

Package ‘scidb’

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Type Package

Title An R Interface to SciDB

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Description An R interface to the 'SciDB' array database <<http://scidb.org>>.

BugReports <https://github.com/Paradigm4/SciDBR/issues>

URL <http://paradigm4.github.io/SciDBR>

VignetteBuilder knitr

Depends R (>= 3.0.0), bit64

Imports curl, data.table, digest, methods, openssl

Suggests Matrix, knitr (>= 1.8)

License AGPL-3

LazyLoad yes

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scidb-package	<i>SciDB/R Interface</i>
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Description

SciDB/R Interface

Package options

```
options(scidb.prefix=NULL) # Default shim port and host. options(scidb.default_shim_port=8080L)
options(scidb.default_shim_host="localhost") # How to download arrays and their coordinates. Set
scidb.unpack=FALSE # to use apply, which can be faster in some cases when used with aio. options(scidb.unpack=FALSE) # Disable SSL certificate host name checking by default. This is im-
portant mostly # for Amazon EC2 where hostnames rarely match their DNS names. If you enable
this # then the shim SSL certificate CN entry *must* match the server host name for the # encrypted
session to work. Set this TRUE for stronger security (help avoid MTM) # in SSL connections. options(scidb.verifyhost=FALSE) # List of special DDL operators options(scidb.ddl=c("create_array",
"remove", "rename"))
```

See Also

[scidb](#), [iquery](#)

aflhelp*Display SciDB AFL operator documentation*

Description

Display SciDB AFL operator documentation

Usage

```
aflhelp(topic, db)
```

Arguments

- | | |
|-------|---|
| topic | an afl object from a SciDB database connection, or optionally a character string name |
| db | optional database connection from scidbconnect (only needed when topic is a character string) |

Value

displays help

Examples

```
## Not run:  
d <- scidbconnect()  
aflhelp("list", d)      # explicitly look up a character string  
help(d$list)           # same thing via R's \code{help} function  
  
## End(Not run)
```

as.R*Download SciDB data to R*

Description

Download SciDB data to R

Usage

```
as.R(x, only_attributes = FALSE, binary = TRUE)
```

Arguments

x	a scidb object (a SciDB array or expression)
only_attributes	optional logical argument, if TRUE do not download SciDB dimensions (see note)
binary	optional logical value, set to FALSE to download data using text format (useful for some unsupported SciDB types)

Value

An R [data.frame](#)

Note

This convenience function is equivalent to running `iquery(db, x, return=TRUE)` for a SciDB connection object db.

The `only_attributes=TRUE` option only works with binary transfers, and if specified will set `binary=TRUE`. Beware of the `only_attributes=TRUE` setting—SciDB may return data in arbitrary order.

SciDB values are always returned as R data frames. SciDB scalar types are converted to corresponding R types as follows:

- double -> double
- int64 -> integer64
- uint64 -> double
- uint32 -> double
- int32 -> integer
- int16 -> integer
- unit16 -> integer
- int8 -> integer
- uint8 -> integer
- bool -> logical
- string -> character
- char -> character
- binary -> raw
- datetime -> Date

See Also

[as.scidb](#)

Examples

```
## Not run:
db <- scidbconnect()
x <- scidb(db, "build(<v:double>[i=1:5], sin(i))")
as.R(x)
## i          v
## 1  0.8414710
## 2  0.9092974
## 3  0.1411200
## 4 -0.7568025
## 5 -0.9589243

as.R(x, only_attributes=TRUE)
##          v
## 0.8414710
## 0.9092974
## 0.1411200
## -0.7568025
## -0.9589243

## End(Not run)
```

as.scidb

Upload R data to SciDB

Description

Upload R data to SciDB

Usage

```
as.scidb(db, x, name, start, gc = TRUE, ...)
```

Arguments

db	a scidb database connection returned from scidbconnect
x	an R data frame, raw value, Matrix, matrix, or vector object
name	a SciDB array name to use
start	starting SciDB integer coordinate index (does not apply to data frames)
gc	set to FALSE to disconnect the SciDB array from R's garbage collector
...	other options, see df2scidb

Value

A scidb object

Note

Supported R objects include data frames, scalars, vectors, dense matrices, and double-precision sparse matrices of class `CsparseMatrix`. Supported R scalar types and their resulting SciDB types are:

- `integer` -> `int32`
- `logical` -> `int32`
- `character` -> `string`
- `double` -> `double`
- `integer64` -> `int64`
- `raw` -> `binary`
- `Date` -> `datetime`

R factor values are converted to their corresponding character levels.

See Also

[as.R](#)

`at_least`

Returns TRUE if version string x is greater than or equal to than version y

Description

Returns TRUE if version string x is greater than or equal to than version y

Usage

`at_least(x, y)`

Arguments

- | | |
|----------------|---|
| <code>x</code> | version string like "12.1", "15.12", etc. (non-numeric ignored) |
| <code>y</code> | version string like "12.1", "15.12", etc. (non-numeric ignored) |

Value

logical TRUE if x is greater than or equal to y

cov,scidb-method	<i>Covariance matrix</i> This function is more limited than R's default cov function. It can only compute a covariance matrix from a data matrix without any missing value handling by the procedure (in R notation) S0 <- sweep(x, 2, colMeans(x)), crossprod(S0)/(nrow(S0) - 1) # (covariance matrix result)
------------------	---

Description

Covariance matrix This function is more limited than R's default cov function. It can only compute a covariance matrix from a data matrix without any missing value handling by the procedure (in R notation)
S0 <- sweep(x, 2, colMeans(x)), crossprod(S0)/(nrow(S0) - 1) # (covariance matrix result)

Usage

```
## S4 method for signature 'scidb'
cov(x, y = NULL, use = "everything", method = c("pearson", "kendall", "spearman"))
```

Arguments

- | | |
|--------|---|
| x | a 2-d scidb array with a single numeric attribute |
| y | UNUSED, limited to correlation matrix in the SciDB case |
| use | UNUSED, limited to "everything" in the SciDB case |
| method | UNUSED, limited to "pearson" in the SciDB case |

Value

covariance matrix of x (as a SciDB array)

dimnames.scidb	<i>Names of array dimensions</i>
----------------	----------------------------------

Description

Names of array dimensions

Usage

```
## S3 method for class 'scidb'
dimnames(x)
```

Arguments

- | | |
|---|--------------------|
| x | scidb array object |
|---|--------------------|

Value

a vector of SciDB array dimension names

getpwd

Simple utility to interactively enter a password without showing it on the screen

Description

Simple utility to interactively enter a password without showing it on the screen

Usage

```
getpwd(prompt = "Password:")
```

Arguments

prompt	a text prompt to display, defaults to "Password:"
---------------	---

help,operator-method *AFL operator help*

Description

AFL operator help

Usage

```
## S4 method for signature 'operator'
help(topic)
```

Arguments

topic	afl operator
--------------	--------------

Value

help summary

iquery*Run a SciDB query, optionally returning the result.*

Description

Run a SciDB query, optionally returning the result.

Usage

```
iquery(db, query, return = FALSE, binary = TRUE, ...)
```

Arguments

db	a scidb database connection from scidbconnect
query	a single SciDB query string or scidb array object
return	if TRUE, return the result
binary	set to FALSE to read result from SciDB in text form
...	additional options passed to <code>read.table</code> when <code>binary=FALSE</code> , or optional result schema when <code>binary=TRUE</code> (see note below).

Note

When `query` is an arbitrary AFL query string and `binary=TRUE`, optionally specify `schema` with a valid result array schema to skip an extra metadata lookup query (see [scidb](#)).

Setting `return=TRUE` wraps the AFL query expression with a SciDB save operator, saving the data on the SciDB server in either binary or text format depending on the value of the `binary` parameter. Please note that some AFL expressions may not be "saved" using the AFL save operator, including for instance the AFL `create_array` operator. Trying to return the result of such a SciDB expression will result in a run-time error.

See Also

[scidb as.R](#)

Examples

```
## Not run:  
db <- scidbconnect()  
iquery(db, "build(<v:double>[i=1:5], sin(i))", return=TRUE)  
## i          v  
## 1  0.8414710  
## 2  0.9092974  
## 3  0.1411200  
## 4 -0.7568025  
## 5 -0.9589243  
  
# Use binary=FALSE and additional options to read.table function:
```

```

iquery(db, "build(<val:string>[i=1:3], '[(01),(02),(03)]', true)",xi
      return=TRUE, binary=FALSE, colClasses=c("integer", "character"))
##   i val
## 1 1 01
## 2 2 02
## 3 3 03

## End(Not run)

```

ls,afl-method *List contents of a SciDB database*

Description

List contents of a SciDB database

Usage

```
## S4 method for signature 'afl'
ls(name)
```

Arguments

name	afl SciDB connection object from scidbconnect
------	---

Value

a `data.frame` listing the contents of the database

names.scidb *SciDB dimension and attribute names*

Description

SciDB dimension and attribute names

Usage

```
## S3 method for class 'scidb'
names(x)
```

Arguments

x	scidb array object
---	--------------------

Value

Character vector of names

operators

Base SciDB operators

Description

Base SciDB operators as of SciDB version 16.9

Usage

```
data(operators)
```

Format

A data frame with 4 variables, name, signature, help.

Source

Paradigm4 <http://paradigm4.com>

print,afl-method

Print a summary of a afl SciDB database connection object

Description

Print a summary of a afl SciDB database connection object

Usage

```
## S4 method for signature 'afl'  
print(x)
```

Arguments

x afl object

Value

printed object summary

print,scidb-method *Print a summary of a scidb object*

Description

Print a summary of a scidb object

Usage

```
## S4 method for signature 'scidb'  
print(x)
```

Arguments

x a scidb object

Value

printed object summary

print.afl *Print a summary of a afl SciDB database connection object*

Description

Print a summary of a afl SciDB database connection object

Usage

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

Arguments

x afl object
... optional arguments (not used)

Value

printed object summary

schema*SciDB array schema*

Description

SciDB array schema

Usage

```
schema(x, what = c("schema", "attributes", "dimensions"))
```

Arguments

- | | |
|------|--|
| x | a <code>scidb</code> array object |
| what | optional schema subset (subsets are returned in data frames; partial argument matching is supported) |

Value

character-valued SciDB array schema

Examples

```
## Not run:
s <- scidbconnect()
x <- scidb(s, "build(<v:double>[i=1:10,2,0,j=0:19,1,0],0)")
schema(x)
# [1] "<v:double> [i=1:10:0:2; j=0:19:0:1]"
schema(x, "attributes")
#   name    type nullable
#1   v    double     TRUE
schema(x, "dimensions")
  name start end chunk overlap
#1   i      1   10      2       j
#2   0      0   19      1       0
## End(Not run)
```

scidb*Create an R reference to a SciDB array or expression.*

Description

Create an R reference to a SciDB array or expression.

Usage

```
scidb(db, name, gc = FALSE, schema)
```

Arguments

<code>db</code>	scidb connection object from <code>scidbconnect</code>
<code>name</code>	a character string name of a stored SciDB array or a valid SciDB AFL expression
<code>gc</code>	a logical value, TRUE means connect the SciDB array to R's garbage collector
<code>schema</code>	optional SciDB array schema, if specified avoid an extra metadata query to determine array schema. Use this option with care, the schema must exactly match the SciDB array result.

Value

a scidb object

`scidbconnect`

Connect to a SciDB database

Description

Connect to a SciDB database

Usage

```
scidbconnect(host = getOption("scidb.default_shim_host", "127.0.0.1"),
            port = getOption("scidb.default_shim_port", 8080L), username, password,
            auth_type = c("scidb", "digest"), protocol = c("http", "https"),
            int64 = FALSE, doc)
```

Arguments

<code>host</code>	optional host name or I.P. address of a SciDB shim service to connect to
<code>port</code>	optional port number of a SciDB shim service to connect to
<code>username</code>	optional authentication username
<code>password</code>	optional authentication password
<code>auth_type</code>	optional SciDB authentication type
<code>protocol</code>	optional shim protocol type
<code>int64</code>	logical value, if TRUE then preserve signed 64-bit SciDB integers as R integer64 values from the bit64 package. Otherwise, 64-bit integers from SciDB are converted to R double values, possibly losing precision.
<code>doc</code>	optional AFL operator/macro documentation (see notes)

Value

A scidb connection object. Use \$ to access AFL operators and macros, ls() on the returned object to list SciDB arrays, and names() on the returned object to list all available AFL operators and macros.

Note

Use the optional username and password arguments with auth_type set to "digest" to use HTTP digest authentication (see the shim documentation to configure this). Digest authentication may use either "http" or "https" selected by the protocol setting. Set auth_type = "scidb" to use SciDB authentication, which only works over "https".

Use the returned SciDB connection object (of class afl) with other package functions to interact with SciDB arrays. Apply R's `ls` function on the returned value to see a list of arrays. The returned value contains a list of available SciDB AFL language operators and macro names. Use the dollar-sign function to access those functions.

The optional doc argument may be a three-column data frame with character-valued columns name, signature, and help containing AFL operator names, function signatures, and help strings, respectively. See `'data("operators", package="scidb")'` for an example.

All arguments support partial matching.

See Also

[scidb_prefix](#)

Examples

```
## Not run:  
db <- scidbconnect()  
  
# SciDB 15.12 authentication example (using shim's default HTTPS port 8083)  
db <- scidbconnect(user="root", password="Paradigm4",  
                    auth_type="scidb", port=8083, protocol="https")  
  
# List available AFL operators  
names(db)  
  
# List arrays  
ls(db)  
  
# Explicitly upload an R matrix to SciDB:  
x <- as.scidb(db, matrix(rnorm(20), 5))  
# Implicitly do the same as part of an AFL expression  
y <- db$join(x, as.scidb(matrix(1:20, 5)))  
print(y)  
  
as.R(y) # Download a SciDB array to R.  
  
## End(Not run)
```

<code>scidb_prefix</code>	<i>Register an AFL prefix expression</i>
---------------------------	--

Description

SciDB AFL statements are normally executed in a stateless query context. Use `scidb_prefix` to create compound AFL expressions useful in some circumstances.

Usage

```
scidb_prefix(db, expression = NULL)
```

Arguments

<code>db</code>	a scidb database connection returned from scidbconnect
<code>expression</code>	a valid AFL expression to be issued prior to, and in the same context as all subsequent query expressions issued to the database corresponding to <code>db</code> . Set <code>expression=NULL</code> to remove the prefix expression.

Value

A new SciDB database connection object with the prefix set.

Note

This is mostly useful for setting namespaces, see the examples.

Examples

```
## Not run:  
library(scidb)  
db <- scidbconnect()  
ls(db)  
new_db <- scidb_prefix(db, "set_role('functionary')")  
ls(new_db)  
  
## End(Not run)
```

show,afl-method	<i>Print a summary of a afl object</i>
-----------------	--

Description

Print a summary of a afl object

Usage

```
## S4 method for signature 'afl'  
show(object)
```

Arguments

object afl object

Value

printed object summary

show,scidb-method	<i>Print a summary of a scidb object</i>
-------------------	--

Description

Print a summary of a scidb object

Usage

```
## S4 method for signature 'scidb'  
show(object)
```

Arguments

object a scidb object

Value

printed object summary

store	<i>Evaluate an expression to scidb or scidb objects</i>
-------	---

Description

Force evaluation of an expression that yields a scidb or scidb object, storing the result to a SciDB array when eval=TRUE.

Usage

```
store(db, expr, name, eval = TRUE, gc = TRUE, temp = FALSE)
```

Arguments

db	scidb connection object from scidbconnect
expr	a quoted SciDB expression scidb object
name	(character) optional SciDB array name to store result to
eval	FALSE do not evaluate the expression in SciDB (leave as a view)
gc	(logical) optional, when TRUE tie result to R garbage collector
temp	(logical, optional), when TRUE store as a SciDB temp array

%as%,scidb-method	<i>AFL arrary aliasing</i>
-------------------	----------------------------

Description

AFL arrary aliasing

Usage

```
## S4 method for signature 'scidb'
x %as% y
```

Arguments

x	an object of class scidb (a scidb array or expression)
y	alias name

Value

a [scidb](#) object

Note

Use the %as% operator in place of the native AFL "as" operator in AFL expressions written in R.

Examples

```
## Not run:  
db <- scidbconnect()  
x <- scidb(db, "build(<v:double>[i=1:2,1,0], i)")  
x %as% y  
  
## End(Not run)
```

`%as%`*AFL array aliasing*

Description

AFL array aliasing

Usage

```
x %as% y
```

Arguments

x	an object of class scidb (a scidb array or expression)
y	alias name

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