Overall average treatment effects and one-variable-at-a-time subgroup analysis: The Scylla and Charybdis of Evidence Based Medicine



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Scylla and Charybdis



Limitations of RCTs for Clinical Decision Making

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Limitations of RCTs for Clinical Decision Making

- While RCTs can determine the better treatment on average, they "do not answer the practicing doctor's question: what is the most likely outcome when this particular drug is given to a particular patient?"
 - Austin Bradford Hill





Evidence Based Medicine

Hill was wrong.



- RCTs should be used by doctors to
 - determine what's best for individual patients.
- "Evidence based medicine is the conscientious, explicit, judicious and reasonable use of current best evidence in making decisions about the care of individual patients."



 EBM proposed to repurpose RCTs from tools to establish causation into tools for *prediction* in single cases.



What's best on average must be best for each individual.

The Fallacy of Division (Wennington's Fallacy)



Starters	PTS
B.J. Armstrong	16
Scottie Pippen	19
Michael Jordan	55
Toni Kukoc	3
Will Perdue	6
Reserves	PTS
Luc Longley	5
Corie Blount	2
Steve Kerr	5
Larry Krystkowiak	0
Bill Wennington	2
Pete Myers	U
Team Totals	113

"Michael and I combined for 57 points" -Bill Wennington, 1995

- It is potentially misleading to draw inferences about individuals based on aggregated characteristics of the (heterogeneous) group to which they belong.
 How do we estimate "individual" treatment
 - How do we estimate "individual" treatment effects?

ACTUAL OUTCOME			
0 = alive 1 = dead			

In	Individual Treatment Effects in a					
D	Deterministic Framework: Four					
ро	ossibilities					
	Without	With				
	Treatment	Treatment				
	0	0	NO EFFECT			
	0	1	HARM			
	1	0	BENEFIT			
	1	1	NO EFFECT			

Subject Name	Without Treatment	With Treatment	
SAM	0		
MARY		0	
вов	0		
BEN		0	
CHRISTINE		0	
NEIL	1		
MOHAMED		1	
JENNIFER		1	
PAUL	0		
NISHA	1		
MIGUEL	1		
LAYLA		0	
PAUL	0		
EMANUEL		1	
CHERYL		0	
PATRICK	0		
OSCAR		1	
JULIANNE	0		
THOMAS	0		
GEORGE		0	

ACTUAL OUTCOME
COUNTER FACTUAL OUTCOME

0 = alive 1 = dead

•••			, in a			
D	Deterministic Framework: Four					
po	ossibilities					
	Without	With				
	Treatment	Treatment				
	0	0	NO EFFECT			
	0	1	HARM			
	1	0	BENEFIT			
	1	1	NO EFFECT			

Individual Treatment Effects in a

Subject Name	Without Treatment	With Treatment	
SAM	0	1	
MARY	0	0	
вов	0	0	
BEN	1	0	
CHRISTINE	1	0	
NEIL	1	1	
MOHAMED	1	1	
JENNIFER	1	1	
PAUL	0	1	
NISHA	1	1	
MIGUEL	1	1	
LAYLA	1	0	
PAUL	0	0	
EMANUEL	1	1	
CHERYL	0	0	
PATRICK	0	0	
OSCAR	1	1	
JULIANNE	0	0	
THOMAS	0	0	
GEORGE	1	0	

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	0	1	HAR			
	1	0	BENE			

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Individual Treatment Effects in a

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Subject Name	Wit	hout Treatment	With Treatme	ent	
SAM		0	1		← HARM
MARY		0	0		
вов		0	0		
BEN		1	0		← BENEFIT
CHRISTINE		1	0		← BENEFIT
NEIL		1	1		
MOHAMED		1	1		
JENNIFER		1	1		
PAUL		0	1		← HARM
NISHA		1	1		
MIGUEL		1	1		
LAYLA		1	0		← BENEFIT
PAUL		0	0		
EMANUEL		1	1		
CHERYL		0	0		
PATRICK		0	0		
OSCAR		1	1		
JULIANNE		0	0		
THOMAS		0	0		
GEORGE		1	0		← BENEFIT



Conventional Approach: One variable at a time subgroup analysis

 This approach relies on the detection of "statistically significant" effect modifiers contrasting *relative* treatment effects across levels of each subgrouping variable, onevariable-at-a-time.



Problems With Conventional Subgroup Analysis

Fail to detect HTE even when its there:

- Low power
 - For covariates with 50% prevalence (e.g. gender): ~4 fold the sample size.
 - For covariates with 20% prevalence (e.g. common comorbidities): ~10 fold the sample size.
- Compare Groups of patients that are more similar than dissimilar.
- Individuals patients belong to many different subgroups.
- Spurious False Positives



Why most subgroup effects are false or overestimated



INITIAL PACE

Kent DM, Steyerberg E, van Klaveren D. BMJ 2018 (In Press).

Why most subgroup effects are false or overestimated





Kent DM, Steyerberg E, van Klaveren D. BMJ 2018 (In Press).

"Positive" subgroup analyses subsequently shown to be false

Observation	Refutation	
Aspirin is ineffective in secondary prevention of stroke in women ^{29,30}	31	



Interim Summary

Determining the best treatment on average (the task of an RCT) is very different from determining the best treatment for an individual (the task of a good clinician). Conventional subgroup analysis of clinical trials are typically inadequate and can also be misleading, and are not consistent with the clinically most important goal of *prediction*.



- The reference class problem is a model selection problem
- The goal of Personalized EBM can be conceived as the identification of an optimal subgrouping scheme, based on all relevant patient characteristics, that yields a more individualized reference class for each patient than the overall trial results.

Why Risk Based Subgroup Analysis Should be Routine



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Why privilege risk-based HTE analysis?

 Risk is a known mathematical determinant of treatment effect.



Common Measures of Treatment Effect

Risk Reduction	Definition
(RR)	
Absolute RR	EER-CER
Relative RR	1 - <u>EER</u> CER
Odds Ratio	EER/(1-EER) CER/(1-CER)
CER=control event rate	

EER=experimental event rate



An Illustration of Scale Dependence of HTE over Baseline Outcome Risk



Dahabreh, Hayward, Kent, IJE, 2016



Why privilege risk-based HTE analysis?

- Risk is a known mathematical determinant of treatment effect.
- When baseline risk heterogeneity is present (and the treatment effect is non-zero), there is always HTE.
- Risk provides a summary measure that takes into account multiple variables that are relevant; provides "patient-centered" evidence.



Figure 1: Distribution of Mortality Risk with Thrombolytic Thearpy in Patients with Acute Myocardial Infarction



Percentile Mortality Risk



Kent DM, et al. J Gen Intern Med 2002; 17:887-94.

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Percentile Mortality Risk



Kent DM, et al. J Gen Intern Med 2002; 17:887-94.



DANAMI-2



Number at risk

TIMI 0-4 Fx 556 533 531 PA 578 546 540 TIMI ≥ 5 Fx 207 154 141 PA 186 150 145

Thune JJ, et al. Circulation 2005,112:2017-2021.



Predicted Risk Distributions in RCTs

PACE



PACE

Predicted Risk Distributions in RCTs



Diabetes Prevention Program (DPP) Randomized Controlled Trial

- <u>Participants</u>: 3060 nondiabetic persons with evidence of impaired glucose metabolism.
- Intervention: Intervention groups received metformin or a lifestyle-modification program.
- <u>Main Outcome Measure</u>: Development of diabetes

The DPP study was conducted by the DPP Investigators and supported by the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK).



DPP Risk Stratified Results



p value = NS

p value = 0.0008



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DPP Risk Stratified Results







Risk based analyses can reveal counter-intuitive findings

- Overall effectiveness results may be driven by a relatively small group of influential (typically high risk) patients;
- The typical (median) risk patient is frequently at considerably lower risk than the overall average;
- The average benefit seen in the summary result often over estimates the benefit (on the RD scale) in most patients (and may obscure harm in many).



METHODOLOGY

Open Access

Assessing and reporting heterogeneity in treatment effects in clinical trials: a proposal **Strials**

David M Kent^{1*}, Peter M Rothwell², John PA Ioannidis^{1,3}, Doug G Altman⁴, Rodney A Hayward⁵

- Evaluate and <u>report on the distribution of risk</u> in the overall study population and in the separate treatment arms of the study by using a risk prediction model or index.
- 2. Primary subgroup analyses should include reporting how relative and absolute risk reduction varies in a **risk-stratified analysis**.
- 3. Any additional <u>primary subgroup analysis should be pre-</u> <u>specified</u> and limited to patient attributes with strong a prior pathophysiological or empirical justification.
- 4. Conduct and <u>report on secondary (exploratory) subgroup</u> <u>analyses separate</u> from primary subgroup comparisons.
- 5. All analyses conducted must be reported and statistical testing of HTE should be done using **appropriate methods** (such as interaction terms) and avoiding over-interpretation.



Risk modeling vs. Effect modeling

Annals of Internal Medicine RESEARCH AND REPORTING METHODS

The Predictive Approaches to Treatment effect Heterogeneity (PATH) Statement: Explanation and Elaboration

Table 2. Equations Corresponding to Risk-Modeling and Effect-Modeling Approaches

Risk modeling

A multivariable regression model *f* that predicts the risk for an outcome based on risk predictors *x_i* is identified or developed:

Equation 1: risk = $f(\alpha + \beta_1 \times x_1 + \ldots + \beta_p \times x_p)$

Variation in the treatment effect across risk can be tested statistically on the relative scale through the interaction between a linear predictor of risk $(lp = \beta_1 \times x_1 + ... + \beta_p \times x_p)$ and treatment assignment *tx*:

Equation 2: risk = $f(\alpha + \beta_{tx} \times tx + \beta_{lp} \times lp + \delta_{lp} \times lp \times tx)$

Including a treatment interaction with the linear predictor of risk permits the relative treatment effect to vary linearly across levels of risk (and permits testing of the statistical significance of this interaction effect, δ_{lp}).

When relative effects across risk strata seem constant, a model with a constant treatment effect may suffice:

Equation 3: risk = $f(\alpha + \beta_{tx} \times tx + \beta_1 \times x_1 + \ldots + \beta_p \times x_p)$,

where the parameter β_{tx} represents a constant risk reduction on the log hazard or log odds scale for treated (tx = 1) versus control (tx = 0) patients.

Effect modeling

A regression model *f* is developed on RCT data with inclusion of risk predictors x_i , a treatment assignment variable *tx*, and potential treatment interaction terms ($x_i \times tx$):

Equation 4: risk = $f(\alpha + \beta_{tx} \times tx + \beta_1 \times x_1 + \ldots + \beta_p \times x_p + \delta_t \times x_1 \times tx + \ldots + \delta_p \times x_p \times tx)$

RCT = randomized controlled trial.



JAMA | Original Investigation

Research

Heterogeneity of Treatment Effects in an Analysis of Pooled Individual Patient Data From Randomized Trials of Device Closure of Patent Foramen Ovale After Stroke

Redevelopment and va individualise decision r surgical revascularisati artery disease: seconda randomised controlled validation

Kuniaki Takahashi, Patrick W Serruys, Valentin Fuster, Jung-Min Ahn, Arie Pieter Kappetein, Stuart J Head, Da David van Klaveren, on behalf of the SYNTAXES, FREEL

Summary

Background Randomised controlled trials ar interventions, and typically report the averavary between patients, basing treatment dec be suboptimal. We aimed to develop an in strategy in patients with complex coronary (David M. Kent, MD, MS; Jeffrey L. Saver, MD; Scott E. Kasner, MD; Jason Nelson, MS; John D. Carroll, MD; Gilles Chatellier, MD; Geneviève Derumeaux, MD; Anthony J. Furlan, MD; Howard C. Herrmann, MD; Peter Jüni, MD; Jong S. Kim, MD; Benjamin Koethe, MS; Pil Hyung Lee, MD; Benedicte Lefebvre, MD; Heinrich P. Mattle, MD; Bernhard Meier, MD; Mark Reisman, MD; Richard W. Smalling, MD, PhD; Lars Soendergaard, MD; Jae-Kwan Song, MD; Jean-Louis Mas, MD; David E. Thaler, MD, PhD

IMPORTANCE Patent foramen ovale (PFO)-associated strokes comprise approximately 10% of ischemic strokes in adults aged 18 to 60 years. While device closure decreases stroke recurrence risk overall, the best treatment for any individual is often unclear.

OBJECTIVE To evaluate heterogeneity of treatment effect of PFO closure on stroke recurrence based on previously developed scoring systems.

DESIGN, SETTING, AND PARTICIPANTS Investigators for the Systematic, Collaborative, PFO Closure Evaluation (SCOPE) Consortium pooled individual patient data from all 6 randomized clinical trials that compared PFO closure plus medical therapy vs medical therapy alone in patients with PFO-associated stroke, and included a total of 3740 participants. The trials were conducted worldwide from 2000 to 2017.

EXPOSURES PFO closure plus medical therapy vs medical therapy alone. Subgroup analyses used the Risk of Paradoxical Embolism (RoPE) Score (a 10-point scoring system in which higher scores reflect younger age and the absence of vascular risk factors) and the PFO-Associated Stroke Causal Likelihood (PASCAL) Classification System, which combines the RoPE Score with high-risk PFO features (either an atrial septal aneurysm or a large-sized shunt) to classify natients into 3 categories of causal relatedness: unlikely, possible.



Summary

- Heterogeneity of outcome risk is ubiquitous.
- Heterogeneity of outcome risk inevitably gives rise to heterogeneity of treatment effect.
- One variable at a time subgroup analyses are inadequate (and prone to spurious false positive results).
- Risk based subgroup analyses can do better.











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