

Pragmatic Clinical Trial Challenges: Lessons Learned from the NIH Collaboratory Biostatistics and Design Core

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Outline



NIH Collaboratory Pragmatic Trial Setting

Common themes across studies

- Study Design
- Analysis/Sample Size
 - Implications of Variable Cluster Size on Estimation and Power
- Randomization

□ Conclusions/Next Steps

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The NIH Collaboratory



□ Supported by The Common Fund (NIH Director's fund)

Goal: improve the way (pragmatic) clinical trials are conducted

Build infrastructure for collaborative research

The NIH Collaboratory



□ Supported by The Common Fund (NIH Director's fund)

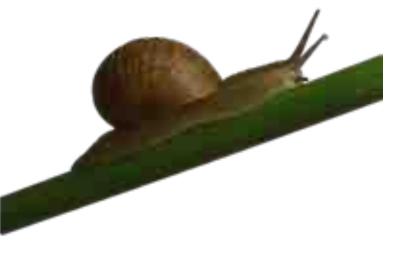
Goal: improve the way (pragmatic) clinical trials are conducted

□ Build infrastructure for collaborative research

•Why was the NIH Collaboratory created????

Challenge #1: Clinical research is slow

- Traditional RCTs are slow and expensive and rarely produce findings that are easily put into practice.
- In fact, it takes an average of 17 years
 before research findings
 lead to widespread
 changes in care.



Health Care Systems Research Collaboratory

Challenge #2: Clinical research is not relevant to practice

- Traditional RCTs study effectiveness of txs for carefully selected populations under ideal conditions.
- Difficult to translate to real world.
- When implemented into everyday clinical practice, often see a "voltage drop" — dramatic decrease in effectiveness.

"If we want more evidencebased practice, we need more practice-based evidence." Green, LW. American Journal

of Public Health, 2006.



Health Care Systems Research Collaboratory

Challenge #3: The evidence paradox

- >18,000 RCTs published each year—plus tens of thousands of other clinical studies.
- Yet systematic reviews consistently find not enough evidence to effectively inform clinical decisions providers and patients must make.

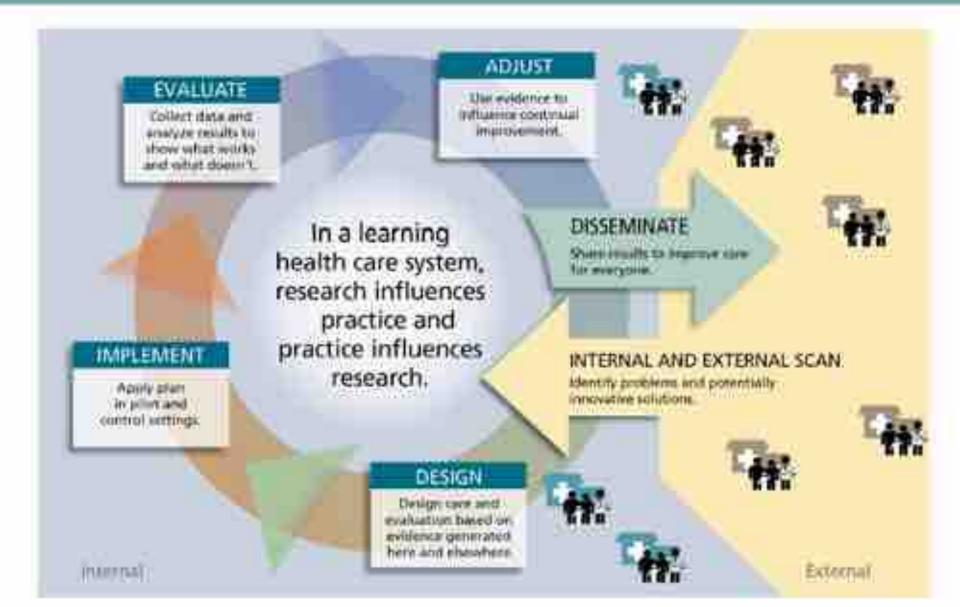




Health Care Systems Research Collaboratory

Learning health care systems





Pragmatic vs. Explanatory Trials



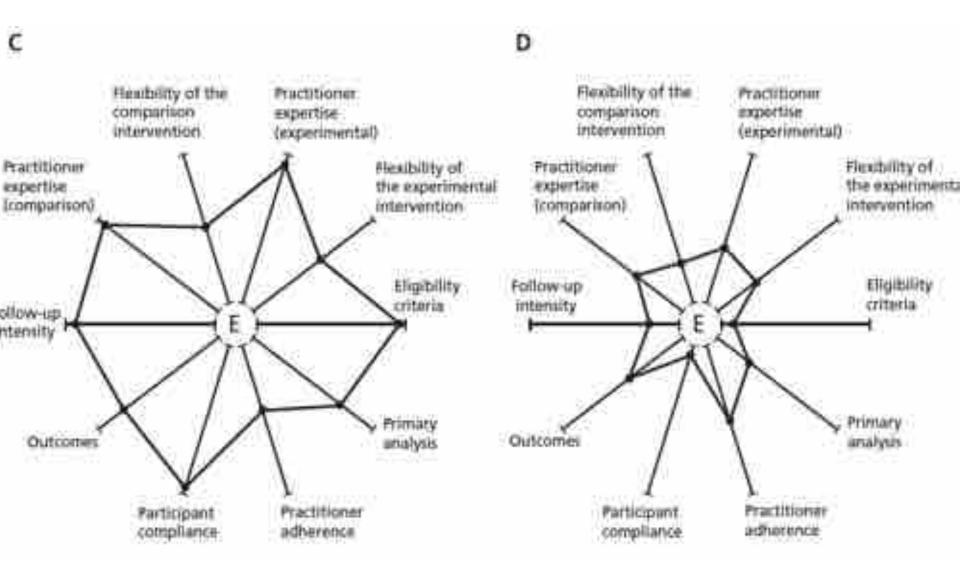


A pragmatic-explanatory continuum indicator summary (PRECIS): a tool to help trial designers

Kevin E. Thorpe MMath, Merrick Zwarenstein MD MSc, Andrew D. Oxman MD, Shaun Treweek BSc PhD, Curt D. Furberg MD PhD, Douglas G. Altman DSc, Sean Tunis MD MSc, Eduardo Bergel PhD, Ian Harvey MB PhD, David J. Magid MD MPH, Kalipso Chalkidou MD PhD

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Pragmatic vs. Explanatory Trials



Key features of most PCTs

Use of electronic health records (EHRs)



• EHRs allow efficient and cost-effective, recruitment, participant communication & monitoring, data collection, and follow up



Randomization at clinic or provider level

 Protocols can be tailored to local sites and can adapt to changes in a dynamic health care environment

Health Care Systems Research Collaboratory

laboratory

Pragmatic Trials Concept



 \Box Size: Large simple trials \rightarrow precise estimates, evaluate heterogeneity

Endpoints: patient oriented usually with minimal adjudication

Setting: integrated into real world

- Non-academic centers
- Leverage electronic data
- Patients as partners

Round 1 Demonstration Projects



Principal Investigator	Institution	Project	
Gloria Coronado	Kaiser Foundation Research Institute	Strategies and Opportunities to Stop Colon Cancer in Priority Populations	
Lynn DeBar	Kalser Foundation Research Institute	Collaborative Care for Chronic Pain in Primary	
Laura Dember	University of Pennsylvania	Pragmatic Trials in Maintenance Hemodialysis	
Susan Huang	University of CaliforniaIrvine	Decreasing Bioburden to Reduce Healthcare- Associated Infections and Readmissions	
Jeffrey Jarvik	University of Washington	A Pragmatic Trial of Lumbar Image Reporting with Epidemiology (LIRE)	
Gary Rosenthal	University of Iowa	Nighttime Dosing of Anti-Hypertensive Medications: A Pragmatic Clinical Trial	
Gregory Simon	Group Health Cooperative	Pragmatic trial of population-based programs to prevent suicide attempt	

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STUDY DESIGN

Study Design: Cluster RCT



□ Mostly Cluster RCTs (except one)

- Randomization Unit:
 - Provider < Panel < Clinic < Region < Site
- □ Average Size of Cluster
 - Initial Proposals: Most large clinic level clusters
 - Goal: Smallest Unit without contamination
 - More clusters are better if possible
 - Smaller number of clusters increase sample size along with estimation issues (GEE)
 - Potential Solutions: Panel-level or physician-level



Cluster

- Randomize at cluster-level
- Most common, but not necessarily the most powerful or feasible
- Advantages:
 - Simple design
 - Easy to implement
- Disadvantages:
 - Need a large number of clusters
 - Not all clusters get the interventions
 - Interpretation for binary and survival outcomes:
 - Mixed models within cluster interpretation problematic
 - GEE marginal estimates interpretation, but what if you are interested in within cluster changes?



□ Cluster with Cross-over

- Randomize at cluster but cross to other intervention assignment midway
- Feasible if intervention can be turned off and on without "learning" happening
- Alternative: baseline period without intervention and then have half of the clusters turn on



5				
	Cluster	Period 1	Period 2	
	1	INT		
Simple	2	UC		
Cluster	3	UC		
	4	INT		
Cluster With Crossover	1	INT	UC	
	2	UC	INT	
	3	UC	INT	
	4	INT	UC	
Cluster With Baseline	1	UC	INT	
	2	UC	UC	
	3	UC	UC	
	4	UC	INT	

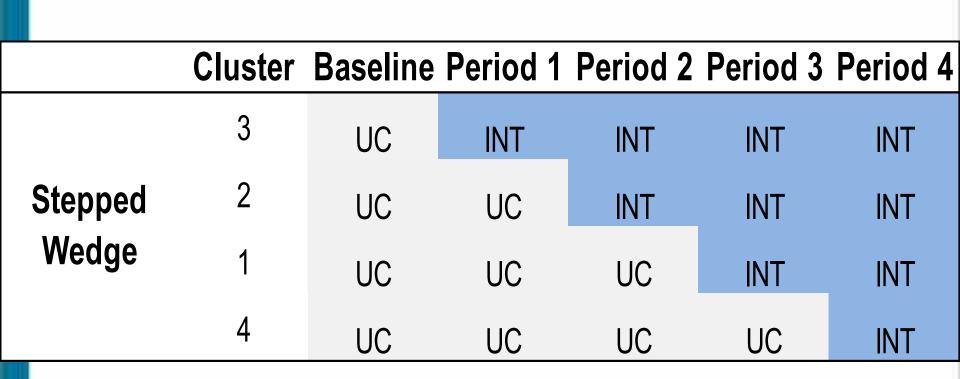


- □ Cluster with Cross-over
 - Advantages:
 - Can make within cluster interpretation
 - Potential to gain power by using within cluster information
 - Disadvantages:
 - Contamination can yield biased estimates especially for the standard cross-over design
 - May not be feasible to switch assignments or turn off intervention
 - Not all clusters have the intervention at the end of the study



□ Stepped Wedge Design

- Randomize timing of when the cluster is turned on to intervention
- Staggered cluster with crossover design
- Temporally spaces the intervention and therefore can control for system changes over time



GroupHealth.



- □ Stepped Wedge Design
 - Advantages:
 - All clusters get the intervention
 - Controls for external temporal trends
 - Make within cluster interpretation if desired
 - Disadvantages:
 - Contamination can yield biased estimates
 - Heterogeneity of Intervention effects across clusters can be difficult to handle analytically
 - Special care of how you handle random effects in the model
 - Relatively new and available power calculation software is relatively limited



ANALYSIS/SAMPLE SIZE



Analysis Implications

- What are you making inference to?
 - Compare intervention across clinics
 - Marginal cluster-level effect
 - Compare within-clinic intervention effect
 - Within-clinic effect
 - Compare intervention effect across patients
 - Marginal patient-level effect
 - Compare an in-between cluster and patient-level effect

DeLong, E, Cook, A, and NIH Biostatistics/Design Core (2014) Unequal Cluster Sizes in Cluster-Randomized Clinical Trials, *NIH Collaboratory Knowledge Repository*.

Cook, AJ, Delong, E, Murray, DM, Vollmer, WM, and Heagerty, PJ (2016) Statistical lessons learned for designing cluster randomized pragmatic clinical trials from the NIH Health Care Systems Collaboratory Biostatistics and Design Core *Clinical Trials* **13(5)** 504-512.



□ What is the scientific question of interest?

- Marginal cluster-level effect
 - "What is the average expected clinic benefit if all clinics in the health system changed to the new intervention relative to Usual Care?"
- □ Within-clinic effect
 - "What is the expected benefit if a given clinic implements the new intervention relative to Usual Care?"
- Marginal patient-level effect
 - "What is the average expected patient benefit if all the clinics in the health system changed to the new intervention relative to Usual Care?"



□ Simplified Example:

- *Y lci* is a binary outcome for patient *i* at clinic *c*
- $n\downarrow c$ is the number of patients at clinic c
- X1c is 1 if clinic c was randomized to intervention or 0
- Estimate a simple marginal clinic-level effect (difference in clinic means amongst those randomized to intervention relative to those not randomized)

 $\Delta \uparrow c = \sum c = 1 \uparrow N \implies \mu \downarrow c X \downarrow c / \sum c = 1 \uparrow N \implies X \downarrow c - \sum c = 1 \uparrow N \implies \mu \downarrow c$ $(1 - X \downarrow c) / \sum c = 1 \uparrow N \implies (1 - X \downarrow c)$

where $\mu \downarrow c = \sum i = 1 \uparrow n \downarrow c \implies Y \downarrow ci / n \downarrow c$ is the mean outcome at clinic *c*



□ Simplified Example:

- *Y lci* is a binary outcome for patient *i* at clinic *c*
- $n\downarrow c$ is the number of patients at clinic c
- *X*1*c* is 1 if clinic *c* was randomized to intervention or 0
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Patients are weighted equally and clustering is really just nuisance in terms of variance and not of interest



□ Some ways to estimate these quantities in practice

- Marginal cluster-level effect
 - □ GEE with weights the inverse of the cluster size with independent correlation structure and robust variance
- □ Compare within-clinic intervention effect
 - GLMM but need to get correlation structure correct but most often just a cluster random effect
- Marginal patient-level effect
 - GEE with no weights with independent correlation structure and robust variance
- □ In-between cluster and patient-level effect
 - GEE with no weights but exchangeable cluster correlation structure and robust variance
 - Exchangeable weights based on statistical information, but not necessarily the most interpretable

Sample Size: Variable Cluster Size



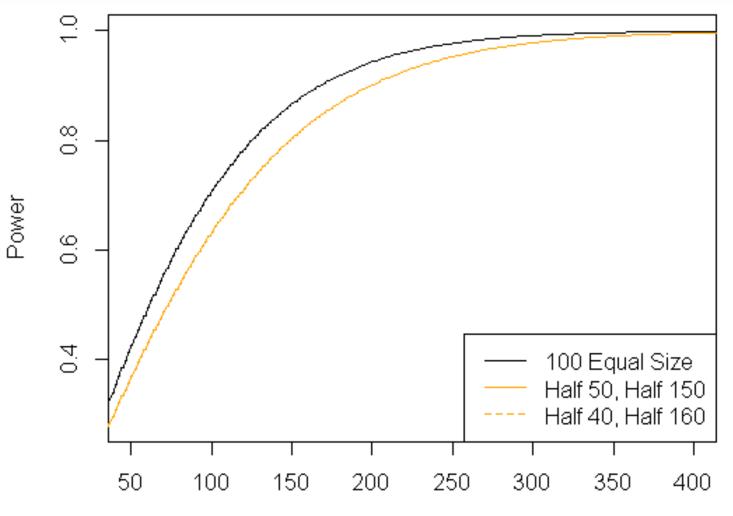
- Sample Size calculations need to take variable cluster size into account
 - Design effects (amount sample size is inflated due to cluster randomization relative to individual patient randomization) are different
 - Depends on the analysis of choice and the estimate of interest
- □ Example: Estimating marginal clinic-level mean difference
 - Design effect:

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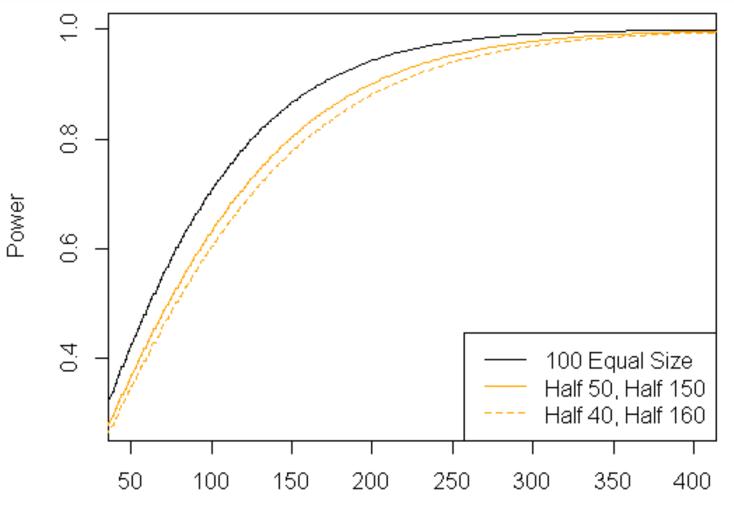
DeLong, E, Lokhnygina, Y and NIH Biostatistics/Design Core (2014) The Intraclass Correlation Coefficient (ICC), *NIH Collaboratory Knowledge Repository*.

Eldridge, S.M., Ashby, D., and Kerry, S. (2006) Sample size for cluster randomized trials: effect of coefficient of variation of size and analysis method. *Int J Epi* **35**:1292-1300.

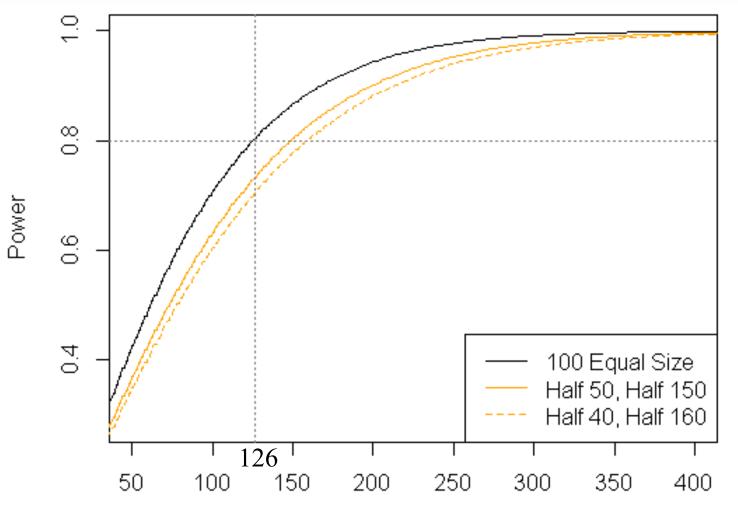














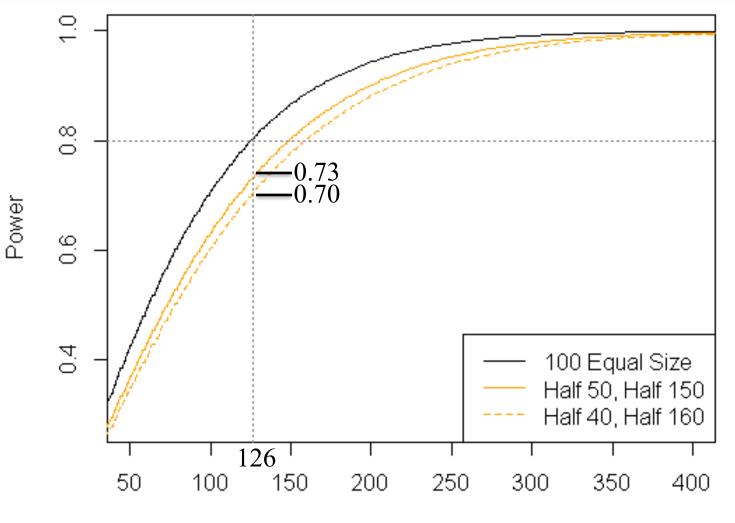
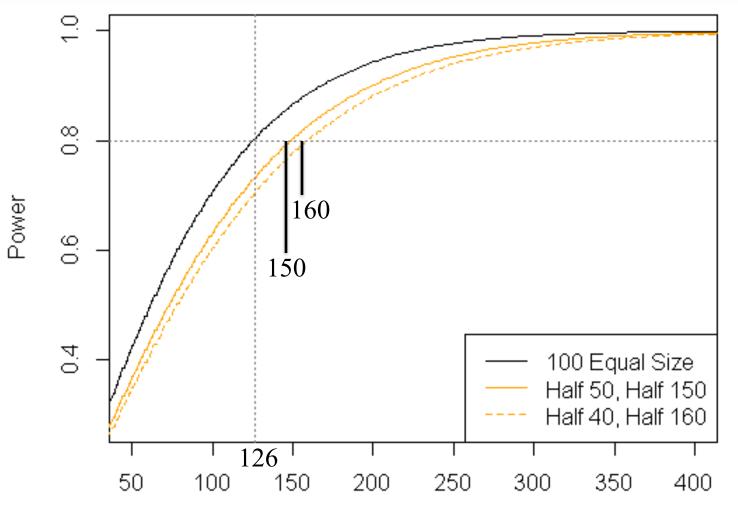


Figure: Power Curve ICC is 0.03 and effect size 0.1σ





Number of Clusters



RANDOMIZATION

Randomization



- Crude randomization not preferable with smaller number of clusters or need balance for subgroup analyses
- □ How to balance between cluster differences?
 - Paired
 - How to choose the pairs best to control for important predictors?
 - Implications for analyses and interpretation
 - Stratification
 - Stratify analysis on a small set of predictors
 - Can ignore in analyses stage if desired
 - Other Alternatives

DeLong, E, Li, L, Cook, A, and NIH Biostatistics/Design Core (2014) Pair-Matching vs stratification in Cluster-Randomized Trials, *NIH Collaboratory Knowledge Repository*.

Randomization: Constrained Randomization



- Balances a large number of characteristics
- □ Concept
 - 1. Simulate a large number of cluster randomization assignments (A or B but not actual treatment)
 - 2. Remove duplicates
 - 3. Across these simulated randomizations assignments assess characteristic balance
 - 4. Restrict to those assignments with balance
 - 5. Randomly choose from the "constrained" pool a randomization scheme.
 - 6. Randomly assign treatments to A or B

Randomization: Constrained Randomization



- Is Constrained randomization better then unconstrained randomization
- How many valid randomization schemes do you need to be able to conduct valid inference?
- Do you need to take into account randomization scheme in analysis?
 - Ignore Randomization
 - Adjust for variables in regression
 - Permutation inference

Randomization: Constrained Randomization



- Is Constrained randomization better then unconstrained randomization
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 - Ignore Randomization
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Conduct a simulation study to assess these properties

Randomization: Constrained Randomization Simulation Design



- □ Outcome Type: Normal
- Randomization Type: Simple versus Constrained
- □ Inference Type: Exact (Permutation) versus Model-Based (F-Test)
- Adjustment Type: Unadjusted versus Adjusted
- Clusters: Balanced designs, but varied size and number
- □ Correlation: Varied ICC from 0.01 to 0.05
- Description Potential Confounders: Varied from 1 to 4

Li, F., Lokhnygina, Y., Murray, D, Heagerty, P., and Delong, ER. (2015) An evaluation of constrained randomization for the design and analysis of group-randomized trials (In Submission).

Randomization: Constrained Randomization Simulation Results



- Adjusted F-test and the permutation test perform similar and slightly better for constrained versus simple randomization.
- □ Under Constrained Randomization:
 - Unadjusted F-test is conservative
 - Unadjusted Permutation holds type I error (unless candidate set size is not too small)
 - Unadjusted Permutation more powerful then Unadjusted F-Test
- Recommendation: Constrained randomization with enough potential schemes (>100), but still adjust for potential confounders

Randomization: Constrained Randomization Next Steps



□ What about Binary and Survival Outcomes??

□ Hypothesized Results (Mine not NIH Collaboratories):

- Constrained Randomization probably still wins
- Binary Outcomes: Likely less of a preference for adjusted versus unadjusted analyses (mean and variance relationship (minimal precision gains))
- Survival Outcomes: Depends on scenario and model choice (frailty versus robust errors)

Conclusions



- Pragmatic Trials are important to be able to move research quickly into practice
- □ Pragmatic Trials add Complication
 - First Question: Can this study be answered using a pragmatic trial approach??
 - Study Design is essential and needs to be flexible
 - Choice of which quantity to estimate should be made based on the scientific question of interest, but statistical trade-offs, including power, must also be considered.
 - Variability in cluster sizes have potentially major implications for power and analysis approach
- □ Lots of open statistical questions still to be addressed



EXTRA SLIDES



OUTCOME ASCERTAINMENT

Outcome Ascertainment



Most trials use Electronic Healthcare Records (EHR) to obtain Outcomes

- Data NOT collected for research purposes
- If someone stays enrolled in healthcare system assume that if you don't observe the outcome it didn't happen
 - In closed system this is likely ok
 - Depends upon cost of treatment (likely to get a bill the more the treatment costs)

Outcome Ascertainment (Cont)



Do you need to validate the outcomes you do observe?

- Depends on the Outcome (PPV, sensitivity)
- Depends on the cost (two-stage design?)
- □ How do you handle Missing Outcome Data?
 - Leave healthcare system
 - Type of Missing Data: Administrative missingness (MCAR), MAR or non-ignorable?
 - Amount of Missing Data: how stable is your population being studied?
 - Depends on the condition and population being studied.

DeLong, E, Li, L, Cook, A, and NIH Biostatistics/Design Core (2014) Key Issues in Extracting Usable Data from Electronic Health Records for Pragmatic Clinical Trials, *NIH Collaboratory Knowledge Repository*