

# Package ‘HotellingEllipse’

October 12, 2022

**Title** Hotelling T-Square and Confidence Ellipse

**Version** 1.1.0

**Description** Functions to compute the semi-axes lengths and coordinate points of Hotelling ellipse. Bro and Smilde (2014) <[DOI:10.1039/c3ay41907j](https://doi.org/10.1039/c3ay41907j)>. Brerton (2016) <[DOI:10.1002/cem.2763](https://doi.org/10.1002/cem.2763)>.

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**Encoding** UTF-8

**LazyData** true

**LazyDataCompression** xz

**RoxygenNote** 7.2.0

**URL** <https://github.com/ChristianGoueguel/HotellingEllipse>

**BugReports** <https://github.com/ChristianGoueguel/HotellingEllipse/issues>

**Imports** dplyr, FactoMineR, ggforce, ggplot2, lifecycle, magrittr, purrr, stats, tibble

**Depends** R (>= 2.10)

**Suggests** rmarkdown, knitr, markdown, testthat (>= 3.0.0), spelling, covr

**VignetteBuilder** knitr

**Config/testthat/edition** 3

**Language** en-US

**NeedsCompilation** no

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**Repository** CRAN

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ellipseCoord	<i>Coordinate Points Of Hotelling Ellipse</i>
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### Description

Get the  $x$  and  $y$  coordinates of Hotelling ellipse.

### Usage

```
ellipseCoord(data, pcx = 1, pcy = 2, conf.limit = 0.95, pts = 200)
```

### Arguments

<code>data</code>	Data frame or tibble of PCA, PLS, and ICA scores, or from other feature projection methods.
<code>pcx</code>	Integer specifying which component is on the x-axis (by default 1).
<code>pcy</code>	Integer specifying which component is on the y-axis (by default 2).
<code>conf.limit</code>	Number between 0 and 1 specifying the confidence level (by default 0.95).
<code>pts</code>	Integer indicating the number of points for drawing the Hotelling ellipse (by default 200).

### Value

Data frame containing the  $x$  and  $y$  coordinate points of the Hotelling ellipse.

### Author(s)

Christian L. Goueguel, christian.goueguel@gmail.com

### Examples

```
## Principal components analysis (PCA)
library(dplyr)
set.seed(123)
pca_mod <- specData %>%
  dplyr::select(where(is.numeric)) %>%
  FactoMineR::PCA(scale.unit = FALSE, graph = FALSE)

## Extract PCA scores
pca_scores <- pca_mod %>%
```

```
purrr::pluck("ind", "coord") %>%
  tibble::as_tibble()

## Get Hotelling ellipse coordinate points
library(HotellingEllipse)
xy_coord <- ellipseCoord(data = pca_scores, pcx = 1, pcy = 2, conf.limit = 0.95, pts = 200)
```

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ellipseParam	<i>Lengths Of The Semi-Axes Of Hotelling Ellipse</i>
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## Description

Compute the lengths of the semi-axes of Hotelling ellipse.

## Usage

```
ellipseParam(data, k = 2, pcx = 1, pcy = 2)
```

## Arguments

data	Data frame or tibble of PCA, PLS, or ICA scores, or from other feature projection methods.
k	Integer specifying the number of components (by default 2).
pcx	Integer specifying which component is on the x-axis (by default 1).
pcy	Integer specifying which component is on the y-axis (by default 2).

## Value

Returns a list that includes:

1. **Tsquare** Data frame containing the T-squared statistic.
2. **Ellipse** Data frame containing the lengths of the semi-minor and semi-major axes.
3. **cutoff.99pct** Number corresponding to the T-square cutoff at 99% confidence level.
4. **cutoff.95pct** Number corresponding to the T-square cutoff at 95% confidence level.

## Author(s)

Christian L. Goueguel, christian.goueguel@gmail.com

## Examples

```
## Principal components analysis (PCA)
library(dplyr)
set.seed(123)
pca_mod <- specData %>%
  dplyr::select(where(is.numeric)) %>%
  FactoMineR::PCA(scale.unit = FALSE, graph = FALSE)

## Extract PCA scores
pca_scores <- pca_mod %>%
  purrr::pluck("ind", "coord") %>%
  tibble::as_tibble()

## Get Hotelling T2-value and the lengths of the ellipse semi-axes
library(HotellingEllipse)
T2 <- ellipseParam(data = pca_scores, k = 2, pcx = 1, pcy = 2)
```

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specData	<i>LIBS spectra of 100 soil samples</i>
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## Description

A dataset containing the emission spectra of 171 soils measured in laboratory conditions. The samples were cleaned, dried, homogenized, sieved (10 Mesh size) and thereafter pelletized prior to LIBS measurements. LIBS spectra were preprocessed by performing baseline removal.

## Usage

```
specData
```

## Format

Data frame of 100 rows (soil samples) and 3152 columns (wavelengths).

## Source

[doi:10.1039/C9JA00090A](https://doi.org/10.1039/C9JA00090A)

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