



An overview and empirical appraisal of recent methodological developments for the dynamic prediction of survival outcomes

Mirko Signorelli

 [mirkosignorelli.github.io](https://github.com/mirkosignorelli)
 [in/signorelli](https://www.linkedin.com/in/signorelli)

Mathematical Institute
Leiden University

Joint work with Sophie Retif

October 8, 2025
65th ISI World Statistics Congress



Universiteit
Leiden

Preprint:  [arXiv:2403.14336v2](https://arxiv.org/abs/2403.14336v2)

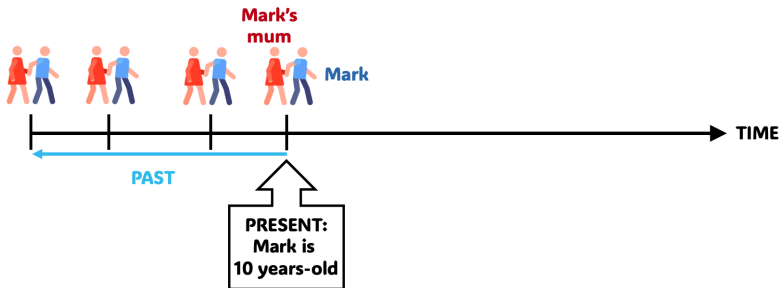
The dynamic prediction problem

Dynamic prediction with numerous longitudinal covariates

Benchmarking study

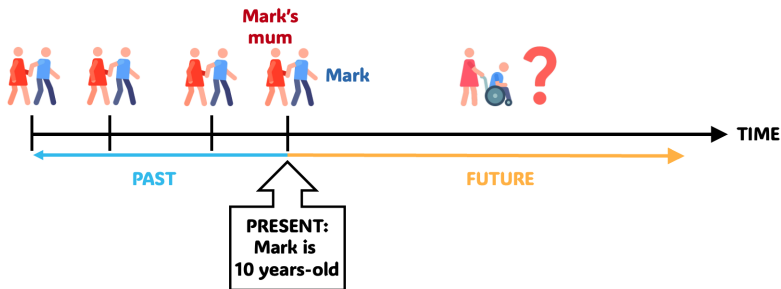
Appendix

Dynamic prediction 101



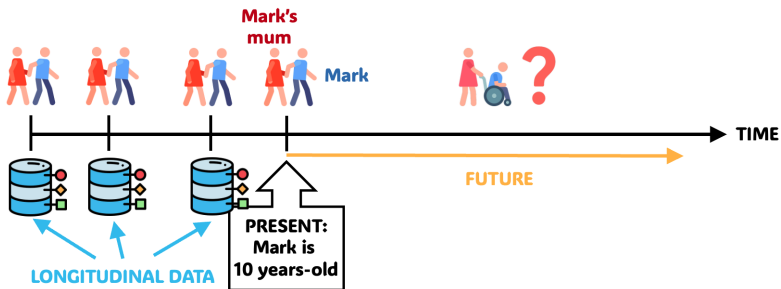
- ▶ Mark suffers from Duchenne muscular dystrophy (DMD)
- ▶ Usually: **loss of ambulation** during adolescence

Dynamic prediction 101

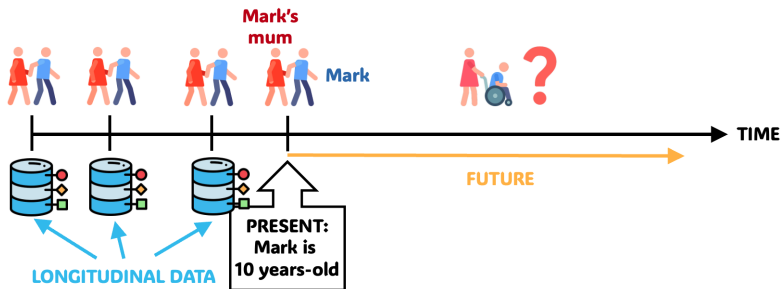


- Mum: $P(\text{wheelchair within 2 years})?$

Dynamic prediction 101



Dynamic prediction 101



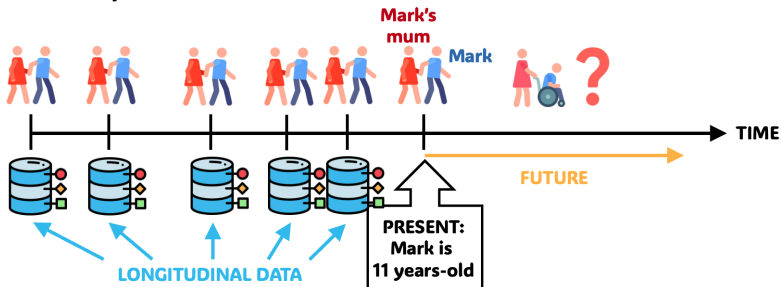
Dynamic prediction goal # 1:

- ▶ use available longitudinal data to predict conditional survival probability

$$S(t|10, Y = \{y_1, y_2, y_3\}) = P(T > t | T > 10, Y = \{y_1, y_2, y_3\}), t > 10$$

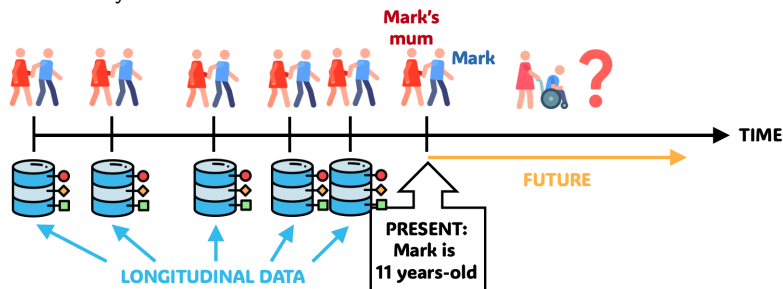
Dynamic prediction 101

► One year later:



Dynamic prediction 101

- ▶ One year later:



Dynamic prediction goal # 2:

- ▶ if still at risk at $t = 11$, exploit new data to update predictions

$$S(t|11, Y = \{y_1, y_2, y_3, y_4, y_5\}), t > 11$$

The problem

► Traditional methods:

1. **landmarking** with Last Observation Carried Forward (LOCF): **discards longitudinal information** + no measurement error correction (important for biomarkers)
2. **joint models**: very **computationally-intensive** (hours / days) + many **estimation errors**. Can't usually be estimated with more than 5-10 longitudinal predictors!

The problem

- ▶ Traditional methods:
 1. **landmarking** with Last Observation Carried Forward (LOCF): **discards longitudinal information** + no measurement error correction (important for biomarkers)
 2. **joint models**: very **computationally-intensive** (hours / days) + many **estimation errors**. Can't usually be estimated with more than 5-10 longitudinal predictors!
- ▶ Nowadays, longitudinal studies can comprise **tens, hundreds, or even thousands** longitudinal predictors ("biomarkers")
- ▶ How to do **dynamic prediction with "many" longitudinal predictors?**

The dynamic prediction problem

Dynamic prediction with numerous longitudinal covariates

Benchmarking study

Appendix

New dynamic prediction methods (2019-2023)

Several solutions proposed over the last 6 years:

1. Multivariate Functional Principal Component Cox model ([MFPC Cox](#), Li & Luo (2019))
2. Penalized Regression Calibration ([pencal](#), Signorelli et al. (2021))
3. Functional Random Survival Forest ([FunRSF](#), Lin et al. (2021))
4. Dynamic Random Survival Forest ([DynForest](#), Devaux et al. (2023))

Modelling approach

Modelling steps:

1. model trajectories of longitudinal predictors over $[0, \ell]^1$
2. obtain subject-specific summaries of the longitudinal variables
3. use baseline covariates and summaries of longitudinal predictors to predict $S(t|\ell)$, $t \geq \ell$

¹Technically, often data gathered after ℓ are also included. This is **problematic** for multi-step methods: selection bias after $\ell \Rightarrow$ lower predictive performance (Gomon et al., 2024) + use future to predict the future! ☹

Methods overview

- In a nutshell:

Method	Approach used to model...	
	longitudinal covariates	survival outcome
MFPCox	multiv. functional PCA	Cox model
FunRSF	multiv. functional PCA	random survival forest
pencal	linear mixed models	penalized Cox model
DynForest	linear mixed models	random survival forest

Software

► Software implementations:

Method	Software	Details
MFPCCox	☹️	
FunRSF	☹️	
pencal	😊	R package on CRAN
DynForest	😊	R package on CRAN

► Software articles:

- `pencal`: Signorelli (2024)
- `DynForest`: Devaux et al. (2024+)

The dynamic prediction problem

Dynamic prediction with numerous longitudinal covariates

Benchmarking study

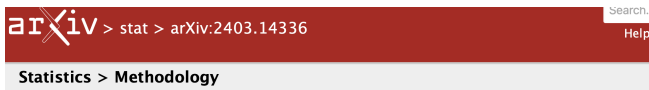
Appendix

Motivation

- ▶ Methods **proposed quite recently**: between 2019 and 2023 \Rightarrow very little knowledge about their (relative) predictive performance, advantages and limitations
- ▶ Let's **compare** them using **real-world data**!

Motivation

- ▶ Methods **proposed quite recently**: between 2019 and 2023 \Rightarrow very little knowledge about their (relative) predictive performance, advantages and limitations
- ▶ Let's **compare** them using **real-world data**!
- ▶ Why benchmarking instead of MC simulations?
 1. Interest in how these methods perform on real datasets with complex and messy data
 2. Difficult to simulate realistic complex longitudinal + survival data
 3. Choices in simulation strategy could unfairly favour some methods over others



[Submitted on 21 Mar 2024 (v1), last revised 17 Apr 2025 (this version, v2)]




Benchmarking multi-step methods for the dynamic prediction of survival with numerous longitudinal predictors

Mirko Signorelli, Sophie Retif

In recent years, the growing availability of biomedical datasets featuring numerous longitudinal covariates has motivated the development of several multi-step methods for the dynamic prediction of time-to-event ("survival") outcomes. These methods employ either mixed-effects models or multivariate functional principal component analysis to model and summarize the longitudinal covariates' evolution over time. Then, they use Cox models or random survival forests to predict survival probabilities, using as covariates both baseline variables and the summaries of the longitudinal variables obtained in the previous modelling step.

Longitudinal studies




- ▶ We considered data from three longitudinal studies:

ROSMAP  <ul style="list-style-type: none">▪ Event: Alzheimer's Disease diagnosis▪ $n = 3293$▪ 5 baseline covariates▪ 30 longitudinal covariates▪ Follow-up: [1, 29] years	ADNI  <ul style="list-style-type: none">▪ Event: diagnosis of dementia▪ $n = 1643$▪ 5 baseline covariates▪ 21 longitudinal covariates▪ Follow-up: [0, 15.5] years	PBC2  <ul style="list-style-type: none">▪ Event: death (primary biliary cirrhosis trial)▪ $n = 312$▪ 3 baseline covariates▪ 8 longitudinal covariates▪ Follow-up: [0, 14] years
---	---	---

- ▶ Methods included: MFPCox, pencal, FunRSF, DynForest + static Cox + LOCF landmarking

Longitudinal studies

- ▶ We considered data from three longitudinal studies:

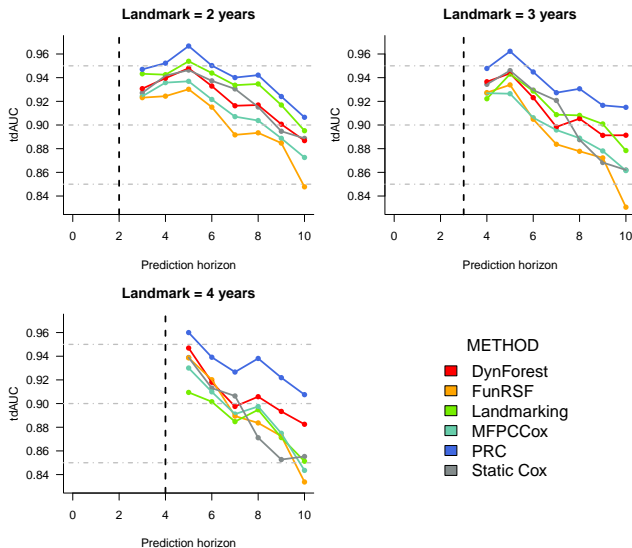
ROSMAP  <ul style="list-style-type: none">▪ Event: Alzheimer's Disease diagnosis▪ $n = 3293$▪ 5 baseline covariates▪ 30 longitudinal covariates▪ Follow-up: [1, 29] years	ADNI  <ul style="list-style-type: none">▪ Event: diagnosis of dementia▪ $n = 1643$▪ 5 baseline covariates▪ 21 longitudinal covariates▪ Follow-up: [0, 15.5] years	PBC2  <ul style="list-style-type: none">▪ Event: death (primary biliary cirrhosis trial)▪ $n = 312$▪ 3 baseline covariates▪ 8 longitudinal covariates▪ Follow-up: [0, 14] years
---	---	---

- ▶ Methods included: MFPCox, pencal, FunRSF, DynForest + static Cox + LOCF landmarking
- ▶ Performance evaluated at multiple landmark times
- ▶ Performance measures: C index, tdAUC, Brier score
- ▶ 10-fold cross-validation, repeated 10 times

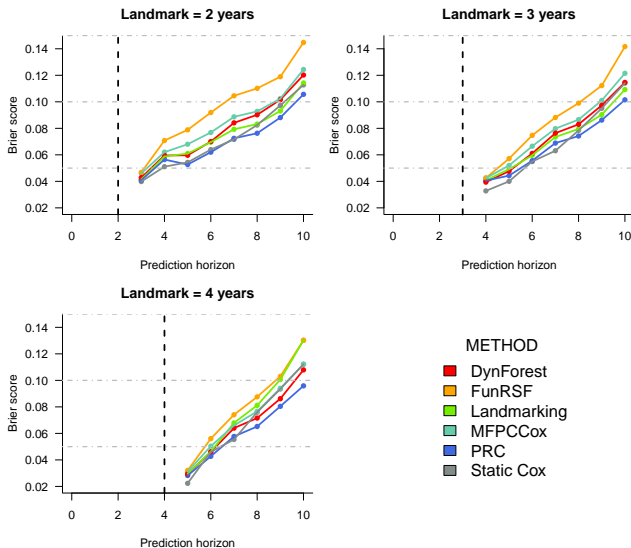
ADNI dataset: C index (higher = better)

Method	Landmark		
	2	3	4
Static Cox	0.901 (0.002)	0.885 (0.006)	0.856 (0.008)
Landmarking	0.906 (0.001)	0.89 (0.004)	0.855 (0.009)
MFPCox	0.889 (0.003)	0.872 (0.009)	0.859 (0.008)
pencal	0.913 (0.001)	0.908 (0.003)	0.904 (0.003)
FunRSF	0.873 (0.004)	0.858 (0.011)	0.845 (0.012)
DynForest	0.891 (0.003)	0.883 (0.005)	0.871 (0.011)

ADNI dataset: time-dependent AUC (higher = better)



ADNI dataset: Brier score (higher = worse)



ADNI dataset: computing time (higher = worse)

- Average computing time per CV fold (in **minutes**):

Method	Landmark			Average
	2	3	4	
Static Cox	0.009	0.007	0.006	0.007
Landmarking	0.010	0.007	0.006	0.008
MFPCox	0.080	0.046	0.046	0.057
pencal	0.776	0.482	0.453	0.571
FunRSF	0.240	0.122	0.125	0.163
DynForest	12.501	9.077	8.099	9.892
<i>n</i> at risk	1226	954	721	

Results overview

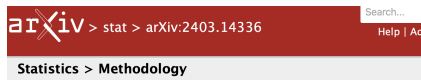
- ▶ Similar results across the 3 datasets
- ▶ `pencal`, `landmarking`, `DynForest` > `MFPCox`, `static Cox`, `FunRSF`
 - ▶ Methods that use LMMs > methods using MFPCA
 - ▶ Conditionally on method used to model longitudinal predictors (`MFPCA` / LMMs), methods that use Cox model > methods that use RSF
- ▶ Relative performance of `landmarking` and `static Cox` worsens with higher landmark & horizon times
- ▶ `DynForest` estimation particularly slow (due to re-estimation of LMMs in each node)
- ▶ More details: [arXiv:2403.14336](https://arxiv.org/abs/2403.14336)

Limitations

- ▶ MFPCA-based methods: regular measurement grid required for MFPCA estimation → unrealistic & unflexible
- ▶ LMM-based methods: only LMMs. Using GLMMs would allow for more modelling flexibility
- ▶ Methods using RSF: need to choose value of multiple tuning parameters
- ▶ Dealing with more complex survival outcomes:
 1. Competing risks: only in DynForest
 2. Interval censoring: none of the methods
- ▶ Software
 1. MFPCox, FunRSF: **no software implementation** 😞
 2. pencial → pencial, DynForest → DynForest

The End

► Preprint:  arXiv:2403.14336v2



[Submitted on 21 Mar 2024 (v1), last revised 17 Apr 2025 (this version, v2)]

Benchmarking multi-step methods for the dynamic prediction of survival with numerous longitudinal predictors

Mirko Signorelli, Sophie Retif

In recent years, the growing availability of biomedical datasets featuring numerous longitudinal covariates has motivated the development of several multi-step methods for the dynamic prediction of time-to-event ("survival") outcomes. These methods employ either mixed-effects models or multivariate functional principal component analysis to model and summarize the longitudinal covariates' evolution over time. Then, they use Cox models or random survival forests to predict survival

🏠: [mirkosignorelli.github.io](https://github.com/mirkosignorelli)
✉: m.signorelli@math.leidenuniv.nl
in: [in/signorelli](https://www.linkedin.com/in/signorelli)

Preprint:  arXiv:2403.14336v2

References I

- Devaux, A., Helmer, C., Genuer, R., & Proust-Lima, C. (2023). Random survival forests with multivariate longitudinal endogenous covariates. *Statistical Methods in Medical Research*, 32(12), 2331–2346.
- Devaux, A., Proust-Lima, C., & Genuer, R. (2024+). *Random Forests for time-fixed and time-dependent predictors: The DynForest R package*. arXiv:2302.02670.
- Gomon, D., Putter, H., Fiocco, M., & Signorelli, M. (2024). Dynamic prediction of survival using multivariate functional principal component analysis: A strict landmarking approach. *Statistical Methods in Medical Research*, 33(2), 256–272.
- Li, K., & Luo, S. (2019). Dynamic prediction of Alzheimer's disease progression using features of multiple longitudinal outcomes and time-to-event data. *Statistics in Medicine*, 38(24), 4804–4818.

References II

- Lin, J., Li, K., & Luo, S. (2021). Functional survival forests for multivariate longitudinal outcomes: Dynamic prediction of Alzheimer's disease progression. *Statistical Methods in Medical Research*, 30(1), 99–111.
- Signorelli, M. (2024). Pencal: An R Package for the Dynamic Prediction of Survival with Many Longitudinal Predictors. *The R Journal*, 16(2), 134–153.
- Signorelli, M., & Retif, S. (2025+). Benchmarking multi-step methods for the dynamic prediction of survival with numerous longitudinal predictors. *arXiv Preprint arXiv:2403.14336*.
- Signorelli, M., Spitali, P., Szgyarto, C. A., The MARK-MD Consortium, & Tsonaka, R. (2021). Penalized regression calibration: A method for the prediction of survival outcomes using complex longitudinal and high-dimensional data. *Statistics in Medicine*, 40(27), 6178–6196.

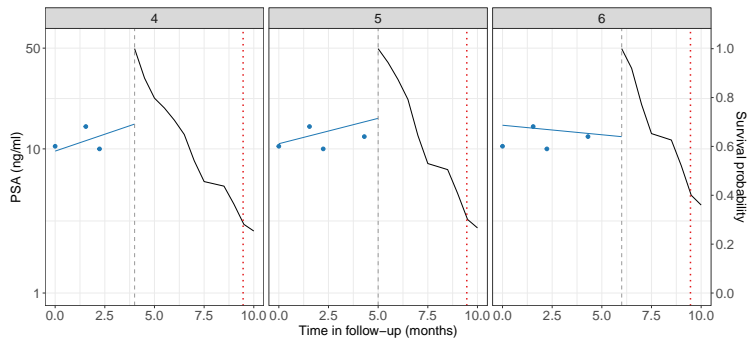
The dynamic prediction problem

Dynamic prediction with numerous longitudinal covariates

Benchmarking study

Appendix

Dynamic prediction example




Strict vs relaxed landmarking with two-step methods

Original Research Article



Dynamic prediction of survival using multivariate functional principal component analysis: A strict landmarking approach

Daniel Gomon¹ , Hein Putter² , Marta Fiocco^{1,2}
and Mirko Signorelli¹

Statistical Methods in Medical Research
2024, Vol. 33(2) 256–272

© The Author(s) 2024



Article reuse guidelines:

sagepub.com/journals-permissions

DOI: 10.1177/09622802231224631

journals.sagepub.com/home/smm



Strict vs relaxed landmarking with two-step methods

262

Statistical Methods in Medical Research 33(2)

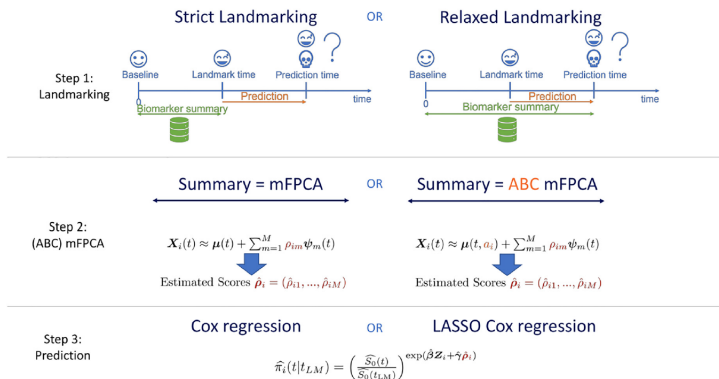


Figure 1. Graphical summary of the methods proposed in Section 2. See also Section 2.6.

Strict vs relaxed landmarking with two-step methods

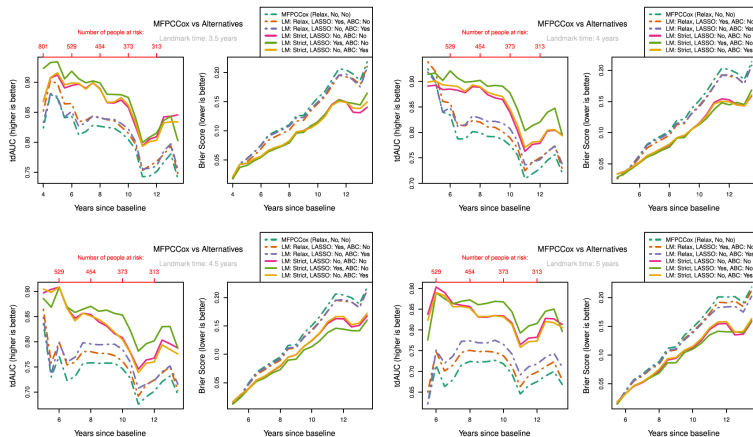
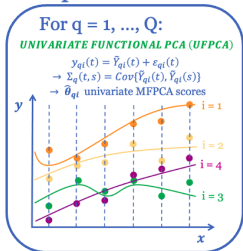


Figure 5. Measure of performance for LM, LASSO regularization (LASSO) and ABC methods at different landmark times on ADNI data. Validation scores were determined by using 20 times repeated 5-fold cross validation. Dashed lines: Relaxed landmarked methods. Solid lines: Strict landmarked methods. MFPCox (LM: Relax, LASSO: No, ABC: No)⁸ used as reference method. (a) Landmark time: 3.5 years; (b) Landmark time: 4 years; (c) Landmark time: 4.5 years; (d) Landmark time: 5 years. LM: landmark; ABC: age-based centered; ADNI: Alzheimer Disease Neuroimaging Initiative.

Multivariate Functional Principal Component Cox model (MFPCCoX, Li & Luo (2019))

Step 1



Step 2

Given the UFPCA scores $\hat{\theta}_{qi}$, compute the **multivariate FPCA scores**

$$H = (I - 1)^{-1} \Theta^T \Theta \rightarrow \hat{\rho}_i.$$

Select MFPCA scores so that they explain 90/95% of original variance

Step 3

Estimate

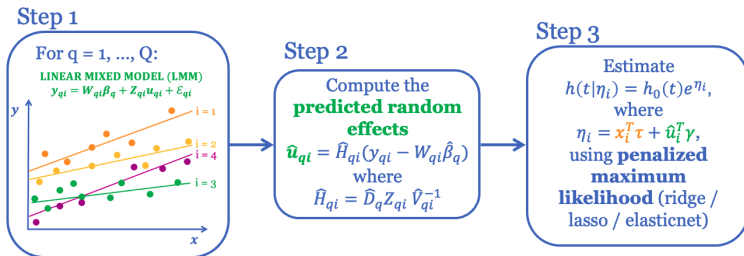
$$h(t|\eta_i) = h_0(t)e^{\eta_i},$$

where

$$\eta_i = \mathbf{x}_i^T \boldsymbol{\tau} + \hat{\rho}_i^T \boldsymbol{\gamma},$$

using **maximum likelihood**

Penalized Regression Calibration (pencal, Signorelli et al. (2021))



- Steps 1-2: multivariate version with MLPMM possible

Functional Random Survival Forest (FunRSF, Lin et al. (2021))

Step 1

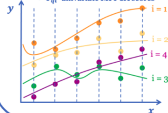
For $q = 1, \dots, Q$:

UNIVARIATE FUNCTIONAL PCA (UFPCA)

$$y_{qi}(t) = \hat{y}_{qi}(t) + \varepsilon_{qi}(t)$$

$$\rightarrow \Sigma_{qi}(t, s) = \text{Cov}\{\hat{y}_{qi}(t), \hat{y}_{qi}(s)\}$$

$$\rightarrow \hat{\theta}_{qi} \text{ univariate MFPCA scores}$$



Step 2

Given the UFPCA scores $\hat{\theta}_{qi}$, compute the **multivariate FPCA scores**

$H = (I - 1)^{-1} \theta^T \theta \rightarrow \hat{\rho}_t$. Select MFPCA scores so that they explain 90/95% of original variance

Step 3

Survival tree:



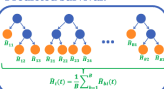
Randomly select k candidate predictors from:

MFPCA **Time-independent**
scores $\hat{\rho}_t$ x_{pt}

Select predictor and threshold that maximize the log-rank test statistic

Repeat for $b = 1, \dots, B$ bootstrap samples

Predicted survival:



Dynamic Random Survival Forest (DynForest, Devaux et al. (2023))

