

# Bayesian nonparametric methods for clustering Part 2: Outcome-guided analysis

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# Summary of Part 1:

### Nonparametric clustering

- Heterogeneous population with unknown clusters
- Cluster number estimated from the data
- Unconstrained number of clusters

### Characterized by

• One or several variables (Gaussian, skew t-distributions...)

#### Bayesian framework

- o Dirichlet process prior
- MCMC estimation

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### Introduction of Part 2:

### Nonparametric clustering

- Heterogeneous population with unknown clusters
- Cluster number estimated from the data
- Unconstrained number of clusters

### Characterized by

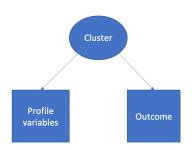
- One or several variables (Gaussian, skew t-distributions...)
- o An outcome used as noisy surrogate for clusters (Bair, 2013)

#### Bayesian framework

- Dirichlet process prior
- MCMC estimation

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### Outcome-guided clustering



- → Subgroups with specific outcome distributions and variable profiles
- ightarrow Independence between profile variables and outcome conditionally on the clusters

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# Motivational example - Dementia study

#### Dementia

Syndrome (chronic or progressive nature)

Deterioration in cognitive functions

Impairment of the social and occupational functioning.



#### Precision medicine

Improve disease prevention and treatment

Refine patient characterisation

Predict disease risk and treatment effect at the individual level

# Profile regression: methodology

#### **Notations:**

 $Y_i$  vector of individual outcome measurements  $W_i = (w_{i1}, ..., w_{iP})^{\top}$  individual vector of profile variables  $c_i$  allocation variable

#### Likelihood:

$$L(\theta|Y,W) = \prod_{i=1}^{N} \sum_{g}^{\infty} P(c_i = g|\theta^c) f(W_i|c_i = g;\theta_g^W) f(Y_i|c_i = g;\theta_g^Y)$$

Independence assumption between W and Y given the cluster allocations.

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# Profile regression: membership probability

#### **Notations:**

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#### Likelihood:

$$L(\theta|Y,W) = \prod_{i=1}^{N} \sum_{g}^{\infty} \underbrace{P(c_i = g|\theta^c)}_{\text{Membership prob.}} f(W_i|c_i = g;\theta_g^W) f(Y_i|c_i = g;\theta_g^Y)$$

Independence assumption between W and Y given the cluster allocations.

#### Sub-model: Dirichlet Process mixture model

$$\mathcal{A} = \sum_{g=1}^{G} \pi_g \ \delta_{\theta_g}$$
  $\pi | \alpha \sim \textit{Dirichlet}(\alpha/G, ..., \alpha/G)$   $c_i | \pi \sim \textit{Multinom}(\pi)$   $\alpha$ : Concentration parameter  $\theta_{c_i} | \mathcal{A}_0 \sim \mathcal{A}_0$   $Y_i, W_i | c_i \sim f(\cdot; \theta_c)$ 

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Independence assumption between W and Y given the cluster allocations.

#### Sub-model: Dirichlet Process mixture model

$\mathcal{A} = \sum_{g=1}^{+\infty} \pi_g  \delta_{\theta_g}$	$m{\pi}   lpha \sim \textit{GEM}(lpha) \ c_i   m{\pi} \sim \textit{Multinom}(m{\pi})$
$\alpha$ : Concentration parameter	$ heta_{c_i}   \mathcal{A}_0 \sim \mathcal{A}_0$
$\mathcal{A}_0$ : Base distribution	$Y_i, W_i   c_i \sim f(\cdot; \theta_c)$

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# Profile regression: profile variables

#### **Notations:**

 $Y_i$  vector of individual outcome measurements  $W_i = (w_{i1}, ..., w_{iP})^{\top}$  individual vector of profile variables  $c_i$  allocation variable

#### Likelihood:

$$L(\theta|Y,W) = \prod_{i=1}^{N} \sum_{g}^{\infty} P(c_i = g|\theta^c) \underbrace{f(W_i|c_i = g;\theta_g^W)}_{\text{Profile variables}} f(Y_i|c_i = g;\theta_g^Y)$$

Independence assumption between W and Y given the cluster allocations.

Sub-model for profile variables: given  $c_i = g$ ,

Discrete case:  $W_i^{(1)} \sim \textit{Multinom}(\phi_{\mathbf{g}})$ Continuous case:  $W_i^{(2)} \sim \mathcal{N}_P(\mu_{\mathbf{g}}; V_g)$ 

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# Profile regression: outcome

#### **Notations:**

 $Y_i$  vector of individual outcome measurements  $W_i = (w_{i1}, ..., w_{iP})^{\top}$  individual vector of profile variables  $c_i$  allocation variable

#### Likelihood:

$$L(\theta|Y,W) = \prod_{i=1}^{N} \sum_{g}^{\infty} P(c_i = g|\theta^c) \ f(W_i|c_i = g;\theta_g^W) \underbrace{f(Y_i|c_i = g;\theta_g^Y)}_{Outcome}$$

Sub-model for outcome: given cluster g,

$$Y_i \sim \mathcal{N}(\theta_g + \beta^\top X_i, \sigma_g^2)$$

Other types of outcomes: Bernoulli, Poisson, Binomial, Categorical, Quantile, Survival

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# Estimated cluster-specific parameters

• MCMC Gibbs sampler:  $\Theta = (\alpha, [\phi]_{\mathfrak{g}}, [\mu]_{\mathfrak{g}}, [V]_{\mathfrak{g}}, c, \beta)$ 

- Posterior similarity matrix S from pairwise co-clustering probability
- Selection of the representative clustering P\*

Partitioning around medoids on dissimilarity matrix  $1-\mathcal{S}$  Average silhouette width

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• MCMC Gibbs sampler:  $\Theta = (\alpha, [\phi]_g, [\mu]_g, [V]_g, c, \beta)$   $\alpha \sim f(\alpha \mid [\phi]_\sigma, [\mu]_\sigma, [V]_\sigma, c, \beta)$ 

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$$\alpha \sim f(\alpha \mid [\phi]_g, [\mu]_g, [V]_g, c, \beta)$$
 
$$[\phi]_g \sim f([\phi]_g \mid \alpha, [\mu]_g, [V]_g, c, \beta) \text{ etc...}$$

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### Estimation

Cluster-specific parameters as the average over the MCMC iterations:

$$\bar{\Theta}_{g}^{(k)} = \frac{1}{N_{g}} \sum_{i \mid c^{*} = g} \Theta_{c_{i}^{(k)}}$$

 $\Theta_{c_i^{(k)}}$  parameter for the cluster subject *i* is allocated to, at iteration *k*  $N_g$  the number of subjects in cluster *g* of  $P^*$ .

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### Estimation

Cluster-specific parameters as the average over the MCMC iterations:

$$\hat{\Theta}_{g} = \frac{1}{K} \sum_{k=1}^{K} \frac{1}{N_{g}} \sum_{i \mid c^{*} = g} \Theta_{c_{i}^{(k)}}$$

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### Estimation

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- $\rightarrow$  Integration of the clustering uncertainty in the parameters estimation.
- ightarrow Empirical credible intervals computed from MC sample of  $ar{\Theta}_{g}^{(k)}$

### Convergence

Difficult to assess convergence based on parameters traces:

- Fixed effects converge quickly
- Cluster-specific parameters not traceable
- If stuck in local mode,  $\alpha$  not good indicator (Hastie et al., 2014)
- $\rightarrow$  Marginal partition posterior p(C|Y,W)
  - identify runs that were significantly different from others, perhaps due to convergence issues
  - which run explored the higher posterior probability regions

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### Application to ADNI data - (Rouanet et al. 2020)

**Objective:** Identify subgroups of the population associated with specific cognitive evolution patterns and brain imaging profiles.

- North American Alzheimer's Disease Neuroimaging Initiative
- o Sample of 199 subjects, 55 years old and over, 8-year follow-up
- o Normalised Mini-Mental State Examination (Philipps et al., 2014)

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- Sample of 199 subjects, 55 years old and over, 8-year follow-up
- Normalised Mini-Mental State Examination (Philipps et al., 2014)
- Profile variables:
  - → Standardised MRI brain volumes: whole brain, ventricles, hippocampus, entorhinal cortex, fusiform gyrus and middle temporal gyrus, standardised by the intracerebral volume
  - → Gender (1 for women, 0 for men) and Education (1 if greater than or equal to 16 years of education, 0 otherwise), APOE4 status (1 if 1 or 2 APOE4 alleles, 0 otherwise)

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# Model specification

### Profile variables, given cluster g:

Gender, Sex, Education:

$$W_i^{(1)} \sim \textit{Bernouilli}(\phi_g)$$
 with  $\phi_g \sim \textit{Dirichlet}(a_j)$ 

Standardized MRI brain volumes:

$$W_i^{(2)} \sim \mathcal{N}_P(\mu_g; V_g)$$
 with  $\mu_g \sim \mathcal{N}_P(\mu_0; V_0)$ ,  $V_g \sim InvWishart(R_0, K_0)$ 

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Longitudinal normalized MMSE, given cluster g:

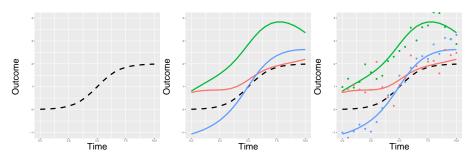
$$Y(t_i) = f_g(t_i) + \epsilon_g(t_i)$$
 with  $\epsilon_g \sim \mathcal{N}(0, L_{1,g})$   
 $f_g(\cdot) \sim GP(\mathbf{0}, \mathcal{K}_g(\cdot, \cdot))$  with  $\mathcal{K}_g(s, t) = L_{g,2} \exp(-\frac{(s-t)^2}{L_{g,3}})$   
 $\exp(L_{g,l}) \sim \mathcal{N}(0, \sigma_l^2)$ 

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### Gaussian Process prior

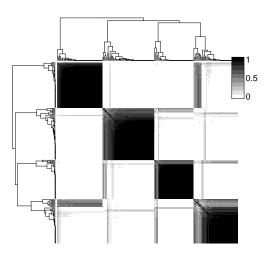
### Given the cluster g:

$$Y_{ij} = f_g(t_{ij}) + \epsilon_{ijg} \text{ with } \epsilon_{ijg} \sim \mathcal{N}(0, L_{g,1})$$
  
 $f_g(\cdot) \sim GP(m_g(\cdot), \mathcal{K}_g(\cdot, \cdot))$ 



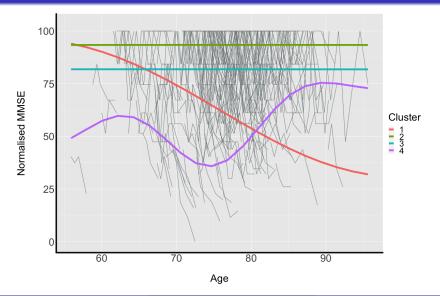
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# Posterior similarity matrix



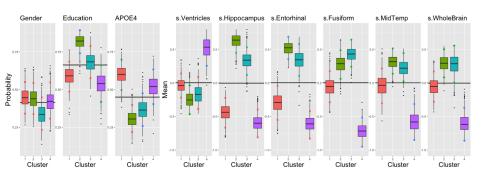
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# Cluster-specific cognitive patterns



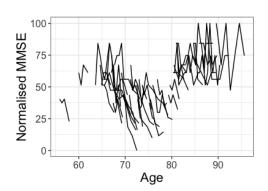
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# Cluster profiles

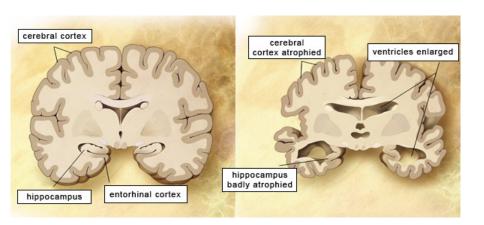


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# Cognitive pattern in cluster 4



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### Discussion

#### Conclusion

Mixture model with unconstrained number of clusters

Outcome-guided clustering analysis

Integration of different types of profile variables

Extension of the package PReMiuM

#### **Practical**

Hands-on exercises on the PReMiuM package (profRegr function)

Data on cognitive decline from lcmm package

Details on DP parameters

### References

Rouanet A, et al. (2020) Benefit of Bayesian clustering of longitudinal data: study of cognitive decline for precision medicine. In Bayesian Methods in Pharmaceutical Research (eds: Lesaffre E, Baio G, and Boulanger B), Chapman & Hall/CRC Biostatistics Series, ISBN: 978-1-138-74848-4 (pp. 223–242).

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