

# A PK-PD Modeling and Simulation Based Strategy for Clinical Translation of Antibody Drug Conjugates: A Case Study with T-DM1

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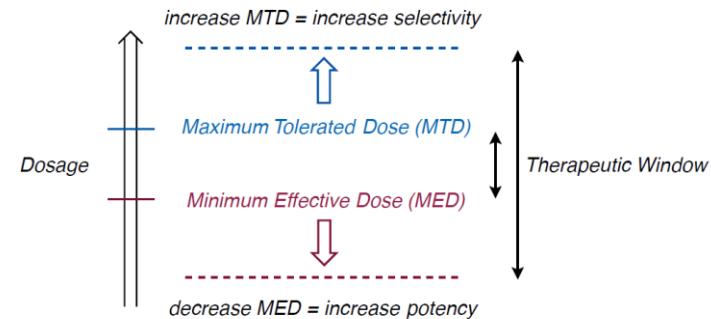
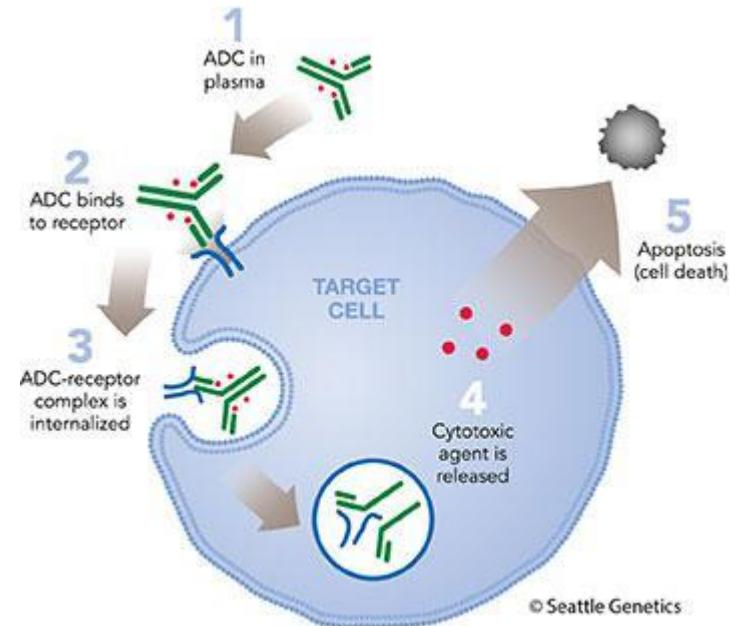
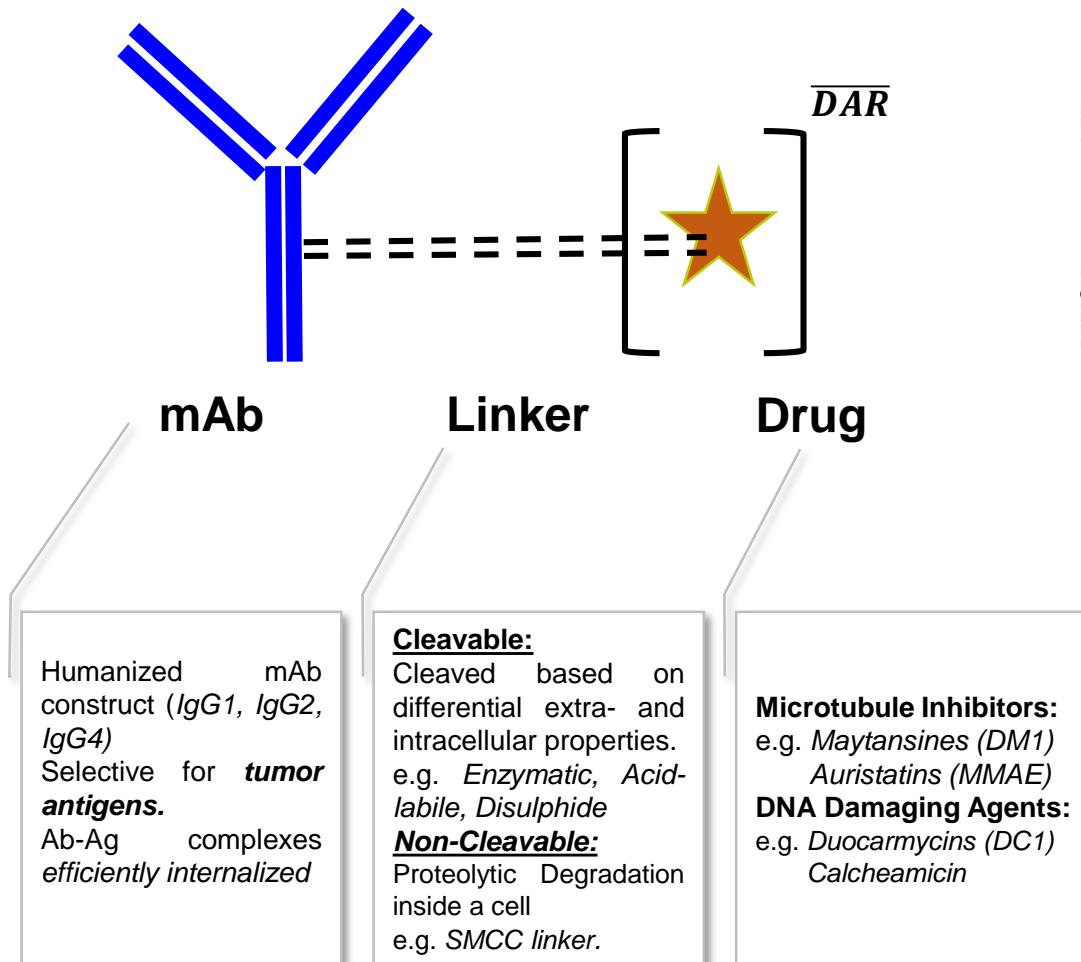
**ROSA Impact Webinar**  
**04/19/2017**



**University at Buffalo** The State University of New York  
School of Pharmacy and Pharmaceutical Sciences

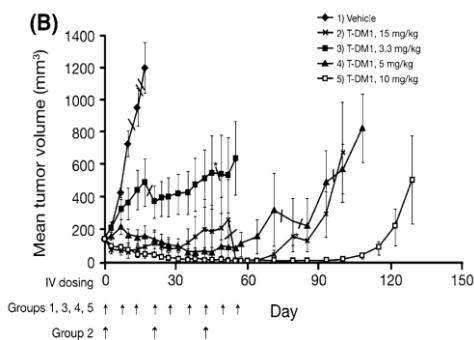
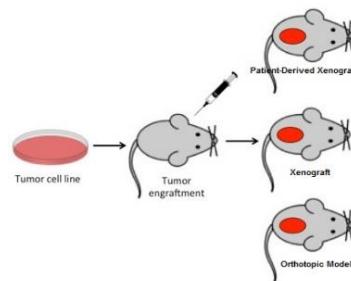
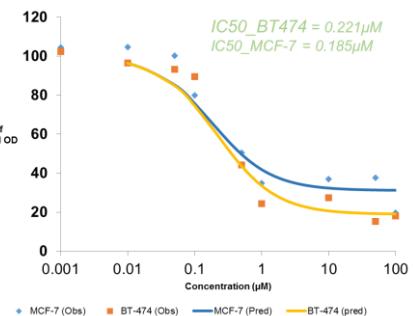
# Background: ADCs

~60 Antibody Drug Conjugates are in clinical trial<sup>1</sup>



Angew. Chem. Int. Ed. 2014, 53, 3796.

# Motivation for Development of a Mechanistic Model



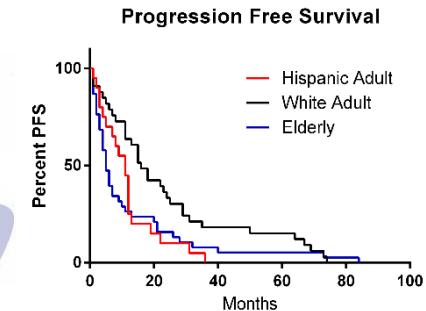
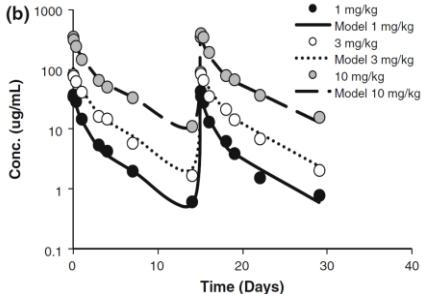
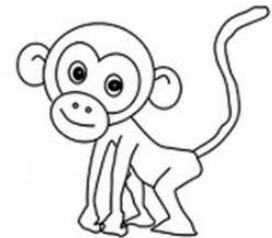
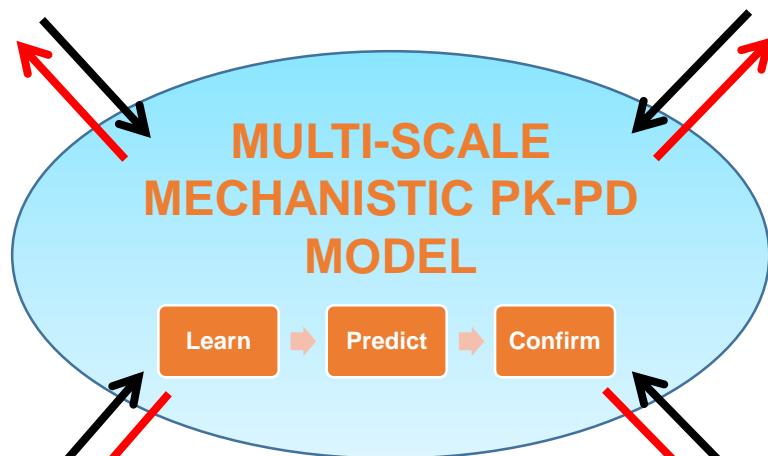
DISCOVERY PHASE

INFORM

LATE PRECLINICAL PHASE

EARLY PRECLINICAL PHASE

CLINICAL PHASE



# Motivation for Development of a Mechanistic Model

Bench to bedside translation of antibody drug conjugates using a multiscale mechanistic PK/PD model: a case study with brentuximab-vedotin

Dhaval K. Shah · Nahor Haddish-Berhane · Alison Betts

Valine-citrulline Linker  
Microtubule Inhibitor

1  
50

DISCOVERY

Preclinical to Clinical Translation of Antibody-Drug Conjugates Using PK/PD Modeling: a Retrospective Analysis of Inotuzumab Ozogamicin

FARI Y PRFCI INICAI

Alison M. Betts,<sup>1,9,10</sup> Nahor Haddish-Berhane,<sup>2</sup> John Tolsma,<sup>3</sup> Paul Jasper,<sup>3</sup> Lindsay E. King,<sup>1</sup> Yongliang Sun,<sup>4</sup> Subramanyam Chakrapani,<sup>5</sup> Boris Shor,<sup>6</sup> Joseph Boni,<sup>7</sup> and Theodore R. Johnson<sup>8</sup>

Acid-labile Linker  
DNA Damaging Agent

PHASE

Application of a PK-PD Modeling and Simulation-Based Strategy for Clinical Translation of Antibody-Drug Conjugates: a Case Study with Trastuzumab Emtansine (T-DM1)

CLINICAL PHASE

Non-cleavable Linker  
Microtubule Inhibitor

Aman P. Singh<sup>1</sup> and Dhaval K. Shah<sup>1,2</sup>

Time (Days)

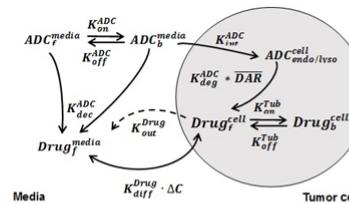
Months

# General PK-PD Based Strategy for Clinical Translation of ADCs

**Step 1**

## *In Vitro* Cellular Disposition Model

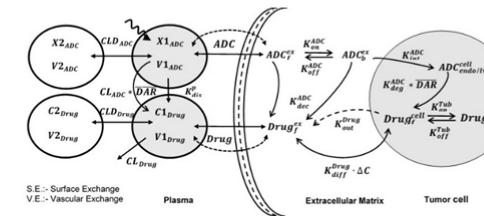
Characterize the internalization, degradation and payload release for ADCs in a tumor cell.



**Step 2**

## *In Vivo* Tumor Distribution Model

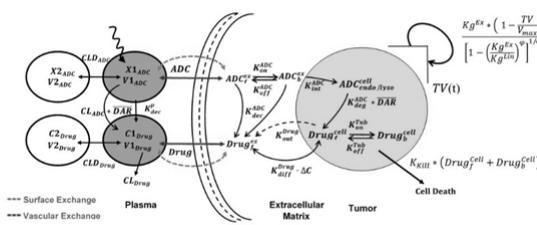
*A Priori* predict plasma and tumor exposures of different analytes of ADC.



**Step 3**

## *In Vivo* Tumor Growth Inhibition (PKPD) Model

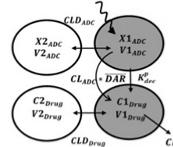
Use tumor concentrations to characterize TGI data and obtain PD parameters



**Step 4**

## *In Vivo* Plasma PK Model in Monkey

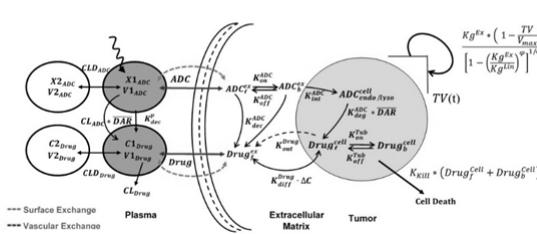
Characterize systemic concentrations of different analytes of ADCs in monkeys



**Step 5**

## Predict Clinical PK from monkey Predict Clinical PD from mouse

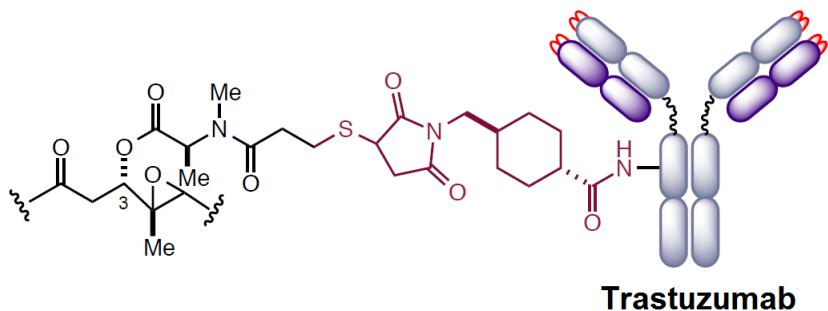
Scale up monkey PK parameters to predict human PK. Use mouse PD parameters to predict Progression Free Survival



# Background- Trastuzumab Emtansine (Kadcyla®)



Average DAR of 3.5 molecules of **DM1** molecules per antibody.



Trastuzumab-smcc-DM1  
(non-cleavable linker)  
(DAR4)

Proteolytic Degradation  
+ Linker Cleavage → Lysine-mcc-DM1  
(charged)

Less  
Permeable

Minimal  
Bystander  
Killing

Ogitani et al,  
Cancer Sci (2016)

## **Trastuzumab** [Herceptin®]

- Humanized **anti-HER2 mAb**
- Indicated for HER2-positive metastatic breast Cancer

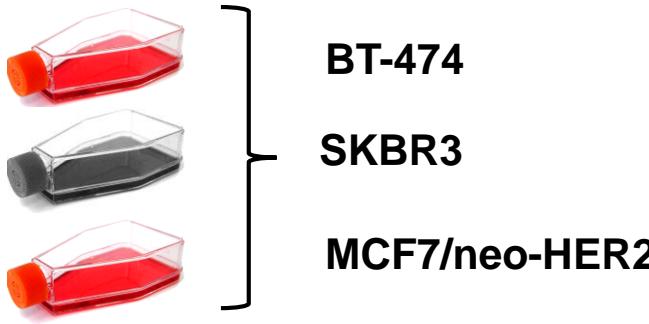
## **Emtansine (DM1)**

- Synthetic Derivative of Maytansine
- **Microtubule Inhibitor**
- Highly potent: IC<sub>50</sub> values of 10-100 pM

## **SMCC Thioether Linker**

- Chemically non-labile (**uncleavable**) linker
- Proteolytic degradation of mAbs leads to formation of metabolites of Drug-linker-amino acid residue

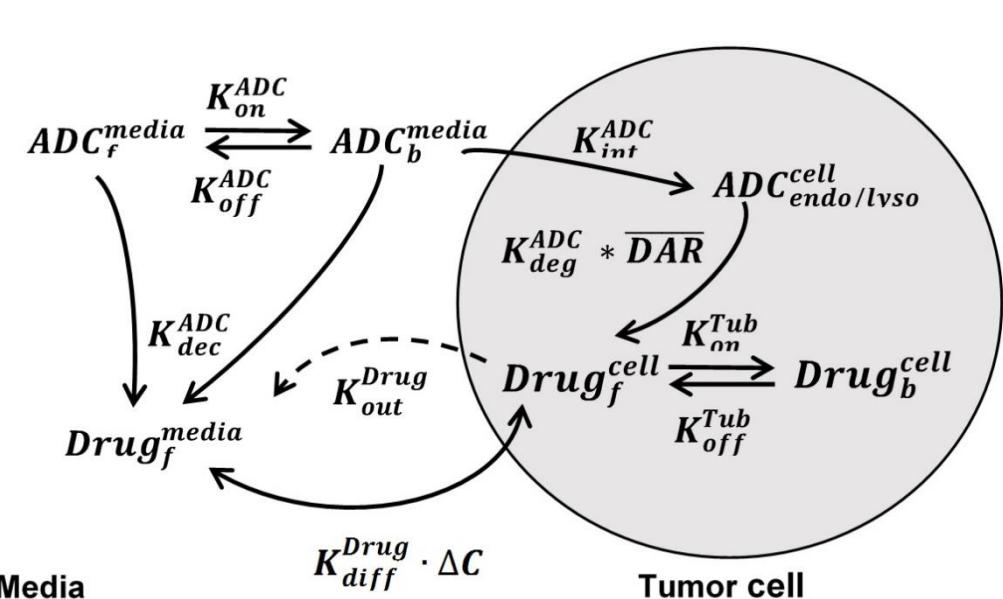
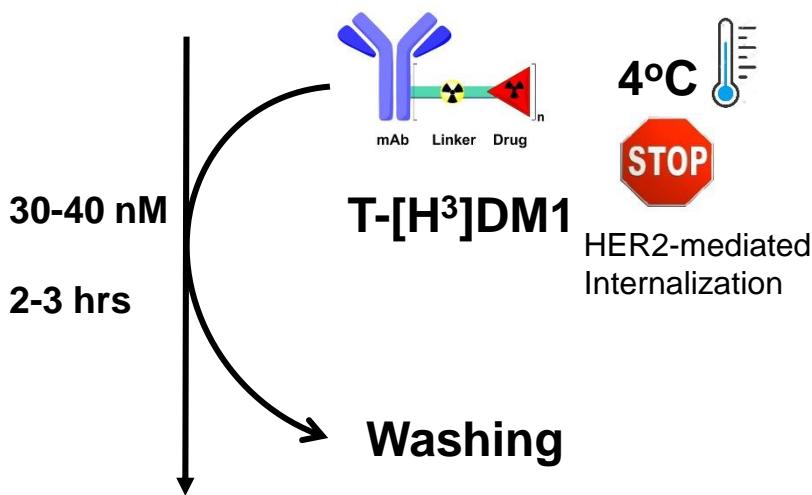
# Cellular Disposition Model for T-DM1



## *In Vitro* Cellular Disposition Model

Characterize the internalization, degradation and payload release for ADCs in a tumor cell.

### Step 1



- Total DM1 Catabolites
- Unconjugated DM1
- Conjugated DM1

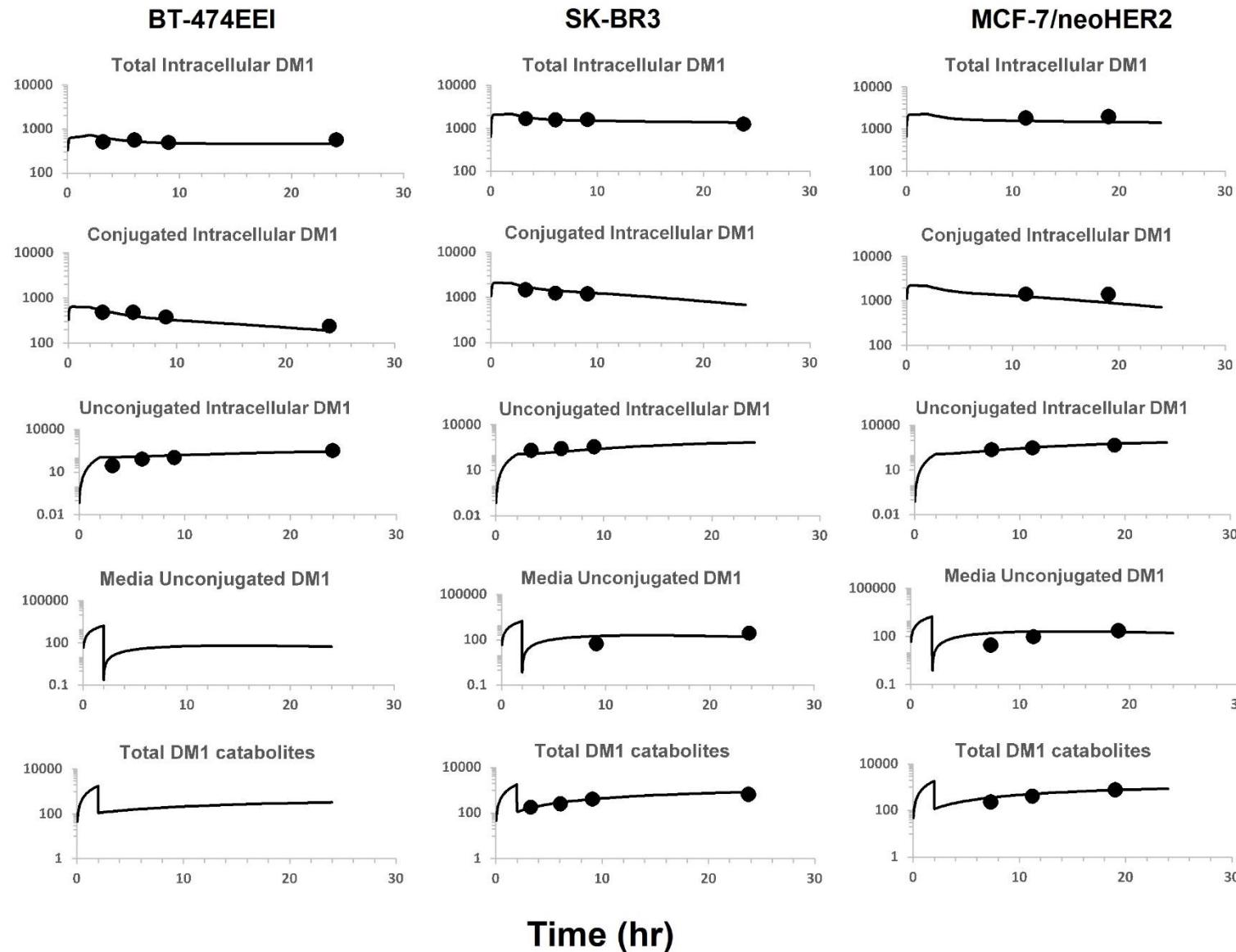
Sampling time= up to 24 hrs

Erickson HK et al, *Mol Cancer Ther.* (2012)

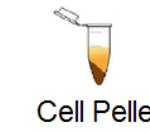
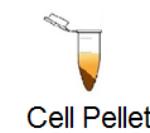
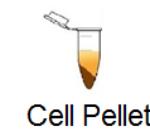
# Cellular Disposition Model Fits-T-DM1



Concentration (nM)



Time (hr)



# Cellular Disposition Model Parameters

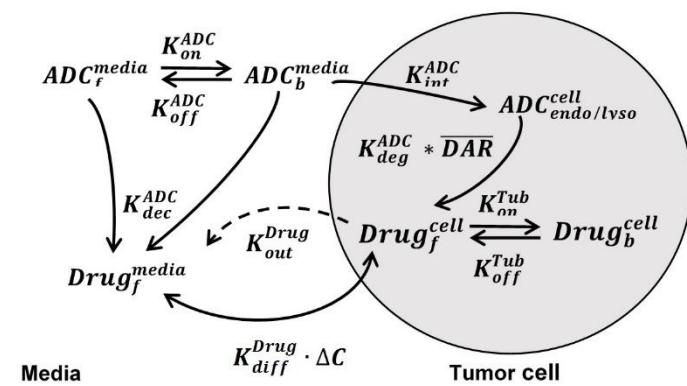
Intracellular Model Parameters	Parameter Value	Units	Reference
$K_{on}^{ADC}$	0.37	1/nM/hr	Maass et al
$K_{off}^{ADC}$	0.014	1/hr	Maass et al
$K_{int}^{ADC}$	0.011	1/hr	Maass et al
$K_{deg}^{ADC}$	0.03	1/hr	Maass et al
$K_{on}^{Tub}$	0.03	1/nM/hr	Shah 2014
$K_{off}^{Tub}$	10.6	1/hr	Bhattacharyya (1977)
$Tub_{total}$	65	nM	Shah 2014
$K_{dec}^{ADC}$	0.0226	1/hr	Bender (2014)
$K_{diff}^{Drug}$	0.092 (17.4%)	1/hr	Estimated
$K_{out}^{Drug}$	0	1/hr	Fixed
$Ag_{total}^{BT-474EEI}$	0.594 (12.4%)	nM	Estimated
$Ag_{total}^{SK-BR3}$	1.6 (11.3%)	nM	Estimated
$Ag_{total}^{MCF-7/neHER2}$	1.96 (11.8%)	nM	Estimated

## In Vitro Cellular Disposition Model

Characterize the internalization, degradation and payload release for ADCs in a tumor cell.



Step 1



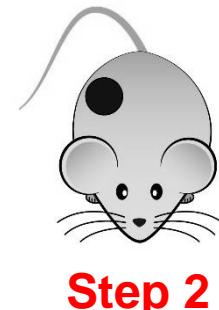
# In Vivo Plasma Pharmacokinetics Model

## Plasma PK Datasets

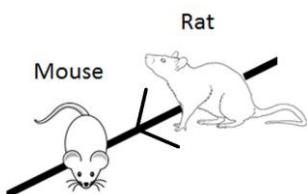
- 1) Erickson et al (2007) in normal mice at two dose levels (2 & 3 mpk) – TTmab and T-DM1
- 2) Jumbe et al (2010) in xenograft mice at 3 dose levels (0.3, 3 and 15 mpk) – T-DM1
- 3) Shen et al (2012) – single dose IV DM1 administration in rats.

**In Vivo** Tumor Distribution Model

*A Priori* predict plasma and tumor exposures of different analytes of ADC.

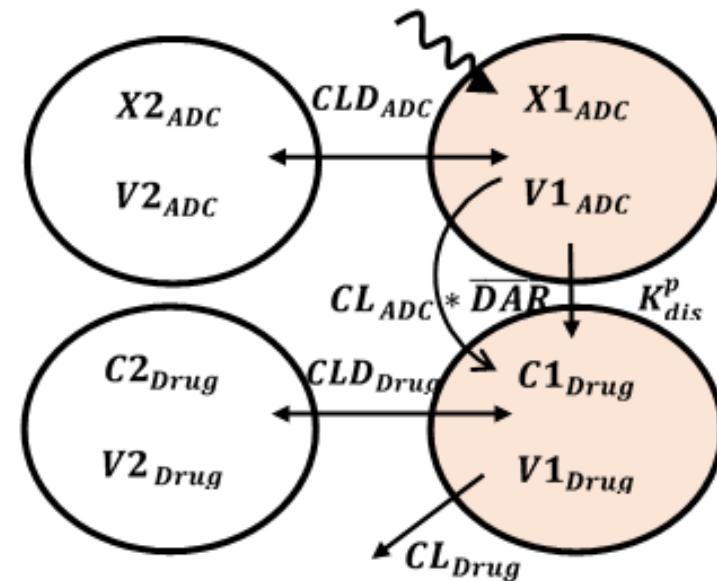


## Allometric Scale Down

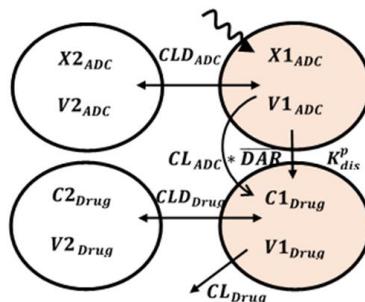


$$CL_{mice} = CL_{rats} \cdot \left[ \frac{BW_{mice}}{BW_{rats}} \right]^{0.75}$$

$$V_{mice} = V_{rats} \cdot \left[ \frac{BW_{mice}}{BW_{rats}} \right]^1$$

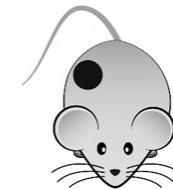


# In Vivo Plasma Pharmacokinetics Model

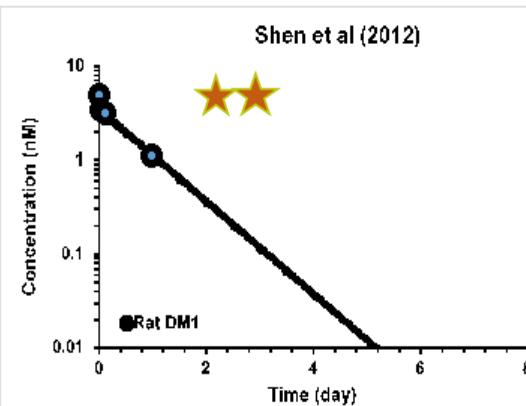
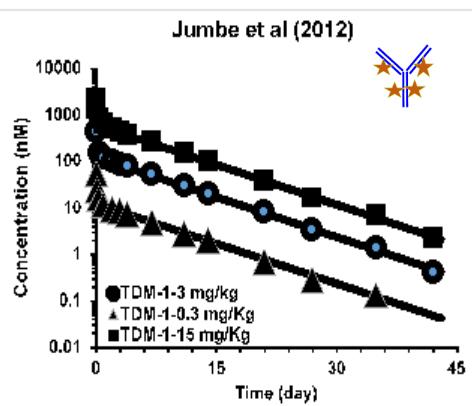
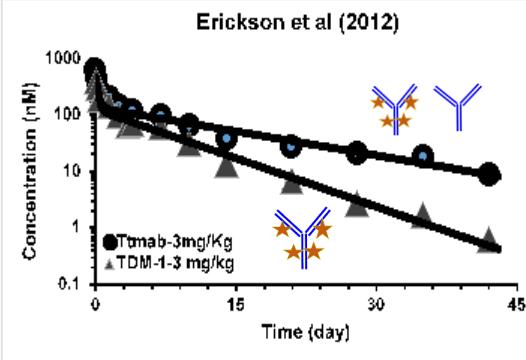
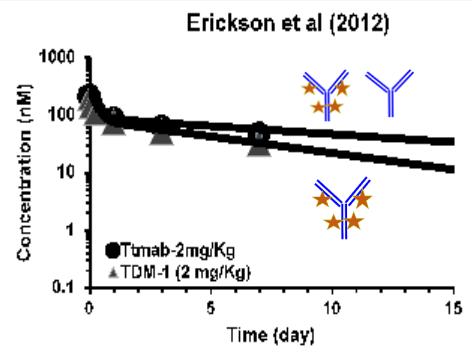


*In Vivo* Tumor Distribution Model

*A Priori* predict plasma and tumor exposures of different analytes of ADC.

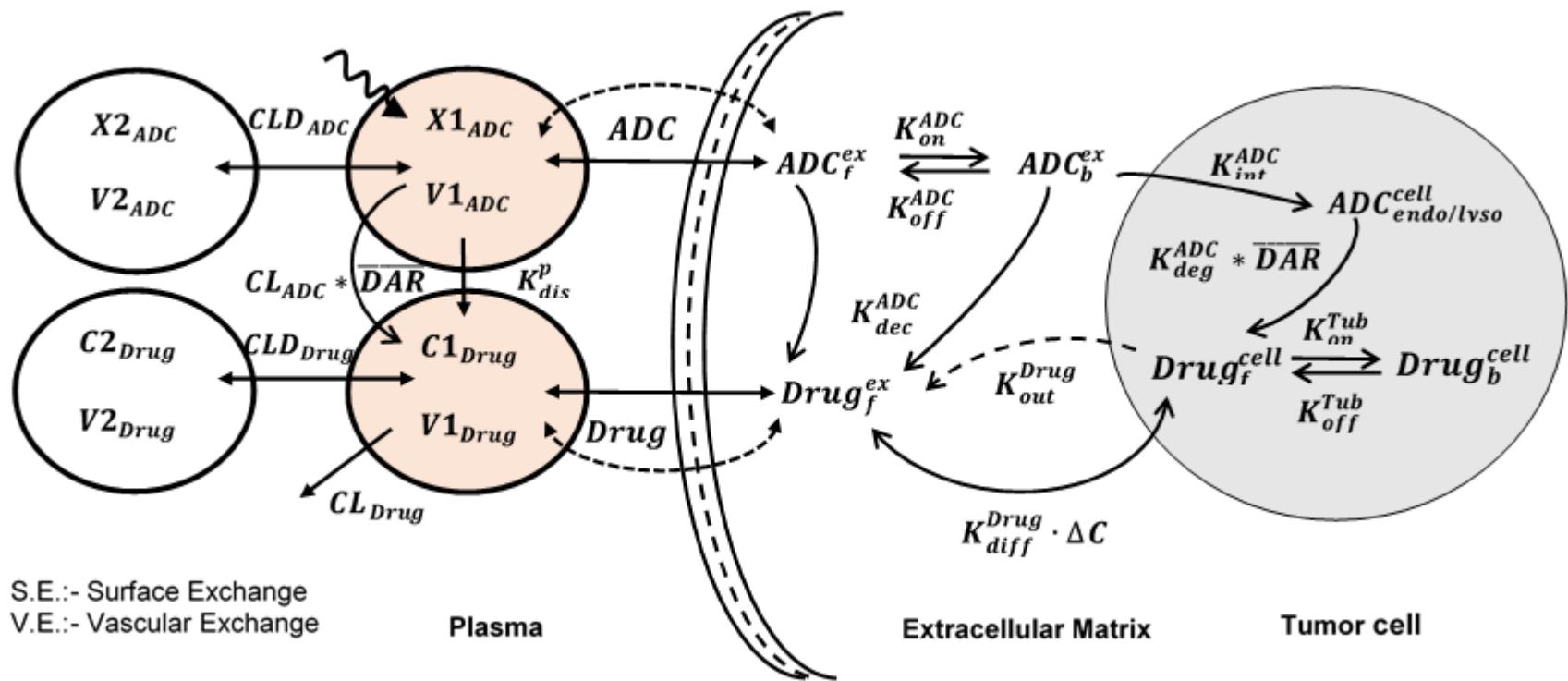


Step 2



Systemic PK Parameters	Parameters	Units	Reference
$CL_{ADC}$	0.0093 (4.4 %)	L/day	Estimated
$CLD_{ADC}$	0.118 (12.6 %)	L/day	Estimated
$V1_{ADC}$	0.043 (7.3 %)	L	Estimated
$V2_{ADC}$	0.0948 (5.2 %)	L	Estimated
$CL_{Drug}$	11.29 (78.2%)	L/day	Estimated
$CLD_{Drug}$	155.4	L/day	Fixed
$V1_{Drug}$	3.30 (48 %)	L	Estimated
$V2_{Drug}$	2.01	L	Fixed
$K^P_{dec}$	0.241 (8.8%)	1/day	Estimated

# In Vivo Tumor Disposition Model for T-DM1



S.E.: Surface Exchange  
V.E.: Vascular Exchange

Plasma

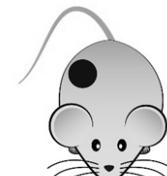
Extracellular Matrix

Tumor cell

Parameters:-	Value	Unit	Reference
$R_{Cap}$	8	$\mu\text{m}$	Shah 2014
$R_{Krogh}$	75	$\mu\text{m}$	Shah 2014
$P_{ADC}$	334	$\mu\text{m}/\text{day}$	Shah 2014
$P_{Drug}$	21000	$\mu\text{m}/\text{day}$	Shah 2014
$D_{ADC}$	0.022	$\text{cm}^2/\text{day}$	Shah 2014
$D_{Drug}$	0.25	$\text{cm}^2/\text{day}$	Shah 2014
$\epsilon_{Drug}$	0.44	Unitless	Shah 2014
$\epsilon_{ADC}$	0.24	Unitless	Shah 2014

**In Vivo** Tumor Distribution Model

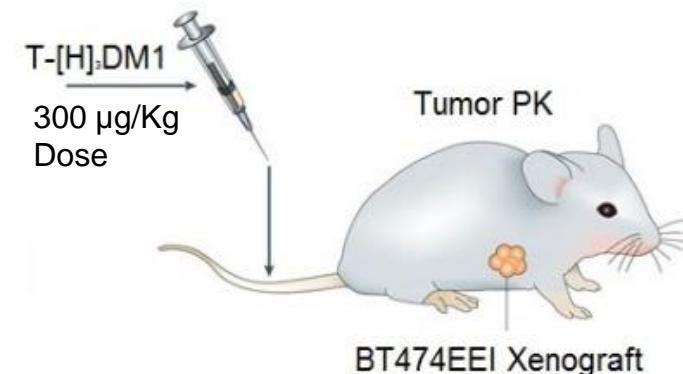
*A Priori* predict plasma and tumor exposures of different analytes of ADC.



Step 2

Thurber GM et al, *Adv. Drug. Deliv Rev.* (2008)  
Thurber GM et al, *Trends Pharmacol Sci.* (2008)

# A Priori Predictions using our Tumor Disposition Model



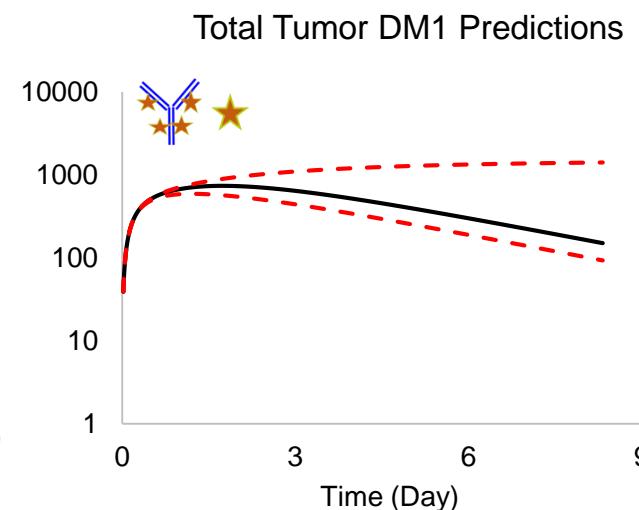
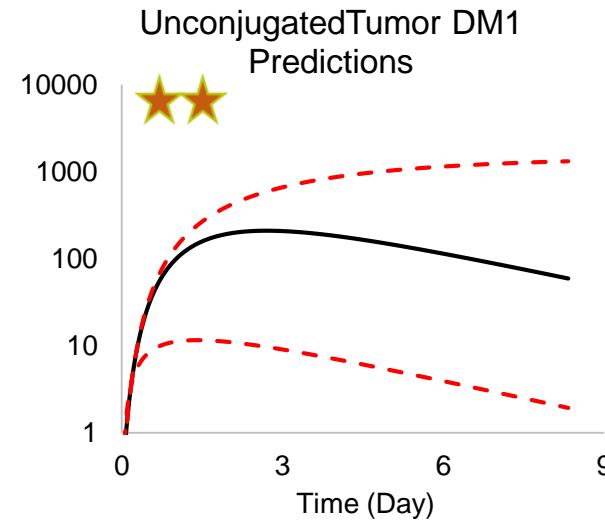
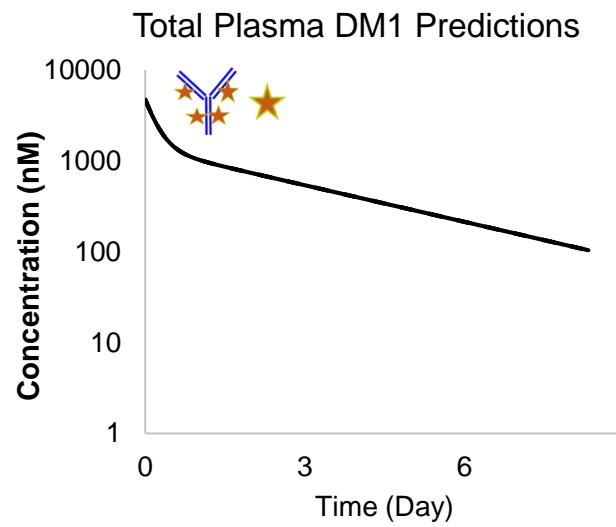
## Plasma and Tumor Homogenates

- Total DM1 Catabolites
- Unconjugated DM1

Sampling time= up to 7 Days

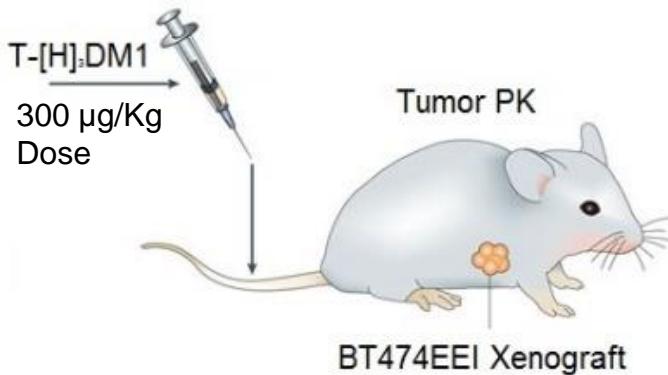
### In Vivo Tumor Distribution Model

*A Priori* predict plasma and tumor exposures of different analytes of ADC.



Monte-Carlo simulations were performed with *IIV* of 17.4 % on  $K_{diff}$  parameter.

# Validation of our Predictions !



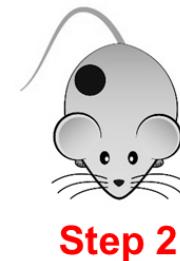
## Plasma and Tumor Homogenates

- Total DM1 Catabolites
- Unconjugated DM1

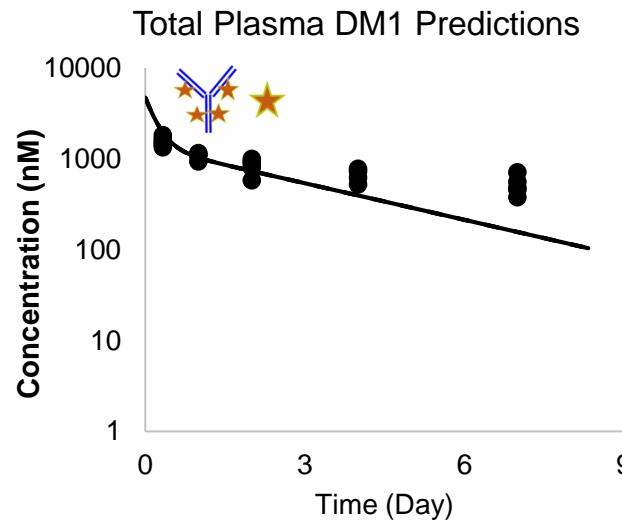
Sampling time= up to 7 Days

### In Vivo Tumor Distribution Model

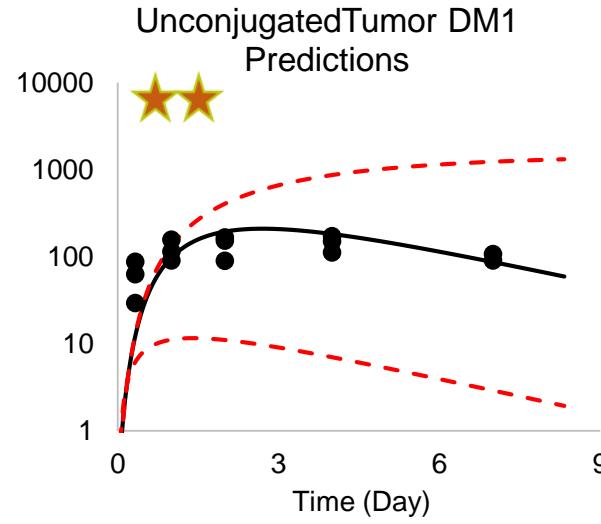
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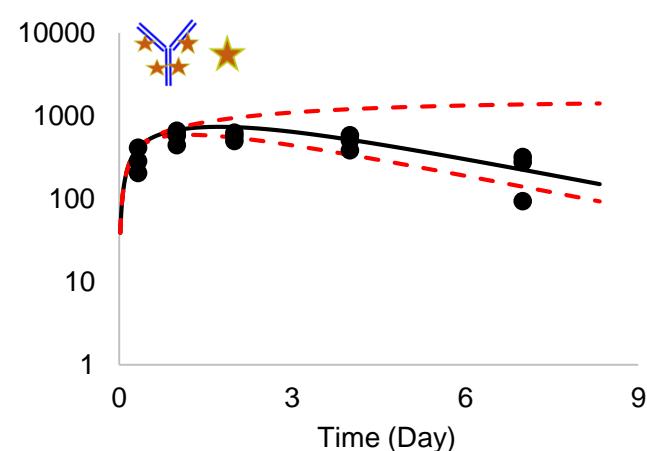
Total Plasma DM1 Predictions



Unconjugated Tumor DM1 Predictions

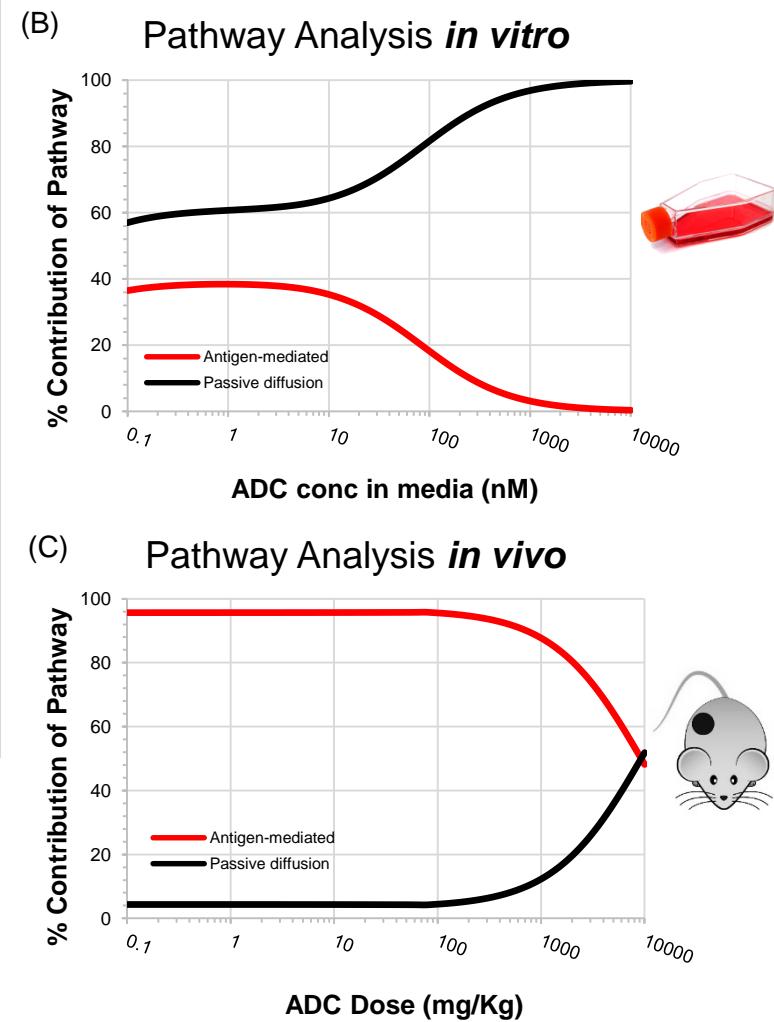
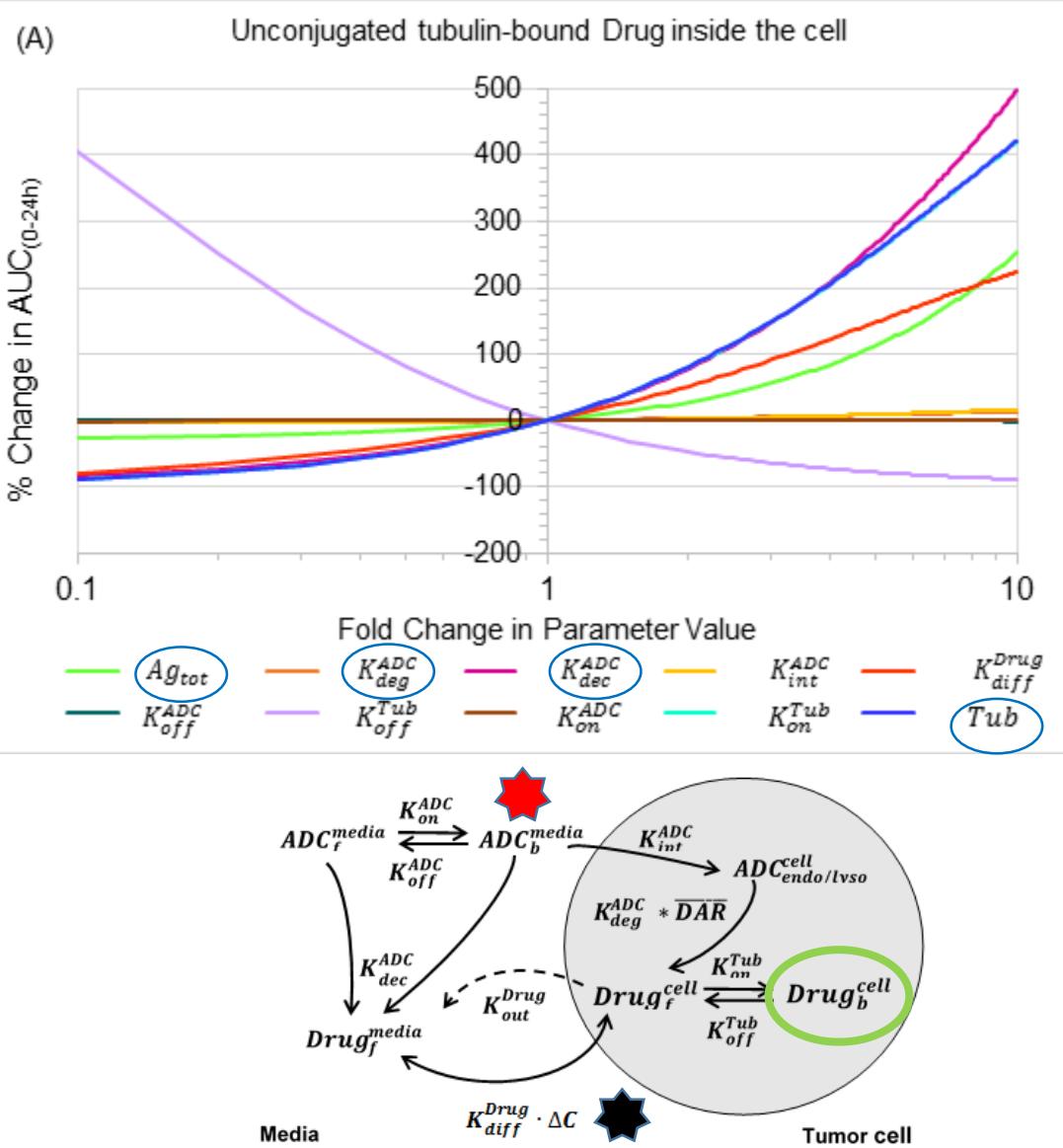


Total Tumor DM1 Predictions



Monte-Carlo simulations were performed with *IIV* of **17.4 % on  $K_{diff}$  parameter.**

# Pathway and Sensitivity Analysis

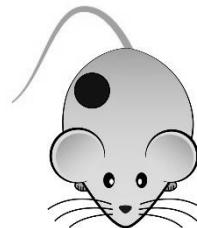


# Integrated Tumor PK-PD model for T-DM1

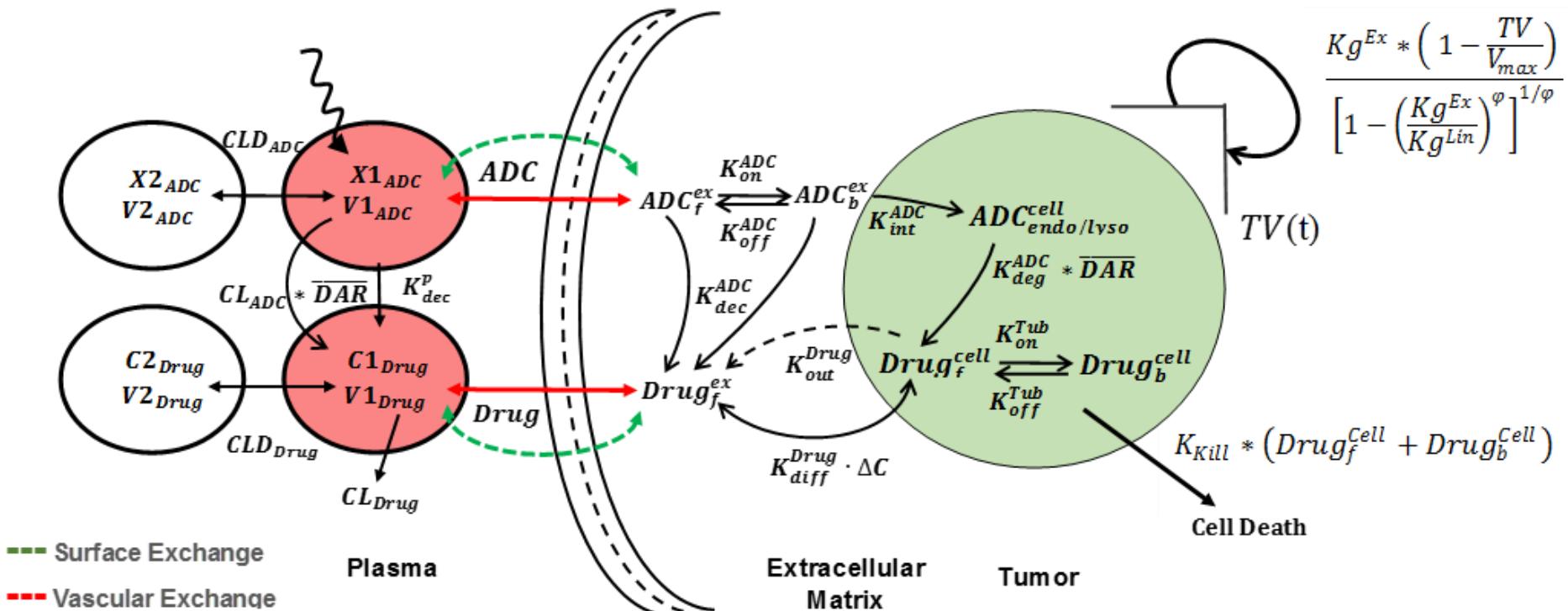
Use the Total DM1 Catabolite concentrations in the tumor to characterize the Tumor Growth Inhibition Data :

**In Vivo** Tumor Growth Inhibition (PKPD) Model

Use tumor concentrations to characterize TGI data and obtain PD parameters



Step 3



# Meta-Analysis on Several HER2 + TGI Datasets

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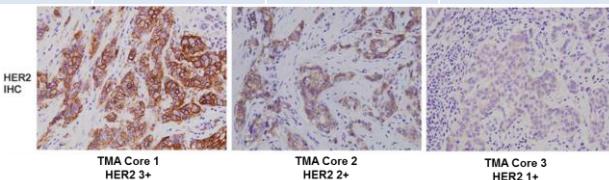
## Preclinical Tumor Growth Inhibition (TGI) Studies of T-DM1

Mouse Models	Model Type	HER2 Status	Dosing Regimen	Reference
BT474EEI	Xenograft	3+	a) 0.3-15 mg/Kg Q3WX3 b) 3.3-18 mg/Kg Q1WX9 c) 6-18 mg/Kg Q2WX5	Jumbe et al (2010)
Fo5	Breast tissue-derived orthotopic and metastatic (BOM) model	3+	a) 1-30 mg/Kg Q3WX3 b) 3.3-10 mg/Kg Q1WX9	Jumbe et al (2010)
Calu-3	Xenograft	3+	a) 1-7 mg/Kg Single Dose b) 15 mg/Kg Q6DX3	a) Lewis Phillips et al (2013) b) Cretella et al (2014)
KPL4	Xenograft	3+	a) 0.3-3 mg/Kg Single Dose b) 15 mg/Kg Single Dose	a) Lewis Phillips et al (2013) b) Lambert et al (2014)
N87	Xenograft	3+	a) 1-10 mg/kg Single Dose b) 5 mg/Kg Single Dose	a) Haddish-Berhane et al (2013) b) Yamashita-Kashima et al (2013)
BT474	Xenograft	3+	0.2-5 mg/Kg Single Dose	Van der Lee et al (2015)
MAXF 1162	PDX	3+	1-10 mg/Kg Single Dose	Van der Lee et al (2015)
HBCx-34 HP	PDX	2+	3-30 mg/Kg Single Dose	Van der Lee et al (2015)
ST313 HP	PDX	2+	3-30 mg/Kg Single Dose	Van der Lee et al (2015)
HBCx-10	PDX	1+	3-30 mg/Kg Single Dose	Van der Lee et al (2015)
MAXF 449 TNBC	PDX	1+	3-30 mg/Kg Single Dose	Van der Lee et al (2015)

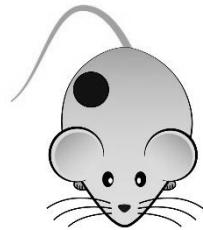
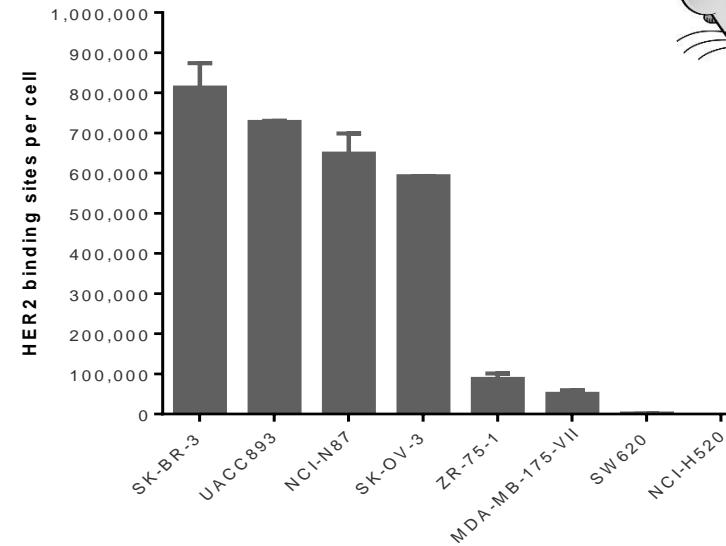
# Assumptions for HER2 Receptor Numbers in different Mouse Models

## HER2 Expression Level

Model	Type	HER2 FISH	HER2 IHC	ER/PR
BT474	CDX	+	3+	+
MAXF 1162	PDX	+	3+	-
ST313	PDX	-	2+	+
HBCx-34	PDX	-	2+	+
MAXF 449	PDX	-	1+	-
MAXF MX1	PDX	-	1+	-
HBCx-10	PDX	-	1+	-

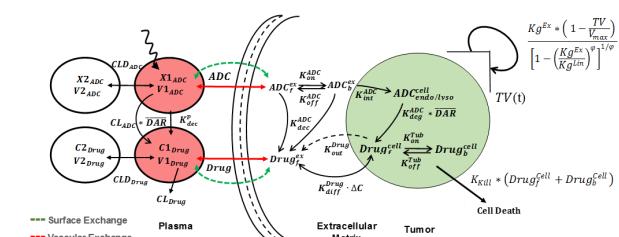


## Figure S3.



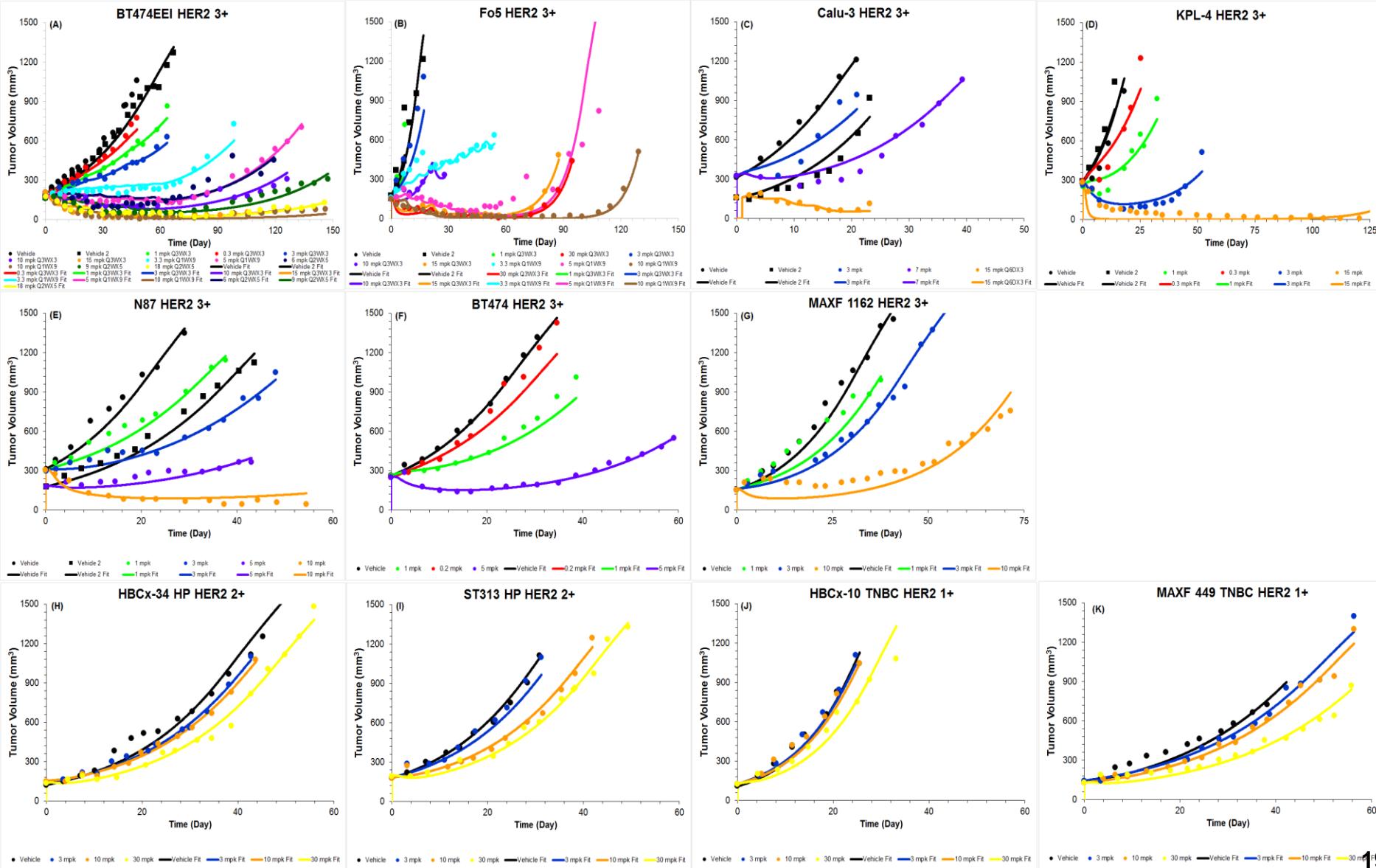
## **Assumptions:-**

1. HER2 3+ ~ 1 million, HER2 2+ ~ 0.5 million and HER2 1+ ~0.1 million Receptors
  2. Rest of all parameters were assumed the same

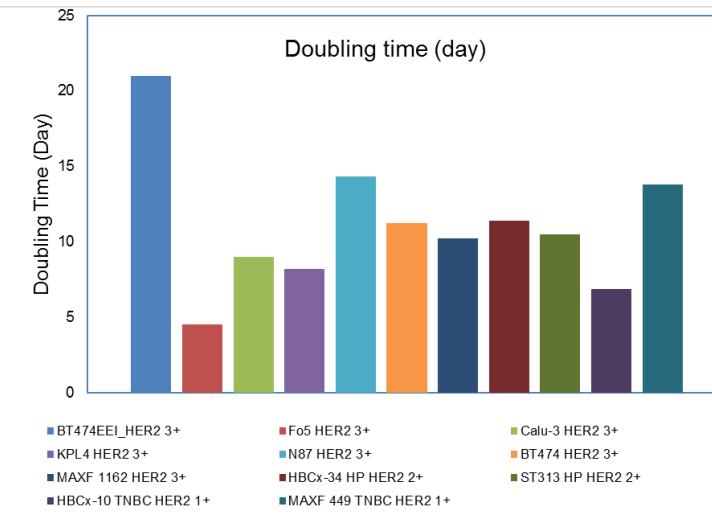
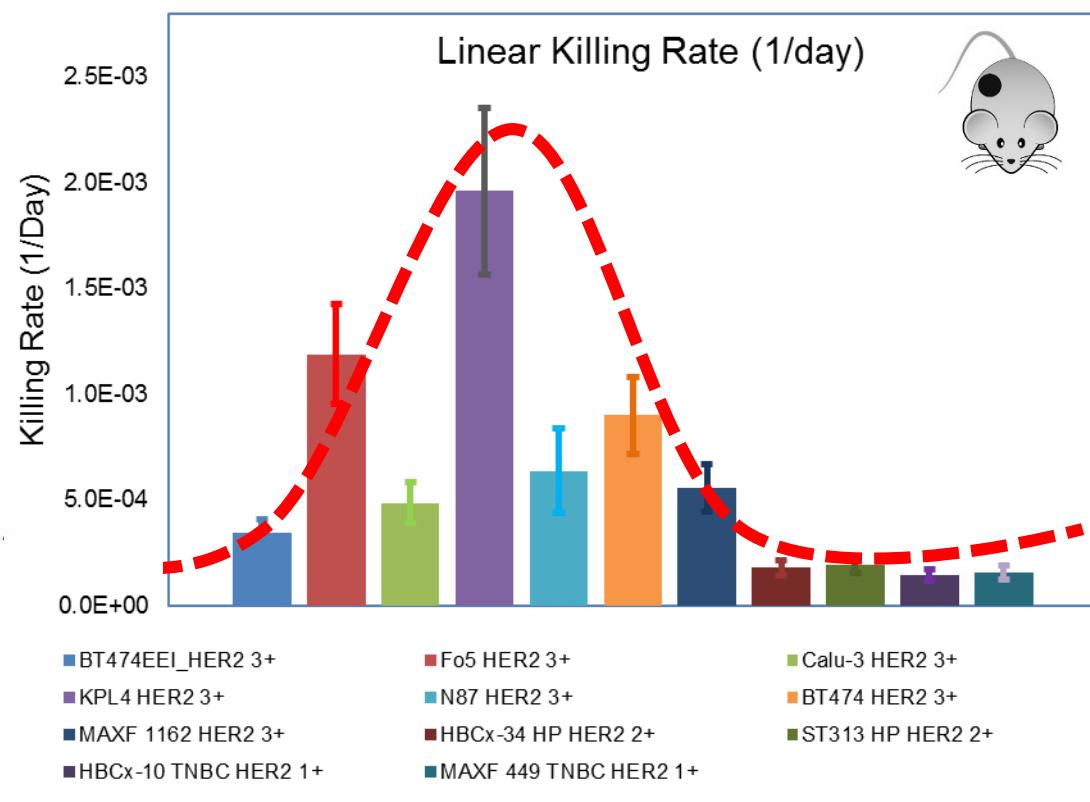


# PK-PD Model Fittings to Different HER2 + TGI Datasets

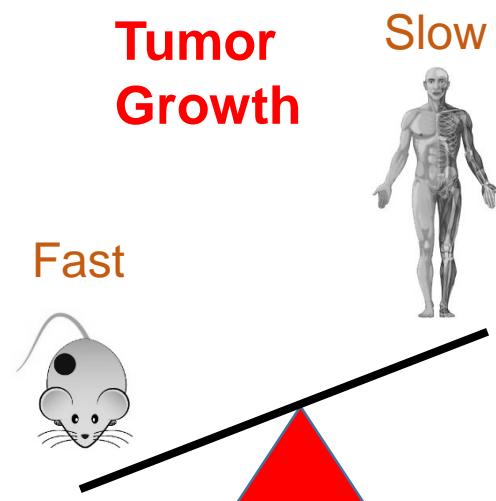
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# Distribution of Growth and Killing Parameters



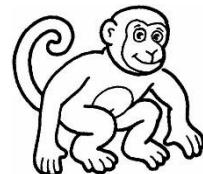
*Resample from this Distribution to Predict Clinical Efficacy*



# Characterization of Plasma Pharmacokinetics in Monkeys

**In Vivo** Plasma PK Model in Monkey

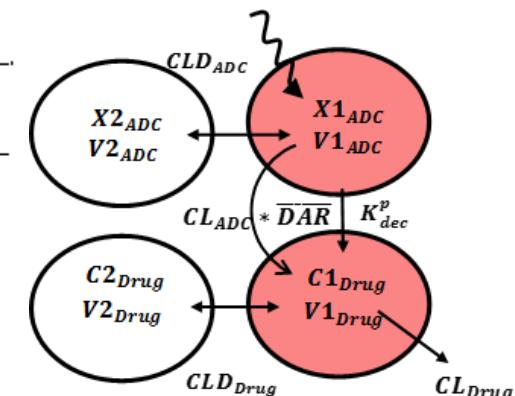
Characterize systemic concentrations of different analytes of ADCs in monkeys



**Step 4**

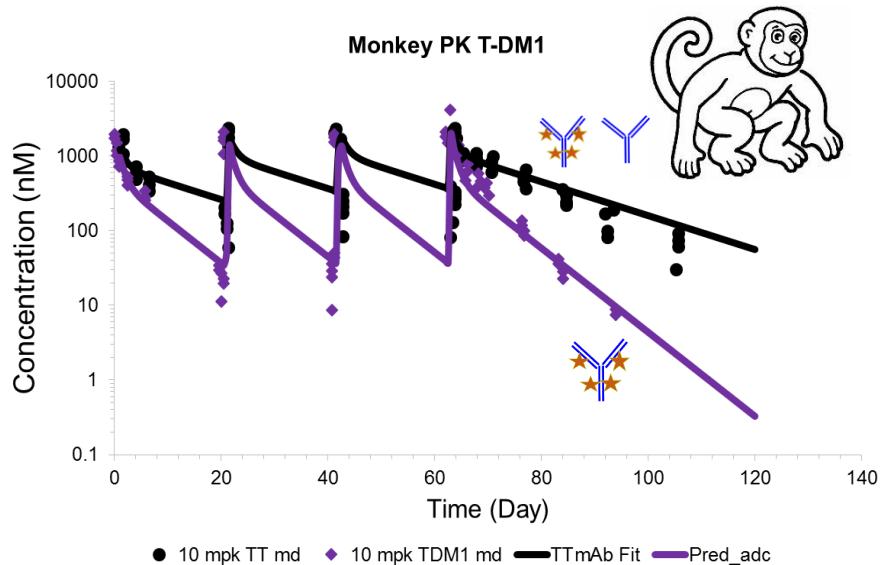
## Pharmacokinetic Studies of T-DM1

Study	Species	Dosing Regimen	Reference
Preclinical (Toxicokinetic)	Cynomolgus Monkeys	10 mg/kg Q3WX4 (TTmab and T-DM1)	a) & b) Bender et al (2014)
		30 mg/Kg Single Dose (TTmAb and T-DM1)	c) Poon et al (2013)
		30 mg/Kg Q3WX4 (TTmab, T-DM1 and DM1)	

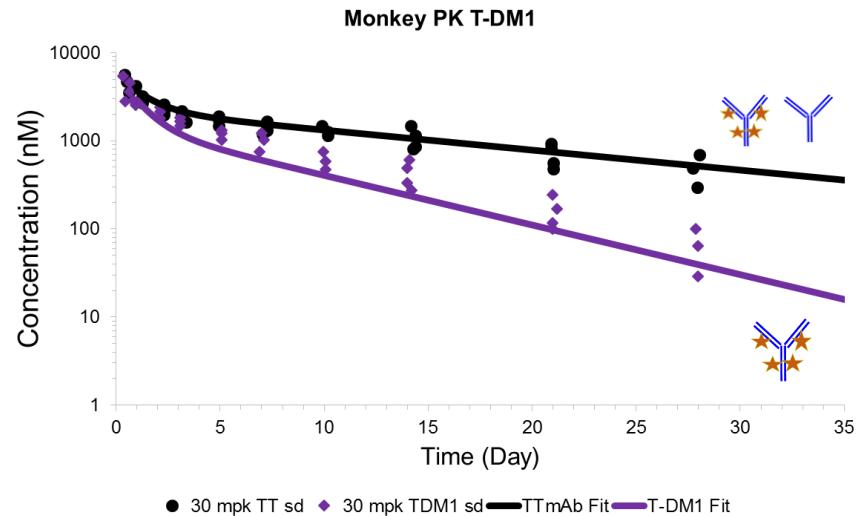
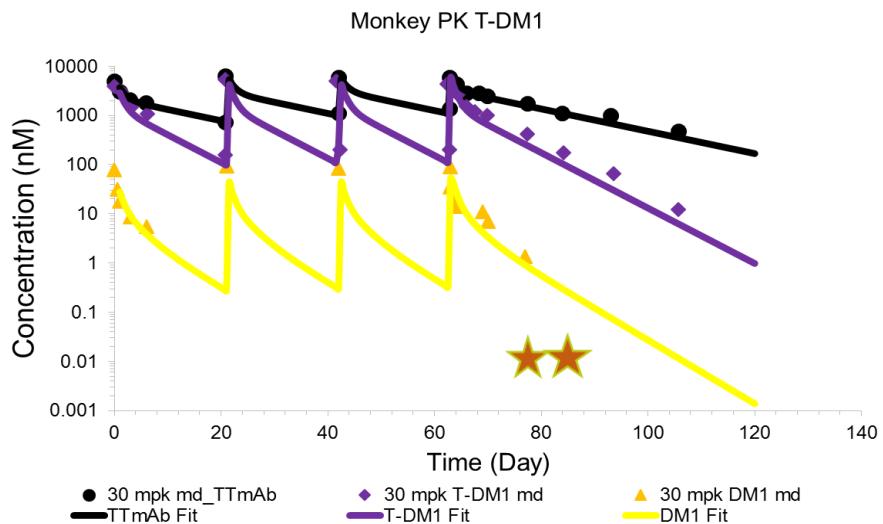


- Disposition of mAbs/ADCs is closer to Humans.
- Hence Non Human Primates (NHPs) are ideal species to predict human PK.

# Plasma Pharmacokinetics Model Fits in Monkeys



Systemic PK Parameters	Parameters	Units	Reference
$CL_{ADC}$	0.0043 (8.2 %)	L/day/Kg	Estimated
$CLD_{ADC}$	0.014 (48 %)	L/day/Kg	Estimated
$V1_{ADC}$	0.034 (14 %)	L/Kg	Estimated
$V2_{ADC}$	0.04 (32 %)	L/Kg	Estimated
$CL_{Drug}$	2.92 (32%)	L/day/Kg	Estimated
$CLD_{Drug}$	1.0 (2.62%)	L/day/Kg	Estimated
$V1_{Drug}$	0.034	L/Kg	Fixed
$V2_{Drug}$	5.0 (7.9%)	L/Kg	Estimated
$K_{dec}^P$	0.241 (8.8%)	1/day	Estimated



# Allometric Scaling of Monkey PK parameters

$$CL^{human} = CL^{monkey} \times \left\{ \frac{BW^{human}}{BW^{monkey}} \right\}^e$$

$$V^{human} = V^{monkey} \times \left\{ \frac{BW^{human}}{BW^{monkey}} \right\}^e$$

For TTmAb and T-DM1:

$e = 1$  for  $CL, CLD, V1$  and  $V2$

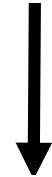
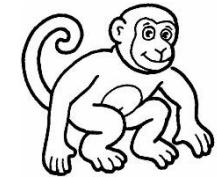
For DM1 Catabolites:

$e = 0.75$  for  $CL, CLD$   
 $1$  for  $V1$  and  $V2$

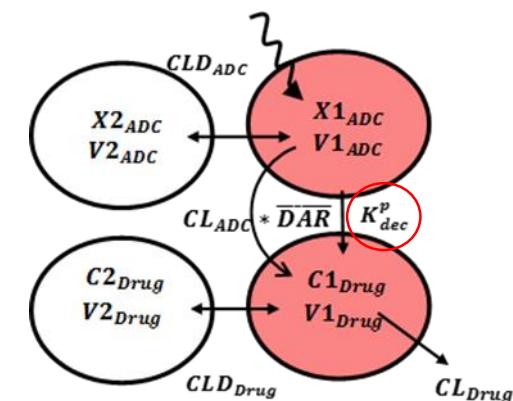
**Predict** Clinical PK from monkey  
**Predict** Clinical PD from mouse

Scale up monkey PK parameters to predict human PK. Use mouse PD parameters to predict Progression Free Survival

**Step 5**

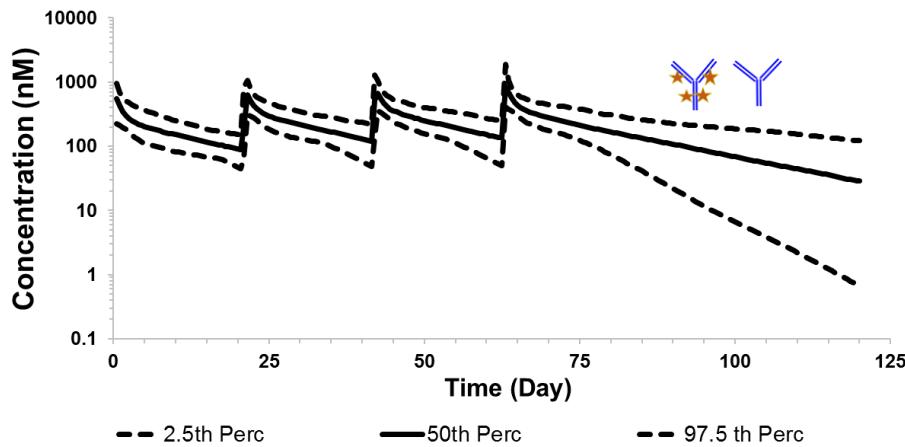


Pharmacokinetic Studies of T-DM1			
Study	Species	Dosing Regimen	Reference
Clinical (Phase-1&2)	HER2-positive Metastatic Breast Cancer Patients	3.6 mg/Kg Q3WX4 (TTmAb, T-DM1 and DM1)	a) Burris et al (2011)
		3.6 mg/Kg Single Dose (TTmAb, T-DM1 and DM1)	b) Agresta et al (2011)

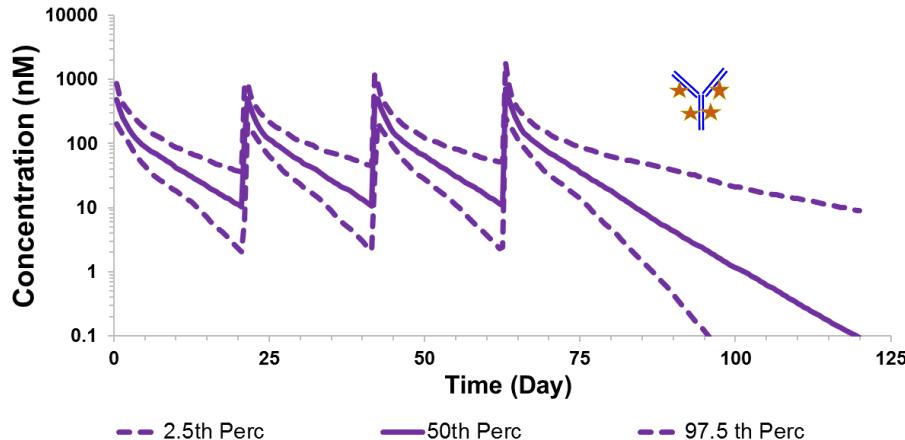


# *A Priori* Predicting Human Pharmacokinetics of T-DM1

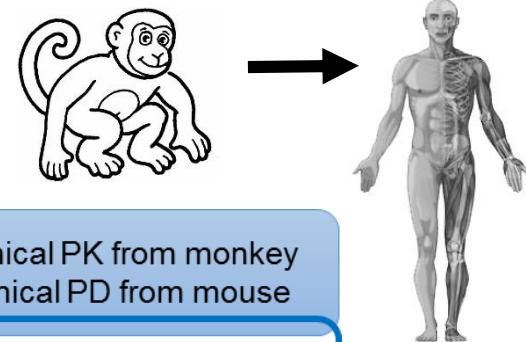
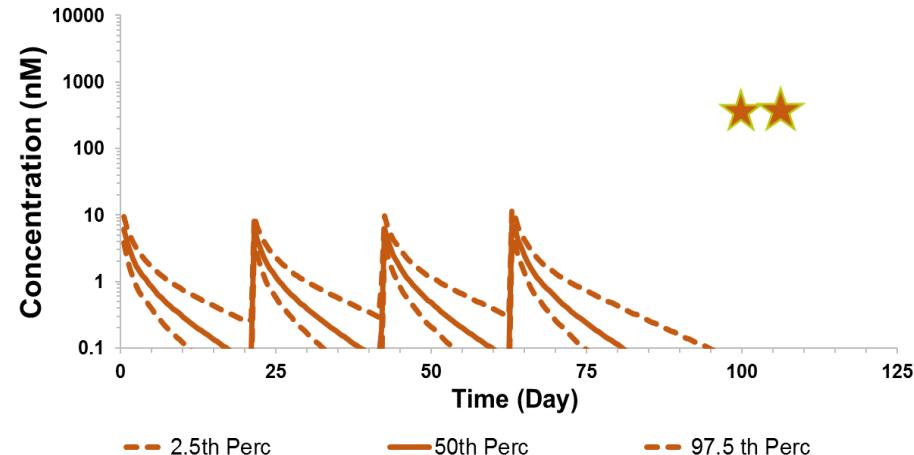
Human PK TTmAB



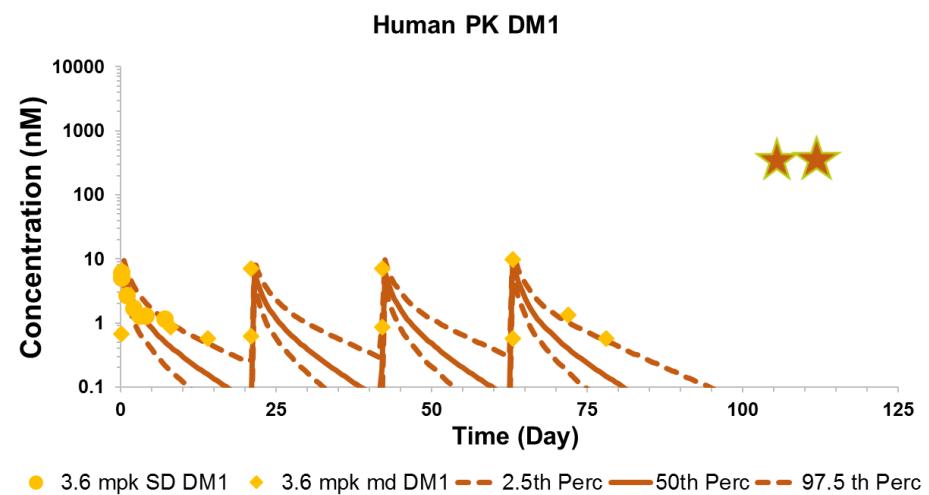
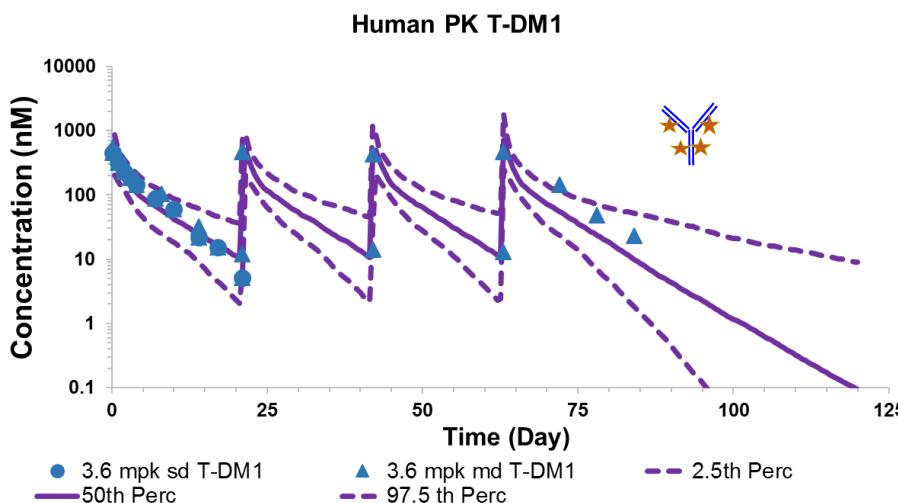
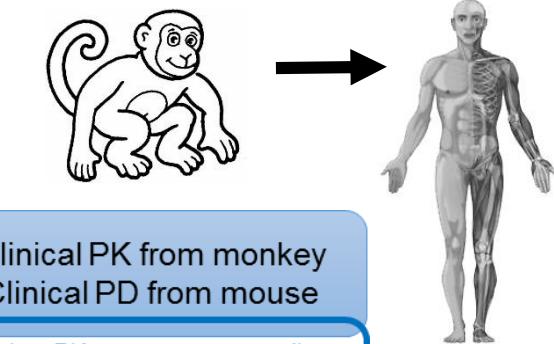
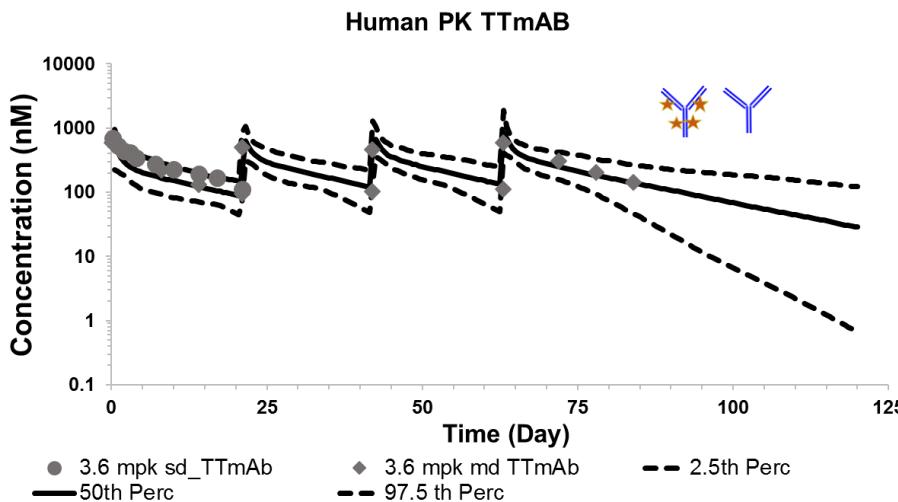
Human PK T-DM1



Human PK DM1



# **Validation** of Our Human PK Predictions for T-DM1

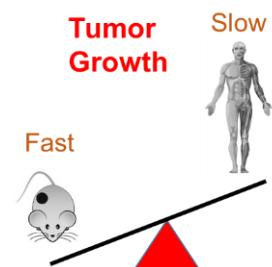


# Preclinical to Clinical Translation:- Clinical Growth Rate

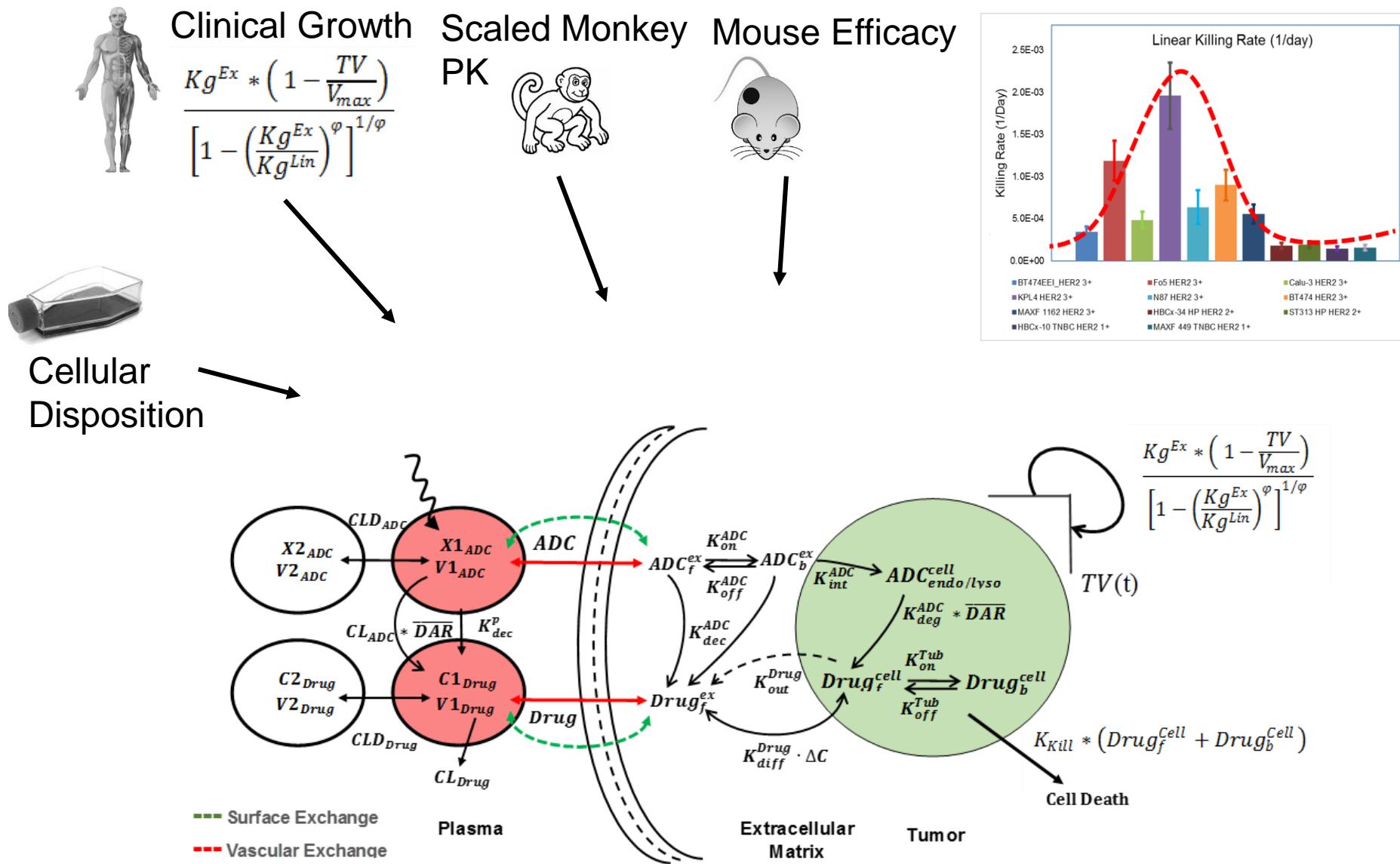
A list of clinically relevant tumor growth parameters for prediction of Progression Free Survival predictions.



Parameters	Definition	Value (CV %)	Units	Source
TV(0)	Initial tumor volume derived using the following expression $TV(0) = \frac{1}{2} * L * B^2$	Initial tumor lesion length = 19 mm (101%) Initial sum of Lengths and Breadth for tumor lesions = 35 mm (125%)	mm <sup>3</sup>	Bernadou et al (2016)
DT <sup>Exp</sup>	Doubling time associated with the exponential growth phase of the tumor	25 (200%)	Day	Pearlman et al (1976)
DT <sup>Lin</sup>	Doubling time associated with the linear growth phase of the tumor	621 (85%)	Day	Weedon-Fekjær et al (2008)
φ	Switch between exponential growth and linear growth phases.	20	Unitless	Haddish-Berhane et al (2014)
V <sub>max</sub>	Maximum achievable tumor volume	523.8	cm <sup>3</sup>	Assumed based on maximum achievable tumor radius to be 5 cm



# Preclinical to Clinical Translation Strategy



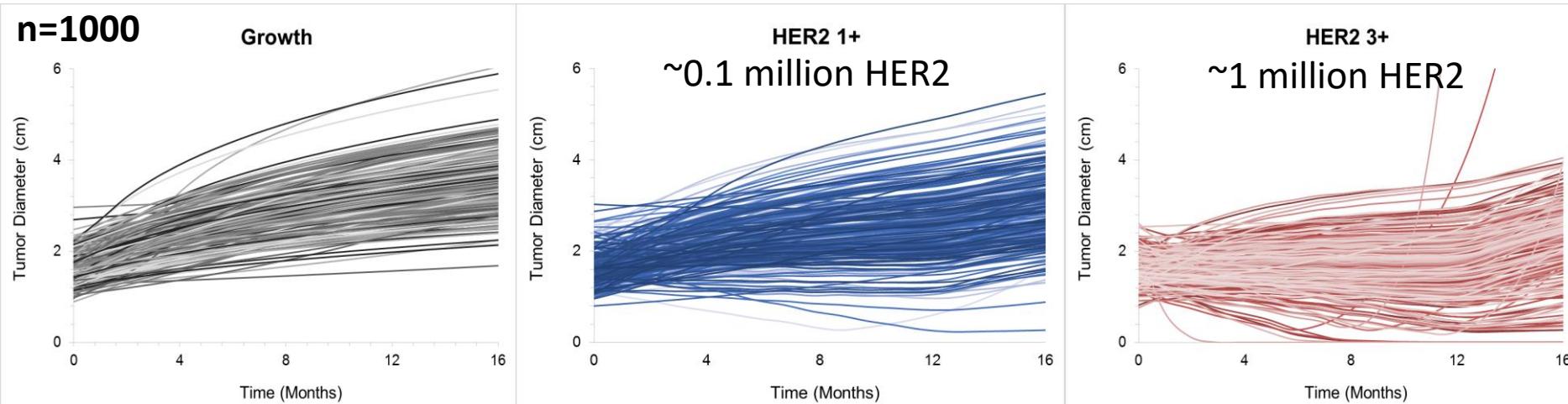
# Preclinical to Clinical Translation: *List of Clinical Trials*

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## Clinical Trials of T-DM1

Study	Patient Population	Phase	Number of Patients	Dosing Regimen	Treatment Arms	Reference
TDM4450	First-Line HER2+ Metastatic Breast Cancer Patients	II	137	3.6 mg/Kg Q3W until Disease Progression	T-DM1 versus Trastuzumab+ docetaxel	Hurvitz et al
TDM4258g	HER2+ Metastatic Breast Cancer Patients with prior treatment with Trastuzumab	II	112	3.6 mg/Kg Q3W up to 12 months	Trastuzumab followed by T- DM1	Burris III et al Krop et al
EMILIA	HER2+ Metastatic Breast Cancer Patients with prior treatment with Trastuzumab and Taxane	III	991	3.6 mg/Kg Q3W until Disease Progression	T-DM1 versus capecitabine + lapatinib	Verma et al

# Preclinical to Clinical Translation: Clinical Trial Simulations



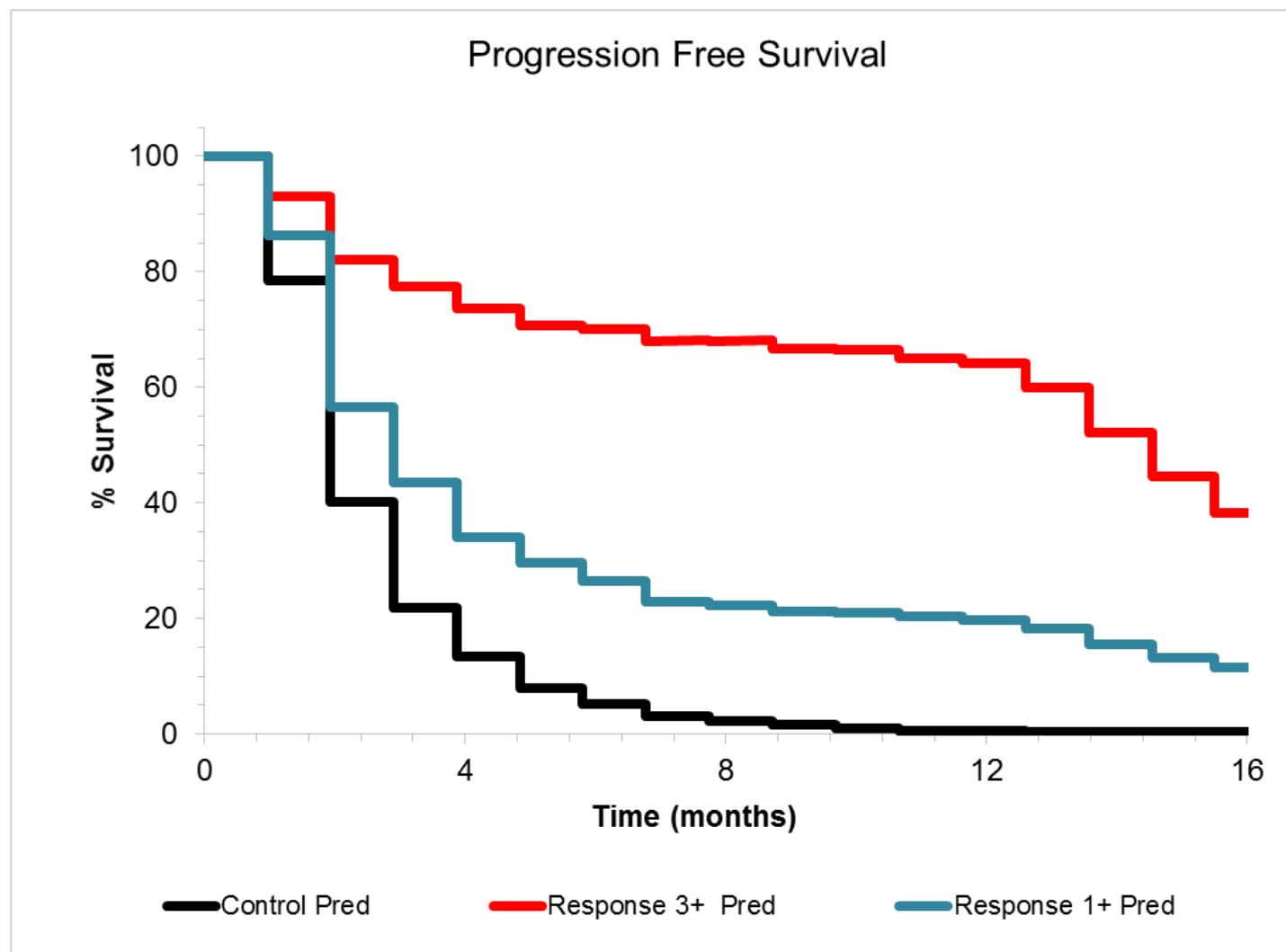
- *T-DM1 administered at a Dosing regimen of Q3W for 1 Year (3.6 mg/Kg Dose)*

## Calculation of PFS

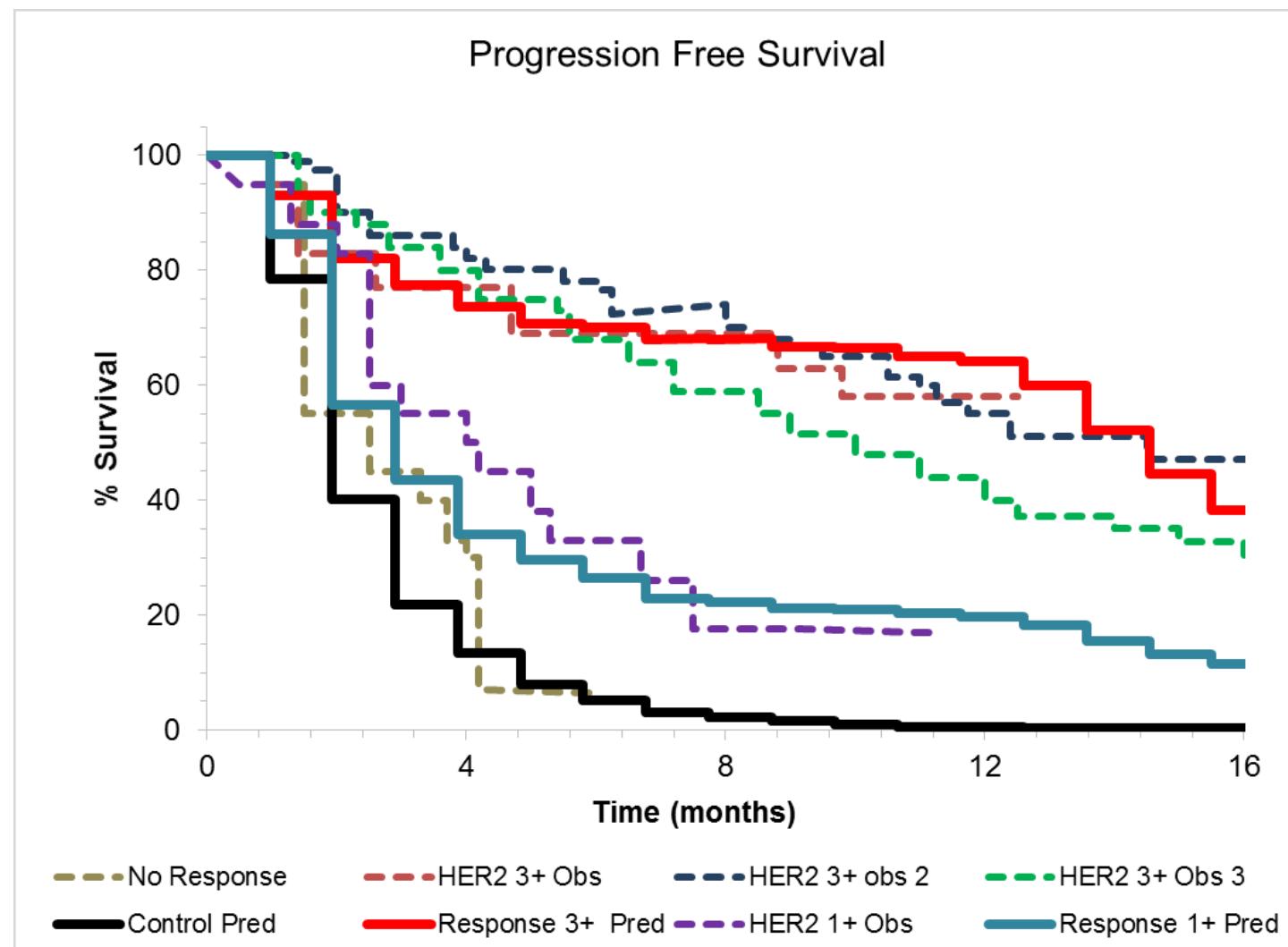
## Response Evaluation Criteria In Solid Tumors :- RECIST

- 1) **Complete Response (CR):** Undetectable <5-10 mm.
- 2) **Partial Response (PR):** At least a **30% decrease** in the sum of diameters of target lesions.
- 3) **Progressive Disease (PD):** At least a **20% increase** in the sum of diameters of target lesions.
- 4) **Stable Disease (SD):** Neither sufficient shrinkage to qualify for PR nor sufficient increase to qualify for PD, (**<20% increase and <30% decrease**).

# Our Predicted Progression-Free Survival Rates

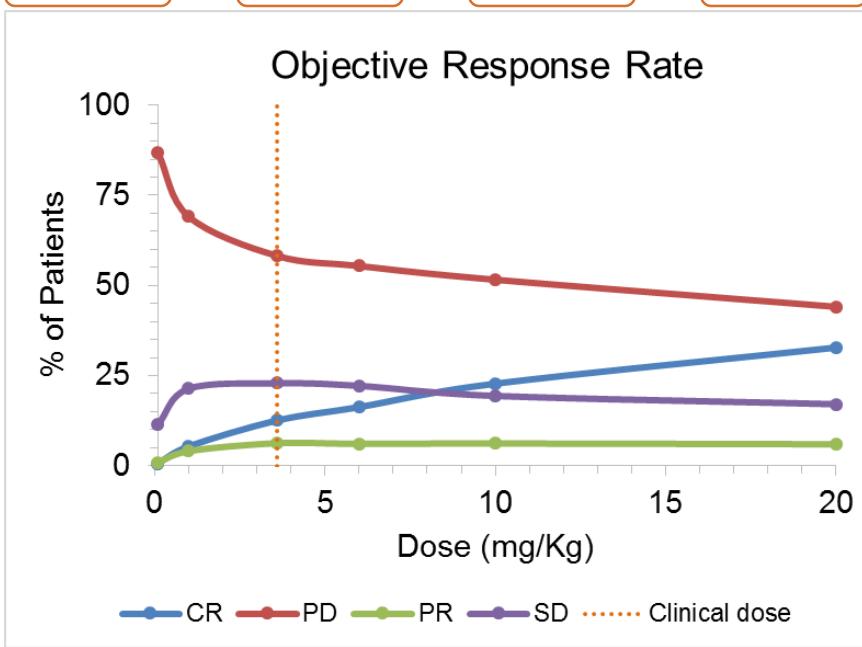


# Our Predicted Progression-Free Survival Rates



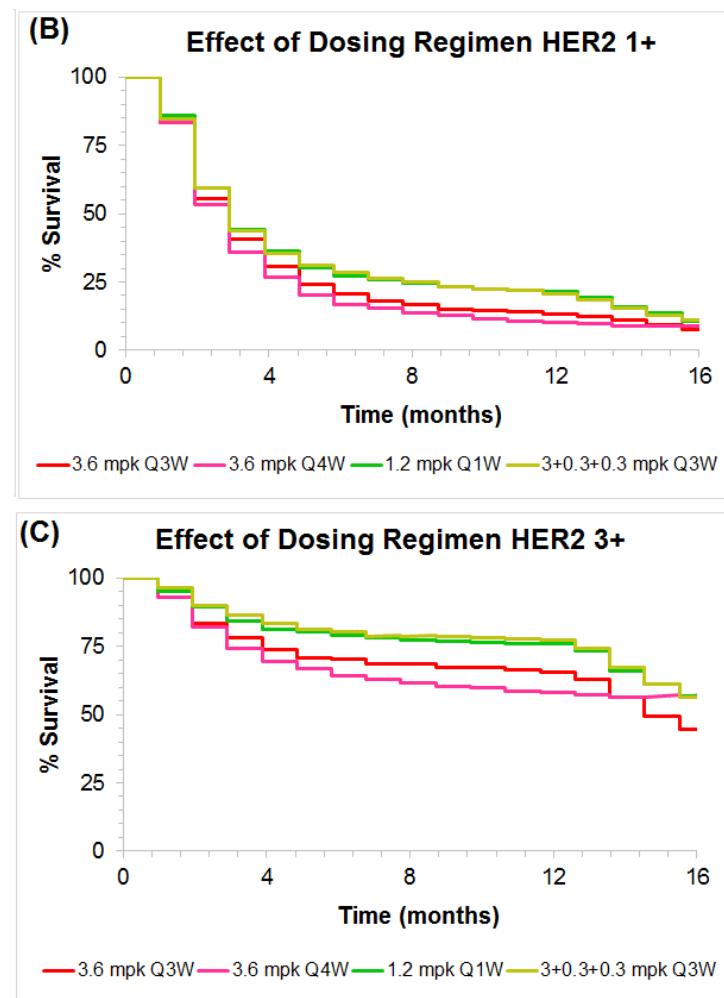
- **Control:-** HER2 Normal Patients non responsive to T-DM1 and Trastuzumab
- **HER2 1+:-** Below Median HER2 Expressing Patients
- **HER2 3+:-** HER2 Over-expressing Patients

# Simulating Objective Response Rates (ORRs) and Alternative Dosing Regimens



ORRs Calculated at the end of 1 Year Therapy

- Not a Significant Improvement in Efficacy with a slight increase in Dose



- Fractionated Dosing Regimen e.g. Front Loading can be more beneficial

# Overall Summary

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- A mechanistic Cellular Disposition Model was developed with more intracellular details (i.e. ADC degradation, passive diffusion and tubulin binding ) to characterize Cellular PK of T-DM1.
- When Combined with Tumor Disposition Model, we were able to *a priori* predict tumor pharmacokinetics of T-DM1
- Later the model was *translated to clinic* where mouse efficacy parameters, clinically observed growth parameters and scaled up human PK from NHPs was able to predict clinical efficacy in *different sub-populations of HER2 + patients*
- Our multi-scale PK-PD Model has been validated on multiple ADC molecules (n=3) and can help inform about the Dose-ORR relationships as well as alternative dosing regimens.
- Presented Translation Strategy can serve as a cornerstone for developing future ADCs.

# Acknowledgements



*Shah Lab Picture*

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**AMGEN**

**abbvie**

**ucb**

**SANOFI**

Roche

NIGMS (NIH)

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4. Professor K. Dane Wittrup (MIT)
5. Katie F. Maass (Genentech®)



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