wrapper function that can be called from a GUI exists as .win.checkADopts.

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#), [readADopts](#)

cleanAD

Clean ADMB-Generated Files from the Working Directory

Description

Detects files in the working directory with the specified `prefix` and removes them all save those with the suffix `.tpl`, `.dat`, and `.pin`.

Usage

```
cleanAD(prefix)
```

Arguments

`prefix` string name prefix of the ADMB project (e.g., "vonb").

Details

Aside from potential garbage files with the specified `prefix`, other files associated with ADMB are detected. Also files `*.tmp` and `*.bak` are displayed. Calling `cleanAD` invokes the hidden function `.cleanUp`, which creates a GUI menu of the potential garbage files. The user can select whichever files s/he wishes for disposal.

Value

Returns nothing. Invokes a GUI menu of potential garbage files.

Note

A wrapper function that can be called from a GUI exists as `.win.cleanAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeAD](#), [runAD](#), [readRep](#)

compAD

Compile C Code

Description

Compile C++ code in '`prefix`'.cpp to create a binary object file '`prefix`'.o.

Usage

```
compAD(prefix, ranef=FALSE, safe=TRUE, dll=FALSE, debug=FALSE, logfile=TRUE, add=TRUE, verbose=TRUE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
raneff	logical: use the random effects model, otherwise use the normal model (currently does not influence the compile stage, but the argument is preserved here for future development).
safe	logical: if TRUE, use safe mode with bounds checking on all array objects, otherwise use optimized mode for fastest execution.
dll	create dll (rather than executable)
debug	compile with debug symbols
logfile	logical: if TRUE, create a log file of the messages from the shell call.
add	logical: if TRUE, append shell call messages to an existing log file.
verbose	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function uses the C++ compiler declared in `.ADopts`. If `logfile=TRUE`, any errors will appear in `'prefix'.log`. If `verbose=TRUE`, they will appear in the R console.

Value

Invisibly returns the shell call and its messages.

Note

A wrapper function that can be called from a GUI exists as `.win.compAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[convAD](#), [linkAD](#), [makeAD](#)

convAD

Convert TPL Code to CPP Code

Description

Convert code in `'prefix'.tpl` to C++ code in `'prefix'.cpp`.

Usage

```
convAD(prefix, raneff=FALSE, safe=TRUE, dll=FALSE, debug=FALSE, logfile=TRUE, add=TRUE, verbose=TRUE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
raneff	logical: if TRUE, use the random effects model executable <code>tpl2rem.exe</code> , otherwise use the normal model executable <code>tpl2cpp.exe</code> .
safe	logical: if TRUE, use safe mode with bounds checking on all array objects, otherwise use optimized mode for fastest execution.
dll	create dll (rather than executable)
debug	compile with debug symbols
logfile	logical: if TRUE, create a log file of the messages from the shell call.
add	logical: if TRUE, append shell call messages to an existing log file.
verbose	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function invokes the ADMB command `tpl2cpp.exe` or `tpl2rem.exe`, if `raneff` is FALSE or TRUE respectively. If `logfile=TRUE`, any errors will appear in `'prefix'.log`. If `verbose=TRUE`, they will appear in R console.

Value

Invisibly returns the shell call and its messages.

Note

A wrapper function that can be called from a GUI exists as `.win.convAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[compAD](#), [linkAD](#), [makeAD](#)

convOS

Convert Text Files to Default OS Format

Description

Convert text files to the default format of the operating system.

Usage

```
convOS(inam, onam = inam, path = getwd() )
```

Arguments

inam	string vector of names specifying files to be converted to the format of the operating system.
onam	string vector of name specifying the output files (the default overwrites the input file).
path	string specifying the path where the input files are located (defaults to current working directory).

Value

Text file(s) formatted in accordance with standards of the operating system.

Note

This function essentially executes a `readLines` command followed by a call to `writeLines`.

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo BC

See Also

[copyFiles](#), `.addQuotes`

copyFiles

Copy System Files

Description

Copy files with specified prefixes and suffixes from one location to another.

Usage

```
copyFiles(prefix, suffix=NULL, srcdir=getwd(), dstdir=getwd(), ask=TRUE)
```

Arguments

<code>prefix</code>	string scalar/vector of potential file prefixes.
<code>suffix</code>	string scalar/vector of potential file suffixes.
<code>srcdir</code>	source directory from which to copy files.
<code>dstdir</code>	destination directory to copy files to.
<code>ask</code>	logical: if TRUE, popup boxes will prompt the user for every instance that a file will be over-written.

Details

This function uses R's `list.files` and `file.copy` functions. The pattern recognition tends not to work when given the wildcard character `*`; however, the user may use this character, and the code will interpret it.

Value

Invisibly returns a Boolean vector with names of files that have been copied or not.

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo, BC

See Also

[editAD](#)

editAD

*Edit ADMB Files***Description**

Edit files associated with specified prefix and suffixes.

Usage

```
editAD(prefix, suffix=c(".tpl", ".cpp", ".log"))
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
suffix	string scalar/vector specifying one or more suffixes.

Value

Invisibly returns Boolean vector with elements TRUE if files exist, FALSE if they do not.

Note

A wrapper function that can be called from a GUI exists as `.win.editAD`.

This function explicitly uses the editor chosen for PBSadmb. PBSmodelling has another function `openFile` that uses Windows file associations or an application specified with `setPBSext`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[editADfile](#), [makeADopts](#)

editADfile

*Edit a File***Description**

Edit a file using the text editor specified in `.ADopts`.

Usage

```
editADfile(fname)
```

Arguments

fname	string name of file in current working directory (or elsewhere if path delimited by / or \).
-------	--

Value

Returns Boolean: TRUE if file exists, FALSE if it does not.

Note

This function explicitly uses the editor chosen for PBSadmb. PBSmodelling has another function `openFile` that uses Windows file associations or an application specified with `setPBSext`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[editAD](#), [makeADopts](#)

installADMB

Install windows ADMB and MinGW binaries

Description

Only applicable for Windows: Downloads and installs the windows ADMB and MinGW binaries.

A user interface is displayed which allows to select to install either ADMB and/or MinGW for either 32bit or 64bit versions of Windows. Default installation directories are suggested for installation; however, a user may choose to install the binaries elsewhere.

A file "pathconfig.txt" is saved in the PBSadmb library (under R), which keeps track of the most recently installed locations.

Usage

```
installADMB()
```

linkAD

Link Object Files to Make an Executable

Description

Links the binary object file 'prefix'.o to the ADMB libraries and produces the executable file 'prefix'.exe.

Usage

```
linkAD(prefix, raneff=FALSE, safe=TRUE, dll=FALSE, debug=FALSE, logfile=TRUE, add=TRUE, verbose=TRUE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
raneff	logical: use the random effects model, otherwise use the normal model.
safe	logical: if TRUE, use safe mode with bounds checking on all array objects, otherwise use optimized mode for fastest execution.
dll	create dll (rather than executable)
debug	compile with debug symbols
logfile	logical: if TRUE, create a log file of the messages from the shell call.
add	logical: if TRUE, append shell call messages to an existing log file.
verbose	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function uses the C++ compiler declared in `.ADopts`. If `logfile=TRUE`, any errors will appear in `'prefix'.log`. If `verbose=TRUE`, they will appear in the R console.

Value

Invisibly returns the shell call and its messages.

Note

A wrapper function that can be called from a GUI exists as `.win.linkAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[convAD](#), [compAD](#), [makeAD](#)

makeAD

Make an Executable Binary File from a C File

Description

Essentially a wrapper function that calls in sequence: `convAD`, `compAD`, and `linkAD`.

Usage

```
makeAD(prefix, ranef=FALSE, safe=TRUE, dll=FALSE, debug=FALSE, logfile=TRUE, add=TRUE, verbose=TRUE)
```

Arguments

<code>prefix</code>	string name prefix of the ADMB project (e.g., "vonb").
<code>ranef</code>	logical: use the random effects model, otherwise use the normal model.
<code>safe</code>	logical: if TRUE, use safe mode with bounds checking on all array objects, otherwise use optimized mode for fastest execution.
<code>dll</code>	create dll (rather than executable)
<code>debug</code>	compile with debug symbols
<code>logfile</code>	logical: if TRUE, create a log file of the messages from the shell call.
<code>add</code>	logical: if TRUE, append shell call messages to an existing log file.
<code>verbose</code>	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function uses the C++ compiler declared in `.ADopts`. If `logfile=TRUE`, any errors will appear in `'prefix'.log`. If `verbose=TRUE`, they will appear in the R console.

Value

Returns nothing. The three functions called by `makeAD` each return the shell call and its messages.

Note

A wrapper function that can be called from a GUI exists as `.win.makeAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[convAD](#), [compAD](#), [linkAD](#), [cleanAD](#)

makeADopts

Creates the ADMB Options List

Description

Creates a global list object detailing the pathways to the ADMB directory, the GCC bin, and the user's preferred text editor.

Usage

```
makeADopts(admbpath, gccpath, editor)
```

Arguments

admbpath	explicit path to the user's ADMB directory.
gccpath	explicit path to the user's GCC bin (C-compiler) directory.
editor	explicit path and program to use for editing text.

Value

Creates a global, hidden list object called `.ADopts`.

Note

A wrapper function that can be called from a GUI exists as `.win.makeADopts`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#), [writeADopts](#)

plotMC	<i>Plot Results of MCMC Simulation</i>
--------	--

Description

Plot results of an ADMB MCMC simulation using various plot methods.

Usage

```
plotMC(prefix, act="pairs", pthin=1, useCols=NULL)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
act	string scalar: action describing plot type (current choices: "pairs", "eggs", "acf", "trace", and "dens").
pthin	numeric scalar indicating interval at which to collect records from the .mc.dat file for plotting.
useCols	logical vector indicating which columns of .mc.dat to plot.

Note

A wrapper function that can be called from a GUI exists as `.win.plotMC`. Use the PBSadmb GUI to explore these plots easily.

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo BC, Canada

See Also

[runMC](#), [showADargs](#)

readADopts	<i>Reads an ADMB Options List into Memory From a File</i>
------------	---

Description

Reads ADMB options into a global, hidden list object called `.ADopts` from an ASCII text file using `PBSmodelling::readList`.

Usage

```
readADopts(optfile="ADopts.txt")
```

Arguments

optfile	string name of an ASCII text file containing ADMB options information.
---------	--

Value

No values returned. Reads the ADMB options into the list object `.ADopts`.

Note

A wrapper function that can be called from a GUI exists as `.win.readADopts`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#), [writeADopts](#)

readRep

Read an ADMB Report into R Memory

Description

Import ADMB-generated report files into R's memory using the names of the report files to name the R-objects.

Usage

```
readRep(prefix, suffix=c(".cor", ".rep", ".std", ".mc.dat"), global=FALSE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
suffix	string scalar/vector specifying one or more suffixes.
global	logical: if TRUE, save the imported reports as objects to global environment using the same names as the report files.

Details

If the report object is one of `c(".cor", ".std", ".mc.dat")`, the report object is a data frame, otherwise it is a string vector. Multiple report objects are returned as a list of objects. A single report object is returned as the object itself.

This function attempts to detect the file format from a number of possibilities. For example, if the file has the special format recognized by `PBSmodelling`, then the function returns a list with named components. The example `vonb` included with this package shows how to write the template to get consistent variable names between ADMB and R. See the User's Guide for complete details.

Value

Invisibly returns the list of report objects. If only one report is imported, a single report object is returned.

Note

A wrapper function that can be called from a GUI exists as `.win.readRep`.

Author(s)

Rowan Haigh, Pacific Biological Station, Nanaimo BC, Canada

See Also

[editADfile](#), `.win.viewRep`

`runAD`*Run an Executable Binary File*

Description

Run the executable binary file 'prefix'.exe that was created by makeAD.

Usage

```
runAD(prefix, argvec="", logfile=TRUE, add=TRUE, verbose=TRUE)
```

Arguments

<code>prefix</code>	string name prefix of the ADMB project (e.g., "vonb").
<code>argvec</code>	string scalar/vector of arguments appropriate for the executable 'prefix'.exe.
<code>logfile</code>	logical: if TRUE, create a log file of the messages from the shell call.
<code>add</code>	logical: if TRUE, append shell call messages to an existing log file.
<code>verbose</code>	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function typically reads the two files 'prefix'.dat and 'prefix'.pin, although in some cases one or both of these files may not be necessary.

If `logfile=TRUE`, output (including error messages, if any) will appear in 'prefix'.log. If `verbose=TRUE`, it will appear in the R console.

Value

Invisibly returns the results of the shell call.

Note

A wrapper function that can be called from a GUI exists as `.win.runAD`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[runMC](#), [makeAD](#), [cleanAD](#)

runMC

*Run an Executable Binary File in MCMC Mode***Description**

Run the executable binary file 'prefix'.exe, created by makeAD, to generate MCMC simulations.

Usage

```
runMC(prefix, nsims=2000, nthin=20, outsuff=".mc.dat",
      logfile=FALSE, add=TRUE, verbose=TRUE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
nsims	numeric scalar indicating number of MCMC simulations to perform.
nthin	numeric scalar indicating the sampling rate or thinning of the nsims MCMC simulations to report.
outsuff	string name suffix of the MCMC output data file.
logfile	logical: if TRUE, create a log file of the messages from the shell call.
add	logical: if TRUE, append shell call messages to an existing log file.
verbose	logical: if TRUE, report the shell call and its messages to the R console.

Details

This function runs 'prefix'.exe twice, first with the arguments -mcmc 'nsims' -mcsave 'nthin' and second with the argument -mceval. By default, output goes to the file 'prefix'.mc.dat, although a user can specify a different output suffix.

To see this function in action, use the PBSadmb GUI with the example vonb or simpleMC.

Value

Invisibly returns the results of the shell call.

Note

A wrapper function that can be called from a GUI exists as .win.runMC.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[runAD](#), [makeAD](#), [cleanAD](#)

setADMBPath	<i>Creates the ADMB Options List</i>
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Description

Creates a global list object detailing the pathways to the ADMB directory, the GCC bin, and the user's preferred text editor.

Usage

```
setADMBPath(admbpath, gccpath, editor)
```

Arguments

admbpath	explicit path to the user's ADMB directory.
gccpath	explicit path to the user's GCC bin (C-compiler) directory.
editor	explicit path and program to use for editing text.

Value

Creates a global, hidden list object called `.ADopts`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#), [writeADopts](#)

showADargs	<i>Show All Arguments for an ADMB Executable</i>
------------	--

Description

Show all arguments available for an ADMB executable in the default text editor.

Usage

```
showADargs(prefix, ed=TRUE)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
ed	logical: if TRUE, write the ADMB arguments to a file and view them with the text editor, else display the arguments on the R console.

Value

Invisibly returns the argument list.

Note

A wrapper function that can be called from a GUI exists as `.win.showADargs`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[editADfile](#), [runAD](#)

startLog	<i>Start a Log File</i>
----------	-------------------------

Description

Start a log file by removing any previous version and appending header information.

Usage

```
startLog(prefix)
```

Arguments

prefix	string name prefix of the ADMB project (e.g., "vonb").
--------	--

Value

No explicit value returned. Writes header lines into a log file `'prefix'.log`.

Note

A wrapper function that can be called from a GUI exists as `.win.startLog`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[appendLog](#), [editADfile](#)

`writeADopts`*Writes the ADMB Options List from Memory to a File*

Description

Writes the global ADMB options list to a file in 'PBS' format (see `PBSmodelling::writeList`).

Usage

```
writeADopts(optfile="ADopts.txt")
```

Arguments

`optfile` string name of the intended output file.

Value

Returns `opts` invisibly. Writes the options list object to an ASCII file.

Note

A wrapper function that can be called from a GUI exists as `.win.writeADopts`.

Author(s)

Jon T. Schnute, Pacific Biological Station, Nanaimo BC, Canada

See Also

[makeADopts](#), [readADopts](#)

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