# Event projection: quantify uncertainty and manage expectations of broader teams

Kaspar Rufibach Department of Biostatistics, Roche Basel Methods, Collaboration & Outreach Group Basel, 28th April 2016



# **Agenda**

- Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tail
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

# **Agenda**

- Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tail
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

# Challenge

#### Clinical trial with time-to-event outcome:

- Portion of patients accrued, some patients still to be accrued.
- Interim or final analysis after pre-specified number of events.

#### Questions:

- **1** When does this number of events happen?
- Can we assign a confidence interval around that timepoint?
- How to manage the potentially large uncertainty in clinical teams (and higher up)?

#### Implications:

- Resource allocation for running the trial.
- Gating of other trials, e.g. early phase combinations.
- Keep investors happy.

# **Example**

Phase 3 trial with time-to-event primary endpoint (PFS).

All n = 1202 patients accrued: FPI 2010-04-06, LPI 2012-11-06.

First interim analysis:

• Clinical cutoff date: 2012-11-21.

• #events: 113.

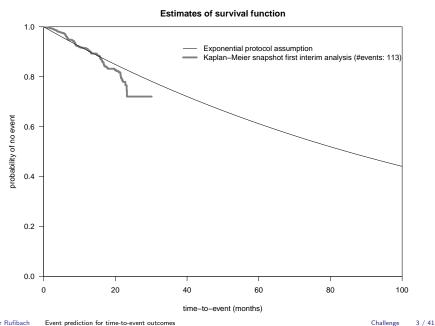
Based on this interim data  $\Rightarrow$  predict time when we see

• 248 events: second interim,

370 events: final analysis.

2 / 41

# Kaplan-Meier estimate based on snapshot



# **Agenda**

- 1 Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tai
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

# **Expected number of events at** T

S: chosen or estimated survival function.

Expected number of events m at T:

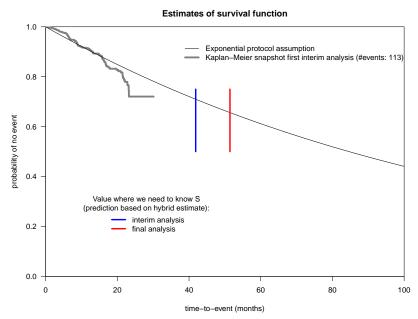
$$m(S) = \sum_{h \in \mathcal{I}_1} \mathbf{1} + \sum_{j \in \mathcal{I}_2} P\Big(\text{event in } (t_j, T] | \text{no event up to } t_j\Big) + \sum_{j \in \mathcal{I}_3} P\Big(\text{event in } (a_i, T]\Big)$$

$$= n + \sum_{j \in \mathcal{I}_2} \frac{S(t_j) - S(T)}{S(t_j)} \mathbf{1}\{T > t_j\} + \sum_{j \in \mathcal{I}_3} \Big(1 - S(T - a_i)\Big) \mathbf{1}\{T > a_i\}.$$

#### Sets of patients:

- $\mathcal{I}_1$ : patients who already had event, n.
- $\mathcal{I}_2$ : patients censored at  $t_i$ .
- $\mathcal{I}_3$ : patients to be recruited at  $a_i$ .

Need to have value for S at least up to T.



### How to estimate *S*?

Nonparametrically via Kaplan-Meier or kernel estimate:

- + No or few assumptions about true  $S \Rightarrow$  unbiased.
- + Can account for the "large steps" due to schedule of assessment.
- High variability, especially in tail.
- No extrapolation beyond last event / censoring time.

#### Fully parametric, e.g. Exponential or Weibull:

- + Efficient, if assumption is true.
- + Accurate estimate of tail, if assumption is true.
- + Can estimate S beyond where we have data.
- Biased if assumption not true.
- Not able to capture particular features of oncological time-to-event data (inspection intervals).

Hybrid approach: Use nonparametric where we have data, complement with parametric tail.

# **Agenda**

- Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tail
- 4 Predicted analysis timepoints
- **(5)** Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

# **Hybrid estimate**

Proposed in Fang and Su (2011). Recipe:

- **①** Choose total number of change points k.
- 2 Estimate piecewise Exponential hazard with k change points.
- 3 Test for "significance" of change points.

Borrows strength of nonparametric and parametric approach.

Allows to capture "big steps" due to schedule of assessments.

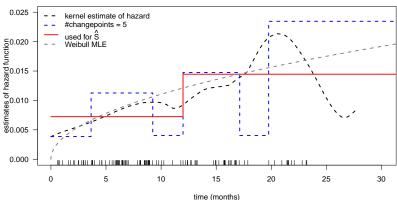
#### Change point:

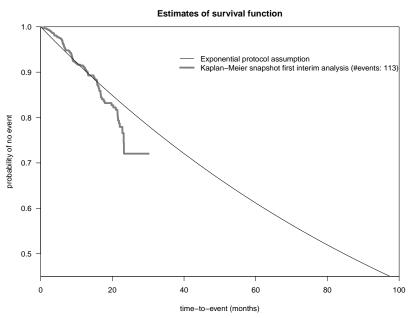
- Sequential test according to Goodman et al. (2011).
- To avoid overfitting correct  $\alpha$  according to Lan and DeMets (1983).
- No change point: pure Exponential fit.
- ullet  $\geq$  1 change point: Kaplan-Meier prior to selected change point, Exponential tail fit beyond selected change point.

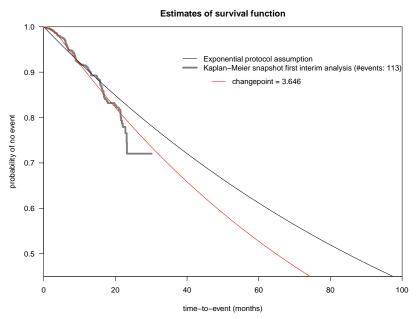
Alternatively: Bagiella and Heitjan (2001), Ying et al. (2004), Ying and Heitjan (2008), Di et al. (2016).

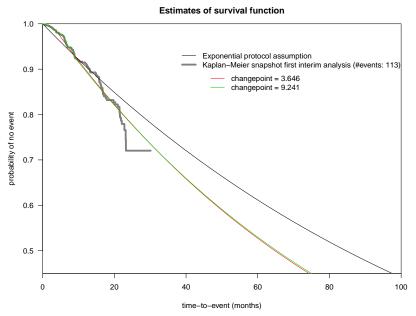
# Apply hybrid approach

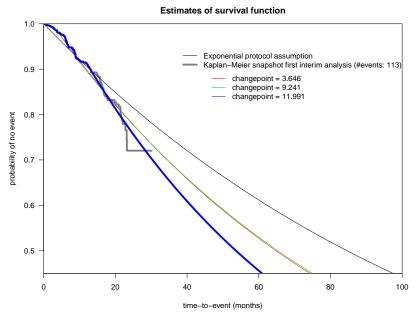


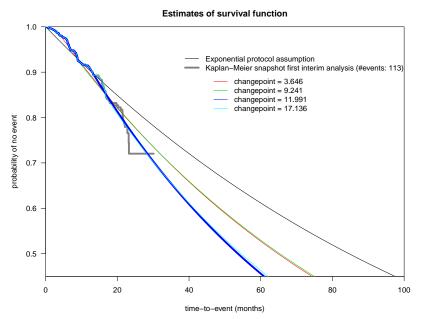


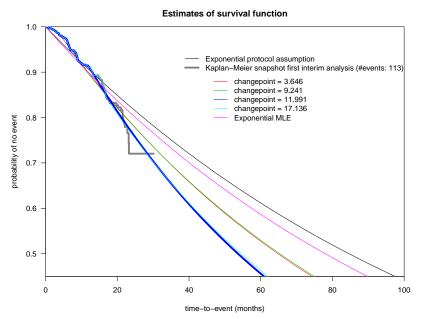


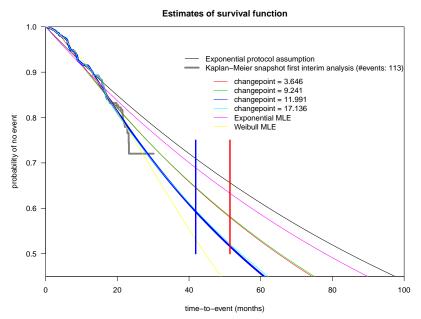








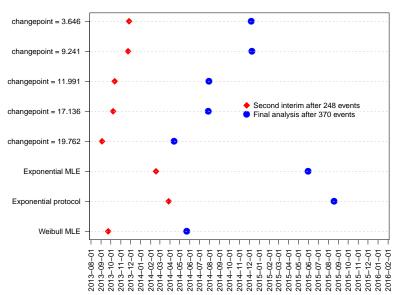




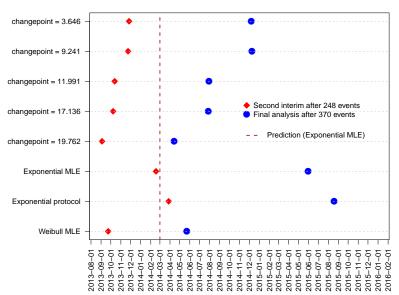
# **Agenda**

- 1 Challenge
- 2 How to predict analysis timepoint in general
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tai
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

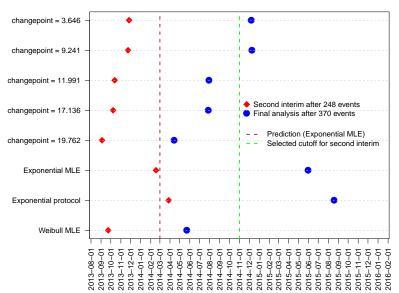
## **Predicted analysis timepoints**



## **Predicted analysis timepoints**



## **Predicted analysis timepoints**



## Shift

Communicated prediction: Exponential MLE  $\Rightarrow$  synthesis of protocol assumption and prediction based on hybrid estimate.

Prediction of second interim analysis:

Data: 113 events in 1202 patients.

Clinical cutoff: 2012-11-21.

• Initial prediction: 2014-03-01.

• Actual cutoff: 2014-11-01.

Shift of cutoff: 8.0 months.

## 8.0 months!



p.s.: 8.0 months earlier would certainly have been ok, but 8.0 months later...

# **Agenda**

- 1 Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tai
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

# Uncertainty in event prediction

Recall formula:

$$m(S) = n + \sum_{j \in \mathcal{I}_2} \frac{S(t_j) - S(T)}{S(t_j)} 1\{T > t_j\} + \sum_{j \in \mathcal{I}_3} (1 - S(T - a_i)) 1\{T > a_i\}.$$

Trial was fully recruited when prediction made  $\Rightarrow$  uncertainty = Sampling error in estimation of S.

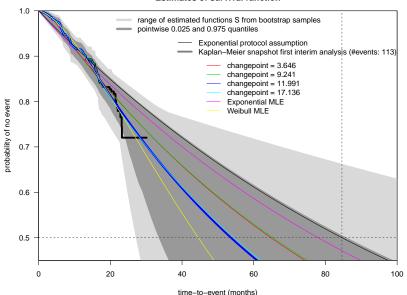
#### Proposal:

- Bootstrap time-to-event data: sample with replacement from  $(X_i, \delta_i)$ .
- Re-estimate S via hybrid Exponential model for each bootstrap sample, choose change point based on sequential test ⇒ S estimated fully automatic.
- Compute analysis timepoint in each sample.
- Compute quantiles of these analysis timepoints ⇒ bootstrap percentile confidence interval for analysis timepoint.

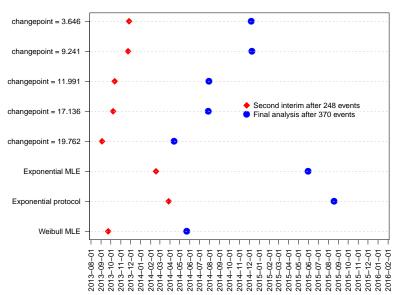
Validity of bootstrap for right-censored data: Efron (1981), see also Akritas (1986).

## **Bootstrapped survival function estimates**

#### Estimates of survival function

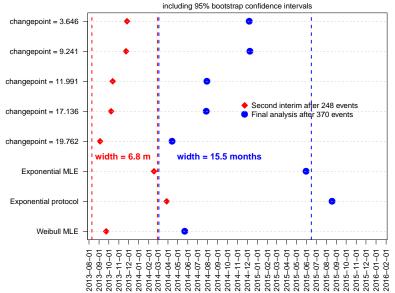


## **Analysis timepoints**



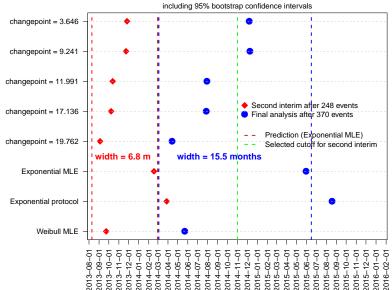
## **Analysis timepoints**





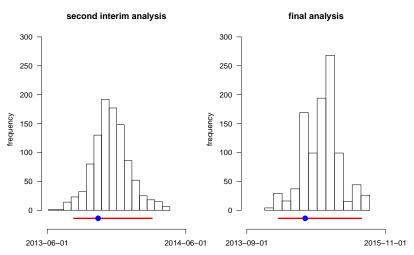
# **Analysis timepoints**





## **Bootstrap distributions**

Histograms of analysis timepoints, based on 1000 bootstrap samples Blue dot: point estimate based on hybrid estimate. CCOD 2012–11–21, 113 events.



## **Confidence intervals**

Statistically sound.

Difficult to communicate to Big Boss. Initially, we did not communicate confidence interval, only gave point estimate.

#### Statistician:

Your interim analysis cutoff will be reached between 2013-08-10 and 2014-03-04, with 95% confidence.

## **Confidence intervals**

Statistically sound.

Difficult to communicate to Big Boss. Initially, we did not communicate confidence interval, only gave point estimate.

#### Statistician:

Your interim analysis cutoff will be reached between 2013-08-10 and 2014-03-04, with 95% confidence.

#### Big boss:

Fine.

## **Confidence intervals**

Statistically sound.

Difficult to communicate to Big Boss. Initially, we did not communicate confidence interval, only gave point estimate.

#### Statistician:

Your interim analysis cutoff will be reached between 2013-08-10 and 2014-03-04, with 95% confidence.

#### Big boss:

Fine. Give me the date.

## **Agenda**

- Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tai
- Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

## Small proportion of events - large variability

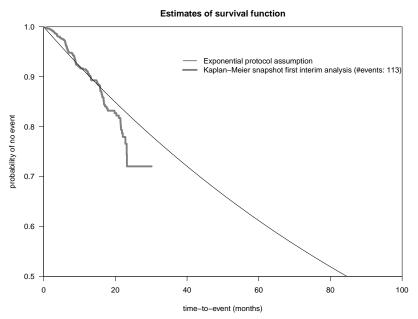
Approx. 1y after initial prediction.

- Data cleaning milestone.
- Based on 193 events became clear we need to shift cutoff.

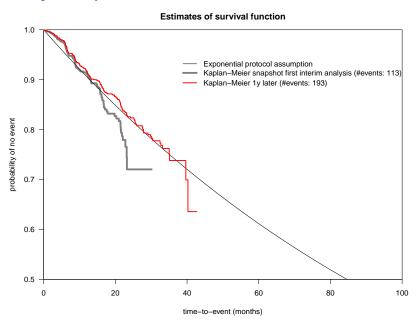
"Root-cause" analysis for 8.0 months shift.

Statistics team explained uncertainty around predictions  $\Rightarrow$  estimates of S quite different based on two snapshots.

## Variability in Kaplan-Meier estimate over time



## Variability in Kaplan-Meier estimate over time



# Big Boss comes back

#### Big Boss:

Seems you had more events in relation to amount of follow-up one year ago? How good was your cleaning back then anyway?

Maybe we had all events, but tumor assessment follow-up for those event-free was not comprehensive?

#### Compare:

- Data Snapshot 1 of first interim analysis: 3892 tumor assessments.
- Snapshot 2 one year later, cut back to first interim analysis cutoff: 3895 tumor assessments.
- Proportion of tumor assessments available in snapshot 1 if Snapshot 2 is considered "full dataset": 3892 / 3895 = 99.923%.

Message: we had everything back then. Difference in estimates

- either due to variability,
- or early events?

## **Agenda**

- 1 Challenge
- 2 How to predict analysis timepoint in general?
- 3 Hybrid estimate of S: Kaplan-Meier with Exponential tai
- 4 Predicted analysis timepoints
- 5 Uncertainty in event prediction
- 6 Small proportion of events
- Remarks

### Remarks

- Hybrid estimate of  $S \Rightarrow$  provides estimated value of S beyond last data point.
- Confidence intervals wide
- In fact used prediction based on Exponential MLE. Could simply have used that standard error.
- Typically: uncertainty driven by #events. But here: need to predict event time for those who had no event yet. Uncertainty driven by proportion of events at prediction (113) to patients (1202)? More research needed?
- Sources of variability not considered: drop-out.
- Not discussed, but essential: do such projections only on cleaned data. Otherwise, you likely have most of the events, but no comprehensive follow-up.
- Communication is key. But difficult.

#### **Technical comments**

- Once k is selected  $\Rightarrow$  automatic method to do event projection  $\Rightarrow$  unambiguity.
- Dependence on  $k \Rightarrow$  sensitivity analysis.
- Point estimate might not lie in center of bootstrap confidence interval. Use alternative function of bootstrap sample (mode, median).
- R package eventTrack, available internally. Everything implemented from scratch
   likely not optimally efficient.
- Confidence interval: 1202 patients and 1000 bootstrap samples: takes about 3h.
   Depends on T, k.

Thank you for your attention.

#### References

- Akritas, M. (1986). Bootstrapping the Kaplan-Meier Estimator. J. Amer. Statist. Assoc. 81 1032-1038.
- ▶ Bagiella, E. and Heitjan, D. F. (2001). Predicting analysis times in randomized clinical trials. Stat. Med. 20 2055-2063.
- ► Carpenter, J. and Bithell, J. (2000). Bootstrap confidence intervals: when, which, what? A practical guide for medical statisticians. Stat. Med. 19 1141-1164.
- ▶ Di, J., Wang, D., Brashear, H. R., Dragalin, V. and Krams, M. (2016). Continuous event monitoring via a bayesian predictive approach. Pharmaceutical Statistics 15 109-122.
- ▶ Efron. B. (1981). Censored Data and the Bootstrap. J. Amer. Statist. Assoc. 76 312–319.
- Fang. L. and Su. Z. (2011). A hybrid approach to predicting events in clinical trials with time-to-event outcomes. Contemp Clin Trials 32 755-759.

#### References

- Goodman, M. S., Li, Y. and Tiwari, R. C. (2011). Detecting multiple change points in piecewise constant hazard functions. J Appl Stat 38 2523–2532.
- ▶ Klein, J. P. and Moeschberger, M. L. (2003). Survival Analysis. 2nd ed. Springer-Verlag.
- Lan, K. K. G. and DeMets, D. L. (1983). Discrete sequential boundaries for clinical trials.
   Biometrika 70 659–663.
- Yao, Y. (1986). Maximum likelihood estimation in hazard rate models with a change-point. Comm. Statist. Theory Methods 15 2455–2466.
- Ying, G. S. and Heitjan, D. F. (2008). Weibull prediction of event times in clinical trials. Pharm. Stat. 7 107–120.
- Ying, G. S., Heitjan, D. F. and Chen, T. T. (2004). Nonparametric prediction of event times in randomized clinical trials. *Clin Trials* 1 352–361.

Backup slides.

### Piecewise constant hazard

Piecewise constant hazard ⇒ piecewise Exponential density or survival function.

"Bridges" parametric and nonparametric model.

Assume for a given number of change points k, hazard function h, and  $0 = \tau_0 < \tau_1 < \ldots < \tau_{k+1} = \infty$ :

$$h(t) = egin{cases} \lambda_1 & t \in [0, au_1) \ \lambda_2 & t \in [ au_1, au_2) \ dots \ \lambda_{k+1} & t \geq au_k. \end{cases}$$

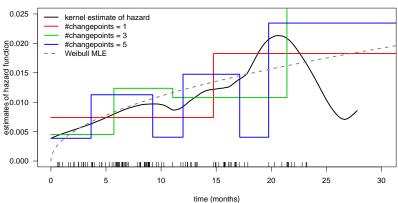
Survival function computed from hazard function:

$$S(t) = \exp(-\int h(t)dt).$$

Estimate  $(\lambda_1, \ldots, \lambda_{k+1}, \tau_1, \ldots, \tau_k)$  incl. standard errors using maximum likelihood (backup for details).

### Piecewise constant hazard

# hazard function



## Maximum likelihood for censored data

General log-likelihood function for censored data:

- We observe  $(T_i, \delta_i) = (\min\{X_i, C_i\}, \delta_i), i = \dots, n$ ,
- $h_{\lambda}$  is the hazard,
- $S_{\lambda}$  the survival function, both depending on  $\lambda$ .

Then (see e.g. Klein and Moeschberger (2003)):

$$\log(L(\lambda)) = \log\left(\prod_{i=1}^{n} h_{\lambda}(T_{i})^{\delta_{i}} S_{\lambda}(T_{i})\right)$$
$$= \sum_{i=1}^{n} \delta_{i} \log h_{\lambda}(T_{i}) + \sum_{i=1}^{n} \log S_{\lambda}(T_{i}).$$

For a constant hazard:  $h_{\lambda}(t) = \lambda$ ,  $S(t) = \exp(-\int \lambda dt) = \exp(-\lambda t)$ . So:

$$\log(L(\lambda)) = \sum_{i=1}^{n} \delta_{i} \log \lambda - \lambda \sum_{i=1}^{n} T_{i}.$$

Taking derivative w.r.t. to  $\lambda$  and setting equal to 0 yields  $\hat{\lambda} = \sum_{i=1}^n \delta_i / \sum_{i=1}^n T_i$ .

## MLE in a piecewise model for the hazard with censored data

Generalize to piecewise function  $\Rightarrow$  again, general recipe.

Piecewise Exponential: see e.g. references in Goodman et al. (2011). We get:

$$\begin{split} \log L(\lambda_1, \dots, \lambda_{k+1}, \tau_1, \dots, \tau_k) &= \\ &= \sum_{j=1}^{k+1} \{ \text{\#events in } (\tau_{j-1}, \tau_j] \} \log \lambda_j - \sum_{i=1}^n \sum_{j=1}^{k+1} \lambda_j \Big( \min \{ T_i, \tau_j \} - \min \{ T_i, \tau_{j-1} \} \Big). \end{split}$$

Closer look at  $\min\{T_i, \tau_j\} - \min\{T_i, \tau_{j-1}\}$ :

$$\min\{T_i, \tau_j\} - \min\{T_i, \tau_{j-1}\} = \begin{cases} 0 & \text{if } T_i < \tau_{j-1} \\ T_i - \tau_{j-1} & \text{if } T_i \in [\tau_{j-1}, \tau_j] \\ \tau_j - \tau_{j-1} & \text{if } T_i > \tau_j. \end{cases}$$

$$= (\min\{T_i, \tau_j\} - \tau_{j-1}) \mathbf{1}\{T_i > \tau_{j-1}\}.$$

Observation *i* contributes to estimation in interval  $(\tau_{j-1}, \tau_j]$  the observation time in that interval.

## MLE in a piecewise model for the hazard with censored data

Assume  $\tau_1, \ldots, \tau_k$  as fixed.

Take derivative of  $\log L(\lambda_1, \ldots, \lambda_{k+1}, \tau_1, \ldots, \tau_k)$  w.r.t. to  $\lambda_1, \ldots, \lambda_{k+1}$ . This gives:

$$\widehat{\lambda}_{j} = \frac{\text{\#events in } (\tau_{j-1}, \tau_{j}]}{\sum_{i=1}^{n} (\min\{T_{i}, \tau_{j}\} - \tau_{j-1}) 1\{T_{i} > \tau_{j-1}\}}.$$

"Usual" MLE for Exponential data, just in interval  $(\tau_{i-1}, \tau_i]$ .

Plug  $\widehat{\lambda}_1, \dots, \widehat{\lambda}_{k+1}$  into log  $L(\lambda_1, \dots, \lambda_{k+1}, \tau_1, \dots, \tau_k)$ , maximize numerically over  $\tau_1, \ldots, \tau_k$  to get estimates  $\widehat{\tau}_1, \ldots, \widehat{\tau}_k$ .

**Profile-likelihood!** Justified since asymptotically,  $\widehat{\lambda}_1, \dots, \widehat{\lambda}_{k+1}$  and  $\widehat{\tau}_1, \dots, \widehat{\tau}_k$  are independent, i.e. asymptotically, estimates are the same irrespective of whether we maximize separately for  $\tau$ 's and  $\lambda$ 's, or jointly, see Yao (1986).

Get standard errors of estimates based on standard maximum likelihood theory. At least I do so.

## Sequential test

Piecewise constant hazard estimate with k = 5 change points:

$$\widehat{h}(t) = \begin{cases} 0.00386 & t \in [0, 3.6) \\ 0.01127 & t \in [3.6, 9.2) \\ 0.00405 & t \in [9.2, 12.0) \\ 0.01475 & t \in [12.0, 17.1) \\ 0.00404 & t \in [17.1, 19.8) \\ 0.02345 & t \ge 19.8 \end{cases}$$

Choose global  $\alpha = 0.05$ .

H <sub>0</sub>	$\alpha_k$	$\chi^2$ -quantile	$X_W$	<i>p</i> -value	reject H <sub>0</sub>	established change point
$\lambda_1 - \lambda_2$	0.0250000	5.02	15.52	< 0.0001	1	3.6
$\lambda_2 - \lambda_3$	0.0125000	6.24	9.90	0.0017	1	9.2
$\lambda_3 - \lambda_4$	0.0062500	7.48	10.91	0.00095	1	12.0
$\lambda_4 - \lambda_5$	0.0031250	8.73	7.19	0.0073	0	
$\lambda_5 - \lambda_6$	0.0015625	10.00	6.98	0.0083	0	

## Final hybrid estimate of S

 $\hat{\tau}$ : last established change point.

Estimate piecewise constant hazard assuming k = 1 with change point  $\hat{\tau}$ . Then:

$$\begin{split} \widehat{S}(t) &= \begin{cases} \widehat{S}_{\mathsf{KM}}(t) & 0 \leq t \leq \widehat{\tau} \\ \widehat{S}_{\mathsf{KM}}(t) \cdot \widehat{S}_{\mathsf{exponential beyond } \widehat{\tau}} & t > \widehat{\tau} \end{cases} \\ &= \begin{cases} \widehat{S}_{\mathsf{KM}}(t) & 0 \leq t \leq \widehat{\tau} \\ \widehat{S}_{\mathsf{KM}}(t) \cdot \exp\left(-\widehat{\lambda}(t-\tau)\right) & t > \widehat{\tau}. \end{cases} \end{split}$$

Plug  $\hat{S}$  in generic formulas to compute m.

### Confidence intervals width decreases with more events

Question: if we get more events - does width of confidence intervals decrease?

Setup: so far, used data with CCOD of 2012-11-21. Cut data back to CCOD of 2011-11-22 and 2012-05-21.

Results for prediction of second analysis, targeted #events = 248:

	CCOD = 2011/11/22	CCOD = 2012/05/21	CCOD = 2012/11/21
#events at CCOD	29	68	113
predicted timepoint	2014-05-14	2014-05-16	2013-10-13
95% confidence interval	[2013/03/14, 2015/02/15]	[2013/05/26, 2014/01/04]	[2013/08/10, 2014/03/04]
width of CI (months)	23.1	7.3	6.8

Dates get "earlier" with later cutoff date  $\Rightarrow$  initial underreporting of events.

### Confidence intervals width decreases with more events

Question: if we get more events - does width of confidence intervals decrease?

Setup: so far, used data with CCOD of 2012-11-21. Cut data back to CCOD of 2011-11-22 and 2012-05-21.

Results for prediction of final analysis, targeted #events = 370:

	CCOD = 2011/11/22	CCOD = 2012/05/21	CCOD = 2012/11/21
#events at CCOD	29	68	113
predicted timepoint	2015-10-18	2015-11-30	2014-07-31
95% confidence interval	[2013/12/09, 2018/06/13]	[2014/01/21, 2016/06/02]	[2014/02/28, 2015/06/16]
width of CI (months)	54.1	28.4	15.5

Dates get "earlier" with later cutoff date ⇒ initial underreporting of events.

# Doing now what patients need next

#### R version and packages used to generate these slides:

R version: R version 3.2.2 (2015-08-14)

Base packages: stats / graphics / grDevices / utils / datasets / methods / base

Other packages: reporttools / xtable / fitdistrplus / MASS / eventTrack / muhaz / survival

This document was generated on 2016-04-24 at 21:13:16.