

Adaptive designs for multiple time-to-event outcomes in Markovian multi-state models-

Moritz Fabian Danzer¹ Andreas Faldum¹ Rene Schmidt¹

¹Institute of Biostatistics and Clinical Research, University of Münster, Germany

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Project

Results of the DFG-funded project "Design and analysis of adaptive-sequential clinical trials with multiple, correlated time-to-event endpoints"

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- ▶ Location: Institute of Biostatistics and Clinical Research, University of Münster
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Aims and scope

Several correlated time-to-event endpoints of interest given

- ▶ Simultaneous assessment of efficacy and toxicity

Enable adaptive design changes which can be based on all of these endpoints

⇒ Extending one-sample methods from Danzer et al. (2022) to a randomized, multi-arm setting

H_0 : Joint distribution of selected time-to-event endpoints in different groups is the same

What's new?

Several time-to-event endpoints may be used to determine interim design changes!

General concern (cf. Bauer and Posch (2004)):

Consideration of surrogate (non-primary) endpoint in interim design changes may inflate type I error!

Existing solutions (patient-wise separation) lead to

- ▶ discarding of information or
- ▶ inevitable worst-case adjustment

Related methods

H_0 : (Marginal) distribution of all selected time-to-event endpoints in different groups is the same

Tests for multiple time-to-event endpoints in Wei and Lachin (1984):

- ▶ Simultaneous log-rank tests for all variables
- ▶ Covariance matrix via multivariate CLT

Group-sequential extension in Lin (1991):

- ▶ Extended use of multivariate CLT for multiple time points
- ▶ component-wise "independent increments structure"

Problem: increments are not independent across components

Compromising the type I error level

Scenario similar to example in Bauer and Posch (2004) to exploit method from Lin (1991) in adaptive design:

- ▶ Two time-to-event endpoints
- ▶ Second endpoint occurs exactly t years after occurrence of first endpoint
- ▶ Exact prediction of second component t years in advance possible
- ▶ Adaptation at interim analysis:
 - ▶ Stop recruitment if forecasted component will exceed decision bound
 - ▶ Else increase sample size

Challenge: Account for "shared information"

Notational conventions

Patients $i \in \{1, \dots, n\}$

Treatment indicators $Z_i \in \{0, 1\}$

States resp. (component) events $j \in \{1, \dots, k\}$

Composite events $E \subset \{1, \dots, k\}$

Event resp. hitting times T_i^E

Counting processes N_i^E

Occupation indicators $Y_i^j \in \{0, 1\}$

Occupation counts $Y^j = \sum_i Y_i^j$ and $Y^{Z=1j} = \sum_i Z_i Y_i^j$

Two notions of time:

- ▶ calendar time t
- ▶ time in trial s

From one- to two-sample methods

Patient-wise martingale

$$M_i^j(s) = N_i^j(s) - \int_0^s \lambda^j(u | \mathcal{F}_i(u)) du$$

as basis for one-sample test

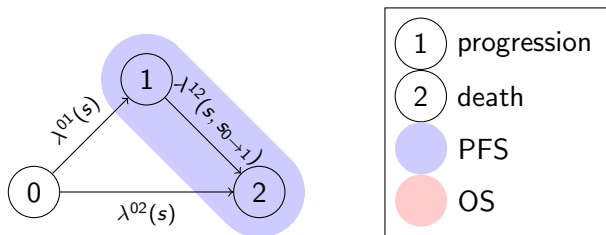
Advantages of one-sample method:

- ▶ Multiple dependence mechanisms applicable, e.g.
 - ▶ (Semi-)Markovianity of underlying multi-state model
 - ▶ Copula or frailty model
- ▶ Joint reference distribution specifies necessary correction (i.e. $\lambda^j(u | \mathcal{F}_i(u))$)

Impose similar assumptions on the data generating process in two-sample case!

Illness-death models

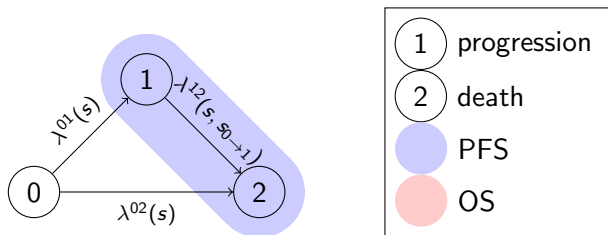
A multi-state model enables a more granular description of the course of disease



Events of clinical interest are given as "hitting times" of a set of nodes

Illness-death models

A multi-state model enables a more granular description of the course of disease

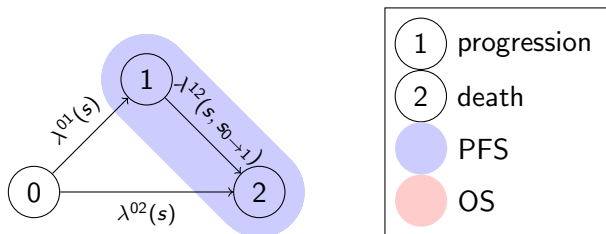


Markov model: $\lambda^{12}(s, s_{0 \rightarrow 1}) = \lambda_M^{12}(s)$

Semi-Markov model: $\lambda^{12}(s, s_{0 \rightarrow 1}) = \lambda_{SM}^{12}(s - s_{0 \rightarrow 1})$

Illness-death models

A multi-state model enables a more granular description of the course of disease



Markov model: $\lambda^{12}(s, s_{0 \rightarrow 1}) = \lambda_M^{12}(s)$

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Approach from Lin (1991)

Test of the hypothesis

$$H_{0,\text{marg}}: F_0^{\text{PFS}} = F_1^{\text{PFS}} \cap F_0^{\text{OS}} = F_1^{\text{OS}}$$

with the classical two-sample log-rank tests for the marginals

- ▶ $U^{\text{PFS}}(t) := \sum_{i=1}^n \int_0^t (Z_i - Y^{0,Z=1}(t,s)/Y^0(t,s)) N_i^{\text{PFS}}(t, ds)$
(\simeq martingale w.r.t. filtration generated by *PFS*-events)
- ▶ $U^{\text{OS}}(t) := \sum_{i=1}^n \int_0^t (Z_i - Y^{Z=1}(t,s)/Y(t,s)) N_i^{\text{OS}}(t, ds)$
(\simeq martingale w.r.t. filtration generated by *OS*-events)

and empirical estimation of covariance from asymptotically equivalent processes

Adjusted approach under Markov assumption

Test of the hypothesis

$$H_{0,\text{joint}}: F_0^{\text{PFS,OS}} = F_1^{\text{PFS,OS}} \quad (\text{Note: } H_{0,\text{joint}} \supset H_{0,\text{marg}})$$

with the multivariate process

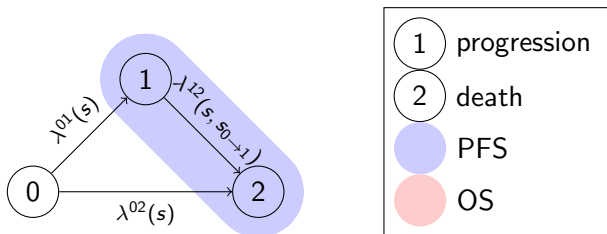
$$\mathbf{U}(t) := \begin{pmatrix} U^{\text{PFS}}(t) \\ U^{\text{OS}}(t) \end{pmatrix}$$

No correction for first component necessary, i.e.

$$U^{\text{PFS}}(t) := \sum_{i=1}^n \int_0^t \left(Z_i - \frac{Y^{0,Z=1}(t,s)}{Y^0(t,s)} \right) N_i^{\text{PFS}}(t, ds)$$

Second component requires consideration of origin of transition!

Distinguishing events by transitions



Decompose $Y_i(t, s) = Y_i^0(t, s) + Y_i^1(t, s)$ where

- ▶ $Y_i^0(t, s) = 1$ if at calendar time t patient i is known to be healthy at trial time s
- ▶ $Y_i^1(t, s) = 1$ if at calendar time t patient i is known to be ill at trial time s

Adjusted approach under Markov assumption (contd.)

Test of the hypothesis

$$H_{0,\text{joint}}: F_0^{\text{PFS,OS}} = F_1^{\text{PFS,OS}} \quad (\text{Note: } H_{0,\text{joint}} \supset H_{0,\text{marg}})$$

with the multivariate process

$$\mathbf{U}(t) := \begin{pmatrix} U^{\text{PFS}}(t) \\ U^{\text{OS}}(t) \end{pmatrix}$$

Adjusted second component:

$$U^{\text{OS}}(t) := \sum_{i=1}^n \int_0^t \left(Z_i - \mathbb{1}_{\{Y_i^0(t,s-) = 1\}} \cdot \frac{Y^{Z=1,0}(t,s)}{Y^0(t,s)} \right. \\ \left. - \mathbb{1}_{\{Y_i^1(t,s-) = 1\}} \cdot \frac{Y^{Z=1,1}(t,s)}{Y^1(t,s)} \right) N_i^{\text{OS}}(t, ds)$$

Asymptotics

Asymptotic equivalence:

$$\frac{1}{\sqrt{n}}(\mathbf{U}(t) - \tilde{\mathbf{U}}(t)) \xrightarrow{\mathbb{P}} \mathbf{0} \quad \forall t \geq 0$$

where $(\tilde{\mathbf{U}}(t))_{t \geq 0}$ is a martingale w.r.t. the filtration generated by PFS- and OS-events!

Central limit theorem yields:

- ▶ $\frac{1}{\sqrt{n}}(\tilde{\mathbf{U}}(t) - \tilde{\mathbf{U}}(s)) \xrightarrow{\mathcal{D}} \frac{1}{\sqrt{n}}(\tilde{\mathbf{U}}_{\infty}(t) - \tilde{\mathbf{U}}_{\infty}(s)) \sim \mathcal{N}(\mathbf{0}, \mathbf{V}(t) - \mathbf{V}(s))$
- ▶ asymptotically independent increments

Sequential testing procedure

For analysis dates $0 =: t_0 < t_1 < t_2$ we get standardized stagewise test statistics

$$\mathbf{z}_1 = \hat{\mathbf{C}}_1^{-1} \begin{pmatrix} U^{\text{PFS}}(t_1) \\ U^{\text{OS}}(t_1) \end{pmatrix} \text{ resp. } \mathbf{z}_2 = \hat{\mathbf{C}}_2^{-1} \begin{pmatrix} U^{\text{PFS}}(t_2) - U^{\text{PFS}}(t_1) \\ U^{\text{OS}}(t_2) - U^{\text{OS}}(t_1) \end{pmatrix}$$

which...

- ▶ ... asymptotically follow a bivariate standard normal distribution
- ▶ ... are asymptotically independent

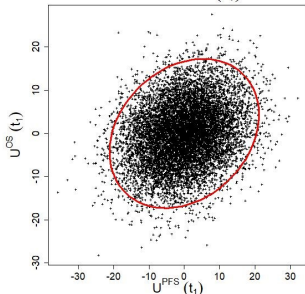
Agreement with asymptotic distribution is already observed for small sample sizes

Stagewise p-values

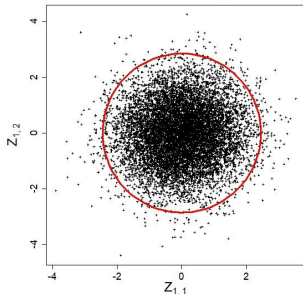
Obtain stagewise p-values by taking a norm of the stagewise test statistic:

L^2 -norm $|Z_1|_2^2$ and $|Z_1|_2^2$ are χ_2^2 -distributed (as in Wei and Lachin (1984))

Joint distribution of $U^{\text{PFS}}(t_1)$ and $U^{\text{OS}}(t_1)$



Joint distribution of $Z_{1,1}$ and $Z_{1,2}$



Can be combined e.g. by inverse normal method

Sample size calculation

Alternative hypotheses in terms of the transition hazards of the model

In the case of our example:

- ▶ Does the therapy prevent from progressions?
- ▶ Is it more or less toxic and thus leading to direct deaths?
- ▶ Is it a good salvage therapy?

Use e.g. transition-wise hazard ratios δ^{01} , δ^{02} and δ^{12}

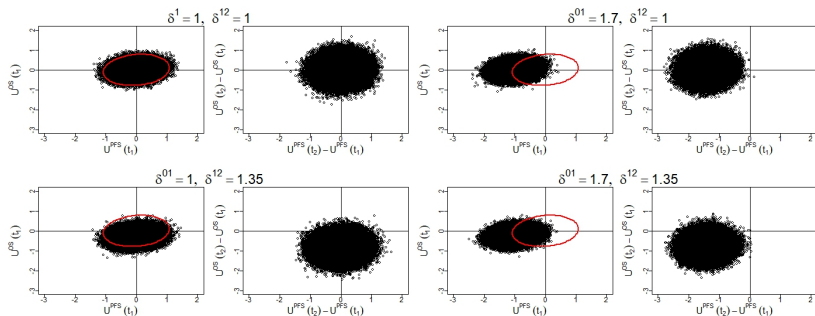
Hazard ratios imply drift function

$$n \cdot d(t; \delta^{01}, \delta^{02}, \delta^{12})$$

in calendar time t .

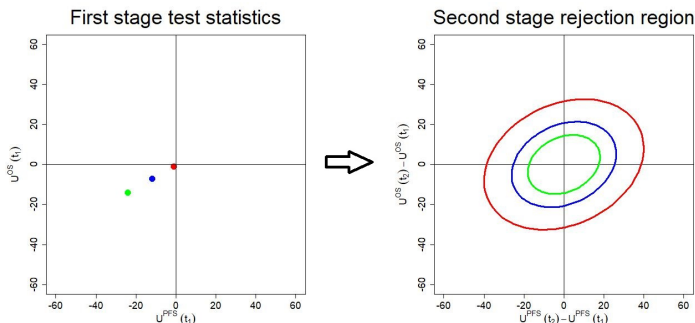
Drift under alternatives

Hazard ratios cause shift of increments of multivariate process \mathbf{U} in corresponding direction:



Adaptations in a two-stage design

First-stage p-value p_1 implies rejection region $\mathcal{R}_2(p_1)$ of second stage



Separate re-assessment of hazard ratios δ^{01} , δ^{02} and δ^{12}
⇒ Manipulation of external parameters of drift function yields required conditional power

Generalized setting

Non-recurrent multi-state model with state space $S = \{0, \dots, k\}$

Transition times $T^{\{1\}}, \dots, T^{\{k\}}$

Clinically relevant composite events $E_1, \dots, E_d \subset \{1, \dots, k\}$ with event times

$$T^{E_m} := \min_{j \in E_m} T^{\{j\}}$$

Component m of test statistic

$$U^{E_m}(t) := \sum_{i=1}^n \int_0^{t \wedge T_i^{E_m}} \left(Z_i - \sum_{l \notin E_m} \mathbb{1}_{\{Y_i^l(t, s-) = 1\}} \frac{Y^{Z=1, l}(t, s)}{Y^l(t, s)} \right) N_i^{E_m}(t, ds)$$

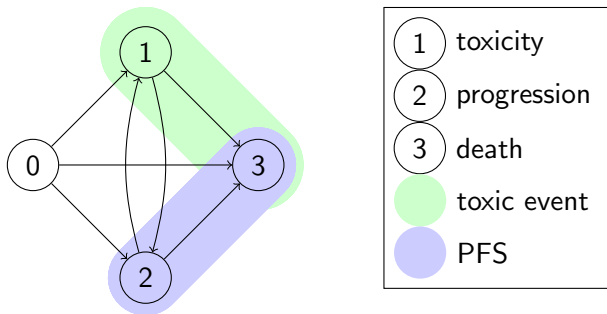
In particular, we generalize

- ▶ the standard log-rank test ($k = 1$, $d = 1$ and $E_1 = \{1\}$)
- ▶ the procedure of Lin (1991) in a competing risks setting

Further application

Simultaneous consideration of efficacy and toxicity

Accounting for death as a competing event



$k = 3$, $d = 2$, $E_1 = \{2, 3\}$ (PFS), $E_2 = \{1, 3\}$ (TFS)

Beyond the Markov assumption

Cox-Markov model:

- ▶ Transition intensity may depend on additional parameters
- ▶ E.g. $t^{\{l\}}$ be the time of transition into current state l

$$\lambda_j(u|\mathcal{F}_i(u)) = \lambda_{lj}(u) \exp(\beta_{lj} t^{\{l\}})$$

- ▶ Estimate and plug in regression parameters as in log-rank test with covariates

Semi-Markov model:

- ▶ Transition intensities given by

$$\lambda_j(u|\mathcal{F}_i(u)) = \lambda_{lj}(u - t^{\{l\}})$$

- ▶ Requires different definition of "at risk sets" and individual counting processes

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Estimation of covariance matrix \mathbf{V}

Non-zero elements of covariance matrix \Leftrightarrow events occur simultaneously

In our case: Deaths without prior illness, i.e.

$$\begin{aligned} & \hat{V}_{12}(t) \\ & := \widehat{\text{Cov}} \left(\frac{1}{\sqrt{n}} U^{\text{PFS}}(t), \frac{1}{\sqrt{n}} U^{\text{OS}}(t) \right) \\ & = \frac{1}{n} \sum_{i=1}^n \int_{[0,t]} \mathbb{1}_{\{Y_i^0(t,s-) = 1\}} \frac{Y^{Z=1,0}(t,s)}{Y^0(t,s)} \left(1 - \frac{Y^{Z=1,0}(t,s)}{Y^0(t,s)} \right) N_i^{\text{OS}}(t, ds) \end{aligned}$$

Compute lower triangular Cholesky factor $\hat{\mathbf{C}}$ of estimate $\hat{\mathbf{V}} = \hat{\mathbf{C}}\hat{\mathbf{C}}^T$

Simulation results: Empirical type I errors

4 different simulation scenarios:

1. time-homogeneous Markov model
2. time-inhomogeneous Markov model
3. Semi-Markov model
4. Frailty model (baseline intensities from 2. with joint Gamma frailty)

Combination of p -values from l^2 norm via inverse normal with equal weights and O'Brien-Fleming rejection bounds

Empirical type I errors ($\alpha = 0.05$, 200,000 runs):

Scenario	$n = 50$	$n = 100$	$n = 200$	$n = 500$	$n = 1000$
1	0.0581	0.0538	0.0517	0.0504	0.0496
2	0.0592	0.0550	0.0516	0.0502	0.0503
3	0.0527	0.0484	0.0472	0.0456	0.0458
4	0.0587	0.0547	0.0527	0.0507	0.0500

One-sided testing

Previously shown methods can only reject $H_{0,\text{joint}}$ in favour of

$$H_{1,\text{joint}} : F_0^{E_1, \dots, E_k} \neq F_1^{E_1, \dots, E_k}$$

Makes further interpretation of results difficult!

Possible choices for one-sided stagewise p -values:

- ▶ Least squares approaches (GLS or OLS, see e.g. Pocock et al. (1987) or Lachin (2014))
- ▶ Approximate Likelihood Ratio test (see Tang et al. (1989))
- ▶ Non-inferiority for all and superiority for at least one endpoint (see Tamhane and Logan (2004) and Perlman and Wu (2004))