

# Package ‘capr’

January 24, 2026

**Title** Covariate Assisted Principal Regression

**Version** 0.2.0

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**Description** Covariate Assisted Principal Regression (CAPR) for multiple covariance-matrix outcomes. The method identifies (principal) projection directions that maximize the log-likelihood of a log-linear regression model of the covariates. See Zhao et al. (2021), ``Covariate Assisted Principal Regression for Covariance Matrix Outcomes'' <[doi:10.1093/biostatistics/kxz057](https://doi.org/10.1093/biostatistics/kxz057)>.

**License** GPL-3

**Encoding** UTF-8

**RoxxygenNote** 7.3.3

**LinkingTo** Rcpp, RcppArmadillo

**Imports** Rcpp, MASS

**Suggests** testthat, roxygen2

**SystemRequirements** C++17

**URL** <https://github.com/rluo/capr>

**BugReports** <https://github.com/rluo/capr/issues>

**NeedsCompilation** yes

**Repository** CRAN

**Date/Publication** 2026-01-24 10:50:02 UTC

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**capr***Covariate Assisted Principal (CAP) Regression***Description**

Fits CAP components sequentially for principal direction vectors  $\gamma^{(k)}$  and regression coefficients  $\beta^{(k)}$ ,  $k = 1, \dots, K$ . Each component is estimated via a flip-flop algorithm with optional orthogonalization of successive directions.

**Usage**

```
capr(
  S,
  X,
  K,
  B.init = NULL,
  Gamma.init = NULL,
  weight = NULL,
  max_iter = 200L,
  tol = 1e-06,
  orth = TRUE,
  n.init = 10L
)
```

**Arguments**

|            |   |
|------------|---|
| S          | Numeric 3D array of size $p \times p \times n$ (for example, a stack of covariance matrices).   |
| X          | Numeric matrix $n \times q$ (design matrix), created for example by <code>model.matrix()</code> .   |
| K          | Integer scalar; number of components ( $K \geq 1$ ).  |
| B.init     | Initial value of the coefficient array $B \in \mathbb{R}^{q \times n.init \times K}$ (default: zero 3D array).  |
| Gamma.init | Initial value of the principal direction array $\Gamma \in \mathbb{R}^{p \times n.init \times K}$ (default: random Gaussian 3D array).                          |
| weight     | Numeric vector of length $n$ (default <code>rep(1, n)</code> ); each element should be proportional to the sample size for the corresponding slice $S[, , i]$ . |
| max_iter   | Integer scalar; maximum flip-flop iterations per component (default 200).   |
| tol        | Positive numeric scalar; convergence tolerance (default 1e-6).  |

|                     |   |
|---------------------|---|
| <code>orth</code>   | Logical scalar; if TRUE (default), enforce orthogonality of successive $\gamma^{(k)}$ . If FALSE, no orthogonalization is performed (which may yield identical components). |
| <code>n.init</code> | Integer scalar; number of random initializations (default 10). If <code>B.init</code> and <code>Gamma.init</code> are both supplied, <code>n.init</code> is ignored.        |

## Details

For component  $k$ , CAP solves

$$\min_{\beta^{(k)}, \gamma^{(k)}} \frac{1}{2} \sum_{i=1}^n (\mathbf{x}_i^\top \beta^{(k)}) T_i + \frac{1}{2} \sum_{i=1}^n \gamma^{(k)\top} S_i \gamma^{(k)} \exp(-\mathbf{x}_i^\top \beta^{(k)})$$

subject to

$$\gamma^{(k)\top} H \gamma^{(k)} = 1$$

and, for  $k > 1$ ,

$$\Gamma^{(k-1)\top} \gamma^{(k)} = \mathbf{0}.$$

Here  $T_i$  denotes the weight for slice  $i$ ,  $S_i$  is the  $i$ -th covariance slice, and  $H$  is the positive definite matrix used for the orthogonality constraint (see Zhao et al., 2021). The algorithm fits  $\gamma^{(k)}$  and  $\beta^{(k)}$  sequentially with multiple random initializations and returns the solution pair that minimizes the negative log-likelihood.

## Value

A list of class `capr` with:

|                      |  |
|----------------------|--|
| <code>B</code>       | numeric matrix $q \times K$ whose $k$ -th column stores $\beta^{(k)}$  |
| <code>Gamma</code>   | numeric matrix $p \times K$ whose $k$ -th column stores $\gamma^{(k)}$ |
| <code>loglike</code> | negative log-likelihood, up to constant scaling and shift              |
| <code>S</code>       | 3D array used for fitting  |
| <code>X</code>       | design matrix used for fitting   |
| <code>weight</code>  | weight values used for fitting   |

## References

Zhao, Y., Wang, B., Mostofsky, S. H., Caffo, B. S., & Luo, X. (2021). "Covariate assisted principal regression for covariance matrix outcomes." *Biostatistics*, 22(3), 629-645.

## Examples

```
simu.data <- simu.capr(seed = 123L, n = 120L)
K <- 2L
fit <- capr(
  S = simu.data$S,
  X = simu.data$X,
  K = K
)
print(fit)
```

---

capr.boot*Bootstrap confidence intervals for CAP coefficients*

---

## Description

Generates bootstrap inference for the CAP regression coefficients while holding the fitted directions  $\Gamma$  fixed. Each replicate samples the covariance slices  $S[, , i]$  with replacement, projects them onto the fixed directions to obtain component-specific variances, and re-solves the  $\beta^{(k)}$  equations. Quantile-based confidence intervals are returned for every predictor/component pair.

## Usage

```
capr.boot(
  fit,
  nboot = 1000L,
  level = 0.95,
  max_iter = 100L,
  tol = 1e-06,
  seed = NULL
)
```

## Arguments

|          |  |
|----------|--|
| fit      | A <code>capr</code> fit containing components B and Gamma. |
| nboot    | Integer; number of bootstrap replicates.                   |
| level    | Confidence level for the returned intervals.               |
| max_iter | Maximum Newton iterations for solving $\beta$ .            |
| tol      | Convergence tolerance for the Newton solver.               |
| seed     | Optional integer seed for reproducibility.                 |

## Value

A list of class `capr.boot` with:

|                    |   |
|--------------------|---|
| beta               | bootstrap average of $\beta$ with dimension $q \times K$      |
| ci_lower, ci_upper | Matrices $q \times K$ with the lower/upper confidence limits. |
| level              | The requested confidence level.                               |

## Examples

```
simu.data <- simu.capr(seed = 123L, n = 120L)
K <- 3L
fit <- capr(
  S = simu.data$S,
  X = simu.data$X,
```

```

K = K
)
capr.boot(fit, nboot = 10L, level = 0.95, seed = 42L)

```

---

**cosine\_similarity**      *Cosine similarity between numeric vectors*

---

### Description

Computes the cosine of the angle between two numeric vectors. Both vectors must have equal length and non-zero Euclidean norms.

### Usage

```
cosine_similarity(a, b, eps = 1e-12)
```

### Arguments

|      |   |
|------|---|
| a, b | Numeric vectors of equal length.  |
| eps  | Non-negative numeric tolerance used to guard against division by zero. Defaults to 1e-12. |

### Value

A scalar double value in [-1, 1] representing the cosine similarity between a and b.

### Examples

```

cosine_similarity(c(1, 2, 3), c(1, 2, 3))
cosine_similarity(c(1, 0), c(0, 1))
cosine_similarity(c(1, 2), c(-1, -2))

```

---

FG

*Flury-Gautschi Common Principal Components*

---

### Description

Implements the Flury & Gautschi (1986) (FG) iterative algorithm and a variant to estimate a common loading matrix across multiple covariance matrices. Each iteration cycles over all ordered pairs of variable indices and updates a (2 x 2) rotation so that the transformed matrices share diagonal structure.

### Usage

```

FG(cov_array, p = NULL, m = NULL, maxit = 30L)

FG2(cov_array, p = NULL, m = NULL, maxit = 30L)

```

## Arguments

|                        |   |
|------------------------|---|
| <code>cov_array</code> | Numeric 3D array of shape $p \times p \times m$ containing covariance matrices in its $m$ slices.       |
| <code>p</code>         | Optional integer specifying the matrix dimension; defaults to <code>dim(cov_array)[1]</code> .          |
| <code>m</code>         | Optional integer specifying the number of matrices/slices; defaults to <code>dim(cov_array)[3]</code> . |
| <code>maxit</code>     | Integer scalar; number of outer iterations of the algorithm.  |

## Details

Two solvers are exported:

`FG()` The original FG algorithm.

`FG2()` An alternative algorithm by Eslami et al. (2013).

## Value

A  $p \times p$  numeric matrix of estimated common loadings.

## References

- Flury, B. N. (1984). "Common Principal Components in k Groups." *Journal of the American Statistical Association*, 79, 892-898.
- Flury, B. N., & Gautschi, W. (1986). "An Algorithm for Simultaneous Orthogonal Transformation of Several Positive Definite Symmetric Matrices to Nearly Diagonal Form." *SIAM Journal on Scientific and Statistical Computing*, 7(1), 169-184.
- Eslami, A., Qannari, E. M., Kohler, A., & Bougeard, S. (2013). "General Overview of Methods of Analysis of Multi-Group Datasets." *Revue des Nouvelles Technologies de l'Information*, 25, 108-123.

## Examples

```
set.seed(1)
p <- 3
m <- 4
mats <- replicate(m,
{
  A <- matrix(rnorm(p * p), p, p)
  crossprod(A)
},
simplify = FALSE
)
cov_cube <- array(NA_real_, dim = c(p, p, m))
for (k in 1:m) cov_cube[, , k] <- mats[[k]]
FG(cov_cube, maxit = 5)
FG2(cov_cube, maxit = 5)
```

`log_deviation_from_diagonality`  
*Log deviation from diagonality*

**Description**

Evaluates the Flury-Gautschi log-deviation criterion for a collection of covariance matrices transformed by a loading matrix.

**Usage**

```
log_deviation_from_diagonality(S_cube, nval, B)
```

**Arguments**

|                     |   |
|---------------------|---|
| <code>S_cube</code> | Numeric 3D array of shape $p \times p \times n$ containing covariance matrices in its slices. |
| <code>nval</code>   | Numeric vector of length $n$ giving weights for each matrix.                                  |
| <code>B</code>      | Numeric $p \times p$ orthonormal matrix applied to the covariance slices.                     |

**Value**

Numeric scalar value equal to  $\sum_i n_i (\log \det \text{diag}(B^\top S_i B) - \log \det(B^\top S_i B)) / (\sum_i n_i)$ .

**Examples**

```
covs <- array(diag(2), dim = c(2, 2, 1))
log_deviation_from_diagonality(covs, 1, diag(2))
```

`plot.capr`

*Plot deviation diagnostics by component count*

**Description**

For a fitted CAP regression, plots two diagnostics across the first  $K$  components: (1) the negative log-likelihood returned by `capr()` and (2) the log deviation-from-diagonality (DfD) for the loading matrix formed by the first  $k$  directions. Both curves help assess the gain from adding components.

**Usage**

```
## S3 method for class 'capr'
plot(x, ...)
```

## Arguments

- x A `capr` object returned by [capr\(\)](#).
- ... Additional arguments passed to [graphics::plot\(\)](#) and applied to both panels (for example, `pch`, `col`, or axis limits).

## Details

The DfD criterion for the first  $k$  directions  $\Gamma^{(k)}$  is

$$\text{DfD}(\Gamma^{(k)}) = \left( \prod_{i=1}^n \nu\left(\Gamma^{(k)\top} S_i \Gamma^{(k)} / T_i\right)^{T_i} \right)^{1/\sum_i T_i},$$

where

$$\nu(A) = \frac{\det\{\text{diag}(A)\}}{\det(A)}$$

for a positive definite matrix  $A$ . The curve shows  $\log \text{DfD}(\Gamma^{(k)})$ . A common choice for  $k$  is the last point before a sudden jump in the negative log-likelihood or log-DfD curve.

## Value

Invisibly returns the numeric vector of log deviation values (one per component).

## See Also

[log\\_deviation\\_from\\_diagonality\(\)](#)

## Examples

```
sim <- simu.capr(seed = 123L, n = 120L)
fit <- capr(S = sim$S, X = sim$X, K = 3L)
plot(fit)
```

**print.capr**

*Print method for CAP regression fits*

## Description

Formats the coefficient matrix  $\hat{B}$  returned by [capr\(\)](#) in a linear-regression style table, showing the estimate for each predictor and component.

## Usage

```
## S3 method for class 'capr'
print(x, digits = max(3L, getOption("digits") - 3L), ...)
```

**Arguments**

- x An object of class `capr`, typically the result of `capr()`.
- digits Number of significant digits to show when printing numeric values.
- ... Additional arguments passed on to `print.data.frame()`.

**Value**

The input object `x`, invisibly.

**Examples**

```
simu.data <- simu.capr(seed = 123L, n = 120L)
K <- 2L
fit <- capr(
  S = simu.data$S,
  X = simu.data$X,
  K = K
)
print(fit)
```

print.capr.boot

*Print method for capr.boot objects***Description**

Displays bootstrap coefficient estimates and their confidence intervals component by component as compact tables.

**Usage**

```
## S3 method for class 'capr.boot'
print(x, digits = max(4L, getOption("digits") - 4L), ...)
```

**Arguments**

- x An object of class `capr.boot`, typically produced by `capr.boot()`.
- digits Number of significant digits to show when printing numeric values.
- ... Additional arguments passed on to `print.data.frame()`.

**Value**

The input object `x`, invisibly.

## Examples

```
simu.data <- simu.capr(seed = 123L, n = 120L)
K <- 2L
fit <- capr(
  S = simu.data$S,
  X = simu.data$X,
  K = K
)
fit.boot <- capr.boot(
  fit = fit,
  nboot = 10L,
  max_iter = 20L,
  tol = 1e-6,
  seed = 123L
)
print(fit.boot)
```

**simu.capr**

*Simulate covariance matrices compatible with capr()*

## Description

Generates a simple synthetic dataset for CAP regression consisting of a covariance cube, design matrix, and the latent orthogonal directions used to build the covariance slices.

## Usage

```
simu.capr(seed = 123L, n = 120L)
```

## Arguments

- |      |  |
|------|--|
| seed | Integer seed used for reproducibility.       |
| n    | Number of observations (slices) to generate. |

## Value

A list with components:

- |         |   |
|---------|---|
| S       | Array of dimension $p \times p \times n$ holding the simulated covariance matrices. |
| X       | Design matrix of size $n \times 2$ with an intercept and a Bernoulli covariate.     |
| Q       | Orthogonal matrix whose columns are the latent directions.                          |
| BetaMat | True coefficient matrix used to form the eigenvalues.                               |
| H       | Average covariance matrix $\frac{1}{n} \sum_i S_i$ .                                |
| p, n    | The dimension and sample size supplied to the generator.                            |

**Examples**

```
sim <- simu.capr(seed = 10, n = 50)
str(sim$S)
```

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