

Package ‘interSAE’

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Type Package

Title Intersectional Small Area Estimation from Survey and Census Data

Version 0.1.0

Description Implements design-aware, margin-constrained estimation of population indicators for geographic areas crossed with demographic subgroups. Complex-survey microdata supply outcomes and association information, while one or more aggregate census tables supply overlapping population margins. A generalized iterative proportional fitting engine reconstructs a coherent latent population table, and an augmented model-assisted estimator produces domain means, proportions, and totals for sampled and unsampled intersections. Tools diagnose non-identification, compute linear-programming sensitivity bounds, propagate sampling and model uncertainty with replicate-weight or multiplier bootstrap procedures, enforce structural zeros, and benchmark estimates to official totals. The framework extends calibration ideas from Deville and Sarndal (1992) <[doi:10.1080/01621459.1992.10475217](https://doi.org/10.1080/01621459.1992.10475217)> and small area estimation ideas from Fay and Herriot (1979) <[doi:10.1080/01621459.1979.10482505](https://doi.org/10.1080/01621459.1979.10482505)>.

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interSAE-package	<i>Intersectional Small Area Estimation from Survey and Census Data</i>
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Description

The **interSAE** package combines complex-survey microdata with overlapping aggregate census margins. It reconstructs a coherent latent population table and produces model-assisted estimates for geographic areas crossed with demographic subgroups.

Details

The package separates survey-design declaration, census-margin validation, constrained population reconstruction, and outcome estimation. Optional tools diagnose non-identification, calculate association-sensitivity bounds, bootstrap the complete fitting procedure, and benchmark estimates.

Author(s)

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References

Deville, J.-C. and Sarndal, C.-E. (1992). Calibration estimators in survey sampling. *Journal of the American Statistical Association*, 87, 376–382. doi:10.1080/01621459.1992.10475217.

Fay, R. E. and Herriot, R. A. (1979). Estimates of income for small places. *Journal of the American Statistical Association*, 74, 269–277. doi:10.1080/01621459.1979.10482505.

benchmark_inter	<i>Benchmark Domain Estimates to an Official Aggregate</i>
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Description

Apply a common adjustment while reproducing an official population-weighted aggregate. Logit adjustment preserves a bounded response range.

Usage

```
benchmark_inter(estimates, target, weight = "population",  
  method = c("logit", "ratio", "additive"), lower = 0, upper = 1)
```

Arguments

estimates	A fitted inter_sae object or estimate data frame.
target	Official weighted mean or proportion.
weight	Benchmarking-weight column.
method	Logit, ratio, or additive adjustment.
lower, upper	Range for bounded logit adjustment.

Value

A data frame with original and benchmarked estimates.

Examples

```
example_data <- inter_example()  
design <- inter_design(example_data$survey, "weight", "area")  
margins <- census_margins(example_data$margins)  
fit <- fit_inter_sae(unemployed ~ sex + age + residence + area,  
  design, margins, family = "binomial")  
official <- with(example_data$survey,  
  weighted.mean(unemployed, weight))  
benchmarked <- benchmark_inter(fit, official)  
attr(benchmarked, "benchmark_achieved")
```

fit_inter_sae

*Fit and Inspect Intersectional Small Area Estimates***Description**

Reconstruct a coherent latent joint population table, fit a survey-weighted outcome model, and calculate augmented model-assisted estimates.

Usage

```
fit_inter_sae(formula, design, margins,
  family = c("gaussian", "binomial"), domain = NULL, prior = 0.5,
  max_iter = 1000L, tol = 1e-8, residual_correction = TRUE,
  max_cells = 1e6, model_args = list())

inter_estimates(object, statistic = c("mean", "total", "both"))

margin_diagnostics(object)

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

## S3 method for class 'inter_sae'
summary(object, ...)

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

## S3 method for class 'inter_sae'
plot(x, y = NULL, compare_direct = TRUE, ...)
```

Arguments

formula	Outcome model formula. Every predictor must occur among the census-margin dimensions.
design	An object returned by <code>inter_design()</code> .
margins	An object returned by <code>census_margins()</code> .
family	Gaussian or binomial response family.
domain	Character vector of output-domain dimensions.
prior	Positive seed added to every feasible latent population cell.
max_iter	Maximum generalized iterative-proportional-fitting iterations.
tol	Relative convergence tolerance.
residual_correction	Add a calibrated survey-residual correction.

max_cells	Maximum permitted latent-table size.
model_args	Optional arguments passed to <code>stats::glm()</code> .
object, x	A fitted <code>inter_sae</code> object.
statistic	Extract means/proportions, totals, or both.
y	Ignored.
compare_direct	Overlay direct survey estimates in the plot.
...	Additional printing or plotting arguments.

Details

For binomial outcomes the working model uses a quasibinomial likelihood so non-integer survey weights are interpreted as analytic weights. Returned estimates remain proportions on the response scale.

Value

`fit_inter_sae()` returns an `inter_sae` object. `inter_estimates()` and `margin_diagnostics()` return data frames. The summary method returns a `summary_inter_sae` object. The plot method returns the plotted data invisibly.

Examples

```
example_data <- inter_example()
design <- inter_design(example_data$survey, "weight", "area",
  "strata", "psu")
margins <- census_margins(example_data$margins)
fit <- fit_inter_sae(unemployed ~ sex + age + residence + area,
  design, margins, family = "binomial", domain = c("area", "sex"))
head(inter_estimates(fit))
max(margin_diagnostics(fit)$relative_error)
summary(fit)
```

identification_bounds *Diagnose Non-Identification and Compute Sensitivity Bounds*

Description

Measure the nullity left by overlapping aggregate margins and, when **lpSolve** is installed, optimize the target over latent population tables that satisfy the margins and optional association restrictions.

Usage

```

check_identifiability(object, tol = 1e-9)

identification_bounds(object, domain = NULL, gamma = Inf,
  statistic = c("mean", "total"))

inter_sensitivity(object, gamma = c(1, 1.25, 1.5, 2, Inf),
  domain = NULL, statistic = c("mean", "total"))

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

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

```

Arguments

object	A fitted <code>inter_sae</code> object.
tol	Numerical-rank tolerance.
domain	Optional named list defining one target domain.
gamma	Association-sensitivity value or vector. One fixes the fitted latent table; infinity imposes census margins only.
statistic	Bound a predicted domain mean/proportion or total.
x	Object to print.
...	Unused.

Details

Mean bounds use a Charnes–Cooper transformation of the linear-fractional target. They concern uncertainty in the latent census table conditional on the fitted cell predictions; they are not replacements for sampling intervals.

Value

`check_identifiability()` returns an `inter_identifiability` object. `identification_bounds()` returns an `inter_bounds` object. `inter_sensitivity()` returns a data frame.

Examples

```

example_data <- inter_example()
design <- inter_design(example_data$survey, "weight", "area")
margins <- census_margins(example_data$margins)
fit <- fit_inter_sae(unemployed ~ sex + age + residence + area,
  design, margins, family = "binomial")
check_identifiability(fit)
if (requireNamespace("lpSolve", quietly = TRUE)) {
  identification_bounds(fit, domain = list(area = "A1"), gamma = 1.5)
}

```

inter_bootstrap	<i>Bootstrap and Summarize Uncertainty</i>
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Description

Refit population reconstruction and the outcome model under each set of weights. Standard errors include sampling and refitting variation. Identification width can be reported separately.

Usage

```
inter_bootstrap(object, R = 199L,
  type = c("multiplier", "replicate"), seed = 123,
  level = 0.95, progress = FALSE)

decompose_uncertainty(object, bootstrap = NULL, bounds = NULL)

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

Arguments

object	A fitted inter_sae object.
R	Number of bootstrap replicates.
type	Multiplier or supplied-replicate-weight bootstrap.
seed	Random seed for multiplier bootstrap.
level	Confidence level.
progress	Print periodic progress messages.
bootstrap	Optional result from inter_bootstrap().
bounds	Optional result from identification_bounds().
x	Object to print.
...	Unused.

Value

inter_bootstrap() returns an inter_bootstrap object containing a summary and replicate matrix. decompose_uncertainty() returns a data frame.

Examples

```
## Not run:
example_data <- inter_example()
design <- inter_design(example_data$survey, "weight", "area", "strata", "psu")
margins <- census_margins(example_data$margins)
fit <- fit_inter_sae(unemployed ~ sex + age + residence + area,
  design, margins, family = "binomial")
```

```
boot <- inter_bootstrap(fit, R = 99)
boot

## End(Not run)
```

inter_design

Declare Survey Design Information and Census Margins

Description

Validate the two inputs required by `fit_inter_sae()`. Column names are supplied as strings to keep programmatic workflows unambiguous.

Usage

```
inter_design(data, weights, area, strata = NULL, psu = NULL,
  replicate_weights = NULL)

census_margins(margins, count = "Freq", structural_zeros = NULL,
  tolerance = 1e-8, reconcile = FALSE)

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

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

Arguments

<code>data</code>	Survey microdata.
<code>weights</code>	Name of the survey-weight column.
<code>area</code>	Name of the geographic-area column.
<code>strata</code>	Optional stratum-column name.
<code>psu</code>	Optional primary-sampling-unit column name.
<code>replicate_weights</code>	Optional character vector naming replicate-weight columns.
<code>margins</code>	A census table or named list of census tables.
<code>count</code>	Common name of the population-count column.
<code>structural_zeros</code>	Optional data frame of impossible category combinations. Missing values act as wildcards.
<code>tolerance</code>	Relative tolerance for equality of census-table totals.
<code>reconcile</code>	Whether inconsistent table totals may be proportionally rescaled.
<code>x</code>	Object to print.
<code>...</code>	Unused.

Value

inter_design() returns an object of class inter_design. census_margins() returns an object of class census_margins.

Examples

```
example_data <- inter_example()
design <- inter_design(example_data$survey, weights = "weight",
  area = "area", strata = "strata", psu = "psu")
margins <- census_margins(example_data$margins)
design
margins
```

simulate_inter_data	<i>Simulate an Intersectional Survey–Census Problem</i>
---------------------	---

Description

Create a reproducible problem with a binary unemployment outcome and partially identified age-by-residence associations within area and sex groups.

Usage

```
simulate_inter_data(seed = 123, n = 500, areas = paste0("A", 1:6))

inter_example()
```

Arguments

seed	Random-number seed.
n	Survey sample size.
areas	Distinct area labels.

Value

A list containing survey microdata, overlapping census margins, and the latent full population table used to generate the data.

Examples

```
example_data <- inter_example()
names(example_data)
head(example_data$survey)
lapply(example_data$margins, head)
```

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