# II-60 Pharmacokinetic interspecies extrapolation of a fully human mAb from animal to human

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### Introduction

Anti-LX is a fully human monoclonal antibody directed against ligand X, which is expressed on a wide variety of cell types and has a key role in down-regulating immune responses. Its anti-tumor effect relies on a dual mode of action :

- 1) Blocking the interaction of ligand X with receptor X on T cells can release T cells from immunosuppression, and lead to elimination of tumor cells;
- 2) 2) Bridging ligand X that is highly expressed on tumor cells and FCGR on leukocytes triggers tumor-directed ADCC.

Blocking ligand X has shown promising results in mouse oncology models. One of the key translational challenges is the prediction of the human pharmacokinetics and thus the selection of the (expected) optimum dose and dosing regimen.

#### Figure 2. Individual Plots: Observations, Individual Predictions and Population Predications



## Objectives

- Describe PK of anti-LX in mouse and monkey.
- Extrapolate the human exposure from the animal profile.
- Support the selection of dosing regimen in first-in-human trial based on target occupancy data from mouse.

## Methods

- Algorithm: FOCE in NONMEM 7.2
- PK and TK data from 48 mice and 21 monkeys were fitted simultaneously.
- PK parameters were scaled between animal species and then to human by allometric relationships to simulate human exposure and its variability.
- EC95 derived from mouse target occupancy data was defined as the efficacy threshold.
- Dosing regimen of achieving at least 95% target occupancy in 70% of population was recommended as a reference for dose escalation in the first-in-human trial.

## Results

- Anti-LX demonstrated pronounced non-linear PK characteristics in mouse and monkey, which was speculated to be target-mediated clearance related.
- Limited data impeded the development of a full mechanistic target-mediated drug disposition (TMDD) model [1].
- A simplification of TMDD model [2], consisting in two-compartment distribution and mixed elimination, well characterized the PK profiles of mice and monkeys (Figure 1, Figure 2).
- Inter-species variability of Km( f(target affinity)) plus IIV in linear CL, Vmax( f (target density)) and Km\_monkey.
- Vc, Vp, Q, CLlin, Vmax for human were scaled up allometrically by body weight with power factor of 1, 1, 0.75, 0.75, 0.75, respectively.

Target occupancy data from mouse were fitted using a receptor binding model (Figure 3).

$$[C] + [R] \xrightarrow{K_{on}} [RG] \xrightarrow{K_{on}} [RG]$$

$$Binding = \frac{B_{max} \cdot [C]}{[C] + K_d}$$

#### Figure 3. Target Occupancy in Mouse: Observations and Predictions



- The target effective concentration was defined as EC95 (58.8  $\mu$ g/ml) derived from the target occupancy model.
- Assuming the same target affinity (Km) in human and monkey, human exposure was simulated, incorporating variability associated with clearance.
- Time above target effective concentration in the predicted human profile was considered to be an important reference when selecting the dose range to be tested in FIM study. Simulation suggested that, in human a dose of 7 mg/kg infused in 90 min can be administered every other week, to maintain a trough plasma level above the target in 70% of the population (Figure 4).
- Sensitivity analysis: 1) As Km << Conc in both monkey and mouse, minor impact of Km to the simulated human exposure was expected; 2) Up to 2 fold alteration of the assumed Vm value had small impact to the simulated trough concentration (Figure 5).



#### Table 1. Population PK Parameter Estimation of Mouse and Monkey

First Order Conditional Estimation with Interaction						
Objective function value:	-468.558					
Theta	Estimate	SE	RSE	[ lower,	init,	upper
1 CLL - linear clearance [L/h]	1.08	0.0879	8.1%	0	1	+I
2 V1 - central volume of distribution [L]	145	5.44	3.8%	0	150	+I
3 Q - intercompartmenal clearance [L/h]	4.03	0.467	11.6%	0	4	+I
4 V2 - peripheral volume of distribution [L]	175	11.9	6.8%	0	100	+I
5 VMAX – maximal TMDD clearance [ug/h]	12.2	3.6	29.5%	0	17	+I
6 KM_ Monkey - Concentration yielding half maximal TMDD clearance [ug/ml]	0.35	0.195	55.7%	0	1	+I
7 KM_ Mouse - Concentration yielding half maximal TMDD clearance [ug/ml]	3.75	1.47	39.2%	0	10	+I
8 Additive residual error	0 FIX	0	%	0	0	
9 Proportional residual error	0.276	0.0255	9.2%	0	0.2	+I
Omega 1 2 6 Etabar (SE) p val Shrinkage						
1.IIV/CLL 0.311 0.008 (0.059) 0.8941 10.9%						
2.IIV/VMAX 0 1.02 0.033 (0.06) 0.5823 50.3%						
6.IIV/KM_Monkey 0 0 1.19 0.002 (0.039) 0.9565 45.4%						
<b>Omega (on SD scale)</b> 1 2 6						
1.IIV/CLL 55.8%						
2.IIV/VMAX 0% 101%						
6.IIV/KM_Monkey 0% 0% 109.1%						
Sigma Shrinkage						
1 SIs as TH 8.5%						

#### Figure 1. Basic Goodness-of-Fit Plots of the Pooled PK Model

#### Figure 4. Simulation of Human Population Exposure with 7mg/kg Biweekly Regimen





## Conclusion / Discussion

- Two-compartment model with mixed elimination, a simplification of TMDD model, well described the PK profiles of anti-LX both in mouse and monkey.
- Linear clearance is dominant at high concentration when TM pathway is saturated; target-mediated clearance is more visible at low concentration resulting in faster elimination.
- In addition to the time independent linear and nonlinear components of the clearance, a time dependency associated with the binding to the anti-drug antibody developed during the late stage of treatment is taken into account for the human extrapolation (working in progress by incorporating SCID mice data).
- Simulations suggested that for maintaining at least 95% target occupancy in human in 70% of the population, a dose of 7 mg/kg can be administered every other week as a 90 min infusion. This supports the selection of the dose range and frequency to be tested in the FIM study, to maintain a trough plasma level for an expected clinical relevant PD response. It is however compounded by uncertainties in the assumptions and importantly so, in allometric scaling, and caution will be exercised with respect to the selection of the starting dose and escalation.



#### References

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