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## Outline

Motivation - use of GLM-trees algorithm in mortality

Q GLM-trees - first use cases and limits

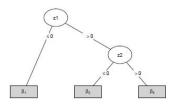
3 Multivariate GLM - single categorical variable approach

The GLM-based tree algorithm [RZ13] is a recursive split of the dataset based on partitioning variables, similar to other tree algorithm, e.g. CART.

A GLM is fitted at each node on a set of explanatory variables.

### Main steps are:

- Fit the GLM on the current sample
- Asses parameter stability for each partition variable
- Choose the best splitting point



Example from [SHZ19]:GLM tree with 2 partition variables

$$g(E(Y_i)) = \left\{ \begin{array}{ll} \beta_1 & \text{if } z_1 \leqslant 0 \\ \beta_2 & \text{if } z_1 > 0 \text{ and } z_2 \leqslant 0 \\ \beta_3 & \text{if } z_1 > 0 \text{ and } z_2 > 0 \end{array} \right.$$

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 $\textbf{Advantages:} \quad \text{Automatic partitioning} \rightarrow \text{enables classification} + \text{increased accuracy}.$ 

**Drawbacks:** Numerical optimization of the MLE  $\rightarrow$  high computational time.

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# Motivation - Reinterpretation of Age-Period-Cohort (M3) model

Classical formulation of APC Model for All-Causes mortality

$$D_{x,t} \sim \mathsf{Poisson}(E_{x,t}\mu_{x,t}) \quad \text{or} \quad D_{x,t} \sim \mathcal{B}(E_{x,t},q_{x,t}), \quad \log(\mu_{x,t}) = \alpha_x + \kappa_t + \gamma_{t-x}$$

Interpretation as a GLM ... with logit or cloglog link [Cur13] (binomial assumption)

$$g(\mathbb{E}[q_{x,t}]) = \alpha_x + \kappa_t + \gamma_{t-x}$$

... with categorical variables

$$z_i^{(1)}$$
 stands for age and takes value in  $\left[x_1;x_{max}
ight]: z_i^{(1),k} = 1_{z_i^{(1)}=x_k}$ 

 $z_i^{(2)}$  stands for year and takes value in  $\left[t_1;t_{max}
ight]:z_i^{(2),k}=1_{z_i^{(2)}=t_k}$ 

$$g(\mathbb{E}[q_{x,t}]) = \sum_{k=x_1}^{x_{max}} z_i^{(1),k} \alpha_k + \sum_{k=t_1}^{t_{max}} z_i^{(2),k} \kappa_k \qquad \text{intercept and single effect}$$
$$+ \sum_{i < l} \sum_{k \neq i} z_i^{(1),k} z_i^{(2),k'} \gamma_{k,k'} \qquad \text{double effect}$$

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• Data source: Human Mortality Database

Explicative variables

• Age: 50-84

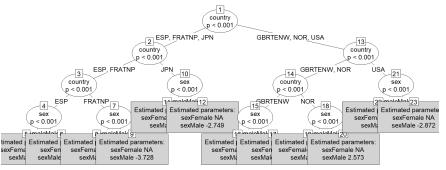
Years: 1990-2019

Partitioning variables

• Countries: France, Japan, Norway, Spain, UK, US

Sex: Male or Female

R functions glmtree (partykit) and glm (stats)



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#### Limitations

- Computation time: 39s
- Convergence: numerical optimization may fail to converge to a solution
  - → known issue with glm.fit in R when data are already well partitioned.
- Potential solutions: find closed-form formulas for the regression estimators.

# **Assumption:** the explanatory variable $z_i$ is a single categorical variable valued in $\{v_1, \ldots, v_r\}$

$$\forall i = 1, \ldots, n, \quad z_{i,k} = 1_{z_i = v_k}$$

### Simplified score formula:

If we denote  $\boldsymbol{b}_k = (\beta_k^{(1)}, \dots, \beta_k^{(d)}) \in \mathbb{R}^d$ 

$$\begin{split} s(\boldsymbol{\beta}) &= \sum_{k=1}^r \left[ \boldsymbol{J} \boldsymbol{h}(\boldsymbol{b}_k)^\mathsf{T} \times \boldsymbol{J} \boldsymbol{\mu}^{-1} (\boldsymbol{h}(\boldsymbol{b}_k))^\mathsf{T} \overline{\boldsymbol{T}}(\boldsymbol{y})^{(k)} \right]_{k,\times d} \\ &- \left[ \boldsymbol{J} \boldsymbol{h}(\boldsymbol{b}_k)^\mathsf{T} \times \boldsymbol{J} \boldsymbol{\mu}^{-1} (\boldsymbol{h}(\boldsymbol{b}_k))^\mathsf{T} \nabla_{\boldsymbol{\theta}} \kappa (\boldsymbol{\mu}^{-1} (\boldsymbol{h}(\boldsymbol{b}_k))) \right]_{k,\times d} \overline{\boldsymbol{\omega}}^{(k)} \\ &= \sum_{k=1}^r \overline{\boldsymbol{\omega}_k} \left[ \overline{\boldsymbol{T}_k}(\boldsymbol{y}) \right] \otimes \boldsymbol{e}_k - \sum_{k=1}^r \overline{\boldsymbol{\omega}_k} \left[ \nabla_{\boldsymbol{\theta}} \kappa (\boldsymbol{b}_k) \right] \otimes \boldsymbol{e}_k \text{ (for the canonical link)} \end{split}$$

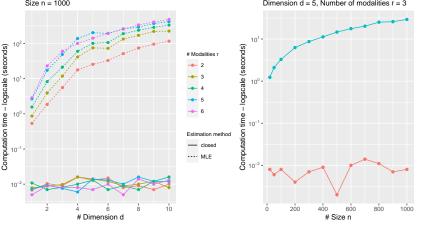
where 
$$\overline{\boldsymbol{T}_k}(\boldsymbol{y}) = \sum_{i=1}^n \frac{\omega_i z_{i,k}}{\overline{\omega_k}} \, \boldsymbol{T}(\boldsymbol{y}_i), \quad \overline{\omega_k} = \sum_{i=1}^n \omega_i z_{i,k}.$$

**Particular cases:** Multinomial and Dirichlet distributions. Univariate distributions were already looked at in [BDR20, BDR22].

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Estimation method closed MLE

## Computation time in function of Dimension d Computation time in function of Size n Size n = 1000Dimension d = 5. Number of modalities r = 310<sup>1</sup> -# Modalities r



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#### New MGLM framework based on APC model

$$(D_{x,t}^{(1)}, \dots, D_{x,t}^{(d)}) \sim \mathcal{M}_d(E_{x,t}, q_{x,t} = (q_{x,t}^{(1)}, \dots, q_{x,t}^{(d)}))$$

$$g(\mathbb{E}\begin{pmatrix} q_{x,t}^{(1)} \\ \vdots \\ q_{x,t}^{(d)} \end{pmatrix}) = \begin{pmatrix} \alpha_x^{(1)} + \kappa_t^{(1)} + \gamma_{t-x}^{(1)} \\ \vdots \\ \alpha_x^{(d)} + \kappa_t^{(d)} + \gamma_{t-x}^{(d)} \end{pmatrix}$$

#### Estimate model parameters:

If  $\eta$  is the single categorical variable estimator,  $\theta = (\alpha_x^{(1)}, \dots, \alpha_x^{(d)}, \kappa_t, \dots, \gamma_c, \dots)$  the vector of parameters of the APC model, such that  $\theta = Q\eta$ , and R a contrast matrix for identifiability, then

$$\tilde{\theta} = (Q^{\mathsf{T}}Q + R^{\mathsf{T}}R)^{-1}Q^{\mathsf{T}}\tilde{\eta}$$

is an estimator of the model parameters.

 Next step: Find a smart way to partition the data according to common trends between causes-of-death.

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## Bibliography I



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