

Probabilistic modelling and functional data analysis of sleep structure

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July 12, 2018

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Sleep and daily behaviour

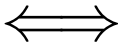


- 146 subjects, European sleep project SIESTA [Klösch et al., 2001]
- EEG recording of two consecutive nights spent in the sleep laboratory
- results of measures representing daily behaviour
 - age, gender
 - subjectively scored sleep and awakening quality
 - level of mood and drowsiness after awakening
 - physiological factors (blood pressure, pulse rate)
 - cognitive tests for working–memory, concentration, fine–motor activity
 - ...

Sleep structure and daily measures

Sleep structure

- total sleep time
- sleep efficiency
- sleep latency
- time spent awake
- time spent in each sleep stage
- ...



Daily measures

- subjectively scored sleep quality
- level of drowsiness, mood
- fine-motor activity test
- working-memory test
- pulse rate
- blood pressure
- ...

correlation coefficient,

linear regression,

prediction,

⋮

→ the dynamics of the sleep process is not taken into account

Probabilistic sleep model (PSM)

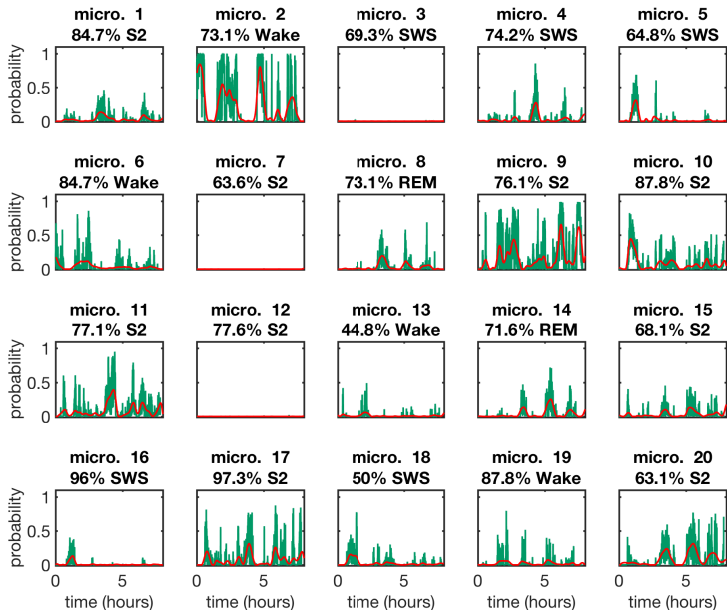
- [Lewandowski et al., 2012]
- 20 sleep microstates
- three-second long time segments of the EEG signal
- time segment $X_t \rightarrow$ AR(10) representation:
$$X_t = \sum_{i=1}^{10} a_i X_{t-i} + e_t$$
- Gaussian mixture model with latent variable Z representing membership to a sleep microstate $z \in \{1, \dots, 20\}$

$$p(a) = \sum_{z=1}^{20} p(z)p(a|z) = \sum_{z=1}^{20} \pi_z \mathcal{N}(a|\mu_z, \Sigma_z)$$

- sleep representation = posterior probability values

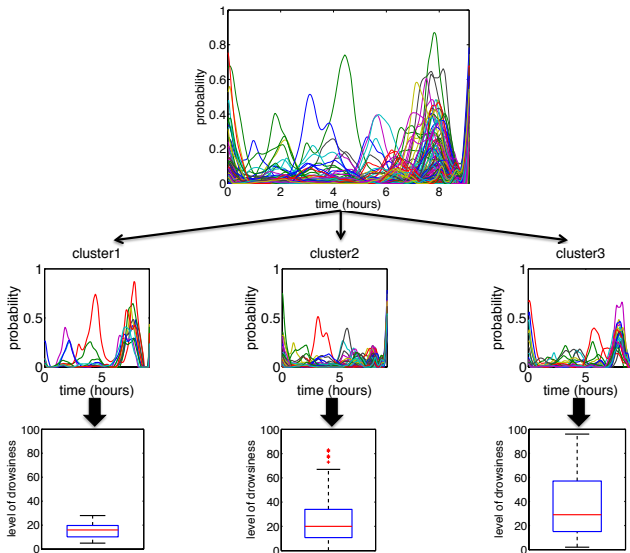
$$p(z|a) = \frac{\hat{p}(z)\hat{p}(a|z)}{\sum_{k=1}^{20} \hat{p}(k)\hat{p}(a|k)}$$

Probabilistic sleep model



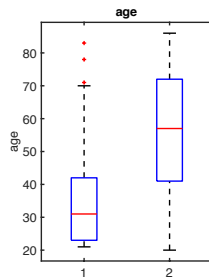
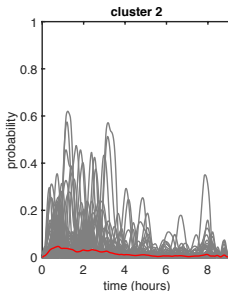
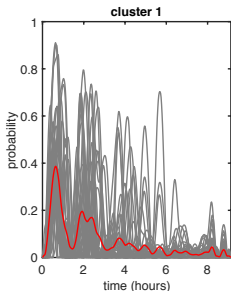
- detection of specific sleep profiles correlated with daily measures
 - cluster analysis of sleep probabilistic curves
 - time alignment of sleep probabilistic curves
- modelling individuality in the sleep pattern
 - Multilevel functional principal component analysis [Di et al., 2009]

I. Cluster analysis of sleep probabilistic curves

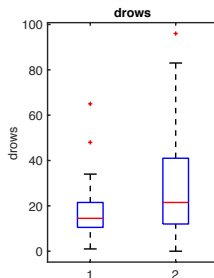
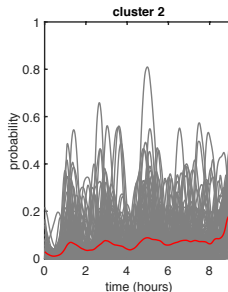
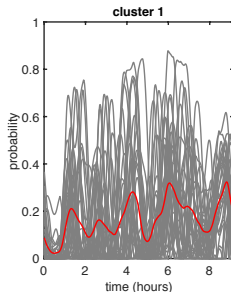


Results

microstate 16
96% SWS

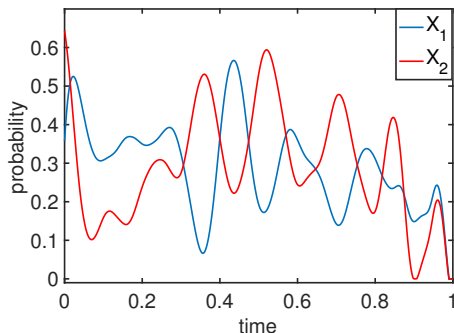


microstate 14
72% REM



Curves misalignment problem

- let X_1, X_2 to be two curves defined on the time interval $T = [0, 1]$



- the curves are **similar in shape**
 - but their important features are **shifted in time**
- **the curves are misaligned in time**

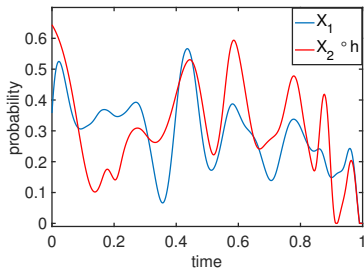
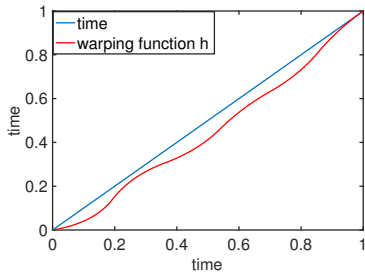
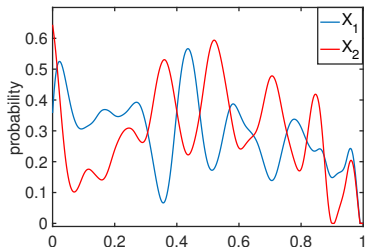
Curves alignment (registration, synchronisation)

- two curves X_1, X_2 are defined over the time interval $T = [0, 1]$
- to align a pair of curves = to find a **warping function** $h^* : T \rightarrow T$

$$h^* \in \operatorname{argmin}_h \int_T (X_1(t) - (X_2 \circ h)(t))^2 dt$$

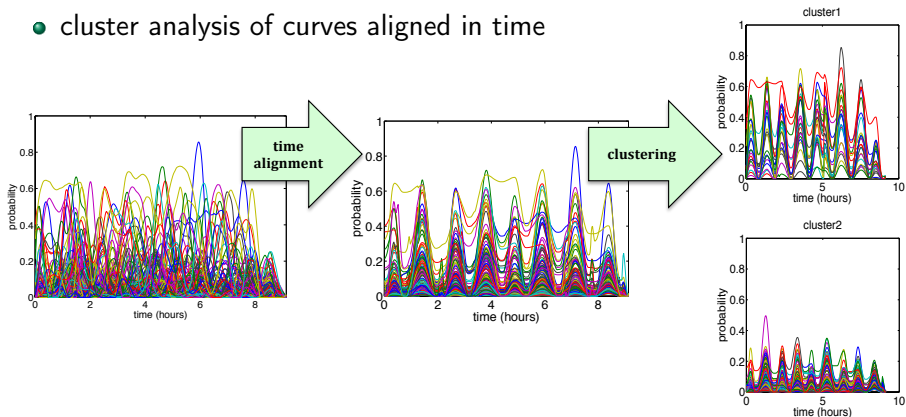
- h^* is strictly increasing,
- $h^*(0) = 0, \quad h^*(1) = 1,$
- $\int_T (h^*(t) - t)^2 dt < \lambda,$
- ...

Example of the curves alignment

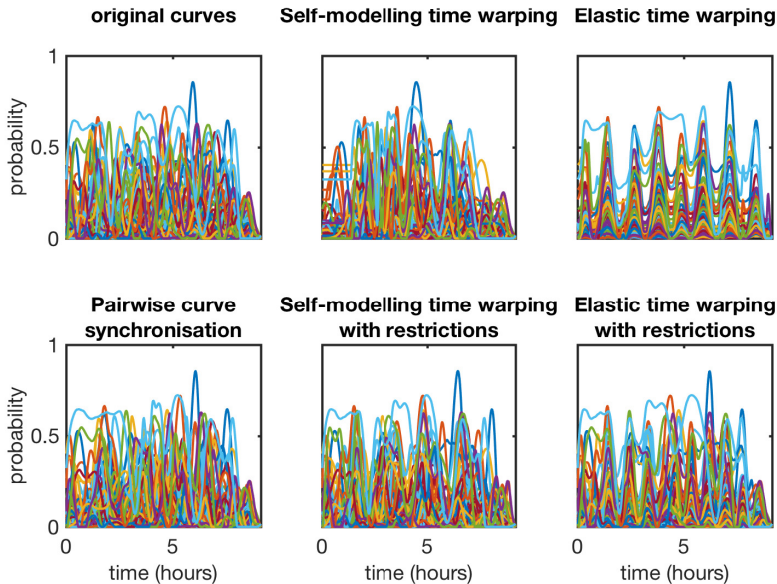


Curves alignment preceding the clustering step

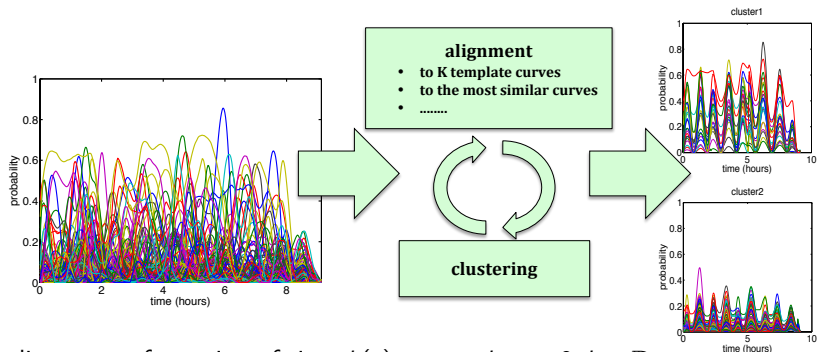
- curves are aligned as the whole dataset
 - Self-modelling time warping (SMTW) [Gervini and Gasser, 2004],
 - Pairwise curve synchronisation (PCS) [Müller and Tang, 2008],
 - Elastic time warping (ETW) [Tucker et al., 2013]
- cluster analysis of curves aligned in time



Curves alignment preceding the clustering step



Simultaneous curves alignment and clustering



- linear transformation of time $h(t) = at + b$, $a > 0$, $b \in \mathbb{R}$ [Gaffney and Smyth, 2005, Sangalli et al., 2010]
⇒ **the assumption of common start and end point is violated**
- truncated Pairwise curve synchronisation [Müller and Tang, 2008]
⇒ **not satisfactory alignment**

2step approach

- initial clustering
 - distance matrix M based on the Dynamic time warping algorithm [Wang and Gasser, 1997, Montero and Vilar, 2014]
 - k -medoids algorithm applied to the matrix M
- k^{th} step: align curves in each cluster separately
- $(k + 1)^{\text{th}}$ step: recompute matrix M for aligned curves, cluster aligned curves . . .

- stopping criteria:
 - maximal number of iterations; changes in cluster membership
 - average dissimilarity L within clusters is small

$$L = \frac{1}{N} \int_0^1 (X_i(h_i(t)) - \mu_{C_i}(t))^2 dt, \quad \mu_{C_i}(t) = \frac{1}{|C_i|} \sum_{j: X_j \in C_i} X_j(h_j(t))$$

Application to the sleep probabilistic curves

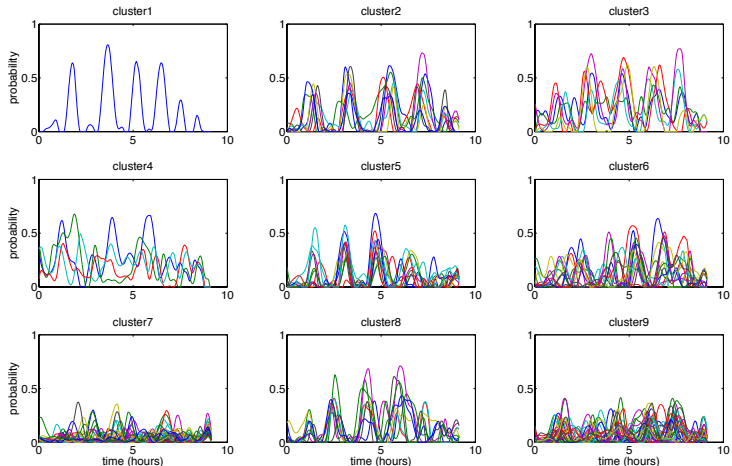


Figure: Microstate 8 (73% REM). Clustering of 146 in time misaligned sleep probabilistic curves into 9 clusters by the k -means algorithm.

Application to the sleep probabilistic curves

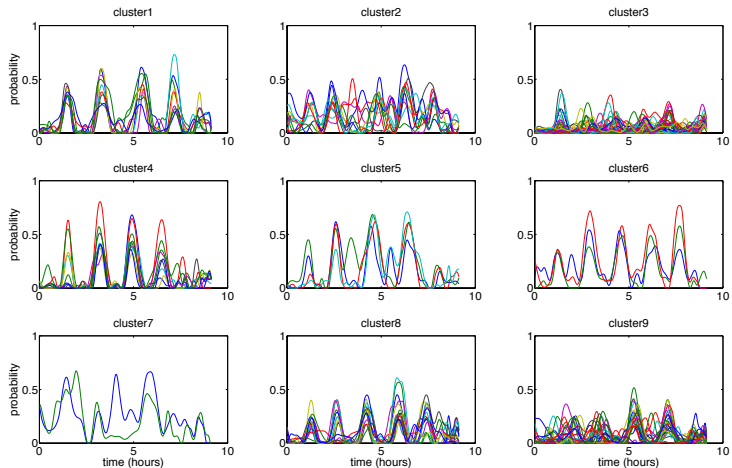


Figure: Microstate 8 (73% REM). Clustering of 146 sleep probabilistic curves into 9 clusters by the 2step approach with the SMTW algorithm used in the alignment step.

Application to the sleep probabilistic curves

Average silhouette	number of clust.	<i>k</i> -means	tPCS	2DTW-SMTW	2DTW-PCS	2DTW-ETW
Microstate 16	8	0.56	0.57	0.64	0.61	0.47
Microstate 8	9	0.33	0.34	0.48	0.44	0.46
Microstate 14	3	0.53	0.54	0.60	0.10	0.60
Microstate 1	2	0.77	0.79	0.79	0.64	0.80
Microstate 6	3	0.62	0.63	0.73	0.71	0.53

<i>L</i> -criterion	number of clust.	<i>k</i> -means	tPCS	2DTW-SMTW	2DTW-PCS	2DTW-ETW
Microstate 16	8	0.40	0.39	0.26	0.33	0.19
Microstate 8	9	0.79	0.77	0.54	0.59	0.34
Microstate 14	3	4.55	4.51	2.71	5.01	1.77
Microstate 1	2	4.07	4.05	3.37	4.34	2.84
Microstate 6	3	1.02	1.01	0.78	0.85	0.73

Results

- ↑ **Microstate 16** (96% *SWS*) ⇒ ↑ drowsiness
- ↑ **Microstate 13** (44% *Wake*) ⇒ ↓ cognitive tests

- ↑ micro. similar to **Wake, S1** ⇒ people above 60 years of age
- ↑ micro. similar to **S2, SWS** ⇒ people under 40 years of age

- ↑ micro. similar to **REM** ⇒ ↑ mood, ↑ drive, ↓ drowsiness
⇒ ↑ subjectively scored sleep quality

- ↑ micro. similar to **Wake** ⇒ ↓ mood, ↓ drive, ↑ drowsiness
⇒ ↓ subjectively scored sleep quality
 - **Microstate 19** (88% *Wake*) ⇒ visible after the alignment
 - **Microstate 6** (85% *Wake*) ⇒ the effect is **lost** after the alignment

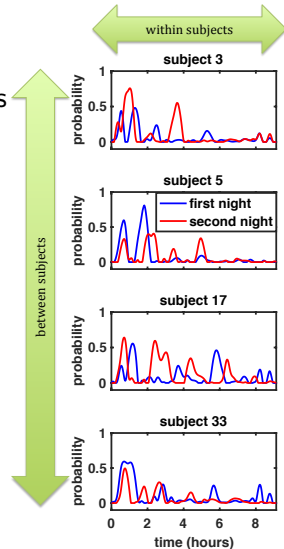
- ...

Subject-specific sleep profiles – open problem

- individual pattern in the subjects' EEG signal
 - [Finelli et al., 2001, De Gennaro et al., 2005, De Gennaro et al., 2008, Lewandowski et al., 2013],...
 - PSM is based on the EEG signal
 - the individual pattern is inherited also in the sleep probabilistic curves
- estimation of the subject-specific profiles
 - need for multiple observations (> 2)
 - Multilevel functional principal component analysis [Di et al., 2009]

Multilevel functional principal component analysis

- MFPCA, [Di et al., 2009]
- functional data observed at multiple visits
- assumptions
 - **balanced design**: the same number of observation for each subject
 - **ordered visits**: observations within subjects have natural order
 - our database
 - 146 subjects with 2 visits
 - first and second night
- two types of variability
 - between subjects
 - within subjects



Multilevel functional principal component analysis

$$X_{ij}(t) = \mu(t) + \eta_j(t) + Z_i(t) + W_{ij}(t)$$

- X_{ij} is the j^{th} sleep curve for the i^{th} subject, $i=1, \dots, 146; j = 1, 2$
- fixed effects
 - **overall mean** μ

$$\hat{\mu}(t) = \bar{X}_{..}(t) = \frac{1}{IJ} \sum_{i=1}^I \sum_{j=1}^J X_{ij}(t), \quad t \in T$$

- the **visit-specific deviation** from the overall mean η_j

$$\sum_{j=1}^J \eta_j(t) = 0, \quad t \in T$$

$$\hat{\eta}_j(t) = \bar{X}_{.j}(t) - \bar{X}_{..}(t) = \frac{1}{I} \sum_{i=1}^I X_{ij}(t) - \bar{X}_{..}(t), \quad j = 1, \dots, J$$

Multilevel functional principal component analysis

$$X_{ij}(t) = \mu(t) + \eta_j(t) + Z_i(t) + W_{ij}(t)$$

- random effects

- the subject-specific deviation from the visit-specific profile Z_i
- residual deviation from the subject- and visit-specific profile W_{ij}

→ uncorrelated stochastic processes

→ $E(Z_i(t)) = E(W_{ij}(t)) = 0, \quad t \in T$

→ covariance functions $R_Z : T \times T \rightarrow \mathbb{R}, R_W : T \times T \rightarrow \mathbb{R}$

$$Z_i(t) = \sum_{l=1}^{\infty} \alpha_{il} \phi_l^{(1)}(t) \quad W_{ij}(t) = \sum_{k=1}^{\infty} \beta_{ijk} \phi_k^{(2)}(t)$$

- level 1 functional principal components $\phi_k^{(1)}, k = 1, 2, \dots$
→ eigenfunctions of R_Z
- level 2 functional principal components $\phi_l^{(2)}, l = 1, 2, \dots$
→ eigenfunctions of R_W

Covariance function estimators

- covariance functions

$$R_T(s, t) = \text{Cov}(X_{ij}(s), X_{ij}(t))$$

$$R_Z(s, t) = \text{Cov}(X_{ij}(s), X_{il}(t)), \quad j \neq l$$

$$R_W(s, t) = R_T(s, t) - R_Z(s, t), \quad s, t \in T$$

- estimators by the method of moments

- $$\hat{R}_T(s, t) = \frac{\sum_{i=1}^I \sum_{j=1}^J (X_{ij}(s) - \hat{\mu}(s) - \hat{\eta}_j(s))(X_{ij}(t) - \hat{\mu}(t) - \hat{\eta}_j(t))}{IJ}$$

- $$\hat{R}_Z(s, t) = \frac{\sum_{i=1}^I \sum_{j \neq l}^J (X_{ij}(s) - \hat{\mu}(s) - \hat{\eta}_j(s))(X_{il}(t) - \hat{\mu}(t) - \hat{\eta}_l(t))}{IJ(J-1)}$$

- $$\hat{R}_W(s, t) = \hat{R}_T(s, t) - \hat{R}_Z(s, t)$$

Algorithm

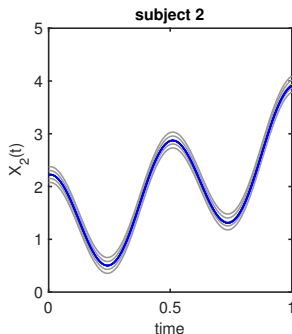
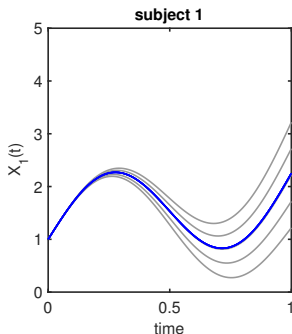
- 1 compute $\hat{\mu}$ and $\hat{\eta}_j$
- 2 compute \hat{R}_T and \hat{R}_Z , $\hat{R}_W = \hat{R}_T - \hat{R}_Z$
- 3 using \hat{R}_Z estimate the level 1 functional principal components
- 4 using \hat{R}_W estimate the level 2 functional principal components
- 5 appropriate number of principal components on both levels
- 6 estimate level 1 and level 2 principal component scores
→ BLUP, MCMC, ...

Example

- 2 subjects, 10 curves

$$X_{1j}(t) = e^t + \sin(2\pi t) + \left(1 - \frac{j}{2}\right) t^2, \quad j = 1, \dots, 5; \quad t \in [0, 1],$$

$$X_{2k}(t) = e^t + \cos(4\pi t) + 0.15 \frac{k}{2}, \quad k = 1, \dots, 5.$$

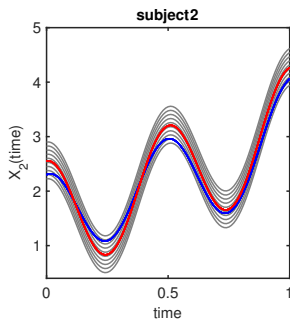
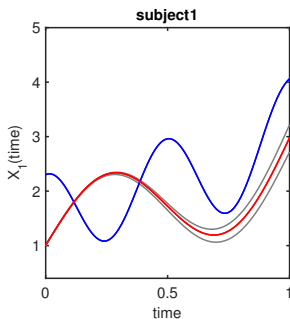


Example

- 2 subjects, 12 curves

$$X_{1j}(t) = e^t + \sin(2\pi t) + \left(1 - \frac{j}{2}\right) t^2, \quad j = 1, 2; \quad t \in [0, 1],$$

$$X_{2k}(t) = e^t + \cos(4\pi t) + 0.15 \frac{k}{2}, \quad k = 1, \dots, 10.$$



- the assumptions of MFPCA are violated
 \Rightarrow the method is not able to properly estimate subject-specific profiles

MFPCA for unbalanced data with unordered visits

- unordered visits $\rightarrow \eta_j \equiv 0$
- estimator for $\mu \rightarrow$ unweighted mean

$$\hat{\mu}(t) = \frac{1}{I} \sum_{i=1}^I \frac{1}{J_i} \sum_{j=1}^{J_i} X_{ij}(t) = \frac{1}{I} \sum_{i=1}^I \bar{X}_i(t)$$

- modified estimator \hat{R}_W

$$\hat{R}_W(s, t) = \frac{1}{2} \sum_{i=1}^I \sum_{j=1}^{J_i} \sum_{l \neq j}^{J_i} w_i (X_{ij}(s) - X_{il}(s)) (X_{ij}(t) - X_{il}(t)),$$
$$\sum_{i=1}^I w_i J_i (J_i - 1) = 1 \quad \text{and} \quad w_i \geq 0, \quad i = 1, \dots, I$$

and \hat{R}_W minimises

$$\mathbb{E} \left(\|\hat{R}_W - R_W\|^2 \right)$$

MFPCA for unbalanced data with unordered visits

- covariance function estimators

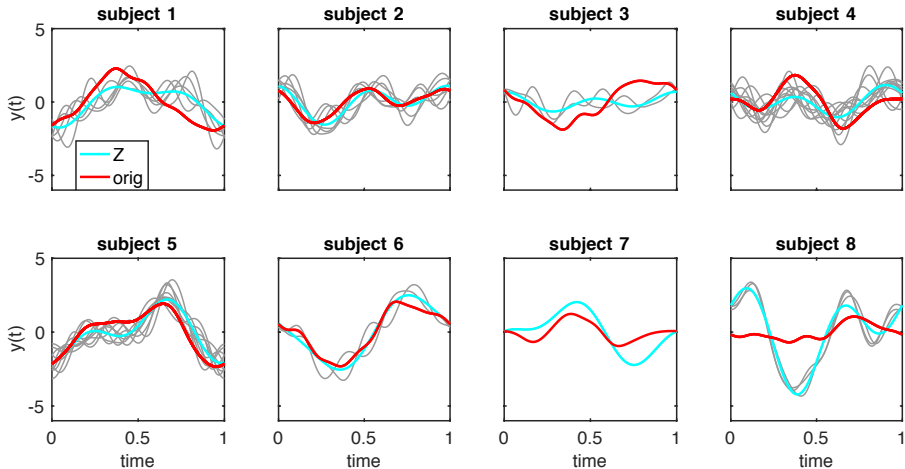
$$\hat{R}_W(s, t) = \frac{1}{2} \sum_{i=1}^l \sum_{j=1}^{J_i} \sum_{l \neq j}^{J_i} \frac{1}{(N_1 - l)J_i} (X_{ij}(s) - X_{il}(s)) (X_{ij}(t) - X_{il}(t))$$

$$\hat{R}_T(s, t) = \frac{1}{N_1} \sum_{i=1}^l \sum_{j=1}^{J_i} (X_{ij}(s) - \hat{\mu}_{uw}(s)) (X_{ij}(t) - \hat{\mu}_{uw}(t))$$

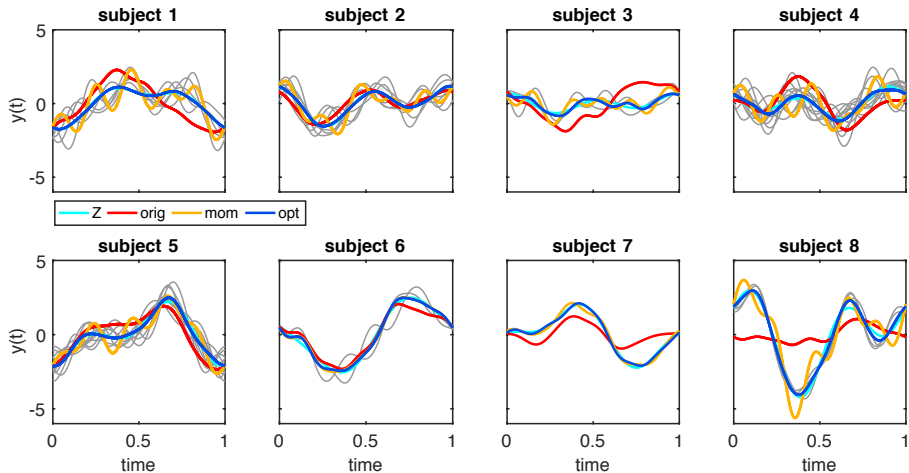
$$\hat{R}_B(s, t) = \frac{l}{l-1} \left(\hat{R}_T(s, t) - \left(1 - \frac{2}{N_1} + \frac{1}{l^2} \sum_{i=1}^l \frac{1}{J_i} - \frac{L}{N_1} \frac{l-2}{l} \right) \hat{R}_W(s, t) \right)$$

- $N_1 = \sum_{i=1}^l J_i$
- $L =$ number of subjects with only one observation ($J_i = 1$)

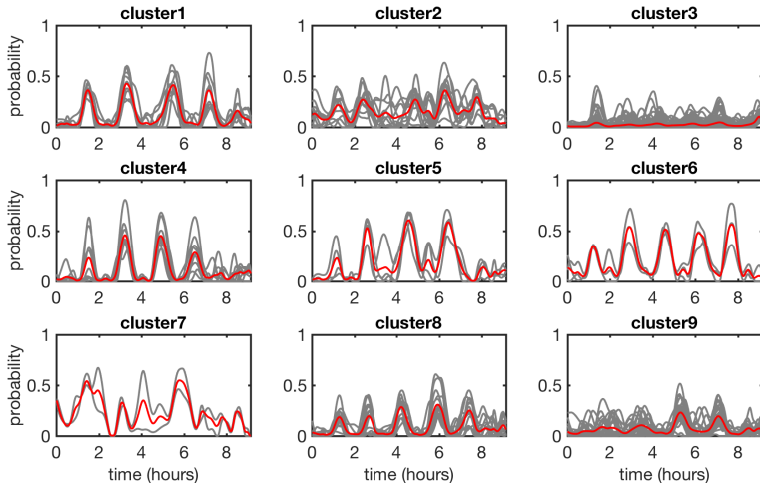
MFPCA for unbalanced data with unordered visits



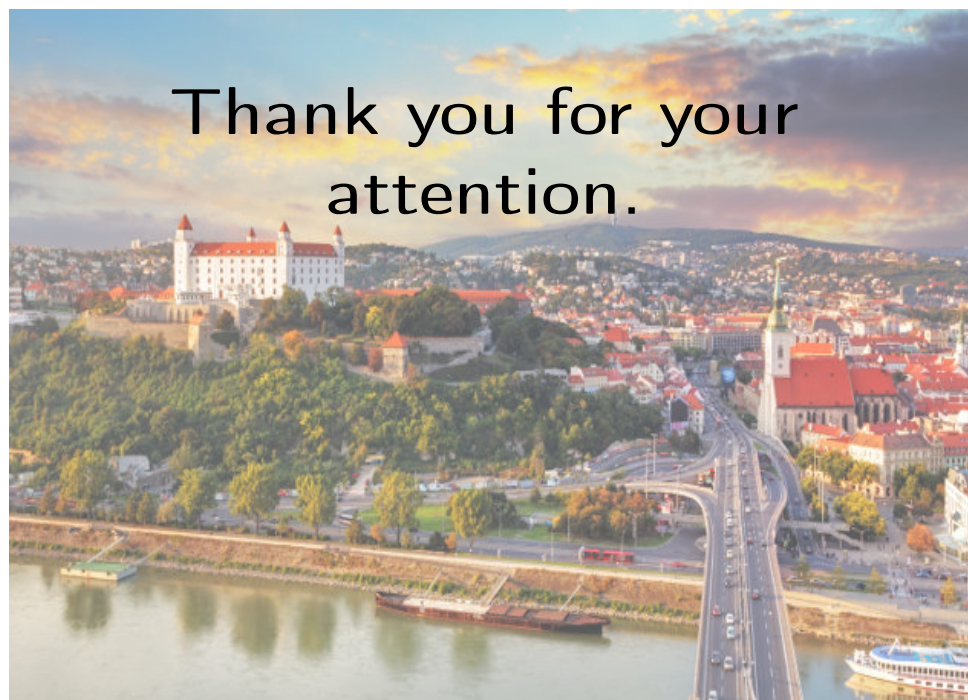
MFPCA for unbalanced data with unordered visits



Application to the results of cluster analysis



Thank you for your
attention.



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