Package 'robustT2'

May 9, 2026

Title Robust Hotelling-Type T² Control Chart Based on the Dual STATIS Approach

Version 0.1.0

Description Implements a robust multivariate control-chart methodology for batch-based industrial processes with multiple correlated variables using the Dual STATIS (Structuration des Tableaux A Trois Indices de la Statistique) framework. A robust compromise covariance matrix is constructed from Phase I batches with the Minimum Covariance Determinant (MCD) estimator, and a Hotelling-type T² statistic is applied for anomaly detection in Phase II. The package includes functions to simulate clean and contaminated batches, to compute both robust and classical Hotelling T² control charts, to visualize results via robust biplots, and to launch an interactive 'shiny' dashboard. An internal dataset (pharma_data) is provided for reproducibility. See Lavit, Escoufier, Sabatier and Traissac (1994) <doi:10.1016/0167-9473(94)90134-1> for the original STATIS methodology, and Rousseeuw and Van Driessen (1999) <doi:10.1080/00401706.1999.10485670> for the MCD estimator.

License MIT + file LICENSE

Encoding UTF-8

LazyData true

RoxygenNote 7.3.2

Imports dplyr, MASS, ggplot2, rrcov, shiny, ggrepel, forcats, Matrix

Depends R (>= 3.5)

URL https://github.com/SergioDanielFG/robustT2

BugReports https://github.com/SergioDanielFG/robustT2/issues

Suggests spelling

Language en-US

NeedsCompilation no

Author Sergio Daniel Frutos Galarza [aut, cre] (ORCID: <https://orcid.org/0009-0007-2961-032X>), Omar Ruiz Barzola [aut] (ORCID: <https://orcid.org/0000-0001-8206-1744>), Purificación Galindo Villardón [aut] (ORCID: <https://orcid.org/0000-0001-6977-7545>)

Maintainer Sergio Daniel Frutos Galarza <sergio_dan88@hotmail.com>

Repository CRAN

Date/Publication 2025-09-10 08:00:02 UTC

Contents

`pharma_data` *Simulated Pharmaceutical Manufacturing Data*

Description

This dataset contains simulated pharmaceutical manufacturing data generated by `simulate_pharma_batches()` with `seed = 780` and `obs_per_batch = 30`.

Usage

```
data("pharma_data")
```

Format

A data frame with 450 rows and 7 variables:

Batch Batch identifier (factor)

Phase Phase indicator: "Phase 1" or "Phase 2" (factor)

Status Batch status: "Under Control" or "Out of Control" (factor)

Concentration Concentration of active ingredient (mg/mL)

Humidity Humidity percentage (% w/w)

Dissolution Dissolution percentage (% released)

Density Density (g/cm^3)

Details

Phase 1 includes 10 under-control batches with natural variability in mean and covariance, without contamination.

Phase 2 includes 2 additional under-control batches and 3 out-of-control batches. The out-of-control batches exhibit shifts in both mean and variability, along with moderate contamination in a portion of their observations.

Each batch contains 30 observations measured across four quantitative quality-control variables.

Source

Simulated using `simulate_pharma_batches` with `seed = 780` and `obs_per_batch = 30`.

`plot_classical_hotelling_t2_chart`
Plot Classical Hotelling T2 Control Chart

Description

Plots the classical Hotelling T2 statistics per batch with a uniform color line. Batches are evaluated against a control threshold obtained from the chi-squared distribution with degrees of freedom equal to the number of variables.

Usage

```
plot_classical_hotelling_t2_chart(
  t2_statistics,
  num_vars,
  title = "Classical Hotelling T2 Control Chart"
)
```

Arguments

`t2_statistics`	A data frame with columns `Batch` and `T2_Stat`.
`num_vars`	Integer. Number of variables used in the multivariate analysis (to compute the Chi² threshold).
`title`	Optional string. Plot title.

Value

A ggplot2 object representing the control chart.

Examples

```
# Simulate pharmaceutical manufacturing batches
sim_batches <- simulate_pharma_batches()

# Phase 1 analysis: use Phase 1 data
phase1_data <- subset(sim_batches, Phase == "Phase 1")

# Apply classical Hotelling T2 methodology
t2_result <- hotelling_t2_phase1(
  data = phase1_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)

# Plot classical Hotelling T2 control chart
plot_classical_hotelling_t2_chart(
  t2_statistics = t2_result$batch_statistics,
  num_vars = 4
)
```

---

plot_classical_hotelling_t2_phase2_chart

Plot Classical Hotelling T2 Control Chart - Phase 2

Description

Plots the classical Hotelling T² statistics per batch for Phase 2 data, using the reference mean and covariance matrix estimated from Phase 1. Batches are color-coded by control status ("Under Control" = blue, "Out of Control" = red).

Usage

```
plot_classical_hotelling_t2_phase2_chart(
  t2_statistics,
  num_vars,
  title = "Classical Hotelling T2 Control Chart (Phase 2)"
)
```

Arguments

`t2_statistics`	A data frame with columns `Batch`, `T2_Stat`, and `Status`.
`num_vars`	Integer. Number of variables used in the multivariate analysis (degrees of freedom for Chi²).
`title`	Optional string. Plot title.

Value

A ggplot2 object with the Phase 2 control chart.

Examples

```
# Simulate pharmaceutical manufacturing batches
sim_batches <- simulate_pharma_batches()

# Split by phase
phase1_data <- subset(sim_batches, Phase == "Phase 1")
phase2_data <- subset(sim_batches, Phase == "Phase 2")

# Fit Phase 1 classical estimators
t2_phase1 <- hotelling_t2_phase1(
  data = phase1_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)

# Evaluate Phase 2 batches
t2_phase2 <- hotelling_t2_phase2(
  new_data = phase2_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density"),
  center = t2_phase1$center,
  covariance = t2_phase1$covariance
)

# Combine with status for plotting
status_info <- phase2_data[!duplicated(phase2_data$Batch), "Status"]
t2_phase2_plot <- cbind(t2_phase2$batch_statistics, Status = status_info)

# Plot Phase 2 control chart
plot_classical_hotelling_t2_phase2_chart(
  t2_statistics = t2_phase2_plot,
  num_vars = 4
)
```

---

plot_statis_biplot_projection

HJ-Biplot Projection - Robust STATIS Dual (Phase 2)

Description

Projects new batches from Phase 2 into the HJ-Biplot space defined by the robust compromise matrix and eigen decomposition from Phase 1.

Usage

```
plot_statis_biplot_projection(phase1_result, phase2_result, dims = c(1, 2))
```

Arguments

`phase1_result`	Result from `robust_statis_phase1()`.

phase2_result	Result from robust_statis_phase2() (must include standardized_data, t2_stats_by_batch and threshold).
dims	Dimensions to plot (default: c(1, 2)).

Details

This implementation follows the HJ-Biplot formulation of Galindo-Villardón (1986). The compromise matrix C, being symmetric and positive semidefinite, is decomposed via an eigen decomposition (not a rectangular SVD). The square roots of eigenvalues are used to build the biplot scaling, consistent with robust STATIS Dual.

Value

A ggplot2 object with the projected HJ-Biplot for Phase 2 batches.

Examples

```
sim_batches <- simulate_pharma_batches()
phase1_data <- subset(sim_batches, Phase == "Phase 1" & Status == "Under Control")
phase2_data <- subset(sim_batches, Phase == "Phase 2")

phase1 <- robust_statis_phase1(
  data = phase1_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)

phase2 <- robust_statis_phase2(
  new_data = phase2_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density"),
  medians = phase1$global_medians,
  mads = phase1$global_mads,
  compromise_matrix = phase1$compromise_matrix,
  global_center = phase1$global_center
)

plot_statis_biplot_projection(phase1, phase2)
```

---

plot_statis_hj_biplot *HJ-Biplot of Robust STATIS Dual Compromise (Galindo-Villardón)*

Description

Generates an HJ-Biplot using the compromise matrix obtained from robust STATIS Dual. Individuals (batch centers) are projected as G = U D, and variables as H = V D, where D is the diagonal matrix of square roots of eigenvalues.

Usage

```
plot_statis_hj_biplot(
  phase1_result,
  dims = c(1, 2),
  color_by = c("none", "weight", "distance"),
  highlight_batches = NULL
)
```

Arguments

phase1_result	Result from robust_statis_phase1().
dims	Dimensions to plot (default: c(1, 2)).
color_by	One of "none", "weight", or "distance" for coloring batches.
highlight_batches	Optional vector of batch names to emphasize.

Value

ggplot2 object with HJ-Biplot.

Examples

```
sim_batches <- simulate_pharma_batches()
phase1 <- robust_statis_phase1(
  data = subset(sim_batches, Phase == "Phase 1" & Status == "Under Control"),
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)
plot_statis_hj_biplot(phase1)
```

---

plot_statis_phase1_chart

Plot Control Chart - Robust STATIS Dual (Phase 1)

Description

Plots the Hotelling T^2 statistic per batch using the robust center and compromise matrix estimated in `robust_statis_phase1()`. The control limit is based on a Chi-squared distribution with degrees of freedom equal to the number of variables.

Usage

```
plot_statis_phase1_chart(
  batch_statistics,
  num_vars,
  title = "Robust STATIS Dual Control Chart - Phase 1"
)
```

Arguments

batch_statistics	A data frame with columns Batch and T2_Stat, typically from phase1_result$batch_statistics.
num_vars	Integer. Number of variables used in the multivariate analysis (to compute the Chi² threshold).
title	Optional string. Plot title.

Value

A ggplot2 object.

Examples

```
sim_batches <- simulate_pharma_batches()

# Phase 1 analysis: select under control batches from Phase 1
phase1_result <- robust_statis_phase1(
  data = subset(sim_batches, Phase == "Phase 1" & Status == "Under Control"),
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)

# Plot the Phase 1 robust control chart
plot_statis_phase1_chart(
  batch_statistics = phase1_result$batch_statistics,
  num_vars = 4
)
```

---

plot_statis_phase2_chart

Plot STATIS Dual Robust Control Chart - Phase 2 Only

Description

Plots the robust Hotelling T² statistics for Phase 2 batches only, using the results from the robust STATIS Dual method.

Usage

```
plot_statis_phase2_chart(
  phase2_result,
  title = "Robust STATIS Dual Control Chart - Phase 2"
)
```

Arguments

phase2_result	A list returned by robust_statis_phase2(), including t2_stats_by_batch with Hotelling T² values and a control threshold.
title	Optional string. Plot title.

Value

A ggplot2 object representing the control chart for Phase 2 batches.

Examples

```
sim_batches <- simulate_pharma_batches()
phase1 <- robust_statis_phase1(
  data = subset(sim_batches, Phase == "Phase 1" & Status == "Under Control"),
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)
phase2 <- robust_statis_phase2(
  new_data = subset(sim_batches, Phase == "Phase 2"),
  variables = c("Concentration", "Humidity", "Dissolution", "Density"),
  medians = phase1$global_medians,
  mads = phase1$global_mads,
  compromise_matrix = phase1$compromise_matrix,
  global_center = phase1$global_center
)
plot_statis_phase2_chart(phase2_result = phase2)
```

---

robust_statis_phase1 *Robust STATIS Dual - Phase 1 (Under Control Batches)*

Description

Applies the Robust STATIS Dual methodology to Phase 1 data (under control batches), using robust batch-wise standardization (median and MAD). Covariance matrices are robustly estimated using the MCD method and used directly (without trace normalization) to construct the compromise matrix.

Usage

```
robust_statis_phase1(data, variables)
```

Arguments

`data`	A data frame containing the process data with batch information.
`variables`	Character vector with the names of the variables to be used in the analysis.

Value

A list containing:

compromise_matrix Robust compromise matrix (without trace normalization)

global_center Global robust center of the batches

batch_statistics Data frame with Batch, T2_Stat (Hotelling-type robust statistic), and Weight

batch_medians List of medians per batch and variable

batch_mads List of MADs per batch and variable

global_medians Global medians per variable (for use in Phase 2)

global_mads Global MADs per variable

robust_means List of robust centers of each batch (estimated by MCD)

standardized_data Data set standardized batch by batch

robust_covariances List of robust covariance matrices per batch

similarity_matrix Hilbert-Schmidt similarity matrix between batches

statis_weights Weights obtained from the first eigenvector of the similarity matrix

first_eigenvector First eigenvector of the similarity matrix (unnormalized)

Examples

```
# Simulate new pharmaceutical manufacturing batches
sim_batches <- simulate_pharma_batches()

# Select only Phase 1 under control batches
phase1_data <- subset(sim_batches, Phase == "Phase 1" & Status == "Under Control")

# Apply robust STATIS Dual methodology
result <- robust_statis_phase1(
  data = phase1_data,
  variables = c("Concentration", "Humidity", "Dissolution", "Density")
)

# View main outputs
result$compromise_matrix
result$batch_statistics
result$robust_covariances
result$similarity_matrix
result$statis_weights
result$robust_means
```

---

run_statis_dashboard *Launch STATIS Dual Robust Dashboard (Shiny)*

Description

Launches an interactive Shiny dashboard that includes:

- Phase 1 control chart (sum of robust Mahalanobis distances)
- Phase 2 control chart (for new batches)
- HJ-Biplot visualization

Usage

```
run_statis_dashboard()
```

Value

No return value, called for side effects (launches a Shiny application).

Examples

```
if (interactive()) {
  run_statis_dashboard()
}
```

`simulate_pharma_batches` *Simulate Pharmaceutical Manufacturing Batches (Realistic Variability)*

Description

Simulates pharmaceutical manufacturing batches across two phases. Phase 1 includes 10 under-control batches, each with natural variability in mean and covariance. Phase 2 includes 2 clean under-control batches and 3 out-of-control batches with shifted mean, increased dispersion, and moderate contamination.

Usage

```
simulate_pharma_batches(obs_per_batch = 30, seed = 780)
```

Arguments

`obs_per_batch`	Integer. Number of observations per batch. Default is 30.
`seed`	Optional integer. If provided, sets a random seed for reproducibility.

Details

The simulated data includes four quality control variables: Concentration, Humidity, Dissolution, and Density.

Value

A data frame with 450 observations and the following columns:

Batch Factor. Batch identifier (Batch_1 to Batch_15).

Phase Factor. Phase of the process: "Phase 1" or "Phase 2".

Status Factor. Control status: "Under Control" or "Out of Control".

Concentration, Humidity, Dissolution, Density Numeric quality control variables.

Index