

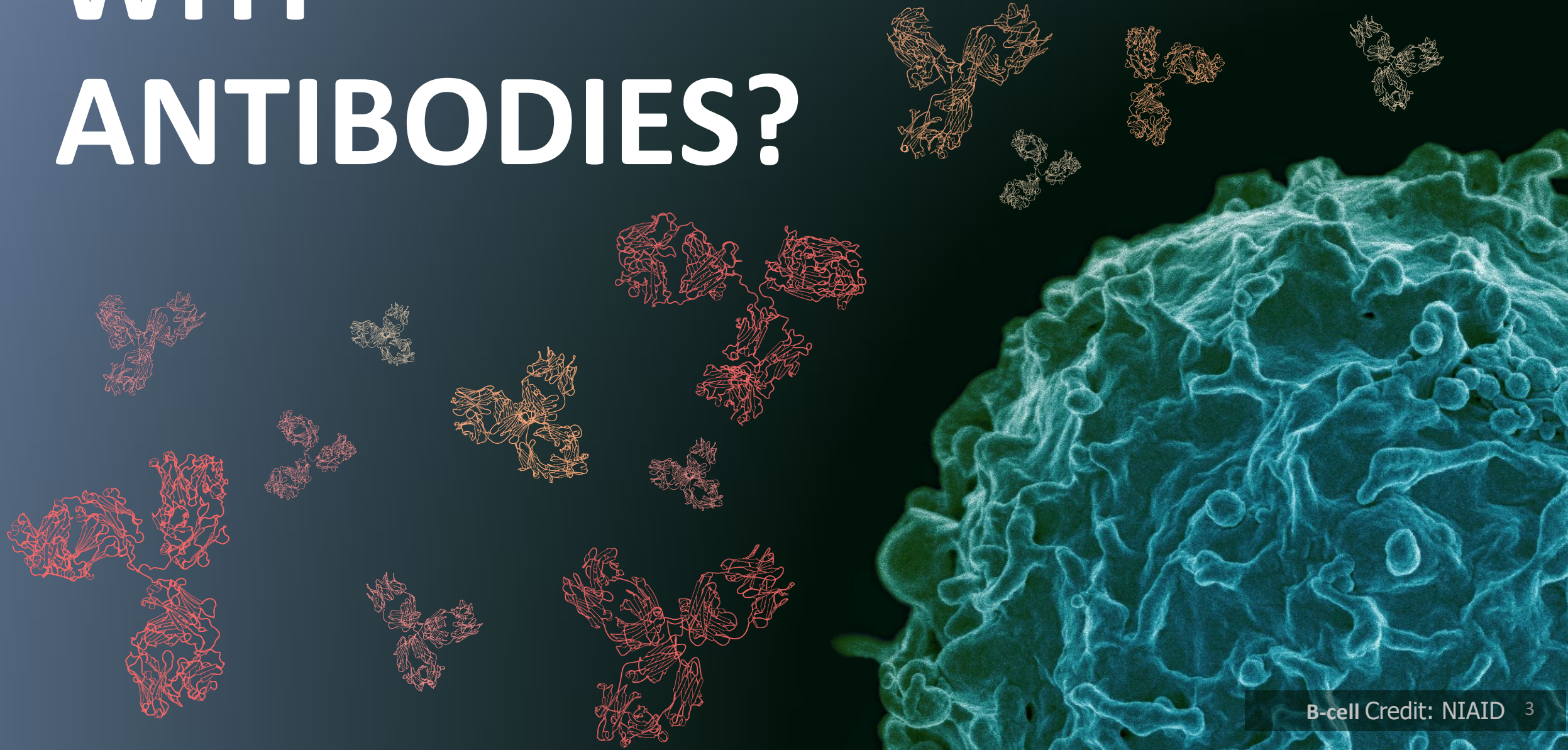
***Bio-Health:
Using AI to Engineer Human Antibodies***

Macromo**tek**
Revolutionizing antibody design

The logo for MacroMOTek features the word "Macro" in a bold, dark blue sans-serif font, followed by "mo" in a thin, red, lowercase sans-serif font, and "tek" in a thin, dark blue, lowercase sans-serif font. A red DNA double helix is positioned between the "mo" and "tek" parts of the logo.

**Monica Berrondo, PhD
CEO and Co-founder**

WHY ANTIBODIES?



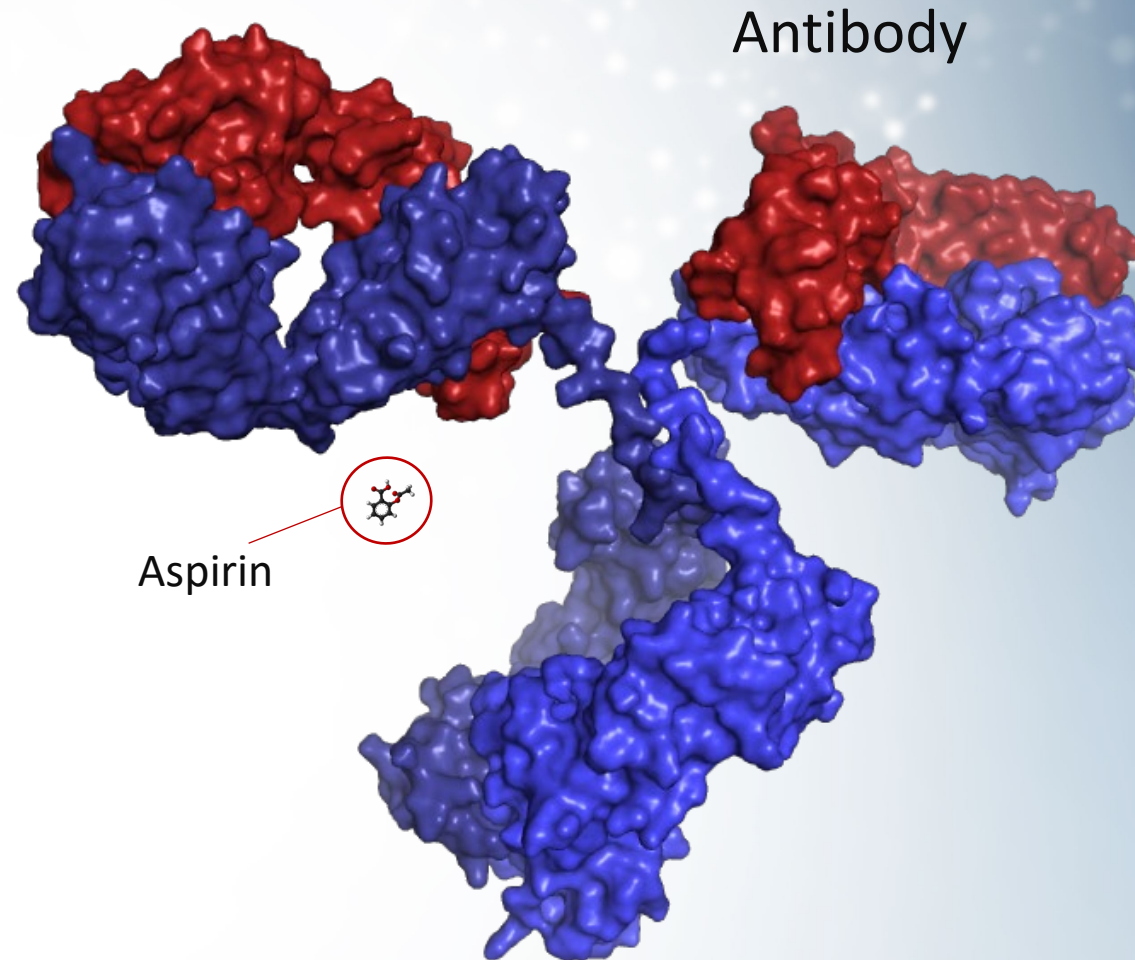
ANTIBODIES ARE PROVEN SUPERIOR DRUGS

Advantages of antibodies over small molecules

- Drugging “undruggable” targets
- Higher specificity and fewer side effects

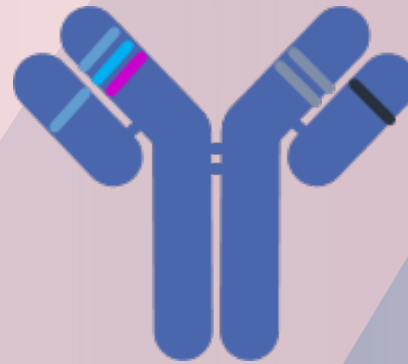
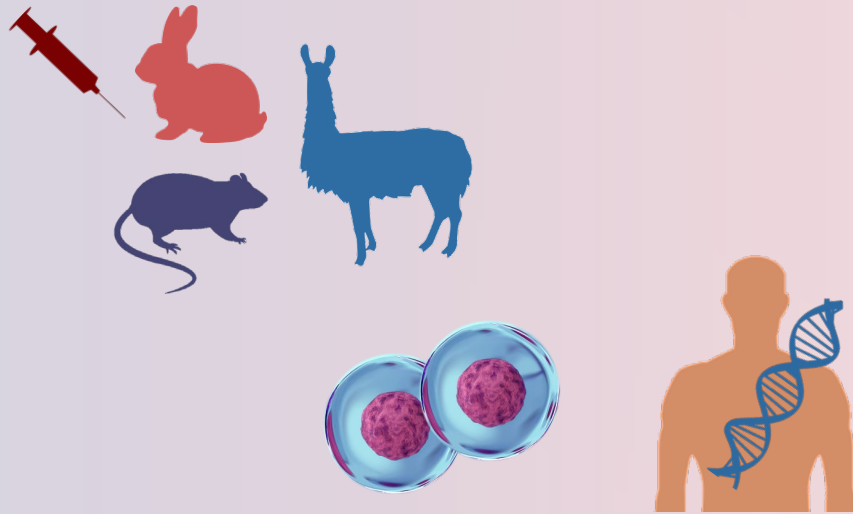
Clinical trial success rates

- 2X small molecule rate
- Highest of any drug modality

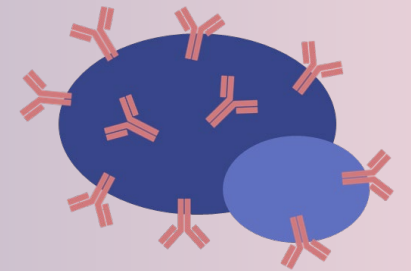
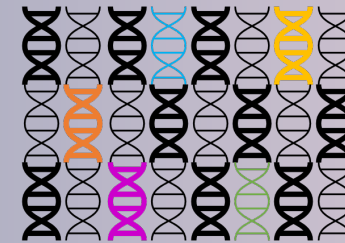


Current antibody discovery comes from two main sources:

Animal immunizations



Display Technologies



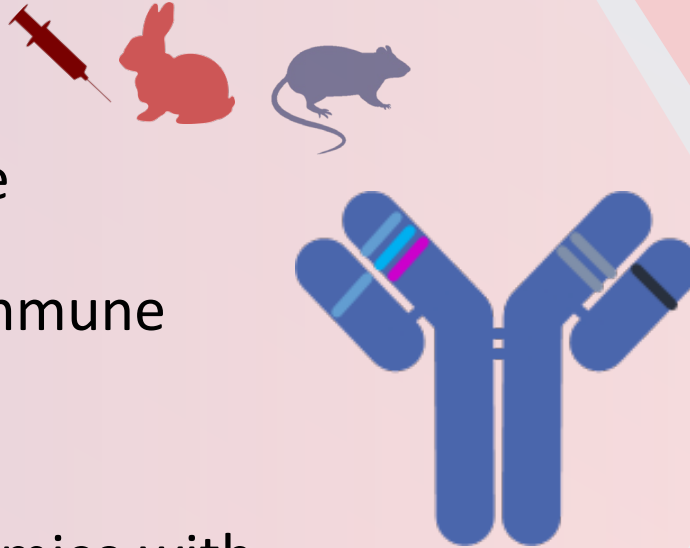
Pitfalls of animal derived antibodies

Animal immunizations

Time, labor, and capital intensive

Antibodies can cause negative immune response in humans

Extremely expensive to produce mice with human immune cells



Pitfalls of display derived antibodies

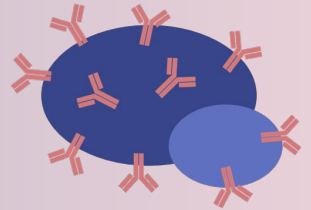


Display Technologies

Difficult to manufacture

Antibody diversity is limited by the cells in which they are produced

Further optimization necessary



DISCOVERY VS *DESIGN*

Discovery Platforms are unable to target specific mechanisms of action

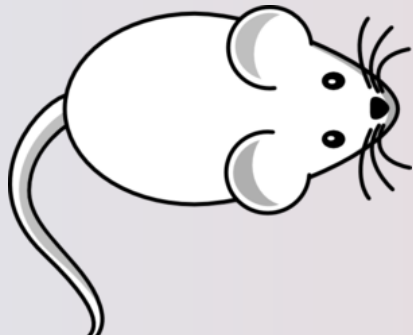
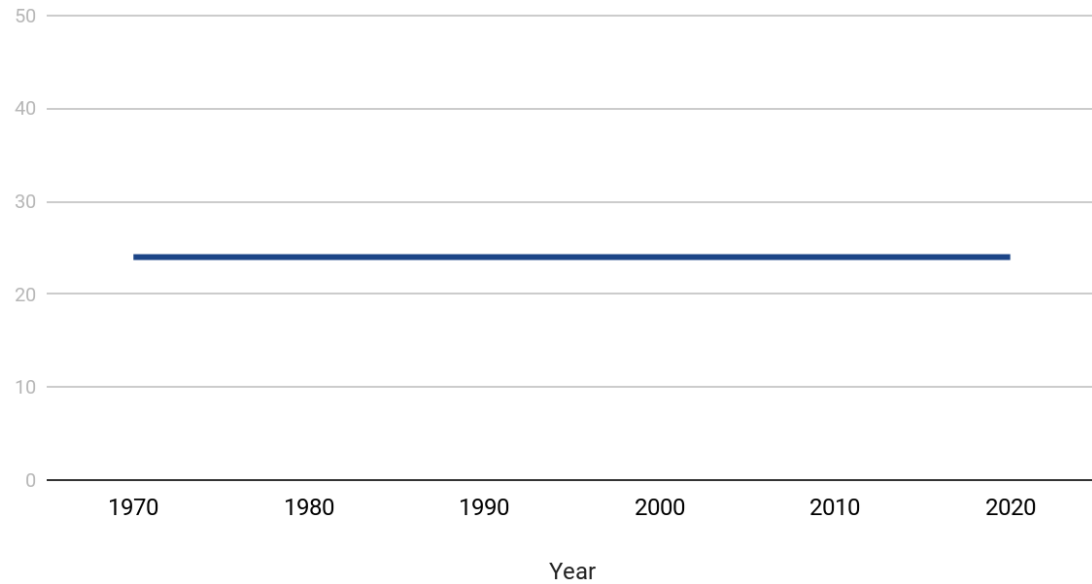
Current platforms blindly generate “hits” to be tested down the pipeline

How the antibody interacts with its target is identified after discovery

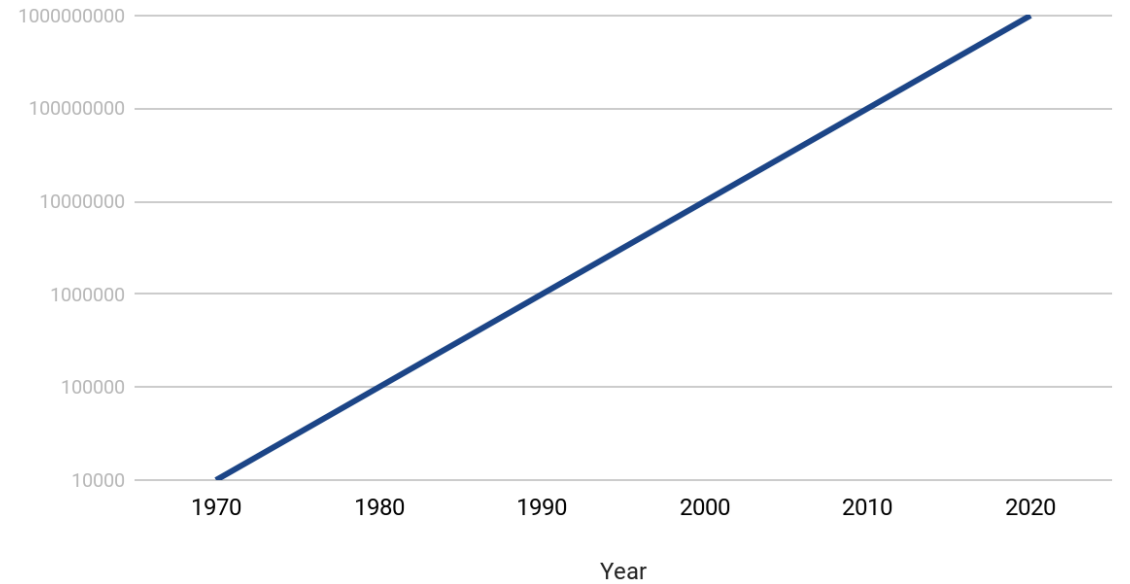
Our DESIGN software uses information about how the target causes disease
to create antibodies that directly address the problem.

Computational Drug Design – Huge Potential

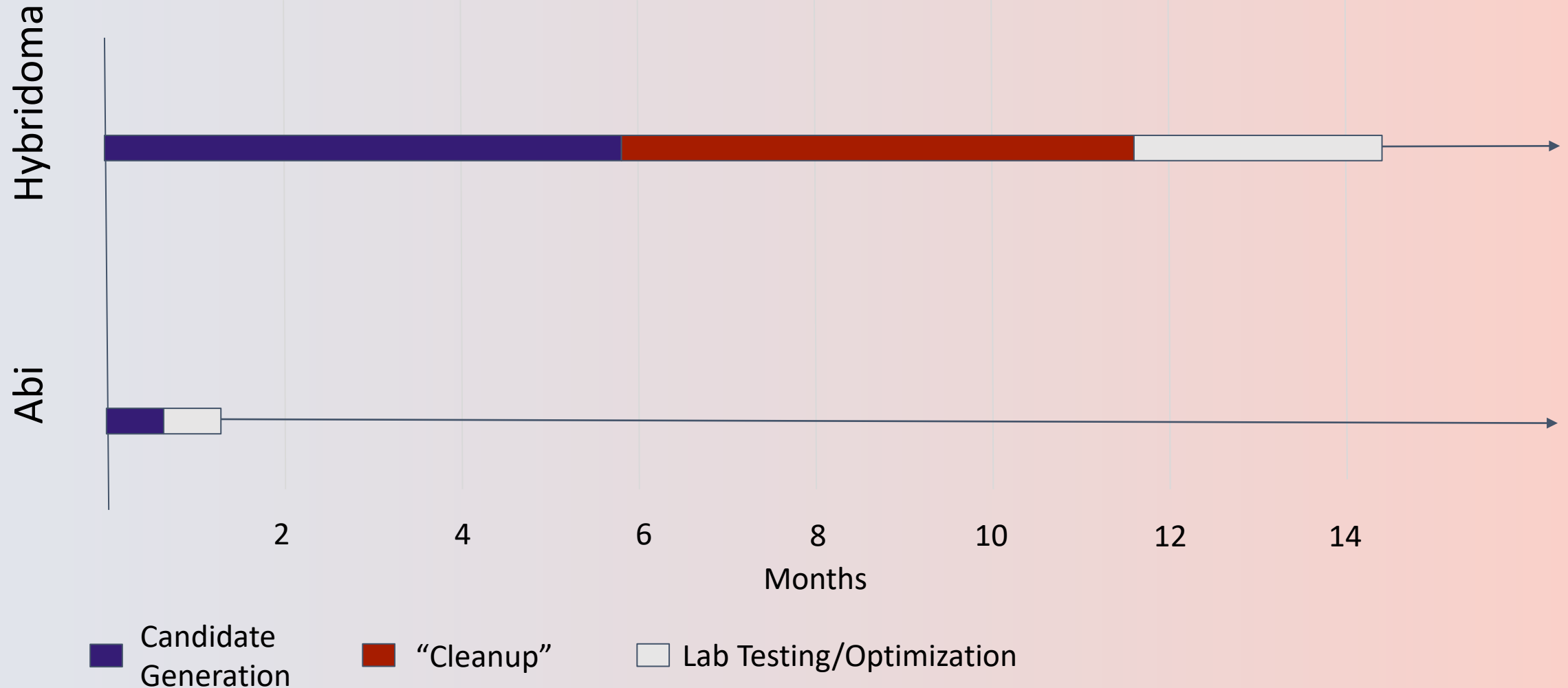
Antibody Processing Power - Mouse



Antibody Processing Power - Computational



Antibody Development Timeline

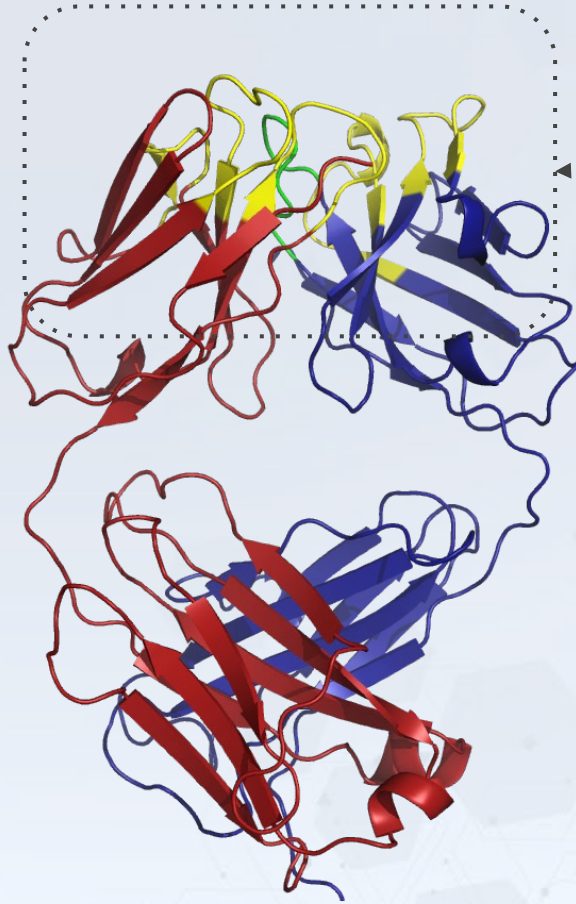


Computational Antibody Design

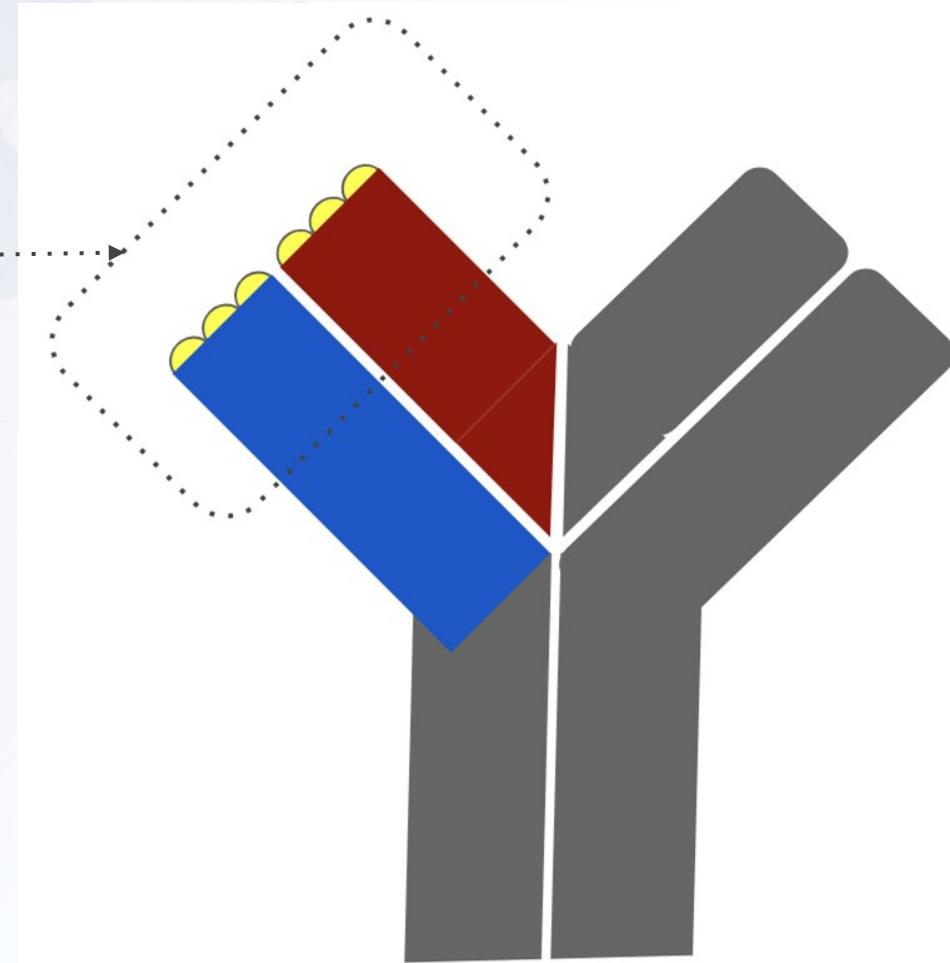
Advantages of Computational Antibody Design

- **Ability to target “undruggable” targets**
 - Inaccessible epitopes, alternate states, autoimmune targets
 - *In silico* design: target can be isolated and designed without interference from the system
- **Antibodies built on human scaffolds**
 - Less likely to be rejected than mouse antibodies
 - No need for extra humanization steps

What is an antibody?



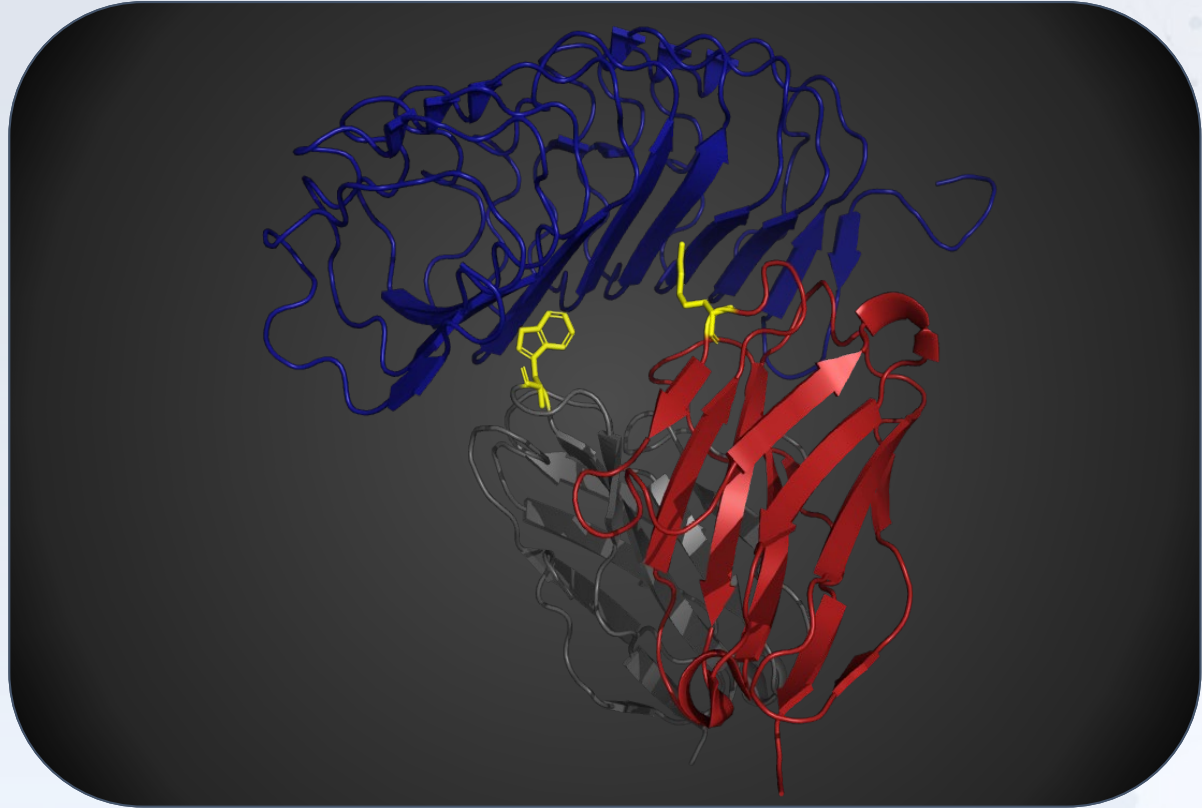
Special
binding
region



Modes of antibody design

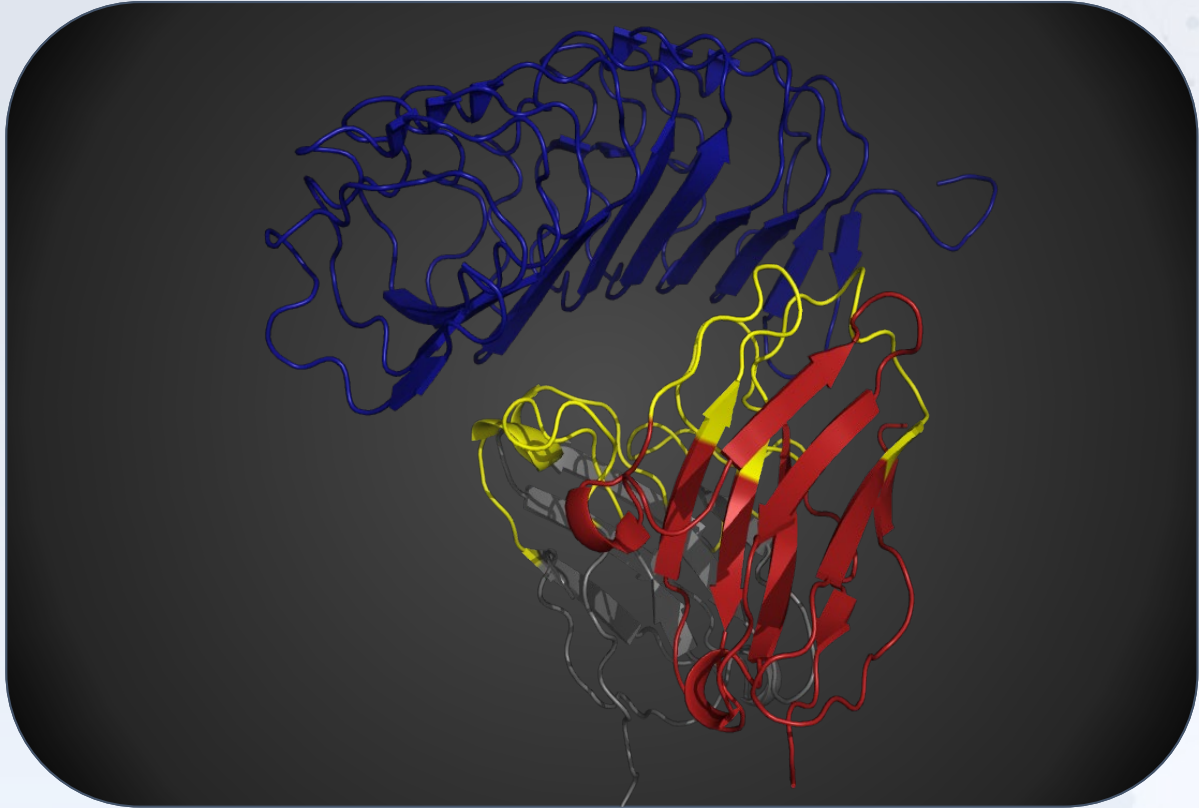
- **Most existing methods focus on improving the chemistry of an *existing* antibody-antigen complex**
- ***De novo* (from scratch) design is more complicated**
 - No starting-point antibody
 - Optimizes chemistry and topology simultaneously
- **New computational power and our proprietary software make it possible to do *de novo* antibody design**

Existing Computational Antibody Design Methods



- Series of individual amino acid changes to improve an existing antibody/antigen pairing
- Changes to make the antibody bind a similar epitope on another antigen
- High information requirement
 - Must know complete antibody and antigen complex structure

De novo Design Against a Selected Epitope



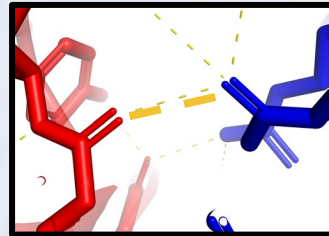
- “Holy Grail” of computational antibody design
- Design an antibody from scratch against a chosen epitope
- Low information requirement: all you need to start is an antigen structure
 - If a similar antigen exists, use homology modeling
 - In this case, only antigen sequence is needed

Key Considerations in Antibody-Antigen Interactions

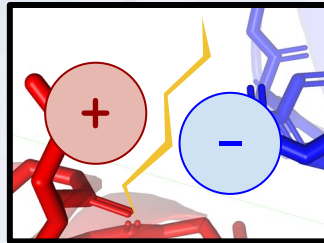
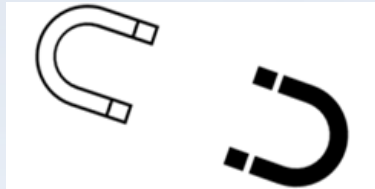
- **Paratope topology: Is its shape complementary to that of the epitope?**
- **Chemistry**
 - Polar interactions: Do the paired amino acids attract or repel each other?
 - Hydrophobics: Are water-soluble amino acids exposed and water-repelling residues hidden?
 - Hydrogen bond formation: Structures which form more hydrogen bonds are more stable.
- **One change can affect everything!**
 - Altering an amino acid to improve chemical interactions might deform the paratope structure
 - Changes in structure can disrupt chemical interactions

The Biochemistry of Antibodies

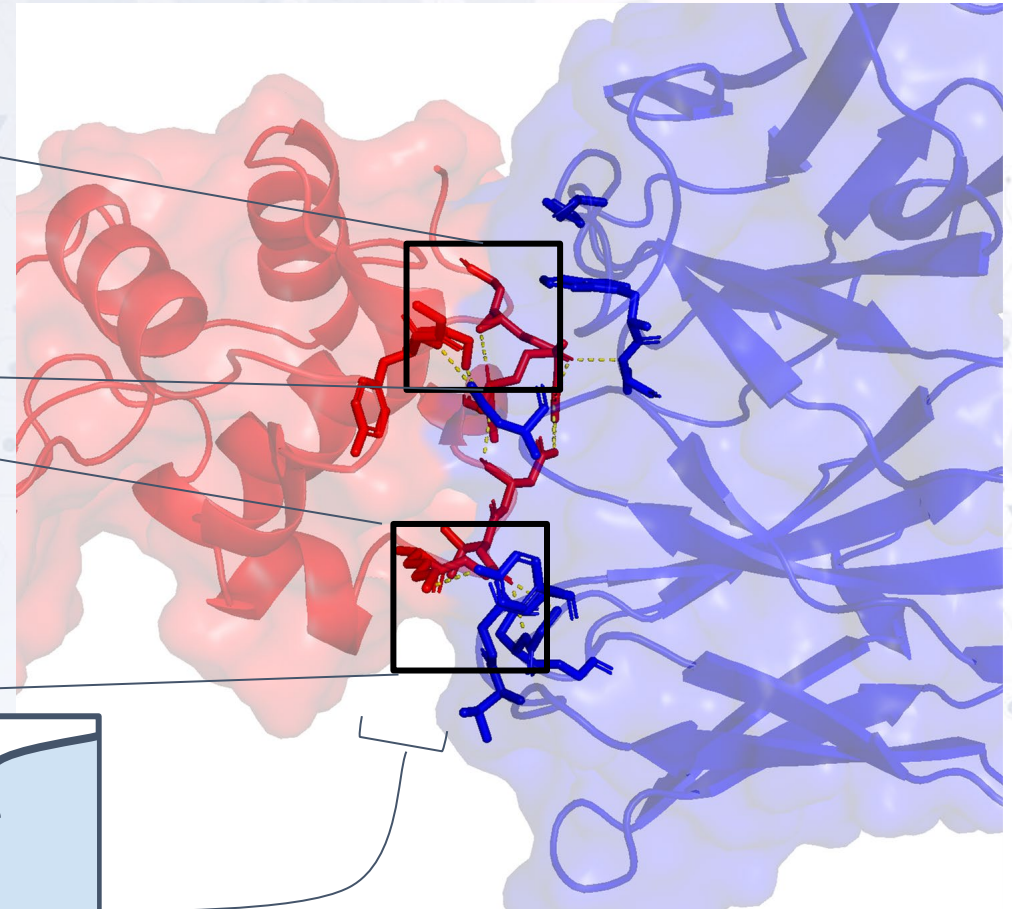
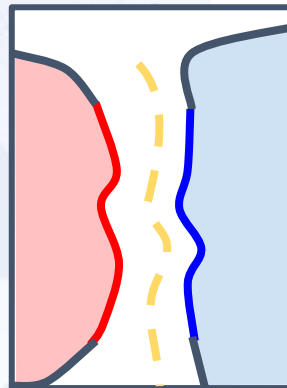
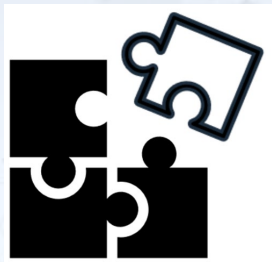
1. Hydrogen bonding



2. Charge interactions

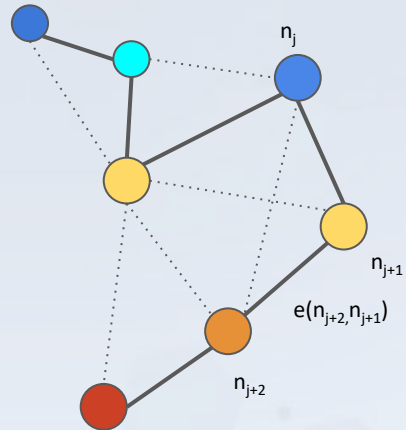


3. Shape complementarity

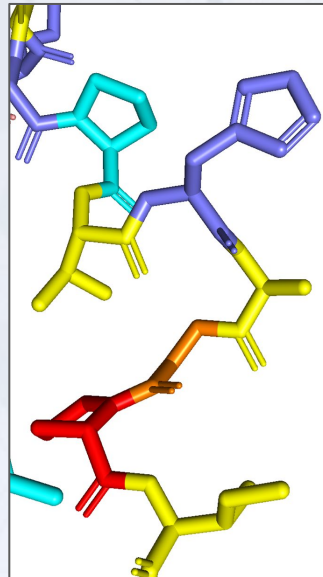


Data Representation

Graph



Protein Structure



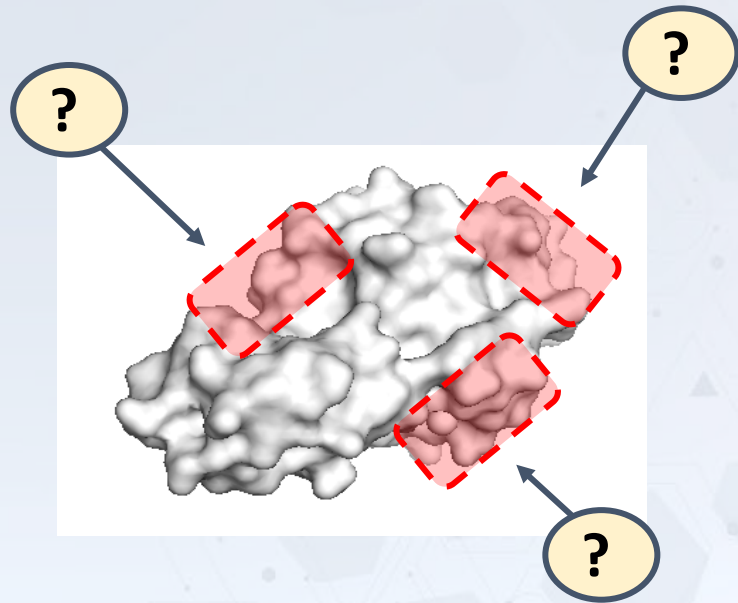
Voxel



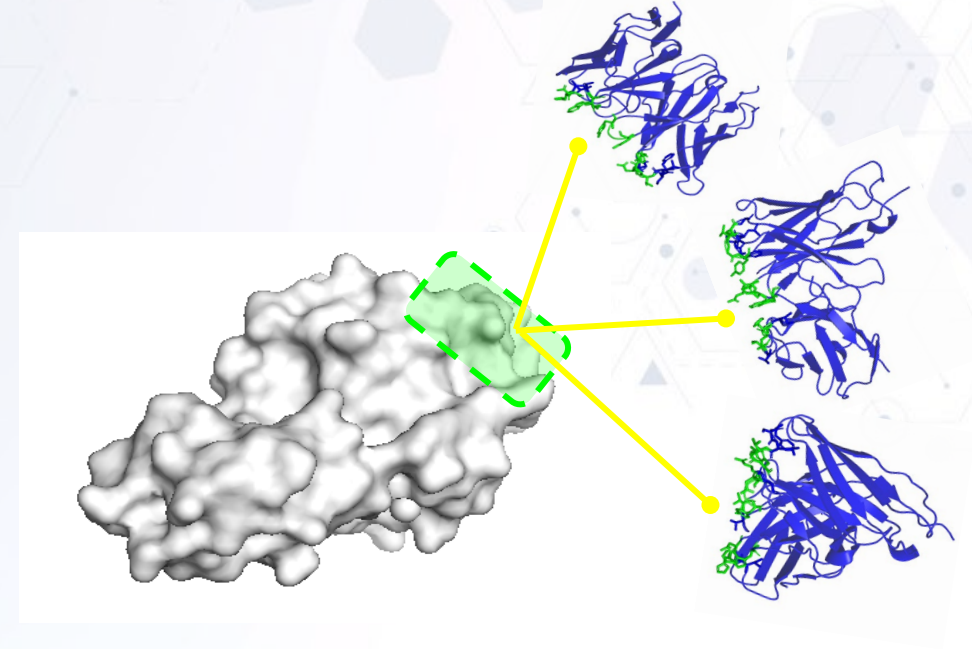
- Transforms into a complex computer vision problem
- Allows us to use build powerful convolutional neural networks
- GPUs are needed to train and infer using these representations

Problems in Computational Design

Where do we target our designs?

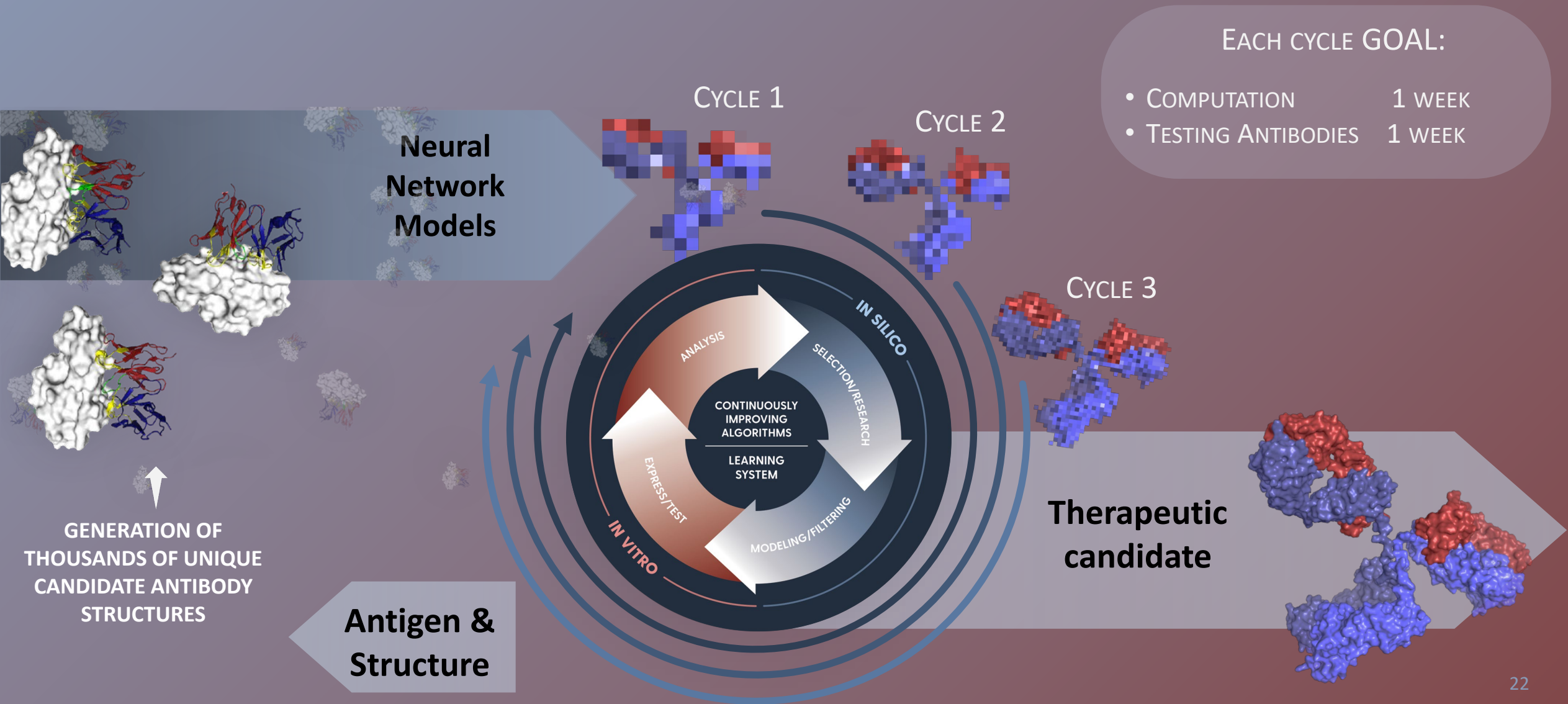


Which designs will work?

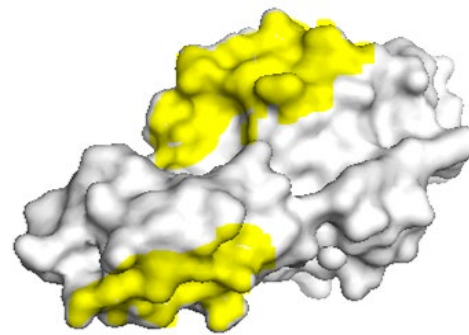
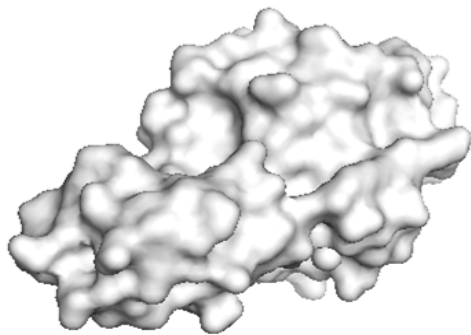


Computational Process

ANTIBODY DESIGN PROCESS TAPS INTO THE FUTURE



STEP 1 - TARGET SELECTION/IDENTIFICATION



TARGET
STRUCTURE

NEURAL
NETWORK
*Proprietary IP

PROBABILITY MAP
OF BEST REGIONS
TO TARGET

*SELECTED REGIONS
SENT TO DESIGN
ALGORITHM*

STEP 2 - RUN DESIGN ALGORITHM

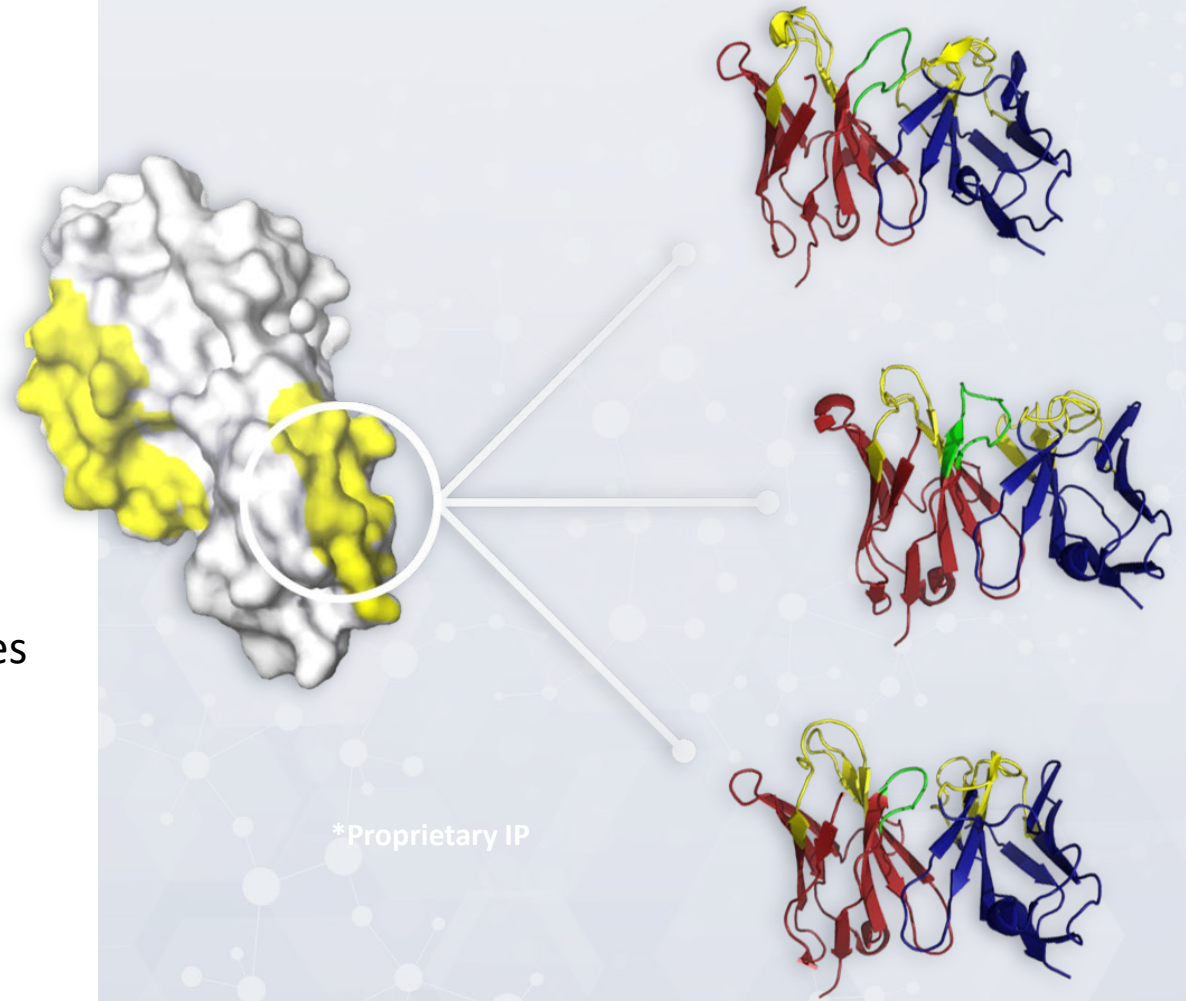
Antibody specific algorithm

Antigen structure is fed into design algorithm

Template antibody is generated

Determines sequence of non-binding antibody regions

Algorithm generates large pool of predicted candidates with unique binding sequence



STEP 3 – DESIGN SELECTION



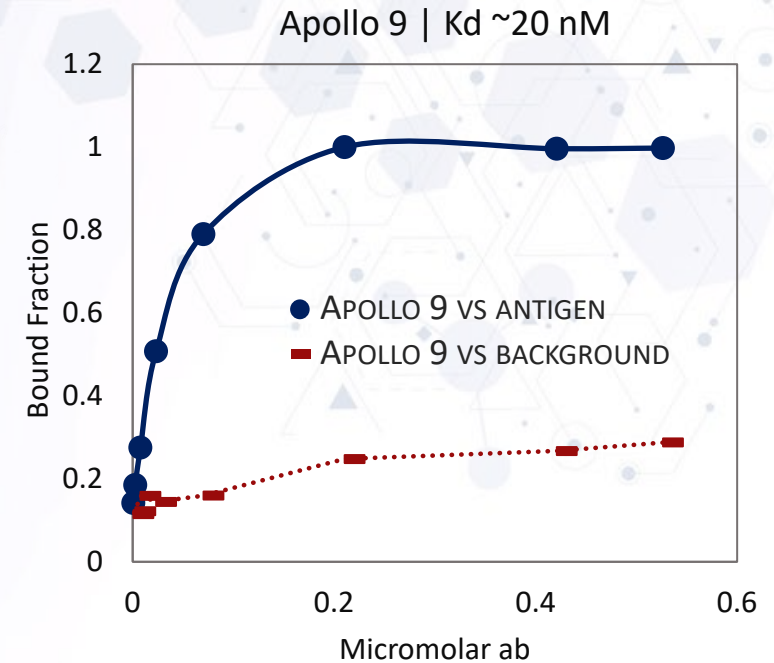
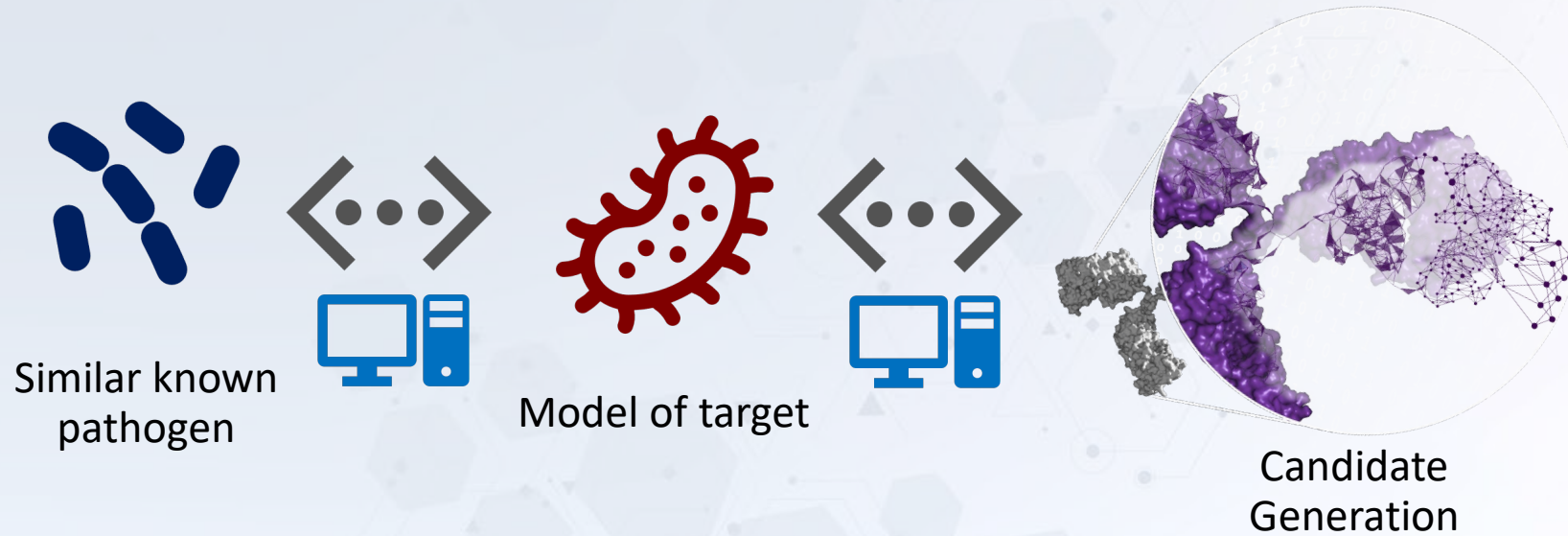
Case Studies

Case Study 1: APOLLO

3 Designs undergoing
biophysical
characterization

TARGETING A **MEMBRANE PROTEIN** OF A **MULTI-DRUG RESISTANT PATHOGENIC BACTERIA**

UNDERGOING **BIOPHYSICAL CHARACTERIZATION** ANTIGEN-ANTIBODY COMPLEX



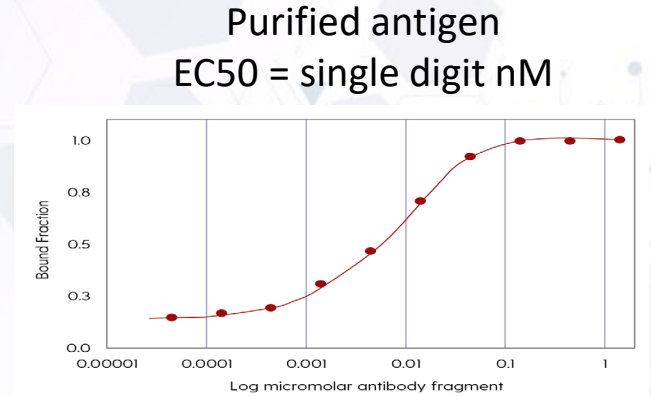
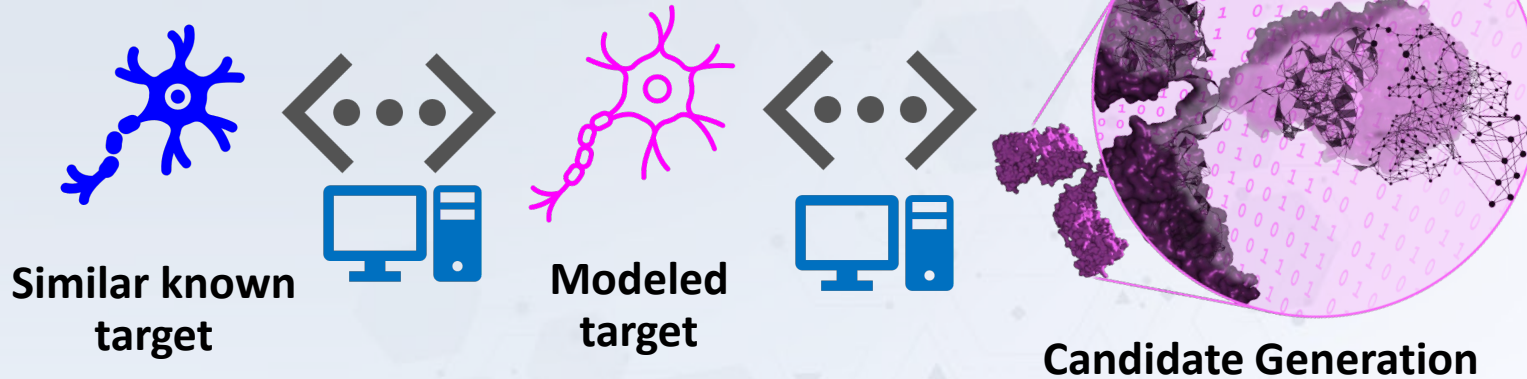
POTENT ANTIBODY DESIGNED USING MODELED TARGET

Case Study 2: SARASWATI

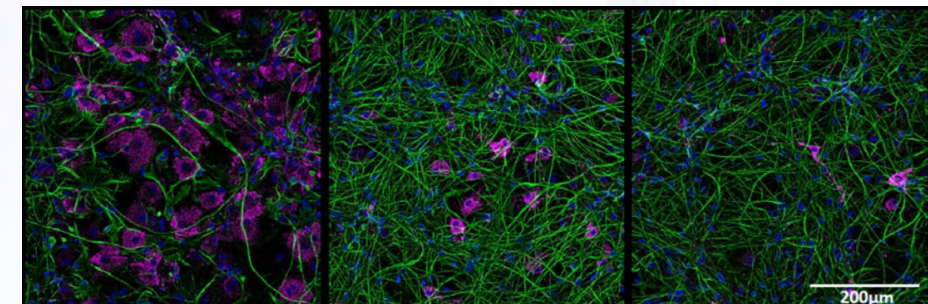
30nM antibody
Designed without preexisting
crystal structure

POTENT ANTIBODY

Tested in tissue culture and progressing toward animal studies



Cell binding at EC50 ~60 nM



Nuclei, Design Antibody, Cells

- Created a model for **understudied disease target**
- Antibody designed **without any pre-existing structural data**
- Currently **undergoing pre-clinical testing in human cells**
- Computational antibody targeting a **neurodegenerative disease**

Case Study 3: OSIRIS

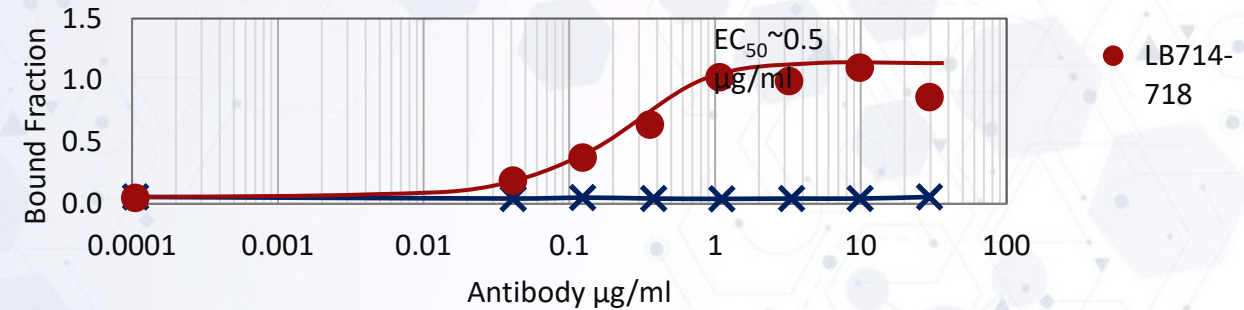
Antibody specifically binding to antigen generated in 2 weeks

**SUCCESSFUL BINDING TO ANTIGEN,
PREVIOUSLY NOT POSSIBLE!**

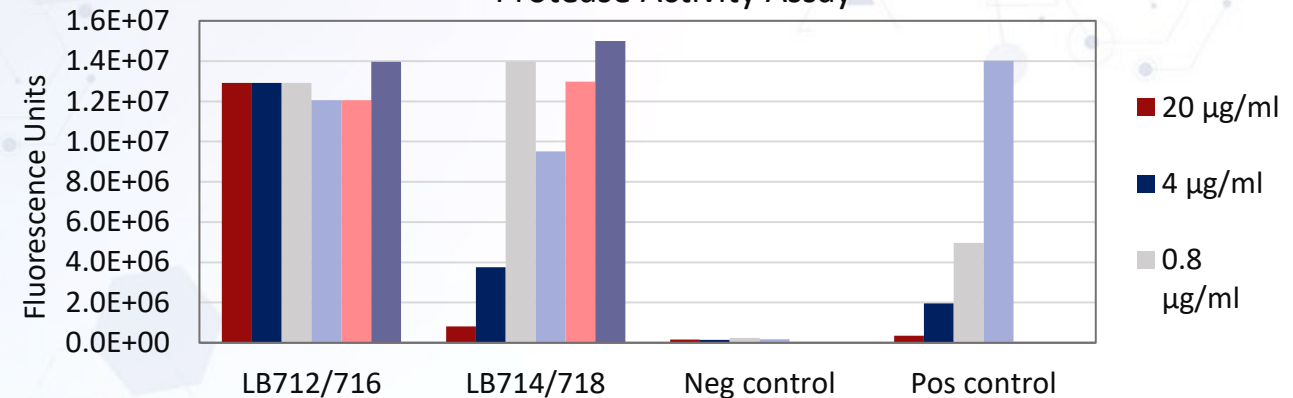
Protease involved
in neonatal skin disease

Biotech partner failed to generate
antibodies with mice or phage display

ELISA binding assay



Protease Activity Assay



WHAT'S NEXT IN AI DIRECTED ANTIBODY DESIGN?

- Increased speed of antibody generation
- Rational binding affinity design *de novo*
- Rational agonism/antagonism design *de novo*
- De-risking immunogenicity and safety issues*
- Harnessing Generative AI
- Bringing order to disorder → antibodies to floppy loops (GPCRs et al.)
- Improving hit rate of library



Macromol**tek**
Revolutionizing antibody design

monica@macromoltek.com

www.macromoltek.com