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# On the Utility of Subgroup Analyses in Confirmatory Clinical Trials

**EMA Expert Workshop on Subgroup Analyses**

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Answers That Matter.

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# Outline

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## Classification of Subgroup Analyses from Confirmatory Clinical Trials

### Proposal for Scenario 1

- General Method
- Implementation Considerations
- Example

### Proposals for Scenarios 2 and 3

### Summary Comments

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# Classifying Subgroup Analyses

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## **Confirmatory Subgroup Analyses**

- Involve well-defined subpopulations and pre-defined analyses yielding valid inference on the subpopulation(s)
- These analyses make up the primary or key (“gated”) secondary analyses (objectives) of the trial
- Strong control of fwer

## **Exploratory Subgroup Analyses**

- Non-confirmatory analyses
- Analyses separate from the primary or key secondary objectives of the trial

### 1: supportive analyses

- Offered to support the primary inference
- Based on some *a priori* hypothesis of subpopulation effects

### 2: discovery analyses

- Used to find potentially viable subpopulations (in a data-driven manner)
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# Classifying Subgroup Analyses

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## Confirmatory Subgroup Analyses

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## Exploratory Subgroup Analyses

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Context determines which of these exploratory subgroup analyses is applicable

# Setting: Scenario 1

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## Quote from Draft Guideline:

The clinical data presented are overall statistically persuasive with therapeutic efficacy demonstrated globally. It is of interest to verify that the conclusions of therapeutic efficacy (and safety) apply consistently across subgroups of the clinical trial population

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# Setting: Scenario 1

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## Quote from Draft Guideline:

The clinical data presented are overall statistically persuasive with therapeutic efficacy demonstrated globally. It is of interest to verify that the conclusions of therapeutic efficacy (and safety) apply consistently across subgroups of the clinical trial population

### TENSION



Assume homogeneity  
unless there is significant  
evidence otherwise

Assume heterogeneity  
unless evidence supports  
homogeneity

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## How may we address consistency across subgroups?

- Goal: provide helpful information for prescribers and patients
  - Requires ...
    - Credibility of the conclusions
    - Must minimize important errors
    - some rigor in the process

# One Proposal

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## Influence Condition

- Introduced for use in confirmatory multipopulation tailoring trials (Millen et al, 2014a,b, 2012)
- May have applicability for the Scenario 1 discussion

## The Principle

- Let  $O = G+ \cup G-$
- Then, given primary inference of beneficial effect (efficacy) in population O, to support a broad indication for the treatment in population O,
- The beneficial effect must not be limited to only the G+ subpopulation

## Important Notes

- Requires *a priori* hypothesis for a marker (subgroup and its complement)
  - There is not a requirement of equivalent effects.
  - The requirement is for each subpop (G+ and G-) to have a positive effect.
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# Influence Condition

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## Application of the Influence Condition

- Looking for evidence of positive effect in the individual subgroups
  - Millen et al (2014a,b) propose metrics for this
  - Bayesian posterior probability
    - $\Pr(\theta_- > \lambda \mid \text{data}) > \gamma$
    - $\theta_-$  : treatment comparison parameter for marker negative subpop or “least benefitted” subpop
    - $\lambda$  : benefit threshold (0, 1, other?)
    - $\gamma$  : evidence threshold
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# Influence Condition

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## Application Details

- Setting the parameters
    - Control of influence error rates (Rothman et al 2012; Millen et al 2014a,b)
      - To what level?
  - For what subpopulation?
    - There should exist a reasonable prior hypothesis of
    - Size of subpopulation relevant?
  - Choice of prior (non-informative vs. informative)
    - informative vs. non-informative
      - based on earlier trials of the drug, external/literature data)?
  - Impact on Trial Design
    - Trial is now sized to meet the multiple objectives of overall effect and influence condition evaluation
      - Feasible?
      - At the trial level? Or at the program level?
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# Motivating Example

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Consider a clinical trial with the following assumptions or details

- 2 treatment arms: Drug vs Control
- Primary Endpoint is the difference of treatment means,  $\theta$ .
- There is a hypothesis that patients in G+ may be better responders to drug than patients in G-
- Apply influence condition as below. Satisfied if

$$\Pr(\theta_+ > 0 \mid \text{data}) \geq 0.75$$

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# Example

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## Operating characteristics

Overall sample size per arm	Relative size of subpop (G+)	$\theta+$	$\theta-$	Influence condition satisfied
133	60%	0.4	0.4	91.6%
200				97.0%
133	85%	0.4	0.4	71.5%
200				80.5%

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# Example

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## Operating characteristics

Overall sample size per arm	Relative size of subpop (G+)	$\theta+$	$\theta-$	Influence condition satisfied
133	85%	0.4	0.4	71.5%
200				80.5%
133	85%	0.418	0.3	58.9%
200				68.4%

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# Application Considerations

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## Evidence threshold

- 3:1, 2:1 generally?

## Size of subpops

- Feasible for small subpops?
- Oversampling/enrichment to increase amount of subpop data available?
  - Complicates reporting overall pop results

## Control of error rates

- False Positive (Influence errors) and False Negative

## Impact on design / feasibility

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# Other General Considerations

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Important that Scenario 1 evaluations are conducted for very **few** (e.g., 1 or 2) potential markers

- False positive concerns with multiplicity
- Impact on design, feasibility
- Should be done where there is prior hypothesis of potentially significant heterogeneity

Sponsor-regulatory alignment

- Pre-defined decision criteria are needed
- Not feasible to have SAWP meeting for every development program. Thus, detailed general guidance will be needed as soon as is practical

Hypotheses of differential effects should be discussed in SAP, rather than in protocol

- Potential to bias investigators when using protocol
  - Ability to be flexible and learn (from external sources) while trial is underway
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Transition

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# Transition

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Supportive



Scenario 2

Scenario 3

Discovery

## Setting: Scenario 3

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Context: Overall population result is negative, but there is a desire to find subpopulation(s) for which there is a positive B-R (and potential regulatory action)

### Proposals

Direct assessment of predefined subpop (if one existed)

Subgroup ID approaches

- Methodology fully predefined and automated (not requiring human intervention/judgment)
- Lilly approach: appendix of SAP
  - Machine learning tools
  - 'honest' estimates

## Setting: Scenario 2

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Context: Overall population result is positive, but there is a desire to find subpopulation(s) for which there is an improved B-R (and potential regulatory action)

### Proposals

Direct assessment of predefined subpop (if one existed)

Assess Subpops of more severe patients (risk)

- Using natural medical/clinical definitions
- Finding a cut-point along a biomarker
- Credibility of the resulting subpopulation is straight-forward

Subgroup ID approaches

- Methodology fully predefined and automated (not requiring human intervention/judgment)
- Lilly approach: appendix of SAP
  - Machine learning tools

# posing Comments

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## Complex issues

ence trial sponsors and [regulators] are put in a difficult position: whether to accept an assumption of homogeneity and disregard ... plausible findings in subgroups, or whether to anticipate some heterogeneity and, with appropriate caution and investigation, attempt to use the results of subgroup analyses as one piece of evidence to inform decision making.” (129- 133)

important to **recognize risks** of subgroup analyses and  
appropriately **limit use**

note that some sections seem to not adhere to this idea

consideration of feasibility, analyses at trial level may not be  
formative (particularly for **small subsets**)

## Research is needed

the proposals offered here require further research to increase understanding of operating characteristics and develop instructive guidance

research into methods in support of the aims of the guidance. The draft

## References

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Wassmann, M. D., Zhang, J. J., Lu, L., Fleming, T. R. (2012).



# BACK-UP SLIDES

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