

Package ‘fmi’

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Title Hierarchical Permutation Tests for Functional Measurement Invariance

Description Provides a suite of functions to test for Functional Measurement Invariance (FMI) between two groups. Implements hierarchical permutation tests for configural, metric, and scalar invariance, adapting concepts from Multi-Group Confirmatory Factor Analysis (MGCF) to functional data.

Methods are based on concepts from:

Meredith, W. (1993) <[doi:10.1007/BF02294825](https://doi.org/10.1007/BF02294825)>,⁵

Yao, F., Müller, H. G., & Wang, J. L. (2005) <[doi:10.1198/016214504000001745](https://doi.org/10.1198/016214504000001745)>, and Lee, K. Y., & Li, L. (2022) <[doi:10.1111/rssb.12471](https://doi.org/10.1111/rssb.12471)>.

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Imports dplyr, ggplot2, knitr, magrittr, methods, purrr, refund, stats, tibble, tidyverse, utils

Suggests fda, rmarkdown

VignetteBuilder knitr

NeedsCompilation no

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build_scree_plot	<i>Build Scree Plot for Configural Invariance</i>
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Description

Creates a ggplot2 scree plot comparing the PVE by FPCs for two groups.

Usage

```
build_scree_plot(f pca_A, f pca_B, groups)
```

Arguments

f pca_A	An f pca.sc object for Group A.
f pca_B	An f pca.sc object for Group B.
groups	A character vector of the two group names.

Value

A ggplot object.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
g_fac <- factor(sim$group_vec)
idx_A <- which(g_fac == levels(g_fac)[1])
# --- FIX: Added Y= and argvals= ---
f pca_A <- refund::f pca.sc(Y = sim$Y_mat[idx_A, ], argvals = sim$argvals, npc = 3)
f pca_B <- refund::f pca.sc(Y = sim$Y_mat[-idx_A, ], argvals = sim$argvals, npc = 3)
build_scree_plot(f pca_A, f pca_B, levels(g_fac))
```

 compare_latent_means *Compare Latent Means*

Description

Performs Welch's t-tests on FPC scores between groups. This function should only be run after scalar invariance is established.

Usage

```
compare_latent_means(fmi_results, group_vec)
```

Arguments

- fmi_results The list object returned by `run_fmi()`.
- group_vec The original group vector used in `run_fmi()`.

Value

A tibble summarizing the t-test results for each FPC.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
fmi_results <- run_fmi(
  Y_mat = sim$Y_mat,
  group_vec = sim$group_vec,
  argvals = sim$argvals,
  n_perms = 9,
  progress = FALSE
)
# This will run only if scalar invariance passed
if (!is.null(fmi_results$scalar) && fmi_results$scalar$passed) {
  mean_results <- compare_latent_means(fmi_results, sim$group_vec)
  print(mean_results)
}
```

 determine_npc *Determine Number of FPCs Automatically*

Description

Determines the number of functional principal components (FPCs) required to meet a target Proportion of Variance Explained (PVE).

Usage

```
determine_npc(Y_mat, argvals, target_pve = 0.95, max_npc = 6L)
```

Arguments

<code>Y_mat</code>	A numeric matrix (N x M) where N is subjects, M is time points.
<code>argvals</code>	A numeric vector of length M listing the observation points.
<code>target_pve</code>	The target cumulative PVE to reach (default: 0.95).
<code>max_npc</code>	The maximum number of components to retain (default: 6L).

Value

A list containing:

<code>npc</code>	The selected number of components.
<code>pooled_fPCA</code>	The <code>fPCA.sc</code> object from the <code>refund</code> package.
<code>scree</code>	A tibble with PVE and cumulative PVE for each component.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
npc_info <- determine_npc(sim$Y_mat, sim$argvals)
print(npc_info$npc)
```

`run_dti_example`

DTI Example Data Wrapper

Description

A helper function to load and pre-process the DTI dataset from the `'refund'` package for use in examples.

Usage

```
run_dti_example(n_perms = 499L)
```

Arguments

<code>n_perms</code>	The number of permutations (default: 499L). Passed to <code>run_fmi</code> .
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Value

Invisibly returns the FMI results list.

Examples

```
# n_perms=9 is for a quick check. Use 499+ for real analysis.
if (requireNamespace("refund", quietly = TRUE)) {
  dti_results <- run_dti_example(n_perms = 9)
}
```

run_fmi

Run Hierarchical Functional Measurement Invariance (FMI) Tests

Description

A wrapper function that performs a sequence of permutation tests for configural, metric, scalar, and (optionally) strict invariance.

Usage

```
run_fmi(
  Y_mat,
  group_vec,
  argvals,
  alpha = 0.05,
  npc = NULL,
  target_pve = 0.95,
  max_npc = 6L,
  n_perms = 499L,
  strict_test = FALSE,
  progress = interactive()
)
```

Arguments

Y_mat	A numeric matrix (N x M) where N is subjects, M is time points.
group_vec	A vector of length N specifying group membership (2 groups).
argvals	A numeric vector of length M listing the observation points.
alpha	Significance level (default: 0.05).
npc	Optional. The number of FPCs. If NULL (default), it is determined automatically by determine_npc.
target_pve	The target PVE if npc is NULL (default: 0.95).
max_npc	The max FPCs if npc is NULL (default: 6L).
n_perms	The number of permutations (default: 499L).
strict_test	Boolean. Whether to perform the strict invariance test (default: FALSE).
progress	Boolean. Show progress bars? (default: interactive()).

Value

A list containing test results for each invariance level, settings, and the pooled FPCA object.

Examples

```
# Use small N and n_perms for a quick example
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
fmi_results <- run_fmi(
  Y_mat = sim$Y_mat,
  group_vec = sim$group_vec,
  argvals = sim$argvals,
  n_perms = 9, # Use 499+ for actual research
  progress = FALSE
)
print(fmi_results)
```

run_growth_example Berkeley Growth Example Data Wrapper

Description

A helper function to load and pre-process the Growth dataset from the 'fda' package for use in examples.

Usage

```
run_growth_example(n_perms = 499L)
```

Arguments

n_perms The number of permutations (default: 499L). Passed to `run_fmi`.

Value

Invisibly returns the FMI results list.

Examples

```
# n_perms=9 is for a quick check. Use 499+ for real analysis.
if (requireNamespace("fda", quietly = TRUE)) {
  growth_results <- run_growth_example(n_perms = 9)
}
```

simulate_fmi_data	<i>Simulate FMI Data</i>
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Description

Generates functional data for two groups with optional differences in latent means or scalar intercepts.

Usage

```
simulate_fmi_data(  
  N_A = 50,  
  N_B = 50,  
  T_points = 51,  
  mean_shift = 0,  
  scalar_shift = 0,  
  eigenvalues = c(4, 1, 0.5, 0.1),  
  noise_sd = 0.5,  
  seed = NULL  
)
```

Arguments

N_A	Sample size for Group A (default: 50).
N_B	Sample size for Group B (default: 50).
T_points	Number of time points (default: 51).
mean_shift	A shift added to FPC 1 scores for Group B (default: 0).
scalar_shift	A shift added to the mean function for Group B (default: 0).
eigenvalues	A vector of eigenvalues for FPCs (default: c(4, 1, 0.5, 0.1)).
noise_sd	Standard deviation of measurement error (default: 0.5).
seed	Optional. A random seed for reproducibility.

Value

A list with `Y_mat`, `group_vec`, and `argvals`.

Examples

```
sim_data <- simulate_fmi_data(N_A = 10, N_B = 10, T_points = 20, seed = 123)  
str(sim_data)
```

test_configural *Test for Configural Invariance*

Description

Permutation test to check if the cumulative PVE is equivalent across groups.

Usage

```
test_configural(Y_mat, group_vec, argvals, npc, n_perms, alpha, progress)
```

Arguments

Y_mat	A numeric matrix (N x M) where N is subjects, M is time points.
group_vec	A vector of length N specifying group membership (2 groups).
argvals	A numeric vector of length M listing the observation points.
npc	Optional. The number of FPCs. If NULL (default), it is determined automatically by determine_npc.
n_perms	The number of permutations (default: 499L).
alpha	Significance level (default: 0.05).
progress	Boolean. Show progress bars? (default: interactive()).

Value

A list with passed (boolean), p_value, statistic, and plot.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
res <- test_configural(sim$Y_mat, sim$group_vec, sim$argvals,
                      npc = 3, n_perms = 9, alpha = 0.05, progress = FALSE)
print(res$p_value)
```

test_metric *Test for Metric Invariance*

Description

Permutation test to check if the eigenfunctions are equivalent across groups.

Usage

```
test_metric(Y_mat, group_vec, argvals, npc, n_perms, alpha, progress)
```

Arguments

<code>Y_mat</code>	A numeric matrix (N x M) where N is subjects, M is time points.
<code>group_vec</code>	A vector of length N specifying group membership (2 groups).
<code>argvals</code>	A numeric vector of length M listing the observation points.
<code>npc</code>	Optional. The number of FPCs. If NULL (default), it is determined automatically by <code>determine_npc</code> .
<code>n_perms</code>	The number of permutations (default: 499L).
<code>alpha</code>	Significance level (default: 0.05).
<code>progress</code>	Boolean. Show progress bars? (default: <code>interactive()</code>).

Value

A list with passed (boolean), `p_value`, and `statistic`.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
res <- test_metric(sim$Y_mat, sim$group_vec, sim$argvals,
                    npc = 3, n_perms = 9, alpha = 0.05, progress = FALSE)
print(res$p_value)
```

test_scalar*Test for Scalar Invariance***Description**

Permutation test to check if the mean functions (intercepts) are equivalent across groups, after accounting for latent factors.

Usage

```
test_scalar(Y_mat, group_vec, argvals, pooled_fpca, n_perms, alpha, progress)
```

Arguments

<code>Y_mat</code>	Numeric matrix (N x M).
<code>group_vec</code>	A vector of length N specifying group membership.
<code>argvals</code>	Numeric vector of length M.
<code>pooled_fpca</code>	The <code>fpca.sc</code> object from the <code>determine_npc</code> function.
<code>n_perms</code>	Integer, number of permutations.
<code>alpha</code>	Significance level (default: 0.05).
<code>progress</code>	Boolean, show progress bar? (default: <code>interactive()</code>).

Value

A list with passed (boolean), p_value, and statistic.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
npc_info <- determine_npc(sim$Y_mat, sim$argvals, max_npc = 3)
res <- test_scalar(sim$Y_mat, sim$group_vec, sim$argvals,
                    pooled_fpca = npc_info$pooled_fpca,
                    n_perms = 9, alpha = 0.05, progress = FALSE)
print(res$p_value)
```

test_strict*Test for Strict Invariance***Description**

Permutation test to check if the residual error variances are equivalent across groups.

Usage

```
test_strict(Y_mat, group_vec, argvals, npc, n_perms, alpha, progress)
```

Arguments

<i>Y_mat</i>	A numeric matrix (N x M) where N is subjects, M is time points.
<i>group_vec</i>	A vector of length N specifying group membership (2 groups).
<i>argvals</i>	A numeric vector of length M listing the observation points.
<i>npc</i>	Optional. The number of FPCs. If NULL (default), it is determined automatically by <i>determine_npc</i> .
<i>n_perms</i>	The number of permutations (default: 499L).
<i>alpha</i>	Significance level (default: 0.05).
<i>progress</i>	Boolean. Show progress bars? (default: <i>interactive()</i>).

Value

A list with passed (boolean), p_value, and statistic.

Examples

```
sim <- simulate_fmi_data(N_A = 20, N_B = 20, T_points = 30)
res <- test_strict(sim$Y_mat, sim$group_vec, sim$argvals,
                    npc = 3, n_perms = 9, alpha = 0.05, progress = FALSE)
print(res$p_value)
```

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