

Package ‘read.dbc’

November 28, 2025

Title Read Data Stored in 'DBC' (Compressed 'DBF') Files

Description Functions for reading and decompressing the 'DBC' (compressed 'DBF') files. Please note that this is the file format used by the Brazilian Ministry of Health ('DATASUS') to publish healthcare datasets. It is not related to the 'Fox-Pro' or 'CANdb' 'DBC' file formats.

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Depends R (>= 3.3.0)

Imports foreign

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URL <https://github.com/danicat/read.dbc>

BugReports <https://github.com/danicat/read.dbc/issues>

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dbc2dbf	<i>Decompress a DBC file</i>
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Description

This function allows you decompress a DBC file. When decompressed, it becomes a regular DBF file.

Usage

```
dbc2dbf(input.file, output.file)
```

Arguments

input.file	The name of the DBC file (including extension)
output.file	The output file name (including extension)

Details

DBC is the extension for compressed DBF files (from the 'XBASE' family of databases). This is a proprietary file format used by the Brazilian government to publish public healthcare data. When decompressed, it becomes a regular DBF file.

Please note that this is the file format is not related to the FoxPro or CANDb DBC file formats.

Value

Return TRUE if succeed, FALSE otherwise.

Author(s)

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Source

The internal C code for dbc2dbf is based on blast decompressor and blast-dbf (see *References*).

References

blast source code in C: <https://github.com/madler/zlib/tree/master/contrib/blast> blast-dbf,
DBC to DBF command-line decompression tool: <https://github.com/eaglebh/blast-dbf>

See Also

[read.dbc](#)

Examples

```
# Input file name
input <- system.file("files/sids.dbc", package = "read.dbc")

# Output file name
output <- tempfile(fileext = ".dbf")

# The call returns TRUE on success
if( dbc2dbf(input.file = input, output.file = output) ) {
  print("File decompressed!")
  # do things with the file
}

file.remove(output) # clean up example, don't do in real life :)
```

dbf2dbc*Compress a standard DBF file into a DBC file (Experimental)*

Description

This function compresses a standard DBF file into a DBC file using the PKWare DCL Implode algorithm. This functionality is experimental and may be subject to change.

Usage

```
dbf2dbc(input, output)
```

Arguments

input	The input DBF file path.
output	The output DBC file path.

Value

TRUE if successful.

Examples

```
## Not run:
# Compress a file
dbf2dbc("data.dbf", "data.dbc")

## End(Not run)
```

`read.dbc`*Read Data Stored in DBC (Compressed DBF) Files*

Description

This function allows you to read a DBC (compressed DBF) file into a data frame.

Usage

```
read.dbc(file, ...)
```

Arguments

<code>file</code>	The name of the DBC file (including extension)
<code>...</code>	Further arguments to be passed to read.dbf

Details

DBC is the extension for compressed DBF files (from the 'XBASE' family of databases). This is a proprietary file format used by the Brazilian government to publish public healthcare data, and it is not related to the FoxPro or CANDb DBC file formats.

The `read.dbc` function will decompress the input DBC file into a temporary DBF file and call [read.dbf](#) from the foreign package to read it into a data frame.

Value

A `data.frame` of the data from the DBC file.

Note

DATASUS is the name of the Department of Informatics of the Brazilian Health System (Sistema Único de Saúde - SUS) and is responsible for publishing public healthcare data in Brazil. Besides the DATASUS, the Brazilian National Agency for Supplementary Health (ANS) also uses this file format for its public data.

This function was tested using files from both DATASUS and ANS to ensure compliance with the format, and hence ensure its usability by researchers.

Neither this project, nor its author, has any association with the Brazilian government.

Author(s)

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See Also

[dbc2dbf](#)

Examples

```
# The 'sids.dbc' file is the compressed version of 'sids.dbf' from the "foreign" package.
file <- system.file("files/sids.dbc", package="read.dbc")
sids <- read.dbc(file)
str(sids)
summary(sids)

# This is a small subset of U.S. NOAA storm database.
file <- system.file("files/storm.dbc", package="read.dbc")
storm <- read.dbc(file)
head(storm)
str(storm)
```

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