### LATEX-TEMPLATE



## PROJECT TITLE

Project Description

## **TITLE**

**rédigé par** Allemand Instable

#### Résumé

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.

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#### contribution

si jamais vous apercevez des fautes dans le polycopié, merci de rédiger une <u>issue</u> sur Github à l'adresse :

correctif



LaTeX-Template/issues

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Notation	Signification
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## Chapitre 1

## **Chapter 1**

#### Contents

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### 1.1 section example

#### 1.1.1 subsection example

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# Chapitre 2

# Chapter 2

Contents	
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2.1

# Chapitre 3

# Chapter 3

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3.1	 3

3.1

## Annexe A

# **Some Appendix**

$\mathbf{A.1}$	with subsection	i
$\mathbf{A.2}$	and another one	i

### A.1 with subsection

### A.2 and another one

### **Annexe B**

## **Code Examples**

#### **B.1** with comments

```
# --- install --- #
install.packages(c("fda", "fda.usc"))
# --- general packages --- #

library(data.table)
# --- FDA packages --- #

library(fda)
library(fda.usc)
```

#### **B.2** Math in code bloc

```
\mid X_1 \mid X_2 \mid \cdots \mid X_p \mid
            date
    # | Jan 1st 12:00 | : | : |
    data <- fread("data.csv")</pre>
    # un individu = une ligne
    # donc pour une série temporelle, il faut transposer les observations et avoir la
    → suite des données disposées sur une ligne.
    fdata_standard_index <- fda.usc::fdata(</pre>
8
        mdata = t(X),
9
        argvals = to_unit_interval(
10
11
        # on doit ramener les dates dans l'intervalle [0,1]
            data[, .(date)]
14
15
```

#### B.3 some generic code

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

#### another code block:

```
fda_optimal_basis <- ...
fdata_obj_temp <- fda_optimal_basis[["fdata.est"]]

fdata_obj <- fda.usc::fdata2fd(fdata_obj_temp)

fpca_result <- fda::pca.fd(

fdobj = fdata_obj,

nharm = 3,

# centrer les données

centerfns = TRUE

)</pre>
```

#### B.4 inline block with math

Regardons désormais à quoi ressemble la sortie :

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

# **Bibliographie**

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