

Joint models for a longitudinal marker and multivariate survival data

Loïc Ferrer

Institut Curie, INSERM U900, Saint-Cloud, France

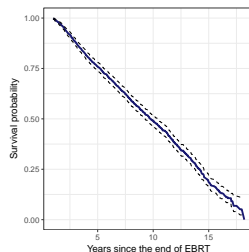
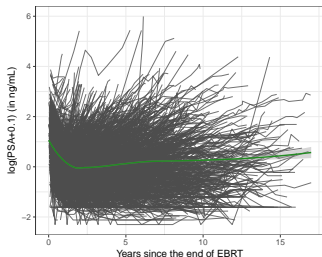
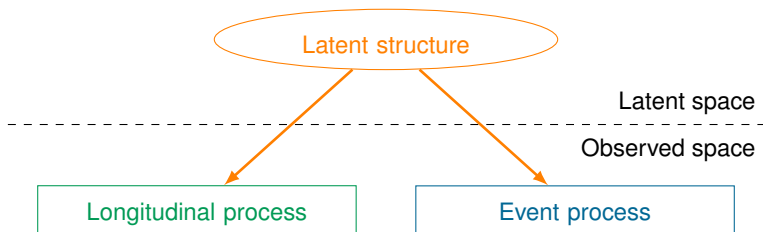
email: loic.ferrer@curie.fr

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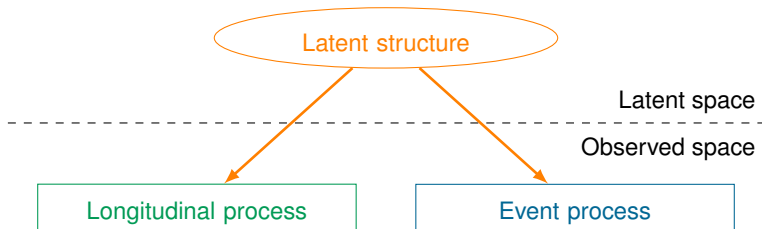
Joint modelling principle

Simultaneous modelling of correlated longitudinal and event processes



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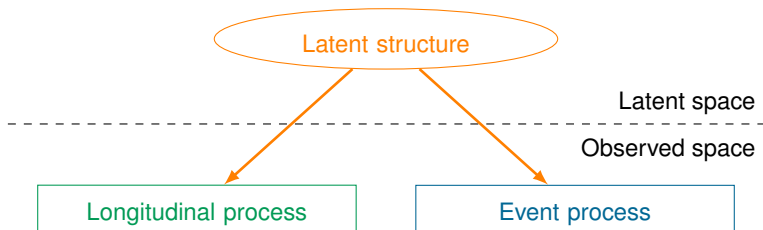


► Objectives:

- Describe the longitudinal process stopped by the event
- Explore the association between the two processes
- Predict the risk of event adjusted for the longitudinal process

Joint modelling principle

Simultaneous modelling of correlated longitudinal and event processes



▶ Latent structure:

- ▶ Function of shared random effects (shared random effect models)¹
 - Homogeneous population
 - Specification and quantification of the association between the two processes
- ▶ Latent classes (joint latent class models)²
 - Heterogeneous population
 - No assumption on the association

1. [Rizopoulos, 2012]

2. [Proust-Lima et al., 2014]

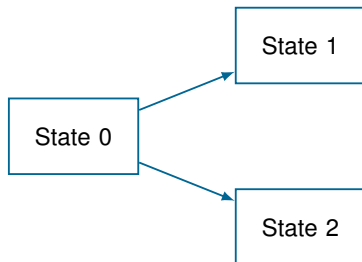
Classical joint modelling

- ▶ Classical joint models are developed for
 - ▶ A Gaussian longitudinal marker
 - ▶ Survival data with one single type of event



Extension of the classical joint models

- ▶ Many applications require joint models with
 - ▶ More complex longitudinal processes
 - ▶ More complex event processes
 - Survival data with competing events
 - Recurrent events
 - Multi-state process with possible multiple transitions



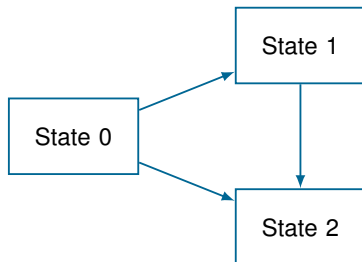
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Joint multi-state models

- ▶ In the literature, very few papers have focused on the succession of events in the joint model setting³
- ▶ Focus on
 - ▶ A joint multi-state model with shared random effects⁴
 - application: link between PSA & multiple clinical progressions in prostate cancer
 - ▶ A joint multi-state model with latent classes⁵
 - application: distinction of profiles of cognitive decline associated with risks of dementia and death in elderly people

3. [Hickey et al., 2018]

4. [Ferrer et al., 2016]

5. [Rouanet et al., 2016]

Research Article

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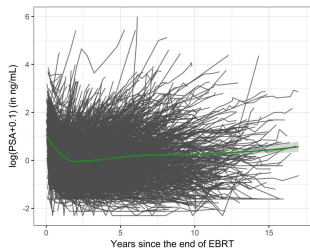
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Joint modelling of longitudinal and multi-state processes: application to clinical progressions in prostate cancer

Loïc Ferrer,^{a*†} Virginie Rondeau,^a James Dignam,^b
Tom Pickles,^c Hélène Jacqmin-Gadda^a and Cécile Proust-Lima^a

Classical modelling in prostate cancer

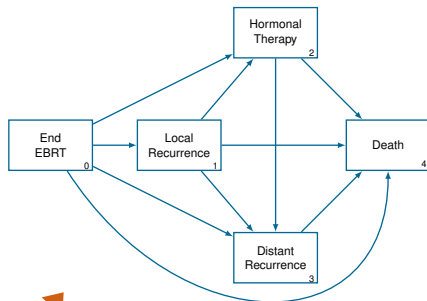
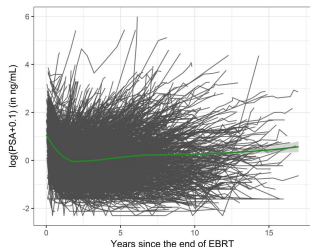
- ▶ Longitudinal PSA & clinical progression



- ▶ No distinction between the clinical progressions
- ▶ No modelling of the full disease progression

Multiple clinical progressions in cancer

► Longitudinal PSA & multiple clinical progressions



► Objectives

- Understanding the PSA evolution during the patient's follow-up
- Distinguishing & evaluating the impact of PSA dynamics and other prognostic factors on disease evolution

Notations

- ▶ Let us consider two observed processes: one longitudinal and one multi-state
- ▶ Multi-state process
 - ▶ $E_i = \{E_i(t), T_{i0} \leq t \leq C_i\}$ non-homogeneous Markovian process
 - $E_i(t)$ with values in the finite state space $S = \{0, 1, \dots, M\}$
 - T_{i0} left truncature time, C_i right censoring time
 - ▶ $T_i = (T_{i1}, \dots, T_{im_i})^\top$ the m_i observed time(s); $T_{ir} < T_{i(r+1)}, \forall r \in S$
 - ▶ $\delta_i = (\delta_{i1}, \dots, \delta_{im_i})^\top$ the vector of indicators of observed transition(s)
- ▶ Longitudinal process
 - ▶ $Y_i = (Y_{i1}, \dots, Y_{in_i})^\top$ the n_i measure(s) of marker collected at time(s) t_{i1}, \dots, t_{in_i} , with $t_{in_i} \leq T_{im_i}$

Joint multi-state model

$$\left\{ \begin{array}{l} Y_{ij} = Y_i^*(t_{ij}) + \epsilon_{ij} \\ \quad = X_i^L(t_{ij})^\top \beta + Z_i(t_{ij})^\top b_i + \epsilon_{ij} \\ \lambda_{hk}^i(t) = \lim_{dt \rightarrow 0} \frac{\Pr(E_i(t+dt) = k | E_i(t) = h)}{dt} \\ \quad = \lambda_{hk,0}(t) \exp(X_{hk,i}^E{}^\top \gamma_{hk} + W_{hk,i}(b_i, t)^\top \eta_{hk}), \text{ for } (h, k) \in S^2, \end{array} \right.$$

- ▶ $b_i \sim \mathcal{N}_q(0, B)$, $\epsilon_i = (\epsilon_{i1}, \dots, \epsilon_{in_i})^\top \sim \mathcal{N}_{n_i}(0, \sigma^2 I)$, $b_i \perp\!\!\!\perp \epsilon_i$
- ▶ $\lambda_{hk,0}(t)$ parametric baseline intensity, $X_{hk,i}^E$ prognostic factors
- ▶ $W_{hk,i}(b_i, t)$ structure of dependence, e.g.
 - ▶ $W_{hk,i}(b_i, t) = Y_i^*(t) \longrightarrow$ (true current level)
 - ▶ $W_{hk,i}(b_i, t) = \partial Y_i^*(t) / \partial t \longrightarrow$ (true current slope)
 - ▶ $W_{hk,i}(b_i, t) = (Y_i^*(t), \partial Y_i^*(t) / \partial t)^\top \longrightarrow$ (both)
 - ▶ ...

Estimation and implementation

- ▶ Maximum likelihood approach

- ▶ Using $Y_i \perp\!\!\!\perp_{b_i} E_i$
- ▶ Likelihood function

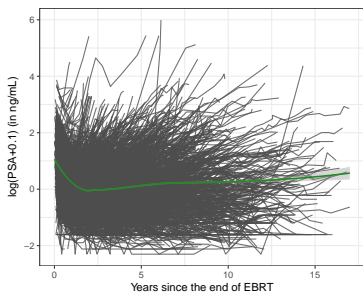
$$L(\theta) = \prod_{i=1}^N \int_{\mathbb{R}^q} f_Y(Y_i|b_i; \theta) f_E(E_i|b_i; \theta) f_b(b_i; \theta) db_i$$

- ▶ Implementation in R

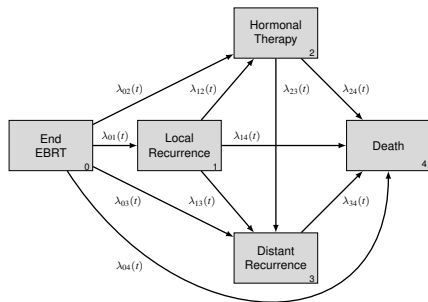
- ▶ Combination and extension of the existing R packages `JM` and `mstate`
- ▶ Codes with detailed examples available at <https://github.com/LoicFerrer/JMstateModel>
- ▶ Likelihood computed and optimised using
 - numerical integration algorithms (Gaussian quadratures: multi-step pseudo-adaptive Gauss-Hermite quadratures for the integral over random effects)
 - optimisation algorithms (EM + quasi-Newton)

Application

- ▶ 2 cohorts of men with localised prostate cancer treated by radiotherapy (N=1474)
- ▶ Longitudinal biomarker: PSA
- ▶ Multi-state representation of the clinical progressions



10, [3–21] measures per patient
(50th, [5th – 95th] %iles)



$$\Upsilon = \begin{pmatrix} 533 & 144 & 227 & 47 & 523 \\ 0 & 20 & 90 & 10 & 24 \\ 0 & 0 & 106 & 33 & 178 \\ 0 & 0 & 0 & 13 & 77 \\ 0 & 0 & 0 & 0 & 802 \end{pmatrix}$$

matrix of direct transitions

Specification of the joint multi-state model

- ▶ Model inspired from the literature with one unique type of event

$$\left\{ \begin{array}{l} Y_{ij} = Y_i^*(t_{ij}) + \epsilon_{ij} \\ = (\beta_0 + X_i^{L0 \top} \beta_{0,\text{cov}} + b_{i0}) + \\ (\beta_1 + X_i^{L1 \top} \beta_{1,\text{cov}} + b_{i1}) \times ((1 + t_{ij})^{-1.2} - 1) + \\ (\beta_2 + X_i^{L2 \top} \beta_{2,\text{cov}} + b_{i2}) \times t_{ij} + \epsilon_{ij} \\ \lambda_{hk}^i(t) = \lambda_{hk,0}(t) \exp \left(X_{hk,i}^E \top \gamma_{hk} + \left(\frac{g(Y_i^*(t))}{\partial Y_i^*(t)/\partial t} \right) \top \begin{pmatrix} \eta_{hk,\text{level}} \\ \eta_{hk,\text{slope}} \end{pmatrix} \right) \end{array} \right.$$

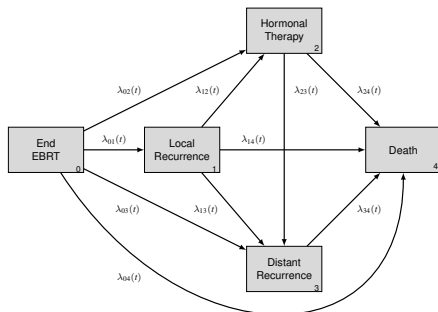
- ▶ $g(Y_i^*(t)) = \text{logit}^{-1}((Y_i^*(t) - 0.71)/0.44)$
- ▶ $\epsilon_i \sim \mathcal{N}(0, \sigma^2 I_{n_i})$
- ▶ $b_i = (b_{i0}, b_{i1}, b_{i2})^\top \sim \mathcal{N}(0, D)$, D unstructured

Results

Estimates of the association parameters between the longitudinal and multi-state processes

	Value	StdErr	<i>p</i> -value
Level : 01	3.32	0.41	< 0.001
Level : 02	4.89	0.39	< 0.001
Level : 03	2.94	0.68	< 0.001
Level : 04	-0.41	0.23	0.071
Level : 12	1.90	0.83	0.023
Level : 13	-2.30	1.32	0.081
Level : 14	-0.07	0.88	0.939
Level : 23	-0.29	1.04	0.778
Level : 24	-0.48	0.62	0.440
Level : 34	-0.02	0.57	0.974
.....			
Slope : 01	1.33	0.33	< 0.001
Slope : 02	1.60	0.24	< 0.001
Slope : 03	1.74	0.54	0.001
Slope : 04	0.59	0.35	0.088
Slope : 12	0.46	0.58	0.336
Slope : 13	3.82	1.07	< 0.001
Slope : 14	0.70	1.02	0.495
Slope : 23	0.23	0.51	0.651
Slope : 24	0.64	0.23	0.005
Slope : 34	-0.56	0.42	0.186

Multi-state representation of the clinical progressions



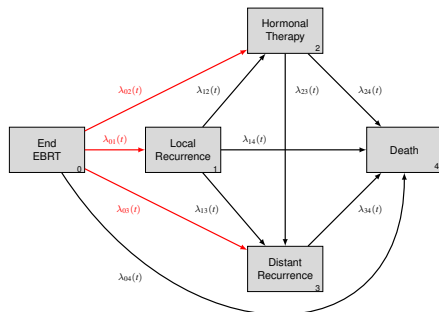
- Prognostic factors: advanced initial stage not always associated with intensities of transitions between health states after adjustment on PSA

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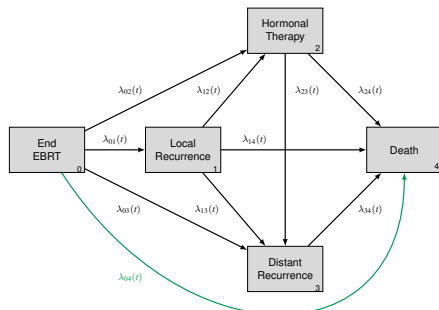
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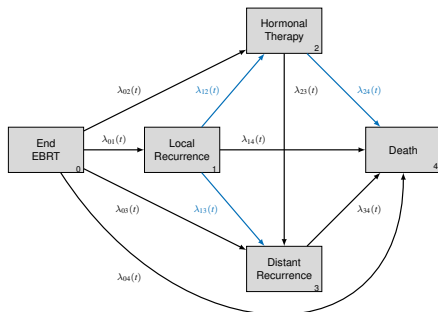
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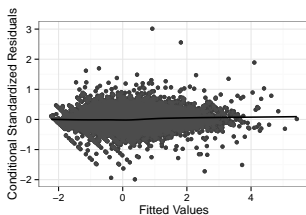


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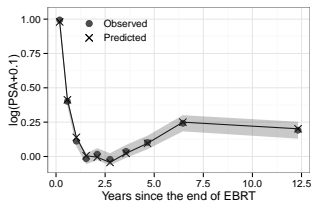
Model diagnostics

► Goodness-of-fit plots for the longitudinal process

- Conditional standardized residuals versus fitted values

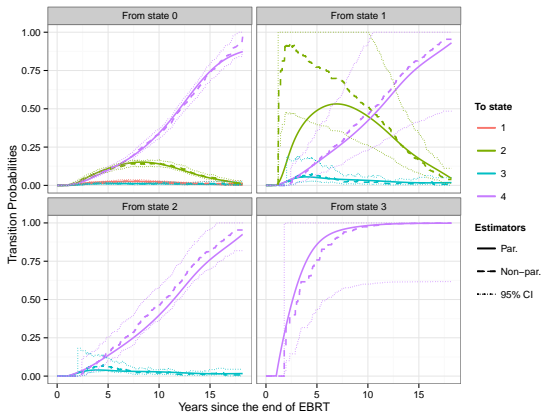


- Observed and predicted values of the biomarker



Model diagnostics (cont'd)

- ▶ Goodness-of-fit plots for the longitudinal process
- ▶ Goodness-of-fit plot for the multi-state process
 - Predicted transition probabilities from the joint multi-state model and non-parametric probability transitions



BIOMETRICS

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Joint Latent Class Model for Longitudinal Data and Interval-Censored Semi-Competing Events: Application to Dementia

Anaïs Rouanet,^{1,2,*} Pierre Joly,¹ Jean-François Dartigues,^{1,2} Cécile Proust-Lima,^{1,2} and
Hélène Jacqmin-Gadda^{1,2}

¹INSERM, Centre INSERM U1912 - Epidemiologie - Biostatistiques, F-33076 Bordeaux, France

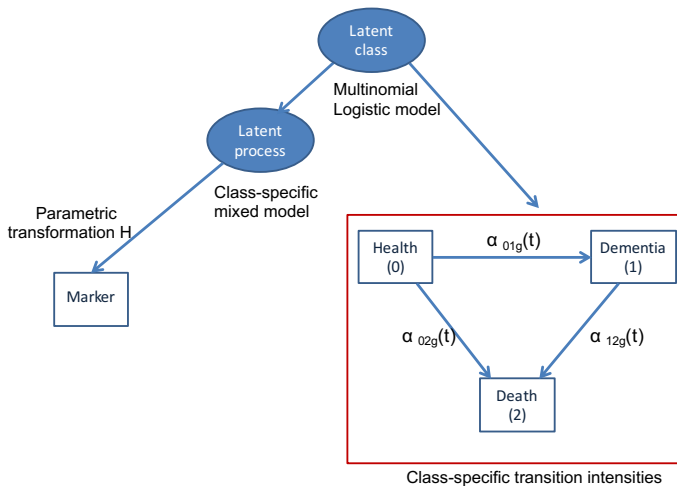
²Université de Bordeaux, ISPED, 146 rue Léo Saignat, F-33076 Bordeaux, France

* *email*: anais.rouanet@isped.u-bordeaux2.fr

Application to dementia

- ▶ **Aim:** To distinguish different profiles of cognitive decline associated with risks of dementia and death
- ▶ **Data:** Paquid Cohort (French prospective cohort: Normal and pathological brain ageing)
 - ▶ 3777 subjects from Dordogne and Gironde, aged 65 and over
 - ▶ Visits every 2/3 years during 25 years
 - ▶ Study of Isaacs Set Test [0-40], verbal fluency
- ▶ **Methodological challenges**
 - ▶ Heterogeneity in cognitive decline
 - ▶ Correlation between cognitive decline and occurrence of dementia
 - ▶ Competing risk of death
 - ▶ Time-to-dementia onset interval-censored

Joint latent class illness-death model



Joint model formulation

- ▶ Membership probability: $p_{ig} = P(c_i = g | X_{pi})$
- ▶ Latent process Λ_i , given the class g :

$$\begin{aligned}\Lambda_i(t_{ij} | c_i = g) &= f_1(X_{ij}; \beta_g) + f_2(Z_{ij}; \beta_g) \mathbf{u}_{ig} \\ &= X_{ij}^T \beta_g + Z_{ij}^T \mathbf{u}_{ig}\end{aligned}$$

f_1, f_2 : (possibly nonlinear) functions of time, covariates

β_g : class-specific parameters

$\mathbf{u}_{ig} \sim \mathcal{N}(0, \sigma_g^2 B)$

- ▶ Transformed gaussian marker \tilde{Y} :

$$\tilde{Y}_{ij} = H(Y_{ij}; \eta) = \Lambda_i(t_{ij}) + \epsilon_{ij} \text{ with } \epsilon_{ij} \sim \mathcal{N}(0, \sigma_e^2)$$

$H(\cdot; \eta)$: Parametric transformation

- ▶ Transition intensity from state k to state l for subject i in class g :

$$\alpha_{klg}(t) = \alpha_{klg}^0(t) e^{X_{ei}^T \gamma_{klg}}$$

α_{klg}^0 : class-specific baseline intensity

γ_{klg} : class-specific regression parameters

Estimation

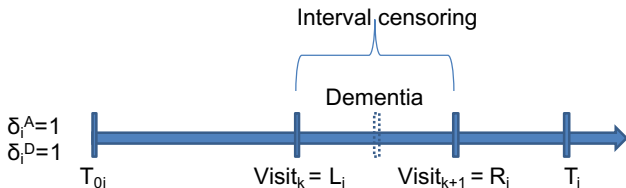
▶ Maximum likelihood approach

▶ Using $Y_i \perp\!\!\!\perp_g D_i$

▶ Log-likelihood function

$$\mathcal{L}(\theta_G) = \sum_{i=1}^N \log \left[\sum_{g=1}^G p_{ig} f(Y_i | c_i = g; \theta_G) P(D_i | c_i = g; \theta_G) \right]$$

$$- \sum_{i=1}^N \log \left[\sum_{g=1}^G p_{ig} e^{-A_{01ig}(T_{0i}; \theta_G) - A_{02ig}(T_{0i}; \theta_G)} \right]$$



- ▶ Marquardt algorithm for a fixed number of latent classes G
- ▶ G chosen by Bayesian Information Criterion (BIC) minimisation

Model specification

- ▶ Mixed model, given latent class g :

$$\Lambda_i(t) = \beta_{0g} + u_{ig}^{(0)} + [\beta_{1g} - \beta_{2g} + u_{ig}^{(1)}] (t - \tau_g) + \beta_3 Educ_i + (\beta_4 - \beta_5) Educ_i (t - \tau_g) + \beta_6 Sex_i \quad \text{if } t \leq \tau_g$$

$$\Lambda_i(t) = \beta_{0g} + u_{ig}^{(0)} + [\beta_{1g} + \beta_{2g} + u_{ig}^{(2)}] (t - \tau_g) + \beta_3 Educ_i + (\beta_4 + \beta_5) Educ_i (t - \tau_g) + \beta_6 Sex_i \quad \text{if } t \geq \tau_g$$

$$u_{ig} = (u_{ig}^{(0)}, u_{ig}^{(1)}, u_{ig}^{(2)})^\top \sim \mathcal{N}(0, \sigma_g^2 B), \epsilon_{ij} \sim \mathcal{N}(0, \sigma_\epsilon^2)$$

$$\tilde{Y}_{ij} = H(Y_{ij}; \eta) = \Lambda_i(t_{ij}) + \epsilon_{ij} \text{ with } H : \text{Beta cumulative distribution function}$$

- ▶ Transition intensities from states $k \rightarrow l$ of the multi-state model:

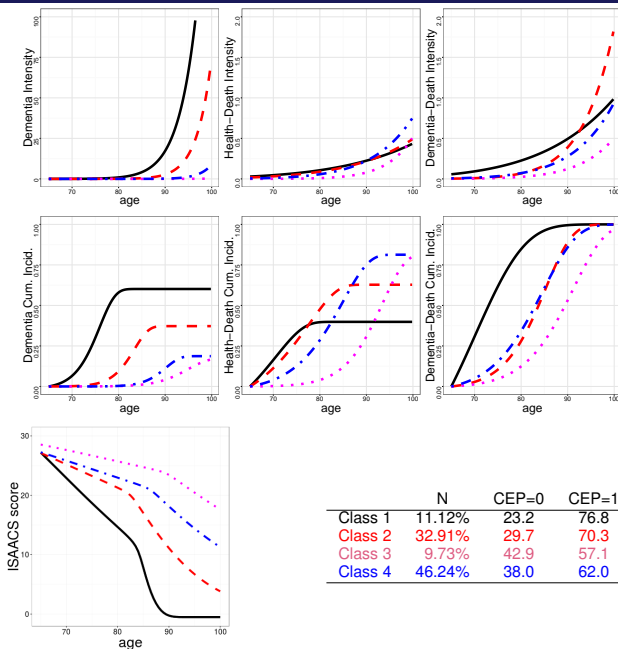
$$\alpha_{klig}(t) = \alpha_{klg}^0(t) e^{\gamma_{kls} Sex_i + \gamma_{kle} Educ_i}$$

Model choice

	BIC	
	Markovian	Semi-markovian
G=1	106901	107055
G=2	106270	106356
G=3	106081	106177
G=4	106005	106091
G=5	106027	106107

Mortality among subjects with dementia depends more on age than on the duration of dementia.

The minimum value of BIC with $G = 4$ classes.



	N	CEP=0	CEP=1	men	women
Class 1	11.12%	23.2	76.8	40.1	59.9
Class 2	32.91%	29.7	70.3	40.7	59.3
Class 3	9.73%	42.9	57.1	42.0	58.0
Class 4	46.24%	38.0	62.0	43.8	56.2

Discrimination of the posterior classification

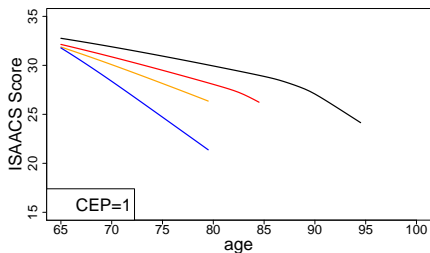
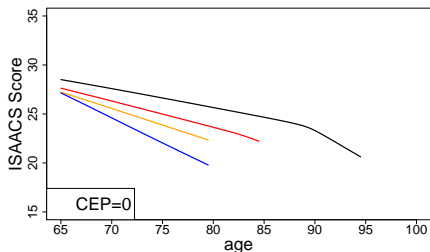
Classification according to:

$$\hat{\pi}_{ig}^{Y,D} = P(c_i = g | Y_i, D_i, \hat{\theta}_G) = \frac{f(Y_i | c_i = g; \hat{\theta}_G) P(D_i | c_i = g; \hat{\theta}_G) P(c_i = g; \hat{\theta}_G)}{\sum_l^G f(Y_i | c_i = l; \hat{\theta}_G) P(D_i | c_i = l; \hat{\theta}_G) P(c_i = l; \hat{\theta}_G)}$$

Class	1	2	3	4
1	71.36	21.92	0.18	6.54
2	12.83	61.54	1.24	24.39
3	0.01	0.47	79.11	20.40
4	0.96	19.11	12.96	66.97

Table 1 – Mean probabilities to belong to each class according to the posterior classification.

Estimated trajectories according to dementia onset & death



Man with a low/high level of education
in 4 different cases:

man alive and dementia-free at 95
 $E(Y(t)|T_i^A > 95, T_i^D > 95, \delta_i^A = 0, \delta_i^D = 0; \hat{\theta}_G)$

man alive and dementia-free at 85

man dead dementia-free at 80

man alive with dementia at 80

Discussion

- ▶ Joint models extended to several kinds of multivariate survival data
 - ▶ See Hickey G.L. et al. (2018) for a full state-of-the-art
- ▶ Ferrer et al. (2016) developed a joint multi-state model with shared random effects to
 - ▶ Model the disease evolution in its whole
 - ▶ Using an easy-to-use function implemented in R
- ▶ Rouanet et al. (2016) developed a joint latent class illness-death model accounting for
 - ▶ Heterogeneity in the data
 - ▶ Competing risk of death
 - ▶ Interval censoring
- ▶ Model diagnostics
 - ▶ Study of the residuals for validating the model assumptions
 - ▶ Graphical comparison of the observations and predictions of the model as goodness-of-fit tool

Discussion (cont'd)

- ▶ Goodness-of-fit assessment in joint multi-state models using a score test for the inclusion of a Gaussian frailty term
 - ▶ for shared random effect models
 - ▶ for joint latent class models
- ▶ Useful to validate the model assumptions and check its goodness-of-fit
 - ▶ Markovian assumption
 - ▶ Incomplete adjustment on covariates
 - ▶ Presence of non-linear covariate effect
 - ▶ ~~Violation of the proportional intensities assumption~~
- ▶ Available at <http://github.com/LoicFerrer/JMstateModel/> for shared random effect models
- ▶ Submitted for publication in a few days

References

- [1.] Rizopoulos D. (2012). *Joint models for longitudinal and time-to-event data: With applications in R*. Chapman and Hall/CRC.
- [2.] Proust-Lima C., Séne M., Taylor J.M. & Jacqmin-Gadda H. (2014). Joint latent class models for longitudinal and time-to-event data: A review. *Statistical Methods in Medical Research*, 23(1):74–90.
- [3.] Hickey G.L., Philipson P., Jorgensen A. & Kolamunnage-Dona R. (2018). Joint models of longitudinal and time-to-event data with more than one event time outcome: a review. *The International Journal of Biostatistics*, 14(1).
- [4.] Ferrer L., Rondeau V., Dignam J., Pickles T., Jacqmin-Gadda, H. & Proust-Lima C. (2016). Joint modelling of longitudinal and multi-state processes: application to clinical progressions in prostate cancer. *Statistics in Medicine*, 35(22):3933–3948.
- [5.] Rouanet A., Joly P., Dartigues J-F., Proust-Lima C. & Jacqmin-Gadda H. (2016). Joint Latent Class Model for Longitudinal Data and Interval-Censored Semi-Competing Events: Application to Dementia. *Biometrics*, 72(4):1123–1135.

Numerical approximation of the integral over the random effects

- ▶ Likelihood function

$$L(\theta) = \prod_{i=1}^N \int_{\mathbb{R}^q} f_Y(Y_i|b_i; \theta) f_E(E_i|b_i; \theta) f_b(b_i; \theta) db_i$$

- ▶ **Adaptive Gauss-Hermite rule** → Centering and rescaling the integral around its modal value at each step of the optimisation algorithm

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- ▶ **Adaptive Gauss-Hermite rule** → Centering and rescaling the integral around its modal value at each step of the optimisation algorithm
 1. **Pseudo-adaptive GH rule**: based on the posterior distribution of the random effects from the LMM

$$\tilde{b}_i = \arg \max_b \{\log f(Y_i, b; \tilde{\theta}_Y)\}$$

and their associated covariance matrix

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 1. **Pseudo-adaptive GH rule**: based on the posterior distribution of the random effects from the LMM

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2. **Multi-step pseudo-adaptive GH rule**: based on the posterior distribution of the random effects from the JM

$$\check{b}_i = \arg \max_b \{ \log f(E_i, Y_i, b; \check{\theta}) \}$$

and their associated covariance matrix

Joint multi-state model – Implementation in R (1/4)

► Example of R code

$$\left\{ \begin{array}{l} Y_{ij} = Y_i^*(t_{ij}) + \epsilon_{ij} \\ = (\beta_0 + \beta_{0,X}X_i + b_{i0}) + (\beta_1 + \beta_{1,X}X_i + b_{i1}) \times t_{ij} + \epsilon_{ij} \\ \lambda_{hk}^i(t|b_i) = \lambda_{hk,0}(t) \exp(\gamma_{hk}X_i + \eta_{hk,\text{level}}Y_i^*(t) + \eta_{hk,\text{slope}}\partial Y_i^*(t)/\partial t) \end{array} \right.$$

where the multi-state process included three states $((h, k) \in \{0, 1, 2\}^2)$ and three transitions $(0 \rightarrow 1, 0 \rightarrow 2, 1 \rightarrow 2)$,

the log-baseline intensities are a linear combination of cubic-splines

```
library(mstate)
library(JM)
source("JMstateModel.R")
load("data.RData")

# Initialisation of the longitudinal sub-part
lmeFit <- lme(fixed = Y ~ 1 + X + time + X:time,
             data = data_long,
             random = ~ (1 + time) | id,
             method = "REML",
             control = list(opt = "optim"))
```


Joint multi-state model – Implementation in R (2/4)

```
# Data preparation to the multi-state framework
tmat <- matrix(NA, 3, 3)
tmat[1, 2:3] <- 1:2
tmat[2, 3] <- 3
dimnames(tmat) <- list(from = c("State_0", "State_1", "State_2"),
                       to = c("State_0", "State_1", "State_2"))

covs <- "X"

data_mstate <-
  msprep(time = c(NA, "time_of_State_1", "time_of_State_2"),
         status = c(NA, "State_1", "State_2"),
         data = data_surv,
         trans = tmat,
         keep = covs,
         id = "id")

data_mstate <- expand.covs(data_mstate, covs,
                          append = TRUE, longnames = FALSE)
```

Joint multi-state model – Implementation in R (3/4)

```
# Initialisation of the multi-state sub-part
coxFit <- coxph(Surv(Tstart, Tstop, status) ~
               X.1 + X.2 + X.3 + strata(trans),
               data = data_mstate,
               method = "breslow", x = TRUE, model = TRUE)
```

Joint multi-state model – Implementation in R (4/4)

```
dForm <- list(fixed = ~ 1 + X,
             indFixed = c(3, 4),
             random = ~ 1,
             indRandom = 2)

# Joint multi-state model
jointFit_both <-
  JMstateModel(lmeObject = lmeFit,
              survObject = coxFit,
              timeVar = "time",
              parameterization = "both",
              method = "spline-PH-aGH",
              interFact = list(value = ~ strata(trans) - 1,
                              slope = ~ strata(trans) - 1,
                              data = data_mstate),
              derivForm = dForm,
              Mstate = TRUE,
              data.Mstate = data_mstate,
              ID.Mstate = "id",
              control = list(GHk = 9, lng.in.kn = 3))
```

Joint multi-state model with an additional frailty

► Model formulation

$$\left\{ \begin{array}{l} Y_{ij} = X_i^L(t_{ij})^\top \beta + Z_i(t_{ij})^\top b_i + \epsilon_{ij} \\ \lambda_{hk}^i(t) = \lambda_{hk,0}(t) \exp \left(X_{hk,i}^E{}^\top \gamma_{hk} + W_{hk,i}(b_i, t)^\top \eta_{hk} + v_i \right), \text{ for } (h, k) \in S^2 \end{array} \right.$$

- $v_i \sim \mathcal{N}(0, \sigma_v^2)$ the frailty term
- $v_i \perp\!\!\!\perp b_i, v_i \perp\!\!\!\perp \epsilon_i$

► Likelihood function $L(\sigma_v^2, \theta)$

$$L(\sigma_v^2, \theta) = \prod_{i=1}^N \int_{\mathbb{R}^{q_b}} f_Y(Y_i | b_i; \theta) f_b(b_i; \theta) \int_{\mathbb{R}} f_E(E_i | b_i, v_i; \theta) f_v(v_i; \theta) \, dv_i \, db_i$$

Score test: score statistic and its variance

$$H_0 : \sigma_v^2 = 0 \quad \text{vs} \quad H_1 : \sigma_v^2 > 0$$

- ▶ Score statistic $U(0, \theta) = U(\sigma_v^2 = 0, \theta)$

$$U(0, \theta) = \sum_{i=1}^N U_i(0, \theta) = \sum_{i=1}^N \frac{\partial \log L_i(\sigma_v^2, \theta)}{\partial \sigma_v^2} \Bigg|_{\sigma_v^2=0}$$

- ▶ Analytic expression
- ▶ Asymptotic variance corrected for the estimation of the nuisance parameters

$$\text{Var} \{U(0, \theta)\} = \left(I_{\sigma_v^2 \sigma_v^2} - I_{\sigma_v^2 \theta} I_{\theta \theta}^{-1} I_{\theta \sigma_v^2} \right) \Bigg|_{\sigma_v^2=0}$$

- ▶ Forward finite difference method

Score test: test statistic

► Test statistic

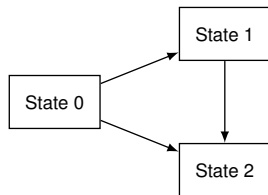
$$T = \begin{cases} 0 & \text{if } U(0, \hat{\theta}_0) \leq 0 \\ \frac{\{U(0, \hat{\theta}_0)\}^2}{\text{Var}\{U(0, \hat{\theta}_0)\}} & \text{if } U(0, \hat{\theta}_0) > 0 \end{cases}$$

- $\hat{\theta}_0$ the model parameters estimated under the null hypothesis
- T follows asymptotically a mixture of chi-square distributions

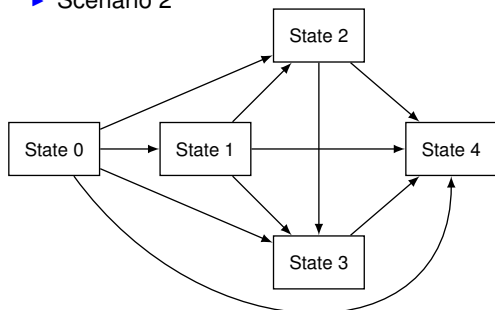
$$T \sim \frac{1}{2}\chi_0^2 + \frac{1}{2}\chi_1^2$$

Assessment by simulation study

► Scenario 1



► Scenario 2



► For each subject $i = 1, \dots, N$ of the 500 replicates,

$$\left\{ \begin{array}{l} Y_i(t) = Y_i^*(t) + \epsilon_i(t) \\ \quad = (\beta_0 + \beta_{0,X}X_i + b_{i0}) + (\beta_1 + \beta_{1,X}X_i + b_{i1}) \times t + \epsilon_i(t) \\ \lambda_{hk}^i(t) = \lambda_{hk,0}(t) \exp(\gamma_{hk}X_i + \eta_{hk,0}Y_i^*(t) + \eta_{hk,1}\partial Y_i^*(t)/\partial t + v_i) \end{array} \right.$$

Simulation study – results

- ▶ Empirical type-I error rate (nominal level of 5%)

	Scenario 1 (3 states) $\sigma_v^2 = 0$ ($\bar{M} = 0.70$)	Scenario 2 (5 states) $\sigma_v^2 = 0$ ($\bar{M} = 2.84$)
$N = 500$	0.008	0.028
$N = 1000$	0.010	0.054
$N = 1500$	0.020	0.060

- ▶ Empirical statistical power (nominal level of 5%)

	Scenario 1 (3 states)		Scenario 2 (5 states)	
	$\sigma_v^2 = 0.5$ ($\bar{M} = 0.74$)	$\sigma_v^2 = 1$ ($\bar{M} = 0.75$)	$\sigma_v^2 = 0.5$ ($\bar{M} = 2.73$)	$\sigma_v^2 = 1$ ($\bar{M} = 2.65$)
$N = 500$	0.278	0.438	0.884	0.990
$N = 1000$	0.568	0.850	0.998	1.000
$N = 1500$	0.846	0.970	1.000	1.000

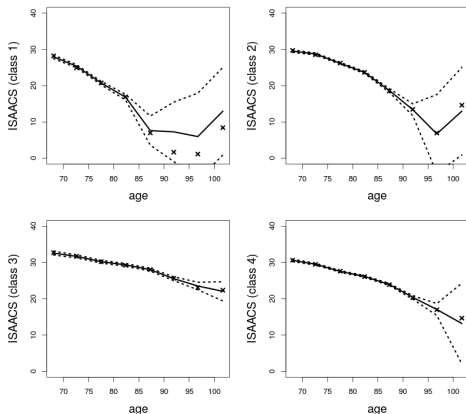
\bar{M} : average number of observed direct transitions per subject

Score statistic

- Score statistic $U(\mathbf{0}, \theta)$

$$\begin{aligned}
 U(\mathbf{0}, \theta) &= \sum_{i=1}^N \frac{1}{2L_i(\mathbf{0}, \theta)} \times \\
 &\quad \int_{\mathbb{R}^{q_b}} f_Y(Y_i | \mathbf{b}_i; \theta) f_E(E_i | \mathbf{b}_i; \mathbf{0}, \theta) f_b(\mathbf{b}_i; \theta) \times \\
 &\quad \left\{ \left[\sum_{r=0}^{m_i-1} (\delta_{i(r+1)} + \Lambda_{E_i(T_{ir}), E_i(T_{ir})}^i(T_{ir}, T_{i(r+1)} | \mathbf{b}_i; \mathbf{0}, \theta)) \right]^2 + \right. \\
 &\quad \left. \sum_{r=0}^{m_i-1} (\Lambda_{E_i(T_{ir}), E_i(T_{ir})}^i(T_{ir}, T_{i(r+1)} | \mathbf{b}_i; \mathbf{0}, \theta)) \right\} d\mathbf{b}_i \\
 &= \sum_{i=1}^N \frac{1}{2} \int_{\mathbb{R}^{q_b}} f_b(\mathbf{b}_i | Y_i, E_i; \mathbf{0}, \theta) \{ \dots \} d\mathbf{b}_i
 \end{aligned}$$

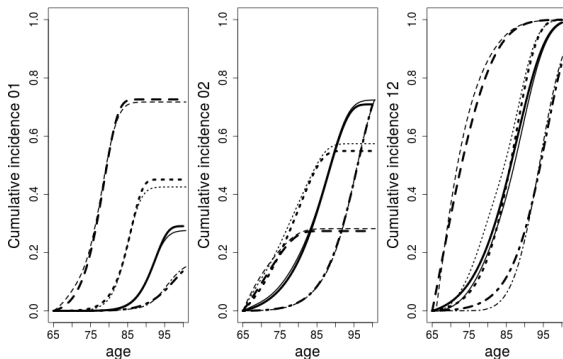
Goodness-of-fit of the longitudinal predictions



$$\hat{\mu}_{gl}^o = \frac{\sum_{(i,j)|\tau_l < t_{ij} < \tau_{l+1}} Y_{ij} P(c_i = g | Y_i, D_i; \hat{\theta}_G)}{\sum_{(i,j)|\tau_l < t_{ij} < \tau_{l+1}} P(c_i = g | Y_i, D_i; \hat{\theta}_G)}$$

$$\hat{\mu}_{gl}^u = \frac{\sum_{(i,j)|\tau_l < t_{ij} < \tau_{l+1}} E(Y_{ij} | c_i = g, \hat{u}_{ig}; \hat{\theta}_G) P(c_i = g | Y_i, D_i; \hat{\theta}_G)}{\sum_{(i,j)|\tau_l < t_{ij} < \tau_{l+1}} P(c_i = g | Y_i, D_i; \hat{\theta}_G)}, \text{ with } \hat{u}_{ig} = E(u_{ig} | Y_i, \hat{\theta}_G)$$

Goodness-of-fit of the illness-death predictions



Class-specific cumulative incidences, marginal on covariates:
 Joint latent class illness-death model **vs.** Semi-parametric illness-death model with
 baseline transition intensities modeled by M-splines.