

# 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 DFM 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](https://doi.org/10.1198/073500102317351921)> and the Generalized Dynamic Factor Model of Forni, Hallin, Lippi and Reichlin (2000) <[doi:10.1162/003465300559037](https://doi.org/10.1162/003465300559037)>; (ii) cointegration analysis à la Johansen (1988) <[doi:10.1016/0165-1889\(88\)90041-3](https://doi.org/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](https://doi.org/10.1103/PhysRev.36.823)> and Vasicek (1977) <[doi:10.1016/0304-405X\(77\)90016-2](https://doi.org/10.1016/0304-405X(77)90016-2)>; (iv) robust inference via 'sandwich' HC/HAC estimators (Zeileis (2004) <[doi:10.18637/jss.v011.i10](https://doi.org/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](https://doi.org/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>
----------------	--

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### 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

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

- 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).

### 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**

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

*Diagnose and prepare data matrices*

**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**

<code>X_matrix</code>	Numeric matrix or data frame of X variables (e.g., Marxist prices).
<code>Y_matrix</code>	Numeric matrix or data frame of Y variables (e.g., market prices/CPI).
<code>verbose</code>	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

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

<code>estimate_factor_OU</code>	<i>Estimate Factor Ornstein-Uhlenbeck model (Stan if available)</i>
---------------------------------	---

## 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

<code>factors_data</code>	List with <code>scores_X</code> , <code>scores_Y</code> .
<code>data_prep</code>	Optional preprocessed data (reserved).
<code>chains</code> , <code>iter</code> , <code>seed</code>	Stan sampling controls.
<code>adapt1</code> , <code>adapt2</code> , <code>mtd1</code> , <code>mtd2</code>	Advanced Stan controls.
<code>verbose</code>	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**      *Extract X innovations from VAR model*

**Description**

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

**Usage**

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

**Arguments**

- |                      |  |
|----------------------|--|
| <code>results</code> | List. Output from main analysis.   |
| <code>p</code>       | Integer. VAR lag order. If <code>NULL</code> , 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`      *Rotation null hypothesis test for factor coupling*

**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**

<code>scores_X</code>	Factor scores from first dataset
<code>scores_Y</code>	Factor scores from second dataset
<code>lag</code>	Number of lags for the model (default: 1)
<code>B</code>	Number of bootstrap iterations (default: 1000)
<code>seed</code>	Random seed for reproducibility (default: 123)
<code>compute</code>	Vector of methods to compute: 'procrustes', 'cca', 'principal', 'dynbeta'
<code>progress</code>	Logical, show progress bar (default: TRUE)
<code>rotate</code>	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***Normalize matrix rows to sum to one*

---

**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

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

<code>factors</code>	List containing PLS-extracted factors ( <code>scores_X</code> , <code>scores_Y</code> ) and related objects ( <code>pls_X</code> , <code>pls_Y</code> , <code>ncomp_X</code> , <code>ncomp_Y</code> ).
<code>dfm</code>	List with DFM estimation results including VAR fit, lag order, diagnostics, and optional impulse responses.
<code>factor_ou</code>	List with Factor-OU model estimates: <code>beta</code> , <code>lambda</code> , <code>sigma</code> , <code>half_life</code> , method used, and optional Stan fit object.
<code>convergence_tests</code>	List with formal convergence test results (if <code>run_convergence_tests</code> = TRUE).
<code>robustness_tests</code>	List with robustness test results (if <code>run_robustness_tests</code> = TRUE).
<code>diagnostics</code>	List with data diagnostics (multicollinearity, stationarity, structural breaks).
<code>bayesian_cpi</code>	List with Bayesian disaggregation results (if CPI paths provided).
<code>metadata</code>	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

<code>results_robust</code>	Output from <code>run_complete_factor_analysis_robust()</code>
<code>X_matrix</code>	Matrix of first set of variables
<code>Y_matrix</code>	Matrix of second set of variables
<code>path_cpi</code>	Path to CPI data (default: NULL)
<code>path_weights</code>	Path to weights data (default: NULL)
<code>sector_names</code>	Vector of sector names (default: NULL)
<code>run_permutation</code>	Logical, run permutation test (default: TRUE)
<code>run_reweighting</code>	Logical, run reweighting test (default: FALSE)
<code>run_jackknife</code>	Logical, run jackknife test (default: TRUE)
<code>run_leadlag</code>	Logical, run lead-lag test (default: FALSE)
<code>run_common_factor</code>	Logical, run common factor test (default: FALSE)
<code>sensitivity_analysis</code>	Logical, run sensitivity analysis (default: TRUE)
<code>verbose</code>	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, ncol(X), 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

<code>factors_data</code>	A list with matrices <code>scores_X</code> and <code>scores_Y</code> ( $T \times K_x$ ) and ( $T \times K_y$ ), respectively.
<code>max_lag</code>	Integer; maximum lag $K$ passed to <code>urca::ca.jo()</code> .
<code>type</code>	Character; Johansen test type, one of "trace" or "eigen". Defaults to "trace".
<code>ecdet</code>	Character; deterministic terms, e.g. "const", "trend", or "none". Defaults to "const".
<code>verbose</code>	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**

<code>factors_data</code>	Data frame with factor information
<code>data_prep</code>	Prepared data object
<code>n_perms</code>	Number of permutations (default: 100)
<code>seed</code>	Random seed for reproducibility (default: 123)
<code>use_stan</code>	Logical, use Stan for estimation (default: TRUE)
<code>chains</code>	Number of MCMC chains (default: 4)
<code>iter</code>	Number of MCMC iterations (default: 2000)
<code>verbose</code>	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.
```

---

<code>to_num_commas</code>	<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

<code>x</code>	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)
```

---

<code>visualize_factor_dynamics</code>	<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**

dfm_result	Result object from DFM analysis
ou_result	Result object from OU estimation
factors_data	Data frame with factor information
save_plot	Logical, save plot to file (default: FALSE)
plot_file	File name for saved plot (default: 'factor_dynamics.pdf')
use_device	Graphics device to use: 'default', 'pdf', 'png' (default: 'default')
verbose	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**

factors_data	List containing factor scores.
output_file	Character string. Optional file path for saving. Default is NULL.
verbose	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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