

Semi-physiologic Population PKPD Model Characterizing the Effect of Bitopertin (RG1678) Glycine Reuptake Inhibitor on Hemoglobin Turnover in Humans

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PAGE 2012, abstract #2553, Venice, 06 June 2012

Quality Teamwork Unity Bench to bedside and back again Passion Patient focus Pride Medical Need Focus Personalized treatment Empowerment Excellence in execution Integrity Aim for Cure Academic collaborations Courage Cutting edge science Quality





Background

- Bitopertin is a selective glycine reuptake inhibitor targeting the glycine type 1 transporter (GlyT1)
 - Currently in late stage development as a potential novel therapeutic approach in schizophrenia
 - In the brain, inhibition of GlyT1 results in increased synaptic glycine levels leading to improved NMDA* receptor function
- Outside the brain, GlyT1 is also localized on erythrocyte precursors in the bone marrow and on circulating reticulocytes
 - Ø Glycine is required for hemoglobin (Hb) synthesis
 - Complete ablation of GlyT1 in knock-out mice results in a 26 % Hb decrease 1
- The **hematological effect of bitopertin** was studied in a 4 month phase 1 study in healthy subjects



Bitopertin Effect on Hemoglobin Synthesis



Proposed mechanism: reduced Hb synthesis is due to reduced glycine uptake in erythrocyte precursors



Objectives



A semi-mechanistic PKPD model for bitopertin was developed in order to

- Characterize the proposed effect of GlyT1 inhibition on the hematological system, taking into account hemoglobin synthesis and red blood cell turnover
- Support drug development with respect to the effect of bitopertin on hemoglobin concentrations as an important clinical parameter





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Phase 1 Study in Healthy Subjects

To assess the hematological effects of 4 month treatment



Population:

- 67 healthy male and female subjects enrolled
- Aged 18 to 45 years; baseline Hb within normal lab range



Hematological Data Observed in Healthy Subjects

Hemoglobin blood concentration

Hb (g/L)



- Dose dependent decrease of Hb
- Reversible effect
- Plateau of effect delayed due to RBC lifespan
- No subject reached Hb discontinuation threshold (100 g/L in females, 110 g/L in males)
- About one week to PK steadystate



Hematological Data Observed in Healthy Subjects

Hb (g/L)





Semi-physiologic PKPD Model for GlyT1 Inhibitory Effect on Hemoglobin



Estimated Parameters:

KinRBC, KinHb, LS, Emax, AUC50, Feedback, SexEff

- Two parallel chains of transit compartments¹ represent the production, senescence and elimination of RBC, with their respective MCH
- AUCss is driving the inhibitory drug effect
- Model fit to MCH and RBC data simultaneously in NONMEM 7

Model Diagnostics: GoF Plots





Model Predictive Performance for Hemoglobin

VPC for hemoglobin, predicted as cumulative RBC x MCH



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Results



PKPD parameter estimates

	Unit	Estimate	% RSE	IIV (% CV)
LS	days	126	4.5	24.9
KinHB	pg/cell/day	0.949	4.8	28.0
KinRBC	10^12/L/day	0.039	4.3	24.3
Feedback		1.38	12.4	
Emax	fraction	*	13.2	
AUC50	mg/L*h	*	23.2	49.2
SexEff		-0.119	8.0	
ERR_RBC	10^12/L	0.183	3.0	
ERR_MCH	pg/cell	0.346	2.7	

Physiological parameters estimates generally in line with values in the literature

	Baseline value	es (derived)		
		Unit	males	females
\rightarrow	MCH ₀	pg/cell	29.8	29.8
	RBC	10^12/L	4.9	4.3
	HB ₀	g/L	147	129

% RSE: Relative Standard Error

Feedback mechanism (FDB):

Scaling parameter to describe the stimulation on RBC production rate as a feedback to Hb reduction KinRBC = KinRBC₀*[1+(HB₀-HB)/HB₀*FDB]

SexEff: Female patients have 11.9 percent lower KinRBC



What is the predicted long term effect on Hb?

Doses up to 20 mg tested in phase 3

Predicted Hb change from baseline (%) for 360 days of treatment



Maximum Hb change from baseline (nadir) expected shortly after one RBC life span Less than 10 % Hb drop expected at projected therapeutic doses

Conclusions



- The physiology of the hematopoietic system together with the inhibitory effect of bitopertin on Hb synthesis were integrated in the semi-physiologic PKPD model to successfully fit the data from healthy subjects
- The effect on hemoglobin, as the key clinical parameter, is best predicted by fitting MCH and RBC data simultaneously
- The model is a useful tool to support bitopertin drug development, and phase 3 trials will provide further information

