Evaluation and optimization of potential for subasumstat (TAK-981) to overcome rituximab resistance via PK/PD and QSP modeling of antibody dependent cell mediated cytotoxicity

Dean Bottino

Takeda Development Center Americas, Inc. (TDCA), Lexington, MA, USA

#### **Acknowledgements**







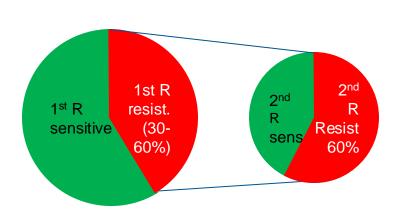


Name	Affiliation
Maria Veronica Ciocanel*	Duke University
Kaitlyn E Johnson*	The University of Texas at Austin
Josua Aponte-Serrano*	Indiana University – Bloomington
Nicolas Bajeux*	University of Manitoba
Fanwang Meng*	McMaster University
Akito Nakamura Allison Berger John Gibbs Dean Bottino*	Takeda Development Center Americas, Inc. (TDCA), Lexington, MA, USA

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<sup>\*</sup>Participants in the Fields Institute Problem Solving Workshop, 2019

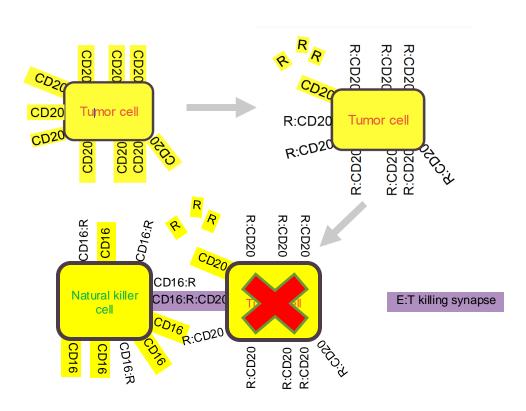
### Rituximab (R) resistance remains an unmet need in Non-Hodgkin's Lymphoma (NHL)



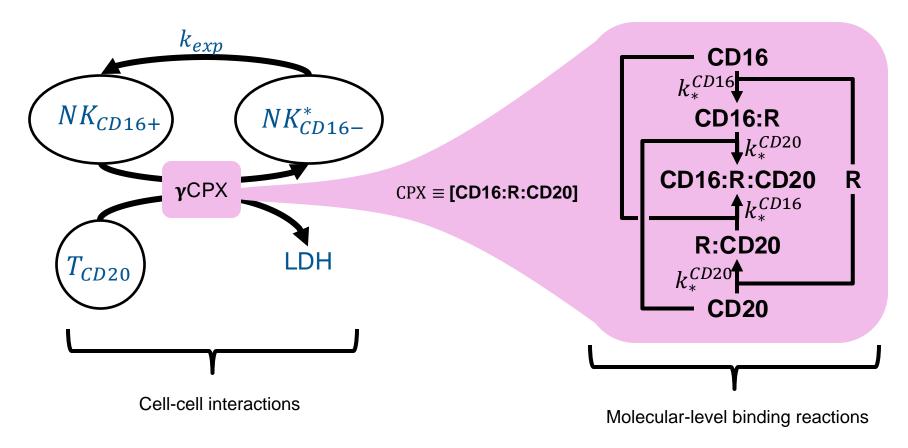
R-CHOP 1<sup>st</sup> line therapy If cure not achieved, options are limited

Binding of R allows interaction with effector cell via CD16, which leads to Antibody Dependent Cell-mediated Cytotoxicity (ADCC)

R resistance mechanisms include CD20 loss, CD16 loss



#### A QSP model of ADCC may be used to simulate mechanisms of RTX resistance and therapeutic mechanisms to overcome resistance

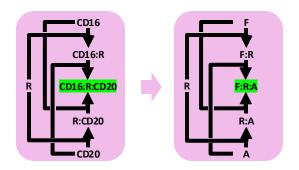


LDH=Lactate DeHydrogenase; CPX = immune synapse ComPleX

#### **Bispecific ODE's**

#### \*New notation:

- CD16 → F ~ Fcg receptor
- Rituximab
- CD20 → A ~ <u>A</u>ntigen



$$\frac{d}{dt}R = -\frac{1}{h} \cdot k_{FR}^{on} \cdot F \cdot R - \frac{1}{h} \cdot k_{RA}^{on} \cdot A \cdot R + \frac{1}{h} \cdot k_{FR}^{off} \cdot FR + \frac{1}{h} \cdot k_{RA}^{off} \cdot RA$$

$$\frac{d}{dt}A = -k_{RA}^{on} \cdot A \cdot R + k_{RA}^{off}RA - \frac{1}{h} \cdot k_{RA}^{on} \cdot A \cdot FR + k_{RA}^{off} \cdot FRA$$

$$\frac{d}{dt}F = -\frac{k_{FR}^{on}}{k_{FR}^{on}} \cdot F \cdot R + k_{FR}^{off} \cdot FR - \frac{1}{h} \cdot \frac{k_{FR}^{on}}{k_{FR}^{on}} \cdot F \cdot RA + k_{FR}^{off} \cdot FRA$$

$$\frac{d}{dt}RA = k_{RA}^{on} \cdot A \cdot R \cdot k_{RA}^{off} \cdot RA - \frac{1}{h} \cdot k_{FR}^{on} \cdot F \cdot RA + k_{FR}^{off} \cdot FRA$$

$$\frac{d}{dt}FR = k_{FR}^{on} \cdot F \cdot R - \frac{1}{h} \cdot k_{RA}^{on} \cdot A \cdot FR - k_{FR}^{off} \cdot FR + k_{FR}^{off} \cdot FRA$$

$$\frac{d}{dt}FRA = \frac{1}{h} \cdot \frac{k_{FR}^{on}}{k_{FR}^{on}} \cdot F \cdot RA + \frac{1}{h} \cdot \frac{k_{RA}^{on}}{k_{RA}^{on}} \cdot A \cdot FR - k_{FR}^{off} \cdot FRA - k_{RA}^{off} \cdot FRA$$

Assuming R is not significantly depleted by the reactions (ie, omitting dR/dt equation) gives a closed-form steady state solution

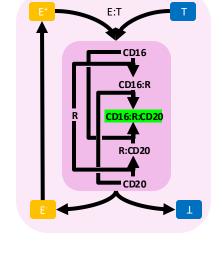
#### **NK ADCC model**

NK cells: 
$$\frac{dN}{dt} = k_{ex}^N (N_0 - N) - r_N (\gamma_N f_N + \delta_N) N^{r_N} T$$

Tumor: 
$$\frac{dT}{dt} = gT - (\gamma_N f_N + \delta_N) N^{r_N} T$$

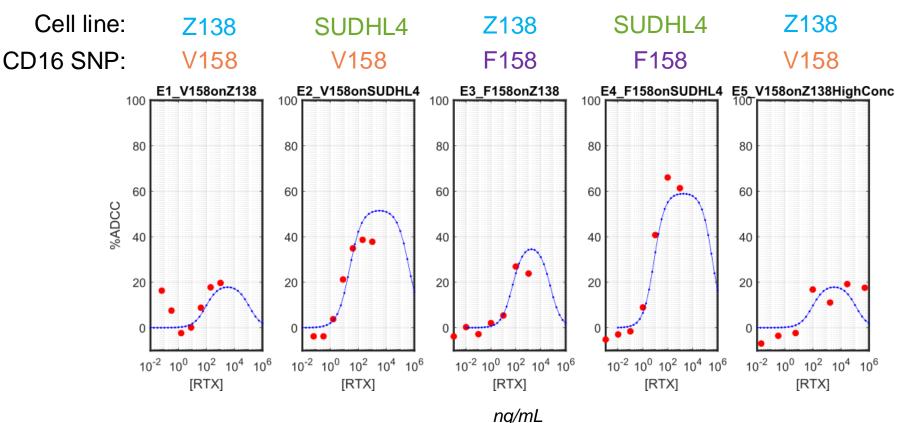
 $f_N = f(RTX, CD16_N, CD20_T)$ , the equilibrium solution to bispecific trimer ODE's

 $\gamma_N = \text{ADCC}$  intensity factor  $= \gamma_N(CD69) = \gamma_N^0$  for unstimulated NKs  $\delta_N = \text{AICC}$  intensity factor  $= \delta_N(CD69) = \gamma_N^0$  for unstimulated NKs



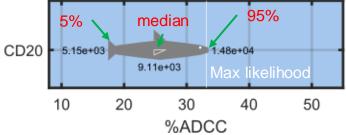
Core ADCC model

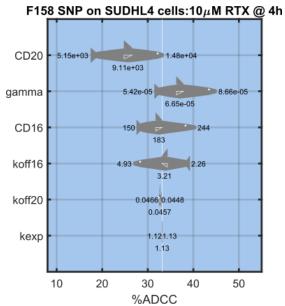
### Model parameters can be calibrated to describe ex vivo data in multiple cell lines and donor CD16 SNPs

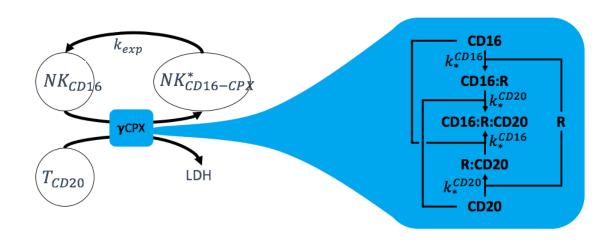


Tumor CD20 is the most sensitive (uncertainty-weighted) driver of

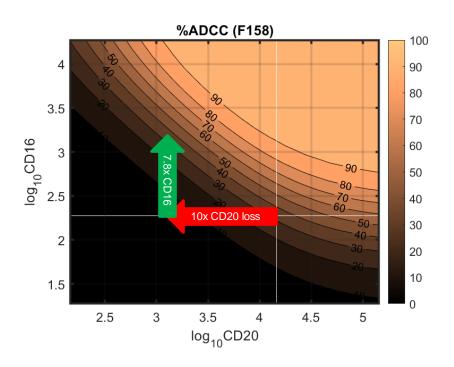
**ADCC** 

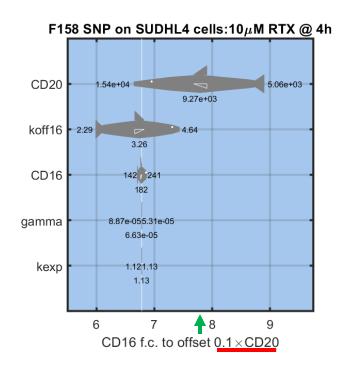




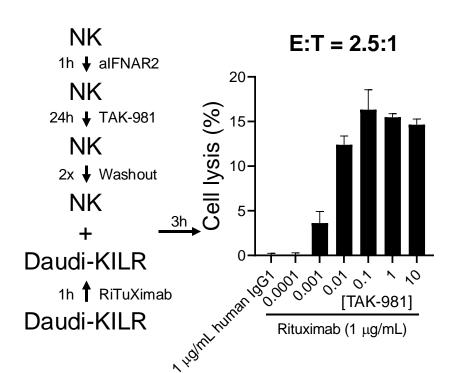


### Model predicts ~8-fold increase in CD16 on NK cells can offset 10-fold CD20 loss





# Questions/motivation for QSP modeling TAK-981 effects on Rituximab combo

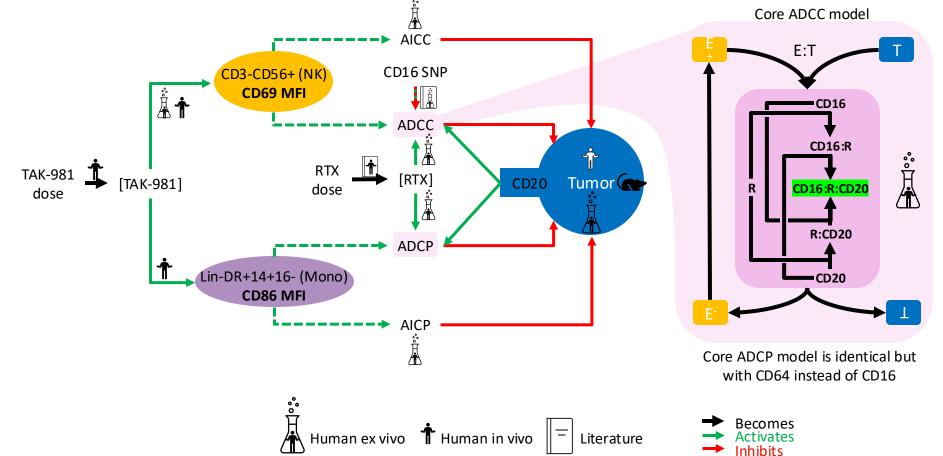


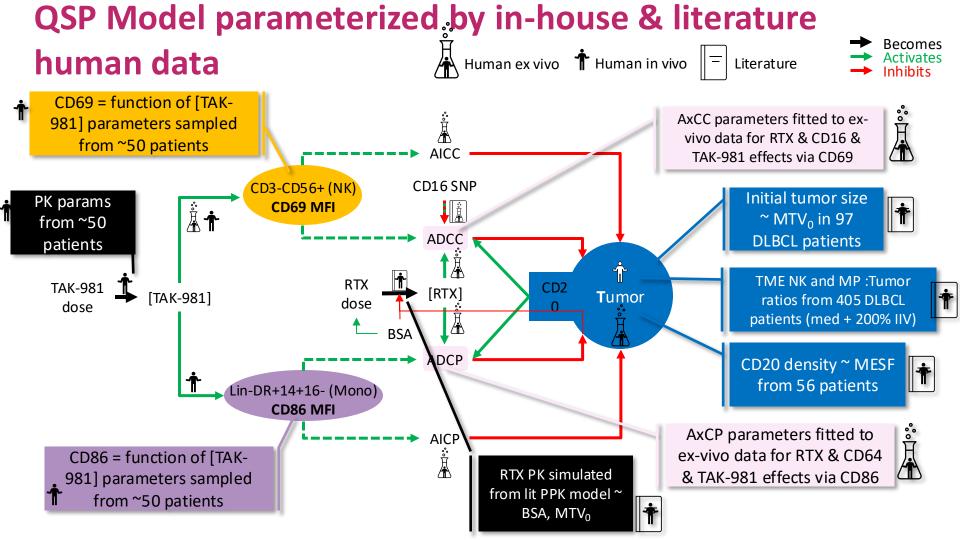
Small molecule SUMOylation inhibitor TAK-981 (subasumstat) enhances NK-mediated ADCC/AICC and Mac-mediated ADCP/AICP in presence / absence of rituximab (RTX) in human ex-vivo experiments.

- In vitro killing → Can TAK-981+RTX deliver clinical responses in RTX R/R?
- 2. Optimal dose and schedule for TAK-981 + RTX?
- 3. Which RTX R/R patients benefit from +TAK-981?

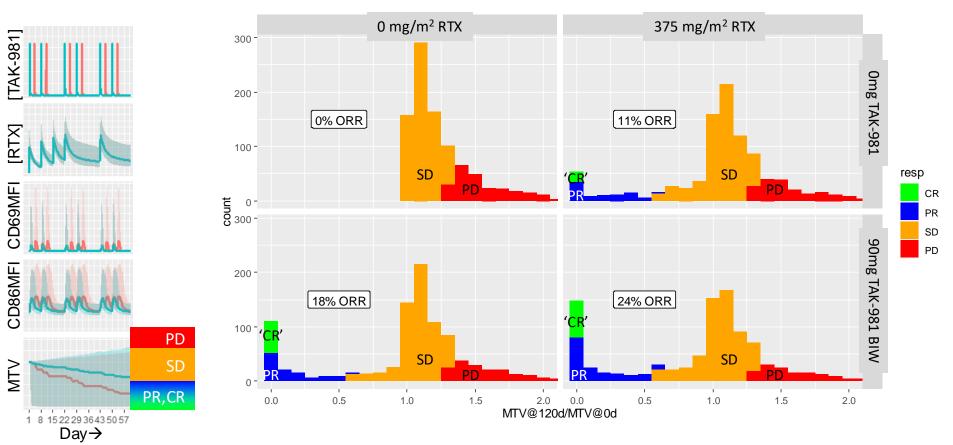
Nakamura et al, Blood 2022

# QSP model combines the ADCC 'core' model with clinical PK/PD models



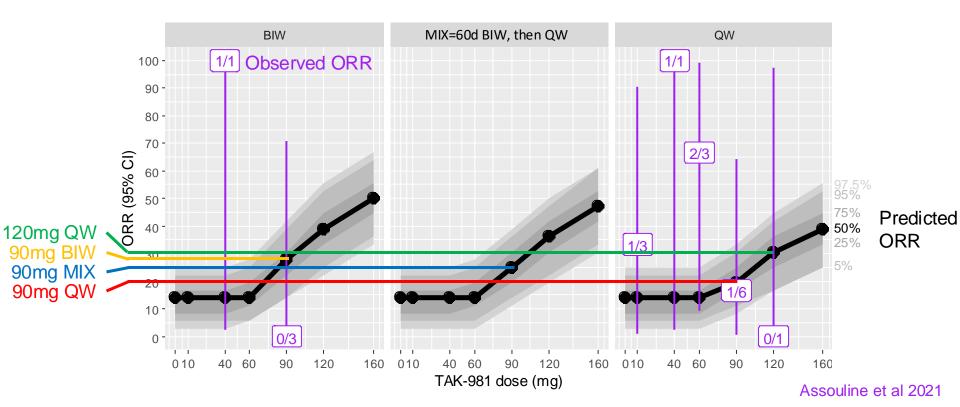


### Without additional parameter calibration, QSP model predicts 0% spontaneous Overall Response Rate (ORR), 11% ORR with Rituximab single agent

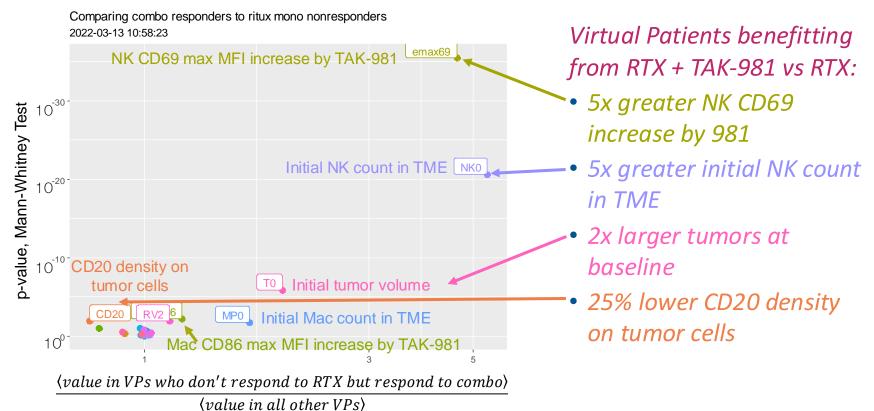


#### Model predicts ORR for 120mg QW > 90mg BIW~90mgMIX > 90mg QW

**Model-predicted ORR** by schedule and dose (50/90/95%PI in a 36-patient cohort) **Observed ORR** by schedule and dose (with 90% Clopper Pearson Cls)



### How do virtual patients (VPs) who benefit from TAK-981 added to rituximab differ from the rest? Synthetic Volcano Plot (preliminary results)



## Questions/predictions for QSP modeling TAK-981 effects on RTX combo

- 1. Based on in vitro activity, could we expect TAK-981/RTX combo to deliver clinical responses in RTX R/R NHL patients?
  - The model predicts TAK-981 at 90mg BIW increases ORR from ~11% to 24%
- 2. What is optimal dose and schedule for TAK-981 in combo with RTX?
  - 120mg MIX (BIW for 60d, then QW)
- 3. Which RTX R/R patients are most likely to benefit from TAK-981 in this context?
  Patients with:
  - High maximum levels of CD69 MFI increase due to TAK-981
  - Higher levels of NK cells in tumor microenvironment at baseline
  - Larger tumors at baseline
  - Lower CD20 on tumor cells

#### **Conclusions**

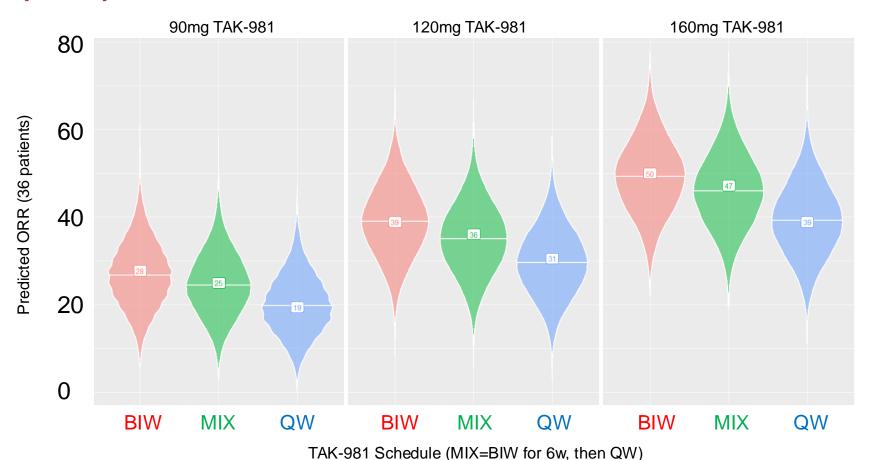
- The 2019 Fields Institute Industry Problem Solving Workshop team
  - Derived steady-state solution for bispecific binding of rituximab to CD20 and CD16
  - Derived an NK-Tumor interaction model & fit to published ADCC data
  - Predicted degree of increase in ADCC factors required to offset loss of rituximab sensitivity
- The academic/industry 'open source' model was then incorporated into a proprietary TAK-981 model to address key program questions:
  - Feasibility
  - Optimal dose and schedule
  - Factors predicting benefit

#### Thank you!

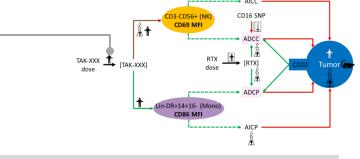
### The model can be used to quantify the magnitudes of therapeutic mechanisms required to offset a given mechanism of RTX resistance

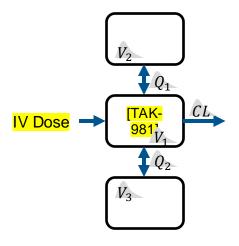
Mechanism of RTX resistance (MoRR)	ADCC fold change due to MoR (90% CI)
10x loss of CD20 on tumor	0.16 (0.13,0.20)
10x loss of RTX exposure	0.17 (0.14,0.21)
10x decrease in CD16 affinity	0.16 (0.13,0.20)

### We can use the QSP model to stimulate overall response rates (ORRs) at various doses and schedules



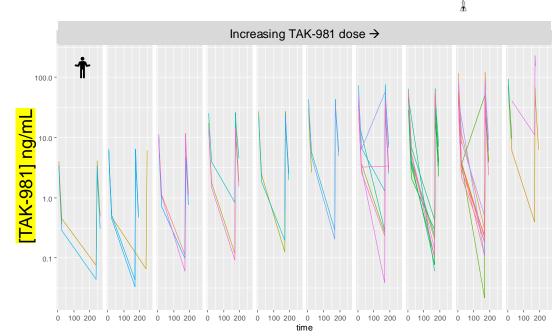
# TAK-981 popPK model based on FIH PK data



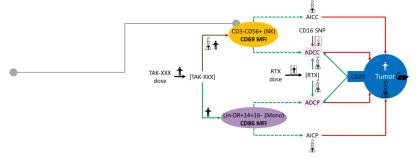


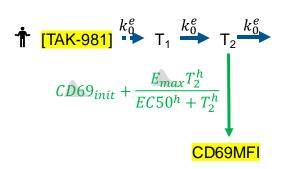
Inter-individual variability

Observation



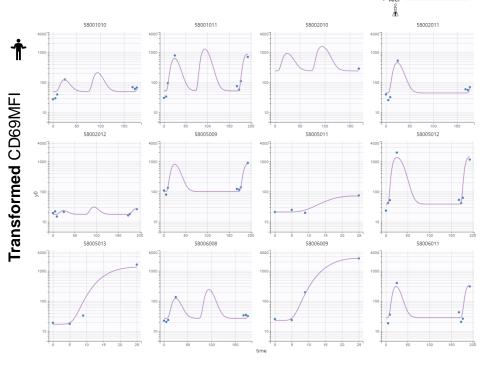
# TAK-981 NK activation (CD69 MFI) module based on patient-level PK/PD data



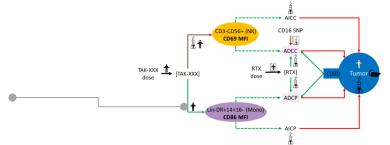


Inter-individual variability

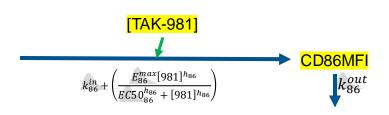
Observation



TAK-981 Mac/monocyte activation (CD86 MFI) based on FIH patient-level PK/PD

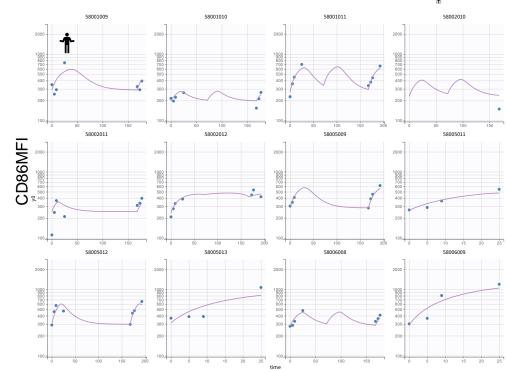


data

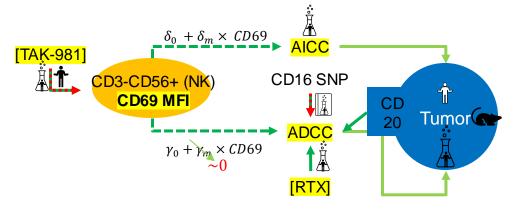


Inter-individual variability

Observation

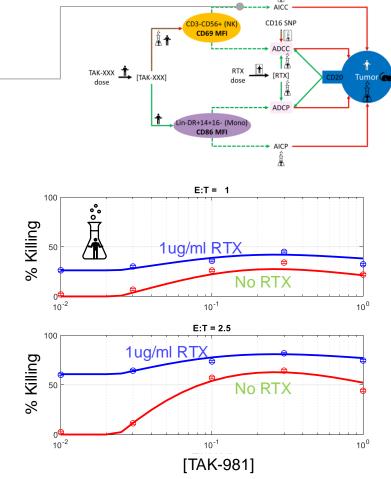


Model of CD69MFI effects on ADCC/AICC calibrated to ex-vivo CD69 & cytotox data

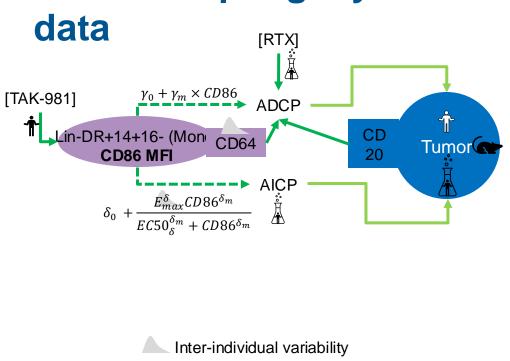


Inter-individual variability

**Observation** 



Model of CD86MFI effects on ADCP/AICP calibrated to ex-vivo phagocytosis



**Observation** 

