

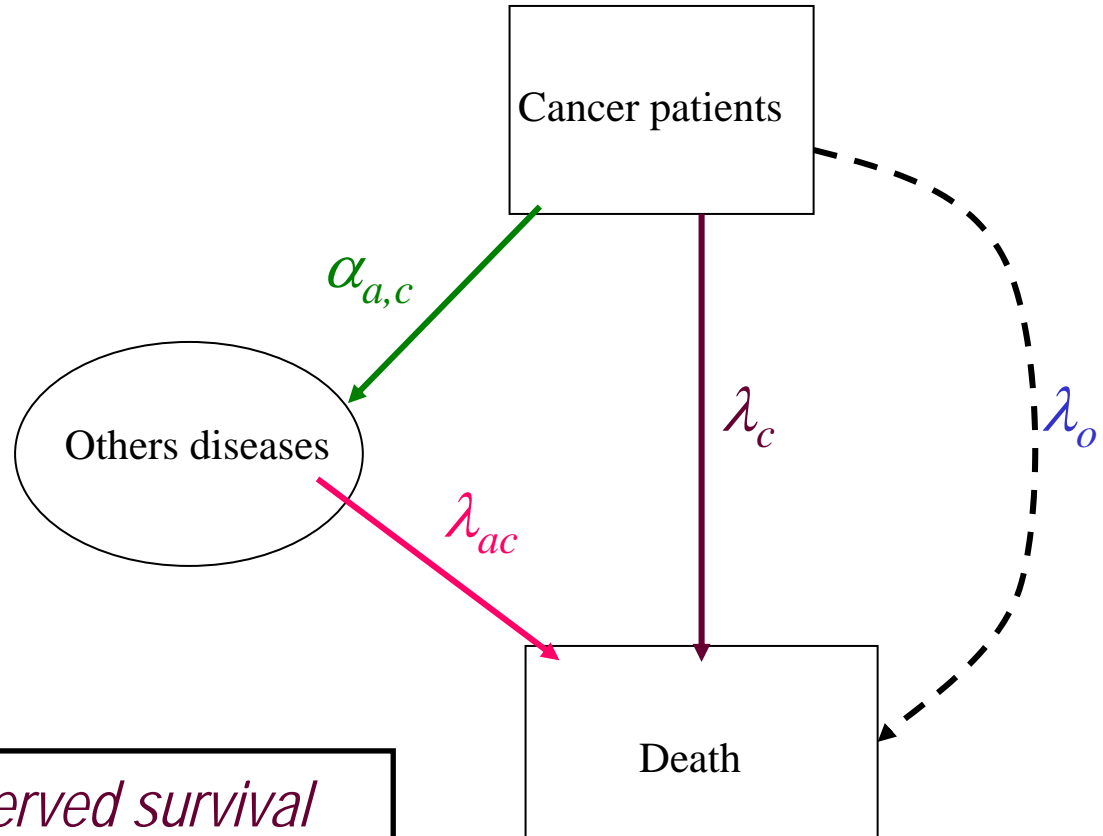
Relative survival vs Net survival definition of concepts and current methods of estimation

PARIS, March 2009

The concept of net survival

α =incidence

λ =lethality



Total Lethality \Leftrightarrow observed survival

$$\lambda_o = \lambda_c + \alpha_{ac} \oplus \lambda_{ac}$$

$$\text{Net survival} = \exp\left[-\int_0^t \lambda_c(u) du\right]$$

= survival which would be observed in absence of other causes of death

- The net survival may be *estimated* from survival data of a given cohort, if the cause of death of each deceased subject is known i.e **cause-specific** survival :
 - An observation is censored at the time of death if the death is not caused by the cancer under study.
- This estimation may be biased due to the subjective determination of the cause of death

The concept of Relative survival

- Relative survival was initially defined as the ratio :
observed survival / expected survival

where expected survival is the survival of a group of the general population with similar characteristics z (sex, age,) than the studied cohort (obtained from the life table)

→ Relative survival coincide in *some* situations with Net survival as

$$S_c = S_o / S_e \Leftrightarrow \lambda_c = \lambda_o - \lambda_e$$

How to estimate relative survival

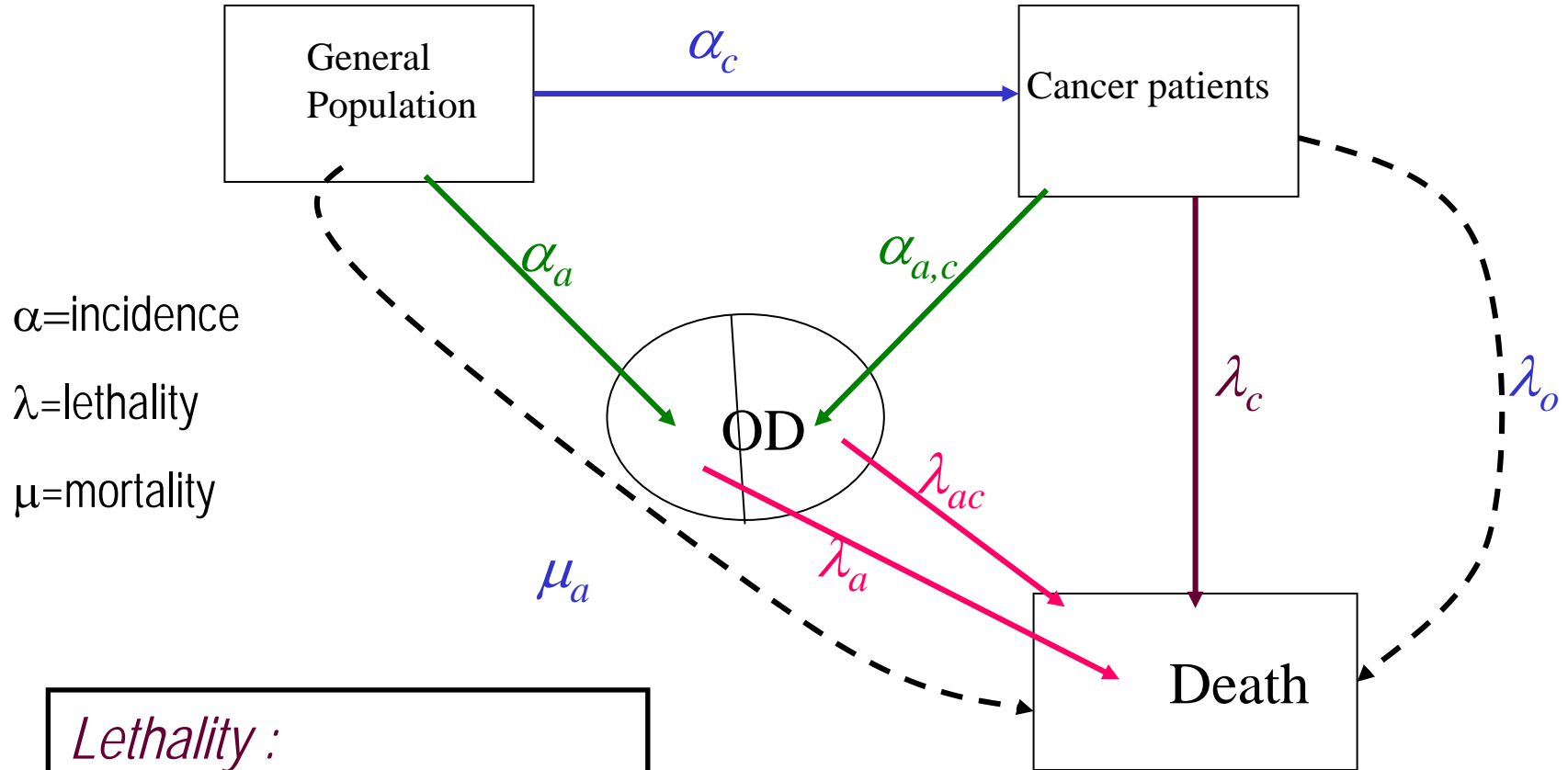
The “ratio estimate”

$$S_c(t) = S_o(t) / S_e(t) \rightarrow \hat{S}_c(t) = \hat{S}_o(t) / \hat{S}_e(t)$$

The relative survival probability is estimated from the observed survival estimate divided by the “known” probability of surviving as obtained from the life table= Plug-in estimate

- Initially the estimation methods relied on the simple calculation of the expected survival from *ad-hoc* life table.
 - The relative survival was then calculated by dividing the “actuarial” or the “KM” survival estimate by this expected survival
- The methods (Ederer I, Ederer II, Hakulinen) differed only in the way the expected survival was computed

How to estimate net survival from excess rate model (1)



Lethality :

$$\lambda_o = \lambda_c + \alpha_{ac} \otimes \lambda_{ac}$$

Hypothesis of excess rate model

$$\lambda_o = \lambda_c + \mu_a$$

How to estimate Net survival (2)

The excess death rate model

- The idea is to estimate the *death rate in excess of the background mortality rate* to obtain the Net relative survival from it (instead of working in survival level)

$$\lambda_o(t) = \lambda_c(t) + \mu_e(x+t, z)$$

Where $\lambda_c(t)$ is a function to be estimated from survival data and μ_e is obtained from vital statistics or life table

- An initial choice for λ_c was the step function:

$$\lambda_c(t) = f(t) = \sum_{k=1}^K \tau_k I_k(t)$$

where the follow-up time has been split in K intervals.

Alternatives for f have been developed.

Problems with relative survival/Net survival (1)

- If the whole group is a **heterogeneous group** relative to variables (age) affecting
 - the cancer death and/or
 - the death for others causes and/or
 - the probability of being censored,

⇒ RS and NS don't necessarily coincide

⇒ RS gives biased estimates of NS

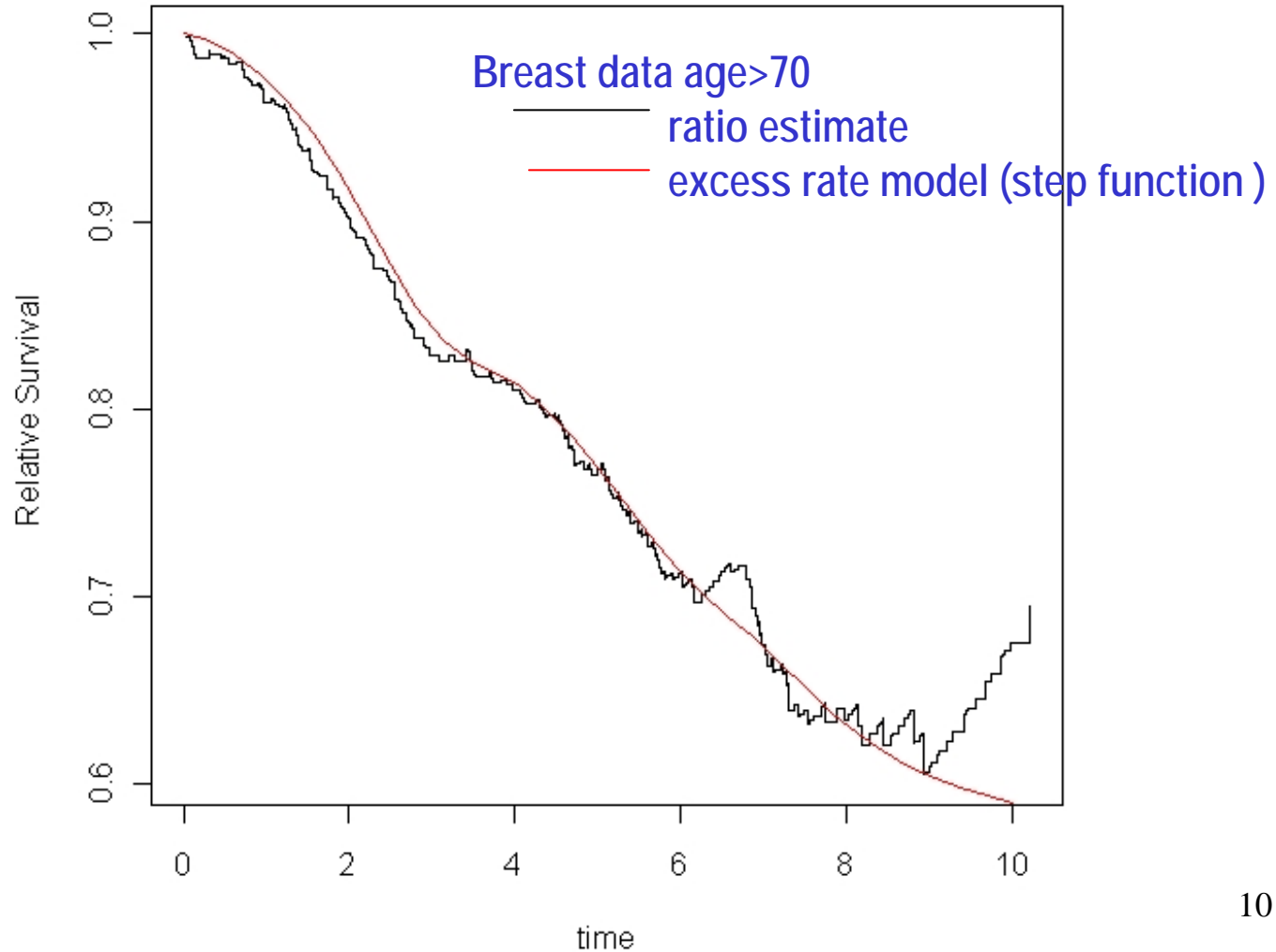
Ex: SR from Ederer 1 coincide to NS (and is an estimator of...) only if cancer death and censoring are independent of age

Problems with relative survival/Net survival (2)

- We need estimate of $S_c(t, \text{age})$ to calculate age standardised relative survival (for worldwide comparisons)
 - The ratio estimate in each strata may be subject to a lack of “robustness” and may show rising relative survival

Example : breast cancer age >70 years

Relative survival estimates by the "ratio estimate" vs Net survival estimates by "excess rate model" in a strata



Estimating Net survival from a (excess rate) model may be more robust by

- 1) Modelling $f(t)$ with a smooth continuous function instead of a step function, in a given strata, or
- 2) An appropriate modelling for age (non linear, non proportional...)...on the whole dataset.

→ Illustrations ...

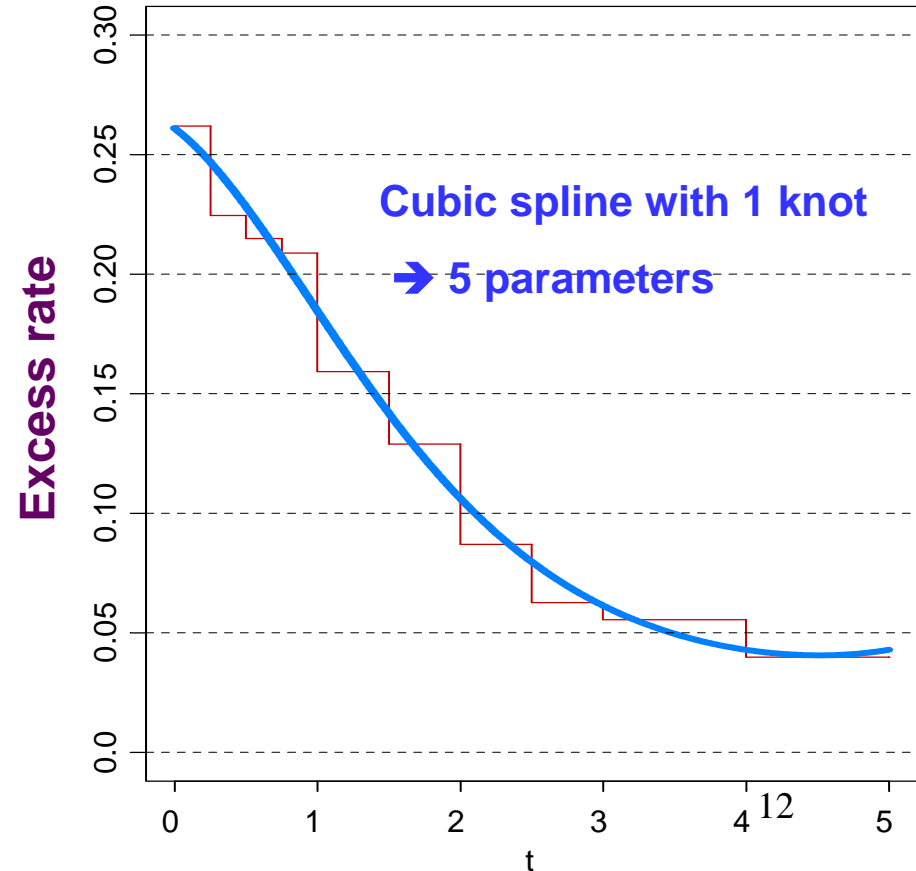
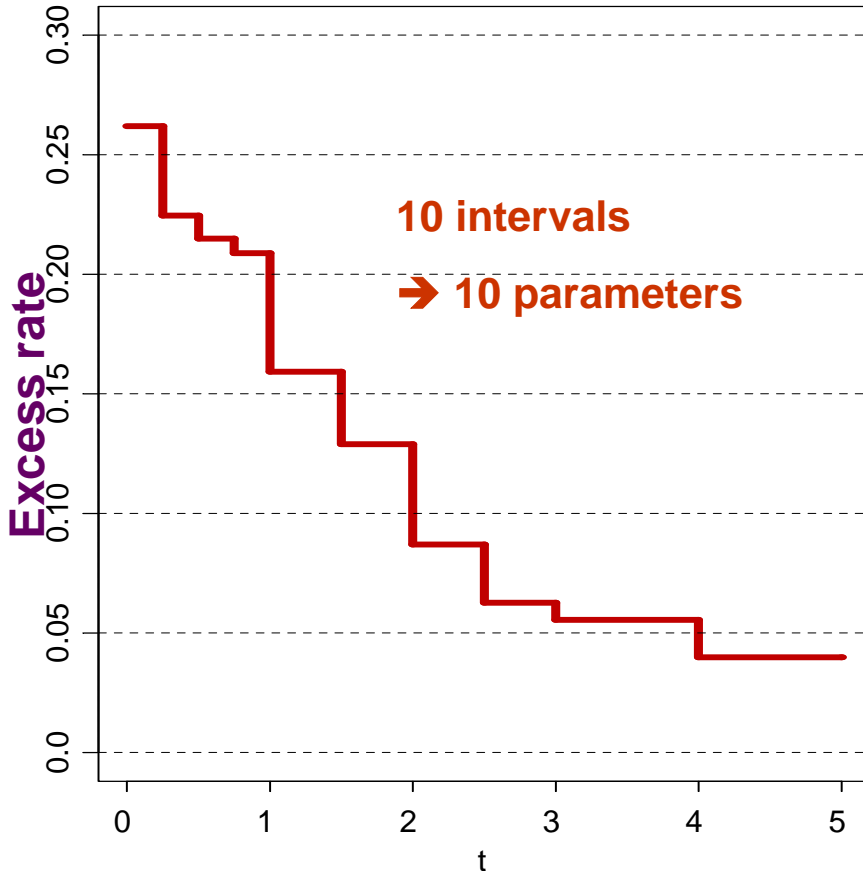
Step versus Smooth function

$$\lambda_c(t, z) = \exp(\beta z) f(t)$$

with $f(t)$ being

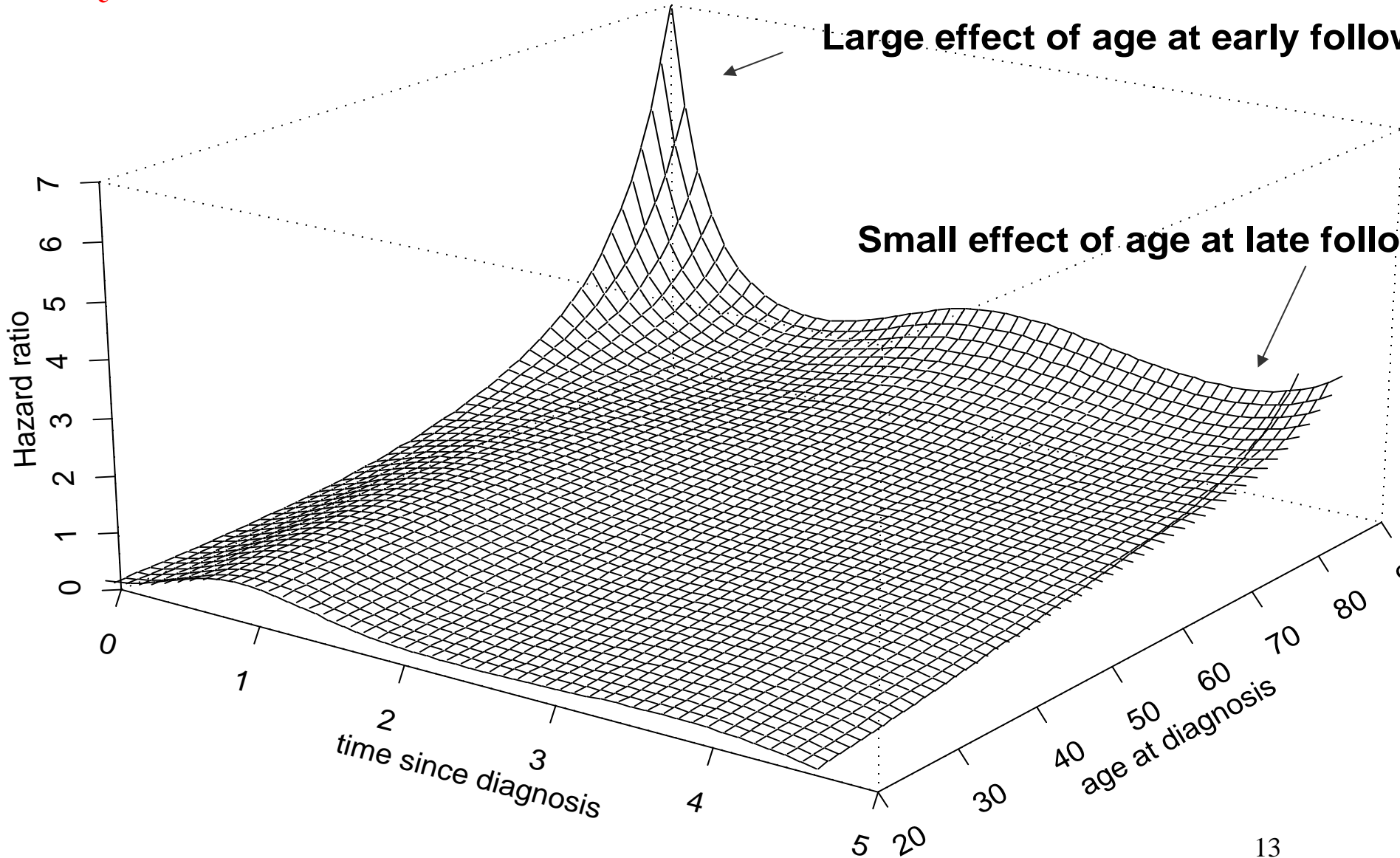
constant within intervals defined a priori

A smoothed parametric function



Time varying age effect (Kidney cancer)

$$\text{Log}[\lambda_c(t, \text{age})] = f(t) + g(\text{age}) + h(t) * \text{age}$$



Review of methods of estimation of
Relative survival / Net survival

and

their availability in usual statistical software (SAS, Stata, S+/R)

➤ Ederer I et al, Hakulinen & al, Biometrics 1982

Methods for relative survival

➤ Hakulinen & al, Applies Stat, 1987

- λ_c =step function
- Grouped data:
 - GLM implementation (Bin, log-log)
 - Categorical explanatory variable

➤ Esteve & al, SIM, 1990

- λ_c =step function
- Full ML on Individual data
- Available in command strel (Stata)
- Available in function relsurv (R-package, Pohar & al)

*Methods for
Net survival (1)*

$$\lambda_c(t) = \lambda_0(t) \exp(\beta X)$$

➤ Dickman, SIM 2004

- λ_c =step function
- Likelihood(survival model)= Likelihood(Poisson model) :
GLM (P, log) on splitted data
- Available in function relsurv (R-package, Pohar & al)

Methods for Net survival (2)

- Bolard & al, JCEP 2002
 - λ_c = quadratic regression splines
 - Non proportionnal hazard

- Giorgi & al, SIM 2003
 - λ_c = quadratic regression splines
 - NPH
 - Available in RSurv (R-package, Giorgi)

- Lambert, SIM 2005
 - λ_c = fractionnal polynomial

- Remontet & al, SIM 2007
 - λ_c = regression splines
 - NPH & NLIN
 - Specific strategy for survival estimate

Flexible parametric of
modelling of $\lambda_0(t)$

$$\lambda_c(t) = \lambda_0(t) \exp(\beta X)$$

$$\lambda_c(t) = \lambda_0(t) \exp(\beta X)$$

Non parametric modelling of $\lambda_0(t)$

➤ Pohar-Perme &al, Biostatistics 2009

- λ_c = not specified = Partial Likelihood
- Estimation procedure by EM algorithm (cause of death treated as missing data)
- Available in R-package relsurv (Pohar &al)

Others approaches that are NOT:

- $\lambda_{\text{obs}} = \lambda_c + \lambda_E$
- $\lambda_c = \lambda_0 \exp(\beta X)$

- Additive model for $\lambda_c(t)$ [Zahl LDA 1998, Cortese SIM 2008]: $\lambda_c(t) = \lambda_0(t) + \beta X$

- Multiplicative model for $\lambda_{\text{obs}}(t)$ [Andersen, Biometrics 1985]: $\lambda_{\text{obs}}(t) = \lambda_c \times \lambda_e$

Project : part 1

theoretical comparison of methods

Several methods **of estimation of relative survival/ Net survival** have been developed in the last decades

→ There is a need for a collaborative work aiming at

- Understanding when the concepts SR/NS coincide
- Understanding what the concepts and the methods of estimation are supposed to do
- Identifying their underlying hypothesis, their robustness
- Evaluating their respective performances in terms of biases and precision using simulation

The MESURE project is an opportunity **for achieving this goal**

Project : part 2

Empirical comparison of methods: Simulations

- Comparison of Net survival estimates (or β) from the different methods with the 'true' simulated net survival using several scenarii:
 - 1) Different patterns of the excess death rate with time elapsed since the diagnosis
 - 2) Effect of covariates (age) on the excess death rate : null, linear, non linear, non proportional...
 - 3) Effect of covariates (age) on expected survival and censoring
 - 4) sample size scenarii leading to sparse data

Bibliography

- PK. Andersen, *Biometrics*, 41, 921-932 (1985)
- P. Bolard et al., *J Cancer Epidemiol.Prev.* 7, 113-122 (2002).
- G. Cortese and T. H. Scheike, *Stat.Med.* 27, 3563-3584 (2008).
- P. W. Dickman, A. Sloggett, M. Hills, T. Hakulinen, *Stat.Med.* 23, 51-64 (2004).
- F. Ederer et al., *Natl Cancer Inst.Monogr.* 6:101-21., 101-121 (1961).
- J. Esteve, E. Benhamou, M. Croasdale, L. Raymond, *Stat Med* 9, 529-538 (1990).
- R. Giorgi et al., *Stat.Med.* 22, 2767-2784 (2003).
- T. Hakulinen, *Biometrics.* 38, 933-942 (1982).
- T. Hakulinen et al. *Applied Statistics*, 36, 309-17 (1987)
- P. Lambert et al, *Stat.Med.* 24, 3871-3885 (2005).
- M. P. Perme, R. Henderson, J. Stare, *Biostatistics.* (2009).
- M. Pohar and J. Stare, *Comput.Methods Programs Biomed.* 81, 272-278 (2006).
- M. Pohar and J. Stare, *Comput.Biol.Med.* 37, 1741-1749 (2007).
- L. Remontet, N. Bossard, A. Belot, J. Esteve, *Stat.Med.* 26, 2214-2228 (2007).
- J. Stare, M. Pohar, R. Henderson, *Stat.Med.* 24, 3911-3925 (2005).
- P. H. Zahl and O. O. Aalen, *Lifetime.Data Anal.* 4, 149-168 (1998).