

Handling Anti-Drug Antibody (ADA) Data for Efficient Analysis

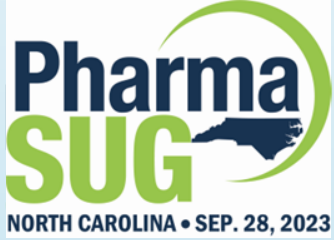
a presentation by
Sabarinath Sundaram



Meet the Speaker



Sabarinath Sundaram has over 10 years of statistical programming experience working in exploratory research studies to Phase III studies, CDISC standards, handling PK/PD/ADA data, and across multiple therapeutic areas. He has a Ph.D. degree in Life Sciences (Biochemistry) and is the Principal Statistical Programmer at **Seagen, Inc.** as well as PK/PD center of excellence lead.



Handling Anti-Drug Antibody (ADA) Data for Efficient Analysis

DISCLAIMER

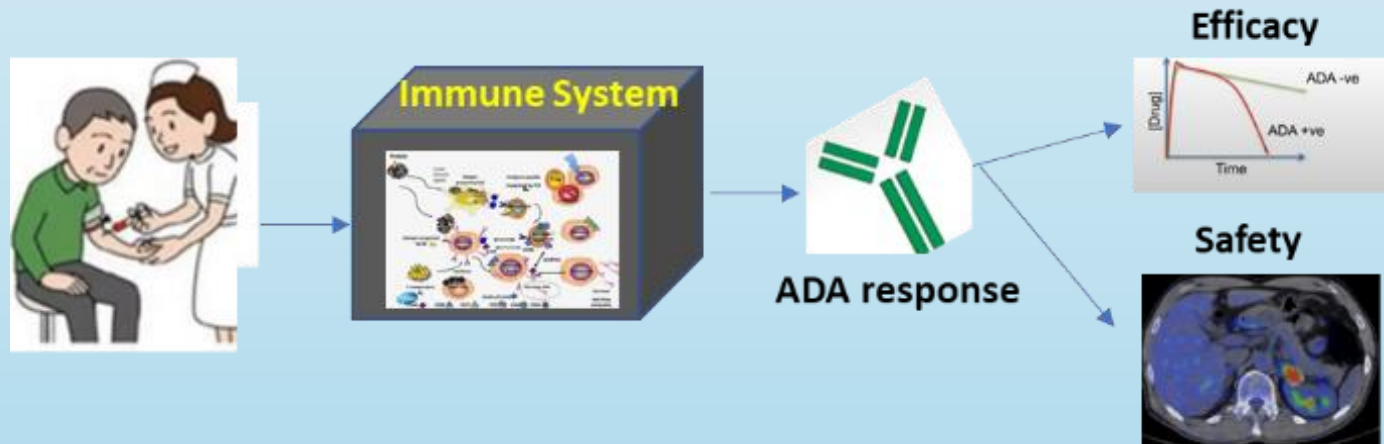
The opinions expressed in this presentation and on the following slides are solely those of the presenters and not necessarily those of Seagen

Agenda

- ▶ Introduction
 - ADA Testing Scheme
- ▶ Mapping Raw ADA data into SDTM IS Domain
 - Sample ADA DTP Template
 - Sample SDTM IS Domain Dataset
- ▶ Analysis Strategy of ADA Results
 - Sample ADaM ADIS Domain Dataset Key parameters & Summary of Anti-Drug Antibody (ADA) Incidence.
- ▶ Conclusion

Overview of Immunogenicity

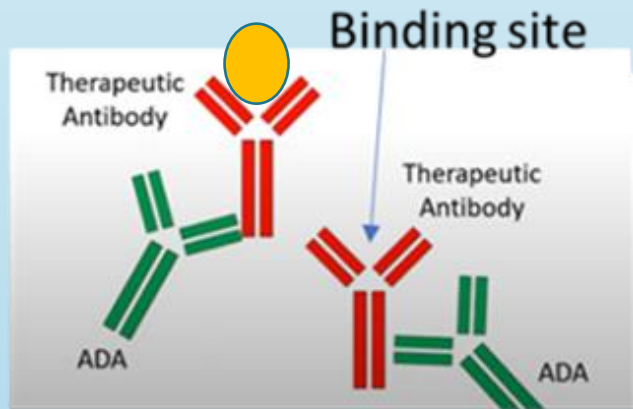
- Large molecules have revolutionized the pharmaceutical industry



- Pharmacokinetics (PK) is how the body responds to a drug.
- Pharmacodynamics (PD) allows us to quantify the relationship between a drug and its pharmacologic or toxicologic effect it has on patients.
- ADA responses can affect the PK, PD, safety, and efficacy of a therapeutic candidate. The clinical effects of ADA formation can be highly variable and may cause adverse events that put the patient at risk.

Interaction of Therapeutic with ADA

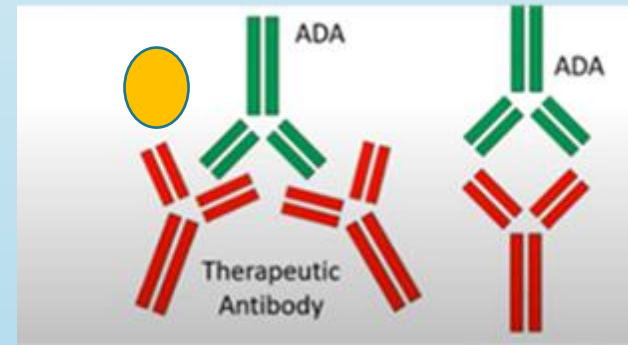
- Two Types, based on Interaction .



(a) Non-neutralizing ADA

Does not affect the binding of the Therapeutic(drug) and its target

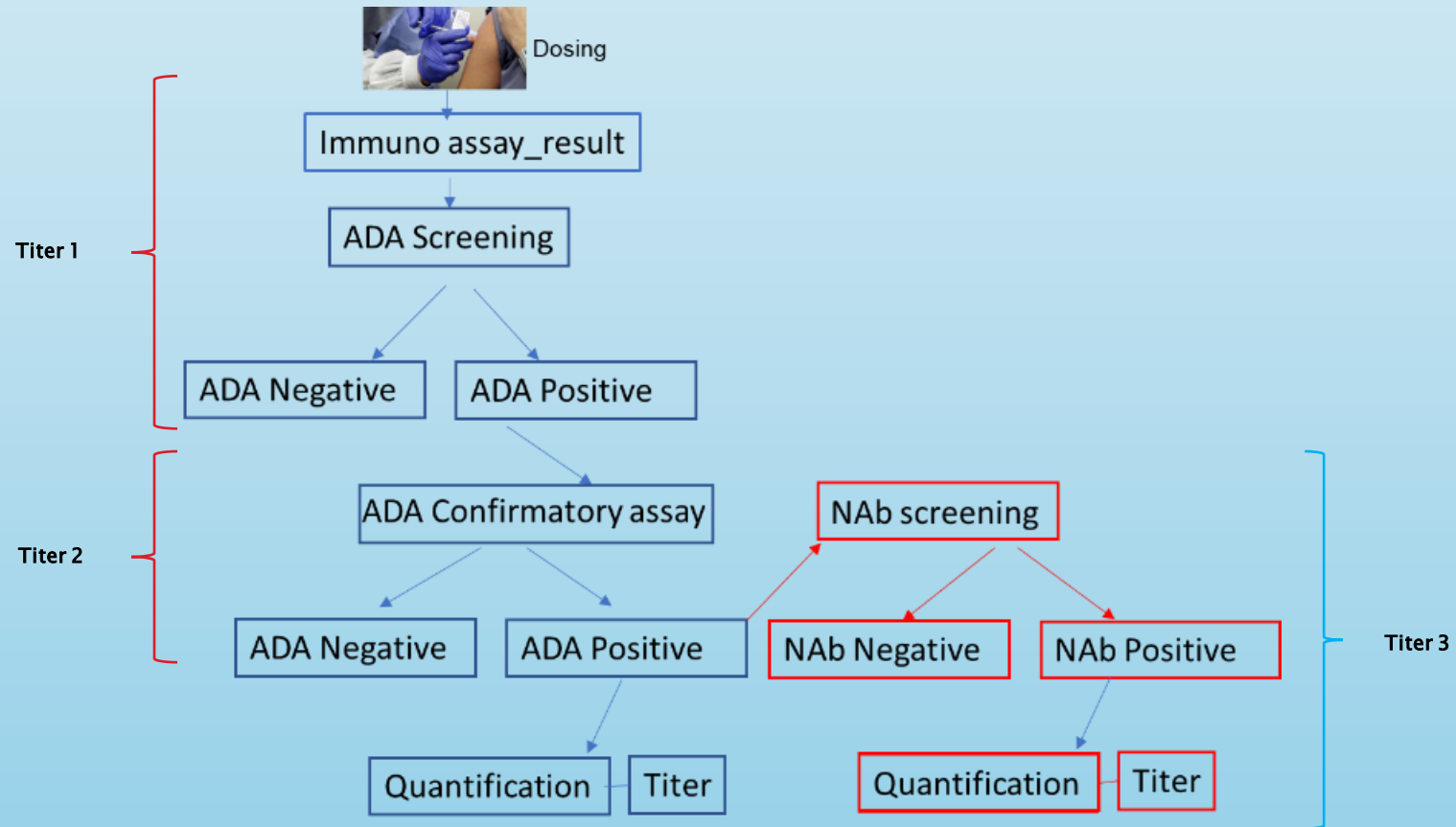
Target

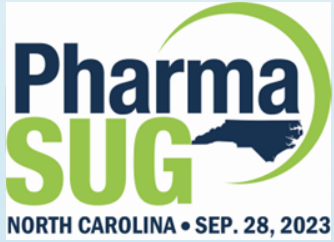


(b) Neutralizing ADA

Prevent interaction between the Therapeutic(drug) and its target

Multi-tiered ADA Testing Scheme





Sample ADA DTP Template & Results

DTP

LBTEST	Description	ISCAT	ISTESTCD	ISTEST	ISORRES	ISORRESU
ADA_Screening	Screening for binding ADA	Binding ADA	ADASCRN	ADA Screening	Positive/Negative	
ADA_Confirm	Confirmation of Binding ADA detection	Binding ADA	ADACONFM	ADA Confirmation	Positive/Negative	
ADA_Quantitative	Quantification of Binding ADA	Binding ADA	ADAQUAN	ADA Quantity	23	RLU
ADA_Titer	Titer ADA	Binding ADA	ADATITER	ADA Titer	<1	dilution
NAB_Screening	Screening for neutralizing ADA	NEUTRALIZING ADA	NABSCRN	NAB ADA Screening	Positive/Negative	
NAB_Confirm	Confirmation of neutralizing ADA detection	NEUTRALIZING ADA	NABCONFM	NAB ADA Confirmation	Positive/Negative	
NAB_Quantitative	Quantification of neutralizing ADA	NEUTRALIZING ADA	NABQUAN	NAB ADA Quantity	23	RLU
NAB_Titer	NAB Titer	NEUTRALIZING ADA	NABTITER	NAB ADA Titer	3	dilution

Raw data

USUBJID	VISIT	LBPT	LBCAT	LBSCAT	SCRQL	T1 SCRQN	SCRQNU	T2 CONFQL	CONFQNU	T3 TITER	TiterU
XXXX-1000-100	CIDI	Predose	immunogenicity	ADA	Negative	xxx.x	RLU				
XXXX-1000-100	CIDI	8 hrs post dose	immunogenicity	ADA	Negative	xxx.x	RLU				
XXXX-1000-101	CIDI	Predose	immunogenicity	ADA	Negative	xxx.x	RLU				
XXXX-1000-101	CIDI	8 hrs post dose	immunogenicity	ADA	potential Positive	xxx.x	RLU	Negative			
XXXX-1000-102	CIDI	Predose	immunogenicity	ADA	Positive	xxx.x	RLU				
XXXX-1000-102	CIDI	8 hrs post dose	immunogenicity	ADA	potential Positive	xxx.x	RLU	Postive	xxx.x	1	dilution

Programming Strategies: SDTM

- ▶ Historically mapped to PC or LB.
- ▶ CDISC guidelines: Mapped to IS (Immunogenicity Specimen Assessments) domain
- ▶ Findings domain: assessments that determine whether a therapy induced an immune response
- ▶ Data structure: One record per test per time point per visit per subject

SDTM Domain IS

DOMAIN	USUBJID	ISSEQ	ISTEST	ISTESTCD	ISTSTOPO	ISORRES	VISIT	ISTPT	ISDTC
IS	1000-100	1	Binding Antidrug Antibody	ADA_BAB	Screen	Negative	Cycle 1 Day 1	Predose	2019-07-08T00:53
IS	1000-100	2	Binding Antidrug Antibody	ADA_BAB	Screen	Negative	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-101	1	Binding Antidrug Antibody	ADA_BAB	Screen	Negative	Cycle 1 Day 1	Predose	2019-04-30T04:39
IS	1000-101	2	Binding Antidrug Antibody	ADA_BAB	Screen	Positive	Cycle 1 Day 1	8 hrs post dose	2019-04-30T12:40
IS	1000-101	3	Binding Antidrug Antibody	ADA_BAB	Confirm	Negative	Cycle 1 Day 1	8 hrs post dose	2019-04-30T12:40
IS	1000-102	1	Binding Antidrug Antibody	ADA_BAB	Screen	Positive	Cycle 1 Day 1	Predose	2019-07-08T00:54
IS	1000-102	2	Binding Antidrug Antibody	ADA_BAB	Screen	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	3	Binding Antidrug Antibody	ADA_BAB	Confirm	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	4	Binding Antidrug Antibody	ADA_BAB	Quantity	234	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	5	Binding Antidrug Antibody	ADA_BAB	Titer	4	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	6	Neutralizing Binding Antidr	ADA_NAB	Screening	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	7	Neutralizing Binding Antidr	ADA_NAB	Confirm	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	8	Neutralizing Binding Antidr	ADA_NAB	Quantity	100	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
IS	1000-102	9	Neutralizing Binding Antidr	ADA_NAB	Titer	2	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54

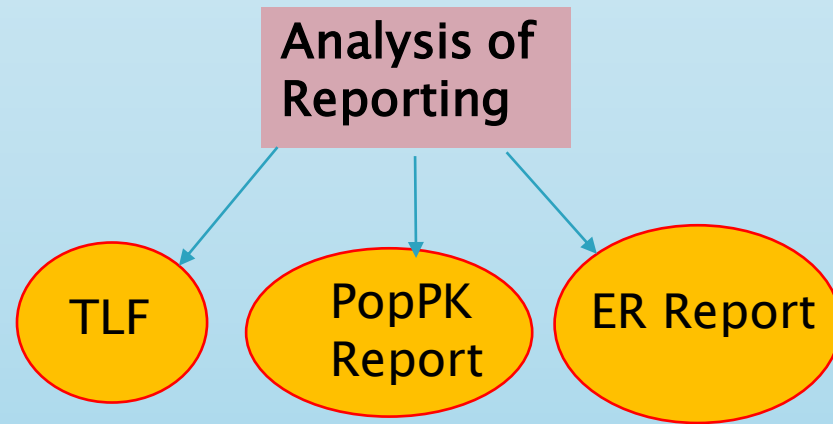
Programming Strategies: ADaM

- ▶ ADIS
- ▶ Class: Basic Data Structure (BDS)
- ▶ One record per analysis parameter per timepoint per visit per subject
- ▶ Additional records - new derived parameters based on ADA results to support the analysis
- ▶ ADSL flag variables for summarization of ADA data:
 - ADA-evaluable flag (ADAFL)
 - NAb evaluable flag (NABFL)

ADaM Domain ADIS

USUBJID	ASEQ	PARCAT	PARAMCD	PARAM	AVALC	AVISIT	ATPT	ADT
XXXX-1000-100	1	Binding ADA	ADASCRN	ADA Screening	Negative	Cycle 1 Day 1	predose	2019-07-08T00:54
XXXX-1000-100	2	Binding ADA	ADASCRN	ADA Screening	Negative	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-101	1	Binding ADA	ADASCRN	ADA Screening	Negative	Cycle 1 Day 1	predose	2019-04-30T04:40
XXXX-1000-101	2	Binding ADA	ADASCRN	ADA Screening	Positive	Cycle 1 Day 1	8 hrs post dose	2019-04-30T12:40
XXXX-1000-101	3	Binding ADA	ADACONFM	ADA Confirm	Negative	Cycle 1 Day 1	8 hrs post dose	2019-04-30T12:40
XXXX-1000-102	1	Binding ADA	ADASCRN	ADA Screening	Positive	Cycle 1 Day 1	Predose	2019-07-08T00:54
XXXX-1000-102	2	Binding ADA	ADASCRN	ADA Screening	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	3	Binding ADA	ADACONFM	ADA Confirm	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	4	Binding ADA	ADAQUAN	ADA Quantity (RLU)	234	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	5	Binding ADA	ADATITER	ADA Titer	4	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	6	Neutraizing ADA	ADASCRN	NAb ADA Screening	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	7	Neutraizing ADA	ADACONFM	NAb Confirm	Positive	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	8	Neutraizing ADA	NABQUAN	NAb Quantity (RLU)	100	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	9	Neutraizing ADA	NABTITER	NAb Titer	2	Cycle 1 Day 1	8 hrs post dose	2019-07-08T08:54
XXXX-1000-102	10	Binding ADA	INDUCADA	Treatment induced ADA	N			
XXXX-1000-102	11	Binding ADA	NOTRTREL	No Treatment related ADA	N			
XXXX-1000-102	12	Binding ADA	TIMEADA	Time to onset of First ADA	42			
XXXX-1000-102	13	Binding ADA	POSTADA	Post baseline ADA	Y			
XXXX-1000-102	14	Binding ADA	LVADA	Last visit ADA	Cycle 3			

Analysis Strategy of ADA Results



Key analysis parameters

- 1) **Treatment-induced ADA:** Incidence of ADA induction is determined if both baseline and post-baseline samples have positive titers, and the subject is only considered positive for the induction of ADA when the post-baseline titer is at least 4-fold greater than the titer prior to initial dosing.
- 2) **No Treatment-related ADA:** Subjects with negative post-baseline ADA results
- 3) **Post-baseline Positive ADA:** Incidence of ADA is determined by counting at least 1 positive result at any timepoint after initial dosing.
- 4) **NAb Negative:** NAb negative at baseline includes patients who are ADA-negative.
- 5) **NAb Incidence:** defined as having at least one positive NAb result at any time, including baseline and/or post baseline.
- 6) **Time to Onset of First ADA:** Select the analysis day of the first record with ADA-positive result after dosing.
- 7) **Last Visit ADA:** The latest visit on which a subject had a positive ADA result after dosing.
- 8) **Persistently Positive ADA:** Defined as ATA positive at 2 or more post-baseline assessments (with \geq 16 weeks between first and last positive) or ATA positive at the last post-baseline assessment.
- 9) **Transiently Positive ADA:** Defined as having at least one post-baseline ATA positive assessment and not fulfilling the conditions of ATA persistently positive.

Sample Summary of Anti-Drug Antibody (ADA) Incidence

	Total (N=xx) n (%)
Subjects with a baseline and at least one post-baseline sample	n
Baseline Negative	n
Negative post-baseline ²	m (%)
Positive post-baseline ³	m (%)
Baseline Positive	n
Negative post-baseline ²	m (%)
Treatment induced ADA ¹	m (%)
NAb negative ⁸ at baseline	n
NAb incidence ⁹	m (%)
Total positive post-baseline	m (%)

The percentage denominators are the first row counts of evaluable subjects, not the analysis set population. Superscript numbers refer to the derived parameters explained in the above section.

Conclusion

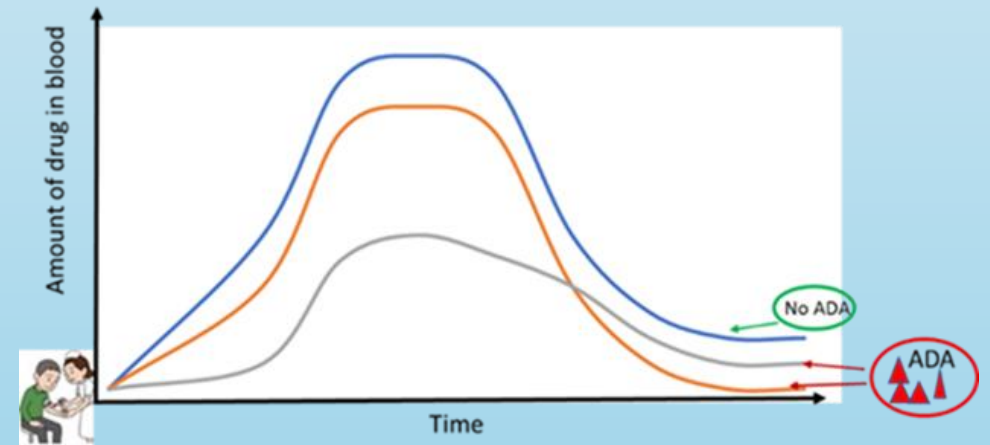
- ❖ Use of ADA Results
- ❖ Explain PK results
- ❖ Clinical Observations (Loss of Efficacy and AE/SAE)
- ❖ Few Important Clinical Development–Stage Questions

Early-stage Development

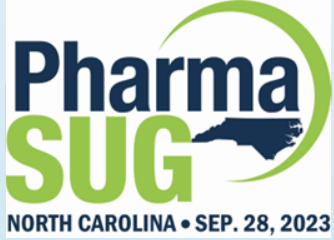
- Do any pre-existing x-reactive antibodies increase after dosing
- Is ADA observed increase or decrease after repeated dosing
- Can any mitigation strategies likely decrease or increase risk profile

Late-stage Development

- Will a dosing regimen, change in manufacturing or Patient population change in immunogenicity profile



Drug concentration by time. Model showing the effect of ADA on PK plot.



Acknowledgement

I would like to thank Johnny Maruthavanan, Shang-Ying Liang, Christine O Day, Daping Zhang, Bala Pitchuka, Shefalica Chand and Michiel Hagendoorn for their feedback, constant support, and guidance.

Thank you for your attention !



AUTHOR DETAILS:

NAME:

Sabarinath Sundaram

AFFILIATION:

Seagen Inc.

E-MAIL:

ssundaram@seagen.com

LINKEDIN:

<https://www.linkedin.com/in/sabari-sundaram-975548b4/>