

Package: regrake (via r-universe)

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Type Package

Title Regularized Survey Raking

Version 1.0.0

Description Calibrates survey weights to known population targets using regularized raking. Constraints are specified with a formula interface (for example, `rr_exact()`, `rr_l2()`, `rr_range()`, `rr_mean()`, `rr_var()`, and `rr_quantile()`). Supports common target formats including autumn-style proportions tables, raw or weighted population microdata, named-list targets (as in 'anesrake'), and 'survey' package design objects. Optimization follows Barratt et al. (2021) https://web.stanford.edu/~boyd/papers/pdf/optimal_representative_sampling.pdf and returns calibrated weights with balance and convergence diagnostics.

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print.raking_formula *Print method for raking_formula objects*

Description

Displays a human-readable representation of a raking formula. This function shows each term in the formula in a structured format, making it easy to understand complex formulas with multiple constraints.

Usage

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

Arguments

x A raking_formula object
 ... Additional arguments passed to other methods

Value

Invisibly returns the object

print.regrake	<i>Print method for regrake objects</i>
---------------	---

Description

Print method for regrake objects

Usage

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

Arguments

x	A regrake object
...	Additional arguments (ignored)

Value

Invisibly returns the object

prox_boolean_reg	<i>Proximal operator for boolean regularizer</i>
------------------	--

Description

Selects top-k weights and assigns them equal weight $1/k$. All other weights are set to zero. This is used for representative sample selection where exactly k samples should be selected.

Usage

```
prox_boolean_reg(w, lam, k)
```

Arguments

w	Input vector of weights
lam	Regularization parameter (unused, kept for interface consistency)
k	Number of samples to select

Value

Vector with k non-zero entries, each equal to $1/k$

prox_equality *Proximal operator for equality constraints*

Description

Proximal operator for equality constraints

Usage

```
prox_equality(x, target, rho)
```

Arguments

x	Input vector
target	Target vector
rho	Proximal parameter (unused for equality constraints)

Value

Projected vector equal to target

prox_equality_reg *Proximal operator for equality regularizer*

Description

Proximal operator for equality regularizer

Usage

```
prox_equality_reg(w, lam)
```

Arguments

w	Input vector
lam	Regularization parameter

Value

Original vector (identity operation)

prox_inequality	<i>Proximal operator for inequality constraints</i>
-----------------	---

Description

Proximal operator for inequality constraints

Usage

```
prox_inequality(x, target, rho, lower, upper)
```

Arguments

x	Input vector
target	Target vector (used for offset)
rho	Proximal parameter (unused for inequality constraints)
lower	Lower bound
upper	Upper bound

Value

Clipped vector within bounds relative to target

prox_kl	<i>Proximal operator for KL divergence loss</i>
---------	---

Description

Proximal operator for KL divergence loss

Usage

```
prox_kl(x, target, rho, scale = 0.5)
```

Arguments

x	Input vector
target	Target vector
rho	Proximal parameter
scale	Scale factor for KL divergence (default 0.5, matching the Python reference)

Value

Updated vector minimizing KL divergence plus proximal term

prox_kl_reg	<i>Proximal operator for KL regularizer</i>
-------------	---

Description

Proximal operator for KL regularizer

Usage

```
prox_kl_reg(w, lam, prior = NULL, limit = NULL)
```

Arguments

w	Input vector
lam	Regularization parameter
prior	Prior weights (default uniform)
limit	Optional upper bound on weight magnitudes

Value

Updated vector minimizing KL divergence plus proximal term

prox_least_squares	<i>Proximal operator for least squares loss</i>
--------------------	---

Description

Proximal operator for least squares loss

Usage

```
prox_least_squares(x, target, tau, diag_weight = 1)
```

Arguments

x	Input vector
target	Target vector
tau	Proximal parameter (1/rho)
diag_weight	Numeric scalar or vector of weights for each element (default 1)

Value

Updated vector minimizing weighted quadratic plus proximal term

prox_sum_squares_reg *Proximal operator for sum squares regularizer*

Description

Proximal operator for sum squares regularizer

Usage

```
prox_sum_squares_reg(w, lam)
```

Arguments

w	Input vector
lam	Regularization parameter

Value

Updated vector minimizing sum squares plus proximal term

regrake *Optimal representative sample weighting*

Description

Optimal representative sample weighting

Usage

```
regrake(
  data,
  formula,
  population_data,
  pop_type = c("raw", "weighted", "proportions", "anesrake", "survey", "survey_design"),
  pop_weights = NULL,
  regularizer = "entropy",
  lambda = 1,
  prior = NULL,
  k = NULL,
  bounds = c(0.1, 10),
  bounds_method = c("soft", "hard"),
  exact_tol = NULL,
  normalize = TRUE,
  control = list(),
  verbose = FALSE,
  ...
)
```

Arguments

<code>data</code>	A data.frame or tibble containing the sample data
<code>formula</code>	A formula specifying the raking constraints (e.g., <code>~ rr_exact(sex) + rr_l2(age)</code>)
<code>population_data</code>	Population data: a data.frame, list, or survey.design object (see <code>pop_type</code>)
<code>pop_type</code>	How population data is specified: <ul style="list-style-type: none"> • "raw": Raw population data (one row per unit) • "weighted": Population data with weights column • "proportions": Direct specification of target proportions (variable, level, target columns) • "anesrake": List of named numeric vectors (anesrake package format) • "survey": Data frame with margin, category, value columns • "survey_design": survey package design object
<code>pop_weights</code>	Column name in <code>population_data</code> containing weights (if <code>pop_type = "weighted"</code>)
<code>regularizer</code>	Regularization method ("entropy", "zero", "kl", or "boolean")
<code>lambda</code>	Regularization strength (default = 1)
<code>prior</code>	Optional prior weights used when <code>regularizer = "kl"</code> . Must be a positive numeric vector of length <code>nrow(data)</code> . If it does not sum to 1, it is normalized internally.
<code>k</code>	Number of samples to select (required for <code>regularizer = "boolean"</code>)
<code>bounds</code>	Numeric vector of length 2 specifying (min, max) allowed weight values. Weights returned sum to <code>n</code> (sample size), so <code>bounds = c(0.3, 3)</code> means each weight is between 0.3 and 3 times the "average" weight of 1. Default is <code>c(0.1, 10)</code> .
<code>bounds_method</code>	How to enforce bounds: <p>"soft" (Default) Uses regularizer clipping. Fast but bounds may be slightly violated when targets conflict with bounds. Asymmetric bounds are approximated as symmetric.</p> <p>"hard" Uses bounded simplex projection. Bounds are strictly enforced but optimization may be slower and targets may be less closely matched when bounds are binding.</p>
<code>exact_tol</code>	Optional tolerance for exact constraints. When non-NULL, all <code>rr_exact()</code> (and <code>rr_mean()</code>) constraints are converted to <code>rr_range()</code> with this margin. For example, <code>exact_tol = 0.02</code> means categorical proportions must be within +/- 2 percentage points of targets, and continuous means within +/- 0.02 of targets. Use <code>rr_range()</code> directly in the formula for per-variable control. Default is NULL (exact constraints enforced strictly).
<code>normalize</code>	Logical. If TRUE (default), continuous variables are automatically scaled by their target value for numerical stability. The achieved values are reported in original units. Set to FALSE to disable this behavior.
<code>control</code>	List of control parameters for the ADMM solver:

	margin_tol Margin-based convergence tolerance (default 1e-4). The ADMM solver scales this by problem size to achieve approximately this level of margin accuracy regardless of sample size or constraint count. For example, <code>margin_tol = 0.001</code> targets ~0.1% max margin error. Internally computes <code>eps = margin_tol / sqrt(m + 2*n)</code> . Set to NULL to use raw <code>eps_abs/eps_rel</code> instead.
	eps_abs Absolute convergence tolerance for ADMM residuals. Only used when <code>margin_tol</code> is NULL. Note: the effective tolerance scales with <code>sqrt(m + 2*n)</code> , so the same value behaves differently at different problem sizes. Default 1e-5.
	eps_rel Relative convergence tolerance for ADMM residuals. Only used when <code>margin_tol</code> is NULL. Default 1e-5.
	rho ADMM penalty parameter (default 50).
	maxiter Maximum ADMM iterations (default 5000).
<code>verbose</code>	Whether to print progress information
<code>...</code>	Additional arguments passed to methods

Value

An object of class "regrake" containing:

<code>weights</code>	The optimal weights (sum to n)
<code>balance</code>	Data frame comparing achieved vs target values with columns: constraint (e.g., "exact_sex"), type ("exact" or "l2"), variable, level, achieved, target, residual
<code>solution</code>	Full solution details from solver
<code>diagnostics</code>	Weight, convergence, and margin matching diagnostics

Examples

```
set.seed(42)
sample_data <- data.frame(
  sex = sample(c("M", "F"), 200, replace = TRUE, prob = c(0.6, 0.4)),
  age = sample(c("young", "old"), 200, replace = TRUE, prob = c(0.7, 0.3))
)
pop_targets <- data.frame(
  variable = c("sex", "sex", "age", "age"),
  level = c("M", "F", "young", "old"),
  target = c(0.49, 0.51, 0.45, 0.55)
)
result <- regrake(
  data = sample_data,
  formula = ~ rr_exact(sex) + rr_exact(age),
  population_data = pop_targets,
  pop_type = "proportions"
)
result
result$balance
```

rr_constraints

Raking Constraint Functions

Description

These functions specify constraint types for raking formulas. They are used within formula specifications passed to `regrake()`.

Usage

```
rr_l2(x)
```

```
rr_kl(x)
```

```
rr_exact(x)
```

```
rr_mean(x)
```

```
rr_var(x)
```

```
rr_quantile(x, p)
```

```
rr_range(x, ...)
```

```
rr_between(x, ...)
```

Arguments

x	Variable name (unquoted) to apply the constraint to
p	For <code>rr_quantile</code> , the quantile probability (0 to 1, e.g., 0.5 for median)
...	For <code>rr_range</code> / <code>rr_between</code> : either a single margin value (or named vector for level-specific margins), or separate lower and upper bounds. Examples: <code>rr_range(x, 0.02)</code> , <code>rr_range(x, c(A=0.01, B=0.02))</code> , <code>rr_range(x, 40, 45)</code> , <code>rr_range(x, lower=40, upper=45)</code>

Details

- `rr_exact()`: Exact equality constraint (weighted sum equals target exactly)
- `rr_l2()`: Soft L2/least squares constraint (penalizes deviation from target)
- `rr_kl()`: KL divergence constraint
- `rr_mean()`: Match the mean of a continuous variable (alias for `rr_exact` on continuous)
- `rr_var()`: Match the variance of a continuous variable
- `rr_quantile()`: Match a specific quantile of a continuous variable

Value

The input variable (these functions are markers for the formula parser)

Examples

```
# Match sex proportions exactly, age proportions with soft constraint
formula <- ~ rr_exact(sex) + rr_l2(age)

# Match categorical variable and continuous mean
formula <- ~ rr_exact(region) + rr_mean(income)

# Match mean and variance of a continuous variable
formula <- ~ rr_mean(income) + rr_var(income)

# Match median income
formula <- ~ rr_quantile(income, 0.5)
```

summary.regrake

Summary method for regrake objects

Description

Summary method for regrake objects

Usage

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

Arguments

object	A regrake object
...	Additional arguments (ignored)

Value

Invisibly returns the object

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