

# Adjusting for Covariates in Randomized Clinical Trials

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# Disclaimer

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# FDA Guidance Documents on Covariate Adjustment



- ICH E9 Statistical Principles for Clinical Trials (1998):  
“Pretrial deliberations should identify those covariates and factors expected to have an important influence on the primary variable(s), and should consider how to account for these in the analysis to improve precision...”
- COVID-19: Developing Drugs and Biological Products for Treatment or Prevention draft guidance (2020)  
“To improve the precision of treatment effect estimation and inference, sponsors should consider adjusting for prespecified prognostic baseline covariates (e.g., age, baseline severity, comorbidities, baseline medications and COVID-19 vaccination status) in the primary efficacy analysis and should propose methods of covariate adjustment.”
- **Adjusting for Covariates in Randomized Clinical Trials for Drugs and Biological Products revised draft guidance (2021)**  
The main focus of the guidance is on the use of prognostic baseline factors to improve precision for estimating treatment effects.

# Adjusting for Covariates, Revised Draft Guidance (2021)

- I. INTRODUCTION
- II. BACKGROUND
- III. RECOMMENDATIONS FOR COVARIATE ADJUSTMENT IN RANDOMIZED TRIALS
  - A. General Considerations
  - B. Linear models
  - C. Nonlinear models
- IV. REFERENCES

# General Considerations



- Adjustment is recommended because it often improves power and precision, and unadjusted analysis remains acceptable as well.
- Prespecification: covariates used for adjustment and mathematical form of the model
- Adjusting for covariates that are prognostic for the outcome leads to the greatest efficiency gains
  - Scientific literature
  - Previous studies (e.g., a Phase 2 trial)
  - Properties of adjustment are best understood when the number of covariates is small relative to the sample size (Tsiatis et al. 2008)
- Stratified randomization
  - Analysis ignoring stratified randomization is likely to overestimate standard errors (SEs) and can be unduly conservative for inference
  - Recommend SE computation account for stratified randomization (Bugni et al. 2018; Ye et al. 2021)

# Linear Model: Estimation

- Estimand is population average treatment effect (i.e., difference in expected outcomes between subjects assigned to treatment and control groups)
- Usual adjusted estimator is least squares fit of treatment coefficient in a regression of the outcome on an intercept, treatment, and baseline covariates
- Can provide valid estimation of the average treatment effect in a randomized trial even when the linear model is misspecified (Lin, 2013)

# Linear Model: Robust Standard Error

- Nominal SEs reported by most packages for generalized linear models can be inaccurate if the model is incorrect
- Otherwise, can be corrected with robust SE
  - Huber-White “sandwich” SEs when model does not include treatment by covariate interaction (Rosenblum and van der Laan 2009, Lin 2013)
  - Other robust SEs for linear model with interactions (Ye et al. 2022)
  - Appropriate nonparametric bootstrap procedure (Efron and Tibshirani 1993)

# Linear Model: Interactions

- The linear model may include treatment by covariate interaction terms
- However, when using this approach, the primary analysis should still be based on an estimate from the model of the average treatment effect (Tsiatis et al. 2008; Ye et al 2021)
- Per ICH E9, interaction effects may be important to assess in supportive analysis or exploratory analysis because differences in treatment effects across subgroups defined by baseline covariates could be relevant to prescribers, patients, and other stakeholders and imply that the average treatment effect gives an incomplete summary of efficacy

# Linear Model: Example

- Estimand: Difference in mean FEV1 at 12 weeks between drug and placebo in patients with moderate-to-severe asthma regardless of adherence to treatment or use of ancillary medications
- Main analysis: ANCOVA of FEV1 at 12 weeks in all randomized patients, adjusting for baseline FEV1, age, and sex, with Huber-White sandwich standard errors

# Nonlinear Model: Collapsibility

- With binary, ordinal, or time-to-event outcomes certain population-level summaries can be non-collapsible even in randomized trials

Table 1: Non-collapsibility of the odds ratio in a hypothetical target population

	Percentage of target population	Success rate		Odds ratio
		New drug	Placebo	
Males	50%	80.0%	33.3%	8.0
Females	50%	25.0%	4.0%	8.0
Combined	100%	52.5%	18.7%	4.8

- As part of the prespecification of the estimand of interest, sponsors should specify whether treatment effect of interest is the unconditional (e.g., 4.8 in the table) or conditional treatment effect (e.g., 8.0 in the table)

# Nonlinear Model: Conditional Treatment Effects

- Nonlinear models such as logistic regression or proportional hazards regression for conditional treatment effects are commonly used in clinical trials
- Advantages:
  - Can provide more personalized information than unconditional treatment effects if assumption holds (and not otherwise)
  - Available in standard statistical software packages
- Disadvantage:
  - When estimating a conditional treatment effect through nonlinear regression, the model will generally not be exactly correct, and results can be difficult to interpret if the model is misspecified and treatment effects substantially differ across subgroups.
- Sponsors should discuss with the relevant review divisions specific proposals in a protocol or statistical analysis plan containing nonlinear regression to estimate conditional treatment effects for the primary analysis

# Nonlinear Model: Unconditional Treatment Effects



- Sponsors can perform covariate adjusted estimation and inference for an unconditional treatment effect in the primary analysis
- The estimand will be the same as in an unadjusted analysis
- The method used should provide valid inference under approximately the same minimal statistical assumptions that would be needed for unadjusted estimation
- Statistically reliable methods
  - Binary outcomes (e.g., Ge et al. 2011)
  - Ordinal outcomes (e.g., Díaz et al. 2016)
  - Time-to-event outcomes (e.g., Tangen and Koch 1999; Lu and Tsiatis 2008)
- SEs or confidence intervals can be formed from the nonparametric bootstrap or formulas justified in the statistical literature (Colantunoni and Rosenblum 2015)

# Example of “Standardized”, “Plug-in”, or G-computation” Estimator for Unconditional Effect with Binary Outcomes (Ge et al. 2011)

For Binary outcome  $Y$ , treatment A (1=treatment, 0=control), covariate B:

- Fit logistic regression model for

$$P(Y = 1 | A, B) = \text{logit}^{-1}(\beta_0 + \beta_1 A + \beta_2 B).$$

- Compute standardized estimators for treatment specific means  $\mu_0, \mu_1$ :

$$\hat{\mu}_0 = \frac{1}{n} \sum_{i=1}^n \text{logit}^{-1}(\hat{\beta}_0 + \hat{\beta}_2 B_i)$$

$$\hat{\mu}_1 = \frac{1}{n} \sum_{i=1}^n \text{logit}^{-1}(\hat{\beta}_0 + \hat{\beta}_1 + \hat{\beta}_2 B_i)$$

- Estimator is contrast of interest between  $\mu_1, \mu_0$ , e.g., risk difference  $\hat{\mu}_1 - \hat{\mu}_0$ .

Same holds for other (unconditional) estimands, e.g., relative risk

# Nonlinear Model: Unconditional Effect Example

- Estimand: Difference in probability of 28-day survival between drug and placebo in severe-to-critical hospitalized COVID-19 patients regardless of adherence to treatment or use of ancillary medications
- Main Analysis: A logistic model in all randomized patients adjusting for age, baseline severity, and COVID-19 vaccination status, with a standardized (plug-in) estimator of the risk difference and SE (Ge et al. 2011) and a Wald test

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# Backup Slides

# Simulation Study Result: difference in restricted mean survival times (RMST) 14 days after hospitalization

n	Estimator	Power	MSE	RE
100	Unadjusted	0.09	53.7	1.00
100	Adjusted	0.15	51.0	0.95
200	Unadjusted	0.33	62.7	1.00
200	Adjusted	0.40	56.4	0.90
500	Unadjusted	0.74	72.9	1.00
500	Adjusted	0.82	62.2	0.85
1000	Unadjusted	0.96	76.5	1.00
1000	Adjusted	0.98	63.5	0.83

n=sample size; RE=relative efficiency (ratio of adjusted vs. unadj. MSE).

# Nonlinear Model: Precision and Efficiency of the Unadjusted, Standardized, and Logistic Coefficient Estimators



	Estimator	Average value of estimator	Empirical standard error	Relative efficiency	Reduction in sample size
<b>Setting 1</b>	Unadjusted	0.13	$4.5 \times 10^{-2}$	1	0
	Standardized	0.13	$3.8 \times 10^{-2}$	1.41	29%
	Logistic coefficient	0.76	0.23	1.31	24%
<b>Setting 2</b>	Unadjusted	0.13	$4.5 \times 10^{-2}$	1	0
	Standardized	0.13	$4.5 \times 10^{-2}$	0.99	-1%
	Logistic coefficient	0.53	0.19	0.94	-7%

Setting 1: Baseline variables prognostic for the outcome; Setting 2: Baseline variables independent of the outcome; For both the unadjusted and standardized estimator, the true unconditional treatment effect is 0.13 in both settings. In setting 2, the true conditional treatment effect on the log odds scale is 0.52. As the logistic regression model is not necessarily correct in setting 1, it is unclear if the true conditional effect is interpretable as a single number.

# Nonlinear Model: Properties of the Unadjusted, Standardized, and Logistic Coefficient Estimators

Estimator	Effect it estimates	Requires regression model assumptions?	Adjusts for baseline variables?
Unadjusted	Unconditional effect	No	No
Standardized	Unconditional effect	No	Yes
Logistic coefficient	Conditional effect	Yes	Yes

Steingrimsson, Hanley, and Rosenblum 2017

# Nonlinear Model: Example of Misspecified Logistic Model

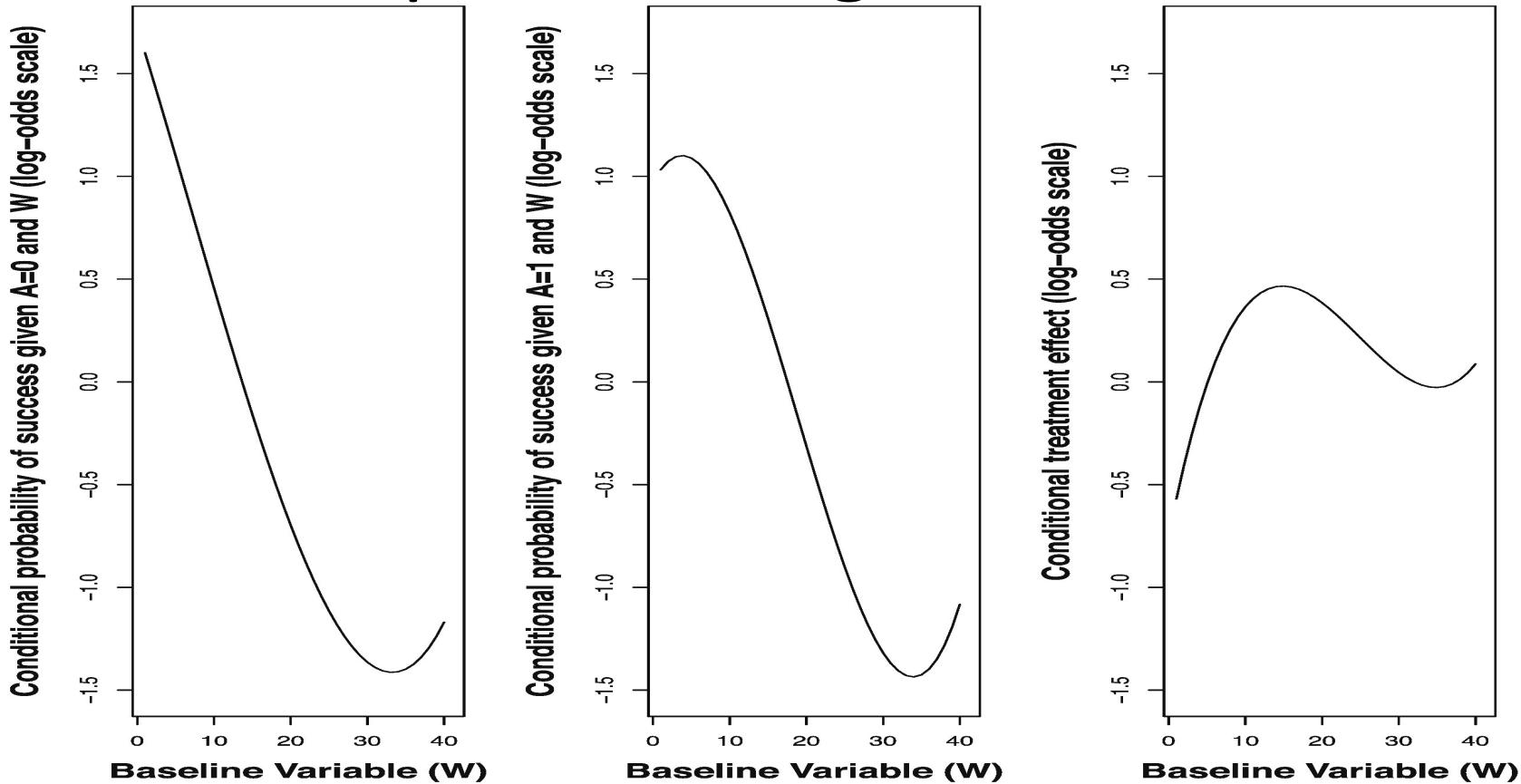
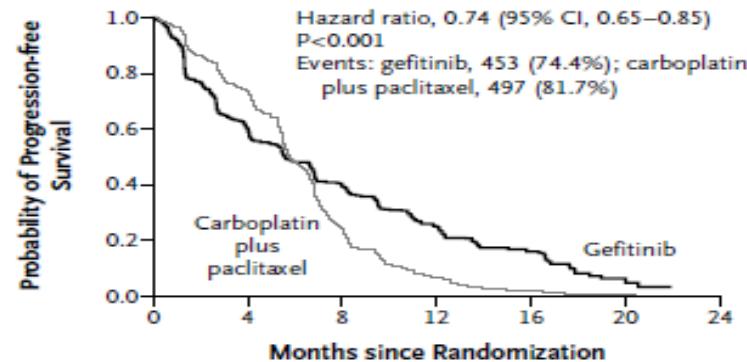


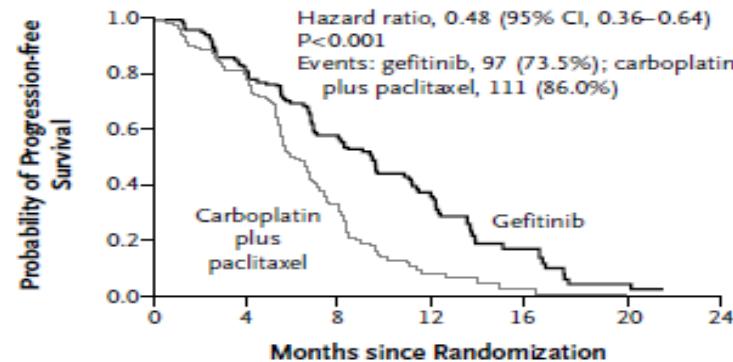
Fig. 1. An example of a conditional effect that depends on the value of the baseline variables and cannot be represented using a single number.

Steingrimsson, Hanley, and Rosenblum 2017

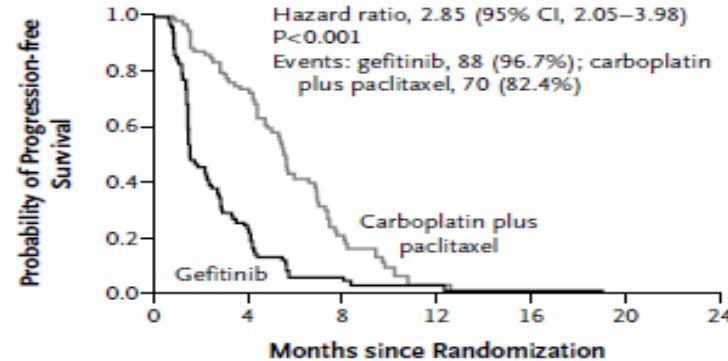
# Non-proportional Hazard Ratio Example

**A Overall****No. at Risk**

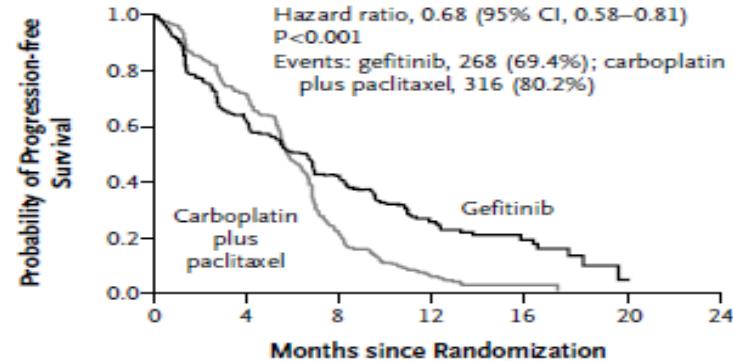
	0	3	6	9	12	15	18	21	24
Gefitinib	609	363	212	76	24	5	0	0	0
Carboplatin plus paclitaxel	608	412	118	22	3	1	0	0	0

**B EGFR-Mutation-Positive****No. at Risk**

	0	3	6	9	12	15	18	21	24
Gefitinib	132	108	71	31	11	3	0	0	0
Carboplatin plus paclitaxel	129	103	37	7	2	1	0	0	0

**C EGFR-Mutation-Negative****No. at Risk**

	0	3	6	9	12	15	18	21	24
Gefitinib	91	21	4	2	1	0	0	0	0
Carboplatin plus paclitaxel	85	58	14	1	0	0	0	0	0

**D Unknown EGFR Mutation Status****No. at Risk**

	0	3	6	9	12	15	18	21	24
Gefitinib	386	234	137	43	12	2	0	0	0
Carboplatin plus paclitaxel	394	251	67	14	1	0	0	0	0

