



**ALCHEMAB**  
THERAPEUTICS

# Parallel Discovery of Therapeutic Antibodies and Novel Targets Using the Antibody Repertoires of Resilient Individuals

PEGS Boston – May 4<sup>th</sup> 2022

# Our vision: To use the power of the human immune system to discover new medicines and diagnostics

Discovering and developing protective, patient-originated therapeutic antibodies



Focus on protective  
antibody responses



Convergent in  
resilient individuals

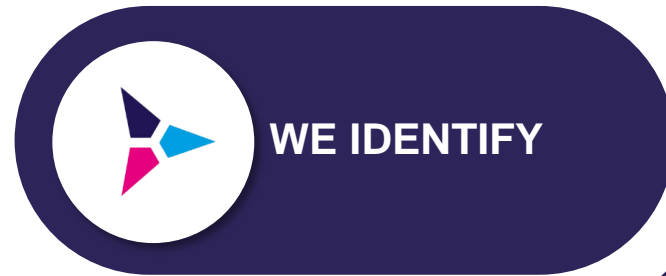


Target agnostic  
approach



Advanced  
computational  
approaches

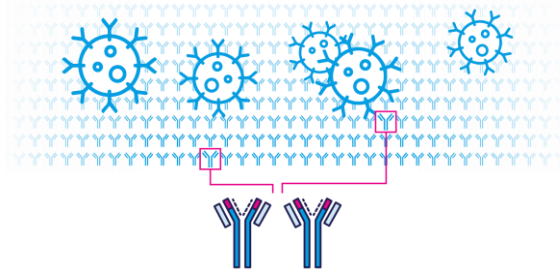
# Alchemab concept



We identify especially resilient individuals – and learn how they overcome or resist disease



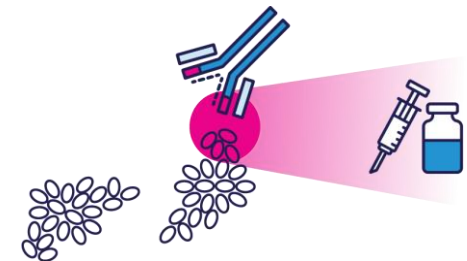
We sequence B cells from the resilient individuals and identify antibodies with similar properties



Antibodies



We discover the binding targets of the antibodies, understand their protective properties and develop candidates that replicate the protective effect



Novel Antibody Therapeutic

**Unbiased platform to identify novel therapeutics**

# Patient-originated, physiologically validated therapeutics



**Our therapeutic antibodies are physiologically validated,  
increasing the likelihood that the  
targets are critical disease modifiers**

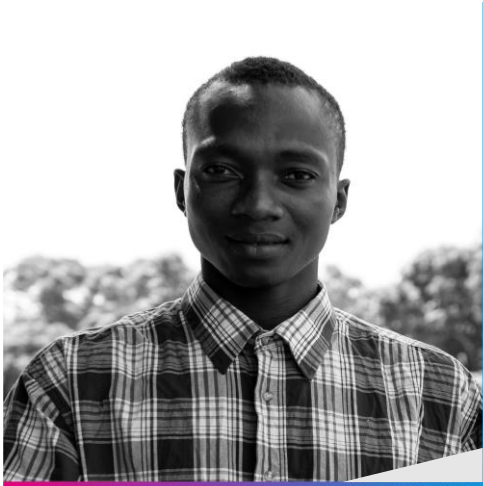
# How do we think about resilience?



Patients who  
survive  
advanced cancer



Patients progressing  
unusually slowly with  
neuro-degenerative disorders



People who survive  
grievous, deadly  
infectious disease



Long-lived,  
healthy individuals

# Humans benefit from naturally-occurring protective autoantibodies in many diseases

SCIENCE TRANSLATIONAL MEDICINE | RESEARCH ARTICLE

## A human-derived antibody targets misfolded SOD1 and ameliorates motor symptoms in mouse models of amyotrophic lateral sclerosis

Marcel Maier<sup>1\*</sup>, Tobias Welt<sup>2\*</sup>, Fabian Wirth<sup>1</sup>, Fabio Montrasio<sup>1</sup>, Daniel Preisig<sup>2</sup>, Jordan McAfoose<sup>2</sup>, Fernando G. Vieira<sup>3</sup>, Luka Kulic<sup>2</sup>, Claudia Späni<sup>2</sup>, Thilo Stehle<sup>4</sup>, Steve Perrin<sup>3</sup>, Markus Weber<sup>5</sup>, Christoph Hock<sup>1,2</sup>, Roger M. Nitsch<sup>1,2</sup>, Jan Grimm<sup>1†</sup>

## Cancer Cell

Article

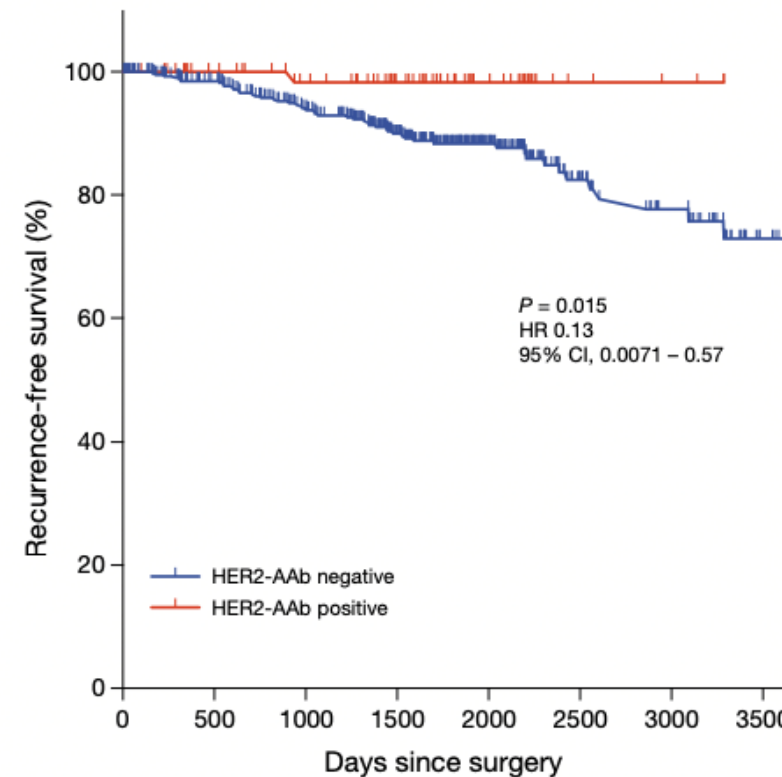
### Intratumoral plasma cells predict outcomes to PD-L1 blockade in non-small cell lung cancer

- Three populations of intratumoral B and plasma cells identified by scRNA-seq in NSCLC
- Plasma cells show the strongest predictive association with overall survival to PD-L1 blockade
- Plasma cell benefits are independent of intratumoral CD8 T cells and PD-L1 expression
- B and plasma cells are present in tertiary lymphoid structures in NSCLC tumors

Breast Cancer Res Treat (2016) 157:55–63  
DOI 10.1007/s10549-016-3801-4

### Protective effect of naturally occurring anti-HER2 autoantibodies on breast cancer

Yukiko Tabuchi<sup>1</sup> · Masafumi Shimoda<sup>1</sup> · Naofumi Kagara<sup>1</sup> · Yasuto Naoi<sup>1</sup> · Tomonori Tanei<sup>1</sup> · Atsushi Shimomura<sup>1</sup> · Kenzo Shimazu<sup>1</sup> · Seung Jin Kim<sup>1</sup> · Shinzaburo Noguchi<sup>1</sup>



# We collaborate with a broad and growing network of institutions



## Neurodegeneration



Accelerating therapeutic development  
for Huntington's disease



## Oncology



UNIVERSITY OF  
OXFORD



CANCER  
RESEARCH  
UK



## Infectious Disease and other areas



Barts Health  
NHS Trust



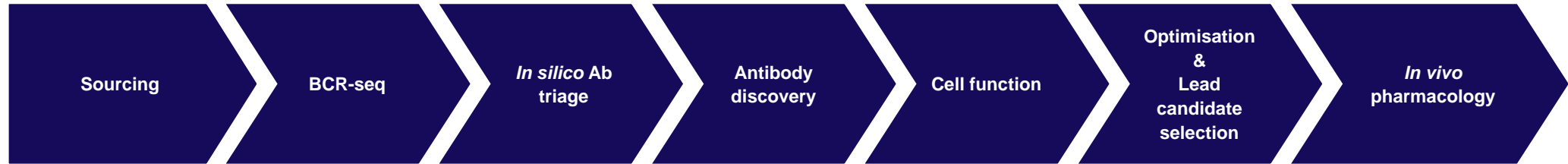
UNIVERSITY OF  
OXFORD



Platform enabling collaborations

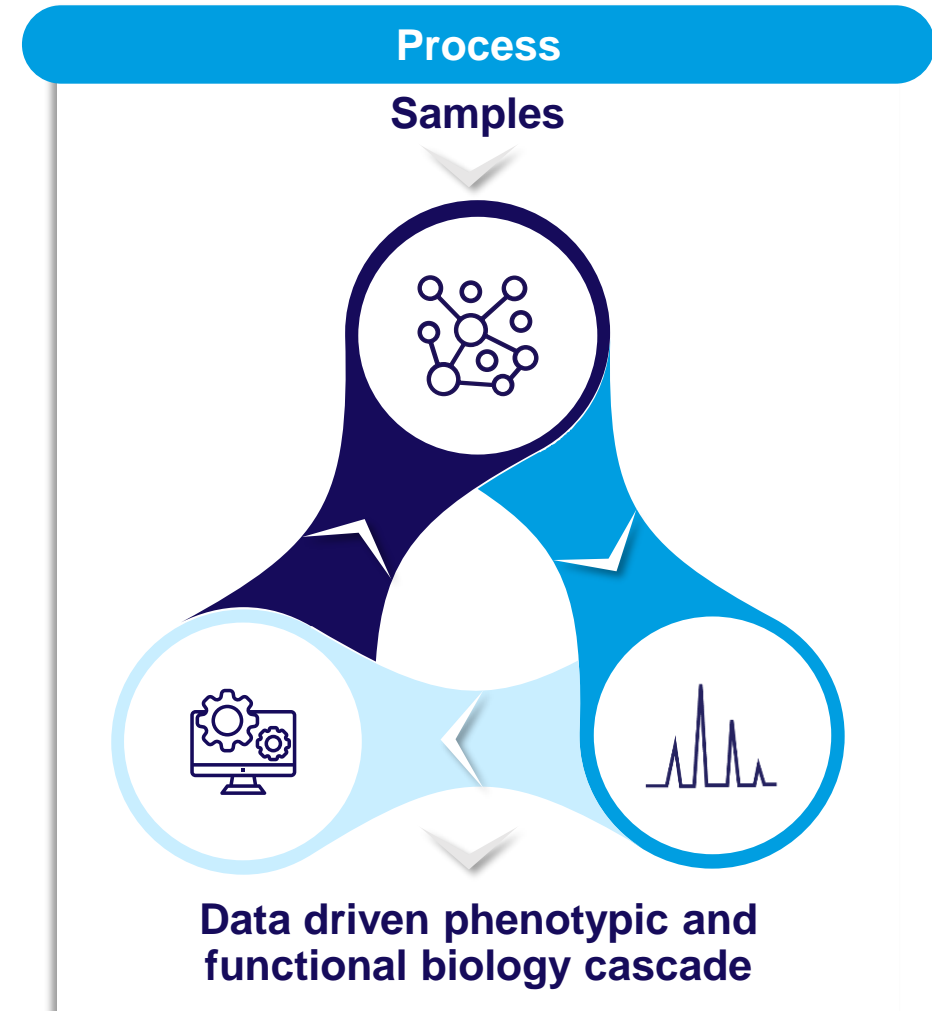
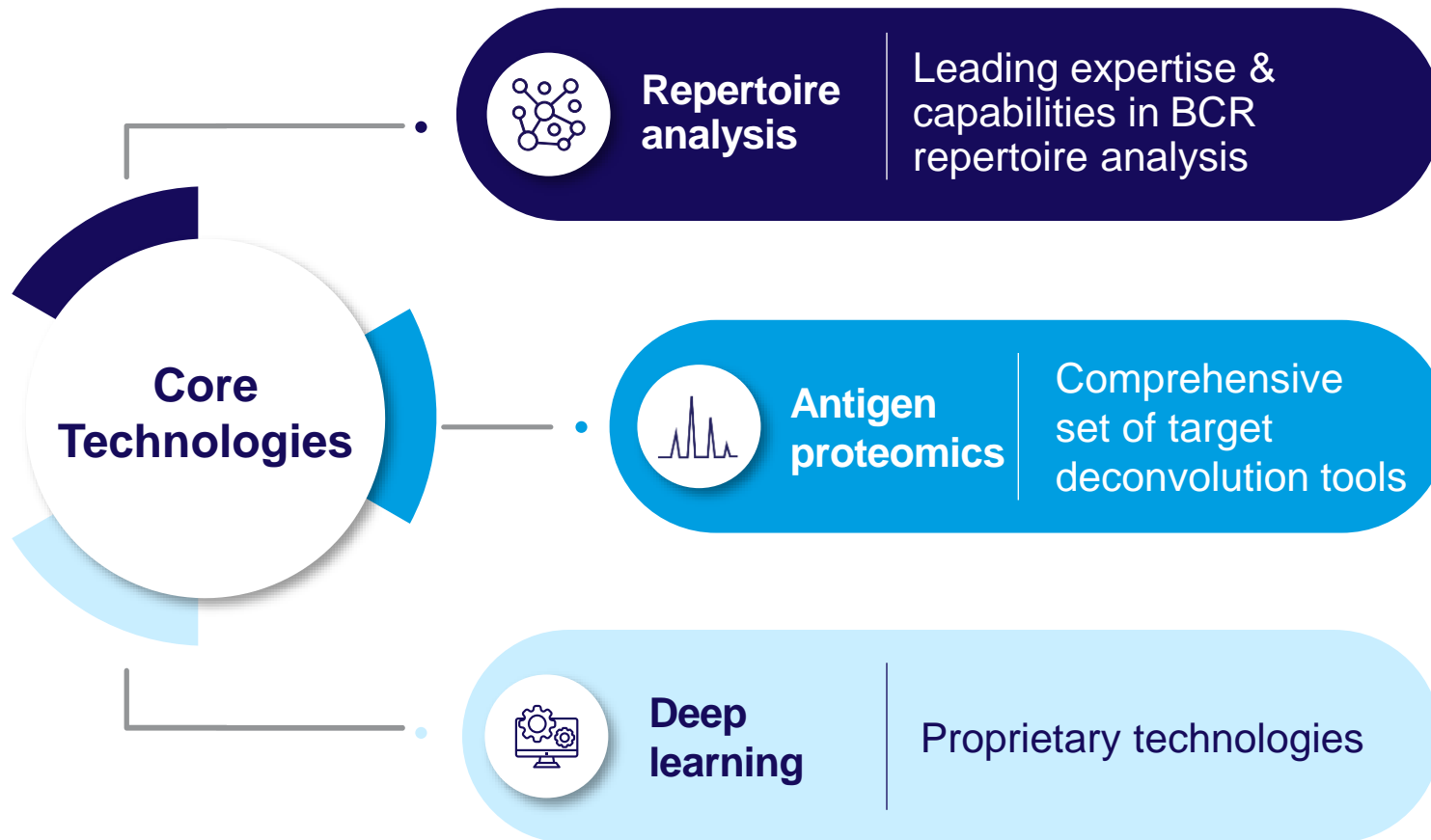


# Our platform has generated numerous program opportunities across a variety of diseases



How do we do what we do?

# Discovery process powered by advanced sequencing, big data, and deep learning



# Convergent autoantibody sequences provide the starting point for discovery

## Population B Cell Repertoire Analysis

Sequences  
of disease-  
free  
population

Sequences  
of resilient  
patients

Sequences  
of disease  
progressors

10's-100's of Individuals;  
Millions of BCR sequences

### Convergence

Shared antibodies that  
do not occur in controls  
or progressors



Convergence reflects similar antibodies that are shared among members of a cohort and directed against a common antigen

Convergence is rare, and therefore meaningful

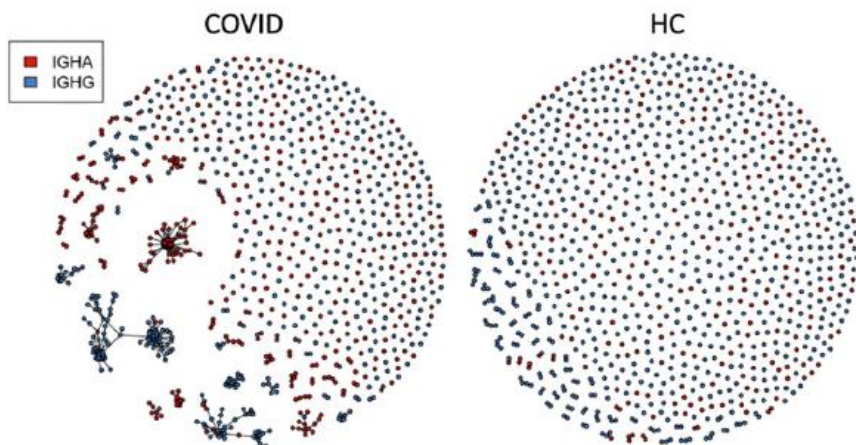
The result is target and pathway agnostic therapeutics

We have a unique deep learning enabled process to express functional antibodies

# Covid-19 study provides proof-of-concept

## Deep Sequencing of B Cell Receptor Repertoires From COVID-19 Patients Reveals Strong Convergent Immune Signatures

Jacob D. Galson<sup>1\*</sup>, Sebastian Schaetzle<sup>1</sup>, Rachael J. M. Bashford-Rogers<sup>1,2</sup>, Matthew I. J. Raybould<sup>3</sup>, Aleksandr Kovaltsuk<sup>3</sup>, Gavin J. Kilpatrick<sup>1</sup>, Ralph Minter<sup>1</sup>, Donna K. Finch<sup>1</sup>, Jorge Dias<sup>1</sup>, Louisa K. James<sup>4</sup>, Gavin Thomas<sup>4</sup>, Wing-Yiu Jason Lee<sup>4</sup>, Jason Betley<sup>5</sup>, Olivia Cavlan<sup>1</sup>, Alex Leech<sup>1</sup>, Charlotte M. Deane<sup>3</sup>, Joan Seoane<sup>6</sup>, Carlos Caldas<sup>7</sup>, Daniel J. Pennington<sup>4</sup>, Paul Pfeffer<sup>4</sup> and Jane Osbourn<sup>1</sup>

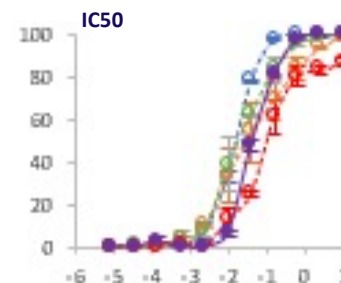


Alchemab

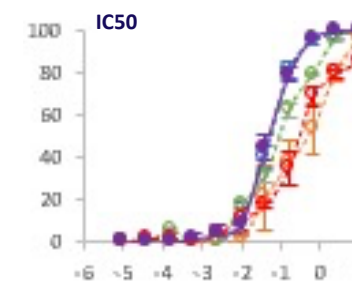
Commercial  
cocktail & control

## Potent multi-strain covid neutralizing antibodies identified

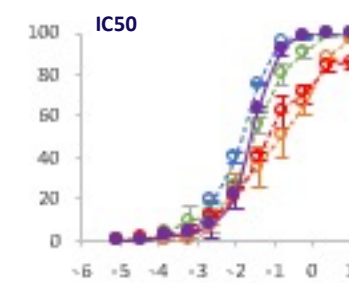
ATL5003



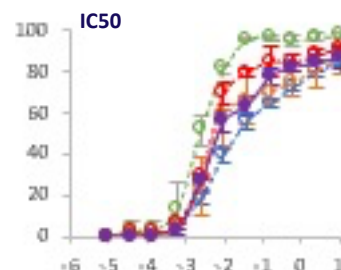
ATL5006



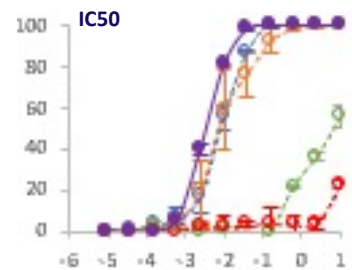
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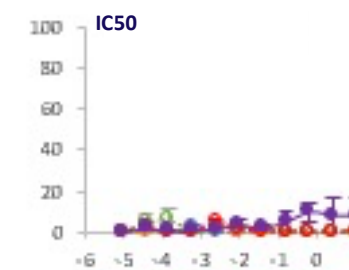
mAb1



mAb2



Isotype control (G1/K)

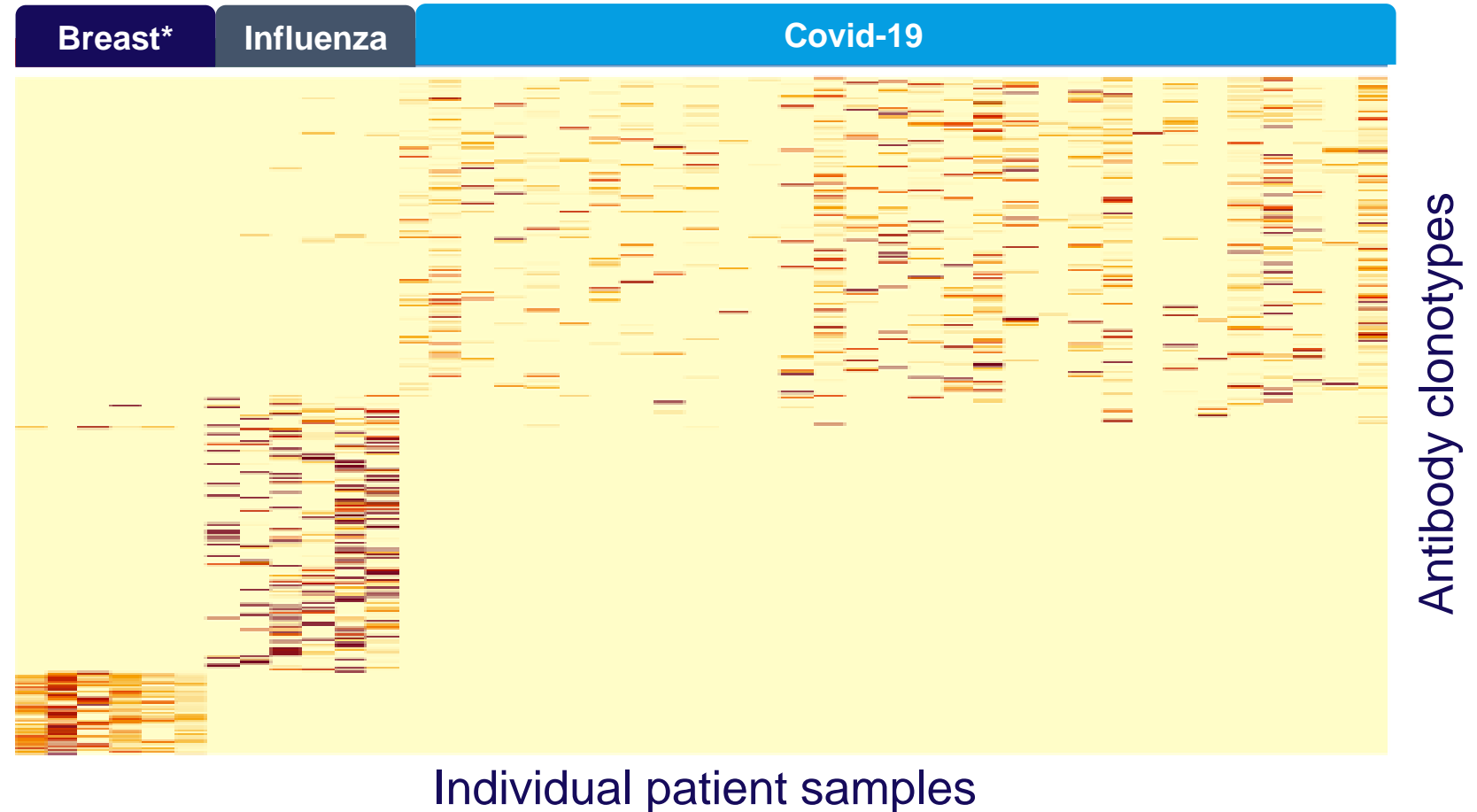


Victoria  
Alpha  
Beta  
Gamma  
Delta

# Alchemab's growing Data Cube can stratify patients into diseases and disease-subtypes

## Disease stratification example

Alchemab data demonstrates that antibody convergence is specific to disease cohorts, suggesting that separation between diseases is possible\*\*

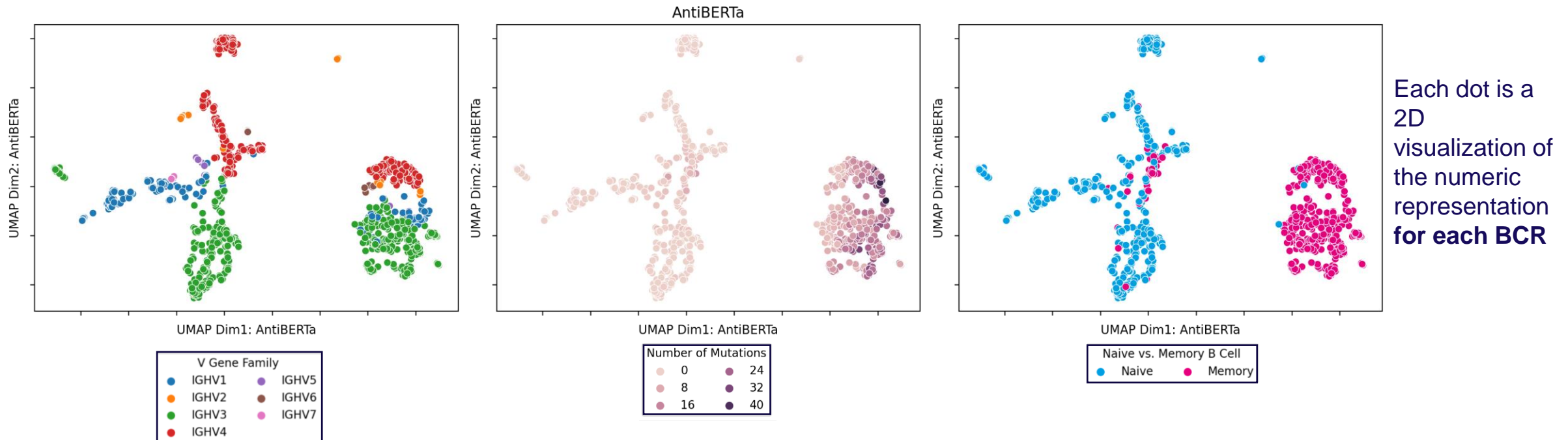


Antibody clonotypes

Individual patient samples

# Representation learning could uncover unique patient signatures

- Discovers hidden patterns in the data
- Alchemab's AntiBERTa – 'learns the language of antibodies'
- It also learns additional information we cannot yet interpret with our current state of biological knowledge



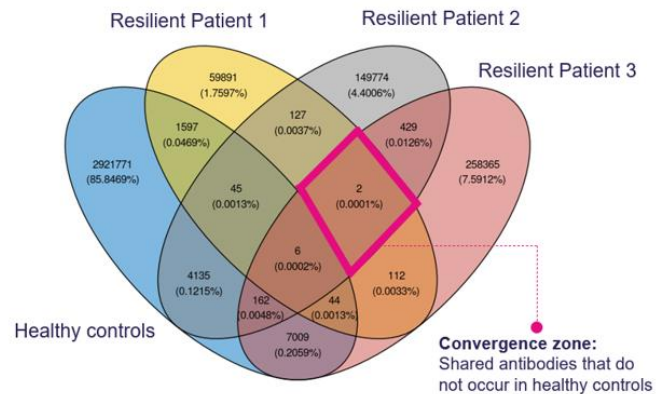
We apply this to our resilient patient antibody data sets and believe that it could be a transformative patient stratification tool

## Oncology – Case study



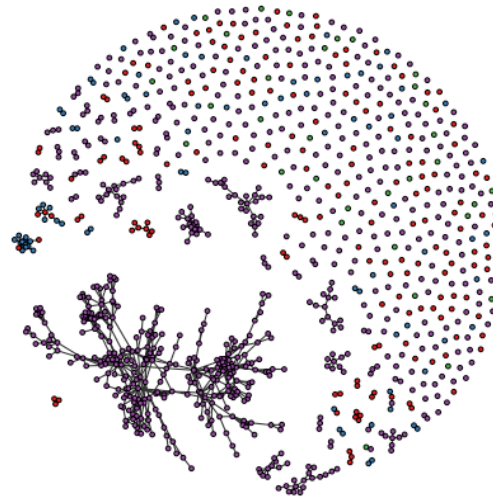
# Convergence: Breadth of opportunities

## Convergence between patients



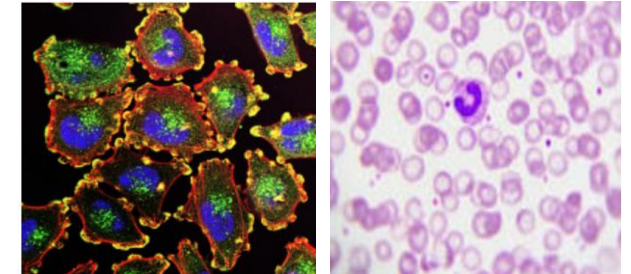
**Shared antibodies may indicate protection at the population level**

## Convergence within patient







**Related antibodies suggest active selection against important targets**

## Convergence between tumor and periphery



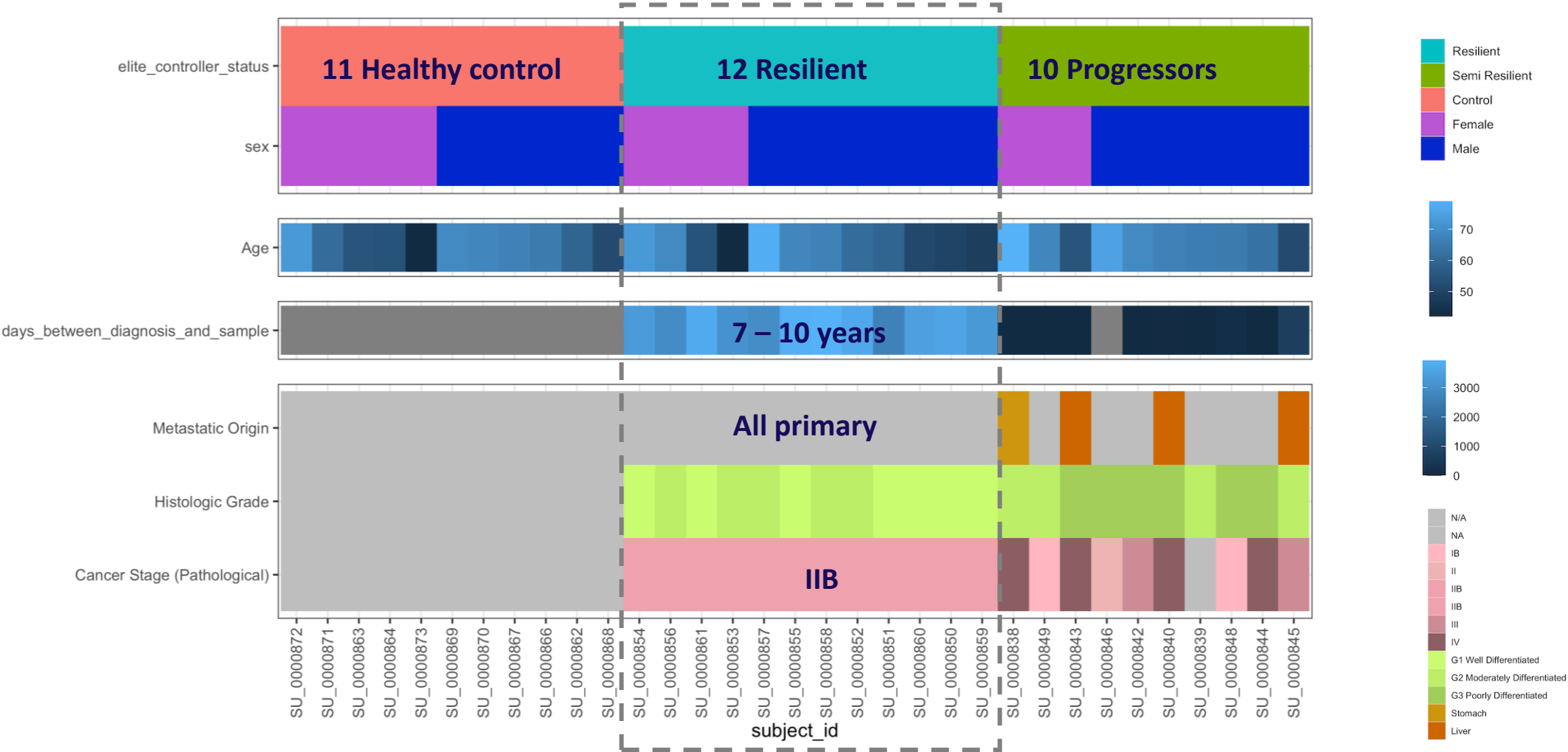
**Tumor-surveilling antibodies may identify targets active in the TME**

# Pancreatic cancer collaborations to date

<div><div></div> RNA<div></div> Plasma/Serum<div></div> Tumour Tissue / B cells</div>			
Collaborator	Type	Donors (#)	Resiliency definition
 UNIVERSITY OF OXFORD	<div></div>	28	Significant B cell infiltration
	<div></div>	31	LTS (5+ years)
	<div></div>		
Case study	<div></div>	36	LTS (7+ years)
	<div></div>		
	<div></div>	30	LTS (5+ years)
	<div></div>		
	<div></div>	40	LTS (5+ years)

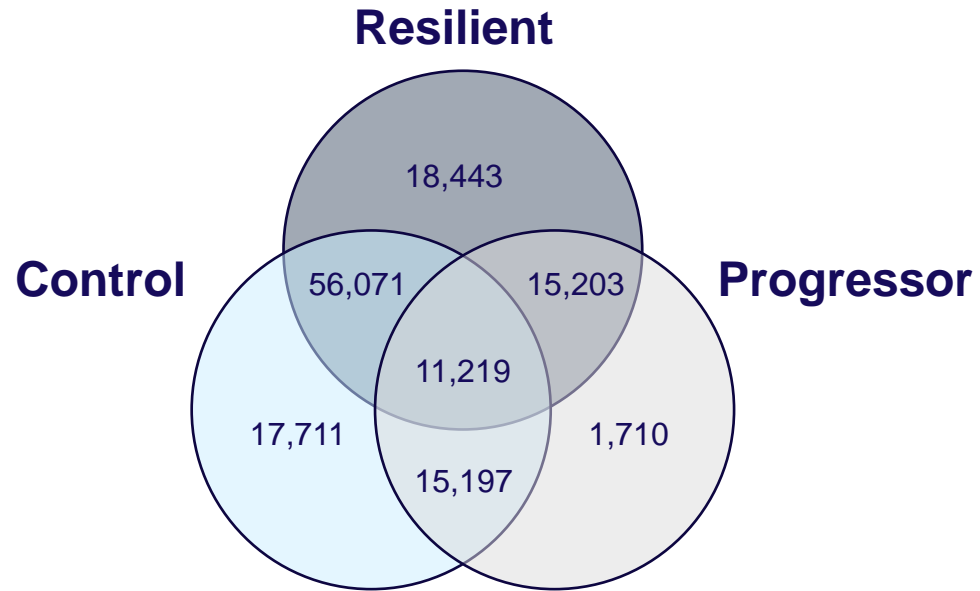
# Pancreatic cancer cohort

All PDAC patients have undergone resection



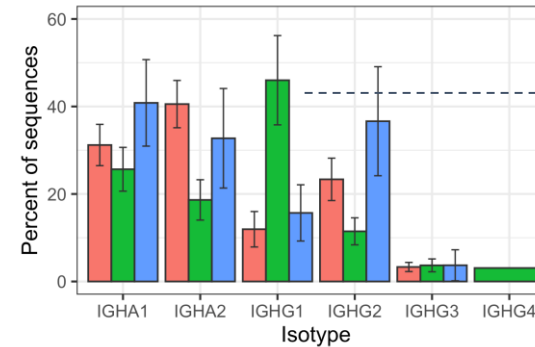
# Early analysis shows high convergence in resilient group

## Convergent clonotypes Total: 135,554



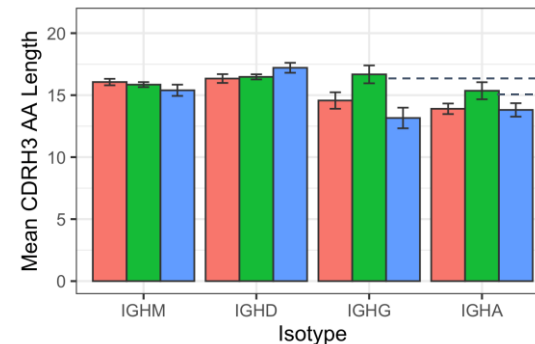
There are many more Resilient-convergent than Progressor-convergent clonotypes

## Class-switch isotype distribution



IGHG1: Increased levels in resilient, & associated with antigen response & improved survival

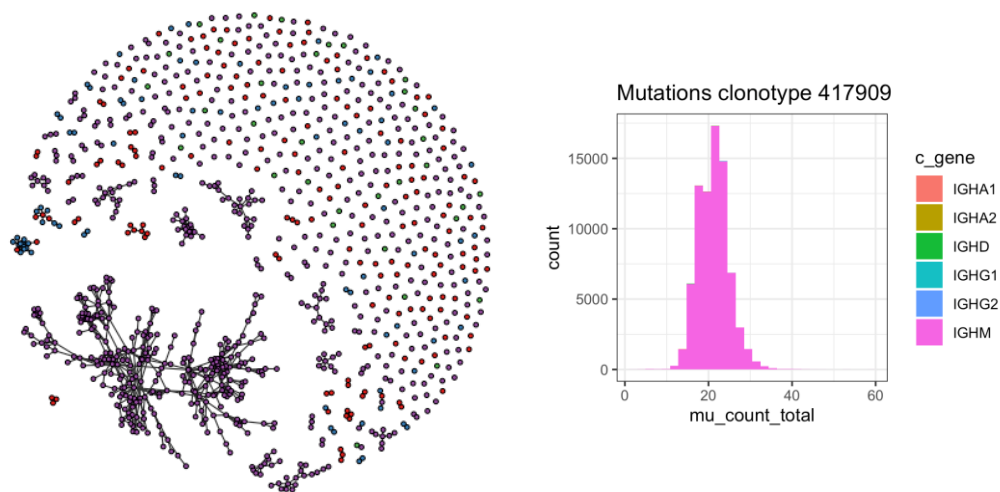
## Mean CDR3 length



Longer loop lengths in class-switched antibodies

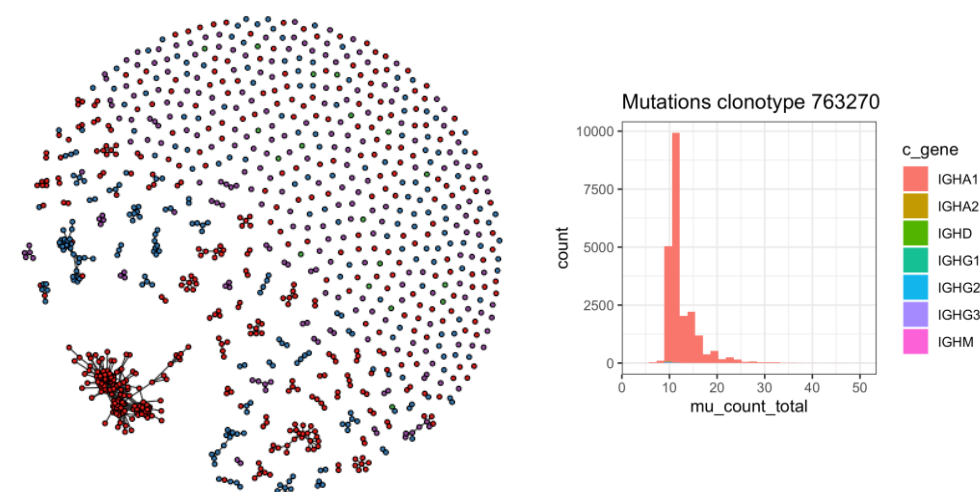
# Two large clonal expansions found in resilient subjects

## SU859-Resilient



29% repertoire

## SU857-Resilient



16% repertoire

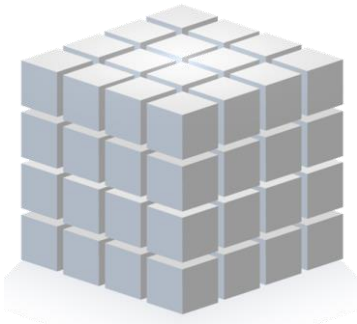
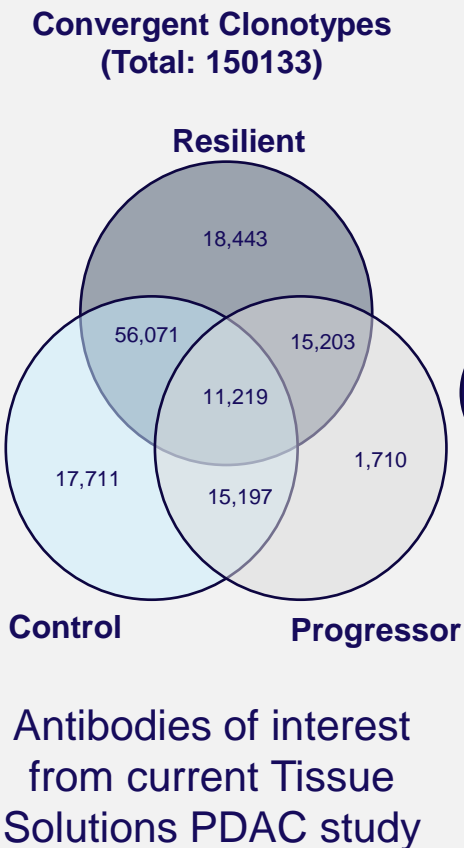
# Highly convergent resilient clonotypes found in multiple individuals

10 clonotypes found with exceptional convergence among resilient individuals **which were not found in progressors**, and evidence for disease relevance:





- Length-independent super-convergence
- Low probability of generation & rare in healthy controls
- Predominantly IgG1

CLUSTER SIZE	CONVERGENCE LEVEL (OUT OF 12)	CDR3 LENGTH	MUTATIONS	GENERATION PROBABILITY	PROP. IN HEALTHY CONTROL	PREVALENT ISOTYPE
151	8	20	1.58	4.35E-12	0.060	IGHG1
40	8	20*	1.65	9.34E-12	0.050	IGHG1
43	8	19*	1.35	4.63E-20	0.186	IGHG1
73	7	16	1.51	2.13E-13	0.055	IGHG1
42	5	16	2.07	1.29E-12	0	IGHG1
73	5	15*	1.41	7.42E-11	0.082	IGHG1
33	5	14*	1.52	7.69E-11	0	IGHG1
17	5	13	1.18	1.14E-11	0.059	IGHG1
16	6	13	17.63	1.01E-11	0	IGHM
22	5	11	25.59	5.20E-10	0.091	IGHA1

\*Similar sequences despite length differences



## Other Alchemab PDAC studies

Collaborator	Type	Donors (#)	Resiliency definition
 UNIVERSITY OF OXFORD		28	Significant B cell infiltration
 LEON BERARD		31	LTS (5+ years)
Case study		36	LTS (7+ years)
 Barts Cancer Institute		30	LTS (5+ years)
 UNIVERSITY OF MANCHESTER		40	LTS (5+ years)

## External datasets of interest

CANCER RESEARCH | TUMOR BIOLOGY AND IMMUNOLOGY

### Tumor-Infiltrating B Lymphocyte Profiling Identifies IgG-Biased, Clonally Expanded Prognostic Phenotypes in Triple-Negative Breast Cancer

Robert J. Harris<sup>1,2,3</sup>, Anthony Cheung<sup>1,2,4</sup>, Joseph C.F. Ng<sup>5</sup>, Roman Laddach<sup>1,2,6</sup>, Alicia M. Chenoweth<sup>1,2,4</sup>, Silvia Crescioli<sup>1,2</sup>, Matthew Fittall<sup>1,2,4</sup>, Diana Dominguez-Rodriguez<sup>1,2</sup>, James Roberts<sup>1,2,6</sup>, Dina Levi<sup>4</sup>, Fangfang Liu<sup>4</sup>, Elena Alberts<sup>1,2,4</sup>, Jelmar Quist<sup>4</sup>, Aida Santaolalla<sup>7,8</sup>, Sarah E. Pinder<sup>9,10</sup>, Cheryl Gillett<sup>9,10</sup>, Niklas Hammar<sup>7</sup>, Sheeba Irshad<sup>9</sup>, Mieke Van Hemelrijck<sup>7,8</sup>, Deborah K. Dunn-Walters<sup>11</sup>, Franca Fraternali<sup>12</sup>, James F. Spicer<sup>9</sup>, Katie E. Lacy<sup>1,2</sup>, Sophia Tsoka<sup>4</sup>, Anita Grigoriadis<sup>4</sup>, Andrew N.J. Tutt<sup>4,12</sup>, and Sophia N. Karagiannis<sup>1,2,4</sup>

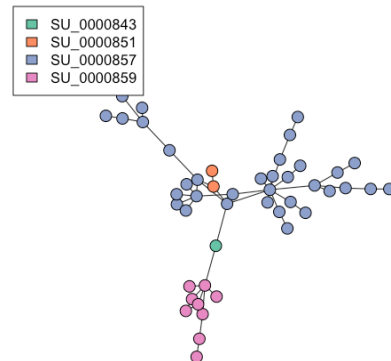
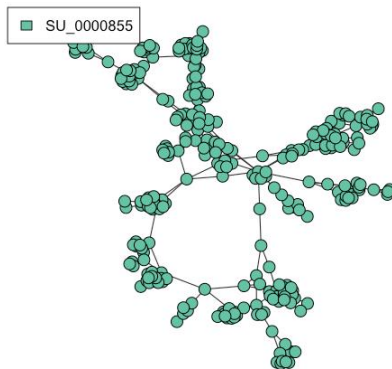
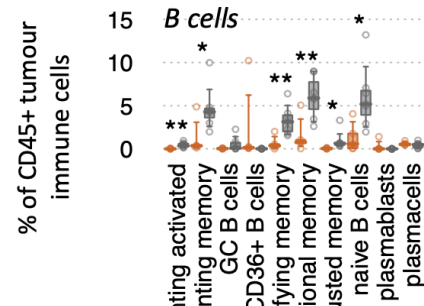
## Alchemab curated antibody database

~10,000 Abs from the literature with known binding properties

# Mining PDAC datasets for convergence

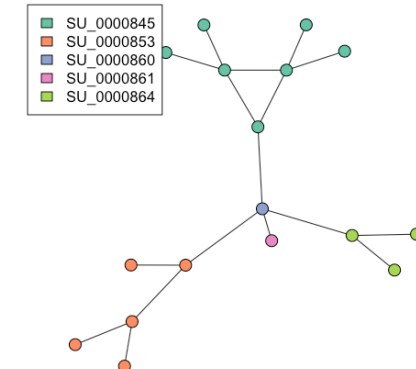
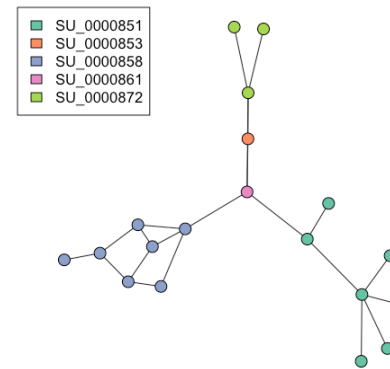
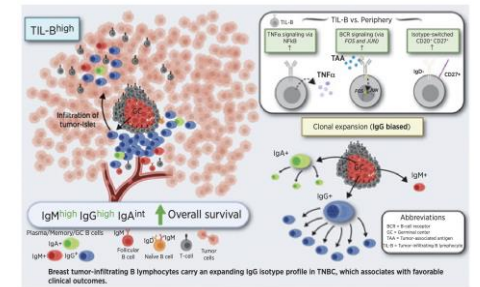
## Other Alchemab PDAC studies

- Alchemab pilot PDAC study comparing high and low B cell infiltrate groups
- Single-cell sequencing from blood and tumor
- Two matches to PDAC case study demonstrating convergence from pilot



## External datasets of interest

- Triple Negative breast cancer study from Harris *et al.* investigating prognostic B cell profiles
- Single-cell sequencing from tumor samples
- Two matches to PDAC case study demonstrating high levels of convergence between different solid tumor studies





# Homology to approved KDR mAb translates to target binding

Ramucirumab is a VEGFR2 (KDR) antibody, approved for treatment of solid tumors

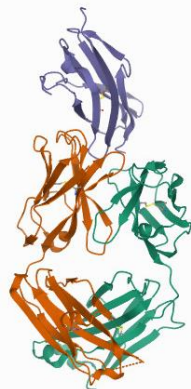
One of Alchemab's convergent clusters is highly homologous to ramucirumab

ARTICLES | VOLUME 383, ISSUE 9911, P31-39, JANUARY 04, 2014

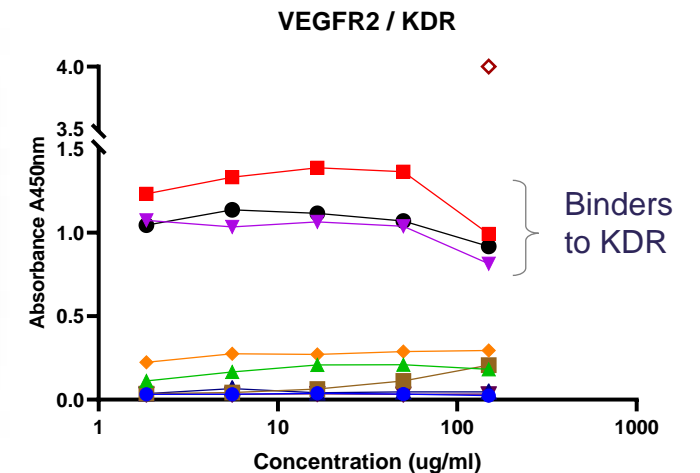
Ramucirumab monotherapy for previously treated advanced gastric or gastro-oesophageal junction adenocarcinoma (REGARD): an international, randomised, multicentre, placebo-controlled, phase 3 trial

Dr Prof Charles S Fuchs, MD • Jiri Tomasek, MD • Prof Cho Jae Yong, MD • Filip Dumitru, MD • Rodolfo Passalacqua, MD • Prof Chanchal Goswami, MD • et al. [Show all authors](#) • [Show footnotes](#)

Published: October 03, 2013 • DOI: [https://doi.org/10.1016/S0140-6736\(13\)61719-5](https://doi.org/10.1016/S0140-6736(13)61719-5) •  Check for updates

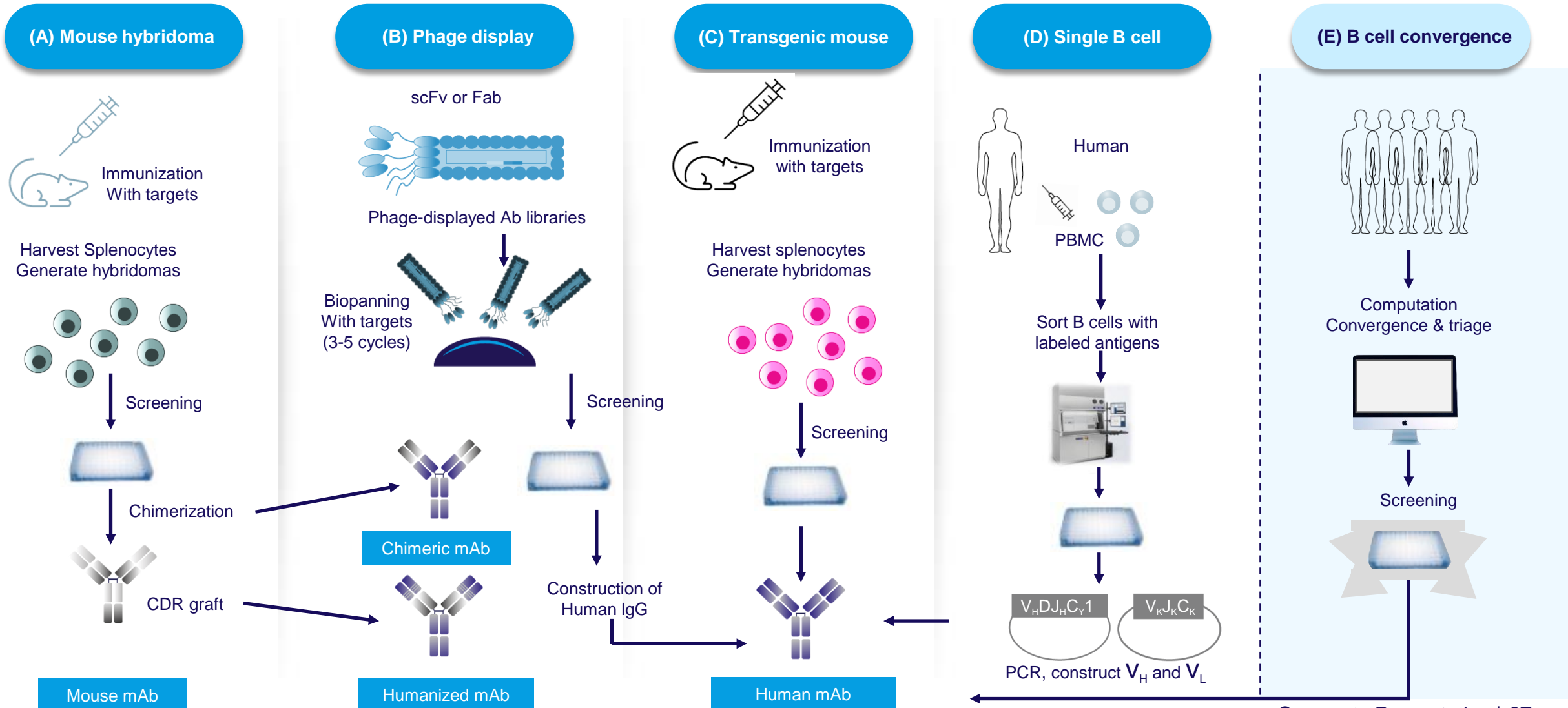


- Assay shows binding for multiple antibodies from one clonotype to KDR



# Summary

# Convergence analysis could be the next wave of antibody generation



# We have proven the concept and we are expanding our capabilities



## **Antibody Therapeutics**

Unique resilient-  
convergent antibodies  
identified from PDAC  
cohort

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Antibodies with  
sequences highly similar  
to known, efficacious  
antibodies identified



## **Target Identification**

Antibodies undergoing  
target deconvolution



## **Patient Stratification**

Early evidence of ability  
to stratify patients into  
disease sub-groups



## **Alchemab Discovery**

World's most advanced  
machine learning model  
evaluating B cell  
repertoires demonstrating  
potential to identify  
new biology

**Thank you!**