

Package ‘childfree’

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Title Access and Harmonize Childfree Demographic Data

Version 0.0.4

Description Reads demographic data from a variety of public data sources, extracting and harmonizing variables useful for the study of childfree individuals. The identification of childfree individuals and those with other family statuses uses Neal & Neal's (2024) ``A Framework for Studying Adults who Neither have Nor Want Children" <doi:10.1177/10664807231198869>; A preprint is available at <doi:10.31234/osf.io/fa89m>.

License GPL-3

Encoding UTF-8

RoxygenNote 7.3.1

Depends R (>= 2.10)

Imports rio, utils, RCurl

Suggests knitr,

VignetteBuilder knitr

URL <https://www.zacharyneal.com/childfree-home>,
<https://github.com/zpneal/childfree>

BugReports <https://github.com/zpneal/childfree/issues>

NeedsCompilation no

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childfree

childfree: Access and harmonize childfree demographic data

Description

Reads demographic data from a variety of public data sources, extracting and harmonizing variables useful for the study of childfree individuals. The identification of childfree individuals and those with other family statuses uses the framework described by Neal & Neal (2024).

Data can be generated from:

- Demographic and Health Surveys data using `dhs()`
- Michigan State University State of the State data using `sooss()`
- US CDC National Survey of Family Growth data using `nsfg()`

An introduction to the package is available using `vignette("childfree")`, and the detailed codebooks generated by these functions are available using `vignette("codebooks")`.

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References

Neal, Z. P. and Neal, J. W. (2024). A framework for studying adults who neither have nor want children. *The Family Journal*, 32, 121-130. Version of record: [doi:10.1177/10664807231198869](https://doi.org/10.1177/10664807231198869)
Preprint: [doi:10.31234/osf.io/fa89m](https://doi.org/10.31234/osf.io/fa89m)

See Also

Useful links:

- <https://www.zacharyneal.com/childfree-home>
- <https://github.com/zpneal/childfree>
- Report bugs at <https://github.com/zpneal/childfree/issues>

dhs	<i>Read and recode Demographic and Health Surveys (DHS) individual data</i>
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Description

Read and recode Demographic and Health Surveys (DHS) individual data

Usage

```
dhs(files, extra.vars = NULL, progress = TRUE)
```

Arguments

files	vector: a character vector containing the paths for one or more Individual Recode DHS data files (see details)
extra.vars	vector: a character vector containing the names of variables to be retained from the raw data
progress	boolean: display a progress bar

Details

The **Demographic and Health Surveys** (DHS) program regularly collects health data from population-representative samples in many countries using standardized surveys since 1984. The "individual recode" data files contain women's responses, while the "men recode" files contain men's responses. These files are available in SPSS, SAS, and Stata formats from <https://www.dhsprogram.com/>, however access requires a **free application**. The `dhs()` function reads one or more of these files, extracts and recodes selected variables useful for studying childfree adults and other family statuses, then returns an unweighted data frame.

Although access to DHS data requires an application, the DHS program provides a **model dataset** for practice. The example provided below uses the model data file "ZZIR62FL.SAV", which contains fictitious women's data, but has the same structure as a real DHS data file. The example can be run without prior application for data access.

Sampling weights

The DHS is collected using a complex survey design. The survey package can be used to perform analyses that take these design features into account, and make it possible to obtain population-representative estimates. In most cases, a `svydesign` object for a single country and wave can be created using `survey::svydesign(data = data, ids = ~cluster, strata = ~strata, weights = ~weight, nest = TRUE)`. Additional information about analyzing DHS data using weights is available [here](#) and in the documentation provided with the downloaded data files.

Non-biological children

Information about non-biological children (e.g., adopted children, foster children, etc.) is not available in the DHS, which means that a respondent with only non-biological children would be classified as a non-parent. This is not exactly match the approach described by the ABC Framework

(Neal & Neal, 2024), and may lead to discrepancies when comparing DHS estimates to estimates derived from other data where information about non-biological children is available.

Additional notes

- The SPSS-formatted files containing data from Gabon Recode 4 (GAIR41FL.SAV, GAMR41FL.SAV) and Turkey Recode 4 (TRIR41FL.SAV, TRMR41FL.SAV) contain encoding errors. Use the SAS-formatted files (GAIR41FL.SAS7BDAT, GAMR41FL.SAS7BDAT, TRIR41FL.SAS7BDAT, TRMR41FL.SAS7BDAT) instead.
- In some cases, DHS makes available individual recode data files for specific regions. For example, women's data from individual states in India from 1999 are contained in files named XXIR42FL.SAV, where the "XX" is a two-letter state code. The `dhs()` function has only been tested using whole-country files, and may not perform as expected for regional files.
- Variables containing women's responses in the individual recode files begin with `v`, while variables containing men's responses in the men recode files begin with `mv`. When applying `dhs()` to both female and male data, these are automatically harmonized. However, if extra variables are requested using the `extra.vars` option, be sure to specify both names (e.g. `extra.vars = c("v201", "mv201")`).

Value

A data frame containing variables described in the codebook available using `vignette("codebooks")`. If you are offline, or if the requested data are otherwise unavailable, NULL is returned.

References

ABC Framework: Neal, Z. P. and Neal, J. W. (2024). A framework for studying adults who neither have nor want children. *The Family Journal*, 32, 121-130. doi:10.1177/10664807231198869

Examples

```
dat <- dhs(files = c("ZZIR62FL.SAV"), extra.vars = c("v201")) #Request data for fictitious country
if (!is.null(dat)) { #If data was available...
  table(dat$famstat)/nrow(dat) #Fraction of respondents with each family status
}
```

nsfg

Read and recode National Survey of Family Growth (NSFG) data

Description

Read and recode National Survey of Family Growth (NSFG) data

Usage

```
nsfg(years, nonbio = TRUE, keep_source = FALSE, progress = TRUE)
```

Arguments

years	vector: a numeric vector containing the starting year of NSFG waves to include (2002, 2006, 2011, 2013, 2015, 2017, 2022)
nonbio	boolean: should non-biological children be included
keep_source	boolean: keep the raw variables used to construct want_cf and famstat
progress	boolean: display a progress bar

Details

The U.S. Centers for Disease Control **National Survey of Family Growth** (NSFG) regularly collects fertility and other health information from a population-representative sample of adults in the United States. Between 1973 and 2002, the NSFG was conducted periodically. Starting in 2006, the NSFG transitioned to continuous data collection, releasing data in multi-year waves (e.g., 2006-2010, 2011-2013). The `nsfg()` function reads the raw data from CDC's website, extracts and recodes selected variables useful for studying childfree adults and other family statuses, then returns an unweighted data frame.

Sampling weights

The NSFG is collected using a complex survey design. The survey package can be used to perform analyses that take these design features into account, and make it possible to obtain population-representative estimates. In most cases, a `svydesign` object for a single wave can be created using `survey::svydesign(data = data, ids = ~cluster, strata = ~strata, weights = ~weight, nest = TRUE)`. Additional information about analyzing DHS data using weights is available [here](#).

Non-biological children

When `nonbio == TRUE` (default), non-biological children (e.g., adopted children, foster children, etc.) are treated the same as biological children when determining a respondent's family status. This matches the approach described by the ABC Framework (Neal & Neal, 2024), and should generally be used. However, non-biological children can be ignored by setting `nonbio = FALSE`, which may be useful when comparing NSFG estimates to estimates derived from other data where information about non-biological children is not available.

Additional notes

- Starting in 2006, "hispanic" was a response option for race, however "hispanic" is not a racial category, but an ethnicity. When a respondent chose this option, their actual race is unknown.
- Partnership status only describes a respondent's status with respect to an opposite-sex partner. Information about current or former same-sex partnerships is not available.
- The NSFG manual explains that "sample sizes for a single year are too small to provide estimates with adequate levels of precision," and therefore recommends avoiding analysis of data from single years. Instead, these data are designed to be analyzed by wave.

Value

A data frame containing variables described in the codebook available using `vignette("codebooks")`

References

NSFG Classification: Neal, J. W. and Neal, Z. P. (2025). Tracking types of non-parents in the United States. *Journal of Marriage and Family*. doi:10.1111/jomf.13097

ABC Framework: Neal, Z. P. and Neal, J. W. (2024). A framework for studying adults who neither have nor want children. *The Family Journal*, 32, 121-130. doi:10.1177/10664807231198869

Examples

```
unweighted <- nsfg(years = 2017) #Unweighted data
table(unweighted$famstat) / nrow(unweighted) #Fraction of respondents with each family status
```

so

Read and recode Michigan State of the State (SOSS) data

Description

Read and recode Michigan State of the State (SOSS) data

Usage

```
so(soss(waves, extra.vars = NULL, progress = TRUE))
```

Arguments

waves	vector: a numeric vector containing the SOSS waves to include (currently available: 79, 82, 84, 85, 86)
extra.vars	vector: a character vector containing the names of variables to be retained from the raw data
progress	boolean: display a progress bar

Details

The **State of the State Survey** (SOSS) is regularly collected by the Institute for Public Policy and Social Research (IPPSR) at Michigan State University (MSU). Each wave is collected from a sample of 1000 adults in the US state of Michigan, and includes sampling weights to obtain a sample that is representative of the state's population with respect to age, gender, race, and education. The `so()` function reads the raw data from IPPSR's website, extracts and recodes selected variables useful for studying childfree adults and other family statuses, then returns an unweighted data frame. Questions necessary for identifying childfree adults have been asked in five waves, which each include unique questions that may be of interest:

- **Wave 79** (May 2020) - Neighborhoods, Health care, COVID, Personality
- **Wave 82** (September 2021) - Trust in government, Critical Race Theory
- **Wave 84** (April 2022) - Trust in scientists, Autonomous vehicles, Morality
- **Wave 85** (September 2022) - Reproductive rights, Race equity

- **Wave 86** (December 2022) - Education, Infrastructure

Sampling weights

The SOSS includes sampling weights that can be incorporated into analyses using the survey package to obtain population-representative estimates. A `svydesign` object for a single wave can be created using `survey::svydesign(data = data, ids = ~1, weights = ~weight)`.

Non-biological children

Non-biological children (e.g., adopted children, foster children, etc.) are treated the same as biological children when determining a respondent's family status. This matches the approach described by the ABC Framework (Neal & Neal, 2024). However, it can lead to discrepancies when comparing SOSS estimates to estimates derived from other data where information about non-biological children is not available.

Additional notes

- Wave 79 did not include a "do not know" option for selected questions. Therefore, it is not possible to identify "undecided" or "ambivalent non-parent" respondents. This may lead other family status categories to be inflated.
- Wave 82 originally included a 500 person oversample of parents, but they are excluded from `nsfg(wave==82)`.
- The provided sampling weights are designed to be used in the analyses of individual waves. Combining data from multiple waves may require using adjusted weights.

Value

A data frame containing variables described in the codebook available using `vignette("codebooks")`. If you are offline, or if the requested data are otherwise unavailable, NULL is returned.

References

ABC Framework: Neal, Z. P. and Neal, J. W. (2024). A framework for studying adults who neither have nor want children. *The Family Journal*, 32, 121-130. [doi:10.1177/10664807231198869](https://doi.org/10.1177/10664807231198869)

Examples

```
dat <- soss(waves = 86) #Request data for December 2022
if (!is.null(dat)) { #If data was available...
  table(dat$famstat) / nrow(dat) #Fraction of respondents with each family status
}
```

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