

# ***Risk Assessment for the measurement of Free and Total Drug and Target***

Lindsay E. King Ph.D.  
PDM NBE, Pfizer Inc.

European Bioanalysis Forum  
18 Nov 2011

# Acknowledgements

---



- Huifen Faye Wang
- AAPS Free Total Focus Group  
White paper
- Chris Grace
- Jie Guo
- Leslie Lorello
- Eve Su
- Rob Webster
- Ellen Wang
- Alison Betts
- Rick Steenwyck
- Derek Winslow
- Tracy Clark
- Joann Wentland
- Jenny Wang
- Guodong Zhang
- Boris Gorovits
- Catherine Soderstrom
- Frank Spriggs



Bioanalytical scientists have become increasingly aware of the potential need to measure Free and Total analyte as well as some of the associated bioanalytical challenges

Despite this awareness there are varied degrees of engagement on this issue by project teams and BA scientists


- Given the rapid progress of projects and need to conserve resources how much effort should be exerted to build and define free/total assays and at what stage?
- What are the risks, how can they be assessed and what are the benefits of doing this?
  - Target Measurement
  - Target/Drug interference
  - Equilibrium Disruption

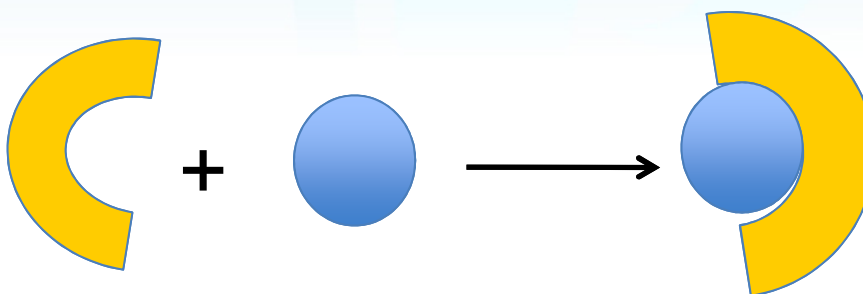
This talk will discuss analytical risk assessment and mitigation using both theoretical and experimental approaches for monoclonal antibody drugs and their targets.

# Introduction

## Free and Total: Clinical Utility

Free D: 

Free L: 



### Bioanalytical Challenges Re Form of Analyte

- Reagent Specificity:
- Free vs. Complex vs. Total:
- Matrix Interference:

Data  
Data  
Data

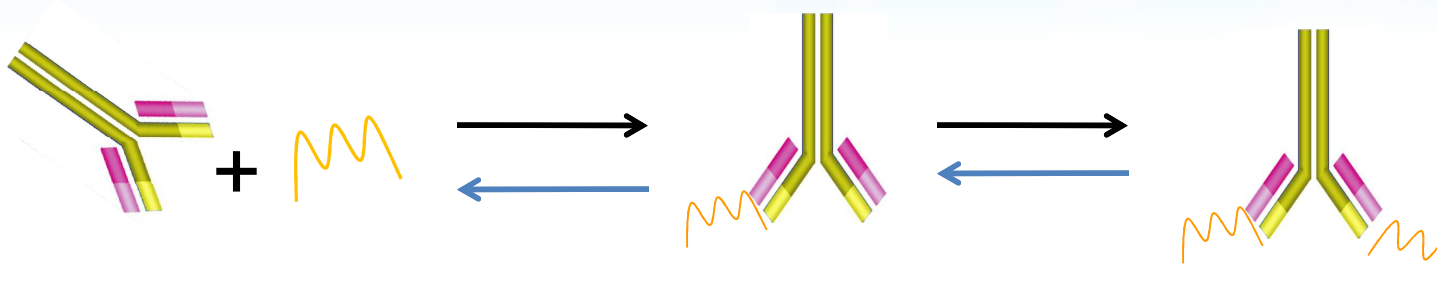
### Utility Challenges: Why are we doing this?

- Normal vs. Disease: Data
- Free less than Y = Normal
- Total Greater than X = Clinical Implications

**Total : Free +  
Complex**

# Introduction

Free and Total: PK : Two site, non-covalent binding



**Free D:**

**Partially Free D:**

**Free L:**

**Total : Free +  
Complex**

## Bioanalytical Challenges

- Reagent Specificity: Data
- Native vs. Cleaved T: ??
- Matrix Interference: Data
- Free vs. Complex vs. Total: ??
- Equilibrium Disruption: ??



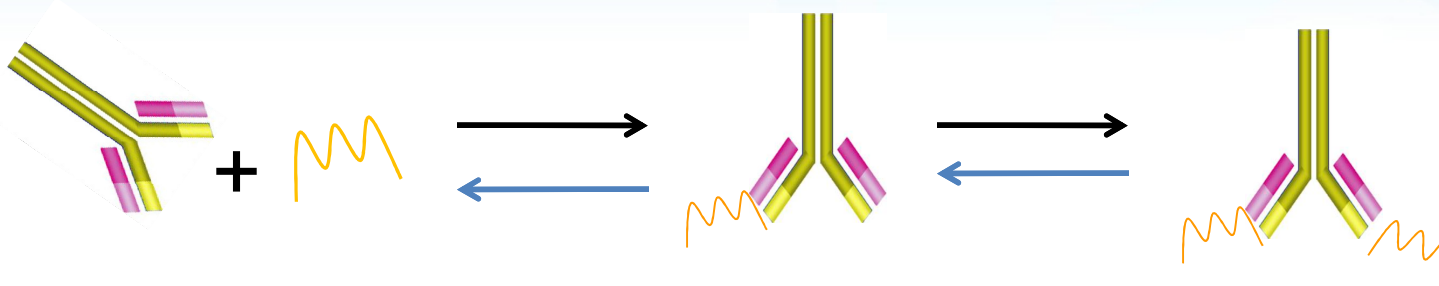
Utility Challenges: Why are we doing this?

How much effort should we expend?



# Introduction

Free and Total: Target :



**Free D:**

**Partially Free D:**

**Free L:**

**Total : Free +  
Complex**

Bioanalytical Challenges Re Form Target

1) Potential for Disulfide bridge formation



2) Trimeric Target

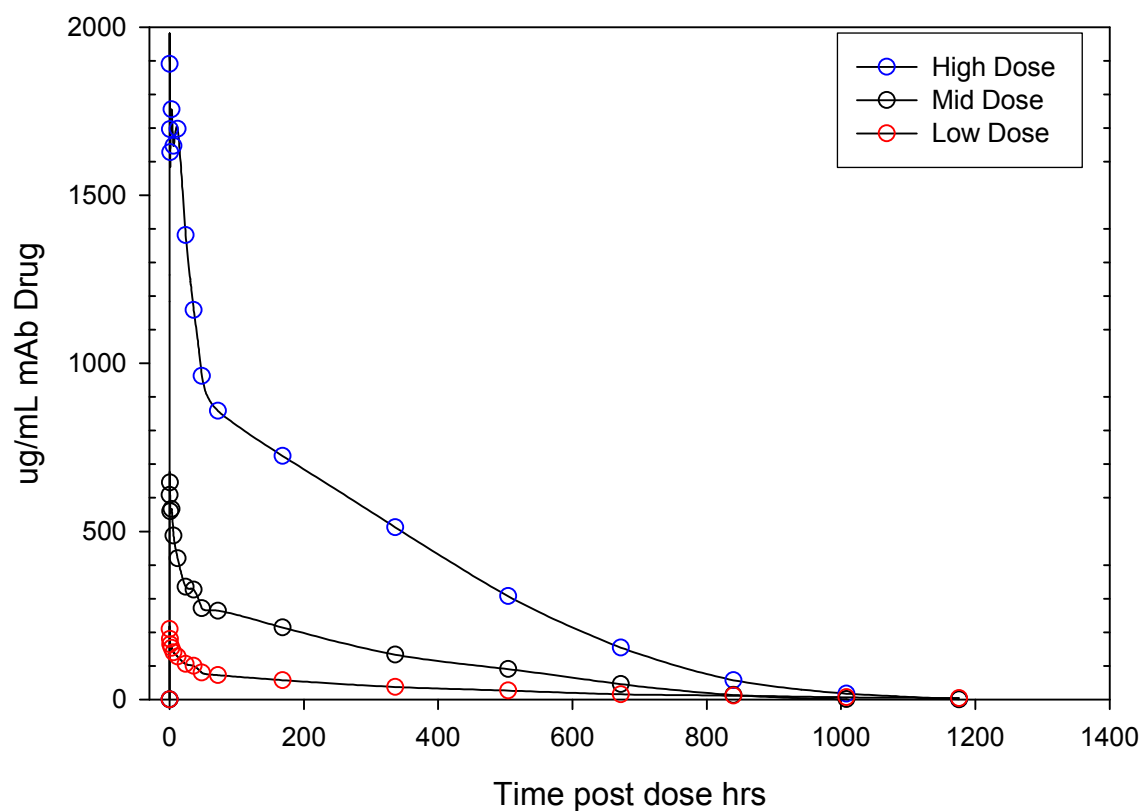


3) Weak bound heterogeneous aggregates

# When Should We Be Concerned ?

Ignorance is Bliss ?

Time vs Concentration Profile



Total or Free Assay?

Clinical Phase  
Question

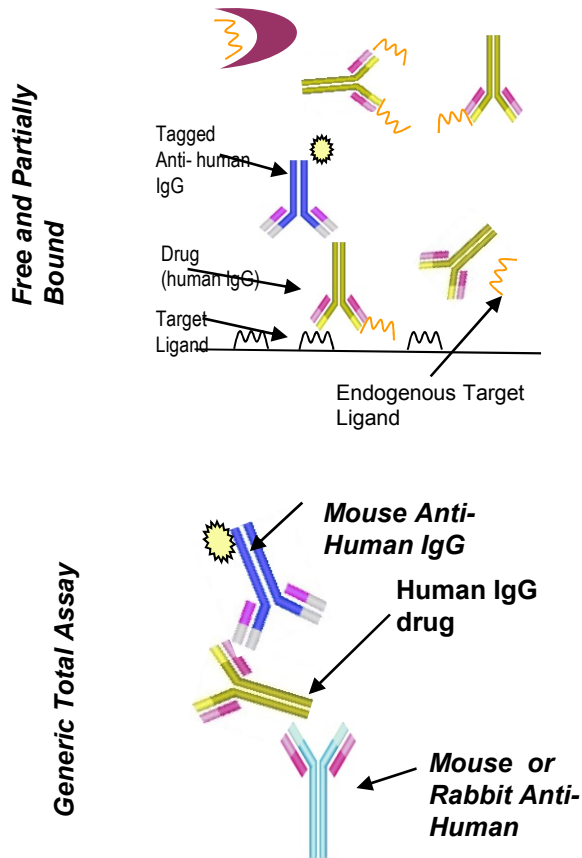
What is [target] ?

Can it interfere in PK  
assay?

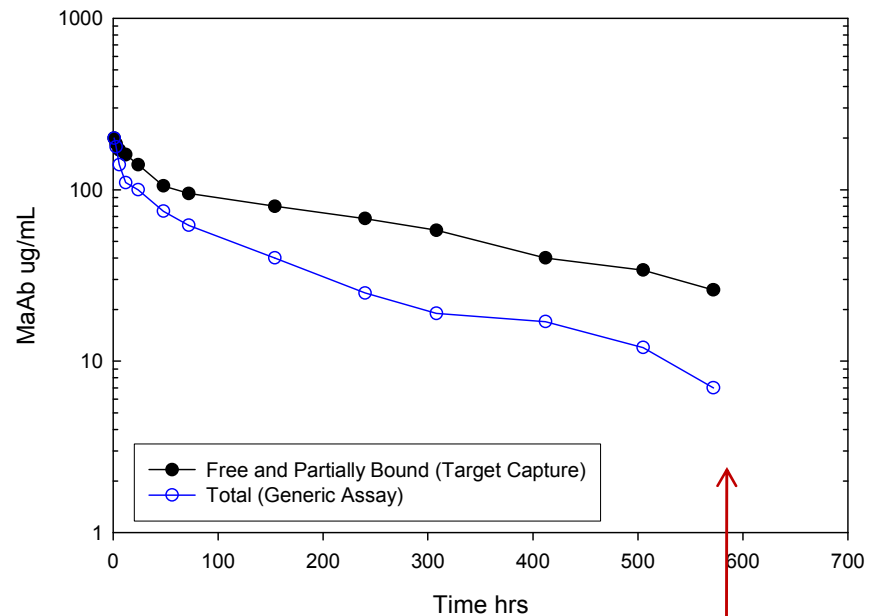
# PK Assay

Impact of lack of confidence in data

## Enlightened PK/ PD Driven Bioanalytical support for IVT study



## Assay format defines form of analyte?



Which data set is correct?

Target interference ?

Complex Dissociation?

Other?



# Potential Target Interference:

## Key Questions



- Is the target is soluble or shed
- What is the target concentration in normal and disease

### PK

- Can the target can interfere in PK assay
- When can the target interfere in PK assay

### Target PD

- Can the drug interfere in PD assay
- When can the drug interfere in PD assay

### PK and PD Assays

- Is equilibrium disruption a real concern

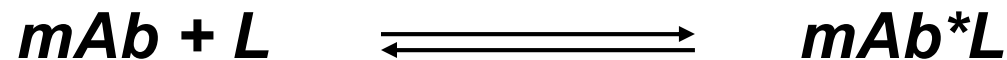


# Binding Equilibrium

*Law of mass action*



Foundation of the binding relationship of the drug (mAb) to the target ligand (L):

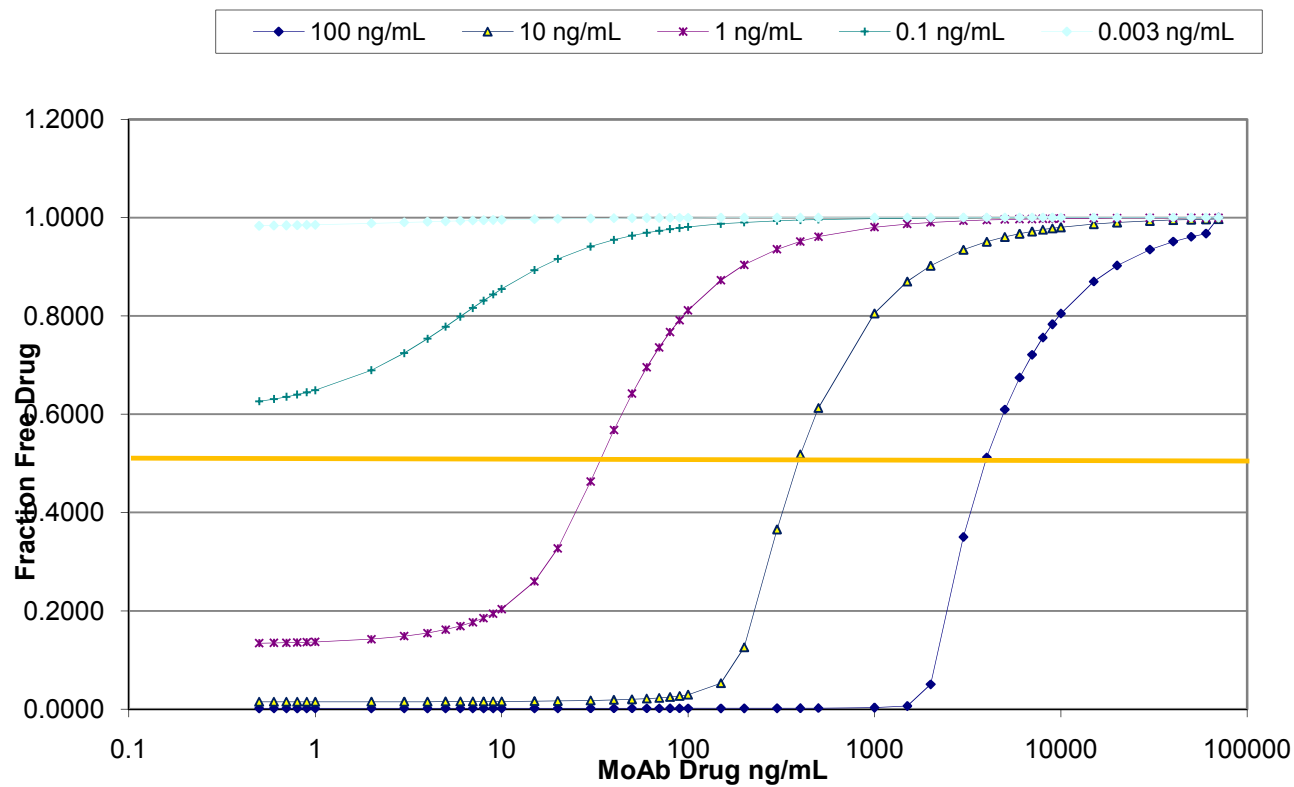


$$K_d = \frac{1}{K_a} = \frac{[mAb]_{\text{free}} \times [L]_{\text{free}}}{[mAb * L]_{\text{bound}}} = k_{\text{off}} / k_{\text{on}}$$

The dynamics of target ligand-mAb binding may shift with dose and time

# Assessment of Target Interference

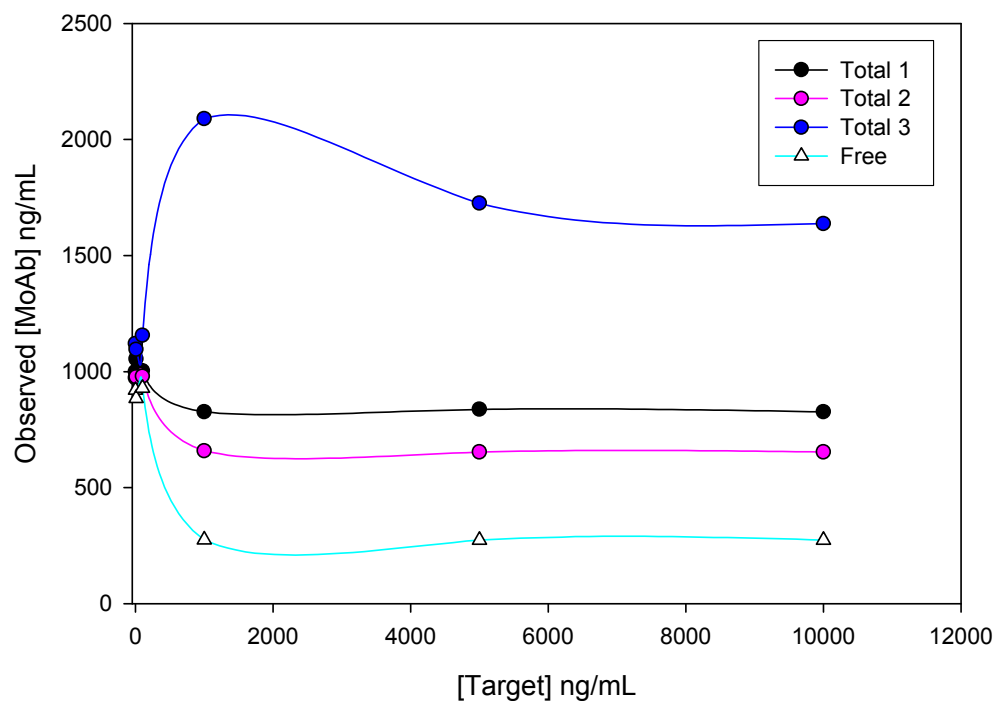
## Free and Partially Bound



Simulation based on Biacore derived Ab – Target KD

# Assessment of Target Interference

Effect of Target on Three Candidate Generic Total and a Free/Partially Bound PK assay



## What we did right:

- Equilibrium
- Defined Edges
- QC neat matrix + control

## What we might do different:

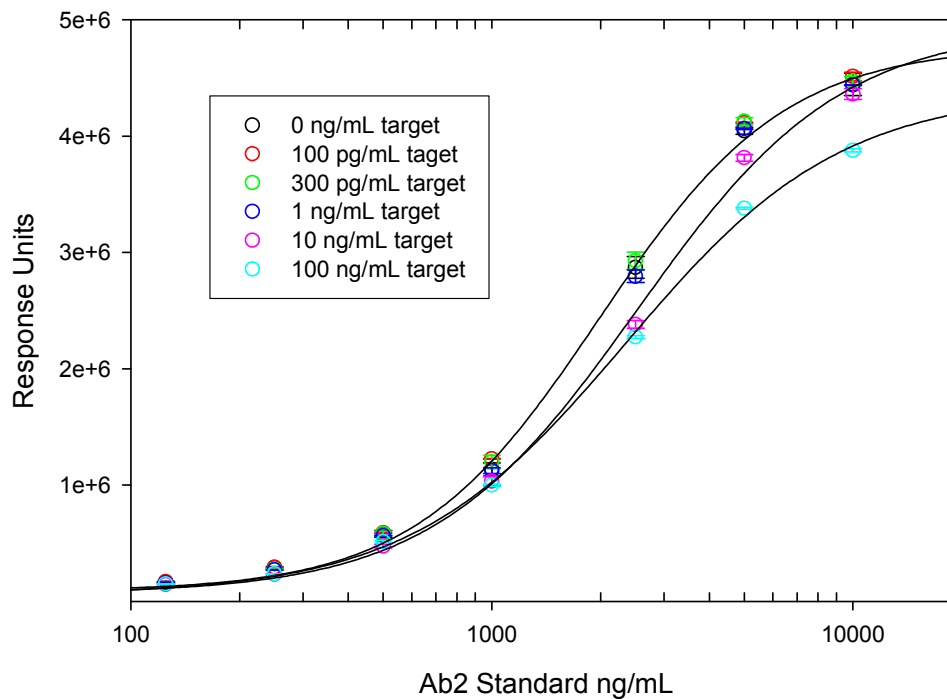
- Conduct interference testing before Sample Analysis: Assay utility?

Guo J, King L, Wang HF (2009) Development of generic total assays for quantitation of therapeutic human IgG2 monoclonal antibodies in nonclinical species using ligand binding assay. AAPS Journal 11(S1), 315.

# Anti-Id “Total” Assay

Unexpected Target Interference

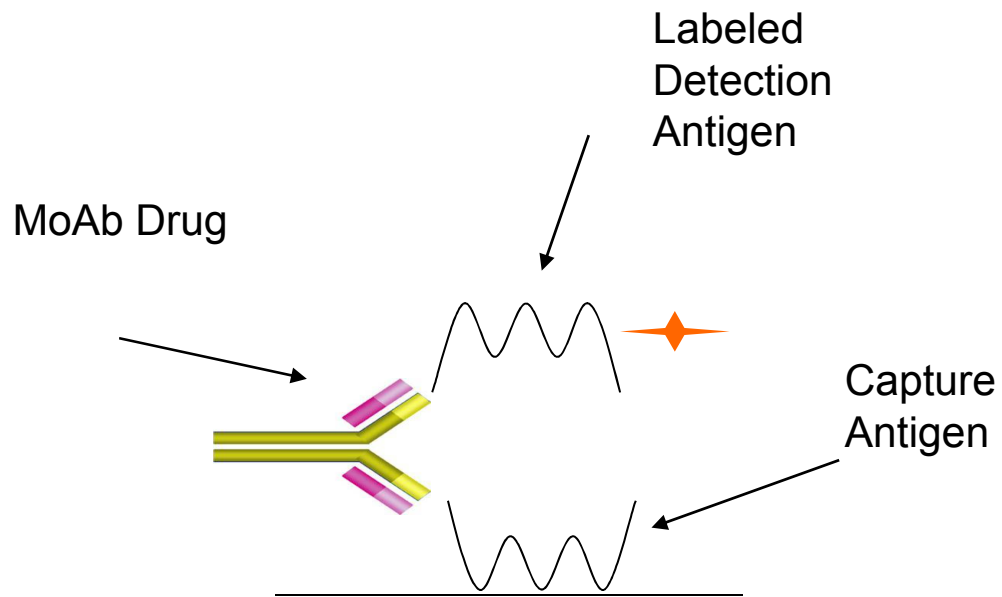
## Non Competing Anti-id based on Octet



# Free PK Bridging ELISA

*In principal assay measures only free drug*

In practice risk of  
overestimating Free?



# Assessment of Target Interference

MoAb Nominal µg/mL	0.5	0.5	0.5	0.5	1.2	1.2	1.2	1.2	1.75	1.75	1.75	1.75
rhTarget Spiked ng/mL	0	1	10	50	0	1	10	50	0	1	10	50
MoAb Observed µg/mL	0.669	0.601	0.529	0.308	1.35	1.26	1.116	1.11	1.85	1.83	1.84	1.52
Percent Recovery From Nominal	133.8	120.2	105.8	61.6	112.5	105.0	93.0	92.5	105.7	104.6	105.1	86.9
Free MoAb Predicted µg/mL	0.5	0.499	0.487	0.433	1.2	1.199	1.186	1.129	1.75	1.749	1.736	1.678
Binding Site Molar Ratio MoAb to target	NA	340	34	6.8	NA	816	81.6	16.3	NA	1190	119	23.8

## Result

***Assay appears to measure total drug at less than 10 ng/mL Target Antigen  
“Apparent Free”***



***Soderstrom et al***

# What data are most important to the team?



- PK data of Ab
- Antigen Baseline concentration measurement
- Turnover rate constant (T1/2 of Antigen)
- Free Antigen level after mAb dosing
- Complex level after mAb dosing
- Total Antigen concentration (=free+bound)

} Can be modeled

- Wide dose range





- Expected PD profile essentially validates assay is fit for purpose
- Free and Total target informs PK/PD relationship
- Drug interference testing needed

Total target data can be used

- to model the potential for target interference in the PK assay
- Trigger PK data review based on target interference data from PK assay validation

Free target can be used

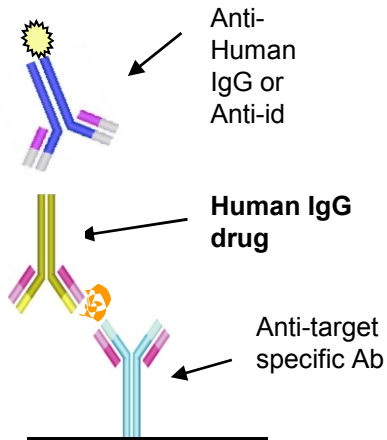
- to define target suppression and subsequent return to baseline after dosing

Equilibrium shifts during sample analysis can result in an overestimation of the amount of free observed

# Total Target as Biomarker

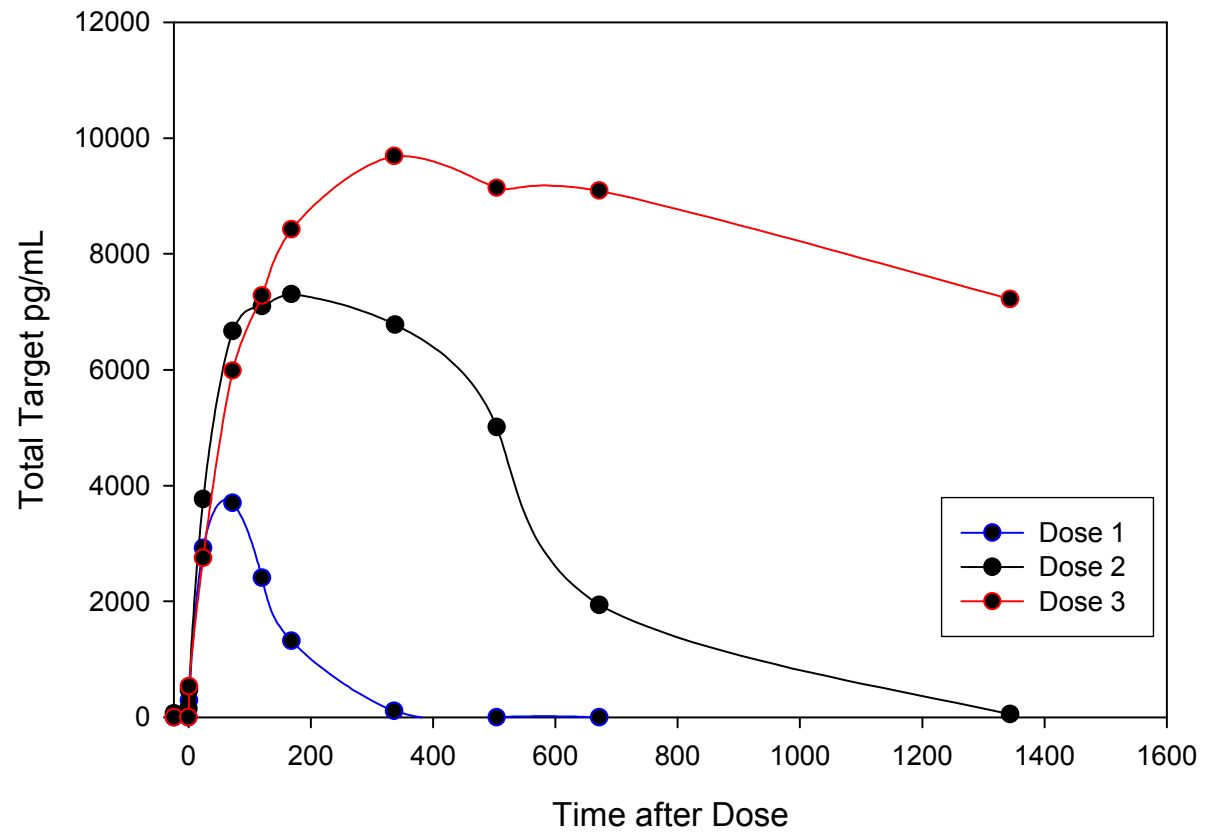
## Fit for Purpose

### Total Target Assay



Drug interference testing conducted

### Effect of Dose on Total Target Levels



Guo et al., 2009

# Sample manipulation and Handling

## Equilibrium Disruption



- Sample dilution can disrupt sample equilibrium
  - MRD in assay buffer 1:10.
  - Dilution in matrix pool; PK samples diluted significantly before the MRD: ULOQ 2 ug/mL
- Incubation time can disrupt sample equilibrium
  - Incubation time after MRD or matrix dilution
  - After sample dilution incubation time on plate
- Antigen coat concentration may contribute to equilibrium shift during plate incubations.
  - Locally high target concentration
  - Avidity contributes (binding of first site facilitates binding of second)
- Reality potentially a combination of above. **How Significant an issue is it?**



## Sample Equilibrium Disruption: Time



Rate of Binding Reaction

$$d[sL]/dt = k_1[s][L] - k_2[sL]$$

[s] = Concentration of binding sites

[L] = Concentration of ligand

$k_1$  = association rate

$k_2$  = dissociation rate

If all free drug and ligand is removed expect exponential decay curve

$$[sL] = [sL]_0 e^{-k_2 t}$$

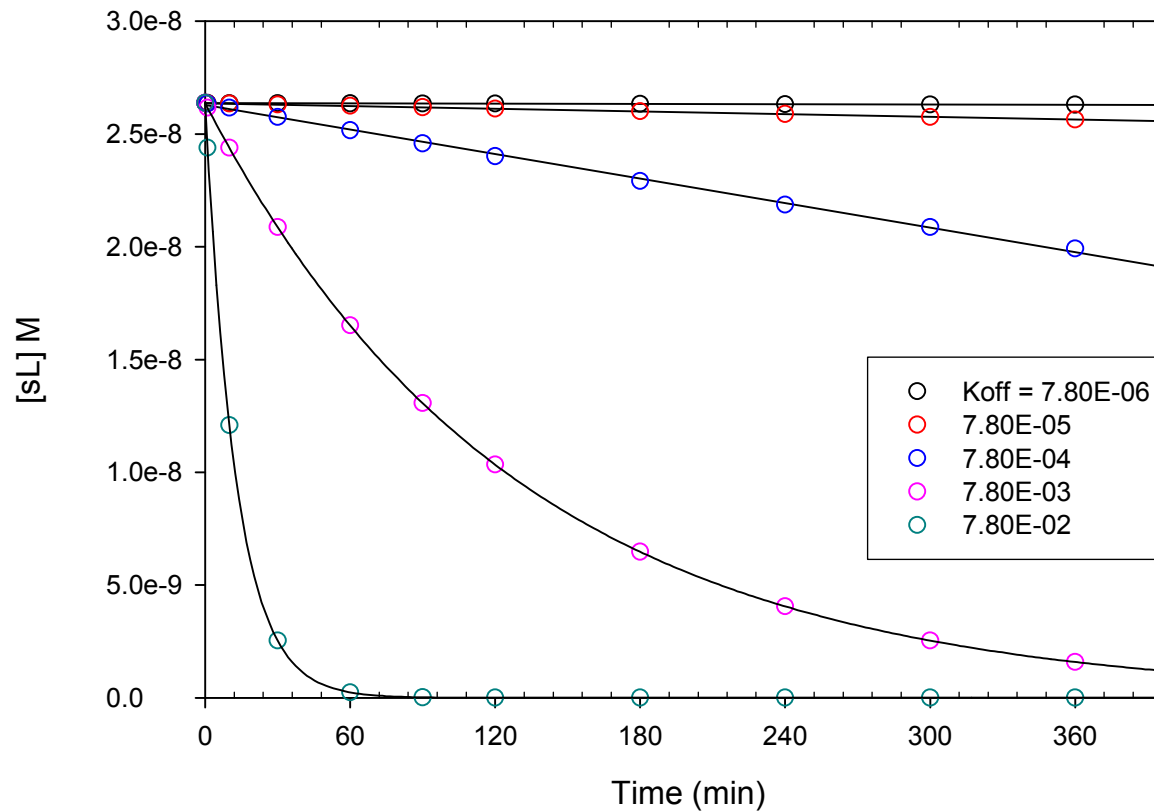
Can use this “worst case scenario” to simulate dilution and time effects on sample that started off at equilibrium ( $t=0$ )



# Simulating the effect of Time

PK Risk Assessment

Dissociation of Complex over time  
Off Rate Simulations



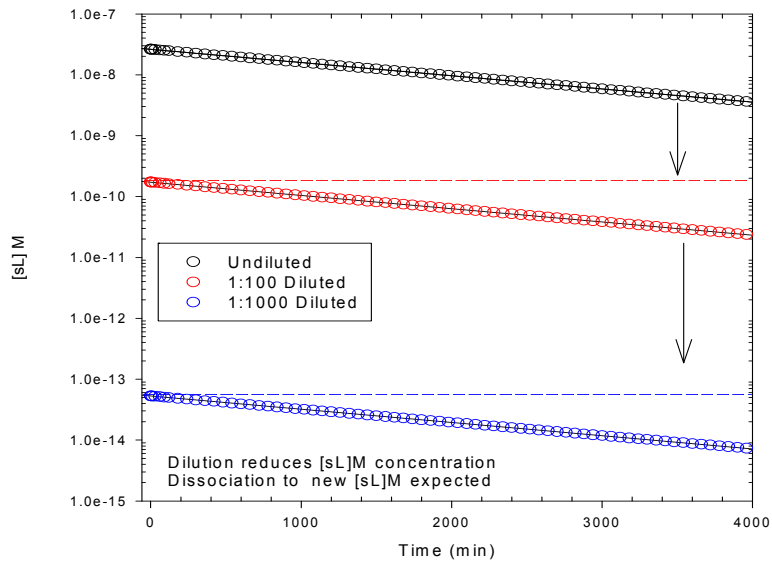
Molar Ratio =  
2.56

Drug Binding  
Sites to Target  
10 µg/mL drug

# Dilution Induced Complex Dissociation

Is it really an issue?

Effect of Dilution on Dissociation of Complex  
 $2 \times 10^4 \text{ sec}^{-1}$  Off Rate  
10 ug/mL Drug LMW target 100 ng/mL



PK Eg.

Experimental: Dilute Drug/Target Interference  
QC

1:5, 1:20, and 1:50

Test within 60 min

Results: No Change in observed result

Conclusion: Complex dissociation not a factor

Experimental data confirm simulation

# What you need to know to better assess risk

*Capture and Detection Ab/Reagents/ Drug-Target Kinetics*



## Assay Format Implications

[Target Ligand]:	10-50 pg/mL	1-50 ng/mL
Target Kinetics:	Unknown	Rapid clearance and synthesis

Structure:	Monomer	Dimer/Trimer/Aggregates
Drug -Target	KD : $\approx 10$ pM	KD $\approx 1$ nM
	On Rate : fast	On Rate : moderate
	Off Rate: slow	Off Rate: moderate

\*\*\*\*\*

Drug -Capture Ab KD :	?	KD $\approx$ nM
	On Rate : ?	On Rate : moderate
	Off Rate: ?	Off Rate: ?

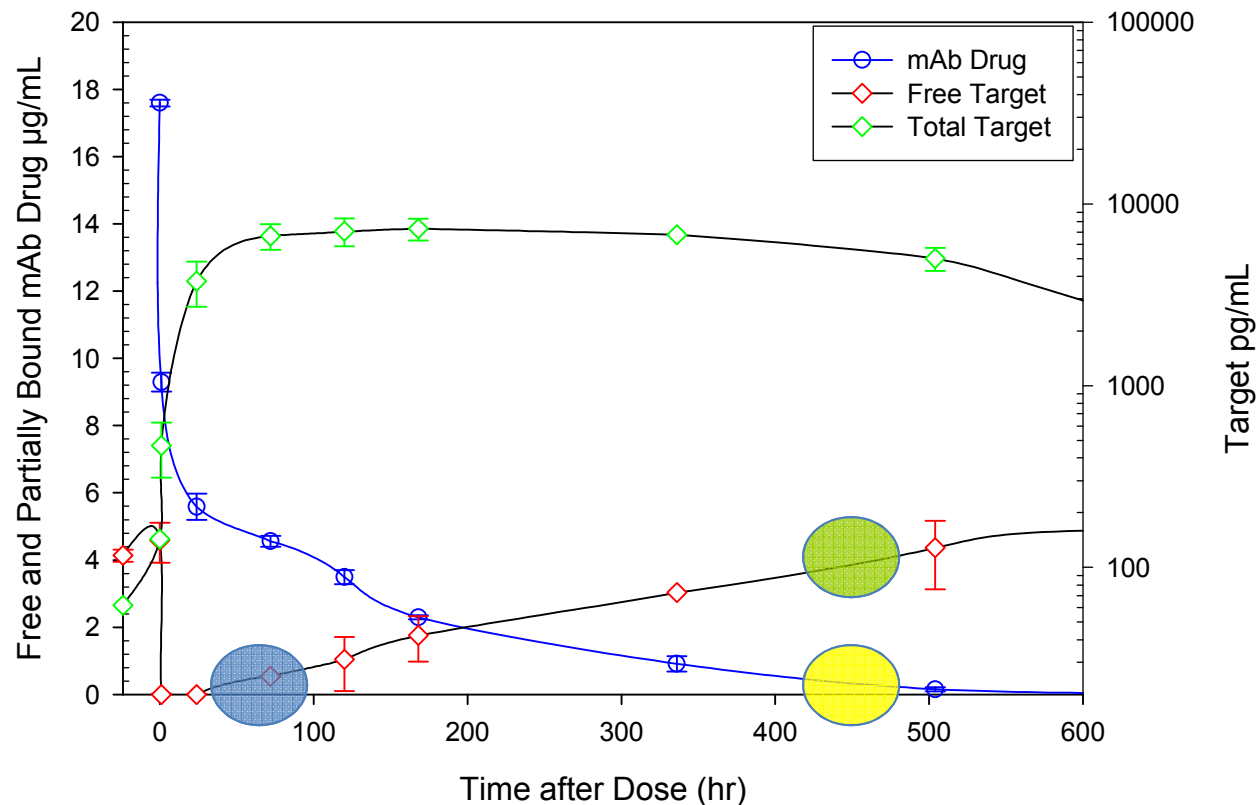
Target-Capture Ab	KD : nM	KD $\approx$ pM
	On Rate : medium	On Rate : moderate
	Off Rate: medium	Off Rate: slow



# When should we be concerned?

PK/PD Efficacy study

Dose vs Time Plasma Levels



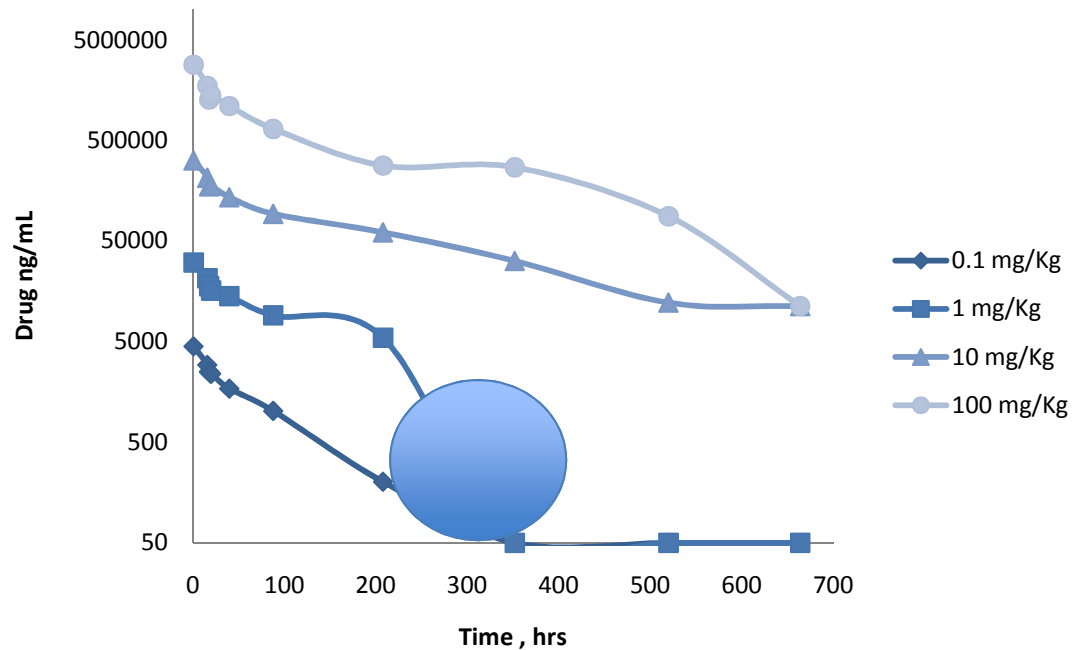
What are the risks ?

- 1) *Lack of sensitivity to measure free target after dosing*
- 2) *Under recover free drug. Target interference*
- 3) *Over recovery of free target*



# PK

## Target interference; when does it matter?



Can target interference alter PK parameters in a meaningful way?  
TMDD, Target Interference or ADA?

# Conclusions

Target /drug interference when does it matter?



Having raised awareness regarding Free and Total now need to find the right balance between

- asking for 5 assays and equilibrium disruption phobia
- doing nothing to assess either issue

When there is a potential for target/drug interference it needs to be experimentally defined so it can be managed

If observed total target level reaches a level that has been shown to interfere in the PK assay then the impact can be evaluated.



# Conclusions



Bioanalytical scientists can have a significant impact on programs by helping the team understand this issue

- It is better to select assays which can generate the best data and define edges than to attempt to measure multiple forms of analyte with less confidence.
- Risk of not assessing interference is potential for a lack of confidence in data, no matter what the cause, that can lead to significant additional bioanalytical investment to define this risk.
- Sample handling (dilution/ time) and assay format can contribute to equilibrium disruption and result in an under or over estimation of Free analyte **but** important not to overstate this risk
- Knowledge of binding kinetics will help put this issue in perspective and simulation is a useful tool to illustrate this for the team

