

Latex-Template



PROJECT TITLE

Project Description

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TITLE

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rédigé par  
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03 Mar 2024

## Abstract

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contribution

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Latex-Template/issues

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Notation	Signification
Category A	
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# List of Algorithms

# Chapter 1

# Chapter 1

## Contents

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## 1.1 section example

### 1.1.1 subsection example

Lorem ipsum dolor sit amet. Ut expedita sunt est delectus quia ad nostrum delectus eum magni dolor. Eos nemo minima sit deleniti porro et necessitatibus minima ab quia necessitatibus in beatae autem et voluptas labore. Lorem ipsum dolor sit amet. Ut expedita sunt est delectus quia ad nostrum delectus eum magni dolor. Eos nemo minima sit deleniti porro et necessitatibus minima ab quia necessitatibus in beatae autem et voluptas labore.

$$\mathbb{Y} = m(x) + \varepsilon$$

## Chapter 2

# Chapter 2

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## 2.1



Chapter 3

Chapter 3

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3.1

Appendix A

Some Appendix

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A.1 with subsection

A.2 and another one

## Appendix B

# Code Examples

### B.1 with comments

```
1 # — install — #
2 install.packages(c("fda", "fda.usc"))
3 # — general packages — #
4
5 library(data.table)
6 # — FDA packages — #
7
8 library(fda)
9 library(fda.usc)
```

### B.2 Math in code bloc

```
1 # |      date      |  $X_1$  |  $X_2$  | ... |  $X_p$  |
2 # | Jan 1st 12:00 | :   | :   |     | :   |
3 data <- fread("data.csv")
4
5
6 # un individu = une ligne
7 # donc pour une série temporelle, il faut transposer les observations
8   ↪ et avoir la suite des données disposées sur une ligne.
9 fdata_standard_index <- fda.usc::fdata(
10   mdata = t(X),
11   argvals = to_unit_interval(
12     #           ↑
13     # on doit ramener les dates dans l'intervalle [0,1]
14     data[, .(date)]
15   )
16 )
```

### B.3 some generic code

```
1 nb_points <- ncol(fdata)
2 nb_ts <- nrow(fdata)
3
4 fda_optim_basis <- fda.usc::optim.basis(
5   fdataobj =
6     ↪ select_representative_observations_for_mean_function_fdata(fdata_ts
7     ↪ = fdata, is_iid = is_iid),
8   type.CV = fda.usc::GCV.S,
9   W = NULL,
10  lambda = lambda_CV_look_list,
11  numbasis = num_basis__seq,
12  type.basis = "bspline",
13  verbose = TRUE
14 )
```

another code block :

```
1 fda_optimal_basis <- ...
2 fdata_obj_temp <- fda_optimal_basis[["fdata.est"]]
3 fdata_obj <- fda.usc::fdata2fd(fdata_obj_temp)
4 fpca_result <- fda::pca.fd(
5   fdataobj = fdata_obj,
6   nharm = 3,
7   # centrer les données
8   centerfns = TRUE
9 )
```

### B.4 inline block with math

Regardons désormais à quoi ressemble la sortie :

$$\text{fpca\_result}\$scores = \downarrow [X_i] \begin{matrix} \xrightarrow{[\phi_k]} \\ \begin{bmatrix} \ddots & \dots & \vdots \\ \vdots & \xi_i^{[k]} = \langle X_i - \mu | \phi_k \rangle & \vdots \\ \dots & \dots & \ddots \end{bmatrix} \end{matrix}$$

## Appendix C

### Article's Appendix

C.1

C.2

# Bibliography

- [1] A. Monfort C. Gourieroux and A. Trognon. Pseudo maximum likelihood methods: Theory. The Econometric Society, 52(3), 1984. pages 681-700. DOI : <https://doi.org/10.2307/1913471>.