Package ‘inlinedocs’

February 20, 2015

Title Convert inline comments to documentation
Type Package
Version 2013.9.3
Description Generates Rd files from R source code with comments.
   The main features of the default syntax are that
   (1) docs are defined in comments near the relevant code,
   (2) function argument names are not repeated in comments, and
   (3) examples are defined in R code, not comments.
   It is also easy to define a new syntax.

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LazyLoad yes
Encoding UTF-8
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Description

Generates Rd files from R source code with comments. The main features of the default syntax are that (1) docs are defined in comments near the relevant code, (2) function argument names are not repeated in comments, and (3) examples are defined in R code, not comments. It is also easy to define a new syntax.

Details

Package: inlinedocs
Title: Convert inline comments to documentation
Type: Package
Version: 2013.9.3
Authors@R: c(person("Toby Dylan","Hocking",role=c("aut","cre"), email="toby@sg.cs.titech.ac.jp"), person("Keith","Ponting"), person("Thomas","Wutzler"), person("Philippe","Grosjean"), person("Markus","Müller"), person("R Core Team",role=c("ctb","cph")))
URL: http://inlinedocs.r-forge.r-project.org
Depends: methods, utils, R (>= 2.9)
License: GPL-2 | GPL-3
LazyLoad: yes
Encoding: UTF-8

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

apply.parsers

Description

Parse code to r objs, then run all the parsers and return the documentation list.

Usage

apply.parsers(code, parsers = default.parsers, verbose = FALSE, ...)

apply.parsers
Arguments

- `code` Character vector of code lines.
- `parsers` List of Parser Functions.
- `verbose` Echo names of Parser Functions?
- ... Additional arguments to pass to Parser Functions.

Value

A list of extracted documentation from code.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

Description

combine lists or character strings

Usage

`combine(x, y)`

Arguments

- `x`
- `y`

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
combine.character

Description
combine character strings by pasting them together

Usage
```r
## S3 method for class 'character'
combine(x, y)
```

Arguments
- **x**
- **y**

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

combine.list

Description
combine lists by adding elements or adding to existing elements

Usage
```r
## S3 method for class 'list'
combine(x, y)
```

Arguments
- **x**
- **y**

Value
A list, same type as x, but with added elements from y.

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
Description
combine NULL objects.

Usage
## S3 method for class 'NULL'
combine(x, y)

Arguments
x
y

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

decomment
decomment

Description
Remove comment prefix and join lines of code to form a documentation string.

Usage
decomment(comments)

Arguments
comments Character vector of prefixed comment lines.

Value
String without prefixes or newlines.

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
default.parsers

Description
List of parsers to use by default with package.skeleton.dx.

Usage
default.parsers

descfile.names
descfile names

Description
Names of Parser Functions that operate on the desc arg.

Usage
descfile.names

do.not.generate
do not generate

Description
Make a Parser Function used to indicate that certain Rd files should not be generated.

Usage
do.not.generate(...)

Arguments
... Character strings indicating Rd files without the .Rd suffix.

Value
A Parser Function that will delete items from the outer Documentation List.

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
Examples

```r
silly.pkg <- system.file("silly", package="inlinedocs")
owd <- setwd(tempdir())
file.copy(silly.pkg,".",recursive=TRUE)

## define a custom Parser Function that will not generate some Rd
## files
custom <- do.not.generate("silly-package","Silly-class")
parsers <- c(default.parsers,list(exclude=custom))

## At first, no Rd files in the man subdirectory.
man.dir <- file.path("silly","man")
dir(man.dir)

## Running package.skeleton.dx will generate bare-bones files for
## those specified in do.not.generate, if they do not exist.
package.skeleton.dx("silly",parsers)
Rd.files <- c("silly-package.Rd","Silly-class.Rd","silly.example.Rd")
Rd.paths <- file.path(man.dir,Rd.files)
stopifnot(all(file.exists(Rd.paths)))

## Save the modification times of the Rd files
old <- file.info(Rd.paths)$mtime

## make sure there is at least 2 seconds elapsed, which is the
## resolution for recording times on windows file systems.
Sys.sleep(4)

## However, it will NOT generate Rd for files specified in
## do.not.generate, if they DO exist already.
package.skeleton.dx("silly",parsers)
mtimes <- data.frame(old,new=file.info(Rd.paths)$mtime)
rownames(mtimes) <- Rd.files
mtimes$changed <- mtimes$old != mtimes$new
print(mtimes)
stopifnot(mtimes["silly-package.Rd","changed"]==FALSE)
stopifnot(mtimes["Silly-class.Rd","changed"]==FALSE)
stopifnot(mtimes["silly.example.Rd","changed"]==TRUE)

unlink("silly",recursive=TRUE)
setwd(owd)
```

Description

The `DocLink` class provides the basis for hooking together documentation of related classes/functions/objects. The aim is that documentation sections missing from the child are inherited from the parent class.
Objects from the Class

Objects can be created by calls of the form `new(DocLink ...)`

Slots

- **name**: (character) name of object
- **created**: (character) how created
- **parent**: (character) parent class or NA
- **code**: (character) actual source lines
- **description**: (character) preceding description block

Methods

No methods defined with class "DocLink" in the signature.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

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extra.code.docs

Extract documentation from code chunks

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Description

Parse R code to extract inline documentation from comments around each function. These are not able to be retrieved simply by looking at the "source" attribute. This is a Parser Function that can be used in the parser list of package.skeleton.dx(). TODO: Modularize this into separate Parsers Functions for S4 classes, prefixes, ##«blocks, etc. Right now it is not very clean!

Usage

`extra.code.docs(code, objs, ...)`

Arguments

- **code**: Code lines in a character vector containing multiple R objects to parse for documentation.
- **objs**: The objects defined in the code.
- **...**: ignored

Value

named list of lists, one for each object to document.
extract.docs.file  

Description

Apply all parsers relevant to extract info from just 1 code file.

Usage

extract.docs.file(f, parsers = NULL, ...)

Arguments

f  
File name of R code to read and parse.

parsers  
Parser Functions to use to parse the code and extract documentation.

...  
Other arguments to pass to Parser Functions.

extract.docs.setClass  

S4 class inline documentation

Description

Using the same conventions as for functions, definitions of S4 classes in the form setClass("classname",...) are also located and scanned for inline comments.

Usage

extract.docs.setClass(doc.link)
extract.file.parse

Arguments

doc.link DocLink object as created by extract.file.parse. Note that source statements are ignored when scanning for class definitions.

Details

Extraction of S4 class documentation is currently limited to expressions within the source code which have first line starting with setClass("classname"). These are located from the source file (allowing also for white space around the setClass and (). Note that "classname" must be a quoted character string; expressions returning such a string are not matched.

For class definitions, the slots (elements of the representation list) fill the role of function arguments, so may be documented by ## comments on the same line or ### comments at the beginning of the following line.

If there is no explicit title on the first line of setClass, then one is made up from the class name.

The class definition skeleton includes an Objects from the Class section, to which any ##details documentation chunks are written. It is given a vanilla content if there are no specific ##details documentation chunks.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

 extract.file.parse  File content analysis

Description

Using the base parse function, analyse the file to link preceding "prefix" comments to each active chunk. Those comments form the default description for that chunk. The analysis also looks for S4 class "setClass" calls and R.oo setConstructorS3 and setMethodS3 calls in order to link the documentation of those properly.

Usage

extract.file.parse(code)

Arguments

code Lines of R source code in a character vector - note that any nested source statements are ignored when scanning for class definitions.
extract.xxx.chunks

Details

If the definition chunk does not contain a description, any immediately preceding sequence consecutive "prefix" lines will be used instead.

Class and method definitions can take several forms, determined by expression type:

- **assignment** (\(<\)-) Ordinary assignment of value/function;
- **setClass** Definition of S4 class;
- **setConstructorS3** Definition of S3 class using R.oo package;
- **setMethodS3** Definition of method for S3 class using R.oo package.

Additionally, the value may be a name of a function defined elsewhere, in which case the documentation should be copied from that other definition. This is handled using the concept of documentation links.

The **setMethodS3** calls introduce additional complexity: they will define an additional S3 generic (which needs documentation to avoid warnings at package build time) unless one already exists. This also is handled by "linking" documentation. A previously unseen generic is linked to the first defining instances, subsequent definitions of that generic also link back to the first defining instance.

Value

Returns an invisible list of .DocLink objects.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

---

**extract.xxx.chunks** *Extract documentation from a function*

Description

Given source code of a function, return a list describing inline documentation in that source code.

Usage

```
extract.xxx.chunks(src, name.fun = "(unnamed function)", ...)
```

Arguments

- **src** The source lines of the function to examine, as a character vector.
- **name.fun** The name of the function/chunk to use in warning messages.
- **...** ignored.
Details

For simple functions/arguments, the argument may also be documented by appending `##<<` comments on the same line as the argument name. Mixing this mechanism with `###` comment lines for the same argument is likely to lead to confusion, as the `###` lines are processed first.

Additionally, consecutive sections of `##` comment lines beginning with `###`<xxx<< (where xxx is one of the fields: alias, details, keyword, references, author, note,seealso, value, title or description) are accumulated and inserted in the relevant part of the Rd file.

For value, title, description and function arguments, these append to any text from "prefix" (^### ) comment lines, irrespective of the order in the source.

When documenting S4 classes, documentation from details sections will appear under a section Objects from the Class. That section typically includes information about construction methods as well as other description of class objects (but note that the class Slots are documented in a separate section).

Each separate extra section appears as a new paragraph except that:

- empty sections (no matter how many lines) are ignored;
- alias and keyword sections have special rules;
- description should be brief, so all such sections are concatenated as one paragraph;
- title should be one line, so any extra title sections are concatenated as a single line with spaces separating the sections.

As a special case, the construct `##describe<<` causes similar processing to the main function arguments to be applied in order to construct a describe block within the documentation, for example to describe the members of a list. All subsequent "same line" `##<<` comments go into that block until terminated by a subsequent `###<<` line.

Such regions may be nested, but not in such a way that the first element in a describe is another describe. Thus there must be at least one `##<<` comment between each pair of `##describe<<` comments.

When nested describe blocks are used, a comment-only line with `##end<<` terminates the current level only; any other valid `###<<` line terminates all open describe blocks.

Value

Named list of character strings extracted from comments. For each name N we will look for N{...} in the Rd file and replace it with the string in this list (implemented in modify.Rd.file).

Note

alias extras are automatically split at new lines.

keyword extras are automatically split at white space, as all the valid keywords are single words.

The "value" section of a Rd file is implicitly a describe block and `##value<<` acts accordingly. Therefore it automatically enables the describe block itemization (`##« after list entries).

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
Description

Copied from R-3.0.1, to support findGeneric.

Usage

findGeneric(fname, envir)

Arguments

fname
envir

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

Description

Copied from R-3.0.1, to support fixPackageFileNames.

Usage

fixPackageFileNames(list)

Arguments

list

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
forall

Description

For each object in the package that satisfies the criterion checked by subfun, parse source using
FUN and return the resulting documentation list.

Usage

forall(FUN, subfun = function(x) TRUE)

Arguments

FUN Function to apply to each element in the package.
subfun Function to select subsets of elements of the package, such as is.function. sub-
fun(x)==TRUE means FUN will be applied to x and the result will be returned.

Value

A Parser Function.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler
[aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

forall.parsers

Description

List of Parser Functions that can be applied to any object.

Usage

forall.parsers
getKnownS3generics

forfun

Description
For each function in the package, do something.

Usage
forfun(FUN)

Arguments
FUN

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

forfun.parsers

Description
Parsers for each function that are constructed automatically. This is a named list, and each element is a parser function for an individual object.

Usage
forfun.parsers

getKnownS3generics

Description
Copied from R-3.0.1, to support getKnownS3generics.

Usage
getKnownS3generics()

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
**getDescription**

Description

Extract a function’s source code.

Usage

ggetSource(fun.obj)

Arguments

fun.obj  A function.

Value

Source code lines as a character vector.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

**get_internal_S3_generics**

Description

Copied from R-3.0.1, to support getKnownS3generics.

Usage

gInternal_S3_generics(primitive = TRUE)

Arguments

primitive

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
get_S3_primitive_generics

Description
Copied from R-3.0.1, to support getKnownS3generics.

Usage
get_S3_primitive_generics(include_group_generics = TRUE)

Arguments
include_group_generics

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

is_primitive_in_base

Description
Copied from R-3.0.1, to support getKnownS3generics.

Usage
is_primitive_in_base(fname)

Arguments
fname

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
Description

Figure out what the whitespace preceding the example code is, and then delete that from every line.

Usage

`kill.prefix.whitespace(ex)`

Arguments

- `ex`: character vector of example code lines.

Value

Character vector of code lines with preceding whitespace removed.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp>, Keith Ponting, Thomas Wutzler, Philippe Grosjean, Markus Müller, R Core Team

Description

Determines whether a function name looks like an S3 generic function

Usage

`leadingS3generic(name, env, ...)`

Arguments

- `name`: name of function
- `env`: environment to search for additional generics
- `...`: ignored here
Details

This function is one of the default parsers, but exposed as possibly of more general interest. Given a function name of the form x.y.z it looks for the generic function x applying to objects of class y.z and also for generic function x.y applying to objects of class z.

Assumes that the first name which matches any known generics is the target generic function, so if both x and x.y are generic functions, will assume generic x applying to objects of class y.z.

Value

If a matching generic found returns a list with a single component:

- `.s3method` a character vector containing generic name and object name.

If no matching generic functions are found, returns an empty list.

Author(s)

Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

Description

List of parser functions that operate on single objects. This list is useful for testing these functions.

Usage

`lonely`

Examples

```r
f <- function # title
### description
(x, ### arg x
 y
### arg y
)(
  ###value<< a list with elements
  list(x=x, ### original x value
       y=y, ### original y value
       sum=x+y) ### their sum
###end<<
)
src <- getSource(f)
lonely$extract.xxx.chunks(src)
lonely$prefixed.lines(src)
```
make.package.and.check

Description
Assemble some R code into a package and process it using R CMD check, stopping with an error if the check resulted in any errors or warnings.

Usage
make.package.and.check(f, parsers = default.parsers, verbose = TRUE)

Arguments
f R code file name from which we will make a package
parsers Parsers to use to make the package documentation.
verbose print the check command line?

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]

modify.Rd.file

Description
Add inline documentation from comments to an Rd file automatically-generated by package.skeleton.

Usage
modify.Rd.file(N, pkg, docs)

Arguments
N Name of function/file to which we will add documentation.
pkg Package name.
docs Named list of documentation in extracted comments.

Author(s)
Toby Dylan Hocking <toby@sg.cs.titech.ac.jp> [aut, cre], Keith Ponting [aut], Thomas Wutzler [aut], Philippe Grosjean [aut], Markus Müller [aut], R Core Team [ctb, cph]
---

**non.descfile.names**  
*non descfile names*

**Description**  
Names of Parser Functions that do NOT use the desc arg.

**Usage**  
non.descfile.names

---

**nondesc.parsers**  
*nondesc parsers*

**Description**  
Parsers that operate only on R code, independently of the description file.

**Usage**  
nondesc.parsers

---

**package.skeleton.dx**  
*Package skeleton deluxe*

**Description**  
Generates Rd files for a package based on R code and DESCRIPTION metadata. After inspecting the specified R code files to find inline documentation, it calls the standard package.skeleton function, which creates bare Rd files. The inline documentation is added to these Rd files and then these files are copied to pkgdir/man, possibly overwriting the previous files there.

**Usage**  
package.skeleton.dx(pkmdir = ".", parsers = NULL, namespace = FALSE, excludePattern = FALSE, ...)

package.skeleton.dx

Arguments

pkgdir Package directory where the DESCRIPTION file lives. Your code should be in pkgdir/R. We will setwd to pkgdir/R for the duration of the function, then switch back to where you were previously.

parsers List of Parser functions, which will be applied in sequence to extract documentation from your code. Default NULL means to first search for a definition in the variable "parsers" in pkgdir/R/inlinedocs.R, if that file exists. If not, we use the list defined in options("inlinedocs.parsers"), if that is defined. If not, we use the package global default in the variable default.parsers.

namespace A logical indicating whether a NAMESPACE file should be generated for this package. If TRUE, all objects whose name starts with a letter, plus all S4 methods and classes are exported.

excludePattern A regular expression matching the files that are not to be processed e.g. because inlinedocs can not handle them yet (like generic function definitions)

... Parameters to pass to Parser Functions.

Author(s)

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Examples

```r
owd <- setwd(tempdir())

## get the path to the silly example package that is provided with
## package inlinedocs
testPackagePath <- file.path(system.file(package="inlinedocs"),"silly")
## copy example project to the current unlocked workspace that can
## be modified
file.copy(testPackagePath,".",recursive=TRUE)

## generate documentation .Rd files for this package
package.skeleton.dx("silly")

## check the package to see if generated documentation passes
## without WARNINGs.
if(interactive()){
  cmd <- sprintf("%s CMD check --as-cran silly",file.path(R.home("bin"), "R"))
  print(cmd)
  checkLines <- system(cmd,intern=TRUE)
  warnLines <- grep("WARNING",checkLines,value=TRUE)
  if(length(warnLines)>0)
    writeLines(checkLines)
    cat("\n\nLines with WARNING:\n")
    print(warnLines)
  ## disable due to bug in R CMD check:
  ## https://bugs.r-project.org/bugzilla3/show_bug.cgi?id=14875
  ## stop("WARNING encountered in package check!")
```
prefixed lines

Description

Prefix for code comments used with grep and gsub.

Usage

prefix

prefixed lines

Description

The primary mechanism of inline documentation is via consecutive groups of lines matching the specified prefix regular expression "## " (i.e. lines beginning with "## ") are collected as follows into documentation sections:

- **description** group starting at line 2 in the code
- **arguments** group following each function argument
- **value** group ending at the penultimate line of the code

These may be added to by use of the ##<< constructs described below.

Usage

prefixed.lines(src, ...)

Arguments

src

... 

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print.allfun

Examples

test <- function
  ### the description
  (x,
   ### the first argument
   y ###<< another argument
  )
  5
  ### the return value
  ###seealso<< foobar
  }
src <- getSource(test)
prefixed.lines(src)
eextract.xxx.chunks(src)

print.allfun

Description

Print method for functions constructed using forall.

Usage

## S3 method for class 'allfun'
print(x, ...)

Arguments

x

...

Author(s)

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replace.one

removeAliasesfromRd.file

Description
remove aliases to methodnames from the Rd file of a class automatically-generated by package.skeleton.

Usage
removeAliasesfromRd.file(N, pkg, code)

Arguments
N Name of function/file to which we will add documentation.
pkg Package name.
code The code of the package

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replace.one

Description
Do find and replace for one element of an inner documentation list on 1 Rd file.

Usage
replace.one(torep, REP, txt)

Arguments
torep tag to find.
REP contents of tag to put inside.
txt text in which to search.

Author(s)
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**Description**

For unit tests, this is an easy way of getting a text representation of the list result of `extract.docs.file`.

**Usage**

```r
save.test.result(f)
```

**Arguments**

- `f`: R code file with inlinedocs to process with `extract.docs.file`.

**Author(s)**

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**Description**

Check an R code file with inlinedocs to see if the `extract.docs.file` parser accurately extracts all the code inside! The code file should contain a variable `.result` which is the documentation list that you should get when you apply `extract.docs.file` to the file. We check for identity of elements of elements of the list, so the order of elements should not matter, and thus this should be a good robust unit test.

**Usage**

```r
test.file(f, verbose = TRUE)
```

**Arguments**

- `f`: File name of R code file with inlinedocs to parse and check.
- `verbose`: Show output?

**Author(s)**

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**See Also**

- `save.test.result`
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