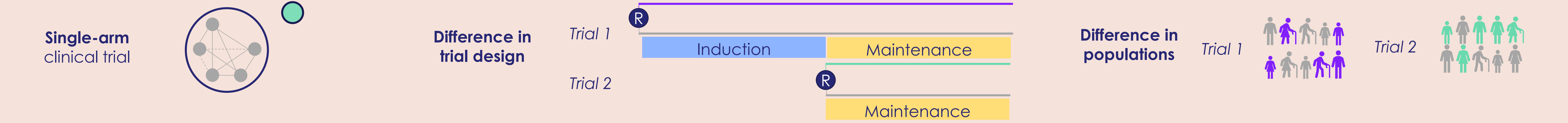


BACKGROUND

- Indirect treatment comparisons (ITCs) have been developed to allow generating **relative treatment effects vs all relevant comparators**, beyond the comparators selected in the pivotal trial.
- Standard ITC methods require **homogeneity across trials** in terms of **treatment effect modifiers**, as well as **connectivity of network of evidence**.
- However, in some cases **data** available to conduct the ITC is **non-comparative**, limiting the use of standard methods:



METHODS & OBJECTIVES

- Recommendations from the NICE Technical Support Documents (TSDs) were reviewed and previous experiences encountered within Amaris were considered.^{1,2}
- To suggest **alternatives** when dealing **with non-comparative data**.
- To provide **guidance** on key **considerations at feasibility assessment** phase to define relevant ITC approach.

Results – Methods to perform ITCs with non-comparative data

- Different **factors** will help identifying the **most appropriate approach** to conduct the ITC, and should be considered during **feasibility assessment phase**:



IPD available for pivotal trial only

Matching-Adjusted Indirect Comparison (MAIC) vs **Simulated Treatment Comparison (STC)**

Objective
Comparing indirectly two treatments not directly compared in clinical trials to overcome heterogeneity or non-connectivity issues.

Principle
Estimating relative treatment effect between two treatments using IPD from one trial and aggregated data for the second one while adjusting for effect modifiers (and prognostic factors depending on anchored status of the ITC)

- MAIC: similar to a propensity score weighting method
- STC: based on outcome model fitted to IPD as function of covariates

Why conduct MAIC and/or STC?

- Unanchored ITC
- High level of heterogeneity between studies

Key assumptions

- All effect modifiers (and prognostic factors for unanchored ITCs) are known and observed in both trials so that they can be adjusted for.
- Assumes that the trial with IPD includes the competitor's trial population.
- Results obtained only valid in the population of comparator's trial.

Additional assumptions for STC

- Use of identity link function, potential bias if done on another scale.
- For time-to-event outcomes: assumes a parametric distribution.
- Assume that parameters of the regression estimated on trial with IPD available are applicable to competitor's population.

What type of data to inform on comparator in MAIC/STC?
Both clinical trials, as well as observational studies, as long as enough data is available

What if comparators' are connected in a network?
Results from MAIC/STC related to one comparator of this network can be used to connect pivotal trial to the global network

IPD available for both trials

Propensity score approaches in brief

Introduced by Rosenbaum and Rubin in 1983³ to address the issue of selection bias in observational research.

Objective
Aim to make treatment groups comparable or to control for the effect of prognostic factors on outcomes.

Principle
The conditional probability of assignment to a particular treatment given a vector of observed covariates.

Key steps

- Estimate propensity score across all patients via a logistic regression
- Ensure sufficient overlap between groups
- Select adjustment method, main options: matching, weighting, stratification, doubly robust

AI to generate propensity scores: our experience

Objective: To compare methods (logistic regression vs machine learning [ML]) to estimate propensity scores when constructing an external control arm using oncology data from Project Data Sphere®

Methods compared

- Logistic regression (standard approach)
- Classification and Regression Trees (CART)
- Naïve Bayes classifier
- Support Vector Machines

Project steps

- Targeted literature review
- Identification of data sources
- Comparison of survival and baseline characteristics
- Methods to evaluate the propensity score
- Comparison of algorithms
- Dissemination of findings

Choose data with sufficient overlap in baseline demographic and clinical characteristics
Using data from the control arm of two data sources in order to compare performance of methods

To identify the most appropriate machine learning algorithms to apply and what prognostic factors should be considered

In terms of covariate balance and homogeneity of survival distributions

AI to generate propensity scores: our findings

Data source
Use of historical clinical trial data generally allows for more clinically relevant variables, not available in larger claims and/or electronic health records databases and more overlap between original populations.

Sample size & variables
Given limited number of variables and small sample size, ML algorithms generally did not perform better than logistic regression to create an external control arm.
Having all relevant prognostic variables available in both data sets is key to a successful model!

Predictive power
More patients were excluded when using ML algorithms due to their high predictive power.

DISCUSSION

- Towards personalized medicine:** new drug development focused on rare diseases and target populations often justifies non-RCT designs – especially in disease areas such as oncology or hematology.
- Alternatives** can be considered to conduct ITCs, should they be conducted in line with recommendations, built on a strong rationale and validated clinically.

