

# Pertussis Vaccination

## CMI-PB Project

Barry Grant  
UC San Diego  
<http://thegrantlab.org>

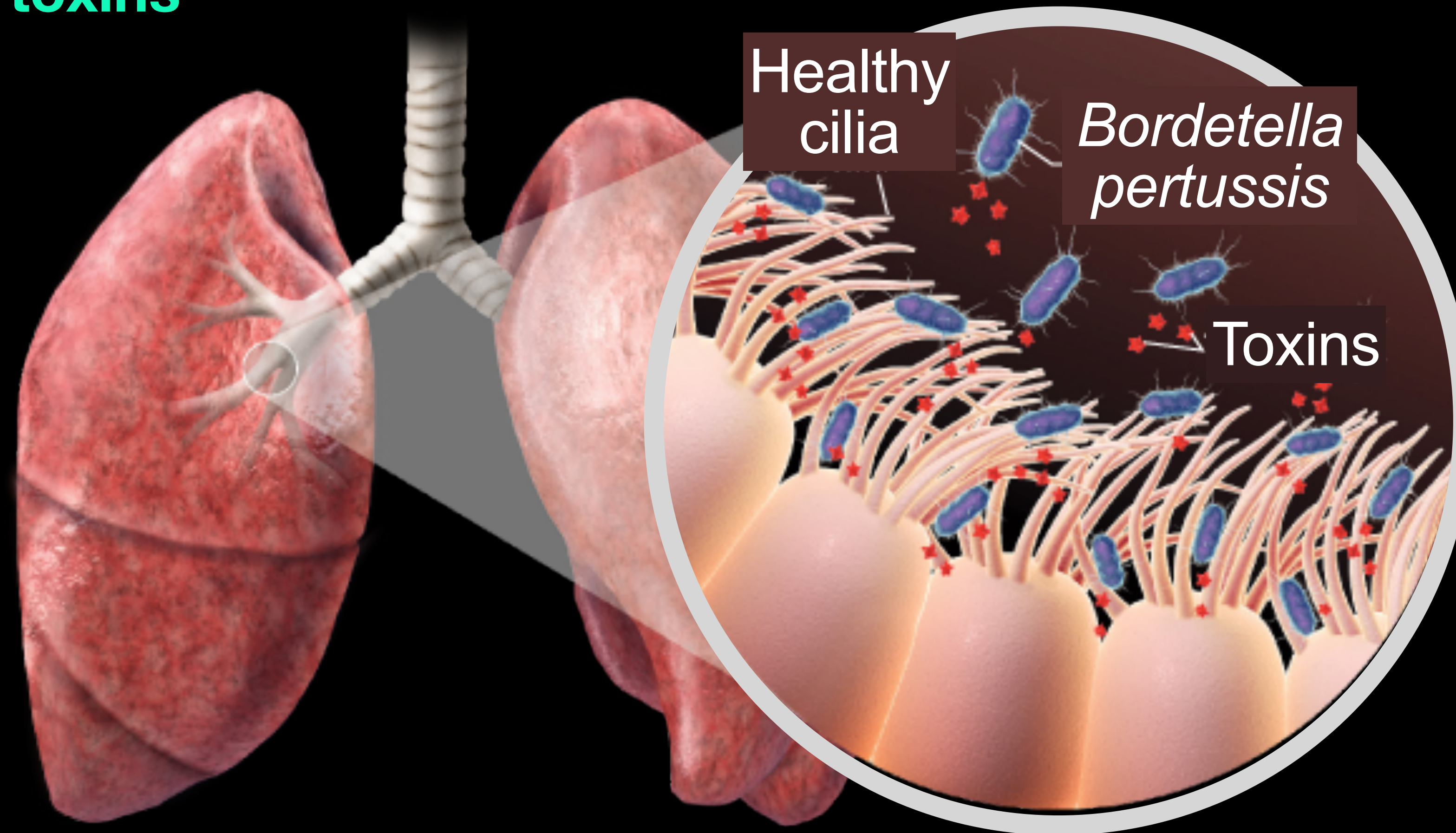
# Pertussis is a leading causes of vaccine-preventable deaths

Pertussis, or **whooping cough**, is a highly contagious lung infection caused by the bacteria *Bordetella pertussis*.

- Estimated 16 million cases and 200,000 associated infant deaths annually.
- Can infect people of all ages but is most severe and life threatening for infants under a year old
- Transmission occurs primarily through bacteria laden **respiratory droplets** produced when an infected individual coughs and sneezes

# *Bordetella pertussis* attacks cells lining the airways

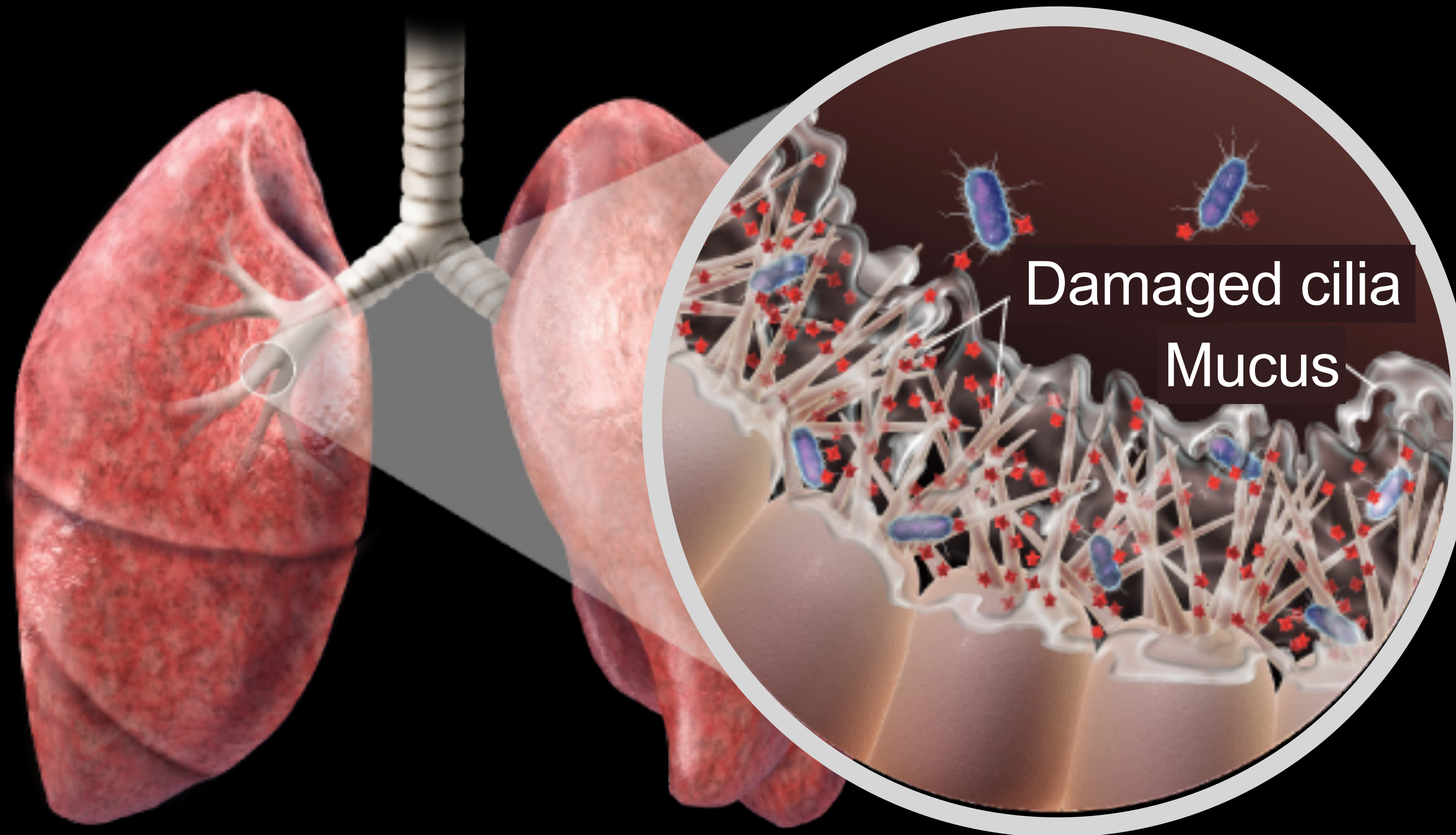
The bacteria use adhesive proteins to stick to **ciliated cells** whilst releasing **toxins**



[More details >](#)

# Pertussis is primarily a toxin-mediated disease

These toxins damage cilia, suppress the immune response and disrupt signaling leading to **inflammation**, **mucus buildup** and **impaired function**



[More details >](#)

# Pertussis develops in three main phases

Time  
(weeks)

1

2

3

4

5

6

7

8

9

10

11

12

13

Initial  
Infection

## Catarrhal Phase

Early Symptoms

(1-2 weeks)

- Runny nose
- Cough
- Mild fever
- Highly contagious

## Paroxysmal Phase

Severe Symptoms

(1-6 weeks but may last up to 10 weeks)

- Paroxysms (uncontrollable coughing fits)
- Inhalatory “whooping” sounds
- Difficulty breathing
- Vomiting
- Exhaustion

## Convalescent Phase

Recovery

(2-3 weeks)

- Gradual lessening of symptoms
- Reduced coughing fits
- Susceptible to other respiratory infections

[More details >](#)

# Fascinating history

CMI-PB  
www.cmi-pb.org



1679

## The Name "Pertussis" First Appears

The name pertussis (from Latin for "intensive cough") was first introduced by the English physician Thomas Sydenham in 1670. This name took over by the end of the decade. Earlier names included hooping cough, tuis perennis, tuisis epidemica infantum, and tuisis quinta.

[Read more](#)

1578

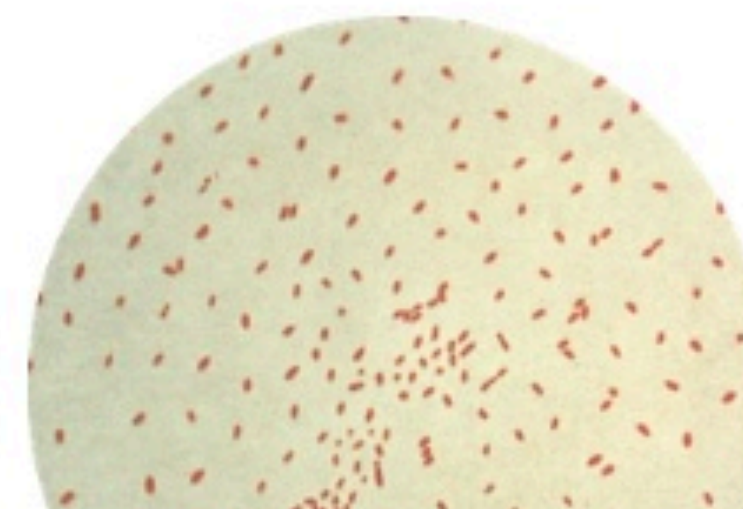


## First Epidemic Reported

The oldest known pertussis epidemic is thought to be the Paris outbreak of 1578. This was documented in detail by the French physician Guillaume de Baillou who described the classic symptoms of the disease.

[Read more](#)

1900



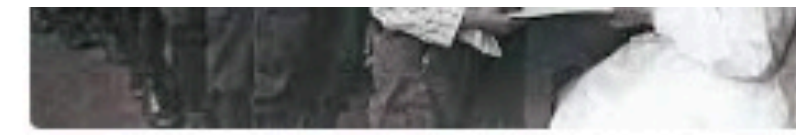
[Timeline >](#)



1942

### First DPT Vaccine

Pearl Kendrick at the Michigan Department of Health combined a refined whole-cell pertussis vaccine with Diphtheria and Tetanus toxoids to create the first combination DPT vaccine.



### First Whole-cell Pertussis Vaccine Tested on a Wide Scale

Danish physician Thorvald Madsen tested a whole-cell pertussis vaccine on a wide scale for the first time reporting promising results.

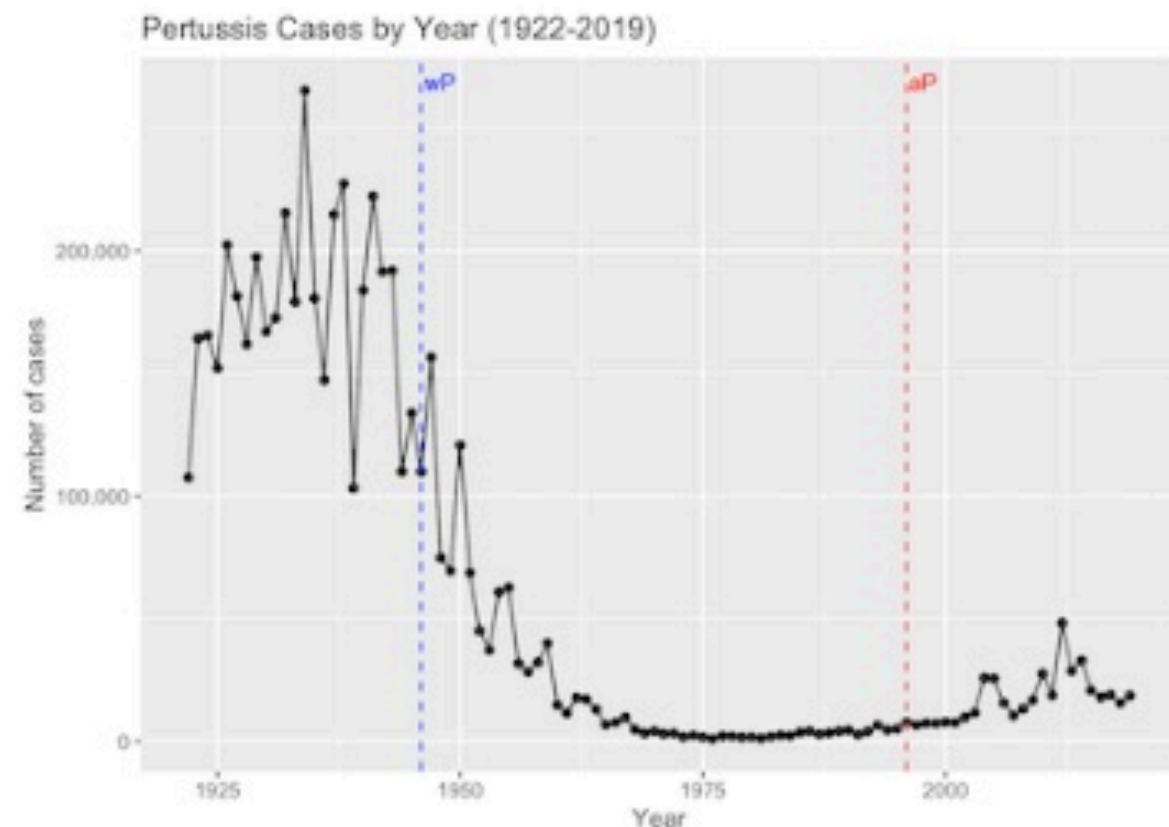
1947



### Routine Vaccination

In 1944, the Committee on Infectious Diseases of the American Academy of Pediatrics suggests routine use of pertussis vaccine and, in 1947, recommends its use in the form of the DPT combination. Routine childhood vaccination begins and is made compulsory in some states by the end of the decade.

1970



### Decline of Whooping Cough

There was a massive decline of pertussis cases in the U.S and other

[Timeline >](#)





### "DPT: Vaccine Roulette"

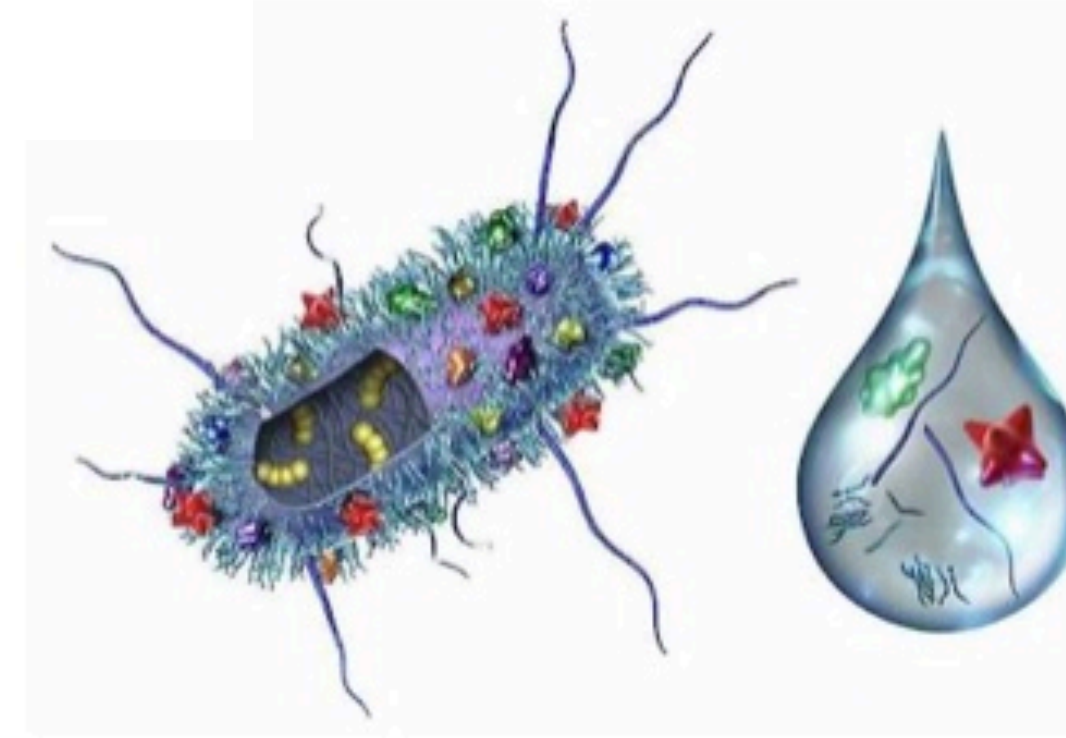
In 1982 negative publicity was encouraged from a documentary called "DPT: Vaccine Roulette", which led to a massive amount of lawsuits against the vaccine manufacturers. This documentary depicted the lives of children whose severe disabilities were **incorrectly** blamed on the DPT vaccine.

[Read more](#)



1985

1981



### Creation of DTaP Vaccine

Japanese scientist Yugi Sato created an acellular pertussis vaccine that contained purified haemagglutinins from *B. Pertussis*. This **aP vaccine** was first used in Japan soon after and was demonstrated to have fewer side effects than the whole-cell (**wP**) vaccine. It was later used in other countries (with additional components of *B. Pertussis*) as the combined DTaP vaccine.

[Read more](#)

1984



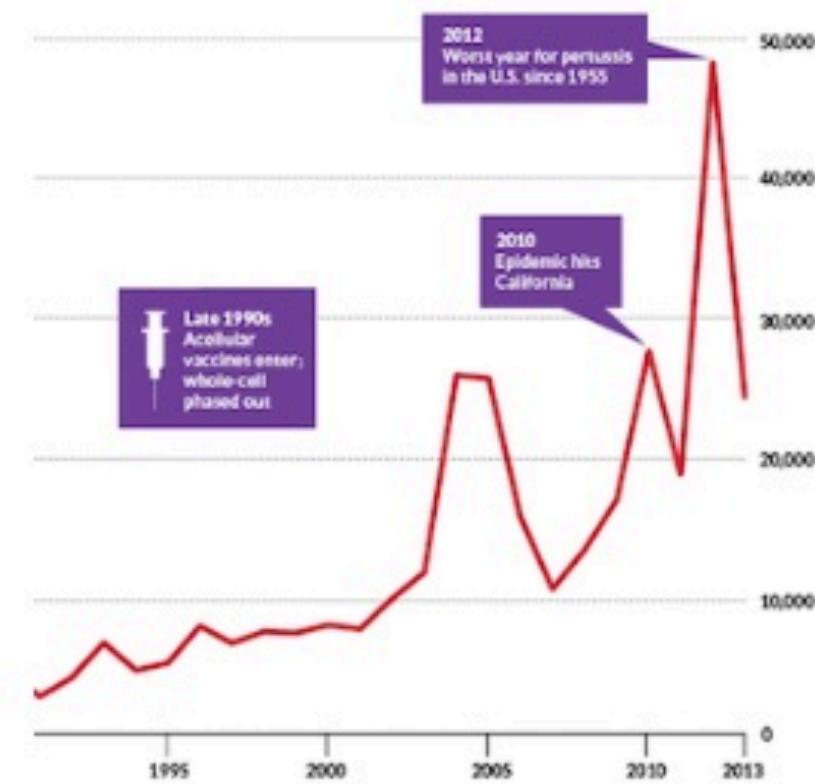
### Liability

By 1984 DPT vaccine manufacturers had a hard time obtaining liability insurance. By the end of the year, only one DPT manufacturer remained. Scientists respond by ramping up development and testing of safer new acellular pertussis vaccines. These would replace the older whole cell vaccine in many countries with a decade.

[Timeline >](#)



Later studies showed that there was no connection between the DPT vaccine and the permanent brain damage. It was in fact called a "Myth" and "Nonsense" by the Journal of American Medical Association in 1990.



### Pertussis Outbreaks

Major pertussis epidemics and outbreaks are once again a major public health concern. With epidemics typically occurring every 3 to 5 years in the U.S. as was evident in the pre-vaccine years. TO FINISH mention CA outbreak.

1992



### aP Vaccine Approved in the U.S.

The acellular pertussis (aP) vaccine was approved in the U.S in 1992, the the older wP formalization was phased out and completely replaced with the DTaP vaccine combination in 1996.

2010

2020



### CMI-PB Project

A new [systems vaccinology project](#) is launched that combines systems biology and genomics to provide a more holistic picture of protective pertussis-specific immune mechanisms. The project provides the scientific community with comprehensive, high-quality, and freely accessible resources related to Pertussis booster vaccination.

These resources, and associated [prediction challenges](#), are geared towards engaging both experts and enthusiasts in developing and improving **computational models** of the immune response to vaccination and in turn informing new intervention strategies to curb the increasing frequency of *B. pertussis* infection.

## Pertussis (Whooping Cough)

CDC > Pertussis Home > Surveillance & Reporting

 [Pertussis Home](#)

[About Pertussis](#) +

[Vaccination](#)

[Pregnancy & Whooping Cough](#)

[Outbreaks](#)

[Clinicians](#) +

[Public Health Professionals](#) +

[Surveillance & Reporting](#) -

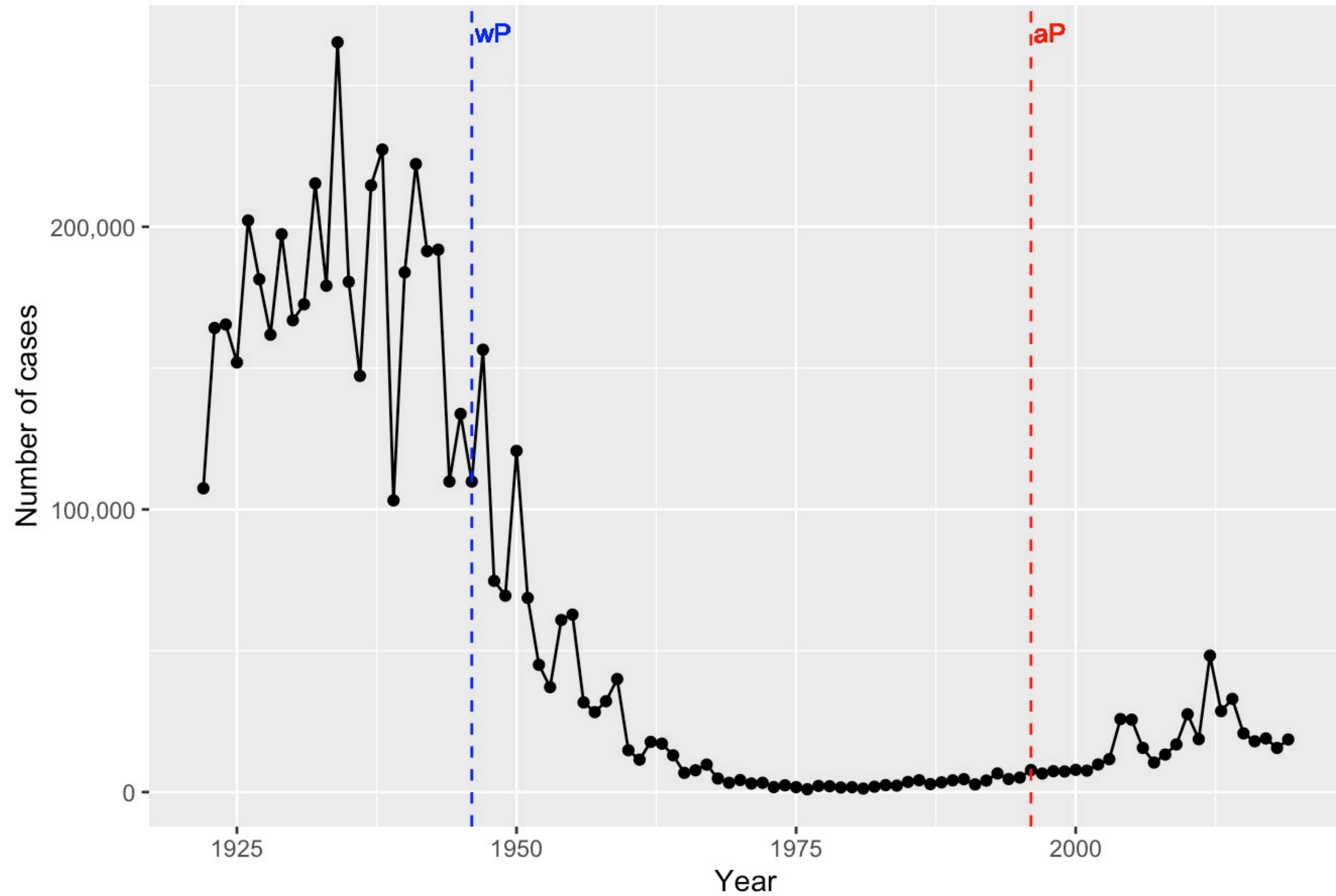
# Pertussis Cases by Year (1922-2019)

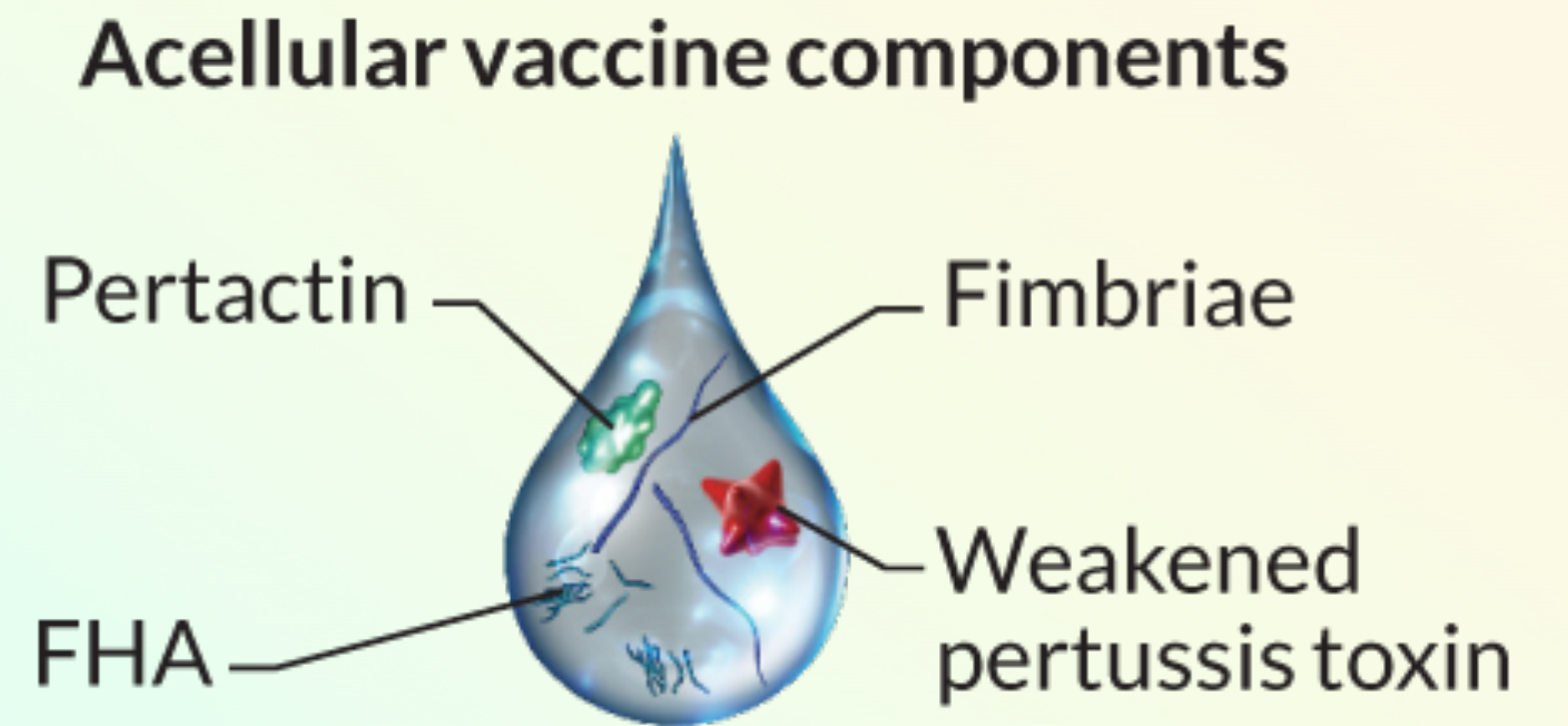
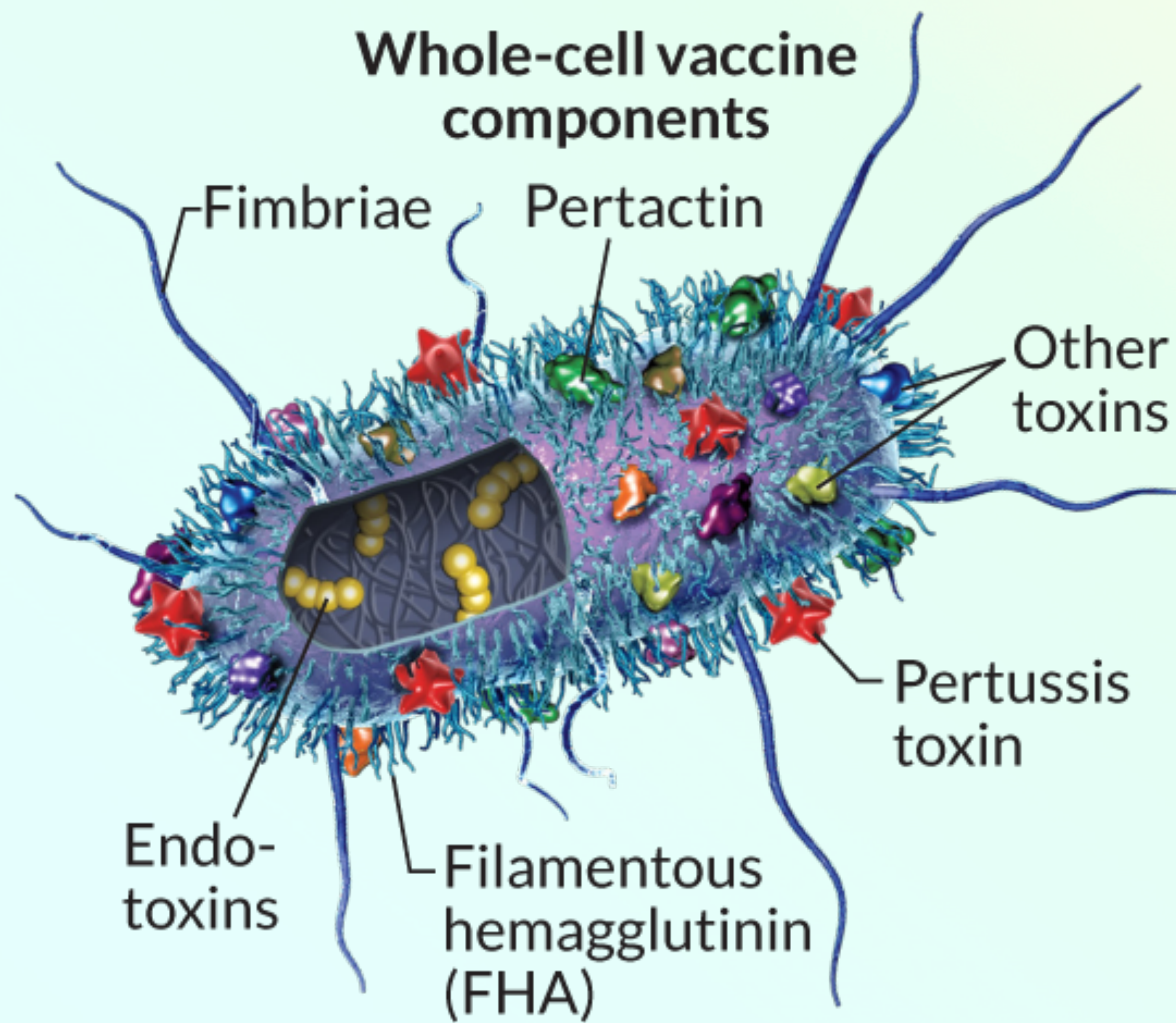
[Print](#)

This table shows reported pertussis cases in the United States since 1922. The related trend charts can be found on the [Surveillance and Reporting](#) page.

| Year | No. Reported Pertussis Cases |
|------|------------------------------|
| 1922 | 107,473                      |
| 1923 | 164,191                      |
| 1924 | 165,418                      |
| 1925 | 152,003                      |
| 1926 | 202,210                      |

Pertussis Cases by Year (1922-2019)





[More details >](#)

# Major aP vaccines (US)

| <b>Vaccine</b> | <b>Trade Name</b>     | <b>Manufacturers</b>               | <b>Components (Concentrations)</b>                                       |
|----------------|-----------------------|------------------------------------|--|
| DTaP           | Daptacel,<br>Infanrix | Sanofi Pasteur,<br>GlaxoSmithKline | Inactivated PT: 10-20 µg, FHA: 5-20 µg,<br>PRN: 3-5 µg, FIM 2+3: 5-10 µg |
| Tdap           | Adacel,<br>Boostrix   | Sanofi Pasteur,<br>GlaxoSmithKline | Inactivated PT: 2.5-8 µg, FHA: 5-8 µg,<br>PRN: 3-5 µg, FIM 2+3: 5-8 µg   |

[More details >](#)

# Major aP vaccines (US)



Tdap  
for pregnant  
women

- During the 27-36th week of each pregnancy



DTaP  
for young  
children

- 2, 4 and 6 months
- 15 through 18 months
- 4 through 6 years



Tdap  
for preteens

- 11 through 12 years



Tdap  
for adults

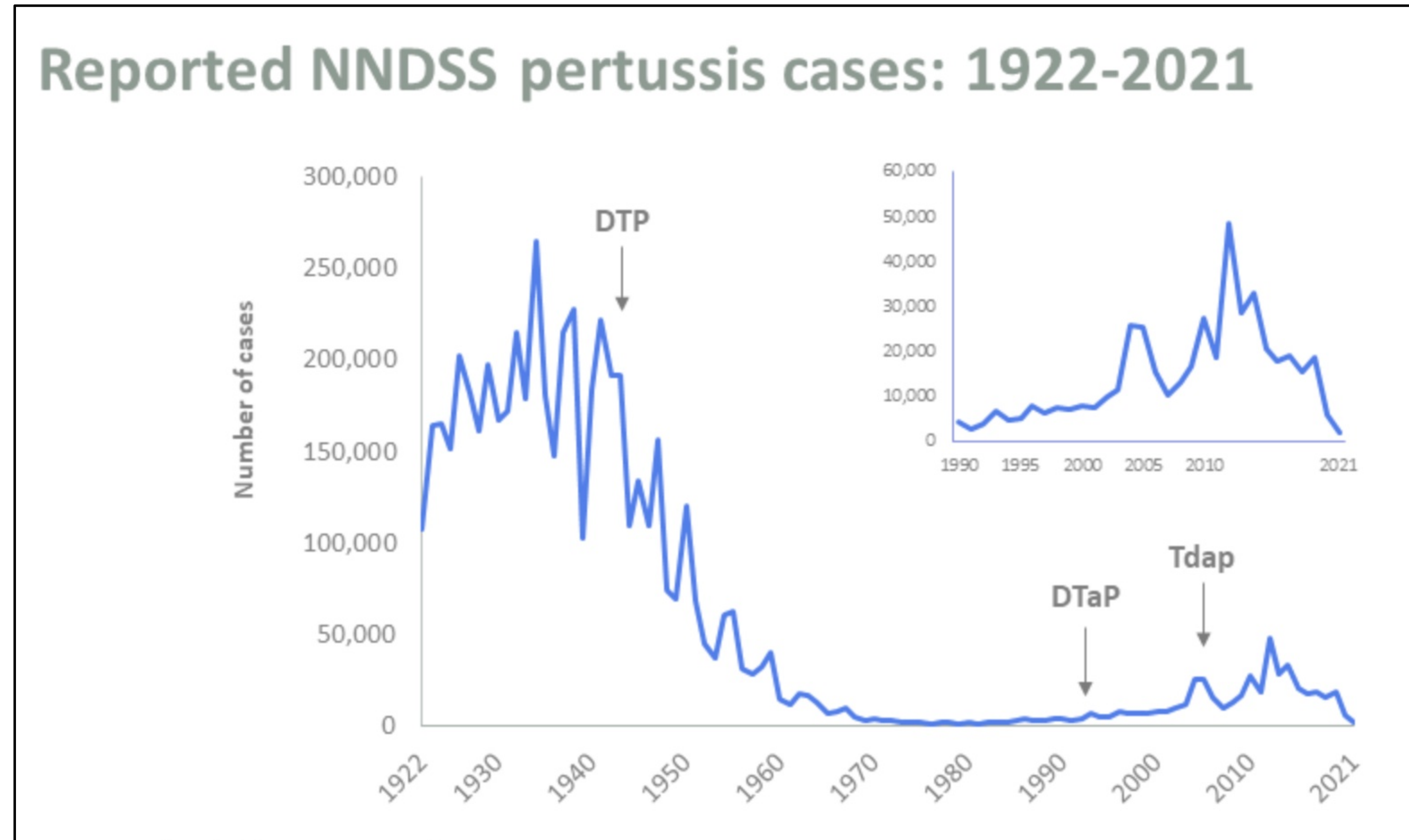
- Anytime for those who have never received it
- Subsequent boosters at 10 year intervals following initial vaccine

Source: Centers for Disease Control

[More details >](#)

## Waning Immunity from aP pertussis vaccination

- 1940s: Introduction of an inactivated whole bacteria PT vaccine (**wP**) dramatically decreased cases
- 1995: Vaccine-related side effects led to a replacement with the acellular PT vaccine (**aP**) in the USA
- aP induced protection wanes faster than wP → **Why?**



Source: National Notifiable Diseases Surveillance System, CDC



[Sign in to CMI-PB](#)

Ab titer

The mission of CMI-PB is to provide the scientific community with a comprehensive, high-quality and freely accessible resource of Pertussis booster vaccination.

### LEARN ABOUT THE PROJECT



The NIH funded CMI network  
What is pertussis vaccination?  
What are the open scientific questions?  
The CMI-PB approach: A community

### UNDERSTAND THE DATA



How do we measure immune responses?  
What data is available?  
Our approach to data standardization  
Browse our terminology

### ACCESS THE DATA



Data statistics  
Use the API in your programs  
Download all data (SFTP)  
More ...





[Sign in to CMI-PB](#)

Ab titer [dropdown]  [Search](#)

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### UNDERSTAND THE DATA



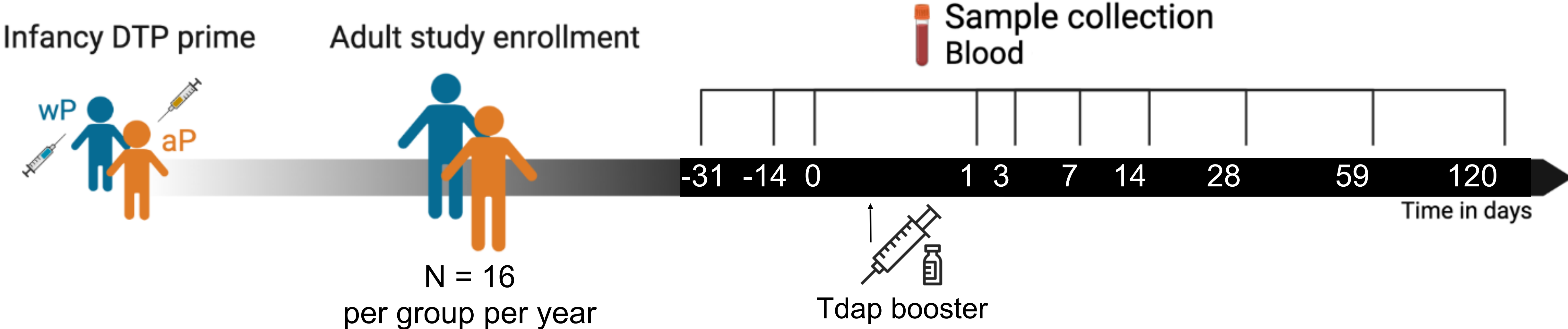
- How do we measure immune responses?
- What data is available?**
- Our approach to data standardization
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### ACCESS THE DATA

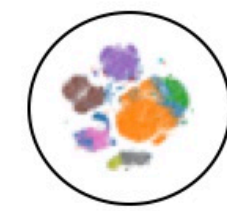
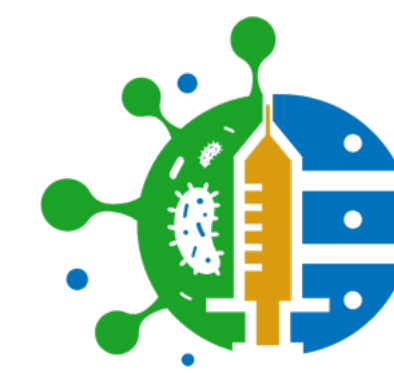


- Data statistics**
- Use the API in your programs
- Download all data (SFTP)
- More ...

# Recruitment Strategy

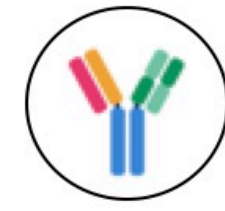


# Characterizing immune responses - Multiomics approach



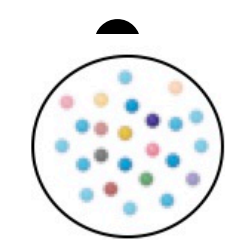
## **PBMC cell frequencies** by flow cytometry

- Total of 37 distinct cell populations



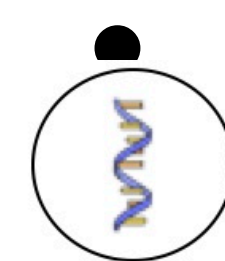
## **Plasma antigen-specific antibody titers** by Luminex

- Antibody Isotypes: IgG, IgG1, IgG2, IgG3, IgG4
- Vaccine Antigens
  - Pertussis Toxin (PT), PRN, FHA, FIM2/3
  - Tetanus Toxoids (TT)
  - Diphtheria Toxoids (DT)
  - OVA (irrelevant control)



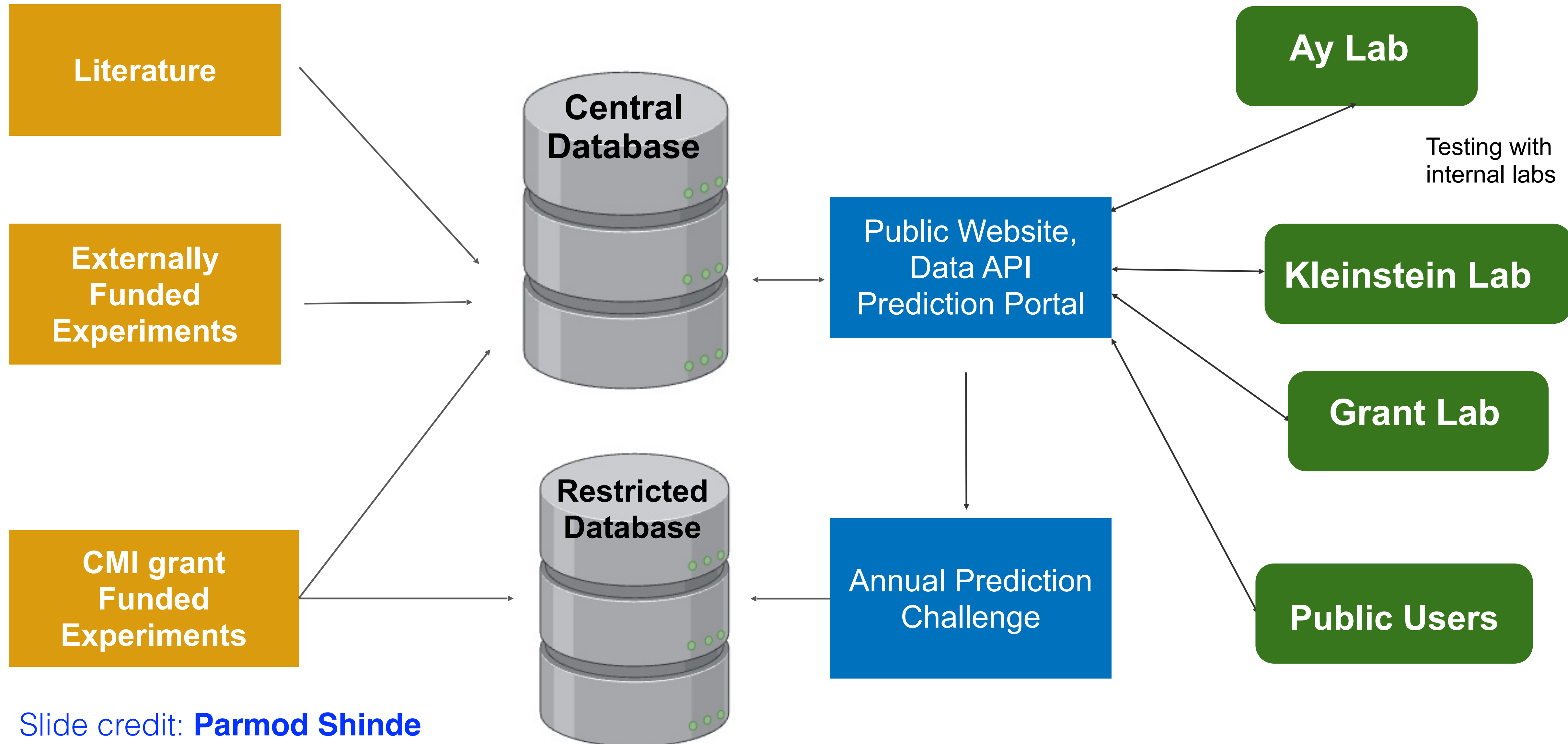
## **Plasma proteomics** by Olink

- Concentration of 45 cytokines

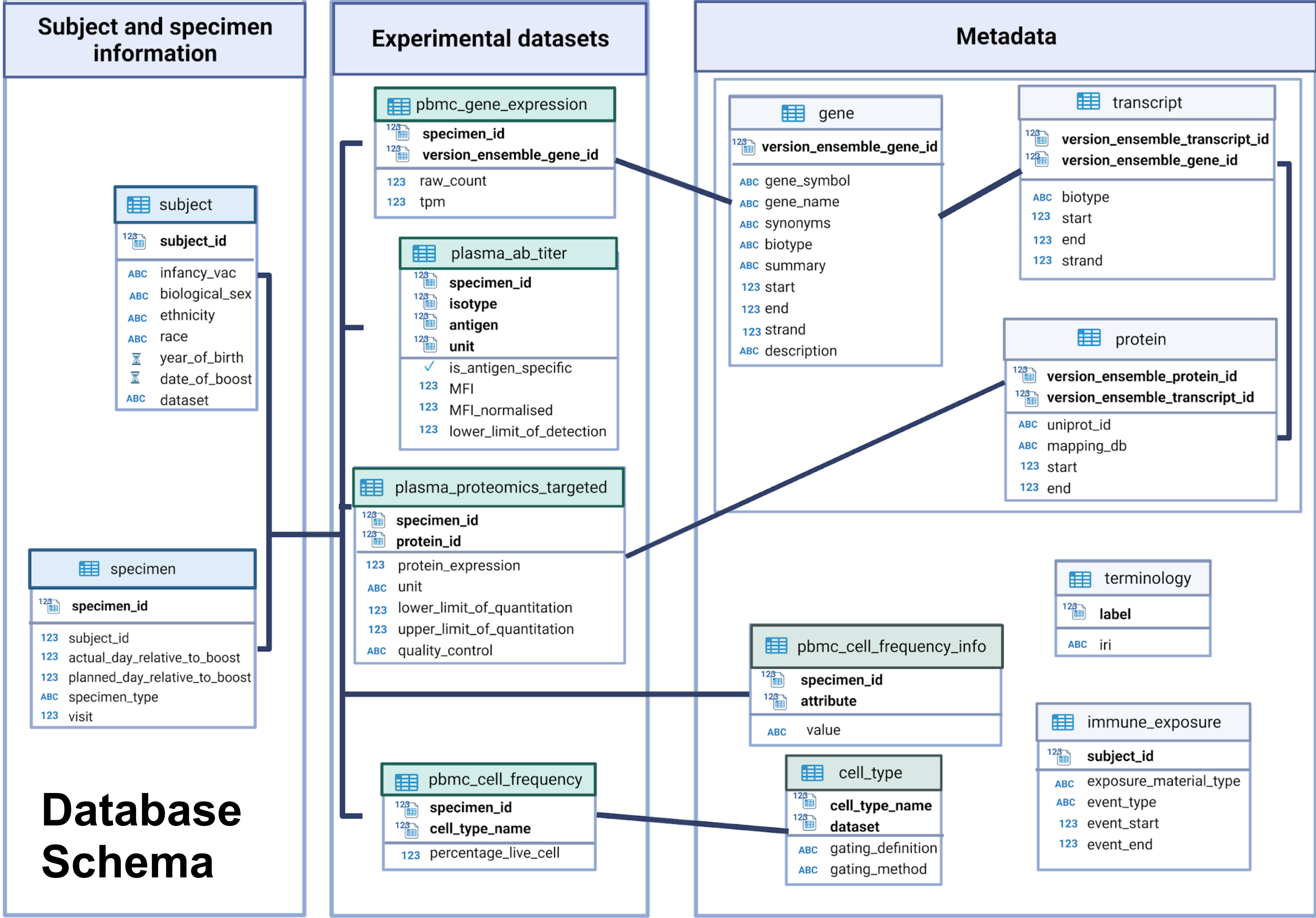


## **Transcriptomics** by bulk RNA-Seq

# d. Databases, Model building, and Prediction Challenge



# e. Providing access to experimental data in a standardized format



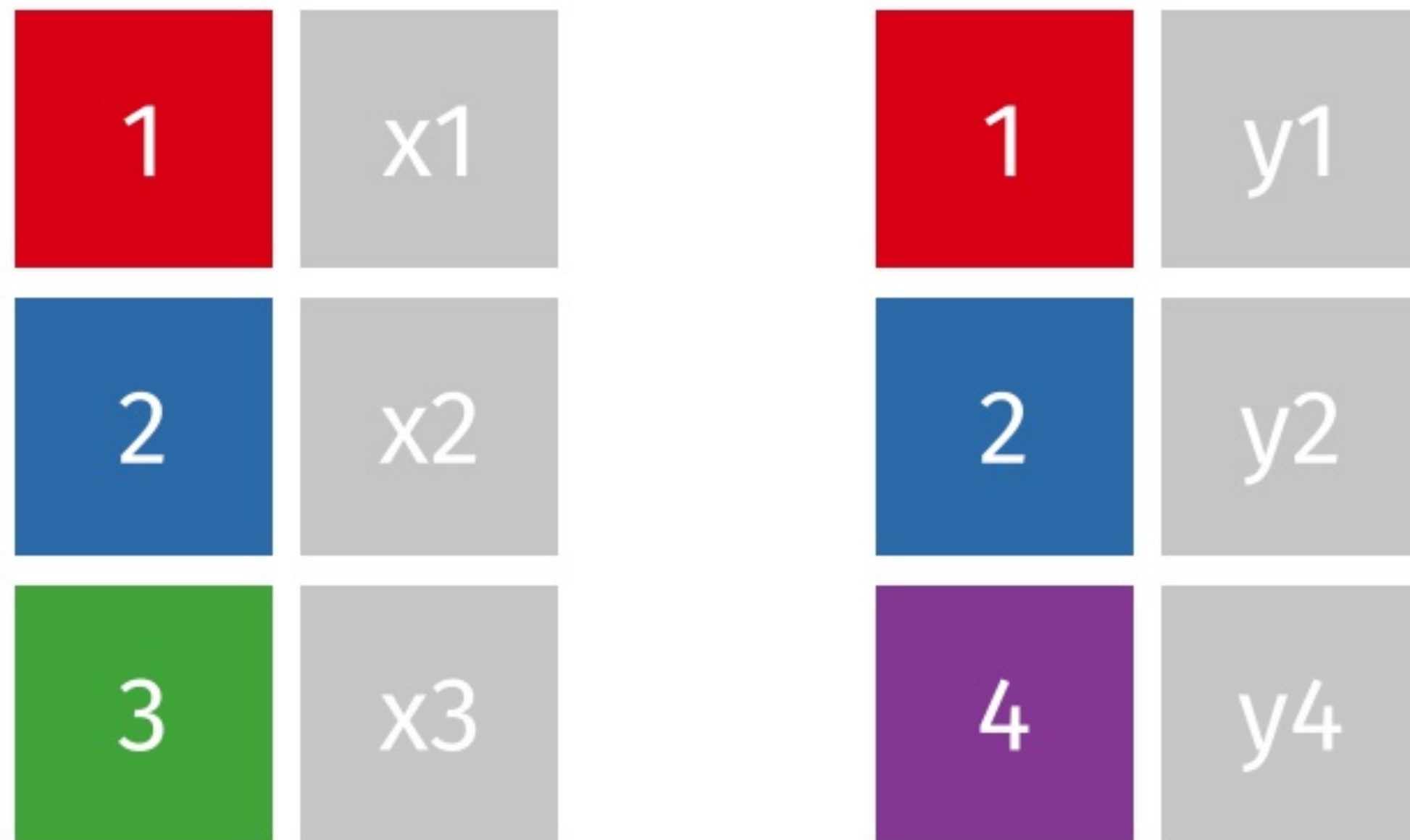
# INFORMATION TABLES

| SUBJECT           |
|-------------------|
| <b>subject_id</b> |
| infancy_vac       |
| biological_sex    |
| ethnicity         |
| race              |
| year_of_birth     |
| date_of_boost     |
| dataset           |

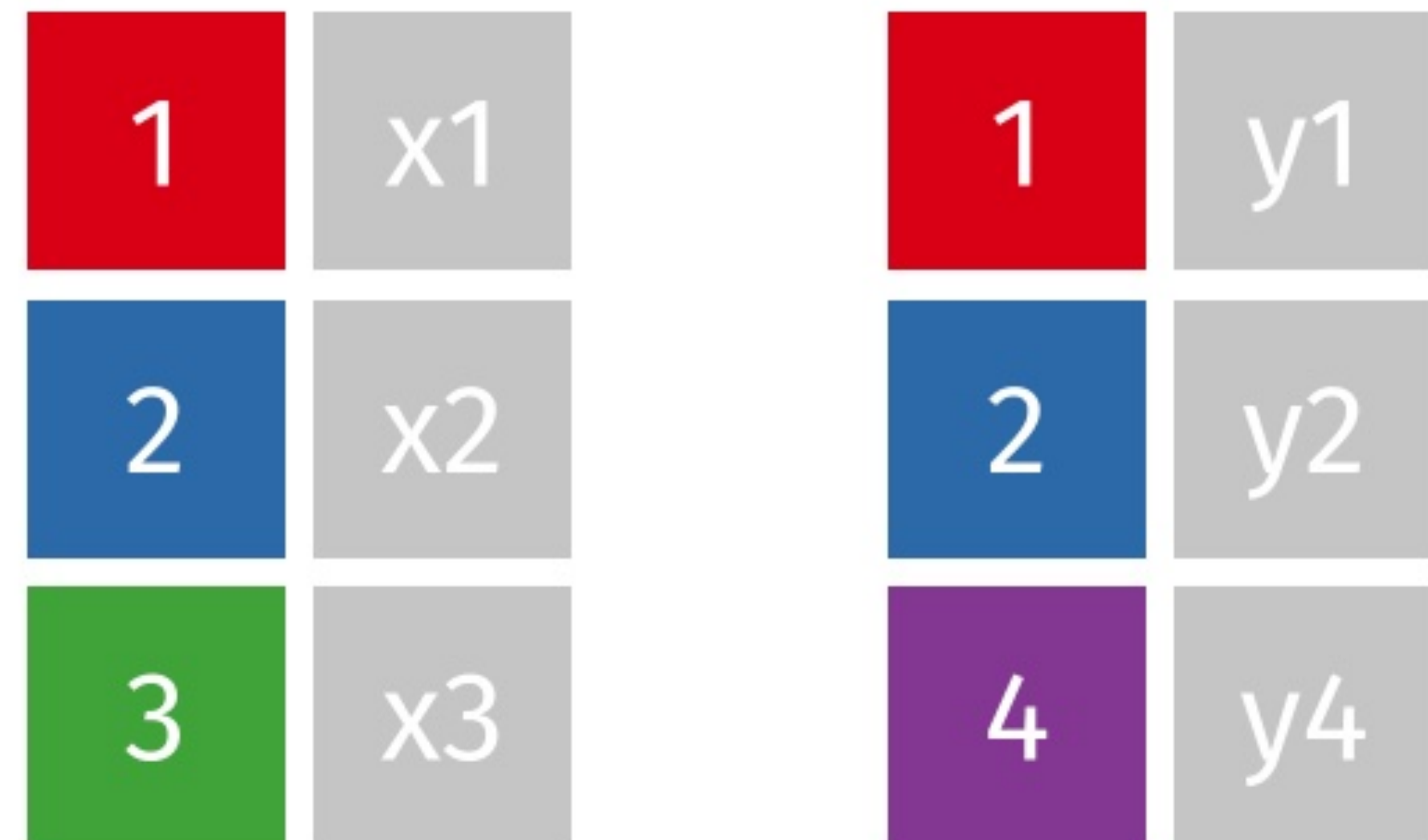
| SPECIMEN                      |
|-------------------------------|
| <b>specimen_id</b>            |
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| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |

# DPLYR **\*\_JOIN()** FUNCTIONS...

`inner_join(x, y)`

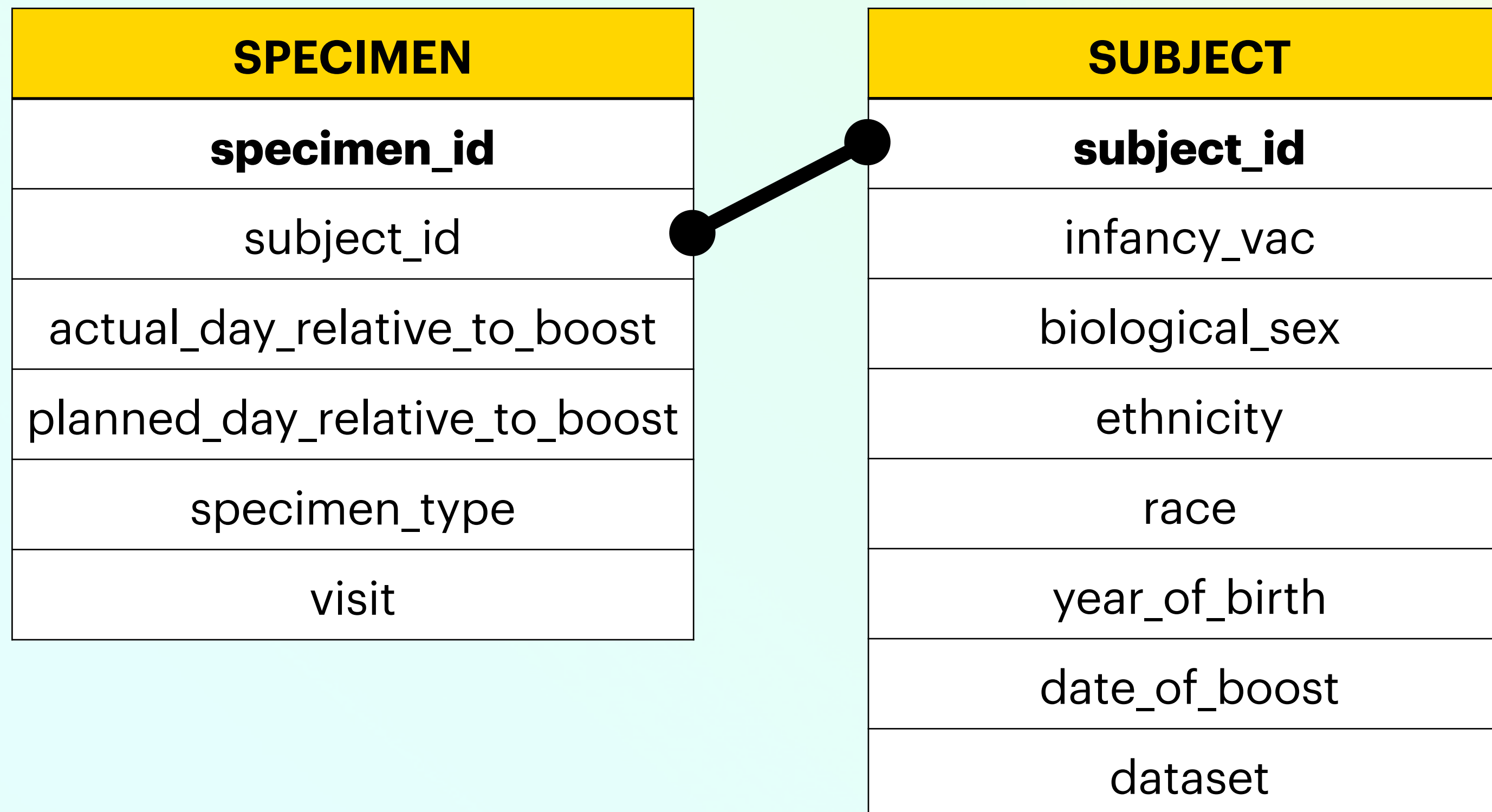


`full_join(x, y)`



# INFORMATION TABLES

CAN BE LINKED BY "SUBJECT\_ID"





# INFORMATION TABLES

CAN BE LINKED BY "SUBJECT\_ID"

| SPECIMEN                      |
|-------------------------------|
| <b>specimen_id</b>            |
| subject_id                    |
| actual_day_relative_to_boost  |
| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |

| SUBJECT           |
|-------------------|
| <b>subject_id</b> |
| infancy_vac       |
| biological_sex    |
| ethnicity         |
| race              |
| year_of_birth     |
| date_of_boost     |
| dataset           |

# WE WANT ONE **META** TABLE

USE DPLYR **\*\_JOIN()** FUNCTIONS...

| SPECIMEN                      |
|-------------------------------|
| <b>specimen_id</b>            |
| subject_id                    |
| actual_day_relative_to_boost  |
| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |



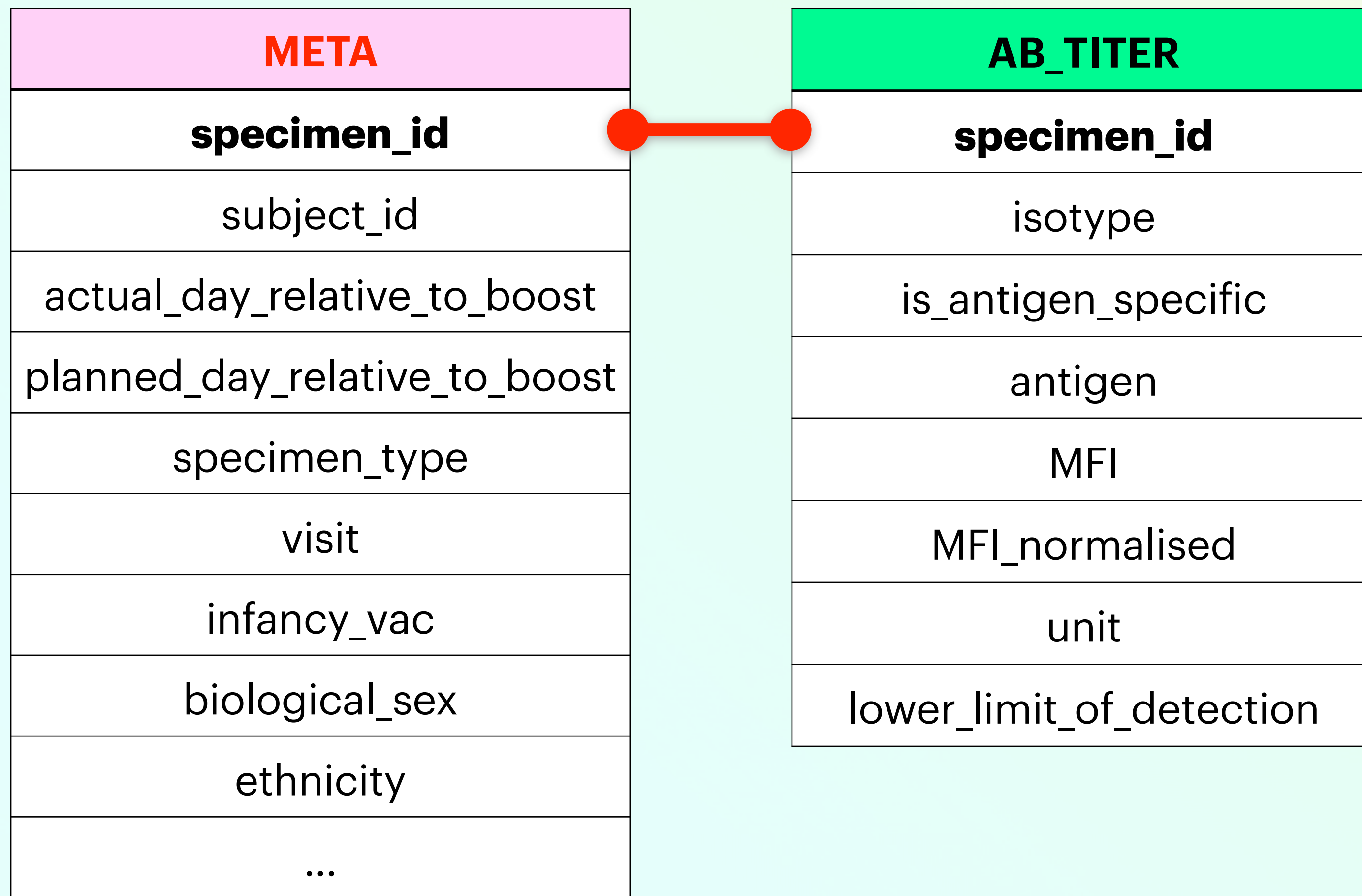
| SUBJECT           |
|-------------------|
| <b>subject_id</b> |
| infancy_vac       |
| biological_sex    |
| ethnicity         |
| race              |
| year_of_birth     |
| date_of_boost     |
| dataset           |

**=**

| META                          |
|-------------------------------|
| specimen_id                   |
| subject_id                    |
| actual_day_relative_to_boost  |
| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |
| infancy_vac                   |
| biological_sex                |
| ethnicity                     |
| race                          |
| year_of_birth                 |
| date_of_boost                 |
| dataset                       |

# JOIN WITH EXPEREMENT TABLES

USE DPLYR `*_JOIN()` FUNCTIONS...



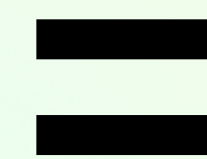
# META + EXPERIMENT

HAS EVERYTHING WE NEED FOR FURTHER ANALYSIS...

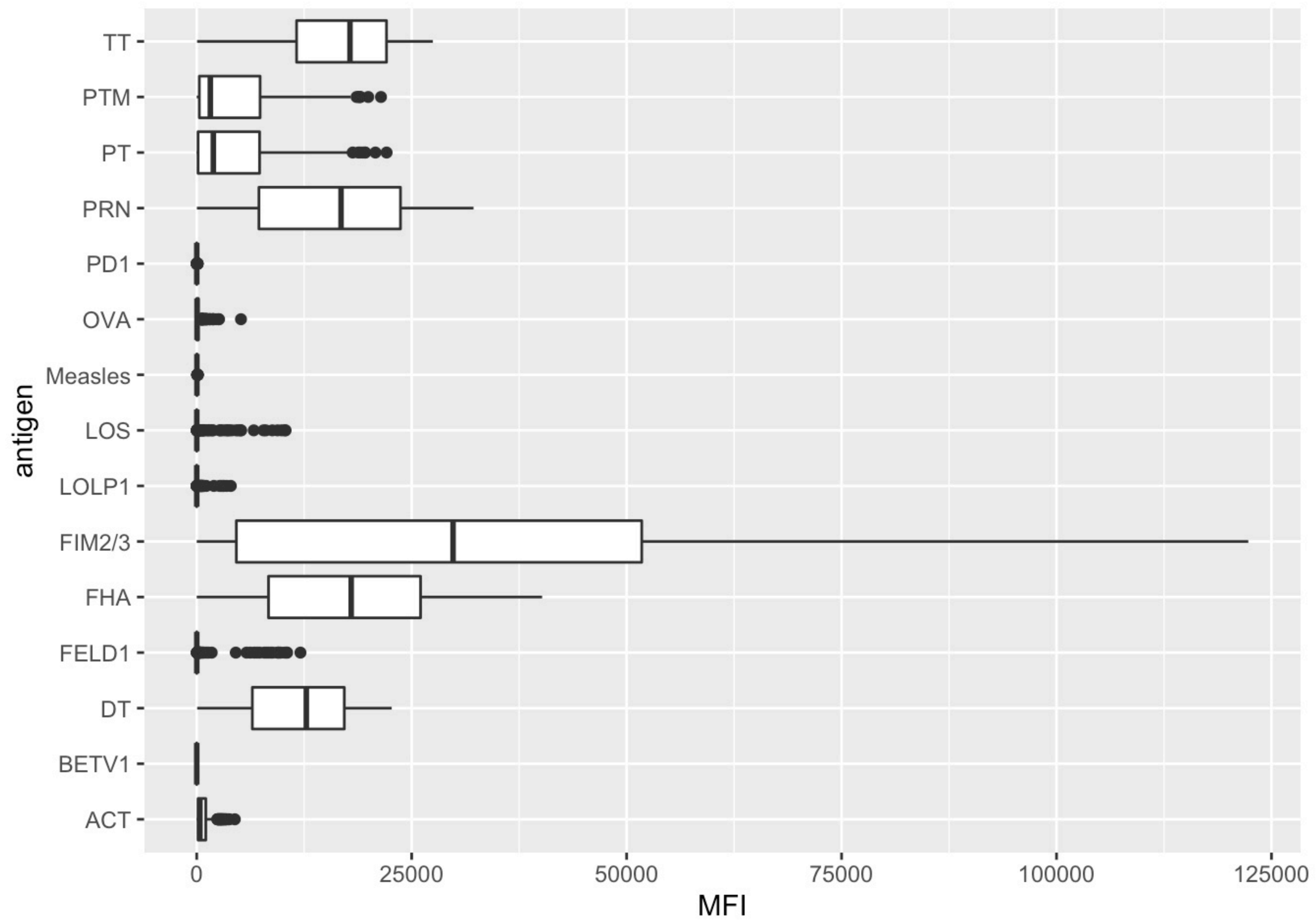
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|-------------------------------|
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| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |
| infancy_vac                   |
| biological_sex                |
| ethnicity                     |
| ...                           |



| AB_TITER                 |
|--------------------------|
| <b>specimen_id</b>       |
| isotype                  |
| is_antigen_specific      |
| antigen                  |
| MFI                      |
| MFI_normalised           |
| unit                     |
| lower_limit_of_detection |



| ABDATA                        |
|-------------------------------|
| <b>specimen_id</b>            |
| subject_id                    |
| actual_day_relative_to_boost  |
| planned_day_relative_to_boost |
| specimen_type                 |
| visit                         |
| infancy_vac                   |
| biological_sex                |
| ethnicity                     |
| race                          |
| year_of_birth                 |
| date_of_boost                 |
| dataset                       |
| isotype                       |
| is_antigen_specific           |
| antigen                       |
| MFI                           |
| MFI_normalised                |
| unit                          |
| lower_limit_of_detection      |





Sign in to CMI-PB

Terminology

FIMbrial protein

Search

Terminology Browser

uniprot:Q518X0

- Fimbrial protein
- Fimbrial protein **Fim3**
- Mixture of **Fim2** and **Fim3**

# Fimbrial prote...

<https://www.uniprot.org/uniprot/Q518X0>

Ontology  
Class  
    material entity  
    molecular entity  
    protein  
        Fimbrial protein  
Annotation Property  
Data Property  
Object Property  
Individual

- label
  - Fimbrial protein
- CMI-PB alternative term
  - fim2
- type
  - Class
- subclass of
  - protein

- Function
- Names & Taxonomy
- Subcellular Location
- Phenotypes & Variants
- PTM/Processing
- Expression
- Interaction
- Structure
- Family & Domains
- Sequence
- Similar Proteins

# Q518X0 · Q518X0\_BORPT

|                       |                               |                                |           |
|-----------------------|-------------------------------|--------------------------------|-----------|
| Protein <sup>i</sup>  | Fimbrial protein              | Amino acids                    | 207       |
| Status <sup>i</sup>   | UniProtKB unreviewed (TrEMBL) | Protein existence <sup>i</sup> | Predicted |
| Organism <sup>i</sup> | Bordetella pertussis          | Annotation score <sup>i</sup>  | 1/5       |
| Gene <sup>i</sup>     | fim2                          |                                |           |

- Entry
- Feature viewer
- Publications
- External links
- History

BLAST, Align, Download, Add, Add a publication, Entry feedback

## Function<sup>i</sup>

### GO Annotations<sup>i</sup>

Slimming set:

generic

Feedback, Help

- Function
- Names & Taxonomy
- Subcellular Location
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Entry Feature viewer Publications External links History

BLAST Align Download Add Add a publication Entry feedback

## Function<sup>i</sup>

## GO Annotations<sup>i</sup>

Slimming set:  
generic



Cell color indicative of number of GO terms

| ASPECT             | TERM  |
|--------------------|---|
| Cellular Component | <b>pilus</b> <a href="#">↗</a> <small>Source:InterPro</small>         |
| Biological Process | <b>cell adhesion</b> <a href="#">↗</a> <small>Source:InterPro</small> |

Expand table

We'd like to inform you that we have updated our [Privacy Notice](#) to comply



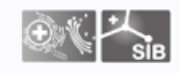
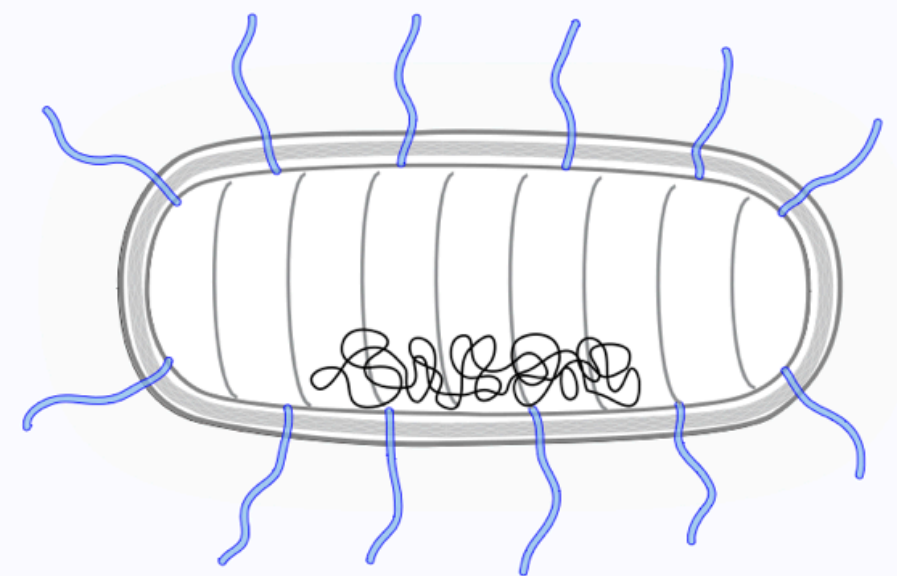
- Function
- Names & Taxonomy
- Subcellular Location**
- Phenotypes & Variants
- PTM/Processing
- Expression
- Interaction
- Structure
- Family & Domains
- Sequence
- Similar Proteins

### Subcellular Location<sup>i</sup>

UniProt Annotation GO Annotation

[pilus](#)

[Complete GO annotation on QuickGO](#)



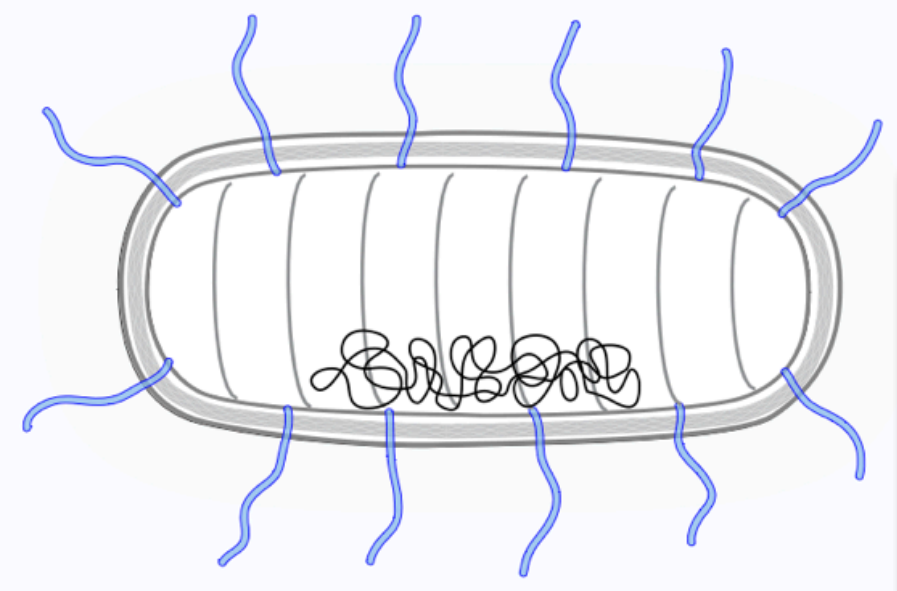
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UniProt Annotation GO Annotation

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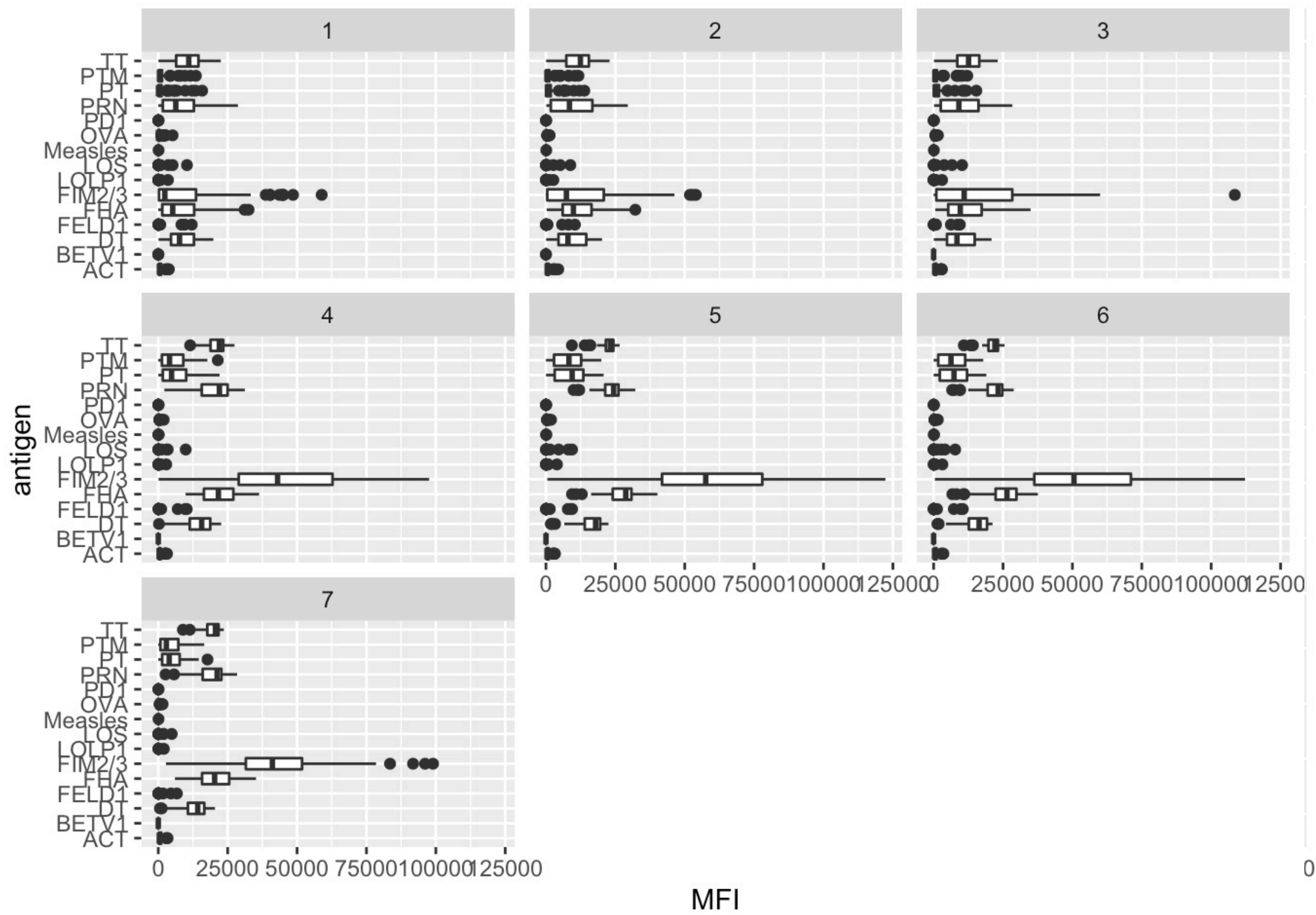
[Complete GO annotation on QuickGO](#)



**Fimbrium**  
A fimbrium or pilus is a hair-like, non-flagellar, polymeric filamentous appendage that extend from the bacterial or archaeal cell surface, such as type 1 pili, P-pili, type IV pili or curli. Pili perform a variety of functions, including surface adhesion, motility, cell-cell interactions, biofilm formation, conjugation, DNA uptake, and twitching motility.

Feedback

Help

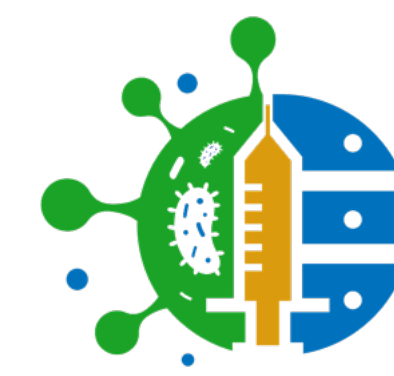


## a. Past and future CMI-PB annual prediction challenges



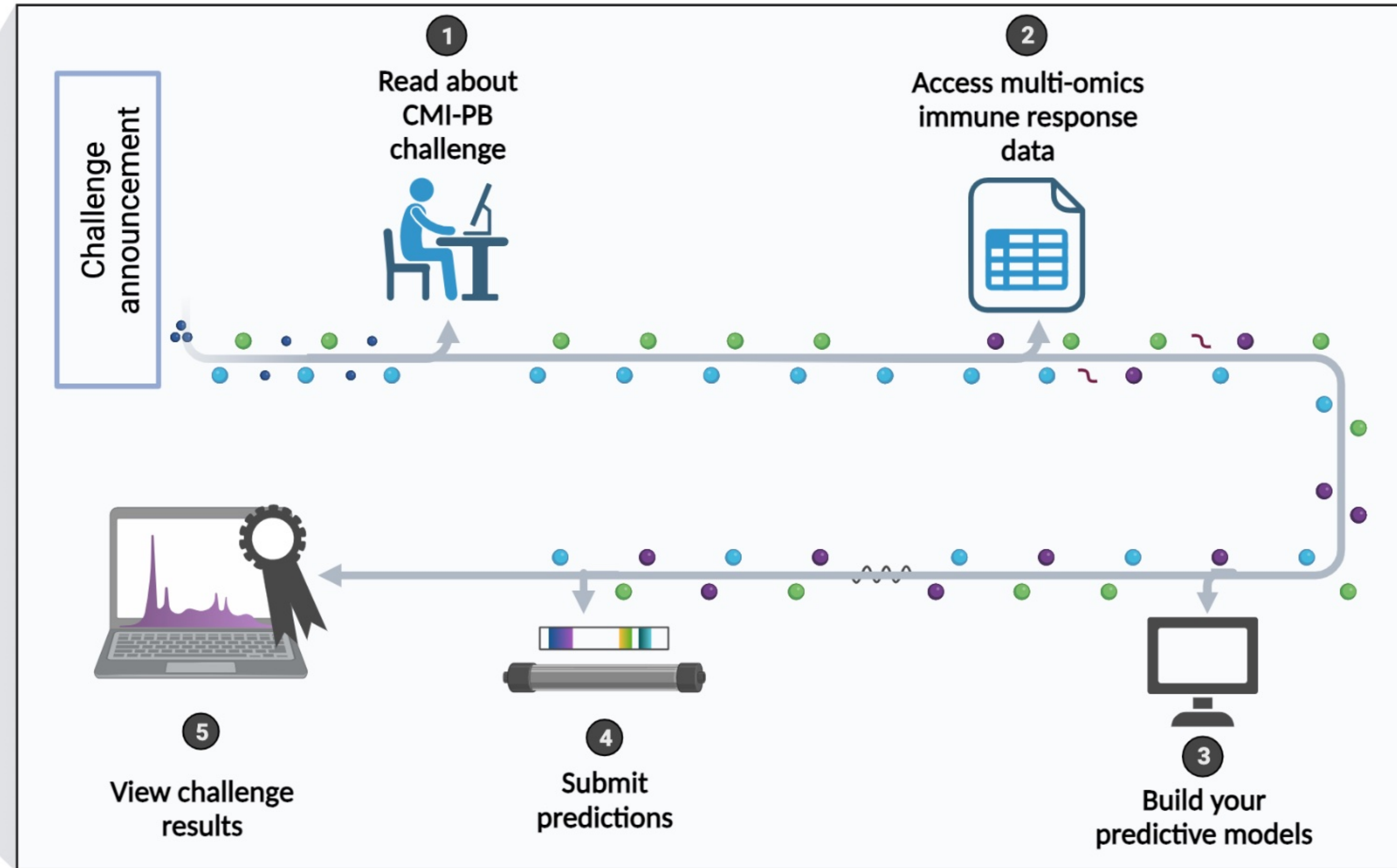
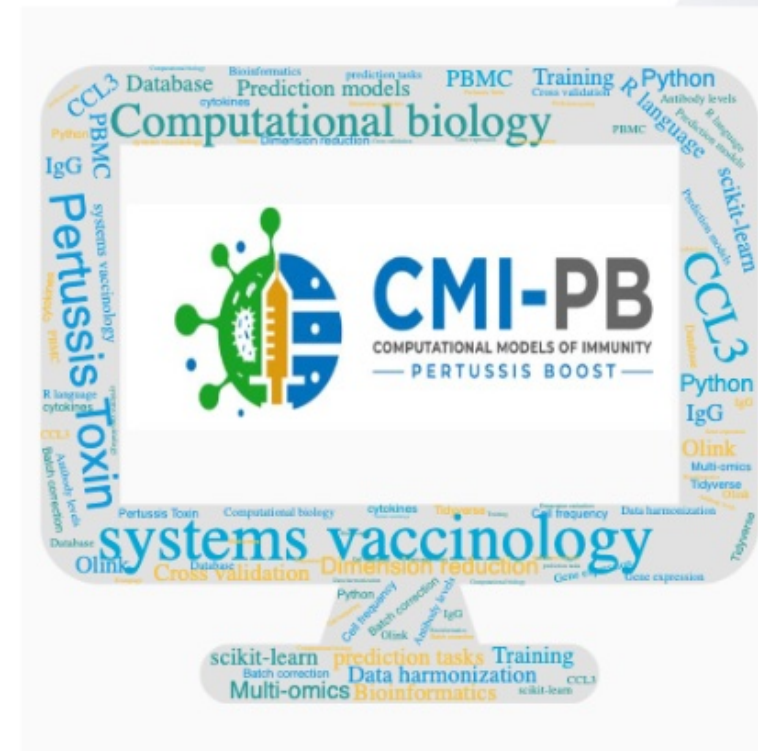
|   | Annual prediction challenge title          | Contestants         | Number of subjects  |                     | Current status                     |
|---|--|---------------------|---------------------|---------------------|------------------------------------|
|   |  |                     | Training dataset    | Test dataset        |                                    |
| 1 | <b>First Challenge:</b> Internal dry run   | CMI-PB consortium   | 60 (28 aP + 32 wP)  | 36 (19 aP + 17 wP)  | May 2022                           |
| 2 | <b>Second Challenge:</b> Invited challenge | Invited contestants | 96 (47 aP + 49 wP)  | 22 (13 aP + 9 wP)   | Announced on September 12, 2023    |
| 3 | <b>Third Challenge:</b> Open Challenge 1   | Public              | 118 (60 aP + 58 wP) | 32 (16 aP + 16 wP)  | Will be announced in April 2024    |
| 4 | <b>Fourth Challenge:</b> Open Challenge 2  | Public              | 150 (76 aP + 74 wP) | 32 (16 aP + 16 wP)* | Will be announced in December 2024 |

# b. Prediction challenge outline



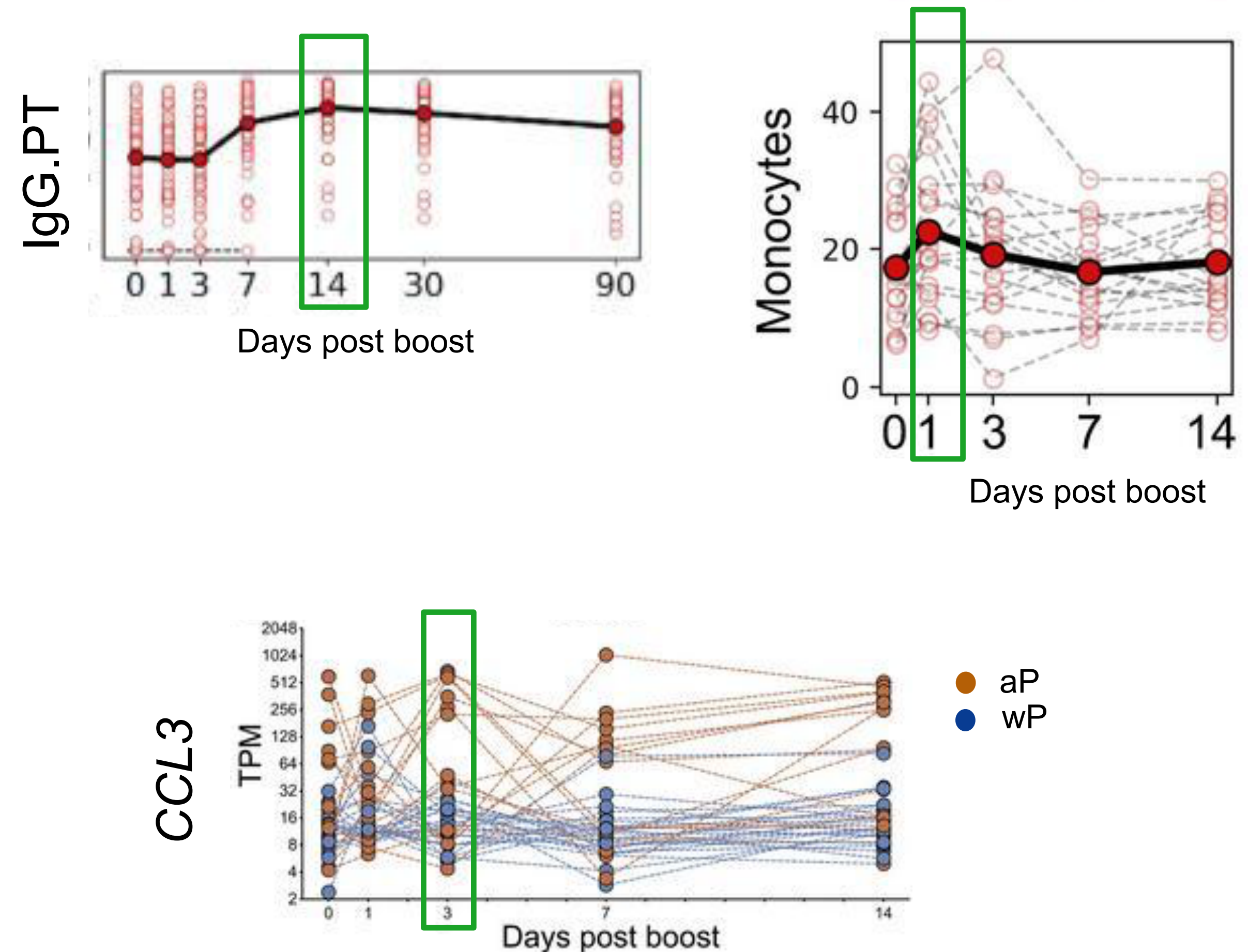
## 2nd CMI-PB Prediction Challenge Outline

*Revolutionizing computational modelling approaches for immune response prediction*



# c. Formulating prediction tasks for CMI-PB Challenge

- Previously identified **vaccine responses** are formulated as prediction tasks\*
- **General vaccine responses:**
  - Plasma IgG levels increased at day 14 post-booster vaccination compared to baseline
  - Increase in the percentage of monocytes on day 1 post-booster than baseline
- **aP/wP specific vaccine responses:**
  - A subset of aP-primed individuals showed an increased expression of proinflammatory genes, including CCL3 at day 3 post-booster vaccination



\* **A system-view of *Bordetella pertussis* booster vaccine responses in adults primed with whole-cell versus acellular vaccine in infancy**

Ricardo da Silva Antunes, ... , Alessandro Sette, Bjoern Peters

JCI Insight. 2021;6(7):e141023. <https://doi.org/10.1172/jci.insight.141023>.

# c. Formulating prediction tasks for CMI-PB Challenge



## List of tasks

### 1) Antibody titer tasks

1.1) Rank the individuals by IgG antibody titers against pertussis toxin (PT) that we detect in plasma 14 days post booster vaccinations.

predicted values

1.2) Rank the individuals by fold change of IgG antibody titers against pertussis toxin (PT) that we detect in plasma 14 days post booster vaccinations compared to titer values at day 0.

predicted fold-change values

### 2) Cell frequencies tasks

2.1) Rank the individuals by predicted frequency of Monocytes on day 1 post boost after vaccination.

2.2) Rank the individuals by fold change of predicted frequency of Monocytes on day 1 post booster vaccination compared to cell frequency values at day 0.

### 3) Gene expression tasks

3.1) Rank the individuals by predicted gene expression of CCL3 on day 3 post-booster vaccination.

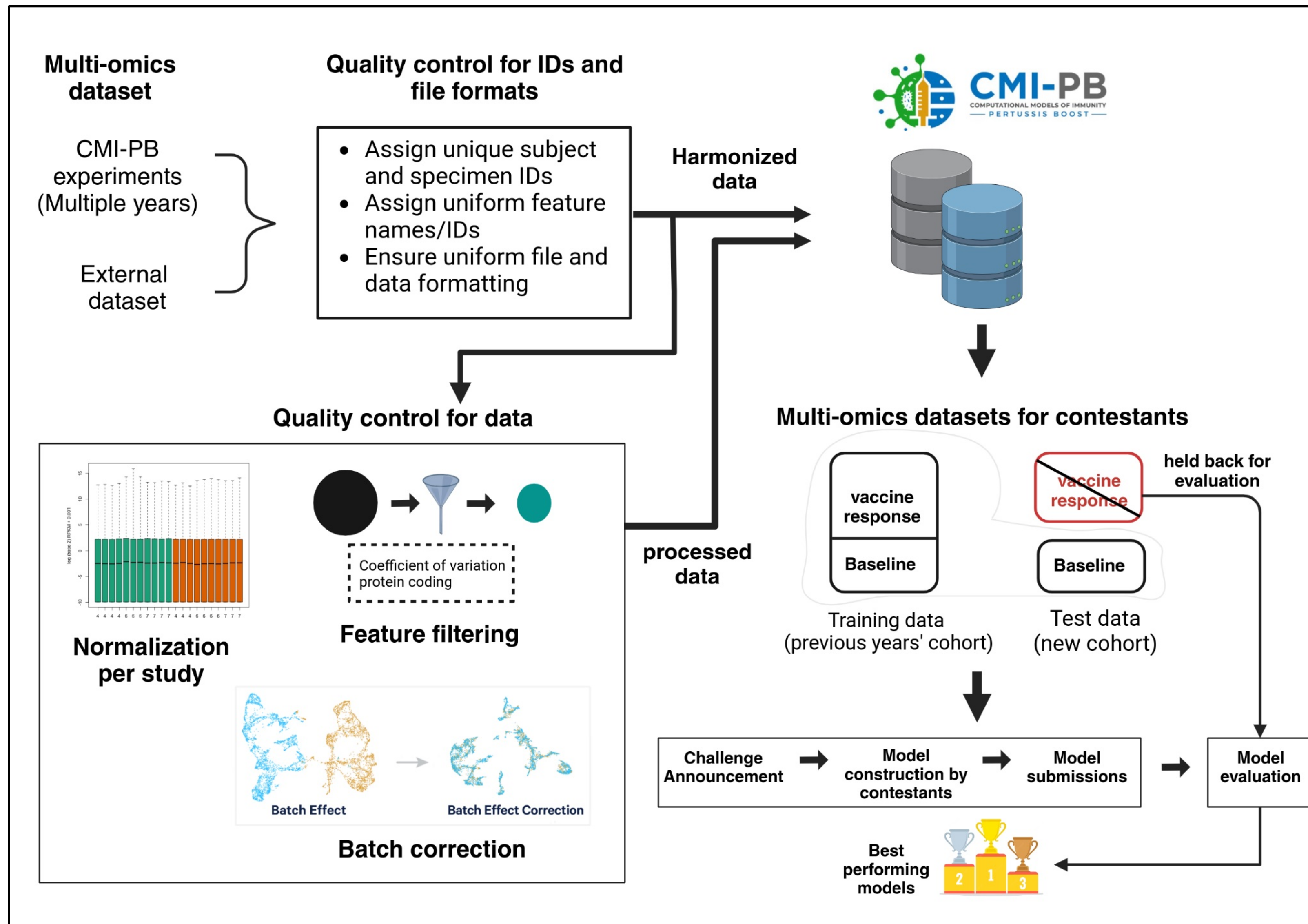
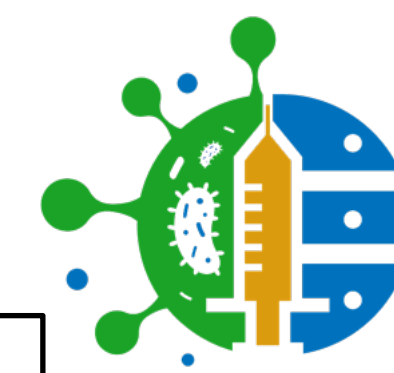
3.2) Rank the individuals by fold change of predicted gene expression of CCL3 on day 3 post booster vaccination compared to gene expression values at day 0.

## Example of Rankings

| Subject ID | Predicted value | Rank     |
|------------|-----------------|----------|
| 101        | 2.9             | 4        |
| 102        | <b>9.1</b>      | <b>1</b> |
| 103        | <b>1.2</b>      | <b>5</b> |
| 104        | 4.5             | 3        |
| 105        | 4.7             | 2        |

The ultimate goal is to model as many of the tasks as possible. However, contestants are not required to submit answers for all tasks.

# d. Overview of the CMI-PB Challenge data



Challenge related information and Data access is provided via the CMI-PB website





# CMI-PB PREDICTION CHALLENGE TIMELINE

  
Challenge begins  
**Sept 12th**

**Oct 6th**  
**1st**  
informational  
meeting




**Dec 1st**  
**2nd**  
informational  
meeting



**Jan 12th**  
**3rd**  
informational  
meeting



  
Final submission  
due date  
**Jan 16th**

Announcement of  
winners + longitudinal  
test data is released

**Feb 2nd**



2023

2024

# The CMI-PB team



## Kleinstein Lab (Yale)



- Expertise: A combination of "big data" analysis and immunology domain.
- Collaborating on data and models being released to the community to support reproducibility and the prediction contest, and also participate in the prediction challenge.

Steven Kleinstein  
Jeremy Gygi  
Leying Guan  
Anna Konstorum

## Grant Lab (UCSD)



- Expertise: the use of computational approaches, based on both biophysics and bioinformatics, to study the structure, function and evolution of key biological macromolecules.
- Dr. Grant will engage and advise over 40 biology graduate students in the CMI-PB Prediction Challenge.

Barry Grant

## Ay Lab (LJI)



- Expertise: Development of bioinformatics tools that utilize high-dimensional and high-throughput datasets to deduce insights into chromatin conformation, genetic variation, and the regulation of gene expression.
- The Ay lab is focused on developing predictive machine learning models, which will serve as examples and baselines for participants in the CMI-PB challenge.

Ferhat Ay  
Joaquin Reyna

## Peters Lab (LJI)



- Expertise: Both experimental and computational studies to better understand human immune responses in the context of infectious diseases, allergy, cancer and vaccines.
- The team is responsible for the generation of experimental data, making it accessible in a central and standardized fashion, and coordinating the creation and coordination of the prediction contest.

Bjoern Peters  
Jason Greenbaum  
James Overton  
Brendan Ha

Pramod Shinde  
Mari Kojima  
Rasteh Haji Kazem Nili

Jiyeun Lee  
Lisa Willemsen  
Shelby Orfield

And thank you to the Sette Lab, Crotty lab, LJI Clinical Core, LJI Bioinformatics Core

# The CMI-PB team members



Bjoern Peters



Steven Kleinstein



Ferhat Ay



Barry Grant



Shane Crotty



Alessandro Sette



Pramod Shinde



Shelby Orfield



Lisa Willemsen



Leying Guan



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Mari Kojima



Ferran Soldevila



Rasteh Nili



Jason Greenbaum



Brendan Ha



Jiyeun Lee



Ricardo De Silva Antunes



Jeremy Gygi



Anna Konstorum