

PAGE 2024, Rome

Virtual Patient Cohorts using Modeling and Simulation to Support Drug Development in Rare Diseases

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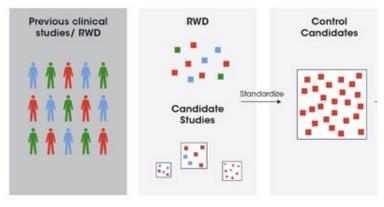


- Increased need for tools to boost drug development and adoption in rare diseases is acknowledged by regulators
- Novartis is dedicated to develop sound approaches to support this effort, PMX group has been actively involved in various Consortia
- For rare diseases, developing methodologies to contextualize the readouts of single arm trials is very challenging also due to small sample size
- We present how PKPD models can be used to generate "virtual" control patients on example of iptacopan in Paroxysmal Nocturnal Hemoglobinuria to support drug approval and potential post-approval clinical adoption

Rare diseases: large, heterogeneous, underserved patient community

- Affect ~350 million people worldwide (about 4% of entire population), 50% are children
- Only 5% of diseases have approved treatments
- High societal and economic burden
- Urgent need for better ways to diagnose and to treat their disease
- Single arm trials →
 - No direct comparison with placebo or an active control
 - Protracted drug approval cycle
 - Limited clinical adoption of new therapies

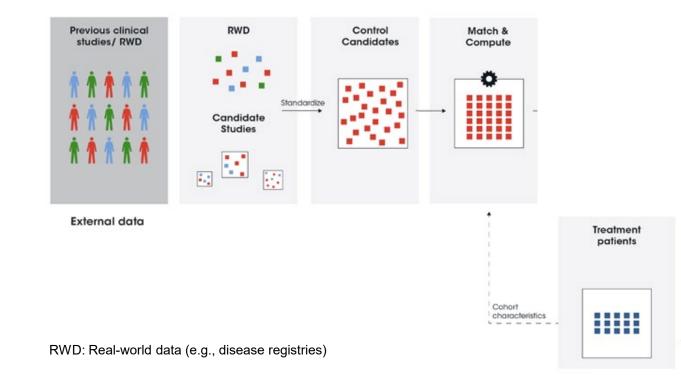
Standard approach: Synthetic Control Arm



External data

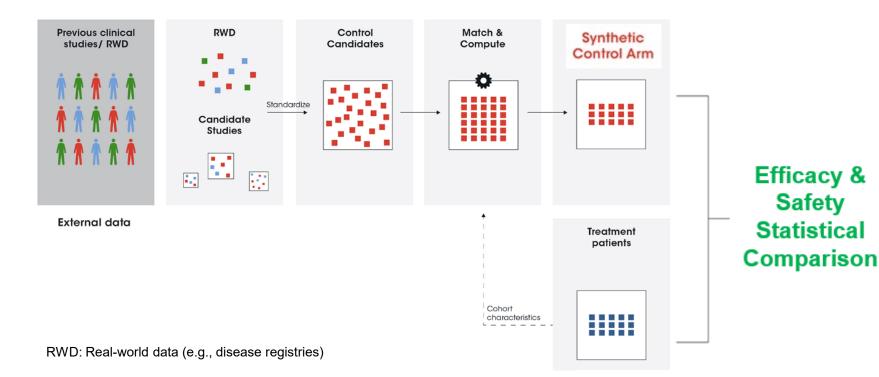
RWD: Real-world data (e.g., disease registries)

Standard approach: Synthetic Control Arm

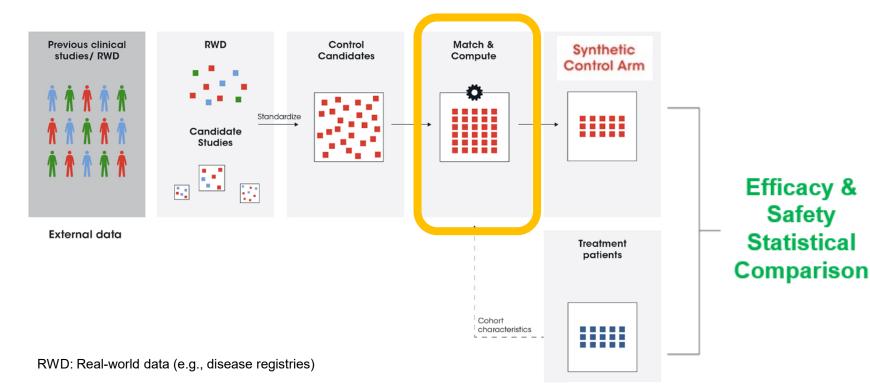


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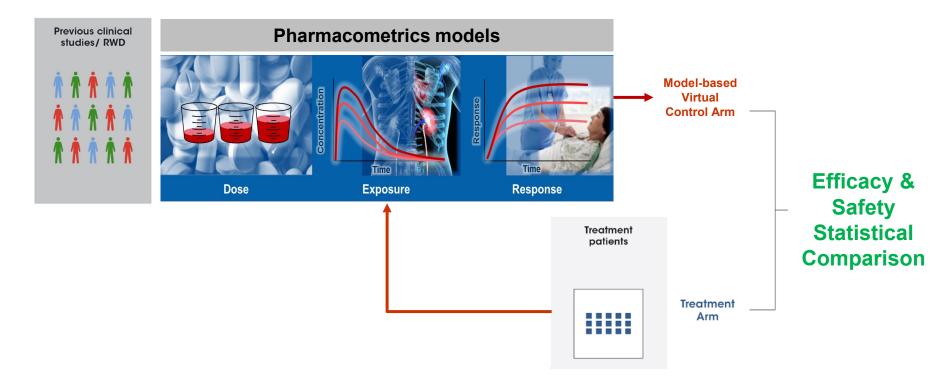
Standard approach: Synthetic Control Arm



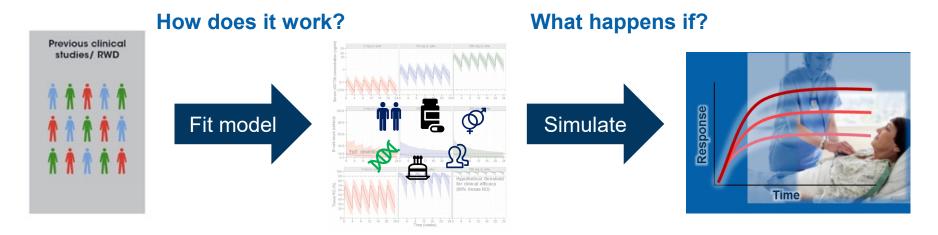
Synthetic Control Arm: several salient limitations



Our objective: generate 'Virtual Patients' using PMX modeling



Advantages of model-based Virtual Arm

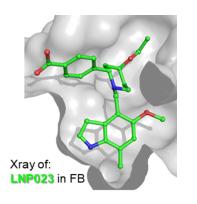


- Disease and response dynamics vs. cross-sectional clinical endpoint evaluation
- All data vs. selected sample
- Account for confounders, reduce bias
- Imputation of missing data from the model
- Scenario evaluation

Case study: iptacopan Pipeline in a pill

Iptacopan/LNP023

 Iptacopan, an oral, selective and potent first-in-class factor B inhibitor targeting the alternative complement pathway and acting upstream of C5



Paroxysmal nocturnal Hemoglobinuria (PNH)

- Genetic (somatic) mutation in HSCs; resulting in red blood cell hemolysis & thrombosis (potentially life-threatening)
- Rare disease: US incidence ~1.3/million
- Standard-of-care (SoC) is anti-C5 therapy (high treatment burden)
- Over 80% of patients suffer anemia (Hb < 10 g/dL) and many remain transfusion-dependent despite anti-C5 therapy [3]



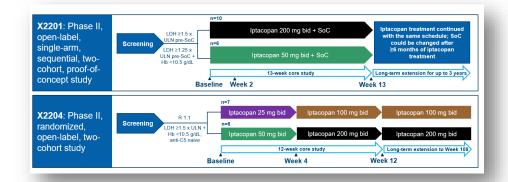
Hemolytic Erythrocyte with complement MAC

Other indications

- IgAN (renal)
- C3G (renal)
- IC-MPGN (renal)
- aHUS (hematology)
- Etc.

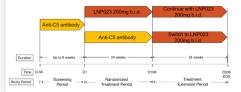
Development program of iptacopan in PNH

- Two Phase 2 trials
 - X2201: Iptacopan 200 mg bid as addon to anti-C5 treatment [4] (NCT03439839)
 - X2204: Iptacopan as monotherapy in treatment-naive patients [5] (NCT03896152)
- Two Phase 3 trials
 - APPLY-PNH: iptacopan 200 mg bid vs anti-C5 in treatment-experienced patients (NCT04558918)
 - APPOINT-PNH: iptacopan 200 mg bid in treatment-naive patients, open label single arm (NCT04820530)



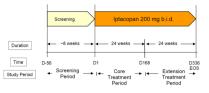
APPLY-PNH: anti-C5 experienced (n = 92)

A randomized, multicenter, active-comparator controlled, open-label trial to evaluate efficacy and safety of oral, twice daily LNP023 in adult patients with PNH and residual anemia, despite treatment with an intravenous anti-C5 antibody





A multicenter, single-arm, open-label trial to evaluate efficacy and safety of oral, twice daily iptacopan in adult PNH patients who are naive to complement inhibitor therapy



PKPD modeling for iptacopan in PNH



- Longitudinal modeling of Hb data (4 PNH studies of iptacopan, n=161)
- Model adequately characterized one of the components of the primary endpoint, haemoglobin (Hb) response
- > Was used to rationalize the registered dose (200 mg bid) in the submission dossier

Iptacopan PKPD model for Hb

$$\frac{dY(t)}{dt} = k_{in} - k_{out}(t)Y(t), \text{ where}$$

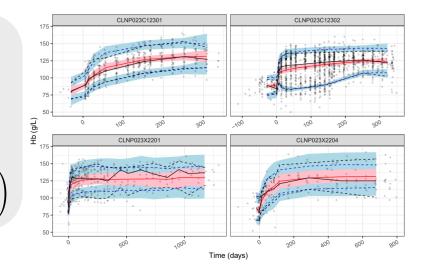
$$k_{out}(t) = \frac{k_{in}}{Y_{base}} \left(1 - \left(\frac{Y_{TRT} - Y_{base}}{Y_{TRT}}\right)S(t)\right)$$

$$S(t) = \left(\frac{C^{\gamma}(t)}{EC_{50}^{\gamma} + C^{\gamma}(t)}\right)$$
At steady state, $Y_{SS} = Y_{Base} + (Y_{TRT} - Y_{Base}) \left(\frac{C_{SS}^{\gamma}}{EC_{50}^{\gamma} + C_{SS}^{\gamma}}\right)$

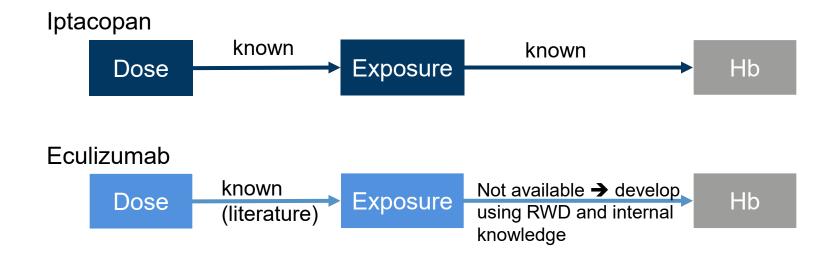
- $Y(0) = Y_{Base}$: Hb level before the start of treatment
- *Y_{TRT}*: maximally improved Hb level on treatment
- k_{in}: production rate of red blood cells (RBCs)
- k_{out}: turnover rate of RBCs
- S(t): drug effect function inhibits elimination
- EC_{50} : iptacopan concentration at which the half-maximal effect is reached

 For more details on the PK-Hb modeling, visit poster I-020





Leveraging internal data and SoC from real world data (RWD)



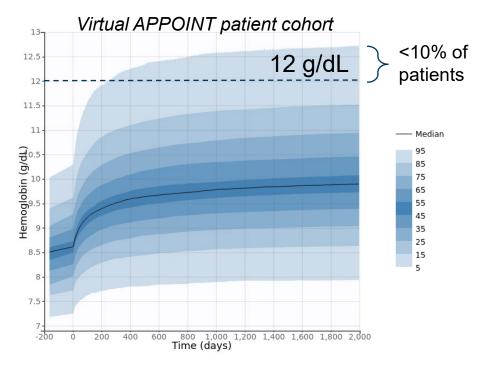
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Exposure-response model for Hb using internal/ external knowledge

- SoC: anti-C5 mAbs eculizumab/ravulizumab
- PKPD model for Hb using published literature data:
 - 1-cpt PK model with CL and V values fixed to values from literature [1, 3]
 - Indirect response model for Hb:
 - ➢ Kin fixed to a value from our internal model for iptacopan
 - Emax and EC50 values derived from RWD

Virtual Patient Cohort generation Simulation of Hb response under SoC

- Simulate the Hb response of patients as in iptacopan APPOINT-PNH single-arm trial (n=40, adjusted for sex and baseline Hb), but as if they were treated with SoC (eculizumab)
- Eculizumab dosing as per label
- APPOINT-PNH readout: iptacopan achieved Hb levels of at least 12 g/dL without the need for red blood cell (RBC) transfusions in 62.8% (95% CI, 47.5%-77.5%) of patients
- Simulation results generate evidence of **better anemia control under iptacopan** as compared to SoC and hence allow to contextualize the results of APPOINT



INVENTS Consortium

Innovative designs, extrapolation, simulation methods and evidence-tools for **rare diseases** addressing regulatory needs <u>https://ecrin.org/projects/invents</u>

- Funding: Horizon Europe (N 101136365)
- **Budget**: €6M
- Coordinator: INSERM
- Duration: 5 years (Jan 2024 Dec 2029)
- Overall objective: To provide clinical trial stakeholders, trialists and regulators with a generalizable framework encompassing methods, workflows and evidence-tools to improve the level of evidence in regulatory decision making in rare diseases.
- Framework to be used to make the most of sparse evidence, to synthesize (seemingly) disparate data, and to robustly bridge between more or less well-understood indications



Conclusions

- Modeling and Simulation techniques using Clinical Trial Data complemented with RWD allow to generate Virtual Patient Cohorts and to substitute SoC information missing in single arm trials
- Major advantages over other techniques are:
 - Allow for indirect comparison while accounting for confounders (differences in MoA, trial design and populations btw. clinical trial and RWD) inference in the situations where data is sparse
 - Allow for knowledge leveraging between clinical trial and RWD
 - Allow for scenario evaluation beyond those observed to address questions of interest to a clinician, regulator, or payer
- Joint work with Regulatory Agencies, HTA bodies, Academia and Pharma will further enhance the acceptance of Virtual Patient Cohorts as a tool for drug approval and post-approval adoption

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- INVENTS Consortium

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Thank you

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