

Generalized pairwise comparisons for right-censored time to event outcomes

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Clinical trials in oncology

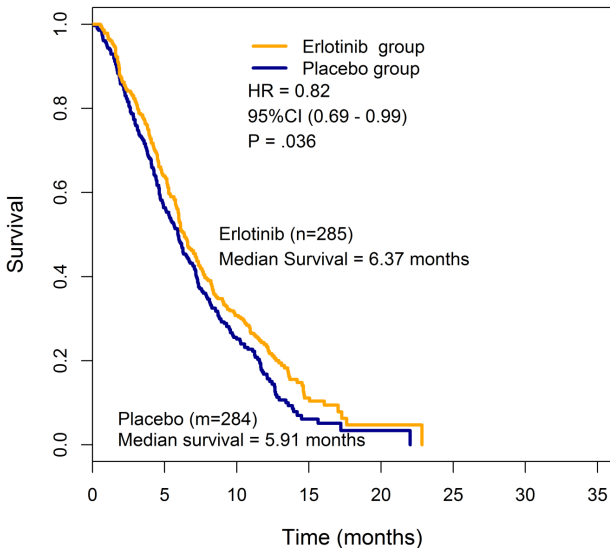
Efficacy/safety can reveal opposite effects of the treatment:

- longer survival
- serious but non-lethal adverse effects

However, efficacy and safety outcomes are usually analyzed and **reported independently**

- *efficacy*: using log-rank test
- *safety*: using Fisher's exact test

Example - Moore et al. (2007)



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Worst grade related adverse event	Erlotinib group (n=282)	Placebo group (n=280)
Grade 1	48 (17%)	69 (24.6%)
Grade 2	118 (41.8%)	89 (31.5%)
Grade 3	72 (25.5%)	47 (16.8%)
Grade 4	11 (3.9%)	6 (2.1%)
Grade 5	4 (1.4%)	3 (1.1%)

Clinical trials - handling multiple endpoints

I don't think there is a good objective approach.

What about a good subjective approach?

Patient preference

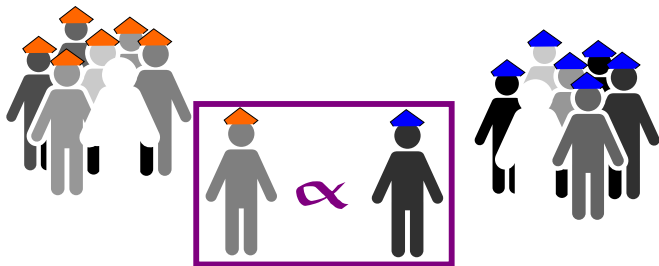
1. increase survival by at least 2 months
2. otherwise, least serious adverse events

Generalized pairwise comparisons (GPC)

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GPC - 1 endpoint

Consider N patients divided into two groups:

- experimental group: m patients with response $(x_i)_{i \in \{1, \dots, m\}}$
- control group: n patients with response $(y_i)_{i \in \{1, \dots, n\}}$

Denote by $\tau \in \mathbb{R}^{+*}$ the smallest difference in response that is clinically relevant.

Our parameter of interest is the **net benefit**:

$$\Delta = \mathbb{P}[X \geq Y + \tau] - \mathbb{P}[Y \geq X + \tau]$$

Point estimation in GPC

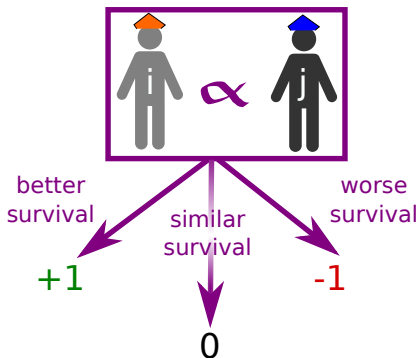
Defining $s_{ij} = \mathbb{1}_{x_i \geq y_j + \tau} - \mathbb{1}_{y_j \geq x_i + \tau}$ the score of the pair i, j :

$$\hat{\Delta} = \frac{1}{nm} \sum_{i=1}^n \sum_{j=1}^m s_{ij}$$

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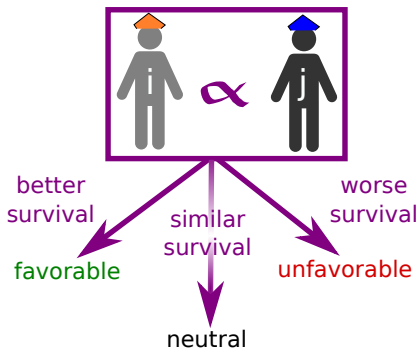
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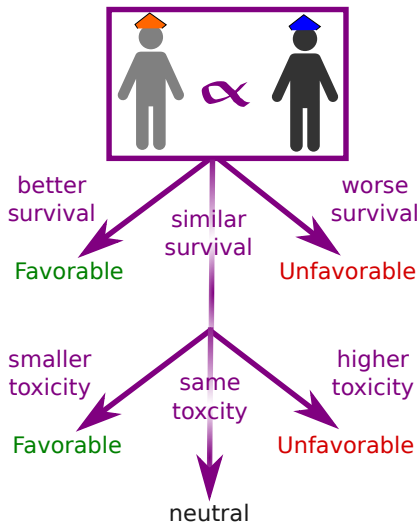
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Generalization to multiple endpoints



i.e. $s_{ij} = \phi(\mathbf{x}_i, \mathbf{y}_j, \tau) - \phi(\mathbf{x}_i, \mathbf{y}_j, \tau)$ where ϕ is a scoring rule.

GPC in presence of censoring

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We only observe:

- right-censored event times $(\tilde{x}_i)_{i \in \{1, \dots, m\}}$ and $(\tilde{y}_j)_{j \in \{1, \dots, n\}}$
- event type indicators $(\delta_{x,i})_{i \in \{1, \dots, m\}}$ and $(\delta_{y,j})_{j \in \{1, \dots, n\}}$.

How can we compute $s_{ij} = \mathbb{1}_{x_i \geq y_j + \tau} - \mathbb{1}_{y_j \geq x_i + \tau}$?

Gehan scoring rule

Set s_{ij} to 0 when the pair cannot be decidedly classified.

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If $\delta_{x,i} = 0$ and $\delta_{y,j} = 1$:

Example 1: $\tilde{x}_i \geq \tilde{y}_j + \tau$

- $s_{ij} = 1$

Example 2: $\tilde{x}_i < \tilde{y}_j + \tau$

- $s_{ij} = 0$: uninformative pair

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 uninformative pairs bias the estimation of Δ toward the null
(Buyse, 2008)

Peron scoring rule - Péron et al. (2018)

Compute

- $p_{ij}^f = \mathbb{P} [X \geq Y + \tau | X \geq \tilde{x}_i, X \geq \tilde{y}_j, \delta_{x,i}, \delta_{y,j}]$
- $p_{ij}^{uf} = \mathbb{P} [Y \geq X + \tau | X \geq \tilde{x}_i, X \geq \tilde{y}_j, \delta_{x,i}, \delta_{y,j}]$
- $s_{ij} = p_{ij}^f - p_{ij}^{uf}$

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If $\delta_{x,i} = 0$ and $\delta_{y,j} = 1$

- $p_{ij}^f = \min \left(\frac{S_X(\tilde{y}_j + \tau)}{S_X(\tilde{x}_i)}, 1 \right)$

where S_X is the survival in the experimental group.

Peron scoring rule - Péron et al. (2018)

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- $p_{ij}^f = \mathbb{P} [X \geq Y + \tau | X \geq \tilde{x}_i, X \geq \tilde{y}_j, \delta_{x,i}, \delta_{y,j}]$
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where S_x is the survival in the experimental group.

Example 1: $\tilde{x}_i \geq \tilde{y}_j + \tau$

- $s_{ij} = 1$

Example 2: $\mathbb{P} [X \geq \tilde{y}_j + \tau | X \geq \tilde{x}_i] = 1$

- $s_{ij} = 1$

Implementation

Package `BuyseTest` (available on CRAN):

- S_X and S_Y estimated using Kaplan-Meier

Example of code:

```
ff <- group ~ tte(survival, censoring = event, threshold = 2) +  
  cont(toxicity)  
  
e.Peron <- BuyseTest(formula = ff,  
                    data = dt.fol,  
                    scoring.rule = "Peron")
```

Implementation

Example of output:

```
summary(e.Gehan)
```

Generalized pairwise comparisons with 2 prioritized endpoints

[...]

> treatment groups: gemcitabine (control) vs. folfirinox (treatment)

[...]

endpoint	threshold	total	favorable	unfavorable	neutral	uninf	delta	Delta
survival	2	100.00	44.74	20.97	12.50	21.79	0.2377	0.2377
toxicity	1e-12	34.29	14.50	8.53	11.27	0.00	0.0597	0.2975

```
summary(e.Peron)
```

Generalized pairwise comparisons with 2 prioritized endpoints

[...]

endpoint	threshold	total	favorable	unfavorable	neutral	uninf	delta	Delta
survival	2	100.00	56.88	26.49	16.61	0.02	0.3039	0.3039
toxicity	1e-12	16.63	6.53	4.41	5.69	0.00	0.0212	0.3251

Limitations & perspectives

The Peron scoring rule requires a consistent estimator for the survival

- e.g. at $\tilde{y}_j + \tau$: may not be available
- remaining uninformative pairs

Ideas:

- lower and upper bound for p^f and p^{uf} (implemented)
- parametric model for S_X and S_Y
- inverse probability of censoring weights

Reference I

- Buyse, M. (2008). Reformulating the hazard ratio to enhance communication with clinical investigators. *Clinical Trials*, 5(6):641.
- Moore, M. J., Goldstein, D., Hamm, J., Figer, A., Hecht, J. R., Gallinger, S., Au, H. J., Murawa, P., Walde, D., Wolff, R. A., et al. (2007). Erlotinib plus gemcitabine compared with gemcitabine alone in patients with advanced pancreatic cancer: a phase iii trial of the national cancer institute of canada clinical trials group. *Journal of clinical oncology*, 25(15):1960–1966.
- Péron, J., Buyse, M., Ozenne, B., Roche, L., and Roy, P. (2018). An extension of generalized pairwise comparisons for prioritized outcomes in the presence of censoring. *Statistical methods in medical research*, 27(4):1230–1239.