

# Validation of a new flexible spline-based relative survival model

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# OUTLINE

- Background
  - Role of relative survival in prognostic studies
  - Limitations of published models
  - Importance of joint modelling of non-linear and time-dependent effects
- New model
  - Validation of our model
    - ✓ Design of simulations
    - ✓ Results
- Conclusions

# BACKGROUND

## ➤ Relative survival

- ✓ Useful methodology in situations where the cause of death is unknown, especially in:
  - in population-based cancer registries
  - for rare diseases with long survival
- ✓ Permits estimation of the covariate effects on disease-specific mortality

# BACKGROUND

## ➤ the Estève's et al relative survival model:

- ✓ Assumes that the observed hazard for total mortality  $\lambda_t$ , is the sum of two hazards:

$$\lambda_t(t|z, a) = \lambda_e(t + a|z_s) + \lambda_c(t|z)$$

- $\lambda_e$  represents the expected hazard function for all-cause mortality in the population at large
  - $\lambda_c$  is the excess hazard due to disease-specific mortality
- ✓ Requires estimation of baseline hazard for disease specific mortality

# BACKGROUND

## ➤ In Estève's et al relative survival model:

- ✓ Disease-related hazard is *a priori* constrained to meet 2 conventional assumptions:
$$\lambda_c = \sum_k \tau_k I_k(t) * \exp(\beta Z)$$
  1. Proportional hazards (PH) → relative risks are constant over time
  2. Log-linearity for continuous covariates (LL) → the logarithm of the hazard increases linearly with increasing value of the covariate

## ➤ Several “Crude” and Relative survival models were proposed to relax one or both of these assumptions:

- ✓ Abrahamowicz and MacKenzie (2007)
- ✓ Giorgi et al (2003)
- ✓ Remontet et al (2007)
- ✓ Lambert (2006)

# BACKGROUND

- Recently, Abrahamowicz & MacKenzie (2007) demonstrated, in crude survival, the importance of modelling and testing both PH and LL hypotheses jointly in order to :
  - ✓ avoid a spurious non-proportional effects due to not accounting for loglinearity (and vice versa)
  - ✓ avoid residual confounding
  - ✓ increase power for identifying new prognostic factors

# OUR JOINT RELATIVE SURVIVAL MODEL

- We propose a flexible extension of the Estève's relative survival model

$$\lambda_t = \lambda_e + \lambda_c$$

- The effect of covariates on  $\lambda_c$  are modelled with quadratic B-spline functions, which relax PH and LL assumptions

# OUR JOINT RELATIVE SURVIVAL MODEL

➤ Therefore, in our model

$$\lambda_c(t|z) = \exp(\gamma(t)) * \exp\left(\sum_i \alpha_i(z_i) * \beta_i(t)\right)$$

- ✓  $\gamma(t)$  represents the baseline hazard
- ✓  $\alpha(z)$  describes how the hazard changes with increasing value of a continuous covariate
- ✓  $\beta(t)$  represents the pattern of changes over time in the impact of the covariate on the risk of death

# OUR JOINT RELATIVE SURVIVAL MODEL

➤ A 3-step iterative alternative conditional full MLE estimation

1. Baseline hazard:  $\gamma(t) = \exp(\sum_{k=-2\dots 2} \gamma_k B_{k,3}(t))$

2. Non-linear dose response:  $\alpha(z) = \sum_{m=-2\dots 1} \alpha_{im} B_{m,3}(z)$

3. Time-dependent effect:  $\beta(t) = \sum_{l=-2\dots 2} \beta_{il} B_{l,3}(t)$

# GENERAL DESIGN OF SIMULATIONS

- We simulated a prognostic study of colon cancer mortality:
- $N = 2000$  subjects
- Major steps in data generation:
  1. Covariate vectors
  2. Three times (for each subject):
    - (i) Cancer-related death (conditional on covariates)
    - (ii) Other-cause death (conditional on age & sex)
    - (iii) Right “administrative” censoring (at 5 yrs)
  3. Final “Observed” time =  $\min \{(i), (ii), (iii)\}$

# COVARIATES GENERATION

## ➤ 5 covariates simulated:

1. Sex ~ Bin(1,0.5)
2. Age:  
Age ~ N(69.8,11.1) for men  
Age ~ N(71.4,12.6) for women
3. Year of diagnosis ~ U(1976,2000)
4. Stage: stage II vs stage III  
Stage II ~ Bin (1,0.61)
5. Location: right colon vs left colon  
Left colon ~ Bin(1,0.62)

# GENERATION OF TIME TO CANCER-RELATED DEATHS

## ➤ Using Permutational Algorithm\*

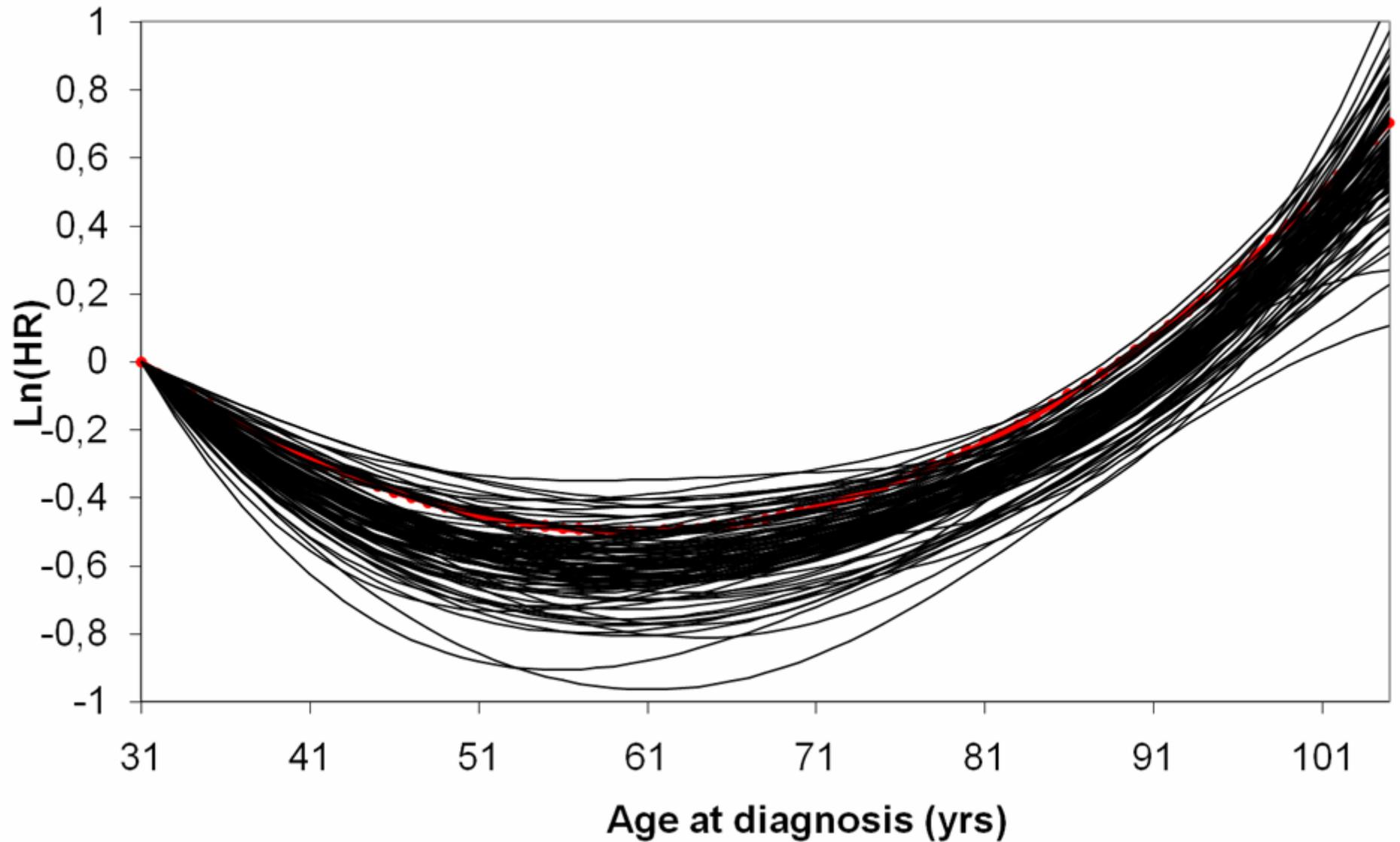
- ✓ (a) Generate N observed times  
Calculate the expected number of deaths related to cancer per month (independent of covariates)
- ✓ (b) Match N covariate vectors with N times generated in step (a) based on the probability derived from the partial likelihood
- ✓ Specifically, we assume the following model for hazard of cancer-related death while assuming non-linear and time-dependent effects of age on the hazard of cancer-related mortality:

$$HR = \exp \left( \begin{array}{l} \log(1.25) * sex + \log(1.2) * location + \log(3) * stage + \\ \alpha_{age}(age) * \beta_{age}(time) \end{array} \right)$$

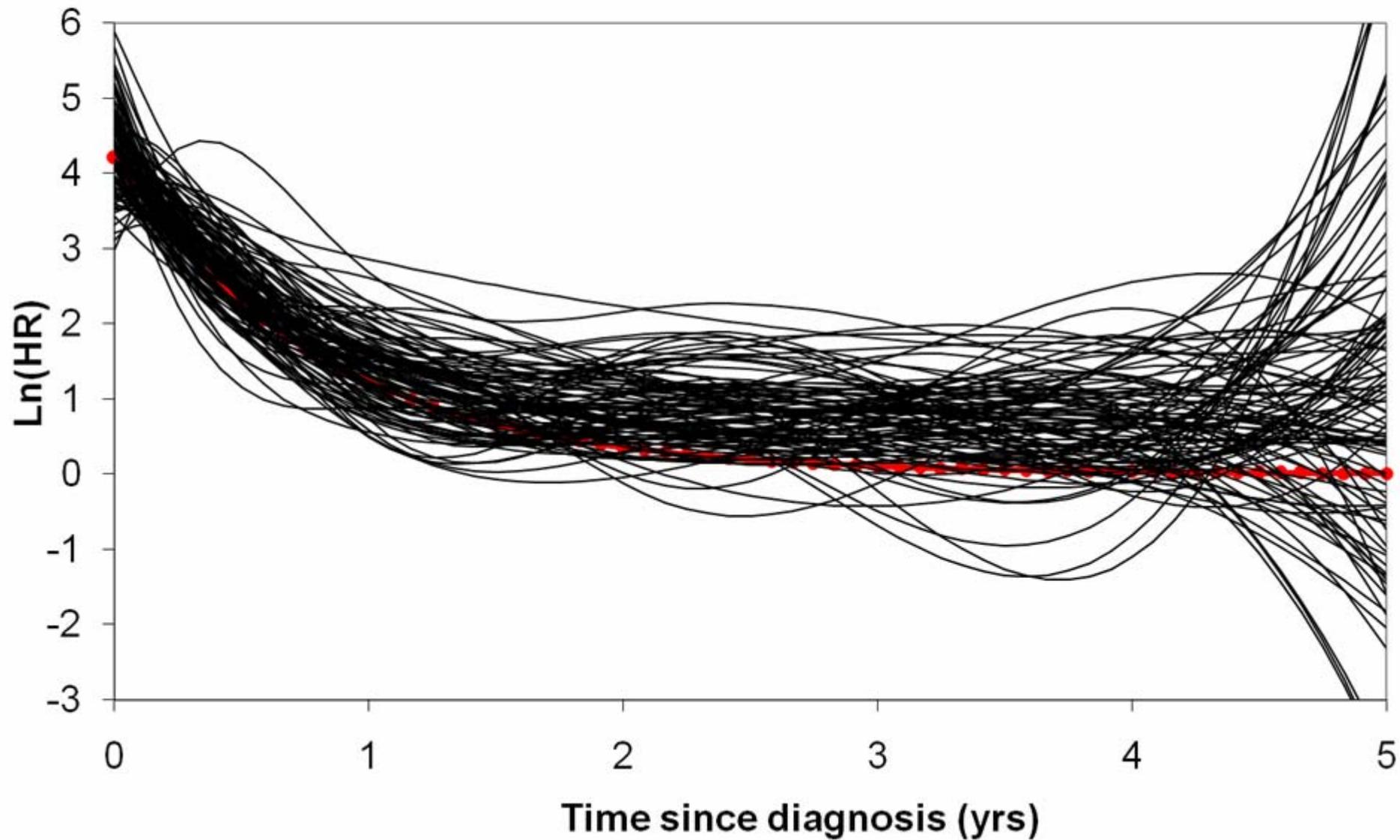
# GENERATION OF “NATURAL” (OTHER-CAUSE) DEATHS

- Time to natural death was generated using the French lifetables, stratified by Sex, Age and Calendar Year

# NON LOG-LINEAR EFFECTS OF AGE



# TIME DEPENDENT EFFECTS OF AGE



# CONCLUSION

- We have implemented a new flexible relative survival model
- This model allows for both simultaneous estimation and testing of time-dependent effects and non-log-linear effects for continuous covariates
- The simulations show
  - ✓ The general shape of both the time dependent and non log linear effects is recovered but there is over-fit bias (especially for TD effects)
  - ✓ The LRT tests show that our model is able to detect a true time dependent effect or a non loglinear effect

# BIBLIOGRAPHY

1. Dickman PW, Sloggett A, Hills M, Hakulinen T. Regression models for relative survival. *Statistics in Medicine* 2004;**23**: 51-64
2. Esteve J, Benhamou E, Croasdale M, Raymond L. Relative survival and the estimation of net survival: elements for further discussion. *Statistics in Medicine* 1990;**9**: 529-38.
3. Giorgi R, Abrahamowicz M, Quantin C, Bolard P, Esteve J, Gouvernet J, Faivre J. A relative survival regression model using B-spline functions to model non-proportional hazards. *Statistics in Medicine* 2003;**22**: 2767-84.
4. Remontet L, Bossard N, Belot A, Esteve J. An overall strategy based on regression models to estimate relative survival and model the effects of prognostic factors in cancer survival studies. *Statistics in Medicine* 2007;**26**: 2214-28.
5. Le Teuff G, Abrahamowicz M, Bolard P, Quantin C. Comparison of Cox's and relative survival models when estimating the effects of prognostic factors on disease-specific mortality: a simulation study under proportional excess hazards. *Statistics in Medicine* 2005;**24**: 3887-909.
6. Quantin C, Abrahamowicz M, Moreau T, Bartlett G, MacKenzie T, Tazi MA, Lalonde L, Faivre J. Variation over time of the effects of prognostic factors in a population-based study of colon cancer: comparison of statistical models. *American Journal of Epidemiology* 1999;**150**: 1188-200.
7. Abrahamowicz M, MacKenzie TA. Joint estimation of time-dependent and non-linear effects of continuous covariates on survival. *Statistics in Medicine* 2007;**26**: 392-408.
8. Lambert PC, Smith LK, Jones DR, Botha JL. Additive and multiplicative covariate regression models for relative survival incorporating fractional polynomials for time-dependent effects. *Statistics in Medicine* 2005; **24**: 3871-85.
9. Ramsay JO, Abrahamowicz M. Binomial regression with monotone splines: A psychometric approach. *Journal of the American Statistical Association* 1989;**84**: 906-915.
10. Sylvestre M.P., Abrahamowicz M. Comparison of algorithms to generate event times conditional on time-dependent covariates. *Statistics in Medicine* 2008;**27**: 2618-2634