

# Package ‘convergenceDFM’

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**Type** Package

**Title** Convergence and Dynamic Factor Models

**Version** 0.1.4

**Description** Tests convergence in macro-financial panels combining Dynamic Factor Models (DFM) and mean-reverting Ornstein-Uhlenbeck (OU) processes. Provides: (i) static/approximate DFMs for large panels with VAR/VECM stability checks, Portmanteau tests and rolling out-of-sample  $R^2$ , following Stock and Watson (2002) <doi:10.1198/073500102317351921> and the Generalized Dynamic Factor Model of Forni, Hallin, Lippi and Reichlin (2000) <doi:10.1162/003465300559037>; (ii) cointegration analysis à la Johansen (1988) <doi:10.1016/0165-1889(88)90041-3>; (iii) OU-based convergence and half-life summaries grounded in Uhlenbeck and Ornstein (1930) <doi:10.1103/PhysRev.36.823> and Vasicek (1977) <doi:10.1016/0304-405X(77)90016-2>; (iv) robust inference via 'sandwich' HC/HAC estimators (Zeileis (2004) <doi:10.18637/jss.v011.i10>) and regression diagnostics ('lmtest'); and (v) optional PLS-based factor preselection (Mevik and Wehrens (2007) <doi:10.18637/jss.v018.i02>). Functions emphasize reproducibility and clear, publication-ready summaries.

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---

choose_var_lag	<i>Select optimal VAR lag order with multiple criteria</i>
----------------	--

---

## Description

Determines the optimal lag order for a Vector Autoregression model using information criteria, stability checks, serial correlation tests, and optional out-of-sample validation.

## Usage

```
choose_var_lag(
  F_combined,
  lag.max = 4,
  type = "const",
  p_pref = c("SC(n)", "HQ(n)"),
```

```

alpha = 0.05,
oos_eval = TRUE,
oos_start = 0.7,
verbose = TRUE
)

```

### Arguments

<code>F_combined</code>	Numeric matrix (T x K) of factor scores to be modeled.
<code>lag_max</code>	Integer. Maximum lag order to consider. Default is 4.
<code>type</code>	Character string. Type of deterministic terms: "const" (default), "trend", "both", or "none".
<code>p_pref</code>	Character vector. Preferred information criteria for initial selection. Default is <code>c("SC(n)", "HQ(n)")</code> .
<code>alpha</code>	Numeric. Significance level for serial correlation test. Default is 0.05.
<code>oos_eval</code>	Logical. Should out-of-sample evaluation be performed? Default is TRUE.
<code>oos_start</code>	Numeric. Proportion of sample to use for training in OOS validation. Default is 0.7.
<code>verbose</code>	Logical; print progress information. Default TRUE.

### Details

The function combines multiple selection criteria: (1) information criteria (AIC, BIC, HQ), (2) VAR stability (eigenvalue modulus < 1), (3) Portmanteau test for serial correlation, and (4) out-of-sample forecast performance. Returns the model that best balances these considerations.

### Value

List with components:

- `p` Selected optimal lag order.
- `fit` Fitted VAR model object of class `varest`.
- `roots_ok` Logical indicating if stability condition is satisfied.
- `serial_ok` Logical indicating if serial correlation test passed.
- `oos_mse` Out-of-sample mean squared error (if `oos_eval = TRUE`).

---

deltaR2\_ou

*Incremental R-squared from X in OU model*

---

### Description

Computes the incremental explanatory power (delta R-squared) contributed by X factors in predicting Y factors, both in-sample and out-of-sample.

**Usage**

```
deltaR2_ou(
  results_robust,
  lag = 1,
  oos = TRUE,
  seed = 123,
  ridge = 1e-08,
  verbose = TRUE
)
```

**Arguments**

results_robust	Output from run_complete_factor_analysis_robust()
lag	Number of lags for the model (default: 1)
oos	Logical, perform out-of-sample validation (default: TRUE)
seed	Random seed for reproducibility (default: 123)
ridge	Ridge regularization parameter (default: 1e-08)
verbose	Logical, print progress messages (default: TRUE)

**Value**

List with components:

in_sample	Data frame with R2_full, R2_baseline, and deltaR2.
per_equation	Vector of deltaR2 for each Y factor equation.
OOS	List with out-of-sample RMSE and deltaR2 (if oos = TRUE).

---

diagnose_data	<i>Diagnose and prepare data matrices</i>
---------------	---

---

**Description**

Performs data validation, missing value imputation, and variance checks on input matrices to prepare them for factor analysis. Handles dimension compatibility, NA values, and zero-variance columns.

**Usage**

```
diagnose_data(X_matrix, Y_matrix, verbose = TRUE)
```

**Arguments**

X_matrix	Numeric matrix or data frame of X variables (e.g., Marxist prices).
Y_matrix	Numeric matrix or data frame of Y variables (e.g., market prices/CPI).
verbose	Logical; print diagnostic information. Default TRUE.

**Details**

The function:

- Converts to matrix format if needed
- Validates dimensional compatibility
- Imputes missing values via interpolation (using zoo if available)
- Adds minimal noise to zero-variance columns
- Reports diagnostic information

**Value**

List with components:

`X_matrix` Cleaned and prepared X matrix.

`Y_matrix` Cleaned and prepared Y matrix.

---

estimate\_DFM

*Estimate Dynamic Factor Model with VAR dynamics*

---

**Description**

Estimates a Dynamic Factor Model by extracting factors via PLS and modeling their dynamics with a Vector Autoregression. Includes automatic lag selection, robust inference, and optional out-of-sample evaluation.

**Usage**

```
estimate_DFM(
  factors_data,
  p = 2,
  compute_oos = TRUE,
  hc_type = "HC3",
  verbose = TRUE
)
```

**Arguments**

<code>factors_data</code>	List containing PLS-extracted factor scores ( <code>scores_X</code> , <code>scores_Y</code> ) and related objects.
<code>p</code>	Integer. VAR lag order. If NULL, selected automatically. Default is 2.
<code>compute_oos</code>	Logical. Should out-of-sample diagnostics be computed? Default is TRUE.
<code>hc_type</code>	Character string. Heteroskedasticity-consistent SE type. Default is "HC3".
<code>verbose</code>	Logical; print progress and diagnostic information. Default TRUE.

**Details**

This function models the joint dynamics of X and Y factors using a VAR. It performs stability checks, tests for serial correlation, computes robust standard errors, and optionally evaluates forecast performance out-of-sample.

**Value**

List with components:

var\_fit Fitted VAR model on combined factors.

p\_used VAR lag order used.

robust\_se Matrices of robust standard errors.

diagnostics List of diagnostic tests (stability, serial correlation).

oos\_metrics Out-of-sample forecast evaluation (if requested).

---

estimate\_factor\_OU      *Estimate Factor Ornstein-Uhlenbeck model (Stan if available)*

---

**Description**

Estimates a multivariate OU with cross-equation coupling  $Y_t$  depending on lagged  $X_{t-1}$  via a  $\beta$  matrix. Uses **cmdstanr** when available, otherwise **rstan**, with a discrete AR(1) fallback.

**Usage**

```
estimate_factor_OU(
  factors_data,
  data_prep = NULL,
  chains = 4,
  iter = 2000,
  seed = 1234,
  adapt1 = 0.98,
  adapt2 = 0.999,
  mtd1 = 12,
  mtd2 = 15,
  verbose = TRUE
)
```

**Arguments**

factors\_data      List with scores\_X, scores\_Y.  
 data\_prep        Optional preprocessed data (reserved).  
 chains, iter, seed  
                   Stan sampling controls.  
 adapt1, adapt2, mtd1, mtd2  
                   Advanced Stan controls.  
 verbose          Logical; print progress/details.

**Value**

A list with posterior medians ( $\phi$ ,  $\mu$ ,  $\beta$ ), half-lives, coupling strength, pseudo-R2, and the fitted Stan object.

**Examples**

```
# Create toy factor data
set.seed(123)
n <- 50
X_scores <- matrix(rnorm(n * 2), n, 2)
Y_scores <- matrix(rnorm(n * 2), n, 2)
factors_data <- list(scores_X = X_scores, scores_Y = Y_scores)

# Estimate OU model (reduce iterations for speed)
ou_result <- estimate_factor_OU(factors_data, chains = 2, iter = 500,
                               verbose = FALSE)

# Check half-lives
print(ou_result$half_lives_Y)
```

---

make_X_innovations	<i>Extract X innovations from VAR model</i>
--------------------	---

---

**Description**

Computes structural shocks (innovations) for X factors by fitting a VAR model and extracting residuals.

**Usage**

```
make_X_innovations(results, p = NULL)
```

**Arguments**

results	List. Output from main analysis.
p	Integer. VAR lag order. If NULL, uses order from DFM estimation.

**Value**

List with components:

shocks Matrix (T x Kx) of structural shocks with NA padding for initial lags.

var\_fit Fitted VAR model object.

lag\_used Integer lag order used.

---

```
plot_error_correction_panel
    Plot error correction panel
```

---

### Description

Creates a two-panel plot showing (1) error correction terms over time and (2) their estimated half-lives, providing visual assessment of convergence speed.

### Usage

```
plot_error_correction_panel(
  results_robust,
  use_contemporaneous_X = TRUE,
  main_left = expression(u[t] == Y[t] - hat(mu)[Y](X[t])),
  main_right = "Half-life de u[t]"
)
```

### Arguments

`results_robust` List. Results from main analysis pipeline.

`use_contemporaneous_X` Logical. Use contemporaneous X (time t) or lagged X (time t-1) for computing equilibrium path? Default is FALSE (lagged).

`main_left` Character string. Title for left panel. Default is "Error Correction Terms u\_t".

`main_right` Character string. Title for right panel. Default is "Half-Lives of Error Correction".

### Value

Invisibly returns a list with components:

`U` Matrix of error correction terms (T x Ky).

`half_lives` Vector of estimated half-lives for each Y factor.

---

```
read_cpi
    Read CPI data from Excel file
```

---

### Description

Reads Consumer Price Index data from an Excel file with robust error handling and data validation.

### Usage

```
read_cpi(path_cpi)
```

**Arguments**

path\_cpi            Path to the CPI data file

**Value**

Data frame with CPI data. Returns NULL if file not found or read operation fails.

---

```
rescue_short_run_channel
      Rescue short-run channel test
```

---

**Description**

Evaluates the short-run causal relationship from X to Y factors using Granger causality tests and out-of-sample forecast comparison.

**Usage**

```
rescue_short_run_channel(
  results_robust,
  lag = 1,
  B = 1000,
  seed = NULL,
  ridge = 0.001,
  oos_start = 0.6,
  verbose = TRUE
)
```

**Arguments**

results\_robust    Output from run\_complete\_factor\_analysis\_robust()  
lag                Number of lags for the model (default: 1)  
B                  Number of bootstrap iterations (default: 1000)  
seed               Random seed for reproducibility (default: NULL)  
ridge              Ridge regularization parameter (default: 0.001)  
oos\_start         Proportion of data to use for training (default: 0.6)  
verbose           Logical; print progress and diagnostic information. Default TRUE.

**Details**

testing. Default is 0.2.

**Value**

List with components:

granger\_p P-values from Granger causality tests.

OOS List with out-of-sample RMSE comparison.

p\_values Bootstrap p-values for RMSE differences.

---

rotation_null_test	<i>Rotation null hypothesis test for factor coupling</i>
--------------------	--

---

**Description**

Tests whether the observed correlation structure between X and Y factor spaces is significantly stronger than would be expected under random orthogonal rotations.

**Usage**

```
rotation_null_test(
  scores_X,
  scores_Y,
  lag = 1,
  B = 1000,
  seed = 123,
  compute = c("procrustes", "cca", "principal", "dynbeta"),
  progress = TRUE,
  rotate = "Y"
)
```

**Arguments**

scores_X	Factor scores from first dataset
scores_Y	Factor scores from second dataset
lag	Number of lags for the model (default: 1)
B	Number of bootstrap iterations (default: 1000)
seed	Random seed for reproducibility (default: 123)
compute	Vector of methods to compute: 'procrustes', 'cca', 'principal', 'dynbeta'
progress	Logical, show progress bar (default: TRUE)
rotate	Which dataset to rotate: 'X' or 'Y' (default: 'Y')

**Details**

(contemporaneous). "spearman", or "kendall".

**Value**

List with components:

observed Observed correlation statistics.

null\_distribution Matrix of statistics under null rotations.

p\_values One-sided p-values for each statistic.

significant Logical indicating significance at alpha = 0.05.

---

row_norm1	<i>Normalize matrix rows to sum to one</i>
-----------	--

---

**Description**

Scales each row of a matrix so that its elements sum to 1. Handles zero-sum rows by leaving them unchanged (avoiding division by zero).

**Usage**

```
row_norm1(M)
```

**Arguments**

M Numeric matrix to be row-normalized.

**Value**

Matrix of the same dimensions as M with each row summing to 1 (except rows that originally summed to zero).

**Examples**

```
M <- matrix(c(1, 2, 3, 4, 5, 6), nrow = 2)
M_norm <- row_norm1(M)
rowSums(M_norm) # Should be c(1, 1)
```

---

`run_complete_factor_analysis_robust`*Complete factor-OU convergence analysis pipeline*

---

### Description

Executes the end-to-end analysis workflow: data preparation, PLS-based factor extraction, DFM estimation, Factor-OU estimation, convergence tests, and robustness checks. This is the main user-facing function.

### Usage

```
run_complete_factor_analysis_robust(  
  X_matrix,  
  Y_matrix,  
  TMG = NULL,  
  COM_matrix = NULL,  
  SPVR_matrix = NULL,  
  CA = NULL,  
  sector_names = NULL,  
  max_comp = 3,  
  dfm_lags = 1,  
  ou_chains = 10,  
  ou_iter = 10000,  
  skip_ou = FALSE,  
  run_convergence_tests = TRUE,  
  path_cpi = NULL,  
  path_weights = NULL,  
  verbose = TRUE  
)
```

### Arguments

<code>X_matrix</code>	Matrix of first set of variables
<code>Y_matrix</code>	Matrix of second set of variables
<code>TMG</code>	Optional TMG matrix (default: NULL)
<code>COM_matrix</code>	Optional COM matrix (default: NULL)
<code>SPVR_matrix</code>	Optional SPVR matrix (default: NULL)
<code>CA</code>	Optional CA parameter (default: NULL)
<code>sector_names</code>	Vector of sector names (default: NULL)
<code>max_comp</code>	Maximum number of components (default: 3)
<code>dfm_lags</code>	Number of lags for DFM (default: 1)
<code>ou_chains</code>	Number of MCMC chains for OU estimation (default: 10)
<code>ou_iter</code>	Number of MCMC iterations (default: 10000)

skip_ou	Logical, skip OU estimation (default: FALSE)
run_convergence_tests	Logical, run convergence tests (default: TRUE)
path_cpi	Path to CPI data (default: NULL)
path_weights	Path to weights data (default: NULL)
verbose	Logical; print progress and diagnostic information. Default TRUE.

### Details

uses column names of `X_matrix`. and reweighting tests). Can be NULL.

This function orchestrates the complete analysis:

1. Data validation and diagnostics
2. Bayesian CPI disaggregation (if applicable)
3. PLS-based factor extraction with optimal component selection
4. Dynamic Factor Model estimation via VAR
5. Factor Ornstein-Uhlenbeck mean-reversion model
6. Formal convergence tests (stationarity, cointegration, speed)
7. Robustness tests (permutation, reweighting, jackknife)

### Value

List with components:

`factors` List containing PLS-extracted factors (`scores_X`, `scores_Y`) and related objects (`pls_X`, `pls_Y`, `ncomp_X`, `ncomp_Y`).

`dfm` List with DFM estimation results including VAR fit, lag order, diagnostics, and optional impulse responses.

`factor_ou` List with Factor-OU model estimates: `beta`, `lambda`, `sigma`, `half_life`, method used, and optional Stan fit object.

`convergence_tests` List with formal convergence test results (if `run_convergence_tests = TRUE`).

`robustness_tests` List with robustness test results (if `run_robustness_tests = TRUE`).

`diagnostics` List with data diagnostics (multicollinearity, stationarity, structural breaks).

`bayesian_cpi` List with Bayesian disaggregation results (if CPI paths provided).

`metadata` List with analysis metadata (timestamp, versions, parameters).

### See Also

[estimate\\_DFM](#), [estimate\\_factor\\_OU](#), [run\\_convergence\\_robustness\\_tests](#), [visualize\\_factor\\_dynamics](#)

## Examples

```
# Basic usage with simulated data
set.seed(123)
X <- matrix(rnorm(100 * 10), 100, 10)
Y <- X + matrix(rnorm(100 * 10, 0, 0.5), 100, 10)

results <- run_complete_factor_analysis_robust(
  X_matrix = X,
  Y_matrix = Y,
  max_comp = 3,
  dfm_lags = 1,
  ou_chains = 4,
  ou_iter = 2000,
  verbose = FALSE
)

# View convergence summary
summary(results$convergence_tests)

# Visualize results
visualize_factor_dynamics(
  dfm_result = results$dfm,
  ou_result = results$factor_ou,
  factors_data = results$factors
)
```

---

run\_convergence\_robustness\_tests

*Run comprehensive robustness test suite*

---

## Description

Executes all available robustness tests (permutation, reweighting, jackknife) and synthesizes results into an integrated assessment.

## Usage

```
run_convergence_robustness_tests(
  results_robust,
  X_matrix,
  Y_matrix,
  path_cpi = NULL,
  path_weights = NULL,
  sector_names = NULL,
  run_permutation = TRUE,
  run_reweighting = FALSE,
```

```

run_jackknife = TRUE,
run_leadlag = FALSE,
run_common_factor = FALSE,
sensitivity_analysis = TRUE,
verbose = TRUE
)

```

### Arguments

**results\_robust** Output from run\_complete\_factor\_analysis\_robust()  
**X\_matrix** Matrix of first set of variables  
**Y\_matrix** Matrix of second set of variables  
**path\_cpi** Path to CPI data (default: NULL)  
**path\_weights** Path to weights data (default: NULL)  
**sector\_names** Vector of sector names (default: NULL)  
**run\_permutation** Logical, run permutation test (default: TRUE)  
**run\_reweighting** Logical, run reweighting test (default: FALSE)  
**run\_jackknife** Logical, run jackknife test (default: TRUE)  
**run\_leadlag** Logical, run lead-lag test (default: FALSE)  
**run\_common\_factor** Logical, run common factor test (default: FALSE)  
**sensitivity\_analysis** Logical, run sensitivity analysis (default: TRUE)  
**verbose** Logical; print progress and diagnostic information. Default TRUE.

### Value

List with components:

**permutation** Results from permutation test.  
**reweighting** Results from reweighting test (if applicable).  
**jackknife** Results from jackknife test (if requested).  
**summary** Data frame summarizing all tests.  
**overall\_robust** Logical indicating if convergence is robust across all tests.

---

```
run_rotation_null_on_results
```

*Run rotation null test on complete analysis results*

---

### Description

Wrapper function that extracts factors from a complete analysis object and runs the rotation null test.

### Usage

```
run_rotation_null_on_results(
  results_robust,
  lag = 1,
  B = 1000,
  seed = 42,
  rotate = "Y",
  compute = c("procrustes", "cca", "principal", "dynbeta")
)
```

### Arguments

results_robust	Output from run_complete_factor_analysis_robust()
lag	Number of lags for the model (default: 1)
B	Number of bootstrap iterations (default: 1000)
seed	Random seed for reproducibility (default: 42)
rotate	Which dataset to rotate: 'X' or 'Y' (default: 'Y')
compute	Vector of methods to compute: 'procrustes', 'cca', 'principal', 'dynbeta'

### Value

List. Output from rotation\_null\_test.

---

```
select_optimal_components_safe
```

*Select optimal number of PLS components with cross-validation*

---

### Description

Determines the optimal number of Partial Least Squares components using leave-one-out or k-fold cross-validation, with robust error handling.

### Usage

```
select_optimal_components_safe(X, Y, max_comp = 15, verbose = TRUE)
```

**Arguments**

X	First data matrix
Y	Second data matrix
max_comp	Maximum number of components to test (default: 15)
verbose	Logical; print progress and diagnostic information. Default TRUE.

**Details**

default) or "CV" (k-fold cross-validation). validation = "LOO".

The function fits PLS models with 1 to max\_comp components, evaluates each via cross-validation, and selects the number that minimizes prediction error while avoiding overfitting.

**Value**

List with components:

ncomp\_optimal Optimal number of components selected.

RMSEP Root Mean Squared Error of Prediction for each component.

R2 R-squared values for each component.

validation\_method Validation method used.

pls\_fit Fitted PLS model object with optimal components.

---

test\_cointegration\_control

*Classical cointegration control (Johansen trace or eigen)*

---

**Description**

Runs Johansen's cointegration test on the first  $\min(2, \text{ncol}(X), \text{ncol}(Y))$  factors from scores\_X and scores\_Y, and counts the number of cointegrating relations at the 5% level.

**Usage**

```
test_cointegration_control(
  factors_data,
  max_lag = 4,
  type = "trace",
  ecdet = "const",
  verbose = TRUE
)
```

**Arguments**

factors_data	A list with matrices scores_X and scores_Y (T x Kx) and (T x Ky), respectively.
max_lag	Integer; maximum lag K passed to <code>urca::ca.jo()</code> .
type	Character; Johansen test type, one of "trace" or "eigen". Defaults to "trace".
ecdet	Character; deterministic terms, e.g. "const", "trend", or "none". Defaults to "const".
verbose	Logical; print progress and a summary of the test. Default TRUE.

**Details**

This function requires the optional package **urca** (declared in Suggests). It does not attempt to install packages at runtime; if **urca** is unavailable, an informative error is thrown.

The 5% critical values are taken from the "5pct" column of the `cval` slot returned by `urca::ca.jo()`.

**Value**

A list with:

- `test`: the `urca::ca.jo` fitted object,
- `n_coint`: integer number of cointegrating relations at 5%,
- `vectors`: matrix of cointegrating vectors (or NULL if none).

**References**

Johansen, S. (1991). Estimation and Hypothesis Testing of Cointegration Vectors in Gaussian Vector Autoregressive Models. *Econometrica*, 59(6), 1551-1580.

Johansen, S. (1995). *Likelihood-Based Inference in Cointegrated Vector Autoregressive Models*. Oxford University Press.

**See Also**

[ca.jo](#)

**Examples**

```
if (requireNamespace("urca", quietly = TRUE)) {
  set.seed(1)
  T <- 120
  X <- cbind(cumsum(rnorm(T)), cumsum(rnorm(T)))
  Y <- cbind(cumsum(rnorm(T)), cumsum(rnorm(T)))
  fd <- list(scores_X = X, scores_Y = Y)
  out <- test_cointegration_control(fd, max_lag = 2, verbose = FALSE)
  str(out)
}
```

---

 test\_jackknife\_sectors

*Jackknife robustness test by sector*


---

### Description

Assesses the influence of individual sectors by systematically dropping each sector and re-estimating convergence parameters. Identifies influential sectors and checks stability.

### Usage

```
test_jackknife_sectors(
  X_matrix,
  Y_matrix,
  sector_names = NULL,
  k_exclude = 3,
  max_comp = 3,
  verbose = TRUE
)
```

### Arguments

X_matrix	Matrix of first set of variables
Y_matrix	Matrix of second set of variables
sector_names	Vector of sector names (default: NULL)
k_exclude	Number of sectors to exclude in each iteration (default: 3)
max_comp	Maximum number of components (default: 3)
verbose	Logical; print progress and diagnostic information. Default TRUE.

### Details

column names.

### Value

List with components:

jackknife\_estimates Matrix of lambda estimates (iterations x factors).

original\_estimate Original lambda from full sample.

bias Estimated jackknife bias.

se Jackknife standard errors.

influential\_sectors Character vector of highly influential sectors.

dfbetas Matrix of influence measures (DFBETAS).

---

test\_permutation\_robustness

*Permutation-based robustness test*

---

### Description

Tests the robustness of factor-OU convergence findings by randomly permuting the Y factor space and re-estimating the model. Generates empirical null distribution for convergence statistics.

### Usage

```
test_permutation_robustness(
  factors_data,
  data_prep,
  n_perms = 100,
  seed = 123,
  use_stan = TRUE,
  chains = 4,
  iter = 2000,
  verbose = TRUE
)
```

### Arguments

factors_data	Data frame with factor information
data_prep	Prepared data object
n_perms	Number of permutations (default: 100)
seed	Random seed for reproducibility (default: 123)
use_stan	Logical, use Stan for estimation (default: TRUE)
chains	Number of MCMC chains (default: 4)
iter	Number of MCMC iterations (default: 2000)
verbose	Logical; print progress and diagnostic information. Default TRUE.

### Details

(too slow for many iterations).

### Value

List with components:

observed_lambda	Original mean-reversion speeds.
null_distribution	Matrix of permutation-based lambda values.
p_values	One-sided p-values for each factor.
significant	Logical vector indicating significance at alpha = 0.05.
effect_size	Standardized effect sizes (z-scores).

---

```
test_reweighting_robustness
```

*Reweighting-based robustness test*

---

## Description

Tests sensitivity of convergence results to alternative sectoral weighting schemes by re-running Bayesian disaggregation and full pipeline with perturbed weights.

## Usage

```
test_reweighting_robustness(  
  path_cpi,  
  path_weights,  
  X_matrix,  
  max_comp = 3,  
  verbose = TRUE  
)
```

## Arguments

path_cpi	Path to CPI data file
path_weights	Path to weights data file
X_matrix	Matrix of variables
max_comp	Maximum number of components (default: 3)
verbose	Logical; print progress and diagnostic information. Default TRUE.

## Value

List with components:

lambda\_distribution Matrix of lambda estimates across schemes.

original\_lambda Baseline lambda from original weights.

robust\_factors Logical vector indicating robust convergence.

sensitivity\_metrics Summary statistics of sensitivity.

---

to_num_commas	<i>Convert European number format to numeric</i>
---------------	--

---

**Description**

Converts character strings in European number format (using comma as decimal separator and period as thousands separator) to numeric values. Already numeric inputs are returned unchanged.

**Usage**

```
to_num_commas(x)
```

**Arguments**

x                    Numeric value or character string in European format (e.g., "1.234,56").

**Value**

Numeric value. Returns NA if conversion fails.

**Examples**

```
to_num_commas("1.234,56")
to_num_commas("1234.56")
to_num_commas(1234.56)
```

---

visualize_factor_dynamics	<i>Visualize factor dynamics comprehensively</i>
---------------------------	--

---

**Description**

Creates a multi-panel visualization summarizing all aspects of the factor model: scores over time, loadings, correlations, OU dynamics, and convergence patterns.

**Usage**

```
visualize_factor_dynamics(  
  dfm_result,  
  ou_result,  
  factors_data,  
  save_plot = FALSE,  
  plot_file = NULL,  
  use_device = "default",  
  verbose = TRUE  
)
```

**Arguments**

<code>dfm_result</code>	Result object from DFM analysis
<code>ou_result</code>	Result object from OU estimation
<code>factors_data</code>	Data frame with factor information
<code>save_plot</code>	Logical, save plot to file (default: FALSE)
<code>plot_file</code>	File name for saved plot (default: 'factor_dynamics.pdf')
<code>use_device</code>	Graphics device to use: 'default', 'pdf', 'png' (default: 'default')
<code>verbose</code>	Logical; print progress and diagnostic information. Default TRUE.

**Details**

Default is NULL (display only).

**Value**

Invisibly returns NULL. Called for side effect of creating plots.

---

visualize\_factor\_dynamics\_simple  
*Simple factor dynamics visualization*

---

**Description**

Creates a simplified multi-panel plot of factor scores over time with minimal dependencies. Useful for quick diagnostics.

**Usage**

```
visualize_factor_dynamics_simple(  
  factors_data,  
  output_file = NULL,  
  verbose = TRUE  
)
```

**Arguments**

<code>factors_data</code>	List containing factor scores.
<code>output_file</code>	Character string. Optional file path for saving. Default is NULL.
<code>verbose</code>	Logical; print progress and diagnostic information. Default TRUE.

**Value**

Invisibly returns NULL.

**Examples**

```
data <- list(scores_X = matrix(rnorm(100), ncol = 2),  
              scores_Y = matrix(rnorm(100), ncol = 2))  
tmp_file <- file.path(tempdir(), "test_plot.pdf")  
visualize_factor_dynamics_simple(data, output_file = tmp_file)
```

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