

MIDD: VACCINE R&D GETS A SHOT IN THE ARM FROM PHARMACOMETRICS

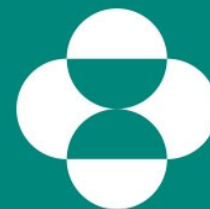
(A.K.A. “VACCINES ARE NOT IMMUNE TO THE CHARMS OF MODELING AND SIMULATION”)

ROSA MIDD Webinar

January 22, 2020

Jeff Sachs

Pharmacokinetics, Pharmacodynamics, and Drug Metabolism
– Quantitative Pharmacology and Pharmacometrics,
Merck & Co., Inc., Kenilworth, NJ, USA



MSD

INVENTING FOR LIFE

Outline

Motivation

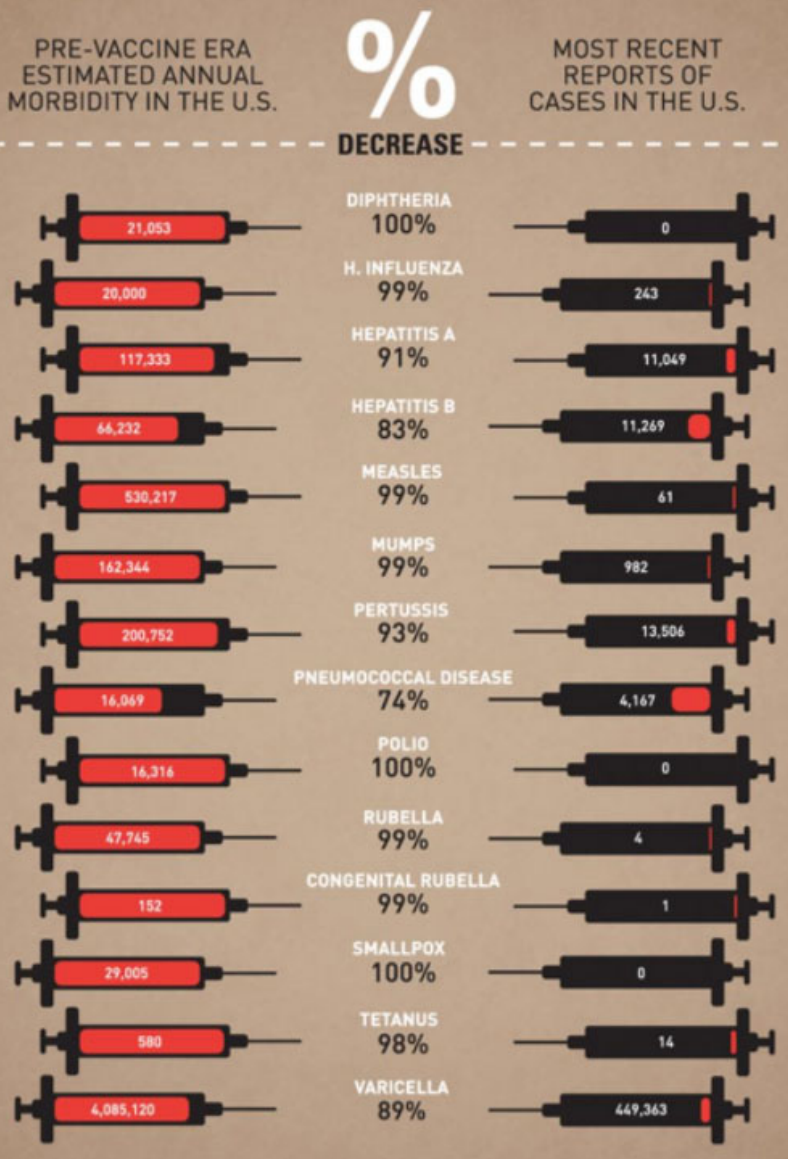
- Motivation – Impact of vaccines
- Potential impact for PMX

Background

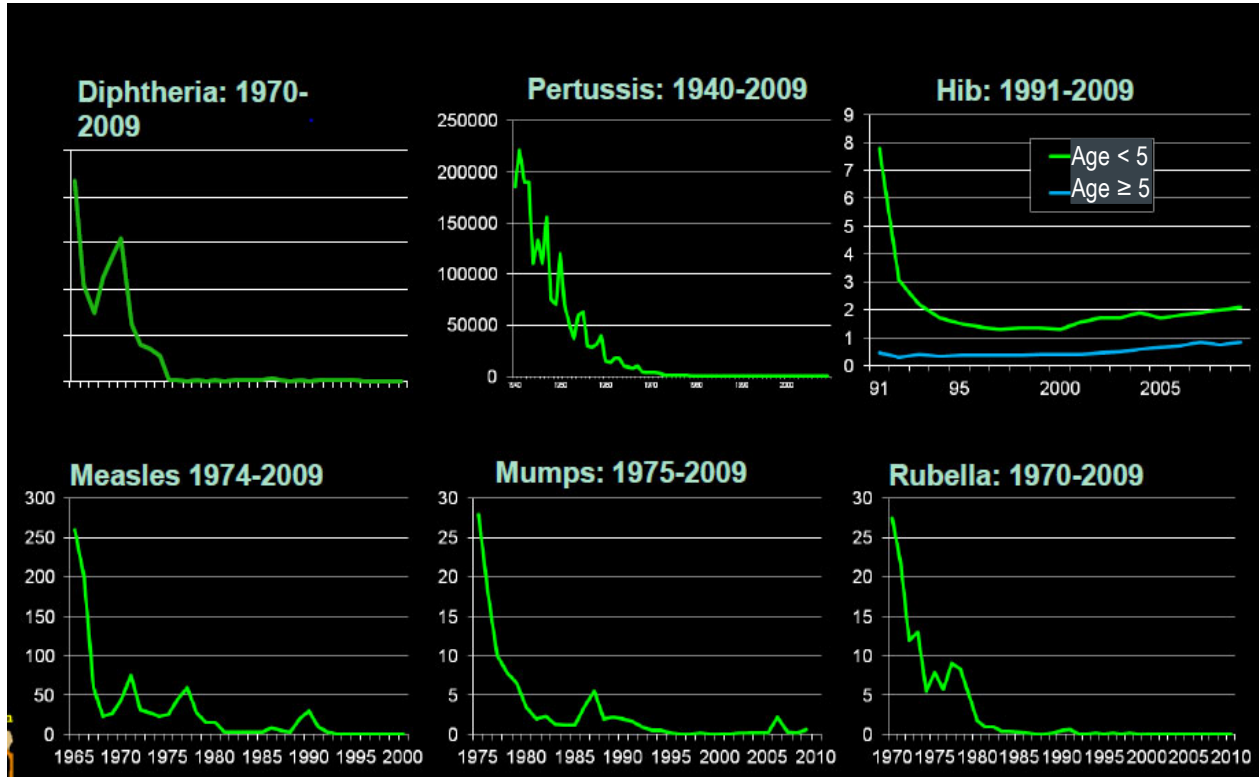
- Vaccines & Immunology

PMX

- Applications
- Their impact



Many Diseases Have Been Prevented



<https://www.forbes.com/sites/matthewherper/2013/02/19/a-graphic-that-drives-home-how-vaccines-have-changed-our-world/-1a58e6dd3302>

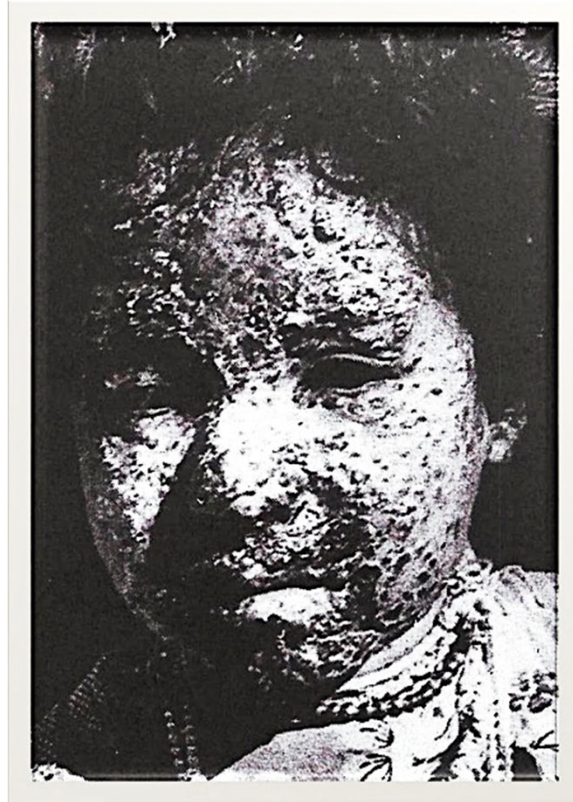
Halsey, NA, *How Vaccines Cause Adverse Events*, ADVAC Course Ancey France 2018



Polio



Historical Perspective: Smallpox



Bazin, H., *Vaccination: a History*

Don't Count Your Children Until The Measles Have Come Through
– African saying



*MMWR / November 11, 2016, 65 (44) 1228-1233

The Modern Toll of Measles



¹ European Region statistics. From “Measles in Europe: record number of both sick and immunized,” WHO Regional Office for Europe, Copenhagen, 7 February 2019

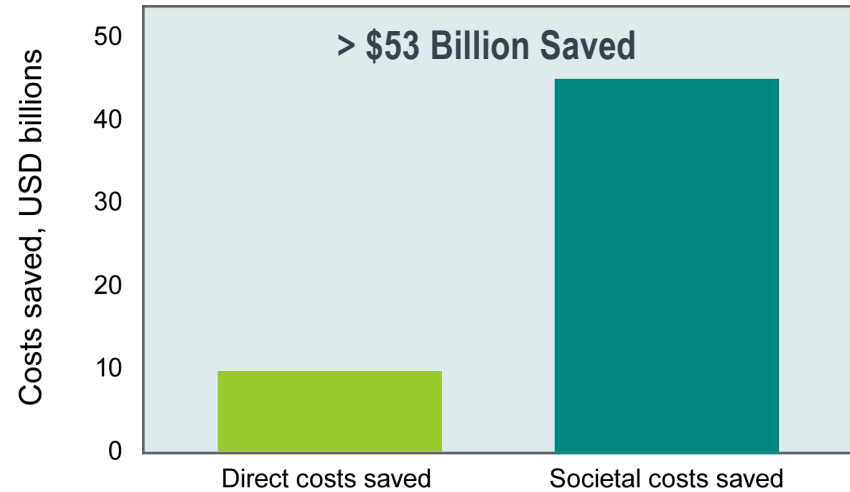
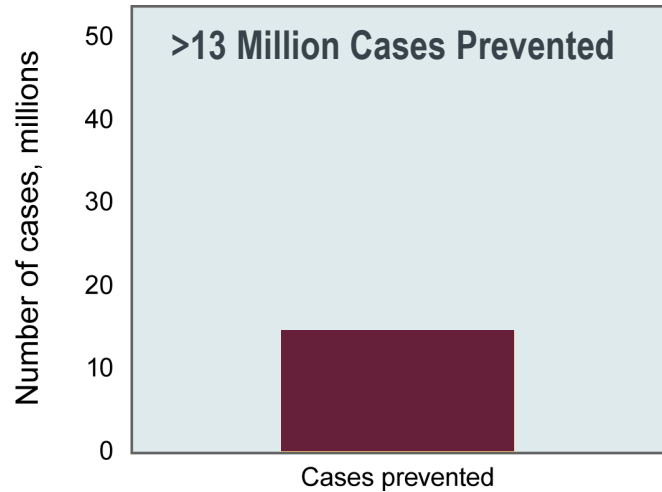
² <http://www.euro.who.int/en/media-centre/sections/press-releases/2019/over-100-000-people-sick-with-measles-in-14-months-with-measles-cases-at-an-alarming-level-in-the-european-region,-who-scales-up-response>

³ <https://www.npr.org/2019/05/19/724747890/measles-outbreak-in-the-philippines>

Health and Economic Impact of Preventing Disease with Vaccination...

Just One Country: USA

Just One Year's Cohort: Children born in 2001

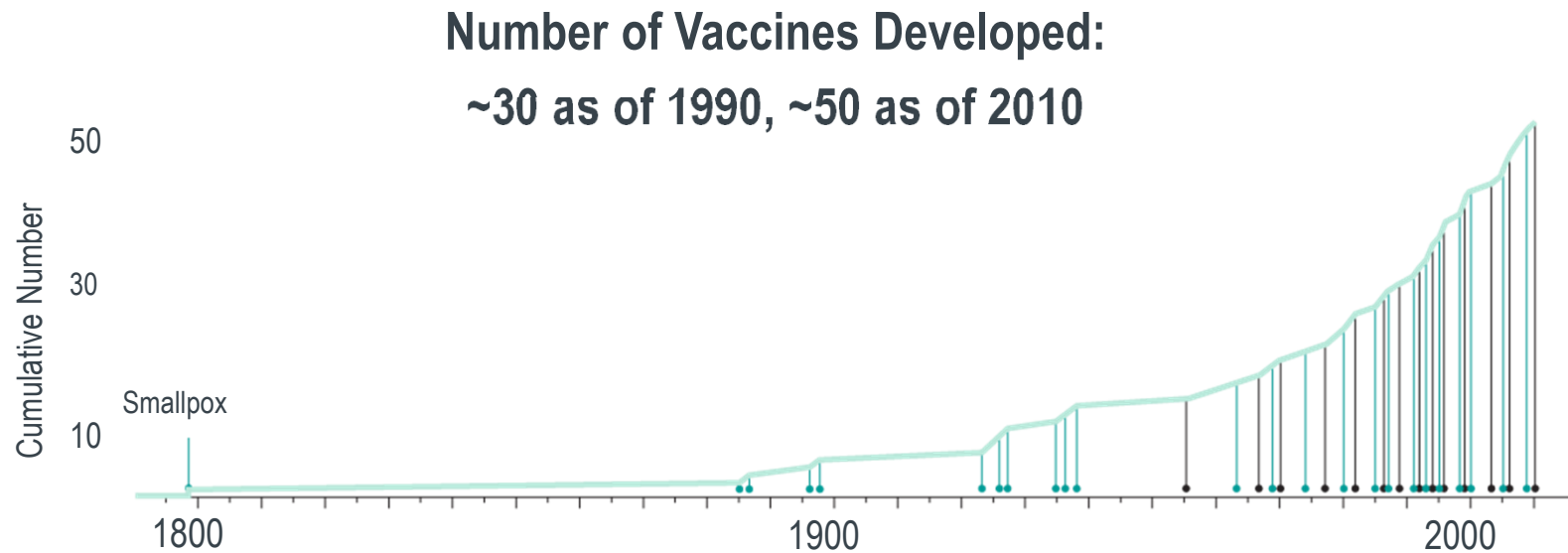


Adapted from: Zhou F et al. Arch Pediatr Adolesc Med. 2005;159:1136-1144.

All costs are given in US dollars (USD).

Direct program costs included vaccines, administration, parent travel, and direct costs for the management of adverse events. Societal costs included direct program costs and parent time lost for vaccination and the management of adverse events.

About 50 Vaccines Developed to Date

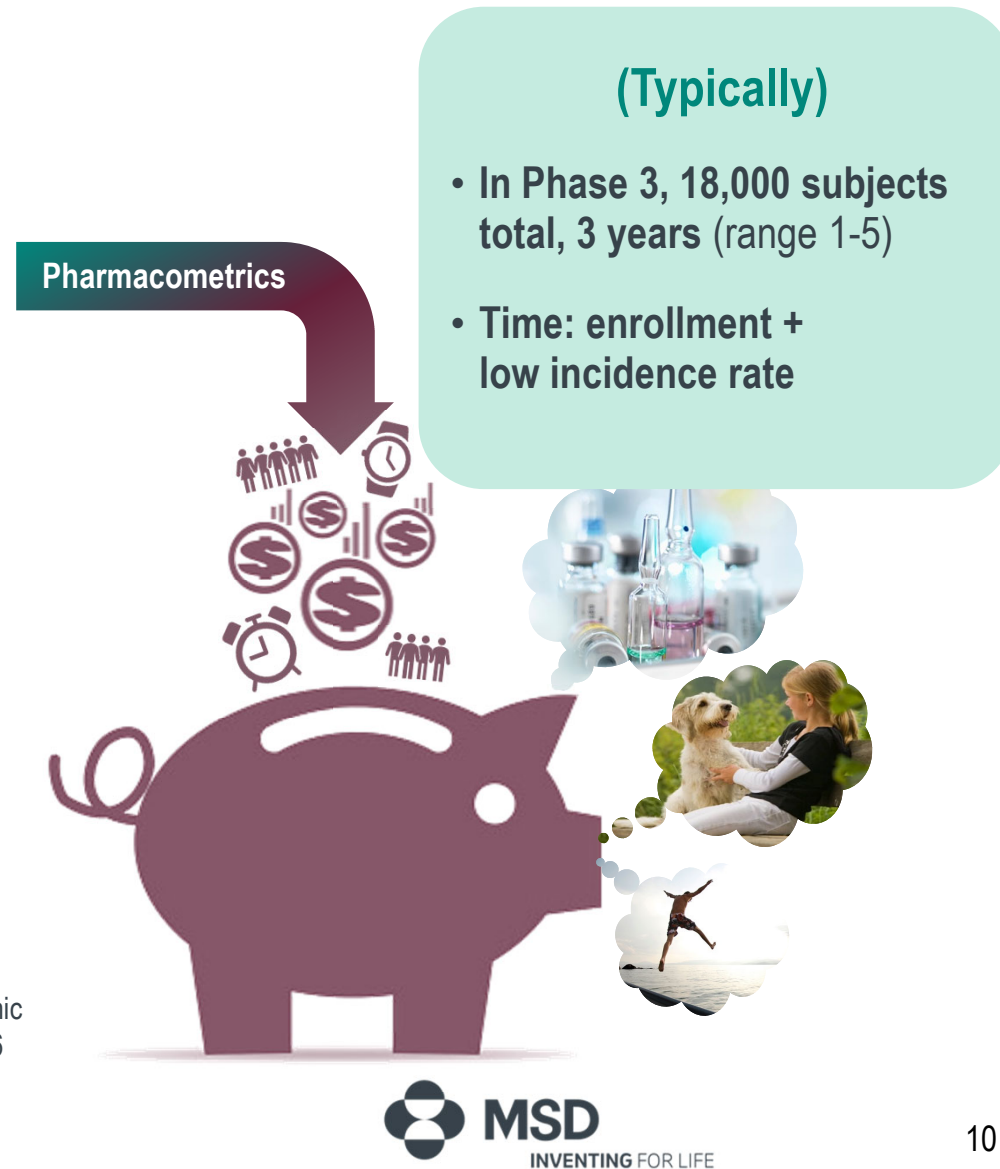


Adapted from: IOM (Institute of Medicine), Ranking vaccines: A prioritization framework: Phase I: Demonstration of concept and a software blueprint. Washington, DC, *The National Academies Press*, 2012, p. 19.

...But Expensive and Takes Too Long

- **Cost** of a vaccine from discovery through Ph. 2a: **\$0.4 Billion** (range \$0.1-1B)*
- **Time** for a vaccine from discovery through Ph. 2a: **7 years** (range 4-15 years)*
- Vaccines too often in development for ~20 years

*Gouglas, D., TT Le, et al., Estimating the cost of vaccine development against epidemic infectious diseases: a cost minimisation study, Lancet Glob. Health, 2018;6:e1386-96



“Vaccine”

MSD



Vaccine (for today): Active Stimulator Of Immune Memory and Antibody Production for Prevention of an Infectious Disease.

~~Therapeutic~~

~~Chemo-Prophylaxis~~



~~Passive Immunization~~

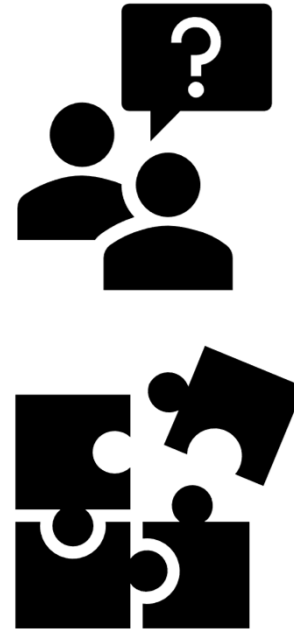
~~Seasonal Flu~~

Jeryl Lynn Hilleman with her sister, Kirsten, in 1966 getting the mumps vaccine developed by their father.

What is Special About Vaccines and Pharmacometrics?

Why were Vaccines not on our radar??

- PK* Rare
 - Little DDI* (concomitant vaccination)
 - Traditional clinical pharmacology analyses not typical
 - except safety & Tox
- not part of our traditional purview.



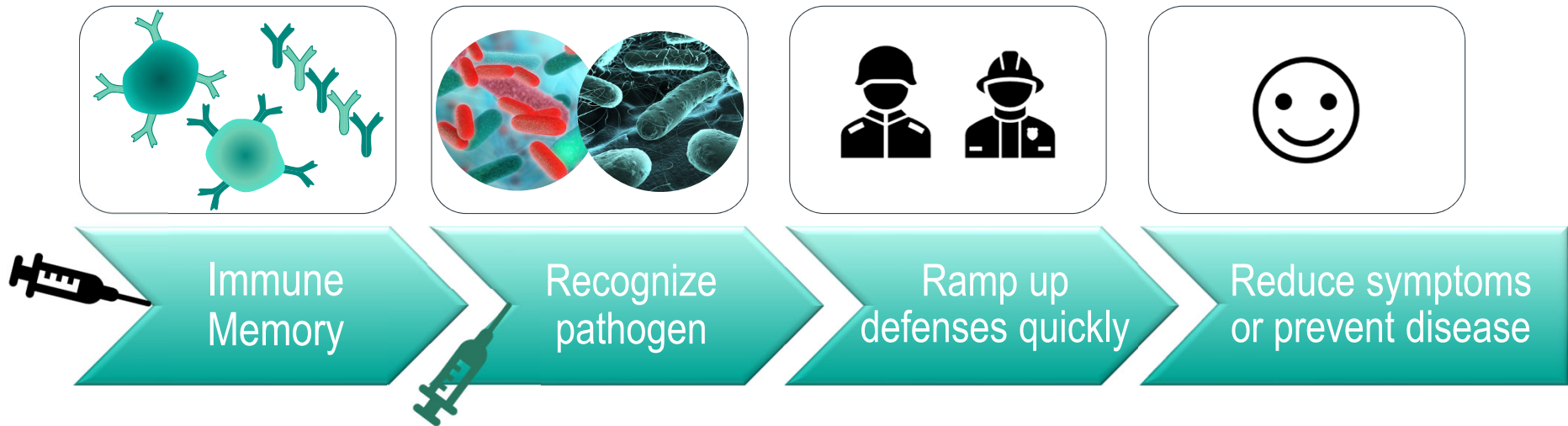
*PK: pharmacokinetics, DDI: drug-drug interaction

Very
The ^vBASICS:
VACCINES and IMMUNOLOGY

MSD



Active Immunization – How it works



- Measuring Immune response: “Titer” ~ Target engagement
 - Quantity and quality of antibodies
- More is better

Immunogenicity \neq efficacy

Overview of PMX and Vax

MSD



Modeling and Simulation in Vaccines

Computational vaccinology:



~~Epidemiology
Health Econ~~

~~PKPD
of mAb
("passive
immu-
nization")~~

~~immuno-
genicity
= $F(\text{antigen
sequence})$~~

~~chem eng. &
SYS BIO for
bioprocess~~

~~QSP/PKPD in
Chemo-
prophylaxis~~

**Vaccine
Pharmacometrics**

(Today's focus)

Rich History of Published Work (not a complete list!)

KEYSTONE SYMPOSIA
on Molecular and Cellular Biology
Accelerating Life Science Discovery

Login/ Create an Account Search meetings by keyword

Conferences Financial Aid Support Us About Us FAQ

Meeting Program
Meeting Summary
Abstract Information
Registration Information

Systems Immunology: From Molecular Networks to Human Biology (A1)
Scientific Organizers: Ronald N. Germain, Aviv Regev, Nir Hacohen and Dana Pe'er
January 10–14, 2016
Big Sky Resort, Big Sky, Montana, USA

al. (2015) Mathematical modeling provides details of the human immune response to infection. *Front Cell Infect Microbiol.* 4:177

Gómez-Mantilla JD, et. al. ADME processes in vaccines and PK/PD approaches for vaccination optimization. In: ADME and tPK/PD. Wiley; 2016.

Rhodes SJ, et al. (2016) The TB vaccine H56 + IC31 dose-

LONDON SCHOOL OF HYGIENE & TROPICAL MEDICINE

Vaccine IS/ID Modelling Consortium

www.lshtm.ac.uk/research/centres-projects-groups/isid#welcome

Welcome About IS/ID Modelling Members Research Publications Discussion Forum Sign up

Vaccine IS/ID Modelling Consortium

Using model-based drug development methods (PK/PD) to accelerate vaccine dose decision making.

Modelling dose responses following vaccination: lessons from the PK/PD field

Rockville, Maryland, May 29, 2014

Organizers:

- BMGF: Steve Kern
- LSHTM: Richard White, Sophie Rhodes
- Imperial College: Gwen Knight
- AERAS: Tom Evans, Lewis Schragar

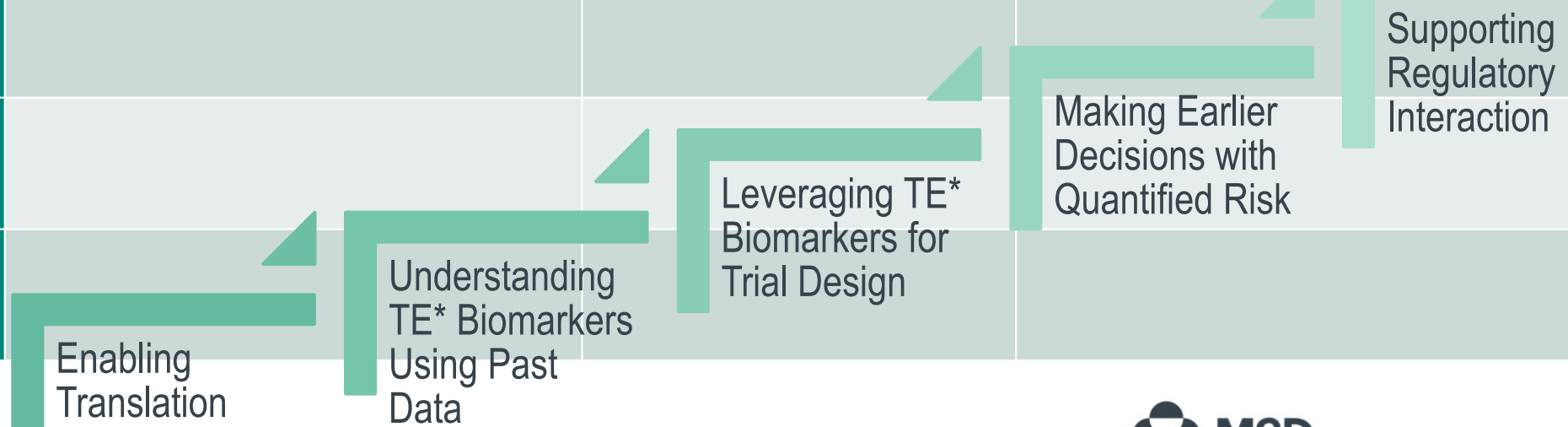
EXAMPLES

MSD



Phase	Key Question	Method(s)	Decision(s) / Impact
-------	--------------	-----------	----------------------

Seven Examples
 Capability Demonstration
 Program Impact



*TE: Target Engagement (stimulating protective immune response)

Phase	Key Question	Method(s)	Decision(s) / Impact
1/2	<p>What N (# subjects) will let us tell if vaccines A and B are different?</p>	<p>Phenomenological model Clinical trial simulation</p>	<p>Trial design, program strategy for sequence of trials</p>
<div style="background-color: #cccccc; border-radius: 20px; padding: 40px; display: inline-block;"> <p>Wrong Question!</p> </div>			

What can we learn about dose-level & formulation impact on immunogenicity?
How can we use past data to inform the trial design?
How can we integrate data across trials in the future?

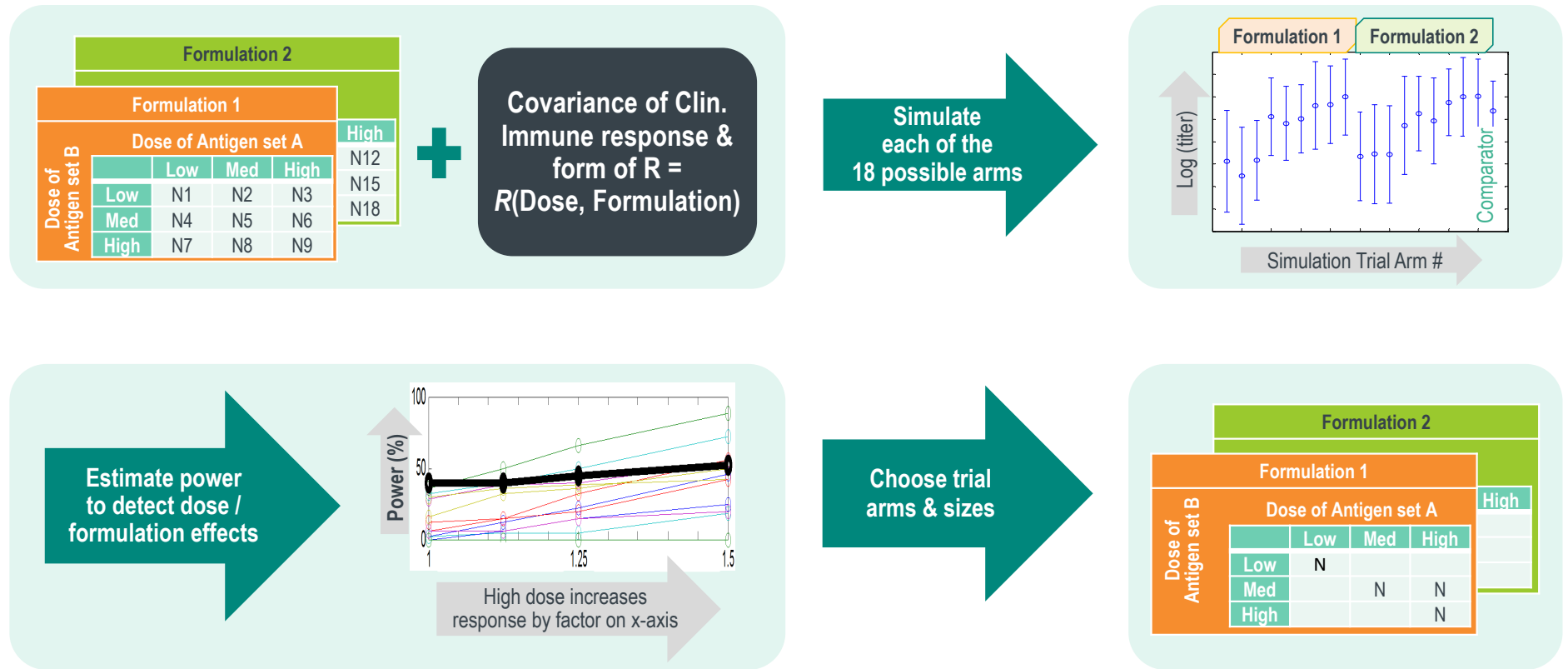
How many arms (and which ones) were needed to address information desired in the first question?

~400 Million

Number of arms the team had planned:

3

Trial Design by *Phenomenological Simulation*



Thanks: Kapil Mayawala, Jon Hartzel

Trial Design by *Phenomenological* Simulation

Impact:

- changed from 2-arm “yes-no” study to 5-arms (same total # subjects), optimized to learn key properties
- Helped plan:
 - (1) next studies, and
 - (2) how to integrate data across studies

Thanks: Kapil Mayawala, Jon Hartzel

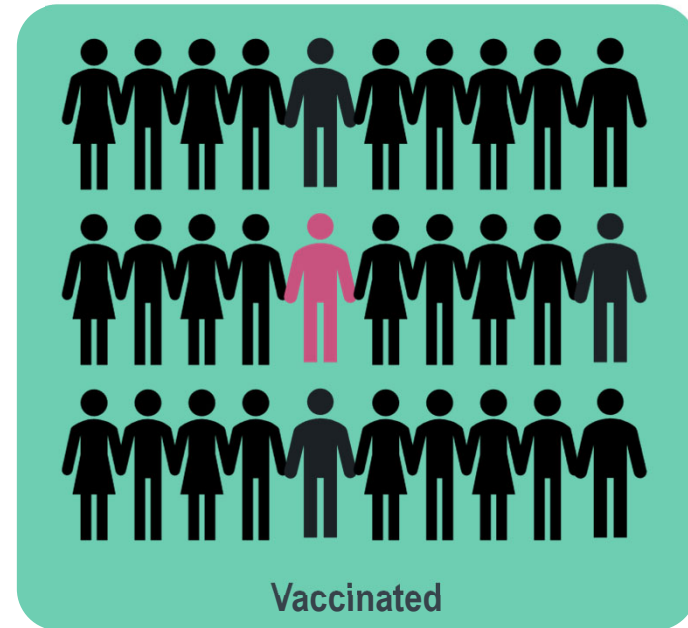
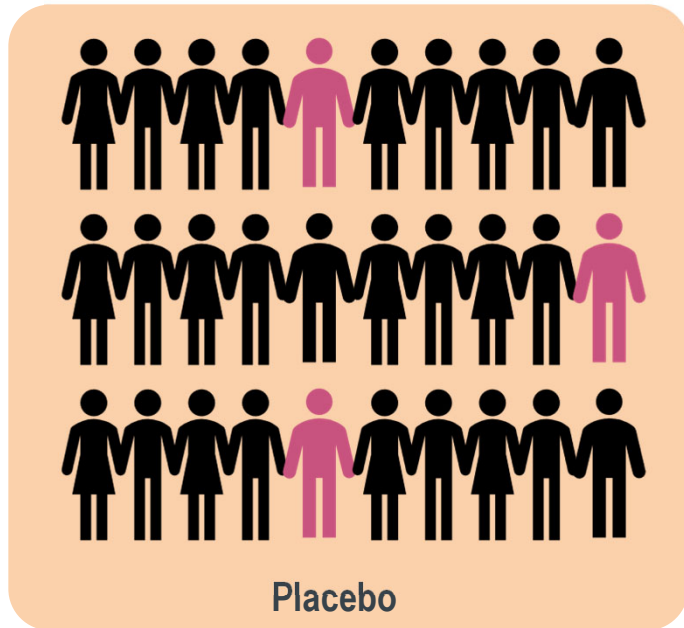
Phase	Key Question	Method(s)	Decision(s) / Impact
2	What disease assay(s) should we use and how often?	QSP and Bayesian probabilistic	Saved \$, increased POS Choice of assay, frequency

- Measuring vaccine efficacy requires counting number of disease cases.
- What happens if the counting (assay) process is not perfect?
- What assays should we use and how often?

Efficacy = Proportional Risk Reduction

10% of placebo subjects get sick,
3% vaccinated subjects get sick

$$\text{Efficacy} = (10\% - 3\%) / 10\% = 0.7 = \text{“70\% efficacy”}$$



Efficacy = Proportional Risk Reduction

10% of placebo subjects get sick,
3% vaccinated subjects get sick

$$\text{Efficacy} = (10\% - 3\%) / 10\% = 0.7 = \text{“70\% efficacy”}$$



Typically need tens/hundred cases (Ph. 2/3, resp.)
→ No efficacy information until Ph. 2b/3



Placebo



Vaccinated

Efficacy = Proportional Risk Reduction

13% ~~10%~~ of placebo subjects get sick,

6% ~~3%~~ vaccinated subjects get sick

~~Efficacy = (10% - 3%) / 10% = 0.7 = "70% efficacy"~~

Efficacy = (13% - 6%) / 13% = 0.54 = "54% efficacy"

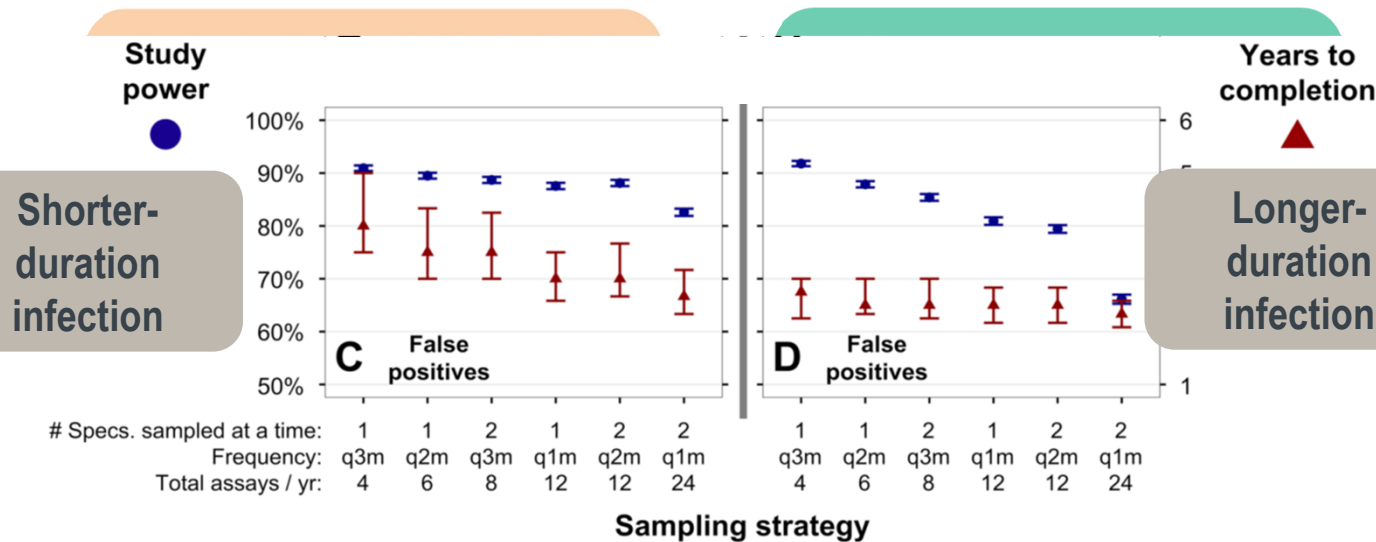
+3%
+3%

Impact of False Positives

Repeat testing → ☹️
Smart testing → 😊

+ Pathogen Dynamics

Ph. 2/3 Design
More assays:
Trade Time for power
Duration of detectability matters



Efficacy = Proportional Risk Reduction

Impact:

- Saved \$,
- increased POS (per subject)
- Choice of assay(s) and their frequency

# Specs. sampled	Frequency:	q3m	q2m	q3m	q1m	q2m	q1m	q3m	q2m	q3m	q1m	q2m	q1m
Total assays / yr:		4	6	8	12	12	24	4	6	8	12	12	24

Sampling strategy

Impact of False Positives

Repeat testing → ☹️
Smart testing → 😊

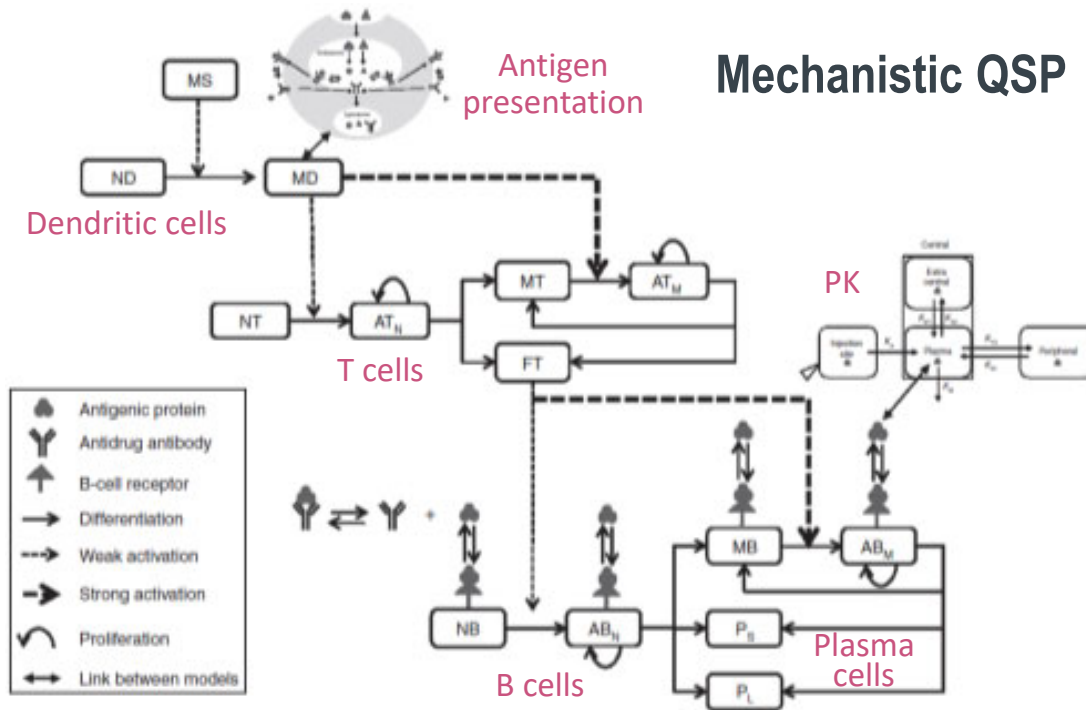
+ Pathogen Dynamics

Ph. 2/3 Design
More assays:
Trade Time for power
Duration of detectability matters

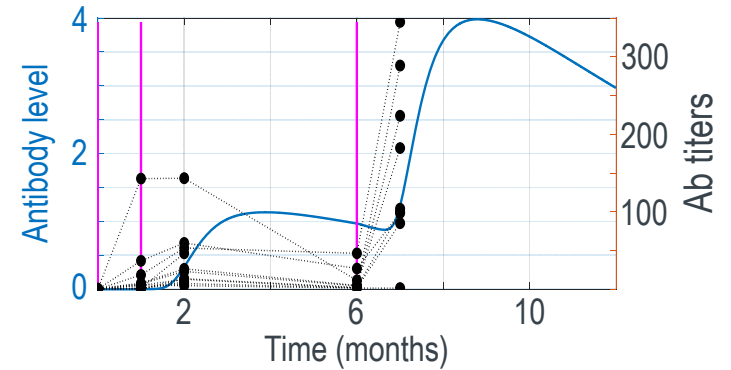
Phase	Key Question	Method(s)	Decision(s) / Impact
(na)	Can we model enough of the immune system to be predictive?	QSP	Capability development

How can we use preclinical data to help design vaccines and to predict the right dose-level or regimen?

Basic Model of Some Immune System Components



Chen et al., CPT PSP 2014



Thanks: Jeff Perley, Josiah Ryman

Basic Model of Some Immune System Components



Chen et al., CPT PSP 2014

Thanks: Jeff Perley, Josiah Ryman

Phase	Key Question	Method(s)	Decision(s) / Impact
4	How many doses of vaccine are needed to confer lasting protection?	QSP	Suggests single dose could provide protective immune memory Mechanistic insight

Can we leverage mechanistic information to help inform regimen?

Hepatitis B: Models for Antigen, Anti-viral Titer and Immune Memory

$$\frac{dV_i(t)}{dt} = -\sigma V_i(t)$$

Antigen

$$\frac{dM_i(t)}{dt} = (\gamma V_i(t) + \beta M_i(t)) \left(1 - \frac{M_i(t)}{N} \right)$$

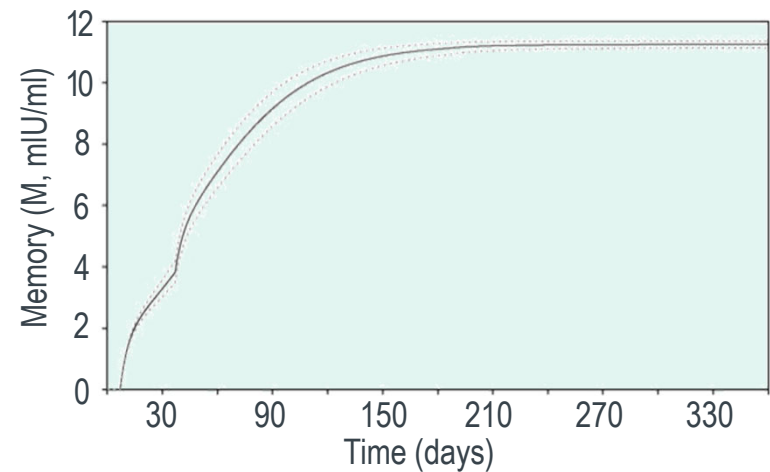
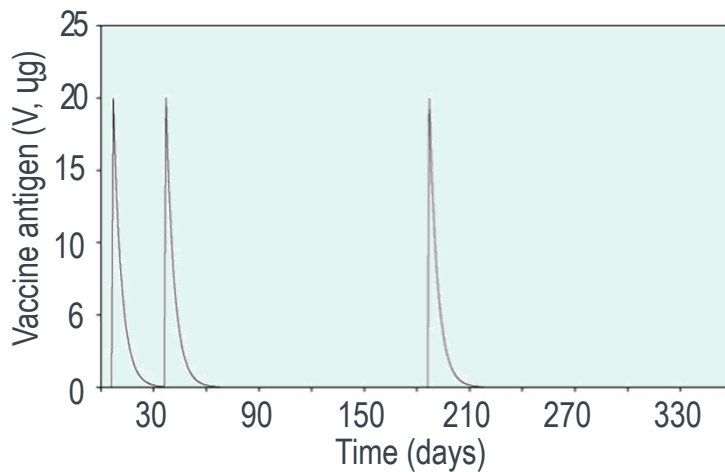
Immune memory

$$\frac{dA_i(t)}{dt} = \delta M_i(t) V_i(t) \left(1 - \frac{A_i(t)}{N} \right) - \frac{\mu A_i(t)}{T_i}$$

Anti-viral titer

- 10,815 anti-viral titres in 1,923 patients
- 2-4 vaccinations in 6-48 month period
- No Immune memory (M_i) or antigen (V_i) measurements

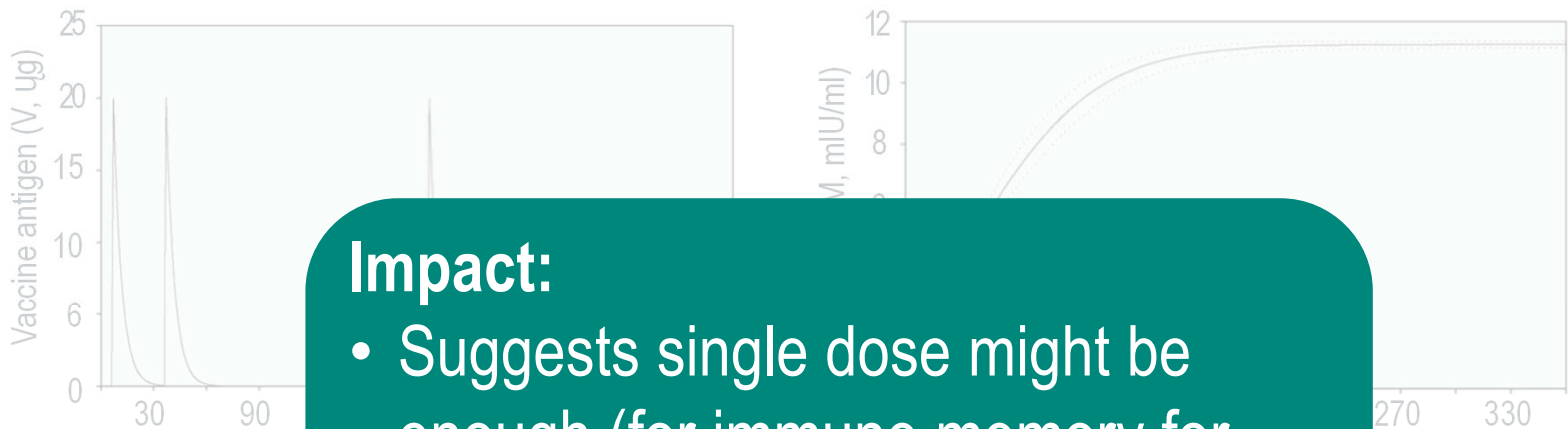
Simulation Results



Authors' Conclusions

- The model demonstrates **significant differences between different vaccines in** both the **time** taken to generate immune memory **and** the **amount of memory** generated.
- The model provides theoretical support for the hypothesis that a **single vaccine dose** can generate protective immune memory.

Simulation Results



Authors' Conclusions

- The model demonstrates that a single dose can generate protective immune memory, suggesting that the **time** taken for the immune response to develop is critical in both the model and in practice.
- The model provides theoretical support for the hypothesis that a **single vaccine dose** can generate protective immune memory.

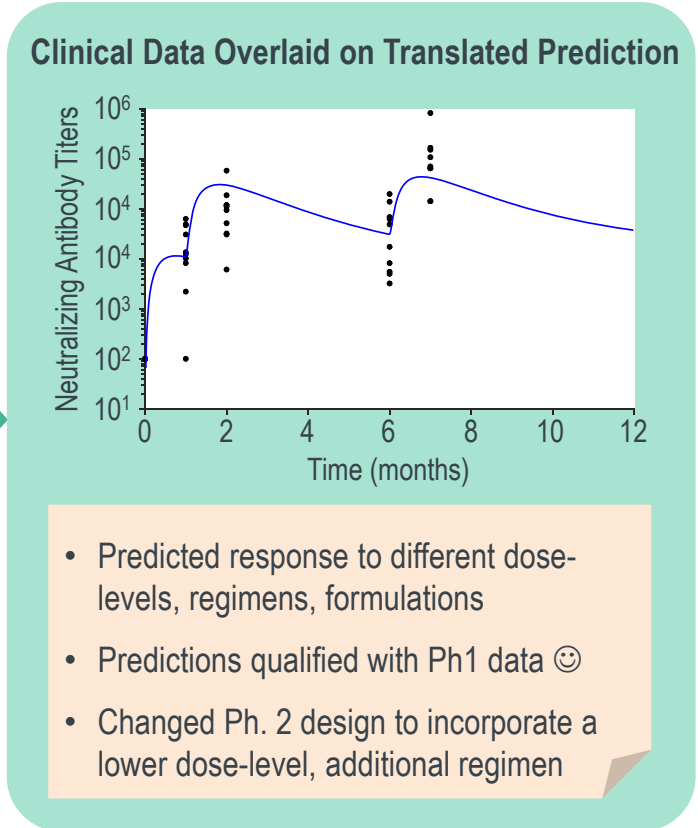
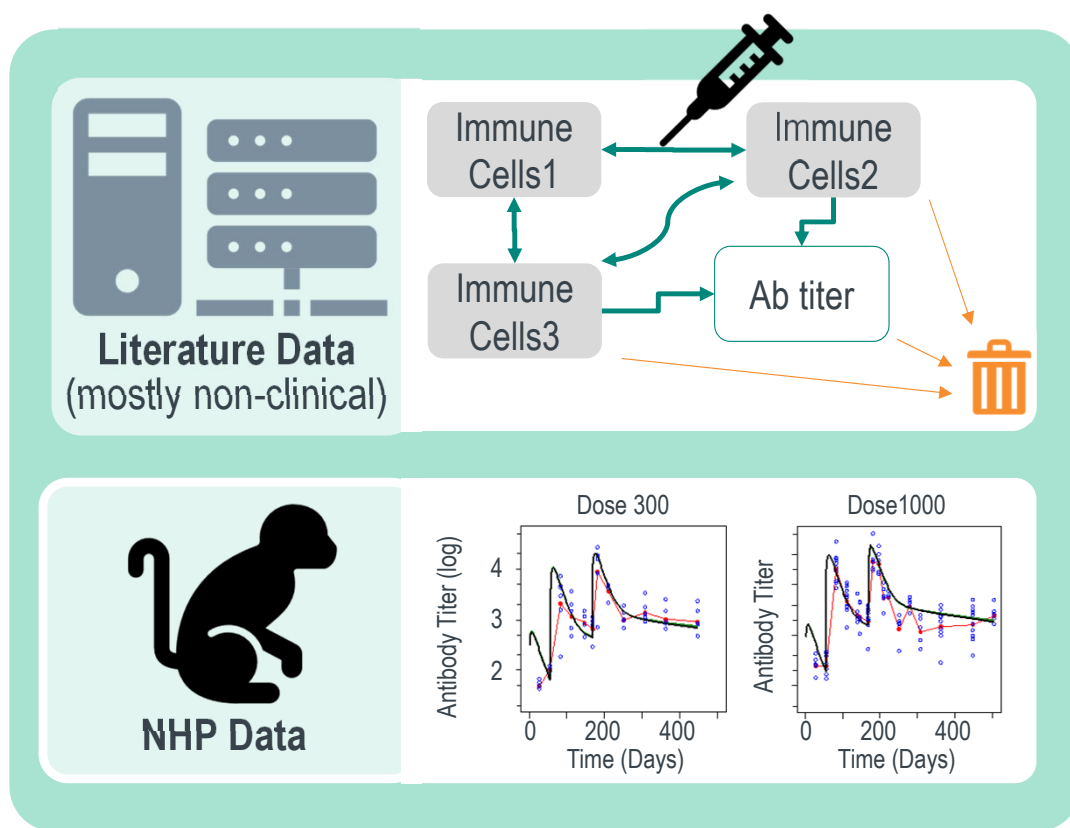
Impact:

- Suggests single dose might be enough (for immune memory for this pathogen and vaccine mechanism)
- Mechanistic insight

Can we leverage mechanistic information to help inform regimen in Ph. 2 trial?

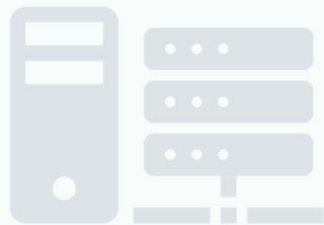
Phase	Key Question	Method(s)	Decision(s) / Impact
2	<p>Which regimens should be tested in Ph. 2 Trial? (Regimen: dose-level, # doses, timing)</p>	QSP	Ph. 2 Trial Design: add new dose level and different regimen

Trial Design by QSP



Thanks: Jeff Perley, Guido Jajamovich, Jos Lommerse (Certara), April Barbour

Trial Design by QSP



Literature Data
(mostly non-clinical)

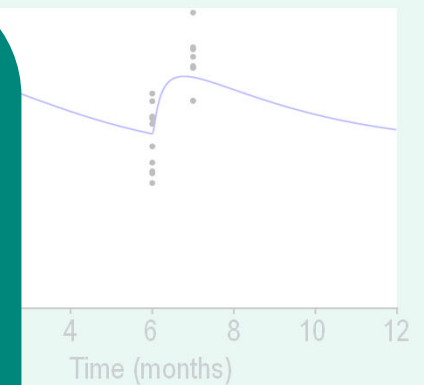


NHP Data

Impact:

- Increased confidence in ability to
 - model regimen-response
 - Translate from non-clinical species
- Changed planned Phase 2
 - dose-levels
 - number of doses

Clinical Data Overlaid on Translated Prediction



response to different dose-levels, formulations

qualified with Ph1 data 😊

n. 2 design to incorporate a dose-level, additional regimen

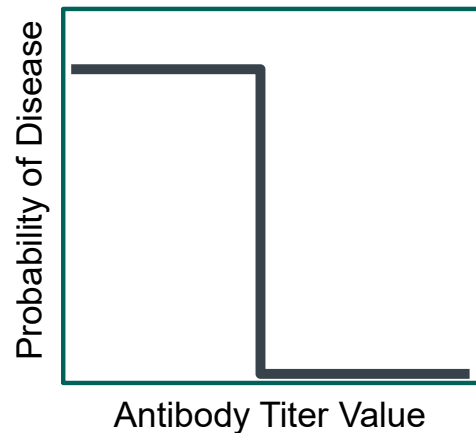
Thanks: Jeff Perley, Guido Jajamovich, Jos Lommerse (Certara), April Barbour

Phase	Key Question	Method(s)	Decision(s) / Impact
2,3	Do we have adequate evidence of efficacy if some pathogens have too few cases?	PoDBA (= Probability of Disease Bayesian Analysis)	Novel Ph. 3 endpoint GNG test criteria

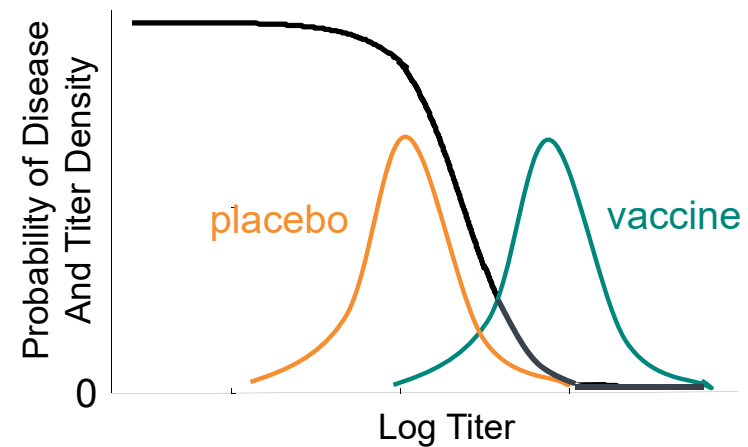
Can we increase POS by mitigating risk of a season with low incidence rate?

PoDBA Method

- Estimate relationship between probability of disease and antibody titer values based on titer values of subjects with and without disease



- Use this relationship and titer values of control and vaccine groups to estimate vaccine efficacy and its confidence interval



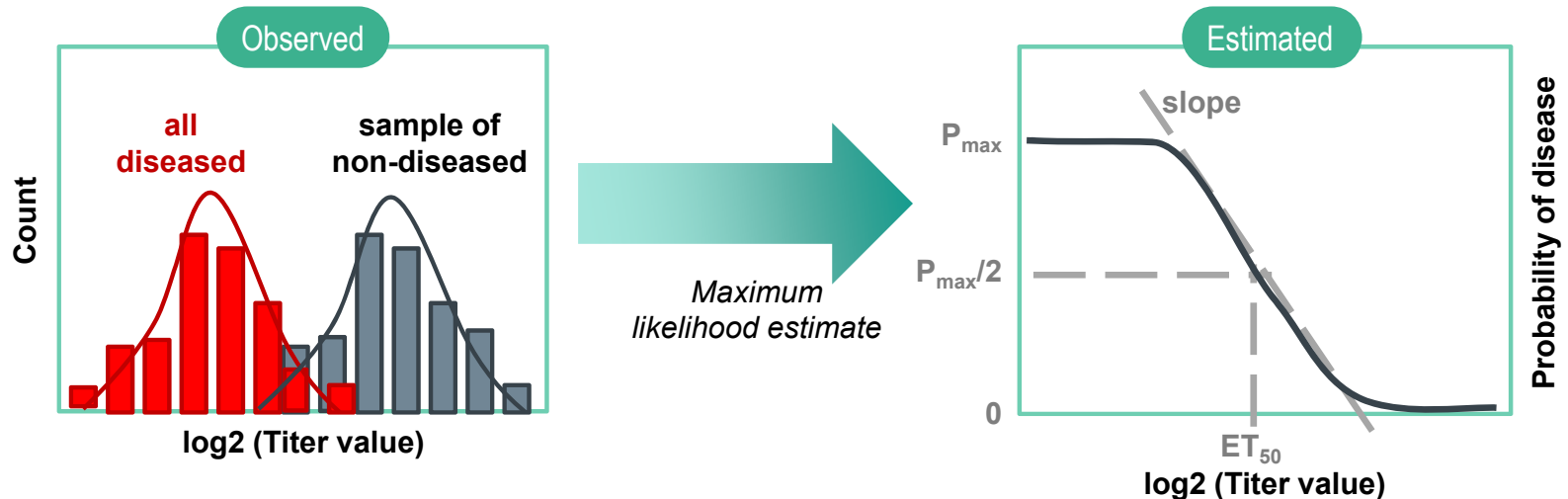
~~Wait for enough cases~~

~~Need many cases~~

$$\text{Efficacy} = 1 - \frac{\text{Expected PoD}(\text{vaccinated})}{\text{Expected PoD}(\text{placebo})}$$

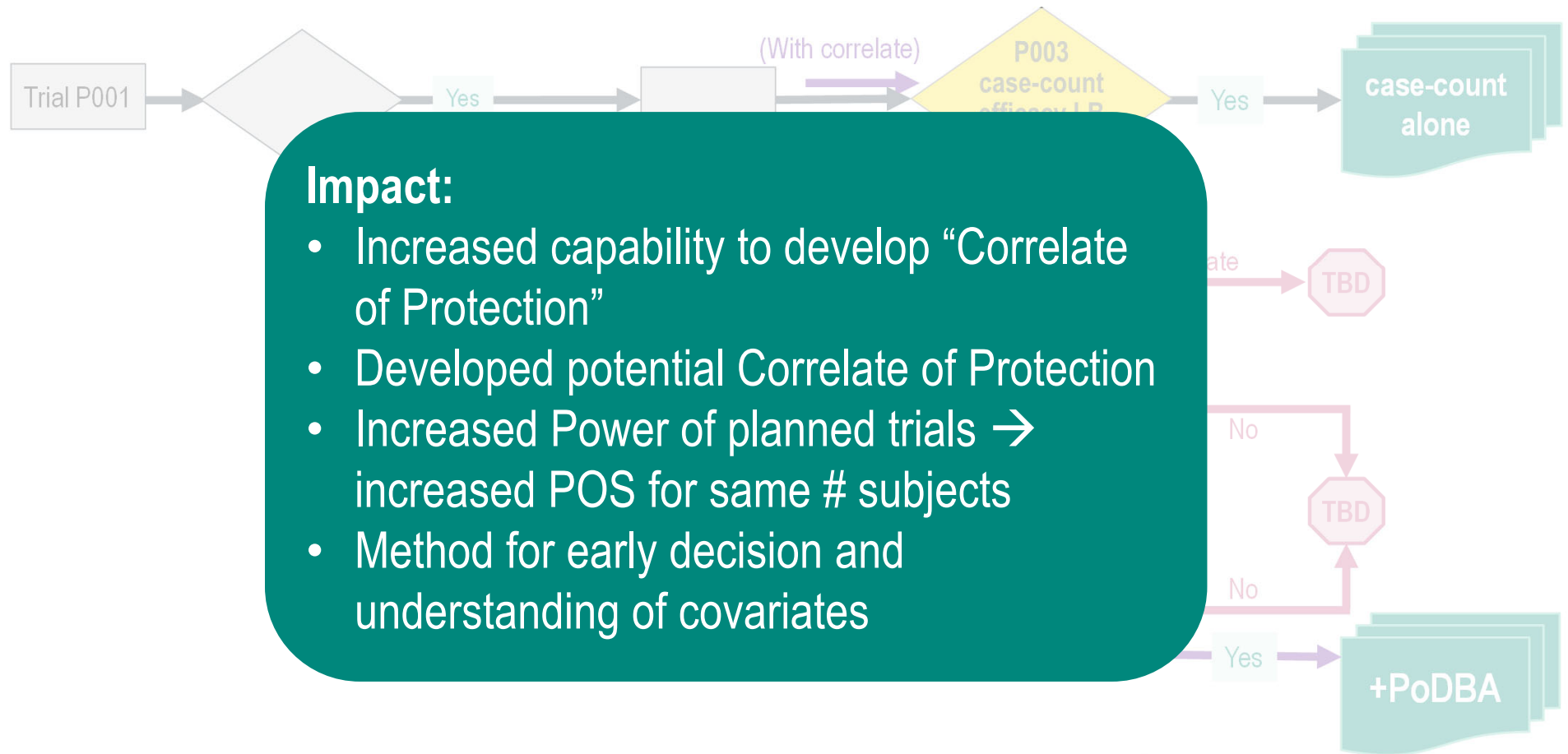
Method: Estimating the “Probability of Disease” Curve

- Use **titer values** measured in infected and non-infected subjects
- Assume that the relationship between titer values and probability of disease follows a **sigmoidal curve**
- **Estimate** the **parameters** of the curve and their **confidence intervals** using standard statistical method (*Maximum likelihood*)



Qualified PoDBA method & Efficacy CI
(demonstrated predictive power) with
published data and simulation

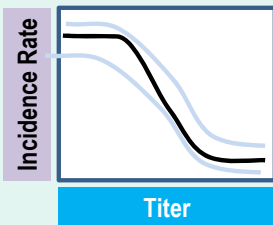
PoDBA → Novel Endpoint (example from a program plan, not agency guidance on different approaches to basis of licensure)



Phase	Key Question	Method(s)	Decision(s) / Impact
2	Is our immunogenicity likely to provide the necessary protection, and at what dose-level?	NLME+ MBMA + PoDBA (Comparator modeling + PoDBA)	No-go, GO, Dose-selection

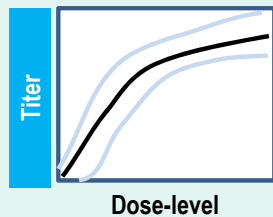
Can we leverage literature data and early immunogenicity data to drive early, objective, risk-based decisions?

Modeling Overview for Supporting Both Go and No-Go Decisions



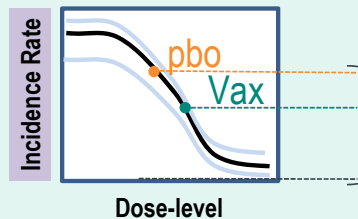
1. Titer → Incidence Rate (“IR”)

- Published clinical data
- Incidence rate for different disease levels
- Data cover various populations



2. Dose-level → Titer

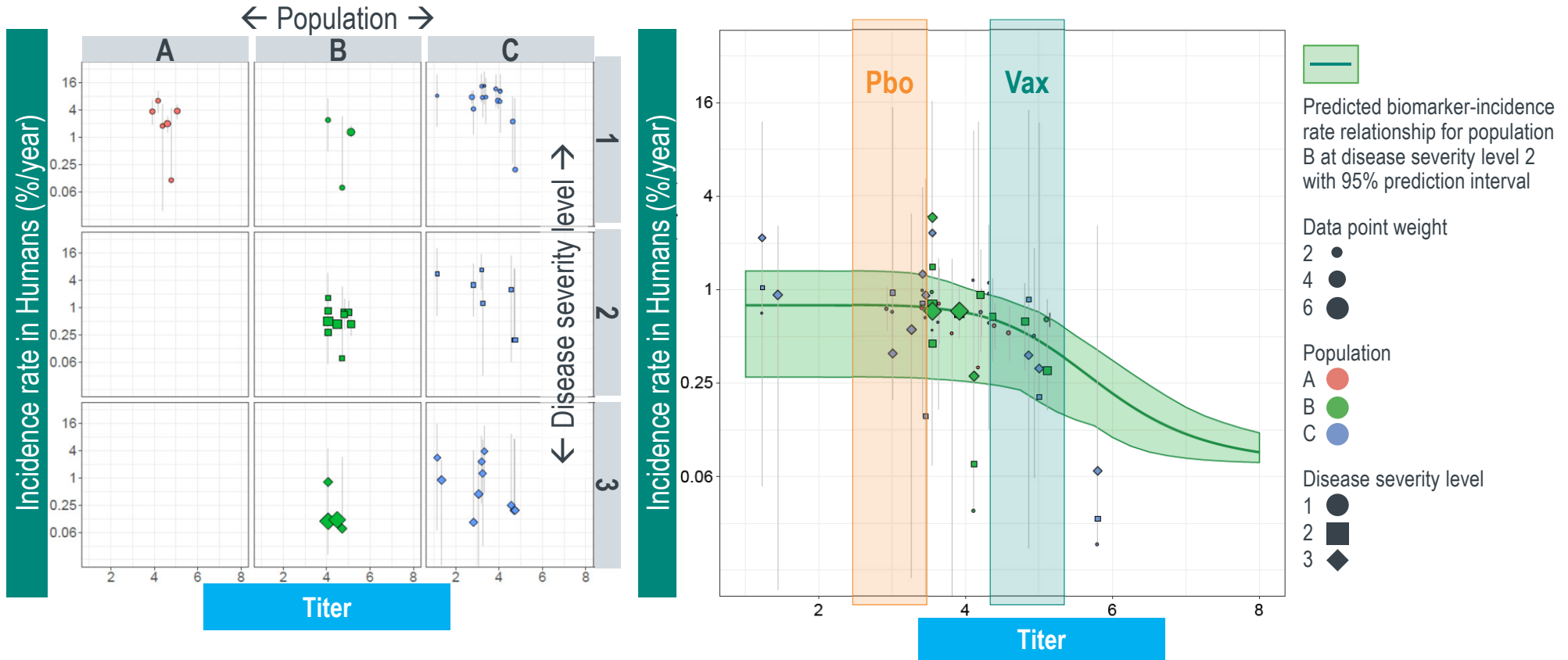
- Relate dose-level to serum neutralization titer response
- FIH Data



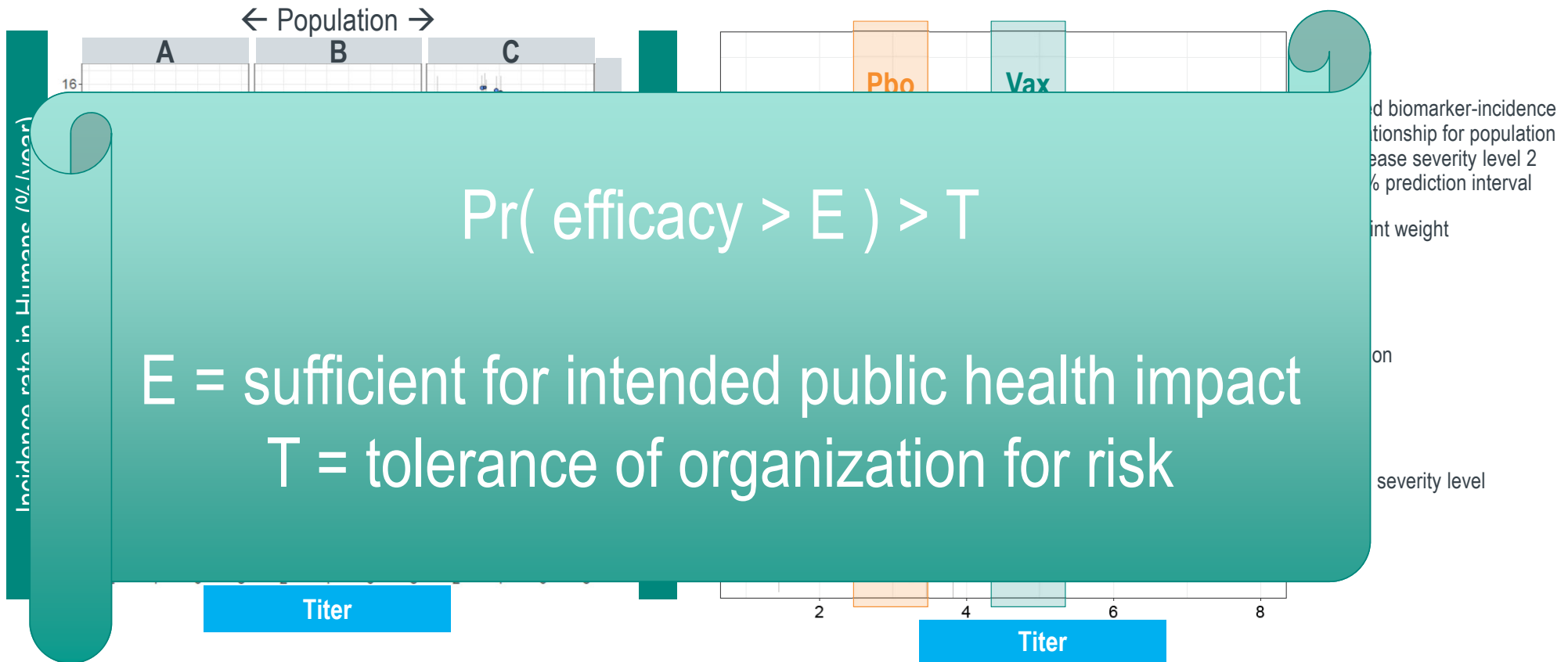
3. Combine 1 & 2 → “integrated” modeling:

Dose-level → Titer → Incidence Rate
Predicted change in incidence rate → efficacy

Visualization Has to Tie Together Data for Different Disease Levels and Populations



Visualization Has to Tie Together Data for Different Disease Levels and Populations



Visualization Has to Tie Together Data for Different Disease Levels and Populations



Impact:
Objective, Quantitative...
No-Go: \$40M trial
Go: \$10M trial
Dose-level, Biological insight, ...

Phase	Key Question	Method(s)	Decision(s) / Impact
2	What N (# subjects) will let us tell if vaccines A and B are different?	Phenomenological model Clinical trial simulation	Trial design, program strategy for sequence of trials
2	What immunogenicity assay(s) should we use and how often?	QSP and Bayesian probabilistic	Saved \$, increased POS Choice of assay, frequency
(na)	Can we model enough of the immune system to be predictive?	QSP	Capability development
4	How many doses of vaccine are needed to confer lasting protection?	QSP	Suggests 1 less dose Mechanistic insight
2	Which regimens should be tested in Ph. 2 Trial? (Regimen: dose-level, # doses, timing)	QSP	Ph. 2 Trial Design: add new dose level and different regimen
2,3	Do we have adequate evidence of efficacy if some pathogens have too few cases?	PoDBA (= Probability of Disease Bayesian Analysis)	Novel Ph. 3 endpoint GNG test criteria
2	Is our immunogenicity likely to provide the necessary protection, and at what dose-level?	NLME+ MBMA + PoDBA (Comparator modeling + PoDBA)	No-go, GO, Dose-selection

Vaccine Pharmacometrics

Today

- Simulation-based trial design to add/save trial arms or subjects
- QSP modeling of the immune system as a platform
- Using dose, regimen, formulation to predict immunogenicity/efficacy
- Translation between preclinical and clinical immunogenicity
- Leveraging literature data
- Establishing new trial endpoints
- Understanding covariate (age, genetics, geography,...) effects on immunogenicity & efficacy

Don't forget also...

- Predicting most effective vaccine platform by mechanistic modeling
- Understanding or predicting safety/toxicity
- Predicting the best route of administration
- Leveraging results of real-world trials
- Prioritizing vaccine candidates
- Prioritizing pathogen candidates

CONCLUSION

MSD



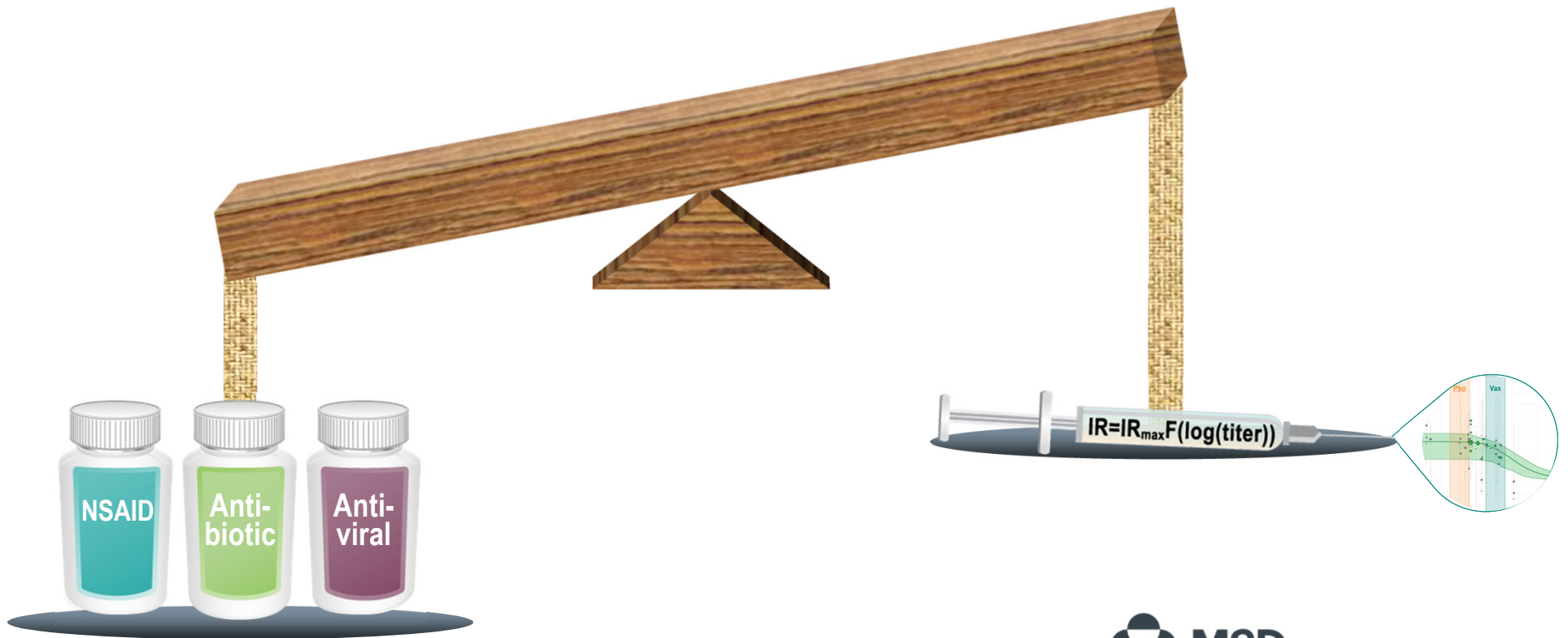
“An ounce of prevention is worth a pound of cure”

Benjamin Franklin



Source: <https://www.ag.ndsu.edu/news/columns/beefstalk/beefstalk-an-ounce-of-prevention-is-worth-a-pound-of-cure/>.

Pharmacometrics
"An ounce of ~~prevention~~ is worth a pound of cure"
Benjamin Franklin



Pharmacometrics
"An ounce of ~~prevention~~ is worth a pound of cure"
Benjamin Franklin

- Vaccines are a key component of public health
- Pharmacometrics useful for vaccine discovery & development
 - QSP, PK/PD, Bayesian, comparator, translational,...
- These (and other) methods have impacted decisions
- Pharmacometrics (we) can impact human health by helping inform vaccine discovery & development *even more*
 - Assumptions, study design & strategy, data interpretation

Acknowledgements

- Subjects (& their Parents)
- MSD
 - Discovery & Development Teams, PPDM-Bioanalytics
 - Vaxmodsim team+:
Luzelena Caro, Carolyn Cho, Julie Dudasova, Pavel Fiser, Jon Hartzel, Sean Hayes, Justina Ivanauskaite, Regina Laube, Brian Maas, Nitin Mehrotra, Jeff Perley, Seth Robey, Radha Railkar, Daniel Rosenbloom, Josiah Ryman
 - Paula Annunziato and Daria Hazuda
 - Dinesh de Alwis & Vikram Sinha, Nancy Agrawal
 - PPDM-QP2
 - Mike Pish & co. @ MCS
 - Skip Irvine, John Grabenstein
- Also...
 - Certara, Inc: Jos Lommerse, Nele Mueller-Plock, Michelle Green, Amy Cheung many others
 - Former MSD: Matt Wiener, Sandy Allerheiligen

Backup

PROPRIETARY ICONS HERE

MSD



Attributes of Vaccine Development

1. Stringent safety
2. Trial size and duration (event frequency)
3. Efficacy = proportional risk reduction
4. Surrogate marker (“Correlate of Protection,” a.k.a. “CoP”) challenges
5. Lack of translational models
6. Need arm with placebo or active comparator
7. Complex biologics, need to be transportable stable, usable

CoP Challenges: Knowledge, Time, Resources, Variability...



Knowledge

- Which measurements?
- Which species?
- Predictive power?



Time

- Knowledge early enough
- Timely availability of clinical data



Resources

- Many samples
- Often multiple species
- Resource-intensive assays



Variability

- Assays often +/- 2-fold
- Large BSV in response
- Variability in CoP predictive power?

BSV: between-subject variability