

# Bivariate Subgroup Analysis for Benefit-Harm Assessment

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# Subgroup Analysis and Heterogeneity in Benefit/Harm

- ▶ In clinical trials, subgroup analyses are regularly performed to investigate the consistency of treatment effect across patient subgroups.
- ▶ While subgroup analyses are frequently used for looking at heterogeneity in treatment effectiveness (HTE), heterogeneity in treatment safety is seldom examined.
- ▶ Even when heterogeneity in treatment-related adverse events (HTAE) is addressed, the subgroup analysis for safety is typically performed separately from the HTE analysis.

## Subgroup Analysis and Heterogeneity in Risk/Benefit

- ▶ From a patient-centered perspective, separate subgroup analyses of HTAE and HTE ignore potentially important relationships between primary and safety outcomes.
- ▶ For example, suppose we have a binary primary event (PE) and binary adverse event (AE) whose joint distribution is given by

	Treatment 1		Treatment 2	
	AE	No AE	AE	No AE
PE	0.1	0	0	0.1
No PE	0.3	0.6	0.4	0.5

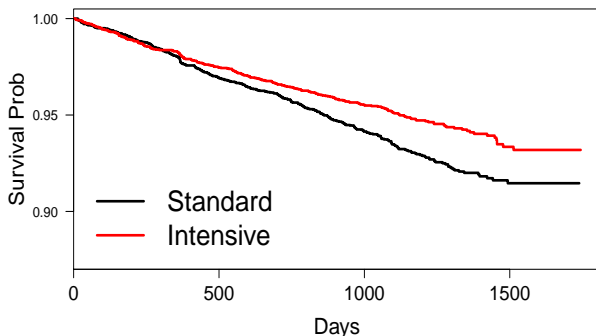
- ▶  $Pr(PE|Trt = 1) = 0.1$        $Pr(PE|Trt = 2) = 0.1$
- ▶  $Pr(AE|Trt = 1) = 0.4$        $Pr(AE|Trt = 2) = 0.4$

# Subgroup Analysis and Heterogeneity in Risk/Benefit

- ▶ Only comparing the marginal distribution of a PE and AE can hide important risk-benefit considerations.
- ▶ For greater relevance to patients, subgroups analysis should assess variation in changes to patients risk-benefit profiles.
- ▶ A “truly” bivariate subgroup analyses would allow us to explore joint patient outcomes and heterogeneity of treatment impact.

# SPRINT Trial

- ▶ The systolic blood pressure intervention (SPRINT) trial (N=9,361) investigated the effect of using a more stringent blood pressure target:  $\leq 120$  mm Hg (intensive) versus  $\leq 140$  mm Hg (standard).
- ▶ At the conclusion of the trial, 243 PEs were observed in the intensive treatment arm and 319 PEs were observed in the standard treatment arm.



## SPRINT Trial

- ▶ Joint counts of PEs and treatment-related serious adverse events (SAEs) in the SPRINT trial.

	Standard Treatment		Intensive Treatment	
	SAE	No SAE	SAE	No SAE
PE	18	301	30	213
No PE	100	4264	190	4245

## Data for a Bivariate Subgroup Analysis

- ▶ **Primary Event:**  $T_i$  - time to the primary event
- ▶ **Safety Event:**  $W_i$  - a binary outcome (an indicator of whether or not patient experienced at least one AE)
- ▶  $Y_i = \min\{T_i, C_i\}$ : duration of follow-up  
 $\delta_i = \mathbf{1}\{T_i \leq C_i\}$ : event indicator
- ▶  $A_i$  - treatment arm assignment
- ▶  $G_i$  - indicator of subgroup membership
- ▶ Subgroup memberships  $G_i$  are for a “fully stratified” subgroup analysis as opposed to the more typical univariate “one variable at-a-time” subgroup analysis.

## Joint model for survival and binary outcomes

- ▶ **Model Parameters:**

$$\alpha, \{\lambda_{awg}\}, \{p_{ag}\}; \quad a = 0, 1; \quad w = 0, 1; \quad g = 1, \dots, G.$$

- ▶ We want to specify the joint distribution of  $(T_i, W_i)$  conditional on treatment arm assignment  $A_i$  and subgroup membership  $G_i$ .
- ▶ This is done by assuming that

$$\begin{aligned} T_i | A_i = a, W_i = w, G_i = g &\sim \text{Weibull}(\alpha, \lambda_{awg}) \\ W_i | A_i = a, G_i = g &\sim \text{Bernoulli}(p_{ag}). \end{aligned}$$

- ▶ The joint distribution  $(T_i, W_i) | A_i = a, G_i = g$  depends on the parameters  $(\alpha, \lambda_{a0g}, \lambda_{a1g}, p_{ag})$ .



## Summary Statistics for the Exponential Model

- ▶ When  $\alpha = 1$ , the time-to-event  $T_i$  follows an exponential distribution.
- ▶ Moreover, when  $\alpha = 1$ , the likelihood only depends on the following summary statistics

$$D_{avg} = \sum_{i=1}^n \delta_i I(A_i = a) I(W_i = w) I(G_i = g)$$

$$U_{avg} = \sum_{i=1}^n Y_i I(A_i = a) I(W_i = w) I(G_i = g)$$

$$V_{ag} = \sum_{i=1}^n W_i I(A_i = a) I(G_i = g)$$

# Summary Statistics for the SPRINT Trial

CKD	Subgroup		Standard Treatment			Intensive Treatment			$N_g$
	Age	Sex	$(D_{01g}, U_{01g})$	$(D_{00g}, U_{00g})$	$V_{0g}$	$(D_{11g}, U_{11g})$	$(D_{10g}, U_{10g})$	$V_{1g}$	
No	< 75	Male	(3, 92.0)	(96, 5546.1)	29	(10, 174.1)	(54, 5446.5)	61	3528
Yes	< 75	Male	(3, 48.4)	(28, 1364.7)	16	(5, 103.2)	(25, 1227.7)	32	858
No	$\geq$ 75	Male	(1, 19.6)	(46, 1277.5)	8	(1, 57.7)	(26, 1265.9)	19	913
Yes	$\geq$ 75	Male	(6, 40.0)	(47, 977.7)	15	(7, 88.3)	(38, 974.8)	31	730
No	< 75	Female	(0, 40.1)	(31, 2641.9)	13	(0, 67.5)	(25, 2705.4)	20	1706
Yes	< 75	Female	(2, 42.4)	(12, 948.7)	14	(4, 50.2)	(16, 1000.3)	16	617
No	$\geq$ 75	Female	(0, 37.8)	(16, 835.1)	12	(1, 60.9)	(18, 778.2)	20	568
Yes	$\geq$ 75	Female	(3, 30.8)	(25, 612.1)	11	(2, 66.4)	(11, 634.9)	21	441

## Modeling Subgroup Parameters (Saturated Model)

- ▶ The distribution of summary statistics  $(D_{awg}, U_{awg}, V_{ag})$  depends on hazard rate parameters  $\lambda_{awg}$  and AE probabilities  $p_{ag}$ .
- ▶ Assume that

$$\log(\lambda_{awg}) = \mathbf{x}_g^T \boldsymbol{\beta}_{aw} \quad \text{and} \quad \text{logit}(p_{ag}) = \mathbf{z}_g^T \boldsymbol{\gamma}_a$$

- ▶ For example, in a **saturated model**, we have

$$\log(\lambda_{awg}) = \beta_{aw,g} \quad \text{and} \quad \text{logit}(p_{ag}) = \gamma_{a,g},$$

where  $\boldsymbol{\beta}_{aw} = (\beta_{aw,1}, \dots, \beta_{aw,G})^T$  and  $\boldsymbol{\gamma}_a = (\gamma_{a,1}, \dots, \gamma_{a,G})^T$ .

## Modeling Subgroup Parameters (Additive Model)

- ▶ In the saturated model, the  $\lambda_{avg}$  and  $\rho_{ag}$  are treated separately with no additional information used to indicate relationships among the subgroups.
- ▶ Subgroups that share much of their characteristics are treated the same as subgroups that are quite different.
- ▶ Some regression structure linking the parameters  $\lambda_{avg}$ ,  $\rho_{ag}$  can induce more sensible correlation.

## Modeling Subgroup Parameters (Additive Model)

- ▶ In the **additive model**,  $\lambda_{awg}$  and  $p_{ag}$  are determined additively from the variables comprising subgroup  $g$ .
- ▶ For example, if we have four subgroups arising from each combination of the variables age (young/old) and smoking behavior (smoker/non-smoker)

(Smoker/Young)	$\log(\lambda_{aw1}) = \beta_{aw,1}$
(Smoker/Old)	$\log(\lambda_{aw2}) = \beta_{aw,1} + \beta_{aw,2}$
(Non-Smoker/Young)	$\log(\lambda_{aw3}) = \beta_{aw,1} + \beta_{aw,3}$
(Non-Smoker/Old)	$\log(\lambda_{aw4}) = \beta_{aw,1} + \beta_{aw,2} + \beta_{aw,3}$

- ▶ We could also use a regression that includes higher-order interactions or a model that makes additional assumptions about how hazards can vary across subgroups:  
 $\lambda_{a1g}/\lambda_{a0g} = \phi$  (proportional hazards)

## Specifying the Prior (Saturated Model)

- ▶ In the saturated model, we assumed

$$\log(\lambda_{awg}) = \beta_{aw,g} \quad \text{for } g = 1, \dots, G.$$

- ▶ For each treatment separately, assume the following for the  $\beta_{aw,g}$

$$\begin{bmatrix} \beta_{a0,1} \\ \beta_{a1,1} \end{bmatrix}, \dots, \begin{bmatrix} \beta_{a0,G} \\ \beta_{a1,G} \end{bmatrix} \Bigg| \boldsymbol{\mu}_a, \boldsymbol{\tau}_a \sim \text{Normal} \left( \begin{bmatrix} \mu_{a0} \\ \mu_{a1} \end{bmatrix}, \begin{bmatrix} \tau_{a0}^2 & 0 \\ 0 & \tau_{a1}^2 \end{bmatrix} \right)$$

- ▶ Place a proper, but “vague” prior on the joint distribution of  $\boldsymbol{\mu}_a$  while allowing for user-specified prior correlation.

## Specifying the Prior (Saturated Model)

$$\log(\tau_a) \sim \text{Normal} \left( \begin{bmatrix} \log(1/2) \\ \log(1/2) \end{bmatrix}, \begin{bmatrix} \sigma_{\tau,a}^2 & \sigma_{\tau,a}^2 \rho_{\tau,a} \\ \sigma_{\tau,a}^2 \rho_{\tau,a} & \sigma_{\tau,a}^2 \end{bmatrix} \right)$$

- ▶ For variance components  $\tau_a = (\tau_{a0}, \tau_{a1})$ , use a “quasi informative” or “weakly informative” prior as a default.
- ▶ Place most prior mass on plausible variation across subgroups.
- ▶ Consider the hazard ratio  $\lambda_{awj}/\lambda_{awk}$  between two subgroups.

$$Pr \left\{ \frac{1}{4} \leq \frac{\lambda_{awj}}{\lambda_{awk}} \leq 4 \mid \tau_a \right\} \geq 0.95 \quad \text{whenever} \quad \tau_{aw} \leq 1/2$$

- ▶ Prior median of  $\tau_{a0}$  and  $\tau_{a1}$  is 1/2. Choose  $\sigma_{\tau,a}^2$  so that

$$Pr\{\tau_a \leq 2\} \approx 0.95$$

- ▶ Prior for correlation:  $\rho_{\tau,a} \sim \text{Uniform}(-1, 1)$ .

# Targets of Inference

- ▶ For each subgroup, we want to target some parameter (or a collection of parameters) which captures important changes in the joint distribution  $(T_i, W_i)$  from treatment  $A_i = 0$  to  $A_i = 1$ .
- ▶ With our Bayesian setup, this is easy for any chosen target because we can just transform the posterior draws of parameters  $\lambda_{awg}$  and  $p_{ag}$  as needed.
- ▶ In our implementation, we consider the following targets:
  - (1) Heterogeneity in joint binary outcomes
  - (2) Heterogeneity in utility gained
  - (3) Heterogeneity in probability of outcome improvement



## Heterogeneity in Utility Gain/Loss

- ▶ Think of the composite score for patient  $i$

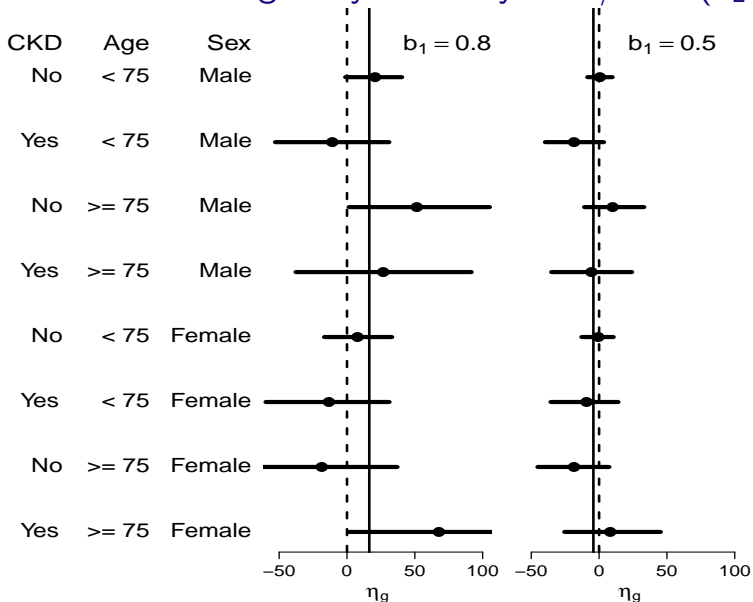
$$H_i = b_1 W_i \min\{T_i, \tau\} + b_2(1 - W_i) \min\{T_i, \tau\}$$

for weights  $b_2 > b_1 > 0$ .

- ▶ Patient  $i$  receives a “score” of  $b_1 T_i$  if surviving to time  $T_i < \tau$  while experience an AE some time in  $(0, T_i)$ .
- ▶ Patient  $i$  receives a “score” of  $b_2 T_i$  if surviving to time  $T_i < \tau$  while never experiencing an AE.
- ▶ For each subgroup  $g$ , the parameter of interest is the expected difference in the composite score

$$\eta_g = E[H_i | A_i = 1, G_i = g] - E[H_i | A_i = 0, G_i = g]$$

# SPRINT Trial: Heterogeneity in Utility Gain/Loss ( $b_2 = 1$ )



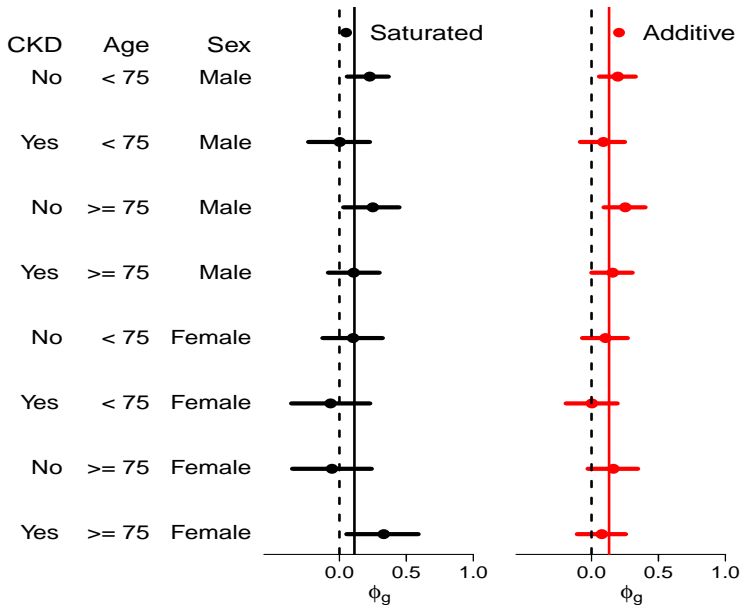
## Heterogeneity in Probability of Outcome Improvement (Assuming $A_i = 1$ and $A_j = 0$ )

	Outcome	Preferred Treatment
$T_i > T_j(1 + \delta)$	$W_i = 1, W_j = 0$	$A = 1$
$T_i \leq T_j(1 + \delta)$	$W_i = 1, W_j = 0$	$A = 0$
$T_j > T_i(1 + \delta)$	$W_i = 0, W_j = 1$	$A = 0$
$T_j \leq T_i(1 + \delta)$	$W_i = 0, W_j = 1$	$A = 1$
$T_i > T_j$	$W_i = 1, W_j = 1$	$A = 1$
$T_i > T_j$	$W_i = 0, W_j = 0$	$A = 1$
$T_i \leq T_j$	$W_i = 1, W_j = 1$	$A = 0$
$T_i \leq T_j$	$W_i = 0, W_j = 0$	$A = 0$

The subgroup-specific parameters of interest are

$$\phi_g = 2 \times \Pr \left\{ \text{outcome } i > \text{outcome } j \mid A_i = 1, A_j = 0, G_i = g, G_j = g \right\} - 1$$

# SPRINT Trial: Outcome Improvement Measure



## Summary

- ▶ Bayesian methods, such as the models proposed here, allow us to undertake patient-centered “joint” benefit-harm assessments
- ▶ Patient-level data is not required - only summaries are required
- ▶ Software implementing the discussed bivariate subgroup analyses is available at <http://hteguru.com/index.php/bbsga/>
- ▶ Software allows one to perform posterior predictive checks and model comparisons.

## References

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4. Evans, S. R. and Follmann, D. (2016), “Using outcomes to analyze patients rather than patients to analyze outcomes: A step toward pragmatism in benefit:risk evaluation”, *Statistics in Biopharmaceutical Research*, **8**(4), 386393.