A Biomarker-Focused QSP Model of Complement Alternative and Terminal Pathways to Evaluate Potential Targets for Therapeutic Impact in Complement-Associated Diseases: Paroxysmal Nocturnal Hemoglobinuria (PNH) as a Case Study

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OBJECTIVE & PURPOSE

- COMPLEMENTsym, QSP model, leveraged Complement overactivity has been implicated in published models¹⁻⁴ and publicly available data multiple diseases, including PNH informing complement pathways and kinetics PNH is associated with overactivity of the
- complement alternative pathway, driven by deficiencies in regulatory proteins
- Therapeutic targeting of complement is complicated by feedback loops, redundant functions, and availability of druggable targets
- Simulated PNH disease state developed including increased convertase dissociation rate as a proxy Here, a QSP model, COMPLEMENTsym[™], focused on for reduced CD55 alternative and terminal pathway complement SimPops (N>2000) NHV incorporates PNH analytes in circulation is described, including disease specific variability in addition to simulated populations (SimPops[®]) of normal healthy modifications, qualified against data for 4 analytes volunteers (NHVs) and PNH patients Exemplars optimized (and validated when possible) • This work aimed to evaluate the impact of exemplar against published analyte or hemolysis data
- treatments to validate the model for predictive use

RESULTS

Eculizumab is a monoclonal antibody binding C5, reducing complement activity by the terminal pathway. suppressing Eculizumab was simulated with average exposure profile for 600 mg weekly for four weeks followed by 900 mg every other week.



Fig 3. Eculizumab representation optimized to free C5 during clinical dose administered to treatment-naïve PNH simulated population⁵.

Pegcetacoplan

Pegcetacoplan targets the C3/C3b binding site, suppressing AP activity. In COMPLEMENTsym, the pegcetacoplan representation is based on clinical data⁷. Simulated pegcetacoplan treatment (270 mg/day) in treatment-naïve PNH SimPops and eculizumab-treated



PNH SimCohorts recapitulated published C3 elevations without optimization^{8,9}.

Fig 5. Simulated C3 concentrations during pegcetacoplan treatment in patients with eculizumab background compared to Day 1 and Day 169 clinical data⁸.

METHODS

NHV SimPops (N>5000) developed based on publicly available data supporting known variability in key parameters, qualified against clinical data ranges for 13 analytes



Fig 4. AP-driven hemolysis representation optimized to reproduce reduced serum lysis during eculizumab administration⁶.

Iptacopan targets the active site of Factor B, suppressing AP activity. Simulated iptacopan in treatmentnaïve PNH SimPops was optimized to reproduce published reductions in Bb¹⁰. A PNH cohort was developed to match sC5b-9 concentrations of PNH patients on eculizumab prior to treatment with iptacopan¹¹. Simulating iptacopan (200 mg twice a day) in this cohort reduces average Bb from 3.29 ug/mL to 1.08 ug/mL over 13 weeks, consistent with published reductions in Bb from 4.871 ug/mL to 1.198 ug/mL¹¹.

Predicted sC5b-9 Outcomes

Treatment-naïve PNH SimPops with simulated exemplars, assuming average exposure, different demonstrates responsiveness as quantified by sC5b-9 concentrations.



Fig 7. Simulated sC5b-9 concentrations for NHV and PNH SimPops baselines compared to post-treatment values for 24 week simulations of treatmentnaïve PNH patients on eculizumab, pegcetacoplan, and iptacopan monotherapies.



CONCLUSION

- SimPops demonstrate consistency with complement analytes in health and disease
- Disease representation is strengthened by reproducing clinical data from PNH patients treated with eculizumab, pegcetacoplan, and iptacopan
- This work establishes a model and framework to evaluate novel targets and compounds for complement-associated diseases

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