

Applicability of novel methods WP5

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Novel methods developed within ASTERIX



Orphan conditions studies and EPARs



WP5 objective



- Determine **applicability, advantages & disadvantages**
 - **added value** of novel methods
- Provide **recommendations on methodology per cluster** of conditions

Six condition clusters

- Determine the drug development d
 1. Acute, single episodes
 2. Acute, recurrent episodes
 3. Chronic, stable/slow progression
 4. Chronic, progressive, one system/organ
 5. Chronic, progressive multiple systems/organs
 6. Chronic, staged conditions

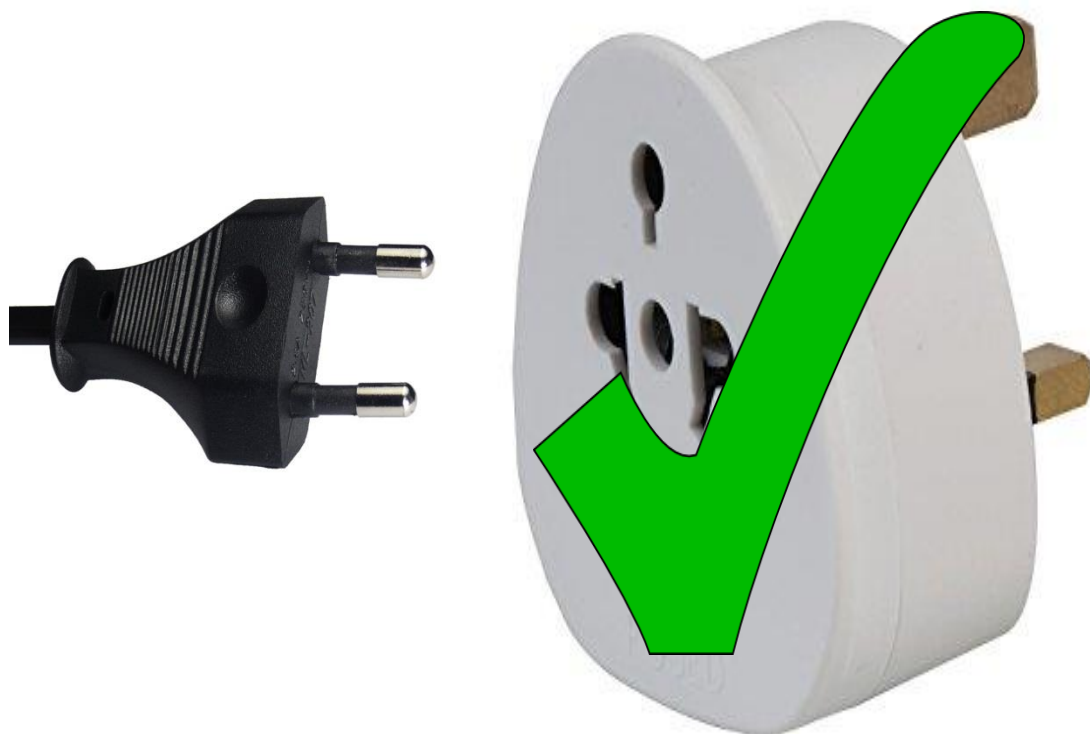
Methods included

- Innovative trial designs
- Study endpoints & statistical analysis
 - **Patient perspective: Goal attainment scaling (GAS)**
- Meta-analysis
- Level of evidence

Testing direct applicability



Testing applicability modifications allowed



Criteria list and pre-requisites

Essential/Critical pre-requisites

All below if applicable to current practice, otherwise NA and not mention/list

SPECIFIC

Context

a.1. Multi-arm group sequential designs with a simultaneous stopping rule

a.1.1. ≥ 3 arms including control (placebo)

number of arms in main trial(s)

a.1.2. ≥ 1 interim analysis

Interim analysis done Y/N, If so, reason: stopping for futility, overwhelming evidence of efficacy, safety.

a.1.3. developed for continuous endpoints, transportable to other types (i.e. Type of endpoint (primary EP) (binary, continuous) Composite? Time to event?

a.1.3.1. Type of (major) secondary endpoints (fill in as above)

a.1.4. Adaptive randomisation

a.1.6. > 1 (time to outcome faster than accrual rate) Delta time= recruitment - assessment (delay) /immediate or delayed response a.2.5.

a.1.7. Recruitment rate

a.1.8. Seamless design?

a.1.9. Adaptive design?

a.1.10. Allocation ratio

a.1.11. Did they allow dropping of arms

a.1.12. What was the control group?

a.1.13. MRCT? Multicentric? If Y, then how many?

a.1.14. Summary of models used in planning (e.g., disease progression, dropout, dose-response)

Study characteristics

- No of studies
- Type of (co-)primary endpoints/key secondary endpoints
- Recruitment pattern
- Time to outcome measurement
- Controlled or not
- Number of arms (if single arm: why?)
- Cross-over or not
- Acute or chronic condition

Example - Fabrazyme EPAR



- Cluster E: chronic, progressive, multiple systems/organs
- Fabry disease (ultra-rare): enzyme deficiency → accumulation of GL3
- Enzyme replacement therapy (exceptional circumstances 2001)
- Drug development:
 - Primary endpoint: reduction of GL3 accumulation (dichotomous)
 - Key secondary:
 - Change in GL3 in endothelium of kidney, skin and heart
 - Score of kidney tissue and urinary GL3 levels
 - 58 randomised patients

Methods' pre-requisites

Multi-arm multi-stage trial with simultaneous stopping rule

1. Time recruitment : Time outcome $> 1 / > > 1$
2. ≥ 1 interim analysis
2. Continuous outcomes (ideally, but transportable to binary)
3. ≥ 3 arms (at least 2 experimental arms + 1 placebo)

Group sequential design for small samples

1. Time recruitment : Time outcome $> 1 / > > 1$
2. Continuous outcomes (ideally, but transportable to binary)
3. Exactly 2 arms (treatment + control)

Method evaluation: 2-step approach



Step 1



- NO adjustments

Step 2



With adjustments
(reasonable)

Methods' pre-requisites – Step 1

Multi-arm multi-stage trial with simultaneous stopping rule

1. Time recruitment : Time outcome $> 1 / > > 1$ ✓
2. ≥ 1 interim analysis ✓
3. Continuous/binary outcomes ✓
4. ≥ 3 arms (at least 2 experimental treatments + 1 placebo) ✗

Group sequential design for small samples

1. T recruitment : T outcome $> 1 / > > 1$ ✓
2. Continuous/binary outcomes ✓
3. Exactly 2 arms (treatment + control) ✓

Methods' pre-requisites – Step 2

Multi-arm multi-stage trial with simultaneous stopping rule

1. Time recruitment : Time outcome $> 1 / > > 1$ ✓
2. ≥ 1 interim analysis ✓
3. Continuous/binary outcomes ✓
4. ≥ 3 arms (at least 2 experimental treatments + 1 placebo) ✓

Group sequential design for small samples

1. T recruitment : T outcome $> 1 / > > 1$ ✓
2. Continuous/binary outcomes ✓
3. Exactly 2 arms (treatment + control) ✓

Match method - study

Step 1:

- Group sequential design for small samples could be applicable immediately (Primary endpoint dichotomised from continuous) ✓
- Multi-arm multi-stage trial with simultaneous stopping rule not applicable (only 2 arms) ✗

Step 2:

- If we choose the continuous form of the primary endpoint (or e.g. urinary level of GL3 [1]) strongly applicable ✓
- Previous *phase I/II study explored multiple doses* for short-term only *but no optimal dose for longer term*, hence **multiple treatment regimens could have been tested in a MAMS** ✓

1. Thurberg BL, Rennke H, Colvin RB, Dikman S, Gordon RE, Collins AB, et al.

Globotriaosylceramide accumulation in the Fabry kidney is cleared from multiple cell types after enzyme replacement therapy. *Kidney Int.* 2002 Dec;62(6):1933–46.

Fabrazyme – potential impact



Potential advantages

- Quicker results
- Decreased placebo and non-inferior treatment exposure
- Control of type I error, maintaining power
- Increased precision for rejection boundaries
- Optimised use of available information

Potential disadvantages

- Increased logistic complexity
- Increased administrative and economic burden
- Sufficient evidence but not overwhelming
- Extra patients in case of effect size overestimation

Adjustments and reasons (1)

- Continuous endpoints preferred over binary
- Some rare conditions are rare versions of non-rare conditions
 - E.g. Tuberculosis – Sirturo
- Rare versions of cancer/tumor conditions
- Use hepatic or pancreatic biomarkers

Adjustments and reasons (2)

- For repurposed drugs a MAMS with a simultaneous stopping rule could be used (NSAIDs for patent ductus arteriosus)
- Multiple endpoints used to capture full clinical efficacy array
- Key secondary endpoints could be used as primary instead of secondary (6MWT/6MWD)

Heatmap of applicability



Heatmap Step 1



Heatmap Step 2

Step 1 - No adjustments

Cluster/topic	Level of Evidence (A)		Meta-analysis (B)		Innovative trial designs C						Study endpoints and statistical analysis (D)			
	Long-short outcomes	Evidence, eminence and extrapolation	Heterogeneity estimators	Prior distributions for variance parameters in sparse-event MA	Delayed-start randomisation	Sample size reassessment and hypothesis testing in adaptive survival trials	Multi-arm group sequential designs with a simultaneous stopping rule	Sequential designs for small samples	Bayesian sample size re-estimation using power priors	Dynamic borrowing through empirical power priors that control type I error	Fallback tests for co-primary endpoints	Optimal exact tests for multiple binary endpoints	Simultaneous inference for multiple marginal GEE models	GAS
Cluster A Acute: single episodes														
Cluster B Acute: recurrent episodes														
Cluster C Chronic: stable/slow progression														
Cluster D Chronic: progressive, one system/organ														
Cluster E Chronic: progressive, multiple systems/organs														
Cluster F Chronic: staged disease														

The proposed method is applicable

The proposed method may be applicable

Limited or no applicability of the method

Step 2 - With adjustments

Cluster/topic	Level of Evidence (A)		Meta-analysis (B)		Innovative trial designs C						Study endpoints and statistical analysis (D)			
	Long-short outcomes	Evidence, eminence and extrapolation	Heterogeneity estimators	Prior distributions for variance parameters in sparse-event MA	Delayed-start randomisation	Sample size reassessment and hypothesis testing in adaptive survival trials	Multi-arm group sequential designs with a simultaneous stopping rule	Sequential designs for small samples	Bayesian sample size re-estimation using power priors	Dynamic borrowing through empirical power priors that control type I error	Fallback tests for co-primary endpoints	Optimal exact tests for multiple binary endpoints	Simultaneous inference for multiple marginal GEE models	GAS
Cluster A Acute: single episodes														
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Cluster C Chronic: stable/slow progression														
Cluster D Chronic: progressive, one system/organ														
Cluster E Chronic: progressive, multiple systems/organs														
Cluster F Chronic: staged disease														

The proposed method is applicable

The proposed method may be applicable

Unfitted or no applicability of the method

Application of methods

Evaluation exercise summary



- First step – some immediate applicability
- Second step - exercise ***flexibility*** – gain applicability + advantages
- There is room for the novel methods to improve the designs for small populations trials
- Recommendations by cluster of conditions

What's next?

Katrien Oude Rengerink

GUIDELINE ON CLINICAL TRIALS IN SMALL POPULATIONS

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Update?

What can we add to the guideline?



- Updated, more specific guidance including novel methods
 - Including literature since 2006
 - ASTERIX
 - IDEAL
 - Inspire
- Use clustering to tailor guidance

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Methodological and statistical considerations

Paragraphs

6.1. Design stage

6.2. Data analysis

6.3. Reporting

Sequential designs

Sequential designs – with a goal to demonstrate ‘statistical significance’ if a treatment is genuinely superior to control - generally reduce the required sample size. There can be several different types of sequential design – all providing valid statistical conclusions but each tailored to specific balances of expected outcomes and patient availability. Some designs are ‘open-ended’ and (in theory) continue to recruit patients until a reliable positive or negative conclusion about the treatments can be made.

Sequential designs, as with response-adaptive designs, require treatment outcomes to be available quickly (relative to the patient recruitment rate). This will almost never be the case if we are looking for long-term survival data, for example, but may be the case if we are looking at shorter term clinical or surrogate/bio-markers

slow because patients are so rare; hence such methods may have more of a place in these situations than in more common diseases. Ultimately, however, the extent to which the sample size can be reduced depends on the size of the effect. Variations on sequential methods are the, so-called, group-sequential methods and adaptive designs. See also draft guidance on adaptive designs (CHMP/2459/02).

		Delayed-start randomisation	Sample size reassessment and hypothesis testing in adaptive survival trials	GAS
Cluster A Acute: single episodes	Savene			
	Pedea			
	Defitelio			
	Sirturo			
Cluster B Acute: recurrent episodes	Ilaris			
	Cayston			
	Xyrem			
	Diacomit			
Cluster C Chronic: stable/slow progression	Revestive			Added value?
	Plenadren			Added value?
	Xagrid			
	Glybera			
Cluster D Chronic: progressive, one system/organ	Soliris			Added value?
	Wilzin			
	Sicklos			
	Glivec			
Cluster E Chronic: progressive, multiple systems/organs	Fabrazyme			
	Orphacol			Added value?
	Tracleer			
	Zavesca			
Cluster F Chronic: staged disease	Afinitor			
	Opsumit			
	Litak			
	Revlimid			

Limitations

- Few EPARs evaluated per cluster
 - If no applicability in heatmap: not impossible
 - If applicable in all cases: not always possible
 - Necessary to combine heatmap with methods pre-requisites
- Clustering helpful, still heterogeneity within clusters

Dis



	EPAR OMP	Delayed-start randomisation	Sample size reassessment and hypothesis testing in adaptive survival trials	GAS
• Cluster A Acute: single episodes	Savene			
	Pedea			
	Defitelio			
	Sirturo			
• Cluster B Acute: recurrent episodes	Ilaris			
	Cayston			
	Xyrem			
	Diacomit			
Cluster C Chronic: stable/slow progression	Revestive			Added value?
	Plenadren			Added value?
	Xagrid			
	Glybera			
Cluster D Chronic: progressive, one system/organ	Soliris			Added value?
	Wilzin			
	Sicklos			
	Glivec			
Cluster E Chronic: progressive, multiple systems/organs	Fabrazyme			
	Orphacol			Added value?
	Tracleer			
	Zavesca			
Cluster F Chronic: staged disease	Afinitor			
	Opsumit			
	Litak			
	Revlimid			

