

Package: rclsp (via r-universe)

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Type Package

Title A Modular Two-Step Convex Optimization Estimator for Ill-Posed Problems

Version 2.0.0

Description Convex Least Squares Programming (CLSP) is a two-step estimator for solving underdetermined, ill-posed, or structurally constrained least-squares problems. It combines pseudoinverse-based estimation with convex-programming correction methods inspired by Lasso, Ridge, and Elastic Net to ensure numerical stability, constraint enforcement, and interpretability. The package also provides numerical stability analysis and CLSP-specific diagnostics, including partial R^2 , normalized RMSE (NRMSE), Monte Carlo t-tests for mean NRMSE, and condition-number-based confidence bands.

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canonize	<i>Construct the canonical design matrix $A = [C S; M Q]$ for CLSP.</i>
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Description

This method assembles the constraint matrix A from user-supplied or internally generated components — C , S , M , and Q — and assigns the corresponding right-hand side vector b . It is a required pre-step before solving a Convex Least Squares Programming (CLSP) problem.

Usage

```

canonize(
  object,
  problem = "",
  C = NULL,
  S = NULL,
  M = NULL,
  Q = NULL,
  b = NULL,
  m = NULL,
  p = NULL,
  i = 1L,
  j = 1L,
  zero_diagonal = FALSE
)

```

Arguments

object	An object of class "clsp".
problem	Character, optional. Structural template for matrix construction. One of: <ul style="list-style-type: none"> 'ap' or 'tm': allocation or tabular matrix problem. 'cmls' or 'rp': constrained modular least squares or RP-type. ' ' or other: General CLSP problems (user-defined C and/or M).
C , S , M	Numeric matrix or NULL. Blocks of the constraint matrix $A = [C S; M Q]$. If C and/or M are provided, the matrix A is constructed accordingly. If both are NULL and A is not yet defined, an error is raised.

Q	Numeric matrix or NULL. Externally supplied residual slack matrix used to adjust inequality constraints in M. Required only when $r > 1$. Encodes the sign pattern of residuals from the previous iteration and is used to construct the $[C S; M Q]$ canonical form. Defaults to a conformable zero matrix on the first iteration.
b	Numeric vector or NULL. Right-hand side vector. Must have as many rows as A . Required.
m, p	Integer or NULL. Dimensions of $X \in \mathbb{R}^{m \times p}$, relevant for allocation problems ('ap').
i, j	Integer, default = 1. Grouping sizes for row and column sum constraints in AP problems.
zero_diagonal	Logical, default = FALSE. If TRUE, enforces structural zero diagonals via identity truncation.

Details

Depending on the specified problem type, it can generate allocation, tabular matrix, or modular constraints and enforce optional diagonal exclusions. All missing blocks are padded to ensure conformability.

Value

An updated object of class "clsp".

Attributes Set

- A Numeric matrix. Canonical design matrix constructed from (C, S, M, Q).
- C_idx Integer vector of length 2 indicating the size of the C block.
- b Numeric vector. Conformable right-hand side vector.

clsp

Convex Least Squares Programming (CLSP) estimator.

Description

The Convex Least Squares Programming (CLSP) estimator solves underdetermined, ill-posed, or structurally constrained least-squares problems using a modular two-step approach. The first step computes a pseudoinverse-based estimate, and the second step applies a convex correction (Lasso, Ridge, or Elastic Net) to ensure numerical stability, constraint enforcement, and interpretability.

Usage

```
clsp(
  problem = "",
  C = NULL,
  S = NULL,
  M = NULL,
```

```

b = NULL,
m = NULL,
p = NULL,
i = 1L,
j = 1L,
zero_diagonal = FALSE,
r = 1L,
Z = NULL,
rcond = FALSE,
tolerance = NULL,
iteration_limit = NULL,
final = TRUE,
alpha = NULL,
cond_tolerance = NULL,
...
)

```

Arguments

problem	character scalar, optional Structural template for matrix construction. One of: <ul style="list-style-type: none"> 'ap' or 'tm': allocation or tabular matrix problem. 'cmls' or 'rp': constrained modular least squares or RP-type. ' ' or other: general CLSP problems (user-defined C and/or M).
C, S, M	numeric matrix or NULL Blocks of the constraint matrix $A = \begin{bmatrix} C & S \\ M & Q \end{bmatrix}$. If C and/or M are provided, the matrix A is constructed accordingly. If both are NULL and A is not yet defined, an error is raised.
b	numeric vector or NULL Right-hand-side vector. Must have as many rows as A . Required.
m, p	integer scalar or NULL Dimensions of $X \in \mathbb{R}^{m \times p}$, relevant for allocation problems ('ap').
i, j	integer scalar, default = 1 Grouping sizes for row and column-sum constraints in AP problems.
zero_diagonal	logical scalar, default = FALSE If TRUE, enforces structural zero diagonals via identity truncation.
r	integer scalar, default = 1 Number of refinement iterations for the pseudoinverse-based estimator. When $r > 1$, the slack block Q is updated iteratively to improve feasibility in underdetermined or ill-posed systems.
Z	numeric matrix or NULL A symmetric idempotent matrix (projector) defining the subspace for Bott–Duffin pseudoinversion. If NULL, the identity matrix is used, reducing to the Moore–Penrose case.
rcond	numeric scalar or logical scalar, default = FALSE Regularization parameter for the Moore–Penrose and Bott–Duffin inverses, providing numerically stable inversion and ensuring convergence of singular values. If TRUE, an automatic tolerance equal to <code>tolerance</code> is applied. If set to a numeric value, it specifies the relative cutoff below which small singular values are treated as zero.

tolerance	numeric scalar or NULL, default = NULL Convergence tolerance for NRMSE change between iterations.
iteration_limit	integer scalar or NULL, default = NULL Maximum number of iterations allowed in the refinement loop.
final	logical scalar, default = TRUE If TRUE, a convex programming problem is solved to refine $\hat{\mathbf{z}}$. The resulting solution \mathbf{z} minimizes a weighted ℓ_1/ℓ_2 norm around $\hat{\mathbf{z}}$ subject to $\mathbf{A}\mathbf{z} = \mathbf{b}$.
alpha	numeric scalar, numeric vector, or NULL, default = NULL Regularization parameter: <ul style="list-style-type: none"> • $\alpha = 0$: Lasso (ℓ_1 norm) • $\alpha = 1$: Ridge (ℓ_2 norm) • $0 < \alpha < 1$: Elastic Net. If a numeric scalar is provided, that value is used after clipping to $[0, 1]$. If a numeric vector is provided, each candidate is evaluated via a full solve, and the α with the smallest NRMSE is selected. If NULL, α is chosen automatically according to $\alpha = \min \left(1, \frac{\text{NRMSE}_{\alpha=0}}{\text{NRMSE}_{\alpha=0} + \text{NRMSE}_{\alpha=1} + \text{tolerance}} \right)$
cond_tolerance	numeric scalar or NULL, default = NULL Singular-value cutoff for the custom condition number function. If NULL, the implementation uses an internal relative cutoff of 1e-14.
...	Optional. Additional arguments passed to the CVXR solver backend.

Details

This estimator unifies pseudoinverse-based least squares with convex programming correction. The pseudoinverse step computes an initial solution $\mathbf{z}^{(r)}$ iteratively via the Moore–Penrose or Bott–Duffin inverse. The convex step then refines \mathbf{z} by minimizing a mixed ℓ_1/ℓ_2 norm under equality constraints $\mathbf{A}\mathbf{z} = \mathbf{b}$. The method supports allocation problems (AP), constrained modular least squares (CMLS), and general CLSP formulations.

To ensure cross-platform reproducibility, all CLSP implementations use a modified condition number function based on singular values, with a relative cutoff equal to $\text{cond_tolerance} * \text{the largest singular value}$.

Value

An object of class "clsp" representing the fitted Convex Least Squares Programming (CLSP) model. The object is a named list containing all initialized fields and solver results. Class-specific methods such as `summary.clsp()`, `corr.clsp()`, and `ttest.clsp()` can be used to extract, analyze, and summarize the results.

See Also

[CVXR](#)

Examples

```

## Not run:
## Example: CMLS (RP) estimation with stationary-point constraints

set.seed(123456789)

# sample (dataset)
k <- 500L                                # number of observations
p <- 6L                                   # number of regressors
c0 <- 1                                  # sum of coefficients

D      <- matrix(NA_real_, nrow = k, ncol = p)
D[, 1] <- 1.0                             # constant
D[, 2:p] <- matrix(rnorm(k * (p - 1)), k, p - 1)

b_true <- rnorm(p)
b_true <- (b_true / sum(b_true)) * c0      # normalize to sum = c

e      <- matrix(rnorm(k), ncol = 1)
y      <- D %>% b_true + e

# build blocks for CLSP (CMLS)
b <- rbind(
  matrix(c0, ncol = 1),                    # sum of coefficients
  matrix(0, nrow = k - 2, ncol = 1),
  matrix(0, nrow = k - 1, ncol = 1),
  matrix(y, ncol = 1)
)

C <- rbind(
  matrix(1, nrow = 1, ncol = p),          # row of ones
  diff(D, differences = 2),               # 2nd differences
  diff(D, differences = 1)               # 1st differences
)

# diagonal sign-matrix for 2nd differences
S <- rbind(
  matrix(0, nrow = 1, ncol = k - 2),
  diag(sign(diff(as.numeric(y), differences = 2))),
  matrix(0, nrow = k - 1, ncol = k - 2)
)

# model
model <- rclsp::clsp(
  problem = "cmls",
  b       = b,
  C       = C,
  S       = S,
  M       = D,
  r       = 1L,                             # no refinement
  alpha   = 1.0                             # MNBLUE solution
)

```

```

# results
print("true beta (x_M):")
print(round(b_true, 4))

print("beta hat (x_M hat):")
print(round(model$x, 4))

print(model)

# bootstrap t-test
tt <- rclsp::ttest(
  model,
  sample_size = 30L,
  seed        = 123456789L,
  distribution = rnorm,
  partial     = TRUE
)

print("Bootstrap t-test:")
print(tt)

## End(Not run)

```

corr

Compute the structural correlogram of the CLSP constraint system.

Description

This method performs a row-deletion sensitivity analysis on the canonical constraint matrix $[C|S]$, denoted as C_{canon} , and evaluates the marginal effect of each constraint row on numerical stability, angular alignment, and estimator sensitivity.

Usage

```
corr(object, reset = FALSE, threshold = 0)
```

Arguments

object	An object of class "clsp".
reset	Logical, default = FALSE. If TRUE, forces recomputation of all diagnostic values.
threshold	Numeric, default = 0. If positive, limits the output to constraints with $\text{RMSA}_i \geq \text{threshold}$.

Details

For each row i in C_{canon} , it computes:

- The Root Mean Square Alignment (RMSA_i) with all other rows $j \neq i$.
- The change in condition numbers $\kappa(C)$, $\kappa(B)$, and $\kappa(A)$ when row i is deleted.
- The effect on estimation quality: changes in NRMSE and in the norms of \hat{z} , z , and x .

Additionally, it computes the total RMSA statistic across all rows, summarizing the overall angular alignment of the constraint block.

Value

A named list containing per-row diagnostic values:

constraint Numeric vector of constraint indices (1-based).

rmsa_i Numeric vector of RMSA_i values.

rmsa_dkappaC Numeric vector of $\Delta\kappa(C)$ after deleting row i .

rmsa_dkappaB Numeric vector of $\Delta\kappa(B)$ after deleting row i .

rmsa_dkappaA Numeric vector of $\Delta\kappa(A)$ after deleting row i .

rmsa_dnormse Numeric vector of ΔNRMSE after deleting row i .

rmsa_dzhat Numeric vector of $\Delta\|\hat{z}\|_2$ after deleting row i .

rmsa_dz Numeric vector of $\Delta\|z\|_2$ after deleting row i .

rmsa_dx Numeric vector of $\Delta\|x\|_2$ or $\Delta\|x\|_F$ after deleting row i .

ttest	<i>Perform bootstrap or Monte Carlo t-tests on the NRMSE statistic from the CLSP estimator.</i>
-------	---

Description

This function either (a) resamples residuals via a nonparametric bootstrap to generate an empirical NRMSE sample, or (b) produces synthetic right-hand side vectors b from a user-defined or default distribution and re-estimates the model. It tests whether the observed NRMSE significantly deviates from the null distribution of resampled or simulated NRMSE values.

Usage

```
ttest(
  object,
  reset = FALSE,
  sample_size = 50L,
  seed = NULL,
  distribution = NULL,
  partial = FALSE,
  simulate = FALSE
)
```

Arguments

<code>object</code>	An object of class "c1sp".
<code>reset</code>	Logical, default = FALSE. If TRUE, forces recomputation of the NRMSE null distribution.
<code>sample_size</code>	Integer, default = 50. Size of the Monte Carlo simulated sample under H0.
<code>seed</code>	Integer or NULL, default = NULL. Optional random seed to override the default.
<code>distribution</code>	Function or NULL, default = NULL. Distribution for generating synthetic b vectors. One of: <code>rnorm</code> , <code>runif</code> , or a custom RNG function. Defaults to standard normal.
<code>partial</code>	Logical, default = FALSE. If TRUE, runs the t-test on the partial NRMSE: during simulation, the C-block entries are preserved and the M-block entries are simulated.
<code>simulate</code>	Logical, default = FALSE. If TRUE, performs a parametric Monte Carlo simulation by generating synthetic right-hand side vectors b. If FALSE (default), executes a nonparametric bootstrap procedure on residuals without re-estimation.

Value

A named list containing test results and null distribution statistics:

p_one_left $P(\text{nmse} \leq \text{null mean})$

p_one_right $P(\text{nmse} \geq \text{null mean})$

p_two_sided 2-sided t-test p-value

nmse Observed value

mean_null Mean of null distribution

std_null Standard deviation of null distribution

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