

# Package ‘srvyr’

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**Type** Package

**Title** 'dplyr'-Like Syntax for Summary Statistics of Survey Data

**Description** Use piping, verbs like 'group\_by' and 'summarize', and other 'dplyr' inspired syntactic style when calculating summary statistics on survey data using functions from the 'survey' package.

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---

as\_survey

*Create a tbl\_svy from a data.frame*


---

### Description

as\_survey can be used to create a tbl\_svy using design information ([as\\_survey\\_design](#)), replicate weights ([as\\_survey\\_rep](#)), or a two phase design ([as\\_survey\\_twophase](#)), or an object created by the survey package.

**Usage**

```
as_survey(.data, ...)

## S3 method for class 'tbl_svy'
as_survey(.data, ...)

## S3 method for class 'data.frame'
as_survey(.data, ...)

## S3 method for class 'tbl_lazy'
as_survey(.data, ...)

## S3 method for class 'survey.design2'
as_survey(.data, ...)

## S3 method for class 'svyrep.design'
as_survey(.data, ...)

## S3 method for class 'twophase2'
as_survey(.data, ...)
```

**Arguments**

<code>.data</code>	a data.frame or an object from the survey package
<code>...</code>	other arguments, see other functions for details

**Details**

See vignette("databases", package = "dplyr") for more information on setting up databases in dplyr.

**Value**

a `tbl_svy`

**Examples**

```
# Examples from ?survey::svydesign
library(survey)
library(dplyr)
data(api)

# stratified sample
dstrata <- apistrat %>%
  as_survey(strata = stype, weights = pw)

# Examples from ?survey::svrepdesign
data(scd)
# use BRR replicate weights from Levy and Lemeshow
scd$rep1 <- 2 * c(1, 0, 1, 0, 1, 0)
```

```

scd$rep2 <- 2 * c(1, 0, 0, 1, 0, 1)
scd$rep3 <- 2 * c(0, 1, 1, 0, 0, 1)
scd$rep4 <- 2 * c(0, 1, 0, 1, 1, 0)

scdrep <- scd %>%
  as_survey(type = "BRR", repweights = starts_with("rep"),
            combined_weights = FALSE)

# Examples from ?survey::twophase
# two-phase simple random sampling.
data(pbc, package="survival")

pbc <- pbc %>%
  mutate(randomized = !is.na(trt) & trt > 0,
         id = row_number())
d2pbc <- pbc %>%
  as_survey(id = list(id, id), subset = randomized)

# dplyr 0.7 introduced new style of NSE called quosures
# See `vignette("programming", package = "dplyr")` for details
st <- quo(stype)
wt <- quo(pw)
dstrata <- apistrat %>%
  as_survey(strata = !!st, weights = !!wt)

```

---

as\_survey\_design

*Create a tbl\_svy survey object using sampling design*


---

## Description

Create a survey object with a survey design.

## Usage

```

as_survey_design(.data, ...)

## S3 method for class 'data.frame'
as_survey_design(
  .data,
  ids = NULL,
  probs = NULL,
  strata = NULL,
  variables = NULL,
  fpc = NULL,
  nest = FALSE,
  check_strata = !nest,
  weights = NULL,
  pps = FALSE,
  variance = c("HT", "YG"),

```

```

    ...
  )

## S3 method for class 'survey.design2'
as_survey_design(.data, ...)

## S3 method for class 'tbl_lazy'
as_survey_design(
  .data,
  ids = NULL,
  probs = NULL,
  strata = NULL,
  variables = NULL,
  fpc = NULL,
  nest = FALSE,
  check_strata = !nest,
  weights = NULL,
  pps = FALSE,
  variance = c("HT", "YG"),
  ...
)

```

### Arguments

<code>.data</code>	A data frame (which contains the variables specified below)
<code>...</code>	ignored
<code>ids</code>	Variables specifying cluster ids from largest level to smallest level (leaving the argument empty, NULL, 1, or 0 indicate no clusters).
<code>probs</code>	Variables specifying cluster sampling probabilities.
<code>strata</code>	Variables specifying strata.
<code>variables</code>	Variables specifying variables to be included in survey. Defaults to all variables in <code>.data</code>
<code>fpc</code>	Variables specifying a finite population correct, see <a href="#">svydesign</a> for more details.
<code>nest</code>	If TRUE, relabel cluster ids to enforce nesting within strata.
<code>check_strata</code>	If TRUE, check that clusters are nested in strata.
<code>weights</code>	Variables specifying weights (inverse of probability).
<code>pps</code>	"brewer" to use Brewer's approximation for PPS sampling without replacement. "overton" to use Overton's approximation. An object of class HR to use the Hartley-Rao approximation. An object of class ppsmat to use the Horvitz-Thompson estimator.
<code>variance</code>	For pps without replacement, use variance="YG" for the Yates-Grundy estimator instead of the Horvitz-Thompson estimator

## Details

If provided a data.frame, it is a wrapper around [svydesign](#). All survey variables must be included in the data.frame itself. Variables are selected by using bare column names, or convenience functions described in [select](#).

If provided a survey.design2 object from the survey package, it will turn it into a srvyr object, so that srvyr functions will work with it

## Value

An object of class tbl\_svy

## Examples

```
# Examples from ?survey::svydesign
library(survey)
data(api)

# stratified sample
dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

# one-stage cluster sample
dclus1 <- apiclus1 %>%
  as_survey_design(dnum, weights = pw, fpc = fpc)

# two-stage cluster sample: weights computed from population sizes.
dclus2 <- apiclus2 %>%
  as_survey_design(c(dnum, snum), fpc = c(fpc1, fpc2))

## multistage sampling has no effect when fpc is not given, so
## these are equivalent.
dclus2wr <- apiclus2 %>%
  dplyr::mutate(weights = weights(dclus2)) %>%
  as_survey_design(c(dnum, snum), weights = weights)

dclus2wr2 <- apiclus2 %>%
  dplyr::mutate(weights = weights(dclus2)) %>%
  as_survey_design(c(dnum), weights = weights)

## syntax for stratified cluster sample
## (though the data weren't really sampled this way)
apistrat %>% as_survey_design(dnum, strata = stype, weights = pw,
                             nest = TRUE)

## PPS sampling without replacement
data(election)
dpps <- election_pps %>%
  as_survey_design(fpc = p, pps = "brewer")

# dplyr 0.7 introduced new style of NSE called quosures
# See `vignette("programming", package = "dplyr")` for details
```

```

st <- quo(stype)
wt <- quo(pw)
dstrata <- apistrat %>%
  as_survey_design(strata = !!st, weights = !!wt)

```

---

as\_survey\_rep

*Create a tbl\_svy survey object using replicate weights*


---

## Description

Create a survey object with replicate weights.

## Usage

```

as_survey_rep(.data, ...)

## S3 method for class 'data.frame'
as_survey_rep(
  .data,
  variables = NULL,
  repweights = NULL,
  weights = NULL,
  type = c("BRR", "Fay", "JK1", "JKn", "bootstrap", "other"),
  combined_weights = TRUE,
  rho = NULL,
  bootstrap_average = NULL,
  scale = NULL,
  rscales = NULL,
  fpc = NULL,
  fpc_type = c("fraction", "correction"),
  mse = getOption("survey.replicates.mse"),
  ...
)

## S3 method for class 'tbl_lazy'
as_survey_rep(
  .data,
  variables = NULL,
  repweights = NULL,
  weights = NULL,
  type = c("BRR", "Fay", "JK1", "JKn", "bootstrap", "other"),
  combined_weights = TRUE,
  rho = NULL,
  bootstrap_average = NULL,
  scale = NULL,
  rscales = NULL,

```

```

    fpc = NULL,
    fpctype = c("fraction", "correction"),
    mse = getOption("survey.replicates.mse"),
    ...
)

## S3 method for class 'svyrep.design'
as_survey_rep(.data, ...)

## S3 method for class 'survey.design2'
as_survey_rep(
  .data,
  type = c("auto", "JK1", "JKn", "BRR", "bootstrap", "subbootstrap", "mrbootstrap",
    "Fay"),
  rho = 0,
  fpc = NULL,
  fpctype = NULL,
  ...,
  compress = TRUE,
  mse = getOption("survey.replicates.mse")
)

## S3 method for class 'tbl_svy'
as_survey_rep(
  .data,
  type = c("auto", "JK1", "JKn", "BRR", "bootstrap", "subbootstrap", "mrbootstrap",
    "Fay"),
  rho = 0,
  fpc = NULL,
  fpctype = NULL,
  ...,
  compress = TRUE,
  mse = getOption("survey.replicates.mse")
)

```

### Arguments

<code>.data</code>	A data frame (which contains the variables specified below)
<code>...</code>	ignored
<code>variables</code>	Variables to include in the design (default is all)
<code>repweights</code>	Variables specifying the replication weight variables
<code>weights</code>	Variables specifying sampling weights
<code>type</code>	Type of replication weights
<code>combined_weights</code>	TRUE if the repweights already include the sampling weights. This is usually the case.
<code>rho</code>	Shrinkage factor for weights in Fay's method

bootstrap_average	For type = "bootstrap", if the bootstrap weights have been averaged, gives the number of iterations averaged over.
scale, rscales	Scaling constant for variance, see <a href="#">svrepdesign</a> for more information.
fpc, fpc_type	Finite population correction information
mse	if TRUE, compute variances based on sum of squares around the point estimate, rather than the mean of the replicates
compress	if TRUE, store replicate weights in compressed form (if converting from design)

## Details

If provided a data.frame, it is a wrapper around [svrepdesign](#). All survey variables must be included in the data.frame itself. Variables are selected by using bare column names, or convenience functions described in [select](#).

If provided a svyrep.design object from the survey package, it will turn it into a srvyr object, so that srvyr functions will work with it

If provided a survey design (survey.design2 or tbl\_svy), it is a wrapper around [as.svrepdesign](#), and will convert from a survey design to replicate weights.

## Value

An object of class tbl\_svy

## Examples

```
# Examples from ?survey::svrepdesign()
library(survey)
library(dplyr)
data(scd)
# use BRR replicate weights from Levy and Lemeshow
scd <- scd %>%
  mutate(rep1 = 2 * c(1, 0, 1, 0, 1, 0),
         rep2 = 2 * c(1, 0, 0, 1, 0, 1),
         rep3 = 2 * c(0, 1, 1, 0, 0, 1),
         rep4 = 2 * c(0, 1, 0, 1, 1, 0))

scdrep <- scd %>%
  as_survey_rep(type = "BRR", repweights = starts_with("rep"),
               combined_weights = FALSE)

# dplyr 0.7 introduced new style of NSE called quosures
# See `vignette("programming", package = "dplyr")` for details
repwts <- quo(starts_with("rep"))
scdrep <- scd %>%
  as_survey_rep(type = "BRR", repweights = !!repwts,
               combined_weights = FALSE)
```

---

as\_survey\_twophase      *Create a tbl\_svy survey object using two phase design*

---

## Description

Create a survey object by specifying the survey's two phase design. It is a wrapper around [twophase](#). All survey variables must be included in the data.frame itself. Variables are selected by using bare column names, or convenience functions described in [select](#).

## Usage

```
as_survey_twophase(.data, ...)

## S3 method for class 'data.frame'
as_survey_twophase(
  .data,
  id,
  strata = NULL,
  probs = NULL,
  weights = NULL,
  fpc = NULL,
  subset,
  method = c("full", "approx", "simple"),
  ...
)

## S3 method for class 'twophase2'
as_survey_twophase(.data, ...)
```

## Arguments

.data	A data frame (which contains the variables specified below)
...	ignored
id	list of two sets of variable names for sampling unit identifiers
strata	list of two sets of variable names (or NULLs) for stratum identifiers
probs	list of two sets of variable names (or NULLs) for sampling probabilities
weights	Only for method = "approx", list of two sets of variable names (or NULLs) for sampling weights
fpc	list of two sets of variables (or NULLs for finite population corrections
subset	bare name of a variable which specifies which observations are selected in phase 2
method	"full" requires (much) more memory, but gives unbiased variance estimates for general multistage designs at both phases. "simple" or "approx" use less memory, and is correct for designs with simple random sampling at phase one and stratified randoms sampling at phase two. See <a href="#">twophase</a> for more details.

**Value**

An object of class `tbl_svy`

**Examples**

```
# Examples from ?survey::twophase
# two-phase simple random sampling.
data(pbc, package="survival")
library(dplyr)

pbc <- pbc %>%
  mutate(randomized = !is.na(trt) & trt > 0,
         id = row_number())
d2pbc <- pbc %>%
  as_survey_twophase(id = list(id, id), subset = randomized)

d2pbc %>% summarize(mean = survey_mean(bili))

# two-stage sampling as two-phase
library(survey)
data(mu284)

mu284_1 <- mu284 %>%
  dplyr::slice(c(1:15, rep(1:5, n2[1:5] - 3))) %>%
  mutate(id = row_number(),
         sub = rep(c(TRUE, FALSE), c(15, 34-15)))

dmu284 <- mu284 %>%
  as_survey_design(ids = c(id1, id2), fpc = c(n1, n2))
# first phase cluster sample, second phase stratified within cluster
d2mu284 <- mu284_1 %>%
  as_survey_twophase(id = list(id1, id), strata = list(NULL, id1),
                    fpc = list(n1, NULL), subset = sub)
dmu284 %>%
  summarize(total = survey_total(y1),
           mean = survey_mean(y1))
d2mu284 %>%
  summarize(total = survey_total(y1),
           mean = survey_mean(y1))

# dplyr 0.7 introduced new style of NSE called quosures
# See `vignette("programming", package = "dplyr")` for details
ids <- quo(list(id, id))
d2pbc <- pbc %>%
  as_survey_twophase(id = !!ids, subset = "randomized")
```

**Description**

Coerce survey variables to a data frame (tibble)

**Arguments**

x                    A `tbl_svy` object

---

cascade	<i>Summarise multiple values into cascading groups</i>
---------	--

---

**Description**

`cascade` is similar to [summarise](#), but calculates a summary statistics for the total of a group in addition to each group.

**Usage**

```
cascade(.data, ..., .dots, .fill = NA)
```

**Arguments**

`.data`,            `tbl` A `tbl_svy` object

`...`             Name-value pairs of summary functions

`.dots`            Used to work around non-standard evaluation. See `vignette("nse", package = "dplyr")` for details.

`.fill`             Value to fill in for group summaries

**Examples**

```
library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata_grp <- dstrata %>%
  group_by(stype)

dstrata_grp %>%
  cascade(api99_mn = survey_mean(api99),
          api00_mn = survey_mean(api00),
          api_diff = survey_mean(api00 - api99))
```

---

collect	<i>Force computation of a database query</i>
---------	--

---

### Description

collect retrieves data from a database query (and when run on a `tbl_svy` object adjusts weights accordingly). Use `collect` when you want to run a function from the survey package on a `srvyr` db backed object. `compute` stores results in a remote temporary table.

---

cur_svy	<i>Get the survey data for the current context</i>
---------	--

---

### Description

This is a helper to allow `srvyr`'s syntactic style. In particular, it tells functions inside of a `summarize` call what survey to use (for the current group with `cur_svy()` or the complete survey for `cur_svy_full()`). In general, users will not have to worry about getting (or setting) the current context's survey, unless they are trying to extend `srvyr`. See `vignette("extending-srvyr")` for more details. `current_svy()` is deprecated, but returns the same value as `cur_svy()`.

### Usage

```
cur_svy()
```

```
cur_svy_full()
```

```
current_svy()
```

### Value

a `tbl_svy` (or error if called with no survey context)

---

get_var_est	<i>Get the variance estimates for a survey estimate</i>
-------------	---

---

### Description

This is a helper to allow `srvyr`'s syntactic style. In general, users will not have to worry about getting survey variance estimates directly unless they are trying to extend `srvyr`. This function helps convert from the result of a survey function into a `data.frame` with an estimate and measures of variance around it in a way that summarize expects. See `vignette("extending-srvyr")` for more details.

**Usage**

```

get_var_est(
  stat,
  vartype,
  grps = "",
  level = 0.95,
  df = Inf,
  pre_calc_ci = FALSE,
  deff = FALSE
)

```

**Arguments**

stat	A survey statistic object, usually the result of a function from the survey package or svyby.
vartype	A vector indicating which variance estimates to calculate (options are se for standard error, ci for confidence interval, var for variance or cv for coefficient of variation). Multiples are allowed.
grps	A vector indicating the names of the grouping variables for grouped surveys (" indicates no groups).
level	One or more levels to calculate a confidence interval.
df	Degrees of freedom, many survey functions default to Inf, but svyr functions generally default to the result of calling degf on the survey object.
pre_calc_ci	Whether the confidence interval is pre-calculated (as in svycirop)
deff	Whether to return the design effect (calculated using survey::deff)

**Value**

a `tbl_svy` with the variables modified

---

groups	<i>Get/set the grouping variables for tbl.</i>
--------	--

---

**Description**

These functions do not perform non-standard evaluation, and so are useful when programming against `tbl` objects. `ungroup` is a convenient inline way of removing existing grouping.

**Arguments**

x data `tbl_df` or `tbl_svy` object.

**See Also**

[groups](#) for information.

---

group_by	<i>Group a (survey) dataset by one or more variables.</i>
----------	---

---

## Description

Most data operations are useful when done on groups defined by variables in the dataset. The `group_by` function takes an existing table (or `svy_table`) and converts it to a grouped version, where operations are performed "by group".

## Arguments

<code>.data</code>	A <code>tbl</code>
<code>...</code>	variables to group by. All <code>tbls</code> accept variable names, some will also accept functions of variables. Duplicated groups will be silently dropped.
<code>add</code>	By default, when <code>add = FALSE</code> , <code>group_by</code> will override existing groups. To instead add to the existing groups, use <code>add = TRUE</code>
<code>.dots</code>	Used to work around non-standard evaluation. See <code>vignette("nse", package = "dplyr")</code> for details.

## Details

See [group\\_by](#) for more information about grouping regular data tables.

On `tbl_svy` objects, `group_by` sets up the object for operations similar to those allowed in [svyby](#).

## See Also

[group\\_by](#) for information about `group_by` on normal data tables.

## Examples

```
# Examples of svy_tbl group_by
library(survey)
data(api)
dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw) %>%
  group_by(stype)

dstrata %>%
  summarise(api_diff = survey_mean(api00 - api99))
```

---

group_map_dfr	<i>Apply a function to each group</i>
---------------	---------------------------------------

---

### Description

group\_map(), group\_walk and group\_map\_dfr are purrr-style functions that can be used to iterate on grouped survey objects (note that group\_map\_dfr replaces dplyr::group\_modify because we are changing the data from a tbl\_svy to a regular tibble).

### Usage

```
group_map_dfr(.data, .f, ..., .keep = FALSE)

## S3 method for class 'tbl_svy'
group_map(.data, .f, ..., .keep = FALSE)

group_map_dfr(.data, .f, ..., .keep = FALSE)
```

### Arguments

.data	A tbl_svy object
.f	A function or purrr-style formula to apply to each group
...	Other arguments passed to .f
.keep	Whether the grouping variables are kept when passed into .f

### Value

For group\_map a list, for group\_map\_dfr a 'tbl\_df', and for group\_walk invisibly the original tbl\_svy.

### Examples

```
data(api, package = "survey")
dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

results <- dstrata %>%
  group_by(both) %>%
  group_map(~survey::svyglm(api00~api99 + stype, .))

# group_map_dfr calls `bind_rows` on the list returned and includes
# grouping variables. This is most useful with a package like `broom`
# but could also be used with survey package functions.
result_coef <- dstrata %>%
  group_by(both) %>%
  group_map_dfr(
    ~data.frame(
      api99_coef = coef(survey::svyglm(api00~api99 + stype, .))["api99"]])
```

```
)
)
```

---

group\_trim

*Single table verbs from dplyr and tidyr*


---

### Description

These are data manipulation functions designed to work on `tbl_svy` objects.

### Details

`mutate` and `transmute` can add or modify variables. See [mutate](#) for more details.

`select`, `rename`, and `rename_with` keep or rename variables. See [select](#) for more details.

`pull` extracts a variable as a vector (whereas `select` returns a `tbl_svy`). See [pull](#) for more details.

`filter` keeps certain observations. See [filter](#) for more details.

`#` `drop_na` drops observations containing missing values. See [drop\\_na](#) for more details.

`arrange` is not implemented for `tbl_svy` objects. Nor are any two table verbs such as `bind_rows`, `bind_cols` or any of the joins (`full_join`, `left_join`, etc.). These data manipulations may require modifications to the survey variable specifications and so cannot be done automatically. Instead, use `dplyr` to perform them while the data is still stored in `data.frames`.

---

set\_survey\_vars

*Set the variables for the current survey variable*


---

### Description

This is a helper to allow `srvyr`'s syntactic style. In general, users will not have to worry about setting variables in a survey object unless they are trying to extend `srvyr`. This function helps convert a vector to a variable in the correct part of a survey object's structure so that functions can refer to it using the survey package's formula notation. See `vignette("extending-srvyr")` for more details.

### Usage

```
set_survey_vars(.svy, x, name = "__SRVYR_TEMP_VAR__", add = FALSE)
```

### Arguments

<code>.svy</code>	A survey object
<code>x</code>	A vector to be included in the variables portion of the survey object
<code>name</code>	The name of the variable once it is added. Defaults to <code>'__SRVYR_TEMP_VAR__'</code> which is formatted weirdly to avoid name collisions.
<code>add</code>	<code>FALSE</code> , the default, overwrite all current variables. If <code>TRUE</code> , will add this variable instead.

**Value**

a `tbl_svy` with the variables modified

---

srvyr	<i>srvyr: A package for 'dplyr'-Like Syntax for Summary Statistics of Survey Data.</i>
-------	--

---

**Description**

The `srvyr` package provides a new way of calculating summary statistics on survey data, based on the `dplyr` package. There are three stages to using `srvyr` functions, creating a survey object, manipulating the data, and calculating survey statistics.

**Functions to create a survey object**

`as_survey_design`, `as_survey_rep`, and `as_survey_twophase` are used to create surveys based on a `data.frame` and design variables, replicate weights or two phase design respectively. Each is based on a function in the `survey` package (`svydesign`, `svrepdesign`, `twophase`), and it is easy to modify code that uses the `survey` package so that it works with the `srvyr` package. See `vignette("srvyr_vs_survey")` for more details.

The function `as_survey` will choose between the other three functions based on the arguments given to save some typing.

**Functions to manipulate data in a survey object**

Once you've created a survey object, you can manipulate the data as you would using `dplyr` with a `data.frame`. `mutate` modifies or creates a variable, `select` and `rename` select or rename variables, and `filter` keeps certain observations.

Note that `arrange` and two table verbs such as `bind_rows`, `bind_cols`, or any of the joins are not usable on survey objects because they might require modifications to the definition of your survey. If you need to use these functions, you should do so before you convert the `data.frame` to a survey object.

**Functions to summarize a survey object**

Now that you have your data set up correctly, you can calculate summary statistics. To get the statistic over the whole population, use `summarise`, or to calculate it over a set of groups, use `group_by` first.

You can calculate the mean, (with `survey_mean`), the total (`survey_total`), the quantile (`survey_quantile`), or a ratio (`survey_ratio`). By default, `srvyr` will return the statistic and the standard error around it in a `data.frame`, but with the `vartype` parameter, you can also get a confidence interval ("ci"), variance ("var"), or coefficient of variation ("cv").

Within `summarise`, you can also use `unweighted`, which calculates a function without taking into consideration the survey weighting.

---

srvyr-se-deprecated     *Deprecated SE versions of main srvyr verbs*

---

## Description

srvyr has updated its standard evaluation semantics to match dplyr 0.7, so these underscore functions are no longer required (but are still supported for backward compatibility reasons). See [se-deprecated](#) or the dplyr vignette on programming (`vignette("programming", package = "dplyr")`) for more details.

## Usage

```
as_survey_(.data, ...)
```

```
as_survey_design_(  
  .data,  
  ids = NULL,  
  probs = NULL,  
  strata = NULL,  
  variables = NULL,  
  fpc = NULL,  
  nest = FALSE,  
  check_strata = !nest,  
  weights = NULL,  
  pps = FALSE,  
  variance = c("HT", "YG")  
)
```

```
as_survey_rep_(  
  .data,  
  variables = NULL,  
  repweights = NULL,  
  weights = NULL,  
  type = c("BRR", "Fay", "JK1", "JKn", "bootstrap", "other"),  
  combined_weights = TRUE,  
  rho = NULL,  
  bootstrap_average = NULL,  
  scale = NULL,  
  rscales = NULL,  
  fpc = NULL,  
  fpc_type = c("fraction", "correction"),  
  mse = getOption("survey.replicates.mse")  
)
```

```
as_survey_twophase_(  
  .data,  
  id,
```

```

    strata = NULL,
    probs = NULL,
    weights = NULL,
    fpc = NULL,
    subset,
    method = c("full", "approx", "simple")
)

cascade_(.data, ..., .dots, .fill = NA)

```

### Arguments

<code>.data</code>	a data.frame or an object from the survey package
<code>...</code>	other arguments, see other functions for details
<code>ids</code>	Variables specifying cluster ids from largest level to smallest level (leaving the argument empty, NULL, 1, or 0 indicate no clusters).
<code>probs</code>	Variables specifying cluster sampling probabilities.
<code>strata</code>	Variables specifying strata.
<code>variables</code>	Variables specifying variables to be included in survey. Defaults to all variables in <code>.data</code>
<code>fpc</code>	Variables specifying a finite population correct, see <a href="#">svydesign</a> for more details.
<code>nest</code>	If TRUE, relabel cluster ids to enforce nesting within strata.
<code>check_strata</code>	If TRUE, check that clusters are nested in strata.
<code>weights</code>	Variables specifying weights (inverse of probability).
<code>pps</code>	"brewer" to use Brewer's approximation for PPS sampling without replacement. "overton" to use Overton's approximation. An object of class HR to use the Hartley-Rao approximation. An object of class ppsmat to use the Horvitz-Thompson estimator.
<code>variance</code>	For pps without replacement, use <code>variance="YG"</code> for the Yates-Grundy estimator instead of the Horvitz-Thompson estimator
<code>repweights</code>	Variables specifying the replication weight variables
<code>type</code>	Type of replication weights
<code>combined_weights</code>	TRUE if the repweights already include the sampling weights. This is usually the case.
<code>rho</code>	Shrinkage factor for weights in Fay's method
<code>bootstrap_average</code>	For <code>type = "bootstrap"</code> , if the bootstrap weights have been averaged, gives the number of iterations averaged over.
<code>scale</code>	Scaling constant for variance, see <a href="#">svrepdesign</a> for more information.
<code>rscals</code>	Scaling constant for variance, see <a href="#">svrepdesign</a> for more information.
<code>fpc_type</code>	Finite population correction information

mse	if TRUE, compute variances based on sum of squares around the point estimate, rather than the mean of the replicates
id	list of two sets of variable names for sampling unit identifiers
subset	bare name of a variable which specifies which observations are selected in phase 2
method	"full" requires (much) more memory, but gives unbiased variance estimates for general multistage designs at both phases. "simple" or "approx" use less memory, and is correct for designs with simple random sampling at phase one and stratified random sampling at phase two. See <a href="#">twophase</a> for more details.
.dots	Used to work around non-standard evaluation. See <code>vignette("nse", package = "dplyr")</code> for details.
.fill	Value to fill in for group summaries

---

summarise	<i>Summarise multiple values to a single value.</i>
-----------	---

---

## Description

Summarise multiple values to a single value.

## Arguments

.data	tbl A <code>tbl_svy</code> object
...	Name-value pairs of summary functions
.groups	Defaults to "drop_last" in <code>srvyr</code> meaning that the last group is peeled off, but if there are more groups they will be preserved. Other options are "drop", which drops all groups, "keep" which keeps all of them and "rowwise" which converts the object to a rowwise object (meaning calculations will be performed on each row).
.unpack	Whether to "unpack" named <code>data.frame</code> columns. <code>srvyr</code> predates <code>dplyr</code> 's support for <code>data.frame</code> columns so it does not treat them the same way by default.

## Details

Summarise for `tbl_svy` objects accepts several specialized functions. Each of the functions a variable (or two, in the case of `survey_ratio`), from the `data.frame` and default to providing the measure and its standard error.

The argument `vartype` can choose one or more measures of uncertainty, `se` for standard error, `ci` for confidence interval, `var` for variance, and `cv` for coefficient of variation. `level` specifies the level for the confidence interval.

The other arguments correspond to the analogous function arguments from the `survey` package.

The available functions from `srvyr` are:

[survey\\_mean](#) Calculate the survey mean of the entire population or by groups. Based on [svymean](#).

[survey\\_total](#) Calculate the survey total of the entire population or by groups. Based on [svytotal](#).

[survey\\_ratio](#) Calculate the ratio of 2 variables in the entire population or by groups. Based on [svyratio](#).

[survey\\_quantile](#) Calculate quantiles in the entire population or by groups. Based on [svyquantile](#).

[survey\\_median](#) Calculate the median in the entire population or by groups. [svyquantile](#).

[unweighted](#) Calculate an unweighted estimate as you would on a regular tbl\_df. Based on dplyr's [summarise](#).

## Examples

```
library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(api99_mn = survey_mean(api99),
            api00_mn = survey_mean(api00),
            api_diff = survey_mean(api00 - api99))

dstrata_grp <- dstrata %>%
  group_by(stype)

dstrata_grp %>%
  summarise(api99_mn = survey_mean(api99),
            api00_mn = survey_mean(api00),
            api_diff = survey_mean(api00 - api99))
```

---

<code>summarise_all</code>	<i>Manipulate multiple columns.</i>
----------------------------	-------------------------------------

---

## Description

See [summarize\\_all](#) for more details. \*\_each functions will be deprecated in favor of \*\_all/\*\_if/\*\_at functions.

---

<code>survey_mean</code>	<i>Calculate the mean and its variation using survey methods</i>
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---

## Description

Calculate means and proportions from complex survey data. A wrapper around [svymean](#), or if `proportion = TRUE`, [svyciprop](#). `survey_mean` should always be called from [summarise](#).

**Usage**

```

survey_mean(
  x,
  na.rm = FALSE,
  vartype = c("se", "ci", "var", "cv"),
  level = 0.95,
  proportion = FALSE,
  prop_method = c("logit", "likelihood", "asin", "beta", "mean"),
  deff = FALSE,
  df = NULL,
  ...
)

survey_prop(
  vartype = c("se", "ci", "var", "cv"),
  level = 0.95,
  proportion = FALSE,
  prop_method = c("logit", "likelihood", "asin", "beta", "mean"),
  deff = FALSE,
  df = NULL,
  ...
)

```

**Arguments**

x	A variable or expression, or empty
na.rm	A logical value to indicate whether missing values should be dropped
vartype	Report variability as one or more of: standard error ("se", default), confidence interval ("ci"), variance ("var") or coefficient of variation ("cv").
level	(For vartype = "ci" only) A single number or vector of numbers indicating the confidence level
proportion	Use methods to calculate the proportion that may have more accurate confidence intervals near 0 and 1. Based on <a href="#">svyciprop</a> .
prop_method	Type of proportion method to use if proportion is TRUE. See <a href="#">svyciprop</a> for details.
deff	A logical value to indicate whether the design effect should be returned.
df	(For vartype = "ci" only) A numeric value indicating the degrees of freedom for t-distribution. The default (NULL) uses <a href="#">degf</a> , but Inf is the usual survey package's default (except in <a href="#">svyciprop</a> ).
...	Ignored

**Examples**

```

library(survey)
data(api)

```

```

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(api99_mn = survey_mean(api99),
            api_diff = survey_mean(api00 - api99, vartype = c("ci", "cv")))

dstrata %>%
  group_by(awards) %>%
  summarise(api00 = survey_mean(api00))

# Leave x empty to calculate the proportion in each group
dstrata %>%
  group_by(awards) %>%
  summarise(pct = survey_mean())

# Setting proportion = TRUE uses a different method for calculating confidence intervals
dstrata %>%
  summarise(high_api = survey_mean(api00 > 875, proportion = TRUE, vartype = "ci"))

# level takes a vector for multiple levels of confidence intervals
dstrata %>%
  summarise(api99 = survey_mean(api99, vartype = "ci", level = c(0.95, 0.65)))

# Note that the default degrees of freedom in srvyr is different from
# survey, so your confidence intervals might not be exact matches. To
# Replicate survey's behavior, use df = Inf
dstrata %>%
  summarise(srvyr_default = survey_mean(api99, vartype = "ci"),
            survey_default = survey_mean(api99, vartype = "ci", df = Inf))

comparison <- survey::svymean(~api99, dstrata)
confint(comparison) # survey's default
confint(comparison, df = survey::degf(dstrata)) # srvyr's default

```

---

survey\_quantile

*Calculate the quantile and its variation using survey methods*

---

## Description

Calculate quantiles from complex survey data. A wrapper around [svyquantile](#). `survey_quantile` and `survey_median` should always be called from [summarise](#).

## Usage

```

survey_quantile(
  x,
  quantiles,
  na.rm = FALSE,

```

```

vartype = c("se", "ci", "var", "cv"),
level = 0.95,
q_method = "linear",
f = 1,
interval_type = c("Wald", "score", "betaWald", "probability", "quantile"),
ties = c("discrete", "rounded"),
df = NULL,
...
)

survey_median(
  x,
  na.rm = FALSE,
  vartype = c("se", "ci"),
  level = 0.95,
  q_method = "linear",
  f = 1,
  interval_type = c("Wald", "score", "betaWald", "probability", "quantile"),
  ties = c("discrete", "rounded"),
  df = NULL,
  ...
)

```

### Arguments

x	A variable or expression
quantiles	A vector of quantiles to calculate
na.rm	A logical value to indicate whether missing values should be dropped
vartype	NULL to report no variability (default), otherwise one or more of: standard error ("se") confidence interval ("ci") (variance and coefficient of variation not available).
level	A single number indicating the confidence level (only one level allowed)
q_method	See "method" in <a href="#">approxfun</a>
f	See <a href="#">approxfun</a>
interval_type	See <a href="#">svyquantile</a>
ties	See <a href="#">svyquantile</a>
df	A number indicating the degrees of freedom for t-distribution. The default, Inf uses the normal distribution (matches the survey package). Also, has no effect for type = "betaWald".
...	Ignored

### Examples

```

library(survey)
data(api)

```

```
dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(api99 = survey_quantile(api99, c(0.25, 0.5, 0.75)),
            api00 = survey_median(api00, vartype = c("ci")))

dstrata %>%
  group_by(awards) %>%
  summarise(api00 = survey_median(api00))
```

---

 survey\_ratio

*Calculate the ratio and its variation using survey methods*


---

## Description

Calculate ratios from complex survey data. A wrapper around [svyratio](#). `survey_ratio` should always be called from [summarise](#).

## Usage

```
survey_ratio(
  numerator,
  denominator,
  na.rm = FALSE,
  vartype = c("se", "ci", "var", "cv"),
  level = 0.95,
  deff = FALSE,
  df = NULL,
  ...
)
```

## Arguments

<code>numerator</code>	The numerator of the ratio
<code>denominator</code>	The denominator of the ratio
<code>na.rm</code>	A logical value to indicate whether missing values should be dropped
<code>vartype</code>	Report variability as one or more of: standard error ("se", default), confidence interval ("ci"), variance ("var") or coefficient of variation ("cv").
<code>level</code>	A single number or vector of numbers indicating the confidence level
<code>deff</code>	A logical value to indicate whether the design effect should be returned.
<code>df</code>	(For <code>vartype = "ci"</code> only) A numeric value indicating the degrees of freedom for t-distribution. The default (NULL) uses <a href="#">degf</a> , but Inf is the usual survey package's default (except in <a href="#">svyciprop</a> ).
<code>...</code>	Ignored

**Examples**

```

library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(enroll = survey_ratio(api00, api99, vartype = c("ci", "cv")))

dstrata %>%
  group_by(awards) %>%
  summarise(api00 = survey_ratio(api00, api99))

# level takes a vector for multiple levels of confidence intervals
dstrata %>%
  summarise(enroll = survey_ratio(api99, api00, vartype = "ci", level = c(0.95, 0.65)))

# Note that the default degrees of freedom in srvyr is different from
# survey, so your confidence intervals might not exactly match. To
# replicate survey's behavior, use df = Inf
dstrata %>%
  summarise(srvyr_default = survey_total(api99, vartype = "ci"),
            survey_default = survey_total(api99, vartype = "ci", df = Inf))

comparison <- survey::svytotal(~api99, dstrata)
confint(comparison) # survey's default
confint(comparison, df = survey::degf(dstrata)) # srvyr's default

```

---

survey\_tally

*Count/tally survey weighted observations by group*


---

**Description**

Analogous to [tally](#) and [count](#), calculates the survey weighted count of observations. `survey_tally` will call [survey\\_total](#) empty (resulting in the count of each group) or on `wt` if it is specified (resulting in the survey weighted total of `wt`). `survey_count` is similar, but calls `group_by` before calculating the count and then returns the data to the original groupings.

**Usage**

```

survey_tally(
  x,
  wt,
  sort = FALSE,
  name = "n",
  vartype = c("se", "ci", "var", "cv")
)

```

```

survey_count(
  x,
  ...,
  wt = NULL,
  sort = FALSE,
  name = "n",
  .drop = dplyr::group_by_drop_default(x),
  vartype = c("se", "ci", "var", "cv")
)

```

### Arguments

x	A <code>tbl_svy</code> object, as created by <code>as_survey</code> and related functions.
wt	(Optional) A variable to weight on (in addition to the survey weights, which are always used). If left unspecified, <code>tally()</code> will use a variable named "n" if one exists, but <code>count()</code> will not. Override this behavior by specifying <code>wt = NULL</code> .
sort	Whether to sort the results (defaults to <code>FALSE</code> )
name	Name of count variable created (defaults to <code>n</code> ). If the variable already exists, will add "n" to the end until it does not.
vartype	What types variation estimates to calculate, passed to <a href="#">survey_total</a> .
...	Variables to group by, passed to <code>group_by()</code> .
.drop	When <code>.drop = TRUE</code> , empty groups are dropped, see <a href="#">group_by</a> documentation for more details.

### Details

If `n` already exists, `tally` will use it as the weight, but `count` will not.

### Examples

```

library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  group_by(awards) %>%
  survey_tally()

dstrata %>%
  survey_count(awards)

```

---

survey_total	<i>Calculate the total and its variation using survey methods</i>
--------------	---

---

### Description

Calculate totals from complex survey data. A wrapper around `svytotal`. `survey_total` should always be called from `summarise`.

### Usage

```
survey_total(
  x,
  na.rm = FALSE,
  vartype = c("se", "ci", "var", "cv"),
  level = 0.95,
  deff = FALSE,
  df = NULL,
  ...
)
```

### Arguments

<code>x</code>	A variable or expression, or empty
<code>na.rm</code>	A logical value to indicate whether missing values should be dropped
<code>vartype</code>	Report variability as one or more of: standard error ("se", default), confidence interval ("ci"), variance ("var") or coefficient of variation ("cv").
<code>level</code>	A single number or vector of numbers indicating the confidence level
<code>deff</code>	A logical value to indicate whether the design effect should be returned.
<code>df</code>	(For <code>vartype = "ci"</code> only) A numeric value indicating the degrees of freedom for t-distribution. The default (NULL) uses <code>degf</code> , but Inf is the usual survey package's default.
<code>...</code>	Ignored

### Examples

```
library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(enroll_tot = survey_total(enroll),
            tot_meals = survey_total(enroll * meals / 100, vartype = c("ci", "cv")))

dstrata %>%
```

```

group_by(awards) %>%
  summarise(api00 = survey_total(enroll))

# Leave x empty to calculate the total in each group
dstrata %>%
  group_by(awards) %>%
  summarise(pct = survey_total())

# level takes a vector for multiple levels of confidence intervals
dstrata %>%
  summarise(enroll = survey_total(enroll, vartype = "ci", level = c(0.95, 0.65)))

# Note that the default degrees of freedom in srvyr is different from
# survey, so your confidence intervals might not exactly match. To
# replicate survey's behavior, use df = Inf
dstrata %>%
  summarise(srvyr_default = survey_total(api99, vartype = "ci"),
            survey_default = survey_total(api99, vartype = "ci", df = Inf))

comparison <- survey::svytotal(~api99, dstrata)
confint(comparison) # survey's default
confint(comparison, df = survey::degf(dstrata)) # srvyr's default

```

---

survey_var	<i>Calculate the population variance and its variation using survey methods</i>
------------	---

---

## Description

Calculate population variance from complex survey data. A wrapper around [svyvar](#). `survey_var` should always be called from [summarise](#).

## Usage

```

survey_var(
  x,
  na.rm = FALSE,
  vartype = c("se", "ci", "var"),
  level = 0.95,
  df = NULL,
  ...
)

survey_sd(x, na.rm = FALSE, ...)

```

## Arguments

`x` A variable or expression, or empty

na.rm	A logical value to indicate whether missing values should be dropped
vartype	Report variability as one or more of: standard error ("se", default) or variance ("var") (confidence intervals and coefficient of variation not available).
level	(For vartype = "ci" only) A single number or vector of numbers indicating the confidence level.
df	(For vartype = "ci" only) A numeric value indicating the degrees of freedom for t-distribution. The default (Inf) is equivalent to using normal distribution and in case of population variance statistics there is little reason to use any other values (see <i>Details</i> ).
...	Ignored

### Details

Be aware that confidence intervals for population variance statistic are computed by package *survey* using *t* or normal (with *df*=Inf) distribution (i.e. symmetric distributions). **This could be a very poor approximation** if even one of these conditions is met:

- there are few sampling design degrees of freedom,
- analyzed variable isn't normally distributed,
- there is huge variation in sampling probabilities of the survey design.

Because of this be very careful using confidence intervals for population variance statistics especially while performing analysis within subsets of data or using grouped survey objects.

Sampling distribution of the variance statistic in general is asymmetric (chi-squared in case of simple random sampling of normally distributed variable) and if analyzed variable isn't normally distributed or there is huge variation in sampling probabilities of the survey design (or both) it could converge to normality only very slowly (with growing number of survey design degrees of freedom).

### Examples

```
library(survey)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(api99_var = survey_var(api99),
            api99_sd = survey_sd(api99))

dstrata %>%
  group_by(awards) %>%
  summarise(api00_var = survey_var(api00),
            api00_sd = survey_sd(api00))

# standard deviation and variance of the population variance estimator
# are available with vartype argument
# (but not for the population standard deviation estimator)
dstrata %>%
  summarise(api99_variance = survey_var(api99, vartype = c("se", "var")))
```

---

svychisq	<i>Chisquared tests of association for survey data.</i>
----------	---

---

### Description

Chisquared tests of association for survey data.

### Arguments

formula	See details in <a href="#">svychisq</a>
design	See details in <a href="#">svychisq</a>
na.rm	See details in <a href="#">svychisq</a>
...	See details in <a href="#">svychisq</a>

---

tbl_svy	<i>tbl_svy object.</i>
---------	------------------------

---

### Description

A `tbl_svy` wraps a locally stored `svydesign` and adds methods for `dplyr` single-table verbs like `mutate`, `group_by` and `summarise`. Create a `tbl_svy` using [as\\_survey\\_design](#).

### Methods

`tbl_df` implements these methods from `dplyr`.

[select](#) or [rename](#) Select or rename variables in a survey's dataset.

[mutate](#) or [transmute](#) Modify and create variables in a survey's dataset.

[group\\_by](#) and [summarise](#) Get descriptive statistics from survey.

### Examples

```
library(survey)
library(dplyr)
data(api)
svy <- as_survey_design(apistrat, strata = stype, weights = pw)
svy

# Data manipulation verbs -----
filter(svy, pcttest > 95)
select(svy, starts_with("acs")) # variables used in survey design are automatically kept
summarise(svy, col.grad = survey_mean(col.grad))
mutate(svy, api_diff = api00 - api99)

# Group by operations -----
```

```
# To calculate survey
svy_group <- group_by(svy, dname)

summarise(svy, col.grad = survey_mean(col.grad),
          api00 = survey_mean(api00, vartype = "ci"))
```

---

tbl_vars	<i>List variables produced by a tbl.</i>
----------	--

---

### Description

List variables produced by a tbl.

### Arguments

x	A tbl object
---	--------------

---

unweighted	<i>Calculate the an unweighted summary statistic from a survey</i>
------------	--

---

### Description

Calculate unweighted summaries from a survey dataset, just as on a normal data.frame with [summarise](#). Though it is possible to use regular functions directly, because the survey package doesn't always remove rows when filtering (instead setting the weight to 0), this can sometimes give bad results. See examples for more details.

### Usage

```
unweighted(x, ...)
```

### Arguments

x	A variable or expression
...	Ignored

### Details

Uses tidy evaluation semantics and so if you want to use wrapper functions based on variable names, you must use tidy evaluation, see the examples here, documentation in [nse-force](#), or the dplyr vignette called 'programming' for more information.

**Examples**

```

library(survey)
library(dplyr)
data(api)

dstrata <- apistrat %>%
  as_survey_design(strata = stype, weights = pw)

dstrata %>%
  summarise(api99_unw = unweighted(mean(api99)),
            n = unweighted(n()))

dstrata %>%
  group_by(stype) %>%
  summarise(api_diff_unw = unweighted(mean(api00 - api99)))

# Some survey designs, like ones with raked weights, are not removed
# when filtered to preserve the structure. So if you don't use `unweighted()`
# your results can be wrong.
# Declare basic clustered design ----
cluster_design <- as_survey_design(
  .data = apiclus1,
  id = dnum,
  weights = pw,
  fpc = fpc
)

# Add raking weights for school type ----
pop.types <- data.frame(stype=c("E","H","M"), Freq=c(4421,755,1018))
pop.schwide <- data.frame(sch.wide=c("No","Yes"), Freq=c(1072,5122))

raked_design <- rake(
  cluster_design,
  sample.margins = list(~stype,~sch.wide),
  population.margins = list(pop.types, pop.schwide)
)

raked_design %>%
  filter(cname != "Alameda") %>%
  group_by(cname) %>%
  summarize(
    direct_unw_mean = mean(api99),
    wrapped_unw_mean = unweighted(mean(api99))
  ) %>%
  filter(cname == "Alameda")

# Notice how the results are different when using `unweighted()`

```

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