

TABLE OF CONTENTS.

- ► EXPOSURE study
- Measures to prevent missing data
- Missing data mechanisms
- Measures to handle missing data



CONTEXT

Pulmonary Arterial Hypertension (PAH):

- rare disease affecting 15–26 cases per million adult inhabitants
- life-threatening condition that gets worse over time
- treatments can help the symptoms so patients can live better and longer with the disease
- 3 treatment pathways, very few naïve patients

Uptravi®:

- new oral prostacyclin receptor agonist approved for PAH treatment
- can be prescribed alone or in combination with other PAH-specific therapies



EXPOSURE

DESIGN, SOURCES FOR MISSING DATA



DESIGN

- Multicentre, prospective, real-world, observational cohort study
 - Post-Authorisation Safety Study listed in the EU-PAS register (ENCePP)
- ► In PAH adult patients newly treated with a PAH-specific therapy
 - Uptravi (n=1450), or
 - any other PAH-specific therapy (n=1850)
- ▶ Patients are followed-up for at least 18 months, or until the earliest of death, withdrawal of consent, loss to follow-up or study end
- Data collection
 - extracted from medical charts of PAH centres
 - includes all visits and hospitalisations during obs. period



OBJECTIVES

- **1.To describe** demographics, disease characteristics and clinical course in PAH patients
- 2.To further characterise Uptravi safety profile
- 3.To compare rates of major adverse cardiovascular events (MACE) and all-cause death
 - between
 - patients newly treated with Uptravi (*Uptravi cohort*), and
 - patients newly treated with any other PAH-specific therapy who were never treated with Uptravi (Other PAH therapy cohort)
 - using an analysis that includes stratification by propensity scores



DATA ANALYSIS

WHY USING PROPENSITY SCORES?

Uptravi treatment is not randomly assigned between cohorts

- Uptravi cohort and other PAH therapy cohort may differ with respect to multiple covariates (e.g., disease severity, type of centre, patient decision, health care system variability) which precludes valid comparison
- Other PAH therapy cohort
 - PAH patients who have not been treated with Uptravi but have had the same opportunity to have been prescribed Uptravi
 - contemporaneous to the *Uptravi cohort*
 - recruited in same countries and centres, all able to prescribe Uptravi



WHY IT MATTERS

Selective statistical procedures are sensitive to the amount of missing data When a data set is incomplete, the data analyst has to decide how to deal with it

Propensity score method requires no missing data for observed influential covariates Missing data can lead to patient exclusion from analysis



SOURCES FOR MISSING DATA

- Demographics /clinical characteristics missing (e.g., gender, weight, disease etiology)
- Clinical / Lab assessment not performed:
 - as per local clinical practice (e.g., 6 Minutes Walking Distance in centres not adapted to this test)
 - due to patient disease severity (e.g., 6 Minutes Walking Distance for patients in wheelchair)
- Clinical or lab assessment performed, result missing / out of range
 - result sheet not available in med chart
 - issue with units
- Other medical speciality assessment/exam missing (e.g., Right Heart Catheterization, ophthalmological exams)
- Lost to follow-up (e.g., patient relocation to another centre)





AT STUDY DESIGN / eCRF DESIGN (1/3)

- Study design
 - Limit the collection of data to those who are participating in the study
- Disease management knowledge
 - State of the art vs clinical practice
 - Discussion with physicians & site personnel to identify clinical / lab tests performed per routine care practice to assess study outcome
 - May have to redefine study outcome (e.g., use of a proxy)



AT STUDY DESIGN / eCRF DESIGN (2/3)

- Check where data are located to identify who should collect them
 - Investigator medical chart only
 - Synergy between several physician specialties
 - Patient reported data
- Reduce the likelihood of "lost to follow-up patients"
 - Include to the protocol a specific process to follow patients with no data collected since X months (e.g., family contact, National death registry)
 - Plan a patient data transfer between sites in case of patient' relocation



AT STUDY DESIGN / eCRF DESIGN (3/3)

- eCRF settings
 - User friendly
 - Follow medical chart flow
 - Use clinical terminology rather than research wording
 - Include "unknown" or "not applicable" option
 - Avoid free text, use preselected answer lists
 - Include potential bridges with other observational studies (e.g., patient ID)
 - Collect raw data, derive needed data (e.g., BMI)
 - Collect lab test results interpretation rather than value (e.g., anemia status vs haemoglobin value)
 - Pre-programmed automatic edit checks (in and between visits collected)
 - Auto-populated fields for Adverse Drug Reaction form
 - Test eCRF in real-world environment



AT STUDY LAUNCH

- Association with networks & registries
 - Establish win-win collaborations: raise investigators interest
 - Investigators options for publications
 - Scientific community interest vs Health Authorities requirements
 - Plan data transfer solutions to avoid double data entry
- Site training
 - Appropriate training on study material and importance of capturing all available data
 - Identify data entry personnel
 - Create exchange events (e.g., site initiation visits, investigator meetings)



DURING STUDY CONDUCT

- Data quality plan
 - Identify critical items with high impact on the robustness of propensity score and study overall
 - Regular review of missing data proportion per site and variable
 - Visualisation tools to portray the pattern of missing data
 - Include scientific/medical plausibility reviews
- Site monitoring
 - Clinical Research Associates assigned per study site
 - Centralised monthly remote monitoring (via EDC system, telephone calls)
 - Additional on-site visits
 - Data entry performance thresholds (time interval of data entry)
 - Follow-up on queries





Three missing data mechanisms:

- Missing Completely At Random (MCAR)
- Missing Not At Random (MNAR)
- Missing At Random (MAR)

Each of the 3 mechanism describes one possible **relationship** between the **propensity of data to be missing** and **values of the data**

Mechanisms generally not directly testable but justifiable through understanding and describing the process that generated the data



MISSING COMPLETELY AT RANDOM (MCAR)

No relationship between the propensity of a data to be missing and any values, observed or missing

Those missing data points are a random subset of the data

There is *nothing* **systematic** going on that makes some data more likely to be missing than others

No bias introduced by data MCAR but potential loss of statistical power



MISSING NOT AT RANDOM (MNAR)

There is a relationship between the propensity of a data to be missing and its values

e.g. - young people without CV disease have no/less blood pressure measurement recorded

Potential bias introduced by data MNAR: Patients with missing data may behave differently from those without missing data



MISSING AT RANDOM (MAR)

The propensity for a data to be missing is not related to the value of the missing data, but is related to some of the *observed* data

Missing data is conditional on another variable

e.g. - if men are more likely to tell you their weight than women, weight is MAR (assumption: weight is not correlated to gender)

Potential bias introduced by data MAR: Specific groups of patients may be excluded from the analysis



Why it is important?

- it affects how much the missing data bias the results
- handling approaches have different assumptions about the mechanism of missingness

If unknown mechanism

- use assumption that data are Missing At Random
- investigate the assumption using sensitivity analysis on the pattern of missing data





Data MCAR: Complete case analysis

Excludes all patients with missing data in at least 1 influential covariate

- the most common decision
- easy to use, default option in most statistical packages
- only results in unbiased estimates if missing data are unrelated to the study treatment or design (MCAR)
- lead to loss of power



Data MAR and MCAR: Multiple imputations

Filling in plausible values for missing data

e.g., in a longitudinal study, if a patient's condition was consistently improving over previous visits and all of a sudden patient drops out, then it might be plausible to assume the patient is doing better (based on previous information)

- Used if missing data may potentially be related to observed covariates but not those unobserved (MAR)
- Can also be used for MCAR if complete case analysis is not powerful enough.
- Multiple Imputation generates multiple simulated datasets (Monte-Carlo technique)
- Propensity score analysis is conducted in each imputed dataset
- Scores across imputations are summarized accounting imputation variability



Data MNAR Pattern mixture model

Retain all the patients in an analysis by grouping them based on observed covariates (patterns of missing data)

- Valid method if there are clearly defined groupings of patients with missing data related to study design (MNAR)
- Propensity score estimation by patterns of missing data
- Separate propensity score model for each distinct missing covariate data pattern
- Patterns may be pooled together if the number of observations is not sufficient to run the propensity score model



CONCLUSION FOR EXPOSURE

Prerequisite to apply Propensity scores to EXPOSURE:

- Maximize efforts to avoid missing data at study design and data collection
- Understand missing data mechanisms in place
- Conduct statistically valid handling method(s) with appropriate mechanisms and assumptions for the missing data
- Apply Propensity score using each handling method selected
- Preplan Sensitivity analysis to check result consistency



THANK YOU.

