SESSION 3: ADDITIONAL TWO-SAMPLE TESTS

Module 9: Survival Analysis in Clinical Trials Summer Institute in Statistics for Clinical Research University of Washington July, 2019

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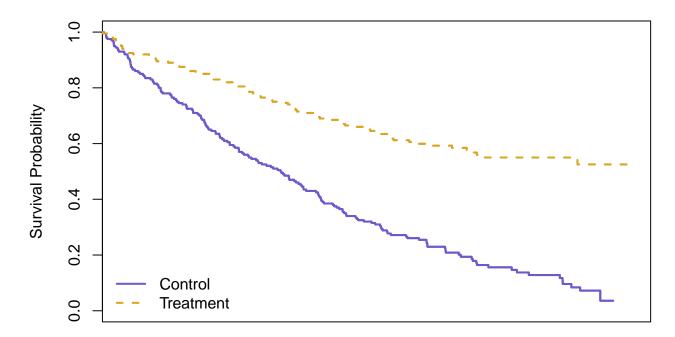
OUTLINE

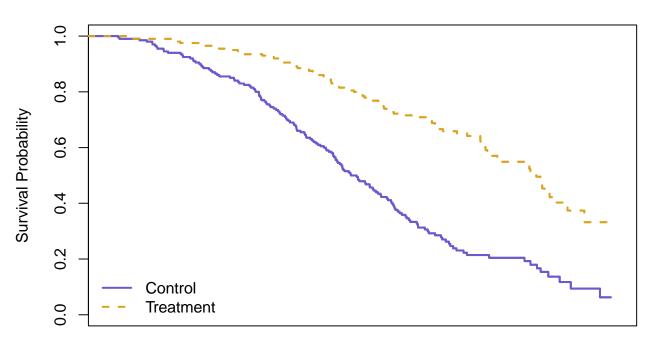
- Limitations of proportional hazards
- Other contrasts based on functionals of S(t)
 - S(t) at fixed time point
 - Quantiles (e.g. median)
 - Mean survival time
 - Restricted mean survival time
- Other metrics to describe the distance between survival curves
 - Weighted difference in S(t)
 - Maximum difference (Kolmogorov Smirnov)
 - Integrated squared difference (Cramér-von-Mises)

OUTLINE

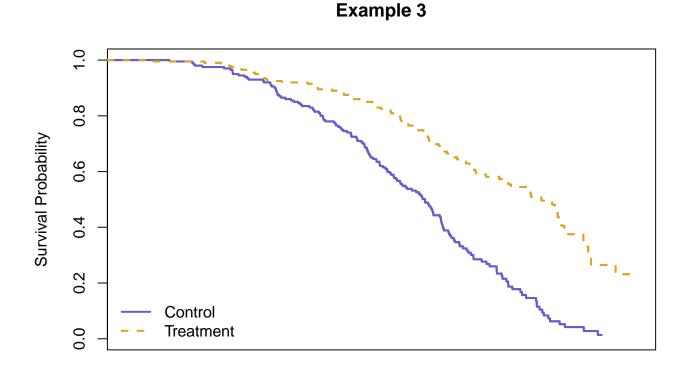
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Example 1



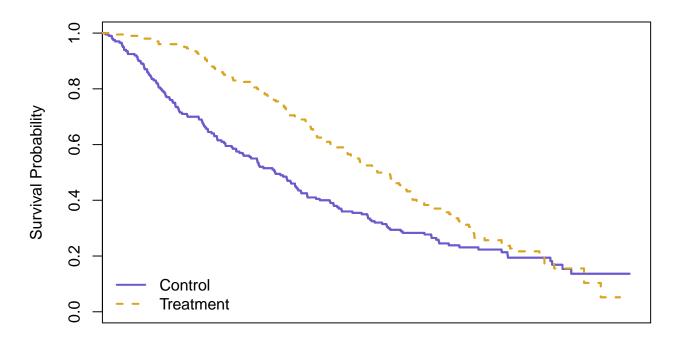


Example 2



Q: Which group has better survival in these examples?A:

Example 4

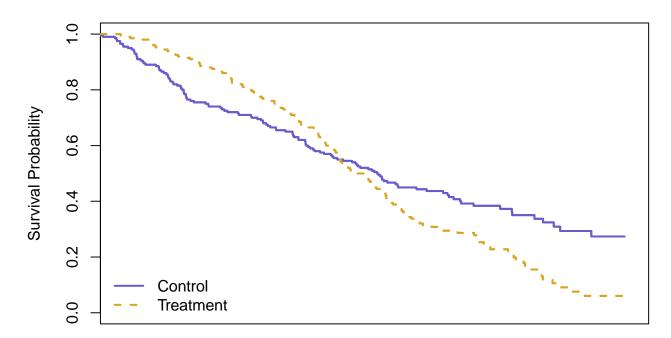


Q: Why does it appear the hazards are not proportional?

A:

Q: Which group has better survival?A:

Example 5



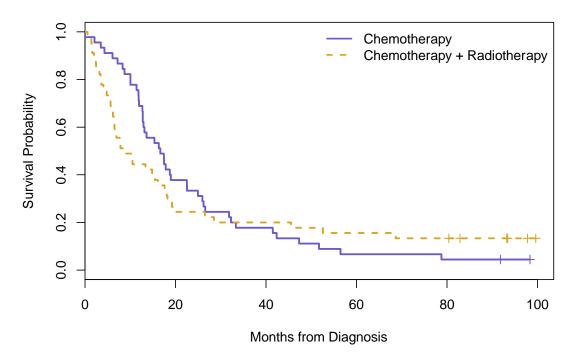
YOUR CHOICE

• Which group has better survival?

 You are a newly diagnosed patient. What would you want to know before choosing whether to take treatment?

REAL DATA

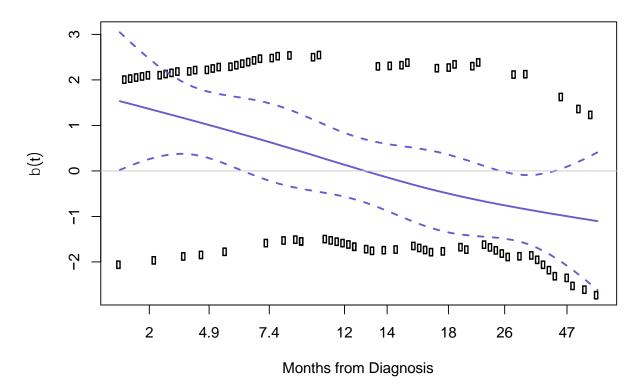
Gastric Cancer



Schein PS, Gastrointestinal Tumor Study Group. A comparison of combination chemotherapy and combined modality therapy for locally advanced gastric carcinoma. <u>Cancer</u>. 1982 May 1;49(9):1771–1777.

HAZARD RATIO

Log Hazard ratio: C+R to C only Based on Schoenfeld Residuals



Test of proportional hazards based on Schoenfeld residuals: P = 0.0003

HAZARD RATIO

	Hazard Ratio	95% CI	P-value
Chemotherapy	1.0 (reference)		
Chemotherapy + Radiotherapy	1.1	(0.72, 1.7)	.63

Assuming hazard ratio is constant...

CROSSING HAZARDS

When the proportional hazards assumption doesn't hold:

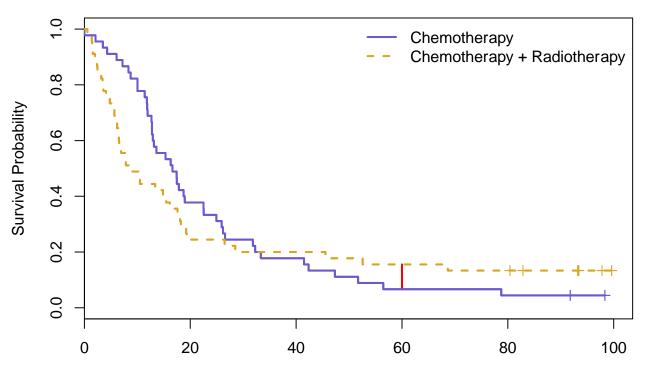
- Cox model will give weighted-average of time-specific hazard ratios (weights depend on censoring distribution)
- log rank test will test whether a weighted-average difference of hazards is zero
 - statistic numerator = $\sum_{j} \frac{n_{1j}n_{2j}}{(n_{1j}+n_{2j})} (\frac{d_{1j}}{n_{1j}} \frac{d_{2j}}{n_{2j}})$
 - More weight at earlier times when number at risk is larger
- May not be the quantity on which you want to base inference (estimation and testing)

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FIVE-YEAR SURVIVAL

Gastric Cancer



FIVE-YEAR SURVIVAL

- Compares only at a single point in time
- Ignores earlier survival differences, which may be important to some patients, given that in this example survival to 5 years in either group is low

S(t) AT A CHOSEN TIME t

- Choose time t for comparison at design stage.
- Compare $\hat{S}_1(t)$ to $\hat{S}_2(t)$ using

$$\frac{\hat{S}_1(t) - \hat{S}_2(t)}{\sqrt{\widehat{\text{var}}(\hat{S}_1(t)) + \widehat{\text{var}}(\hat{S}_2(t))}}$$

where $\widehat{var}(\widehat{S}_2(t))$ is computed using Greenwood's formula or another large-sample formula such as the one based on the complementary log-log of $\widehat{S}(t)$.

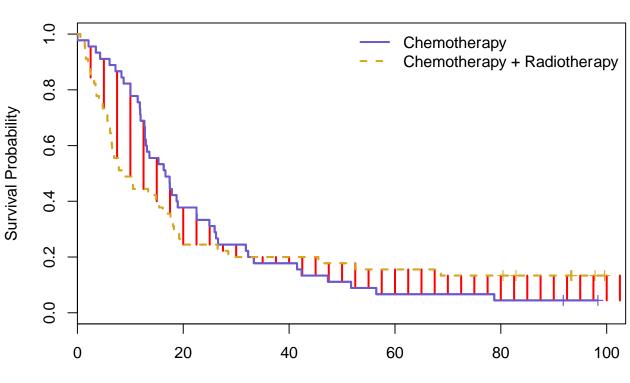
FIVE-YEAR SURVIVAL DIFFERENCE

Gastric Cancer

Difference	se(Difference)	Z Statistic	P-value
.0889	.0656	1.36	.1753

COMPARISON AT MORE THAN ONE TIME

Gastric Cancer



AVERAGE DIFFERENCES

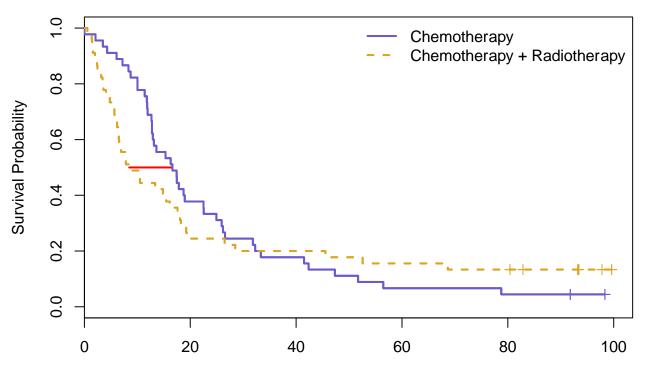
- Average difference between survival curves over time might be of interest
- In gastric cancer example, differences are of different signs at different times, so there would be cancellation
- Allows poorer survival after survival curves cross to detract from better survival before
- Interpretation?
- Also related to average quantile difference

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MEDIAN SURVIVAL

Gastric Cancer



MEDIAN SURVIVAL

- Compares only a single quantile
- Hard for some patients to interpret the difference in medians

MEDIAN TEST

Idea: Define \hat{M}_1 and \hat{M}_2 to be the median survival times in the two samples.

Then let the overall median survival time be defined by the weighted average.

$$\hat{M} = \frac{N_1}{N}\hat{M}_1 + \frac{N_2}{N}\hat{M}_2$$

A test of $H_0: M_1 = M_2$ can be performed by testing

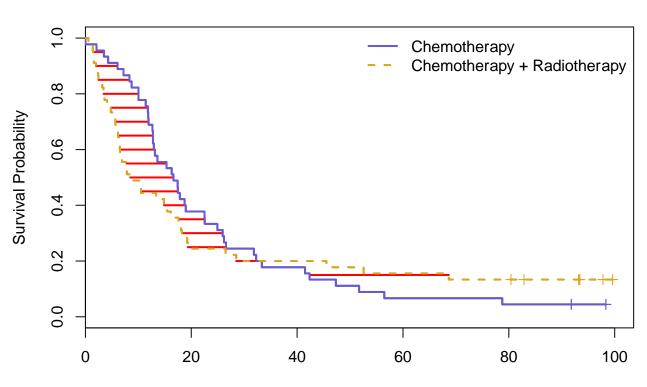
$$H_0: S_1(\hat{M}) = S_2(\hat{M})$$

Reference distribution based on joint asymptotic distribution of $(S_1(\hat{M}), S_2(\hat{M}))$.

Brookmeyer R, Crowley J. JASA 1982;77(378):433–440.

MORE THAN ONE QUANTILE

Gastric Cancer



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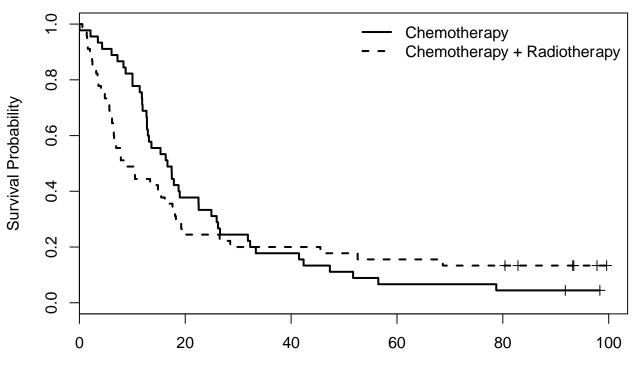
Useful Fact:
$$\int_0^\infty S(t)dt = E(T) = \int_0^\infty tf(t)dt$$

Proof: $\int_0^\infty S(t)dt = S(t)t|_0^\infty - \int_0^\infty t(-f(t))dt = \int_0^\infty tf(t)dt$

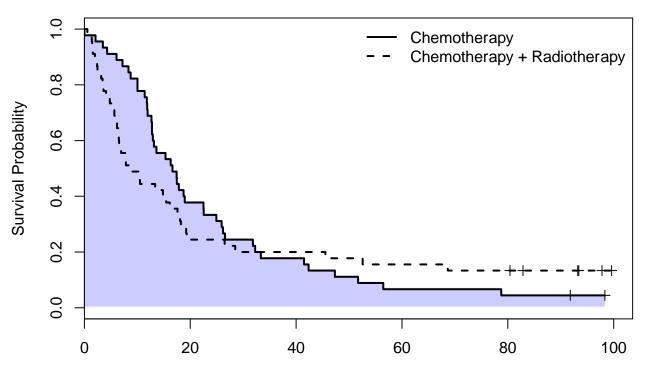
by integration by parts and

the fact that $E(T) < \infty \Rightarrow tS(t) \xrightarrow{t \to \infty} 0$.

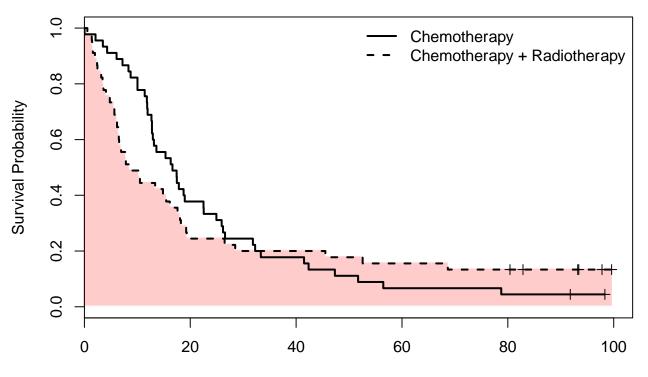
Gastric Cancer



Gastric Cancer



Gastric Cancer

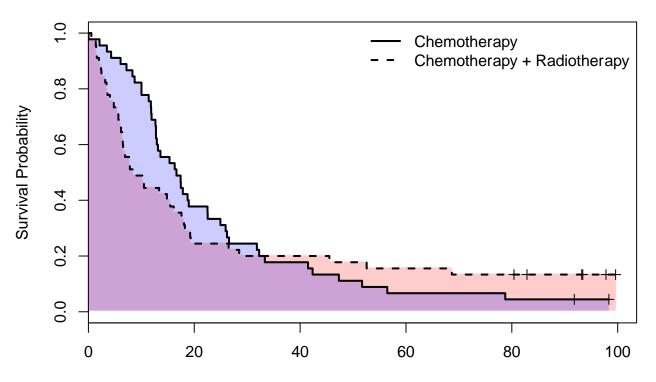


- Mean survival time $\mu = \int_0^\infty S(t) dt$
- Large sample (asymptotic) distribution proved by Gill in The Annals of Statistics. 1983;11(1):49–58.
- In finite samples, can be infinite if last time is a censoring
 - Integrate to last failure time only
 - Integrate to last observed time only

	Mean Survival*	SE
Chemotherapy	24.1 months	3.3 months
Chemotherapy + Radiotherapy	24.3 months	4.8 months

* Up to 99.6 months (last observed time in either group)

Gastric Cancer



MEAN SURVIVAL TIME DIFFERENCE

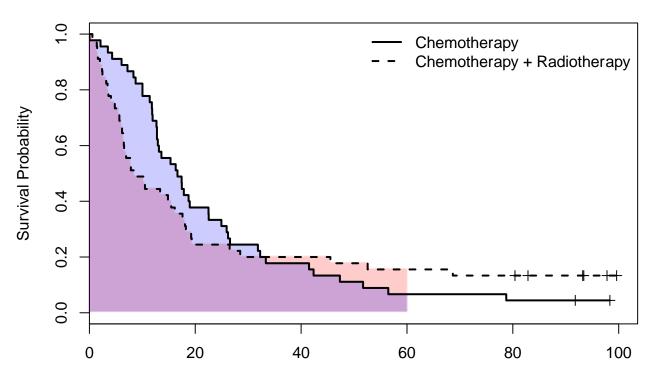
- Average of survival function differences over time
- Average of survival quantile differences over quantiles
- Allows cancellation
- Not much information at late times where few are at risk.
- <u>Infinite</u> estimate if KM curve doesn't descend to zero
- May want to truncate to a shorter interval, restricting to times where *S*(*t*) estimates are precise

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RESTRICTED MEAN SURVIVAL TIME

Gastric Cancer



Months from Diagnosis

MOTIVATION

- Clinically Interpretable ("over the next five years, patients like you live, on average, 13 months longer")
- Power/precision depends on length of observation time as well as number of events. Can achieve enough power/precision for meaningful comparisons with smaller studies.
- May be better measure for non-inferiority safety studies where events are rare. (Uno H et al. <u>Ann Intern Med</u> 2015; 21;163(2):127–134.) "Average number of days out of n event free."
- Excellent motivation when survival curves do not cross.

RESTRICTED MEAN SURVIVAL TIME

- Interpretation: average time lived in the interval $[0, \tau]$.
- Interpretation for differences: on average, the amount more time lived in $[0, \tau]$ on treatment A than on treatment B.
- Some asymptotically equivalent ways to estimate it:

$$- \hat{\mu} = \int_0^\tau \hat{S}(t) dt$$

- $\frac{1}{n} \sum_{i=1}^{n} \frac{d_i y_i}{\hat{s}_{c(y_i)}}$ where $\hat{S}_{c(y_i)}$ is the KM estimated survival function of the censoring distribution
- Using pseudo-observations based on the jackknife.

$$\hat{\mu} = \sum_{i=1}^{n} \hat{\mu}_i,$$

RESTRICTED MEAN SURVIVAL DIFFERENCE

• Standard estimation and testing:

$$-\hat{\mu}_k = \int_0^\tau \hat{S}_k(t) dt$$

-
$$\widehat{\operatorname{var}}(\hat{\mu}_k) = \sum_{j=1}^{J} \left[\int_{t_j}^{\tau} \hat{S}_{K}(t) dt \right]^2 \frac{D_{jk}}{N_{jk}(N_{jk} - D_{jk})}$$

– Compare test statistic:

$$T = \frac{\hat{\mu}_1 - \hat{\mu}_2}{\sqrt{\widehat{\operatorname{var}}(\hat{\mu}_1) + \widehat{\operatorname{var}}(\hat{\mu}_2)}}$$

to standard normal distribution (asymptotic).

RESTRICTED MEAN SURVIVAL TIME

$$E[\min(T,\tau)] = \widehat{E[Y]} = \int_0^\tau \widehat{S}(t) dt$$

Several approaches to variance estimation:

- Asymptotic
- Random perturbation resampling method (Tian L, Zhao L, Wei LJ. Predicting the restricted mean event time with the subject's baseline covariates in survival analysis. Biostat. 2014 Apr 1;15(2):222–233.)
- Variance of pseudo observations

PSEUDO OBSERVATIONS

- There are a number of other less direct ways to estimate $\mu_k = \int_0^{\tau} \hat{S}_k(t) dt$ that make generalizing to regression models easier.
- One appealing method uses pseudo-observations based on the jackknife.
 - Group means computed in the usual way from pseudoobservations
 - Standard errors computed from pseudo-observations in the usual way.
 - Test statistic based on two-sample t-test (unequal variances) with pseudo-observations.

PSEUDO OBSERVATIONS

Estimation of μ using pseudo-observations based on the jackknife.

$$\hat{\mu} = \sum_{i=1}^{n} \hat{\mu}_i,$$

where $\hat{\mu}_{i} = n\hat{\mu} - (n-1)\hat{\mu}_{-i}$.

- $\hat{\mu}$ is computed by the first method from the pooled sample, and
- $\hat{\mu}_{-i}$ is computed the same way but leaving out the *i*th observation.
- Andersen et al. Lifetime Data Anal. 2004;10(4):335–350.
- Functions available in Stata, R and SAS.

RESTRICTED MEAN SURVIVAL TIME

	Restricted Mean Survival (2000 days)	SE
Chemotherapy	673	77.8
Chemotherapy + Radiotherapy	599	101.1

Comparison Method	P-value
Asymptotic	.560
Pseudo observations	.566

DESIGN AND INFERENCE ISSUES

- Not much information / precision available at late times when few subjects are at risk
 - If a restricted mean over an interval [0, τ] is of interest, important to follow subjects enough longer than τ to have an adequate number still at risk at time τ.

EXAMPLE

- Schermerhorn et al. (2015) compared survival in a matched cohort of 39,966 pairs of Medicare patients who received either endovascular or open repair of an abdominal aortic aneurism.
 - Perioperative mortality and complication rates were higher in those given open repair: 5.2% vs 1.6% for mortality and 12.9% vs 3.8%
 - The estimated hazard ratio for death comparing endovascular to open repair varied over time:
 - HR = .32 (95% CI: .29 .35) for the first 30 days
 - HR = .64 (95% CI: .58 -.71) for 30 90 days
 - HR = 1.17 (95% C: | 1.13 1.21) for 90 days 4 years
 - HR = 1.05 (95% CI: 1.00 1.09) after 4 year.

Schermerhorn ML, Buck DB, O'Malley AJ et al. NEJM 2015 Jul 23;373(4):328–338.

EXAMPLE

- Because of non-proportional hazards they estimated differences in restricted mean survival using the pseudo observation approach of Andersen et al with the matchedpair data.
 - Over the first 4 years, the endovascular group lived an average of 12.4 days longer (95% Cl 9.0 15.6)
 - Over the first 7 years, the endovascular group lived an average of 8.2 days longer (95% CI: 1.5-14.4)
 - The authors concluded that the advantage of endovascular repair persisted to 7 years.
- The pseudo-observation approach makes it easy to accommodate the matched design.

SCREENING TRIAL

- 202,546 women 50-72 years of age, England, Wales, Northern Ireland
- Randomized to one of three arms in 1:1:2 ratio between June 1, 2001 and Oct 21, 2005.
 - Annual multimodal screening (serum CA 125 + algorithm)
 - Annual transvaginal ultrasound
 - No screening
- Screening ended Dec 31, 2011.
- Not blinded
- Primary outcome: death from ovarian cancer (by end of 2014) Jacobs IJ, Menon U, Ryan A, et al. (2016) The Lancet. 387(10022):945–956.

OVARIAN CANCER SCREENING TRIAL

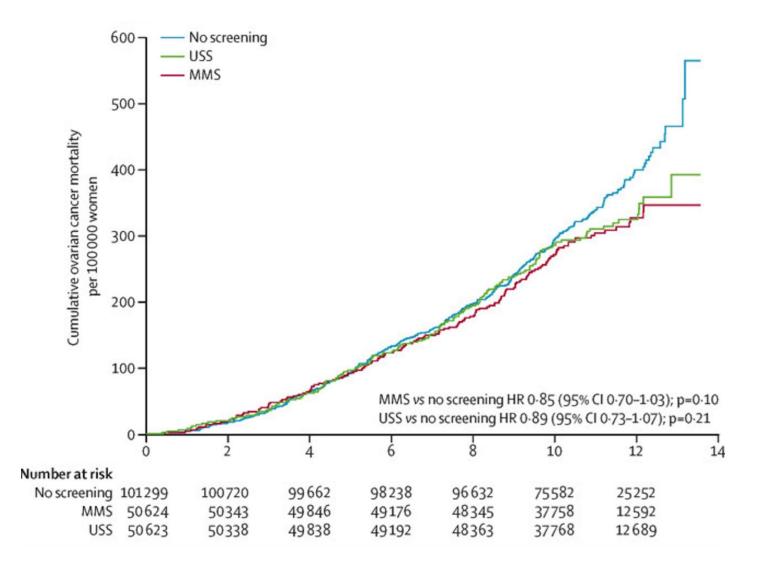
- Primary analysis: Cox regression (proportional hazards)
 - MMS vs. no screening: Mortality reduction =

(1 - HR)100 = 15% (95% CI: -1% - 33%) P = .10

- USS vs. no screening: Mortality reduction =

(1 – HR) 100 = 11% (95% CI: -7% - 27%) P = .21

OVARIAN CANCER SCREENING TRIAL



OVARIAN CANCER SCREENING TRIAL

- Secondary analyses, excluding prevalent cases:
- Post-hoc Weighted* logrank test:
 - MMS mortality reduction = 22% (3-38%) P = .023
 - USS mortality reduction = 20% (0 35%) P = .049

* by pooled cumulative mortality

SURVEY

- Trinquart et al. (JCO. 2016; 20; 34(15):1813–1819) surveyed oncology RCTs reported in five journals during the last six months of 2014.
 - 54 trials, 33,212 patients
 - Reconstructed data
 - 13 (24%) had evidence of non-proportional hazards
 - Compared tests based on HR treatment effect with tests based on ratio and difference of RMST.
 - Statistical significance in agreement between HRbased and RMST-based tests for 53 out of 54 trials.

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ANOTHER OPTION: METRICS

- Tests based on detecting consistent differences between survival curves or hazard across time lose power when the hazards or survival curves cross.
- Weighting can focus on a time period when direction of differences is consistent.
- Other metrics can measure distance between survival functions or hazard functions in a way that does not require the direction of differences to be consistent
- Tests based on them can have more power to detect a difference when survival functions or hazards cross. (Need to think about whether the difference detected is of interest.)

METRICS

 Weighted difference between Kaplan-Meier estimates (Pepe MS, Fleming TR. <u>Biometrics</u>. 1989;497–507).
Choose weights based on toxicity profile, for example.

$$\sqrt{\frac{n_1n_2}{n}} \int_0^\infty \hat{w}(t) [\hat{S}_2(t) - \hat{S}_1(t)] dt$$

 Weighted difference between Kaplan-Meier estimates with adaptively chosen weights (Uno et al. <u>Statistics in</u> <u>Medicine</u>, 2015; 34(28):3680–3695).

Hard to know what parameter is being compared.

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METRICS

• Supremum: Tests based on the supremum of a difference of cumulative weighted hazard functions over [0, *t_m*]:

$$\sup_{t \in [0, t_m]} \sum_{i: t_i < t} W_i \frac{n_{1i} n_{2i}}{n_{1i} + n_{2i}} (\frac{d_{1i}}{n_{1i}} - \frac{d_{1i}}{n_{1i}})$$

- Gill, R.D. (1980). Censoring and stochastic integrals. Math. Centre Tracts 124, Mathematisch Centrum Amsterdam.
- Fleming TR, O'Fallon JR, O'Brien PC, Harrington DP. Biometrics. 1980;36(4):607–625.
- Fleming TR, Harrington DP, O'Sullivan M. JASA. 1987;82(397):312–320.

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METRICS

• l^2 : Tests based on the integrated squared difference of survival or cumulative hazard functions over $[0, t_m]$:

$$\sum_{t_i:t_i \leq t_m, \delta_i = 1} (\hat{S}_2(t_i) - \hat{S}_1(t_i))^2 d(-\hat{S}(t_i))$$

or

$$\sum_{t_i:t_i \le t_m, \delta_i = 1} ((\hat{S}_2(t_i) - \hat{S}_1(t_i))W_i)^2 d(\hat{H}(t_i))$$

where the weight function W_i and H are functions of the asymptotic covariance of the cumulative hazard estimator at different times.

- Koziol Biom. J. 1978;20(6):603–608.
- Koziol, Yuh . Biom. J. 1982;24(8):743-750.
- Schumacher. International Statistical Review 1984;52(3):263–281.

ISSUE

- Hard to think of a good scientific hypothesis that specifies which of these metrics and associated tests is consistent with the hypothesis.
- Large temptation to choose the type of test <u>after</u> looking at the data and noticing crossing hazards or crossing survival functions in the search for a powerful test.
- Scientific hypotheses more likely to be consistent with a difference between functionals of the survival function S(t).

OTHER POSSIBILITIES

- Test based on Cox model with time-dependent interaction terms (time-dependent coefficients). Some on this tomorrow.
- Test based on specific richer model for how hazard ratio depends on time (Yang S, Prentice R. Biometrika. 2005;92(1):1–17).

$$\frac{\lambda_2(t)}{\lambda_1(t)} = \frac{\theta_0 \theta_\infty}{\theta_0 + (\theta_\infty - \theta_0)S_1(t)}$$

parameterized by θ_0 , the limiting hazard ratio as $t \rightarrow 0$ and θ_{∞} , the limiting hazard ratio as $t \rightarrow \infty$

TO WATCH OUT FOR

- Base quantity to be compared (weighted sum for logrank, time, quantile or restricted mean) on what would be meaningful in the context of the trial.
- Important to choose it <u>before</u> looking at the data.