OmniAb

Opening The Barn Door To Antibody Diversity

Bill Harriman, PhD
SVP Antibody Discovery

Antibody Engineering and Therapeutics
December 6, 2022



The OmniAb Technology Suite

OmniAb



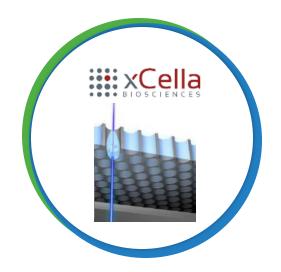














The only platform leveraging four species

Robust solutions for bispecific antibodies

Human frameworks with ultralong CDR-H3s

Industry-leading broadest offering

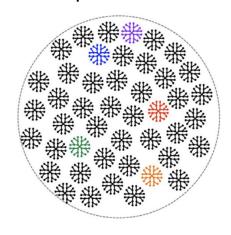
Proven success



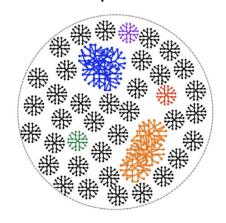
Platforms to Generate Custom Antibody Repertoires

BIOLOGICAL INTELLIGENCE™: INTERPLAY BETWEEN RATIONAL GENETIC DESIGN AND POWERFUL IN VIVO PROCESSES

Naïve repertoire



Immune repertoire



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Building the Animal System

V gene building blocks Structural attributes Diversification architecture Transgene design Immunological robustness

Repertoire Shaping

Antigen design
Host immune recognition
Immunization protocols
Campaign strategy
Immune response monitoring

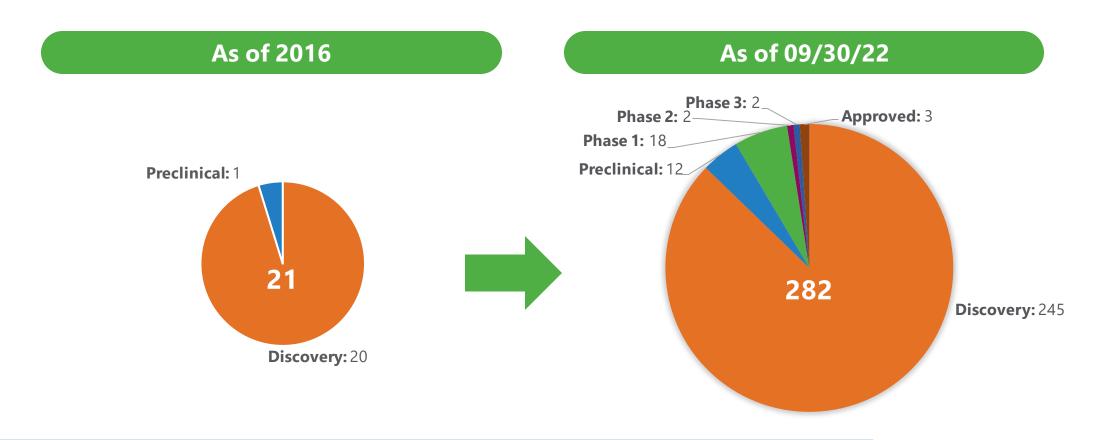
Repertoire Mining

Phenotypic screening
Clonal sampling/sequencing
Antibody characterization
Clone & Repertoire ranking
Selective "Deep Dives"
NGS hit expansion



OmniAb Program Count Continues to Grow

PROGRESSION AND PERFORMANCE IN PROGRAMS BY STAGE OF DEVELOPMENT



Substantial progress in all phases, increase in discovery programs expected to feed growth in new clinical programs and future approvals



OmniAb

Animal Platforms

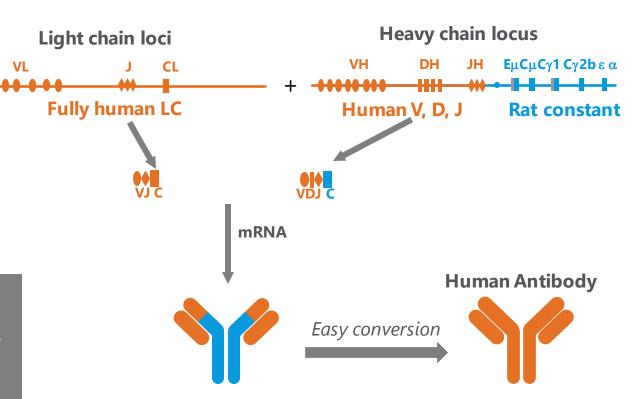


Rodent Platforms



- Endogenous Ig genes inactivated
- Expression of full human V gene diversity
- Streamlined conversion into fully human molecule

Well-validated transgene design utilizes rodent constant regions for robust immune responses from the B-cell repertoire

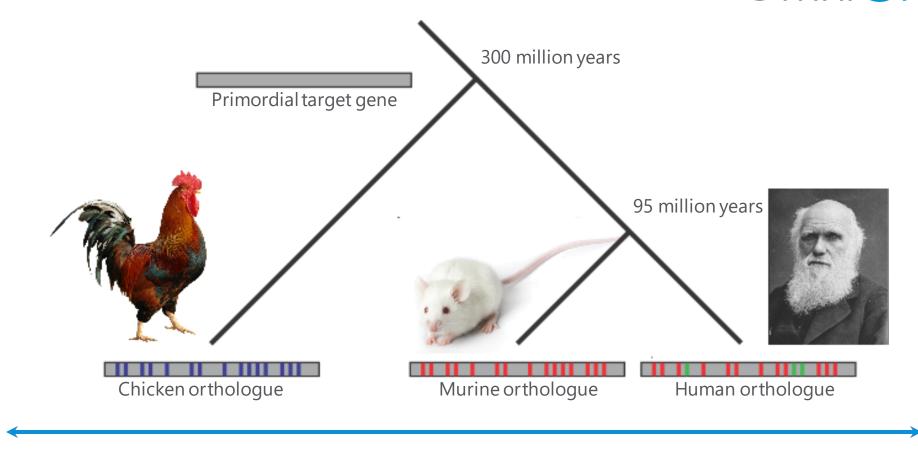




OmniChicken®

POWERED BY EVOLUTION





Greater evolutionary distance yields greater immunogenicity and more antibody diversity



Engineering of Ig Loci

ADAPTATION TO CHICKEN GENE CONVERSION PROCESS

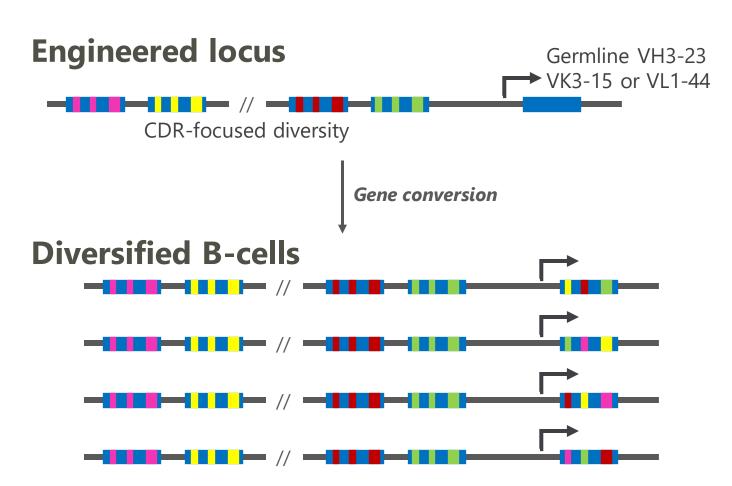


Gene conversion



Human V's selected for:

- High expression level, stability, ubiquity
- High sequence diversity in CDRs
- Low sequence diversity in FWs



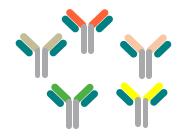


Common Light Chain Platforms

STANDARD IGG FORMAT TO DE-RISK DOWNSTREAM DEVELOPMENT¹ OF BISPECIFIC MABS



Rearranged human VK3-15 light chain combined with diversifying heavy chain



Simple reformatting from monospecific into bispecific for efficient production



Bispecific IgG



"Germlining" human VK3-15 light chain combined with diversifying heavy chain



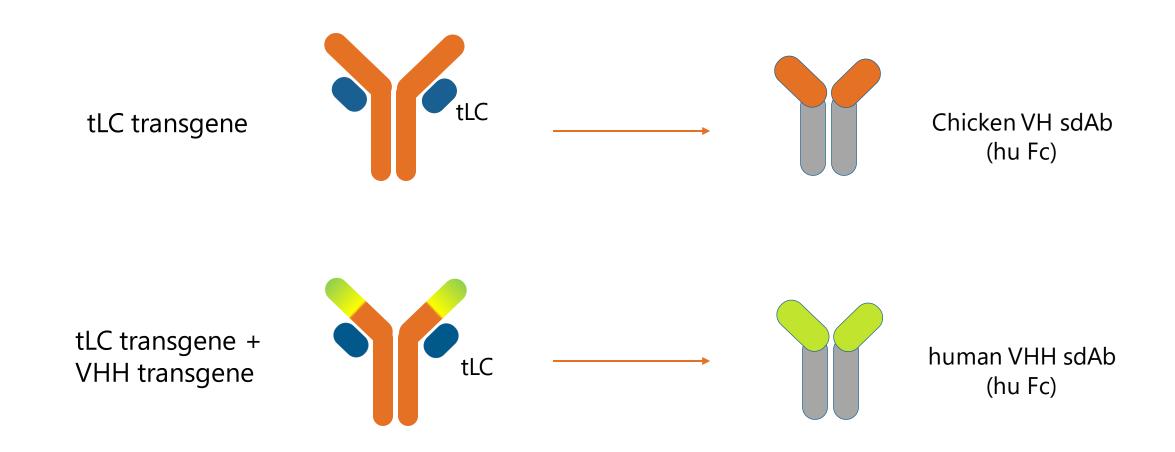
Monospecific IqG

Common light chain for OmniFlic® and OmniClic® allows interchangeability between the platforms



OmnidAb[™]: Heavy Chain Only Transgenic Chickens

HCO STRATEGY USING TRUNCATED LIGHT CHAIN (TLC)



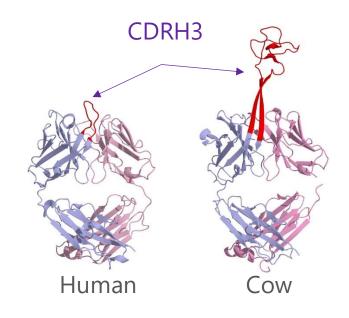
- Normal chicken heavy chain can express as VH alone
- VHH transgene in development



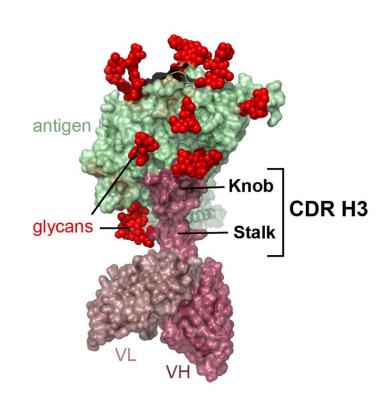
OmniTaur[™]: Ultralong CDRH3s Create Novel Binding Domains



UNIQUE STRUCTURAL FEATURES OF ULTRALONG H3 ANTIBODIES



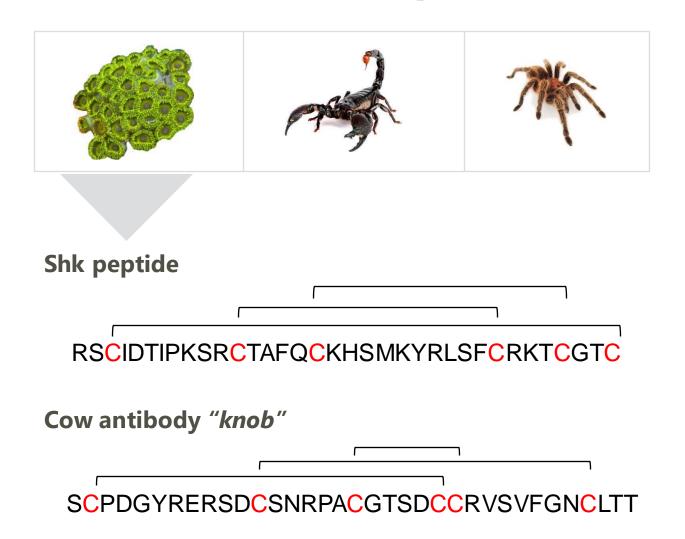
- Novel structure may enable targeting epitopes unreachable by standard antibodies
- Long H3 domains can be expressed on human VH framework, or alone as ~5kD Picobodies[™]

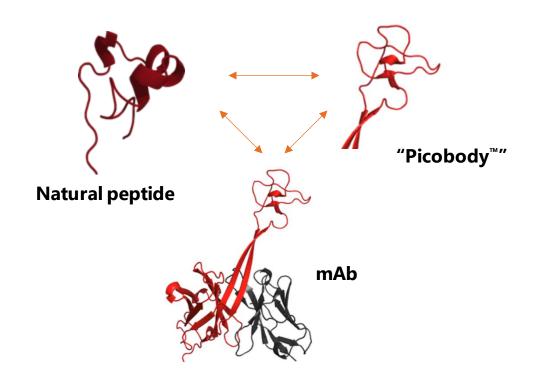


Stanfield, et.al. Sci Adv (2020) 6(20): eaba0468.



OmniTaur™ mAbs Share a Structural Theme with Bioactive Natural Peptides



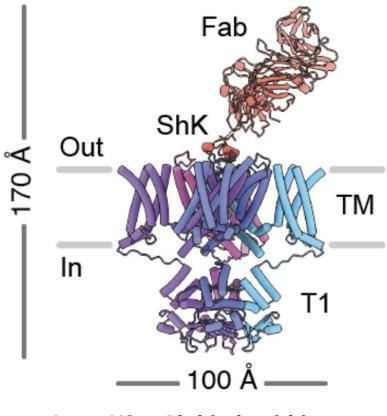


Long H3 mAbs potentially combine high biological potency with high target specificity

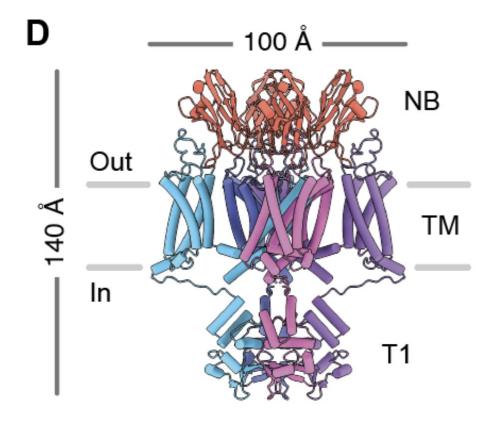


mAb Binding Modes to Ion Channel Kv1.3

H3 LENGTH IMPACTS TYPES OF MOLECULAR INTERACTION CRYO-EM ~3.5A



Long H3 mAb binds within pore



Nanobodies (~15KD) bind turret loops



OmniAb Antibody Repertoires

UNSURPASSED OPTIONS AVAILABLE TO ADDRESS DIVERSE PARTNER OBJECTIVES

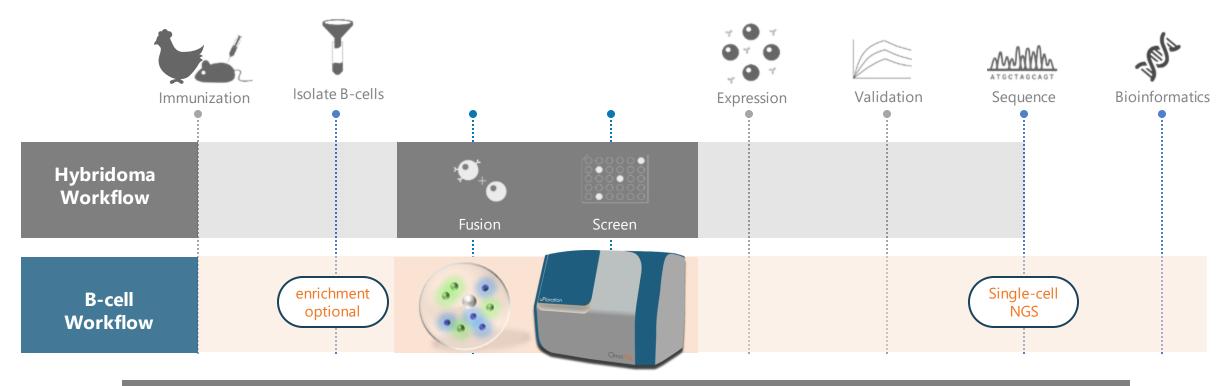
Host	V genes	Structural and immunological features	Benefits for therapeutics discovery and development
5 OmniMouse	Full human V gene diversity Choice of light chain isotype	Diverse V gene usage and mixed genetic backgrounds	Widely accessible and flexible workflows
53 OmniRat [®]	Full human V gene diversity Choice of light chain isotype	Diverse V gene usage and mixed genetic backgrounds Distinctive target recognition	Industry standard Widely accessible and flexible workflows Extensive track record
Omni Chicken	Single framework VH3/VK3 or VH3/VL1	Evolutionarily divergent host system for robust immune responses	Diverse and new epitope coverage High homology targets Excellent physical properties
53 OmniFlic	Full human VH gene diversity with non-diversifying VK3	Fixed light chain for bispecific applications	Bispecific applications leveraging standard IgG format
Omni Č lic*	Single framework VH3/non-diversifying VK3	Fixed light chain for bispecific applications	Diverse epitope coverage Excellent physical properties Ease of manufacturing
Omni dAb ™	Single camelized human VH framework with truncated LC	Domain antibody of the "VHH" type	Diverse and new epitope coverage from human single-domain format, 12-15kD Building blocks for multispecific molecules
OmniTaur™	Single framework VH4/VL1	Ultralong CDR-H3's for enormous structural diversity	Access cryptic epitopes Unique modalities (picobodies™) Building blocks for multispecific molecules



OmniAb

Screening Platforms

Discovery Platforms



Our powerful single B-cell screening technologies, **xPloration**® **and GEM assay**, **bypass bottlenecks of hybridoma workflows**

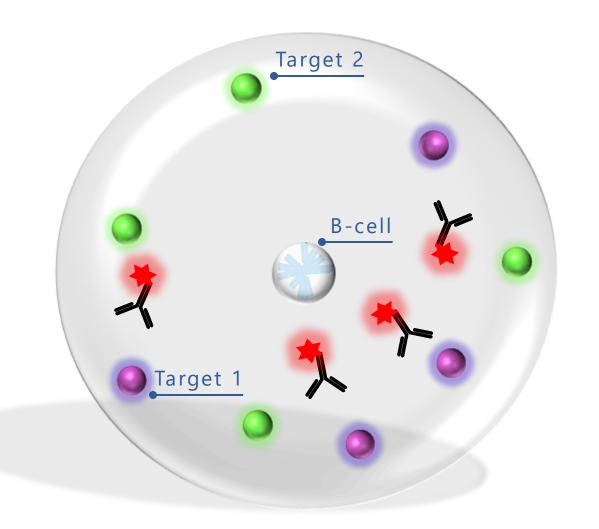
Al-driven multi-parameter screening of tens of millions of cells in hours instead of weeks

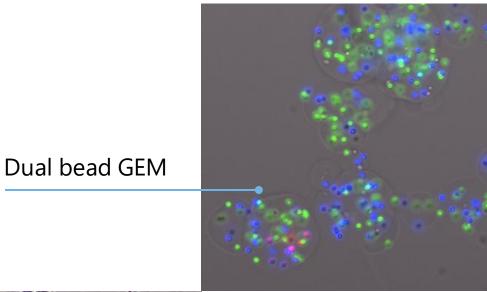
Technologies enable **screening against difficult targets**: GPCRs, ion channels and surface antigens

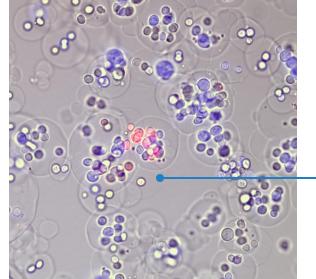


GEM Assay

GEL ENCAPSULATED MICROENVIRONMENT





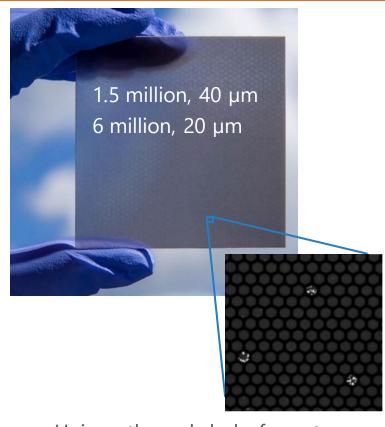


Cell-based GEM



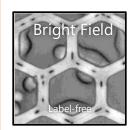
xPloration[®]

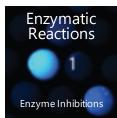
1 | Loading



Unique through-hole format Workflows for OmniAb B-cells

2 | Assay + Machine Vision





















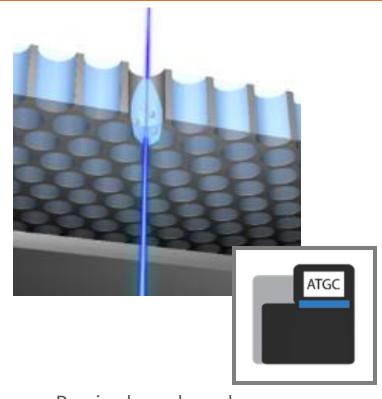






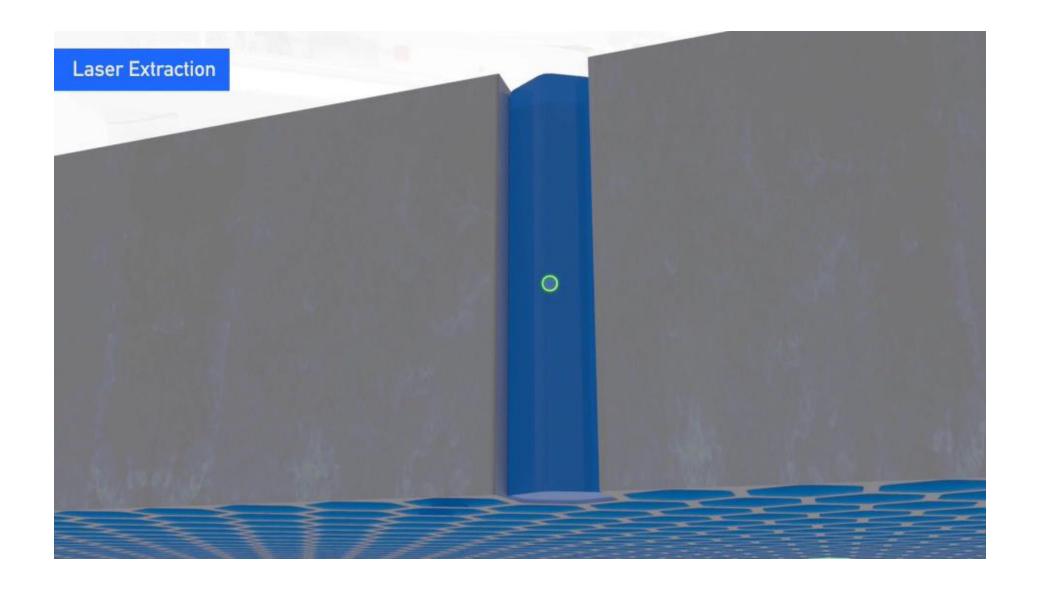
Al-driven hit detection

3 | Recovery & Single-Cell NGS



Precise laser-based recovery

1 cell/sec (single-cell mode), single-cell barcoding

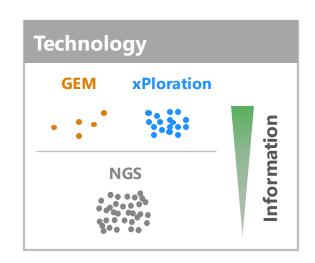




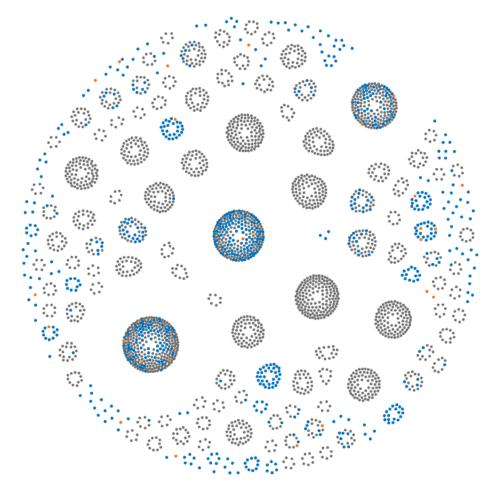
Deeper Characterization Identifies New Clonotype Families

Each dot represents an antibody





- xPloration® expands on majority of clonotypes identified by GEM assay
- Multiple new clonotype families identified
- Next-generation sequencing (NGS) adds support to new clusters and reveals even more diversity





Leveraging Biological Intelligence with Computational Tools

Biological Intelligence OmniChicken Omni Taur Omni**Č**lic Differentiated sources of antibody sequences

Model-aided optimization

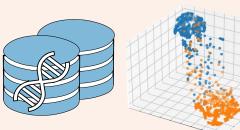
- Antibody homology modeling
- Computational antibody optimization

In silico developability



- Sequence liability assessment
- Comparison to clinical antibodies
- Structure-based calculations

Sequence databases and Bioinformatics



See posters #67 & 68

- Customized cloud-based antibody sequence databases
- Large-scale repertoire analysis

The OmniAb Platform

Create Diverse Antibody Repertoires

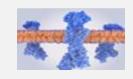
Screen Antibody Candidates

Identify the Right Antibody

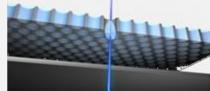
Create Diverse Pools of High-Quality Naturally Optimized Antibodies

Screen Millions of Cells to Find Potential Therapeutic Candidates

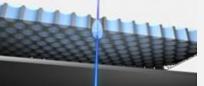
Further Characterize, Select & *Optimize the Right Antibody*



Computational Antigen Design & **Proprietary Reagents**



xPloration High-Throughput Single Cell Screening





Custom

Bioinformatics

Next Generation

Hit Expansion

Sequencing (NGS)

- Comprehensive **Functional** Characterization
 - Proprietary Ion Channel Assays





Generation



Cow-inspired Antibodies for **Difficult Targets**



Gel Encapsulated Microenvironment (GEM) Single Cell Screening

Technology offering addresses the most critical challenges of antibody discovery



OmniAb

THANK YOU TO THE OMNIAB TEAM!

www.OmniAb.com