

Package ‘censable’

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Title Making Census Data More Usable

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URL <https://www.christophertkenny.com/censable/>,
<https://github.com/christopherkenny/censable>

BugReports <https://github.com/christopherkenny/censable/issues>

Description Creates a common framework for organizing, naming, and gathering population, age, race, and ethnicity data from the Census Bureau. Accesses the API <https://www.census.gov/data/developers/data-sets.html> via the package tidycensus. Provides tools for adding information to existing data to line up with Census data.

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Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Suggests roxygen2, testthat (>= 3.0.0)

Imports magrittr, dplyr (>= 1.0.4), rlang (>= 0.1.2), tidycensus, sf (>= 1.0.0), tibble, stringr, memoise

Depends R (>= 2.10)

Config/testthat/edition 3

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add_r_environ	<i>Add Entry to Renviron</i>
---------------	------------------------------

Description

Adds a value to the Renvironment of the form name=value. Designed for flexibly adding API keys for future sessions. Defaults are set up for entering a Census API key to work with tidycensus.

Usage

```
add_r_environ(  
  value,  
  name = "CENSUS_API_KEY",  
  overwrite = FALSE,  
  install = FALSE  
)
```

Arguments

value	Character. Value to add.
name	Defaults to CENSUS_API_KEY. Character. Name to give value.
overwrite	Defaults to FALSE. Boolean. Should existing item with name name in Renviron be overwritten?
install	Defaults to FALSE. Boolean. Should this be added '~/.Renviron' file?

Value

value, invisibly

Examples

```
## Not run:  
add_r_environ('1234', 'SECRET_API_KEY')  
  
## End(Not run)
```

breakdown_geoid *Breakdown Census GEOID into Components*

Description

Breakdown Census GEOID into Components

Usage

```
breakdown_geoid(.data, GEOID = "GEOID", area_type = "spine")
```

Arguments

.data	dataframe, tibble, or sf tibble
GEOID	Column in .data with Census GEOID
area_type	String, default is 'spine' with type of GEOID. Options are 'spine' for states, counties, tracts, block groups, and blocks. 'shd' for lower state legislative districts, 'ssd' for upper state legislative districts, 'cd' for congressional districts, or 'zcta' for zip code tabulation areas.

Value

.data with added identifying columns based on area_type

Examples

```
data(mt_county)
mt_county <- mt_county %>% breakdown_geoid()
```

build_acs *Build Data from the Decennial Census*

Description

Creates a dataset, using the decennial census information, with the standard variables used for redistricting. Creates a stable base for getting data from tidycensus for common calls in redistricting.

#' # Output columns are:

- GEOID: Geographic Identifier
- NAME: Name of County
- pop: total population
- pop_white: total population, Non-Hispanic White
- pop_black: total population, Non-Hispanic Black
- pop_hisp: total population, Hispanic

- pop_aian: total population, Non-Hispanic American Indian and Alaskan Native
- pop_asian: total population, Non-Hispanic Asian
- pop_nhpi: total population, Non-Hispanic Native Hawaiian and Pacific Islander
- pop_other: total population, Non-Hispanic Other
- pop_two: total population, Non-Hispanic Two Plus Races
- vap: voting age population
- vap_white: voting age population, Non-Hispanic White
- vap_black: voting age population, Non-Hispanic Black
- vap_hisp: voting age population, Hispanic
- vap_aian: voting age population, Non-Hispanic American Indian and Alaskan Native
- vap_asian: voting age population, Non-Hispanic Asian
- vap_nhpi: voting age population, Non-Hispanic Native Hawaiian and Pacific Islander
- vap_other: voting age population, Non-Hispanic Other
- vap_two: voting age population, Non-Hispanic Two Plus Races
- geometry: sf geometry

Arguments for geography are not checked, so will fail with `tidycensus` errors if invalid. This is by design to avoid blocking usage that could become valid, especially following the 2020 Census data release.

Currently valid options for geography:

- 'state'
- 'county'
- 'tract'
- 'block group'
- 'block'
- 'county subdivision'
- 'zcta'
- 'congressional district'
- 'state legislative district (upper chamber)'
- 'state legislative district (lower chamber)'
- 'school district (unified)'

Full options for geography that may or may not be valid depending on year and geometry are listed at: [Kyle Walker's tidycensus site](#).

Usage

```
build_acs(
  geography,
  state,
  county = NULL,
  geometry = TRUE,
  year = 2010,
  survey = "acs5",
  groups = "all"
)
```

```
mem_build_acs(
  geography,
  state,
  county = NULL,
  geometry = TRUE,
  year = 2010,
  survey = "acs5",
  groups = "all"
)
```

Arguments

geography	Required. The geography level to use.
state	Required. Two letter state postal code.
county	Optional. Name of county. If not provided, returns blocks for the entire state.
geometry	Defaults to TRUE. Whether to return the geometry or not.
year	year, must be 2000, 2010, or 2020 (after August 2021)
survey	whether the get estimates from the 5-year ('acs5'), 3-year ('acs3'), or 1-year ('acs1') survey. Default is 'acs5'.
groups	defaults to 'all', which gets pop and vap. If 'pop', only gets pop. If 'vap', only gets vap. Any other strings default to 'all'.

Value

tibble with observations for each observation of the geography in the state or county. Data includes up to 3 sets of columns for each race or ethnicity category: population (pop), voting age population (vap), and citizen voting age population (cvap)

Examples

```
## Not run:
# uses the Census API
tb <- build_dec(geography = 'block', state = 'NY', county = 'Rockland', geometry = TRUE)

## End(Not run)
```

Description

Creates a dataset, using the decennial census information, with the standard variables used for redistricting. Creates a stable base for getting data from tidycensus for common calls in redistricting.

Usage

```
build_dec(  
  geography,  
  state,  
  county = NULL,  
  geometry = TRUE,  
  year = 2020,  
  groups = "all"  
)
```

```
mem_build_dec(  
  geography,  
  state,  
  county = NULL,  
  geometry = TRUE,  
  year = 2020,  
  groups = "all"  
)
```

Arguments

geography	Required. The geography level to use.
state	Required. Two letter state postal code.
county	Optional. Name of county. If not provided, returns blocks for the entire state.
geometry	Defaults to TRUE. Whether to return the geometry or not.
year	year, must be 2000, 2010, or 2020 (after August 2021)
groups	defaults to 'all', which gets pop and vap. If 'pop', only gets pop. If 'vap', only gets vap. Any other strings default to 'all'.

Value

tibble with observations for each observation of the geography in the state or county. Data includes 2 sets of columns for each race or ethnicity category: population (pop) and voting age population (vap)

Output columns are:

- GEOID: Geographic Identifier
- NAME: Name of County
- pop: total population
- pop_white: total population, Non-Hispanic White
- pop_black: total population, Non-Hispanic Black
- pop_hisp: total population, Hispanic
- pop_aian: total population, Non-Hispanic American Indian and Alaskan Native
- pop_asian: total population, Non-Hispanic Asian
- pop_nhpi: total population, Non-Hispanic Native Hawaiian and Pacific Islander
- pop_other: total population, Non-Hispanic Other
- pop_two: total population, Non-Hispanic Two Plus Races
- vap: voting age population
- vap_white: voting age population, Non-Hispanic White
- vap_black: voting age population, Non-Hispanic Black
- vap_hisp: voting age population, Hispanic
- vap_aian: voting age population, Non-Hispanic American Indian and Alaskan Native
- vap_asian: voting age population, Non-Hispanic Asian
- vap_nhpi: voting age population, Non-Hispanic Native Hawaiian and Pacific Islander
- vap_other: voting age population, Non-Hispanic Other
- vap_two: voting age population, Non-Hispanic Two Plus Races
- geometry: sf geometry

Arguments for geography are not checked, so will fail with tidycensus errors if invalid. This is by design to avoid blocking usage that could become valid, especially following the 2020 Census data release.

Currently valid options for geography:

- 'state'
- 'county'
- 'tract'
- 'block group'
- 'block'
- 'county subdivision'
- 'zcta'
- 'congressional district'
- 'state legislative district (upper chamber)'
- 'state legislative district (lower chamber)'
- 'school district (unified)'

Full options for geography that may or may not be valid depending on year and geometry are listed at: [Kyle Walker's tidycensus site](#).

Examples

```
## Not run:
# uses the Census API
tb <- build_dec(geography = 'block', state = 'NY', county = 'Rockland', geometry = TRUE)

## End(Not run)
```

collapse4	<i>Collapse Full Race Categories into 4 Categories</i>
-----------	--

Description

Collapses Other, AIAN, Asian, NHPI, and Two+ into other, by prefix.

Usage

```
collapse4(.data, prefix)
```

Arguments

.data tibble, data.frame, or sf tibble
 prefix The prefix(es) for the race categories. Must be a character vector.

Value

.data with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse4(prefix = c('pop_', 'vap_'))
```

collapse4_pop	<i>Collapse Population Race Categories into 4 Categories</i>
---------------	--

Description

Collapses Other, AIAN, Asian, NHPI, and Two+ into other.

Usage

```
collapse4_pop(.data, prefix = "pop_")
```

Arguments

.data tibble, data.frame, or sf tibble
 prefix Default is pop_. The prefix for the race categories.

Value

.data with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse4_pop()
```

collapse4_vap	<i>Collapse Voting Age Population Race Categories into 4 Categories</i>
---------------	---

Description

Collapses Other, AIAN, Asian, NHPI, and Two+ into other.

Usage

```
collapse4_vap(.data, prefix = "vap_")
```

Arguments

.data	tibble, data.frame, or sf tibble
prefix	Default is vap_. The prefix for the race categories.

Value

.data with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse4_vap()
```

collapse5	<i>Collapse Full Race Categories into 5 Categories</i>
-----------	--

Description

Collapses Other, AIAN, NHPI, and Two+ into Other, by prefix.

Usage

```
collapse5(.data, prefix)
```

Arguments

`.data` tibble, data.frame, or sf tibble
`prefix` The prefix(es) for the race categories. Must be a character vector.

Value

`.data` with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse5(prefix = c('pop_', 'vap_'))
```

collapse5_pop	<i>Collapse Population Race Categories into 5 Categories</i>
---------------	--

Description

Collapses Other, AIAN, NHPI, and Two+ into other.

Usage

```
collapse5_pop(.data, prefix = "pop_")
```

Arguments

`.data` tibble, data.frame, or sf tibble
`prefix` Default is pop_. The prefix for the race categories.

Value

`.data` with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse5_pop()
```

collapse5_vap	<i>Collapse Voting Age Population Race Categories into 5 Categories</i>
---------------	---

Description

Collapses Other, AIAN, NHPI, and Two+ into other.

Usage

```
collapse5_vap(.data, prefix = "vap_")
```

Arguments

.data	tibble, data.frame, or sf tibble
prefix	Default is vap_. The prefix for the race categories.

Value

.data with columns collapsed

Examples

```
data(mt_county)
mt_county <- mt_county %>% collapse5_vap()
```

construct_geoid	<i>Create GEOID from Default Columns</i>
-----------------	--

Description

Create GEOID from Default Columns

Usage

```
construct_geoid(
  .data,
  area_type,
  state = "state",
  county = "county",
  tract = "tract",
  block_group = "block group",
  block = "block",
  cd = "cd",
  shd = "shd",
  ssd = "ssd",
  zcta = "zcta"
)
```

Arguments

.data	dataframe, tibble, or sf tibble
area_type	Defaults to creating the smallest possible with 'spine' for states, counties, tracts, block groups, and blocks. You can also pass one of the on spine geographies to create that specific level. Other options are 'shd' for lower state legislative districts, 'ssd' for upper state legislative districts, 'cd' for congressional districts, or 'zcta' for zip code tabulation areas.
state	name of column with state component
county	name of column with county component
tract	name of column with tract component
block_group	name of column with block group component
block	name of column with block component
cd	name of column with cd component
shd	name of column with shd component
ssd	name of column with ssd component
zcta	name of column with zcta component

Value

.data with new column GEOID

Examples

```
data(mt_county)
mt_county <- mt_county %>% breakdown_geoid()
mt_county <- mt_county %>% dplyr::select(-dplyr::all_of('GEOID'))
mt_county <- mt_county %>% construct_geoid()
```

custom_geoid

Create a GEOID from Columns

Description

Create a GEOID from Columns

Usage

```
custom_geoid(.data, ...)
```

Arguments

.data	dataframe, tibble, or sf tibble
...	columns of .data in the order you want to make the GEOID

Value

.data with new column GEOID

Examples

```
data(mt_county)
mt_county <- mt_county %>% custom_geoid(GEOID)
```

fips_2000

Counties FIPS 2000

Description

Contains three columns:

- state: state FIPS
- county: county FIPS
- name: county name

Usage

```
data('fips_2000')
```

Value

tibble

Examples

```
data('fips_2000')
```

fips_2010

Counties FIPS 2010

Description

Contains three columns:

- state: state FIPS
- county: county FIPS
- name: county name

Usage

```
data('fips_2010')
```

Value

tibble

Examples

```
data('fips_2010')
```

fips_2020

Counties FIPS 2020

Description

Contains three columns:

- state: state FIPS
- county: county FIPS
- name: county name

Usage

```
data('fips_2020')
```

Value

tibble

Examples

```
data('fips_2020')
```

join_abb_ansi

Join Abb by ANSI

Description

Adds a column with state abbreviation joining by a column with state ansi

Usage

```
join_abb_ansi(.data, .ansi)
```

Arguments

.data data.frame or tibble
.ansi column with state ansi

Value

.data with column .ansi replaced with state abbreviation

Examples

```
data('stata')
stata %>% join_abb_ansi(ansi)
```

join_abb_fips	<i>Join Abb by FIPS</i>
---------------	-------------------------

Description

Adds a column with state abbreviation joining by a column with state fips

Usage

```
join_abb_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state abb

Examples

```
data('stata')
stata %>% join_abb_fips(fips)
```

join_abb_name	<i>Join Abb by Name</i>
---------------	-------------------------

Description

Adds a column with state abbs joining by a column with state names

Usage

```
join_abb_name(.data, .name)
```


Arguments

.data data.frame or tibble
.name column with state name

Value

.data with column .name replaced with abbreviation

Examples

```
data('stata')  
stata %>% join_abb_name(name)
```

join_ansi_abb	<i>Join ANSI by Abb</i>
---------------	-------------------------

Description

Adds a column with state ansi joining by a column with state abbreviation

Usage

```
join_ansi_abb(.data, .abb)
```

Arguments

.data data.frame or tibble
.abb column with state abbreviation

Value

.data with column .abb replaced with state ansi

Examples

```
data('stata')  
stata %>% join_ansi_abb(abb)
```

join_ansi_fips	<i>Join ANSI by FIPS</i>
----------------	--------------------------

Description

Adds a column with state ansi joining by a column with state fips

Usage

```
join_ansi_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state ansi

Examples

```
data('stata')  
stata %>% join_ansi_fips(fips)
```

join_ansi_name	<i>Join ANSI by Name</i>
----------------	--------------------------

Description

Adds a column with state ansi joining by a column with state name

Usage

```
join_ansi_name(.data, .name)
```

Arguments

.data	data.frame or tibble
.name	column with state name

Value

.data with column .name replaced with ansi

Examples

```
data('stata')
stata %>% join_ansi_name(name)
```

join_fips_abb	<i>Join FIPS by Abb</i>
---------------	-------------------------

Description

Adds a column with state fips joining by a column with state abbreviation

Usage

```
join_fips_abb(.data, .abb)
```

Arguments

.data	data.frame or tibble
.abb	column with state abbreviation

Value

.data with column .abb replaced with state name

Examples

```
data('stata')
stata %>% join_fips_abb(abb)
```

join_fips_ansi	<i>Join FIPS by ANSI</i>
----------------	--------------------------

Description

Adds a column with state fips joining by a column with state ansi

Usage

```
join_fips_ansi(.data, .ansi)
```

Arguments

.data	data.frame or tibble
.ansi	column with state ansi

Value

.data with column .ansi replaced with state fips

Examples

```
data('stata')
stata %>% join_fips_ansi(ansi)
```

join_fips_name	<i>Join FIPS by Name</i>
----------------	--------------------------

Description

Adds a column with state fips joining by a column with state name

Usage

```
join_fips_name(.data, .name)
```

Arguments

.data	data.frame or tibble
.name	column with state name

Value

.data with column .name replaced with fips

Examples

```
data('stata')
stata %>% join_fips_name(name)
```

join_name_abb	<i>Join Name by Abb</i>
---------------	-------------------------

Description

Adds a column with state name joining by a column with state abbreviatio

Usage

```
join_name_abb(.data, .abb)
```

Arguments

.data data.frame or tibble
.abb column with state abbreviation

Value

.data with column .abb replaced with state name

Examples

```
data('stata')  
stata %>% join_name_abb(abb)
```

join_name_ansi	<i>Join Name by ANSI</i>
----------------	--------------------------

Description

Adds a column with state name joining by a column with state ansi

Usage

```
join_name_ansi(.data, .ansi)
```

Arguments

.data data.frame or tibble
.ansi column with state ansi

Value

.data with column .ansi replaced with state name

Examples

```
data('stata')  
stata %>% join_name_ansi(name)
```

join_name_fips	<i>Join Name by FIPS</i>
----------------	--------------------------

Description

Adds a column with state name joining by a column with state fips

Usage

```
join_name_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state name

Examples

```
data('stata')
stata %>% join_name_fips(fips)
```

match_abb	<i>Try to Match to State Abbreviation</i>
-----------	---

Description

Searches for an exact match and offers the best match if no exact match

Usage

```
match_abb(state)
```

Arguments

state	character with state FIPS, Abbreviation, Name, or ANSI
-------	--

Value

Abbreviation if a match is found or character(0) if no match is found

Examples

```
match_abb('NY')
match_abb('01')
```

match_ansi	<i>Try to Match to State ANSI</i>
------------	-----------------------------------

Description

Searches for an exact match and offers the best match if no exact match

Usage

```
match_ansi(state)
```

Arguments

state character with state FIPS, Abbreviation, Name, or ANSI

Value

ANSI if a match is found or character(0) if no match is found

Examples

```
match_ansi('NY')  
match_ansi('01')
```

match_fips	<i>Try to Match to State FIPS</i>
------------	-----------------------------------

Description

Searches for an exact match and offers the best match if no exact match

Usage

```
match_fips(state)
```

Arguments

state character with state FIPS, Abbreviation, Name, or ANSI

Value

FIPS code if a match is found or character(0) if no match is found

Examples

```
match_fips('NY')  
match_fips('01')
```

match_name	<i>Try to Match to State Name</i>
------------	-----------------------------------

Description

Searches for an exact match and offers the best match if no exact match

Usage

```
match_name(state)
```

Arguments

state	character with state FIPS, Abbreviation, Name, or ANSI
-------	--

Value

Name if a match is found or character(0) if no match is found

Examples

```
match_name('NY')
match_name('01')
```

mt_county	<i>Montana County Data</i>
-----------	----------------------------

Description

- GEOID: Geographic Identifier
- NAME: Name of County
- pop: total population
- pop_white: total population, Non-Hispanic White
- pop_black: total population, Non-Hispanic Black
- pop_hisp: total population, Hispanic
- pop_aian: total population, Non-Hispanic American Indian and Alaskan Native
- pop_asian: total population, Non-Hispanic Asian
- pop_nhpi: total population, Non-Hispanic Native Hawaiian and Pacific Islander
- pop_other: total population, Non-Hispanic Other
- pop_two: total population, Non-Hispanic Two Plus Races
- vap: voting age population
- vap_white: voting age population, Non-Hispanic White

- vap_black: voting age population, Non-Hispanic Black
- vap_hisp: voting age population, Hispanic
- vap_aian: voting age population, Non-Hispanic American Indian and Alaskan Native
- vap_asian: voting age population, Non-Hispanic Asian
- vap_nhpi: voting age population, Non-Hispanic Native Hawaiian and Pacific Islander
- vap_other: voting age population, Non-Hispanic Other
- vap_two: voting age population, Non-Hispanic Two Plus Races
- geometry: sf geometry

Usage

```
data('mt_county')
```

Value

sf tibble with one observation for each county in Montana

Examples

```
data('mt_county')
```

recode_abb_ansi	<i>Recode Abb by ANSI</i>
-----------------	---------------------------

Description

Replaces state ansi with state abbreviation

Usage

```
recode_abb_ansi(.data, .ansi)
```

Arguments

.data	data.frame or tibble
.ansi	column with state ansi

Value

.data with column .ansi replaced with state abbreviation

Examples

```
data('stata')
stata %>% recode_abb_ansi(ansi)
```

recode_abb_fips	<i>Recode Abb by FIPS</i>
-----------------	---------------------------

Description

Replaces state fips with state abb

Usage

```
recode_abb_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state abb

Examples

```
data('stata')  
stata %>% recode_abb_fips(fips)
```

recode_abb_name	<i>Recode Abb by Name</i>
-----------------	---------------------------

Description

Replaces state name with state abbreviation

Usage

```
recode_abb_name(.data, .name)
```

Arguments

.data	data.frame or tibble
.name	column with state name

Value

.data with column .name replaced with abbreviation

Examples

```
data('stata')
stata %>% recode_abb_name(name)
```

recode_ansi_abb	<i>Recode ANSI by Abb</i>
-----------------	---------------------------

Description

Replaces state abbreviation with state ansi

Usage

```
recode_ansi_abb(.data, .abb)
```

Arguments

.data	data.frame or tibble
.abb	column with state abbreviation

Value

.data with column .abb replaced with state ansi

Examples

```
data('stata')
stata %>% recode_ansi_abb(abb)
```

recode_ansi_fips	<i>Recode ANSI by FIPS</i>
------------------	----------------------------

Description

Replaces state fips with state ansi

Usage

```
recode_ansi_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state ansi

Examples

```
data('stata')
stata %>% recode_ansi_fips(fips)
```

recode_ansi_name	<i>Recode ANSI by Name</i>
------------------	----------------------------

Description

Replaces state name with state ansi

Usage

```
recode_ansi_name(.data, .name)
```

Arguments

.data	data.frame or tibble
.name	column with state name

Value

.data with column .name replaced with ansi

Examples

```
data('stata')
stata %>% recode_ansi_name(name)
```

recode_fips_abb	<i>Recode FIPS by Abb</i>
-----------------	---------------------------

Description

Replaces state abbreviation with state fips

Usage

```
recode_fips_abb(.data, .abb)
```

Arguments

.data data.frame or tibble
.abb column with state abbreviation

Value

.data with column .abb replaced with state name

Examples

```
data('stata')  
stata %>% recode_fips_abb(abb)
```

recode_fips_ansi	<i>Recode FIPS by ANSI</i>
------------------	----------------------------

Description

Replaces state ansi with state fips

Usage

```
recode_fips_ansi(.data, .ansi)
```

Arguments

.data data.frame or tibble
.ansi column with state ansi

Value

.data with column .ansi replaced with state fips

Examples

```
data('stata')  
stata %>% recode_fips_ansi(ansi)
```

recode_fips_name	<i>Recode FIPS by Name</i>
------------------	----------------------------

Description

Replaces state name with state fips

Usage

```
recode_fips_name(.data, .name)
```

Arguments

.data	data.frame or tibble
.name	column with state name

Value

.data with column .name replaced with fips

Examples

```
data('stata')  
stata %>% recode_fips_name(name)
```

recode_name_abb	<i>Recode Name by Abb</i>
-----------------	---------------------------

Description

Replaces state abbreviation with state name

Usage

```
recode_name_abb(.data, .abb)
```

Arguments

.data	data.frame or tibble
.abb	column with state abbreviation

Value

.data with column .abb replaced with state name

Examples

```
data('stata')
stata %>% recode_name_abb(abb)
```

recode_name_ansi	<i>Recode Name by ANSI</i>
------------------	----------------------------

Description

Replaces state ansi with state name

Usage

```
recode_name_ansi(.data, .ansi)
```

Arguments

.data	data.frame or tibble
.ansi	column with state ansi

Value

.data with column .ansi replaced with state name

Examples

```
data('stata')
stata %>% recode_name_ansi(name)
```

recode_name_fips	<i>Recode Name by FIPS</i>
------------------	----------------------------

Description

Replaces state fips with state name

Usage

```
recode_name_fips(.data, .fips)
```

Arguments

.data	data.frame or tibble
.fips	column with state fips

Value

.data with column .fips replaced with state name

Examples

```
data('stata')
stata %>% recode_name_fips(fips)
```

stata

stata (State Data)

Description

tibble with columns:

- fips: Federal Information Processing Standards codes
- abb: two letter postal abbreviations
- name: title case state name
- ansi: American National Standards Institute codes
- region: Census Regions (for 50 states and D.C.)
- division: Census Divisions (for 50 states and D.C.)

Usage

```
data('stata')
```

Value

tibble with state identifying information

Examples

```
data('stata')
```


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