



# **APEX<sup>TM</sup> Antibody Discovery and Production Technology**

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PEP Talk Conf.  
San Diego  
24Jan20

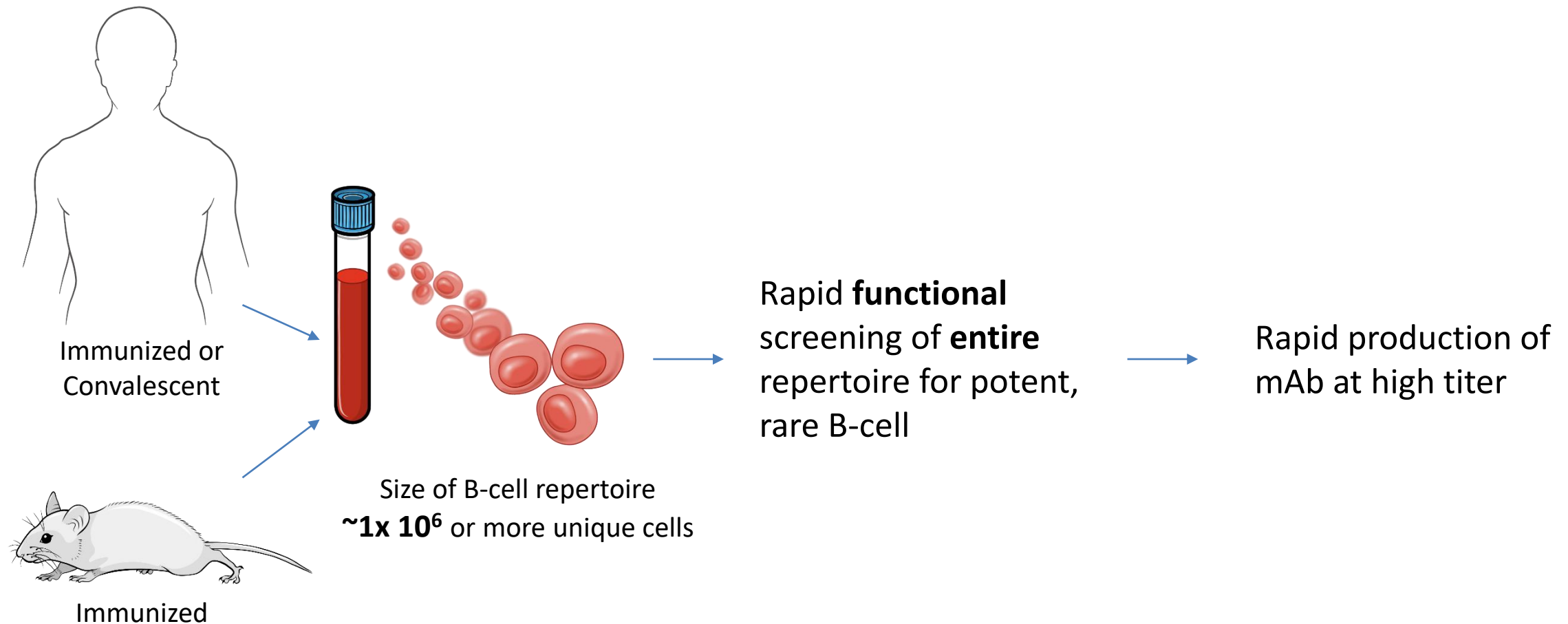
# Outline

- **Single cell cloning & mAb discovery**
- **B-cell immortalization using cell fusion**
- **Enhanced mAb production of established cell lines**
  - **CRISPR-enabled activation of transcription regulatory genes**
- **mAb production CHO cell line technology**

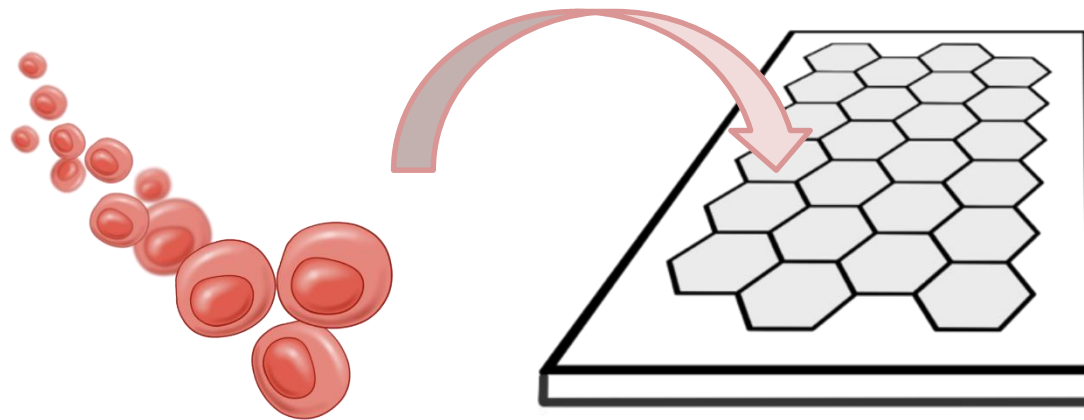
# Outline

- **Single cell cloning & mAb discovery**
- B-cell immortalization using cell fusion
- Enhanced mAb production of established cell lines
  - CRISPR-enabled activation of transcription regulatory genes
- mAb production CHO cell line technology
- **COMING SOON: Transposons-mediated gene amplification**

# Antibody Discovery and Production Challenges



# Single Cell Screening & Clonal Selection Using NanoArrays

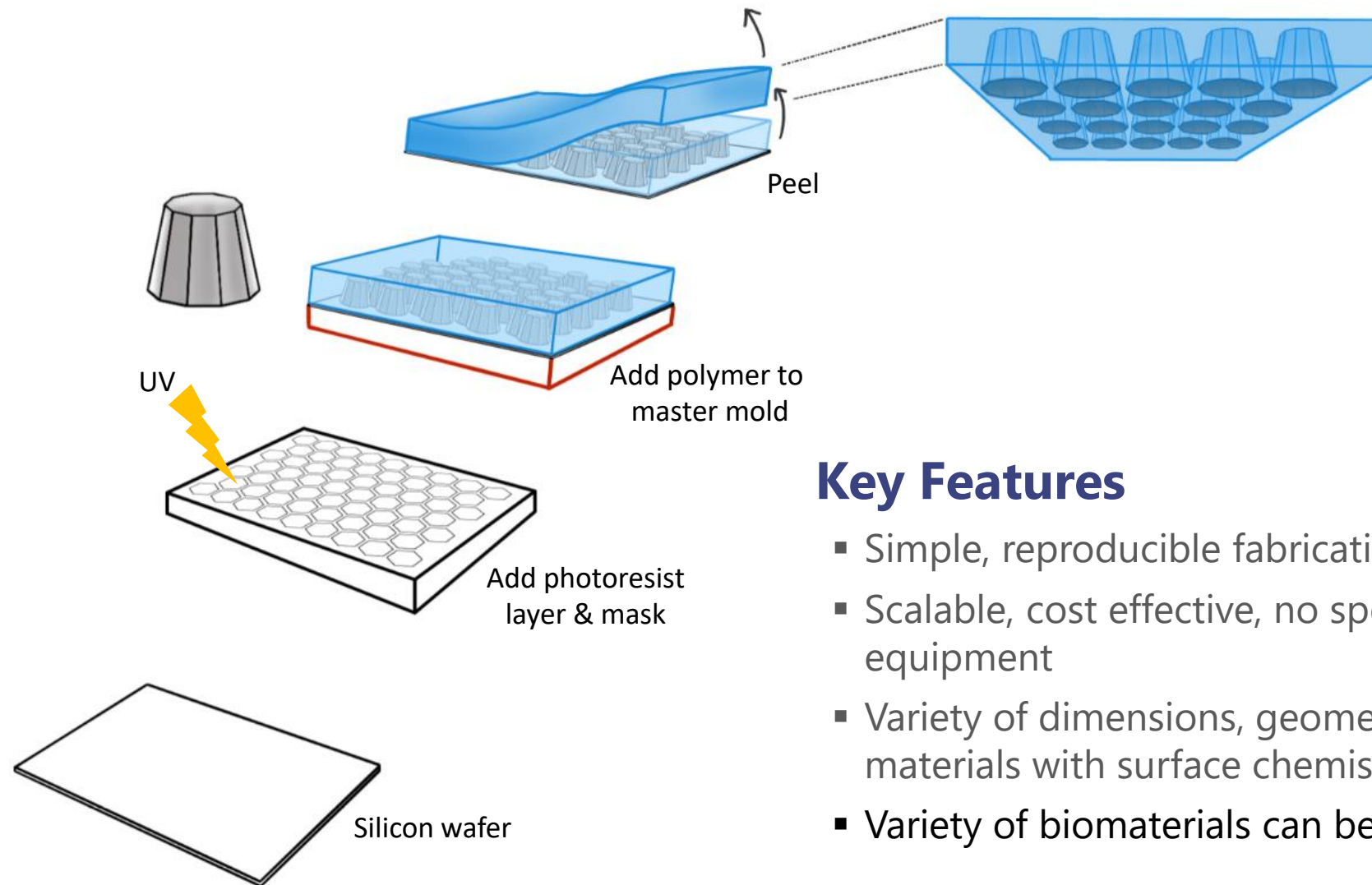


**APEX™  
Nanoculture Arrays**

## Key Features

- Enables comprehensive, **functional screening at single cell level** of  $10^6$ - $10^7$  cells simultaneously
- **~100,000x minimization** of tissue culture wells. **0.5 to 3 million wells** can be accommodated in a 96-well plate surface area
- **Increase protein concentration** by ~5-orders of magnitude enables sub-picogram mAb detection
- Enables simultaneous **single cell cloning and plucking** by microneedles for expansion
- Proteins, viruses, bacteria can be adsorbed as antigen for detection of mAb
- **Highest binders can be identified in ~1-2 days**

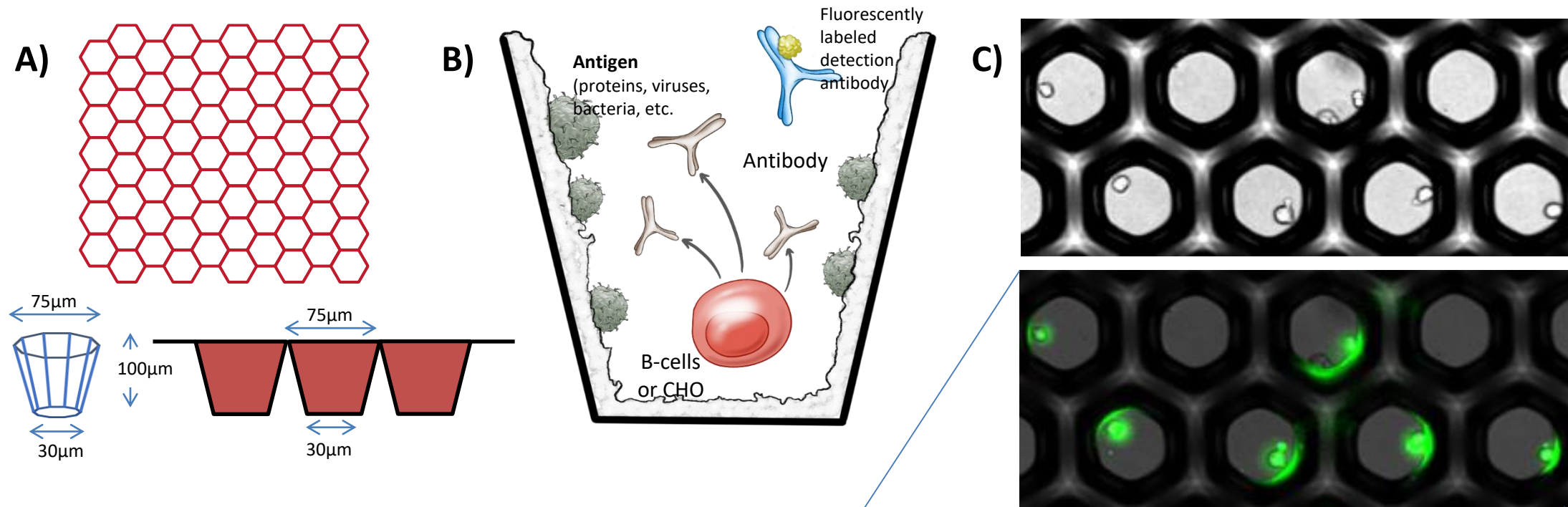
# Fabrication of Nanoculture Arrays



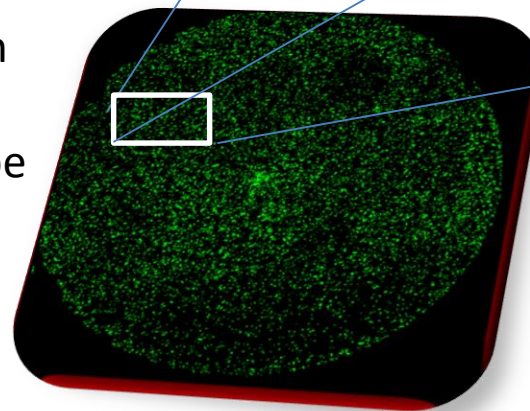
## Key Features

- Simple, reproducible fabrication process
- Scalable, cost effective, no specialized companion equipment
- Variety of dimensions, geometries, biocompatible polymer materials with surface chemistries
- Variety of biomaterials can be adsorbed

# B-cell Repertoire Screening, Single Cell Analysis & Single Cell Cloning with APEX™ Nanoarrays



**APEX™ Nanoarrays:** A) Schematic top view of honeycomb shape arrays with no gaps in-between the wells, which allows seeding of all cells in solution. Packing density of 1.4 million wells can be accommodated per 96-well surface area (75µm opening diameter). C) High content fluorescent imaging of B-cells secreting mAbs in nanoarrays

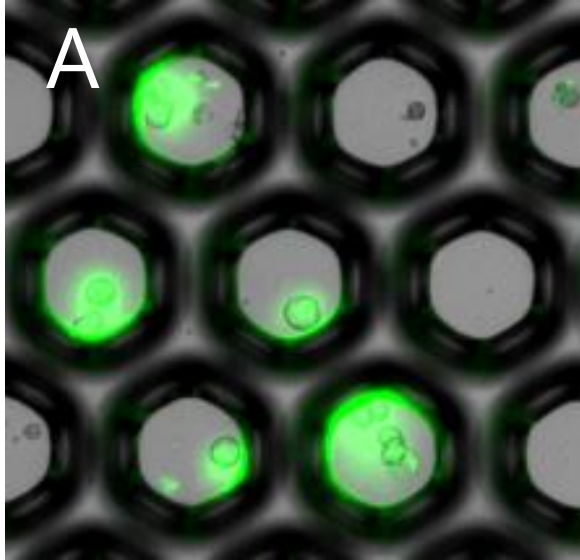




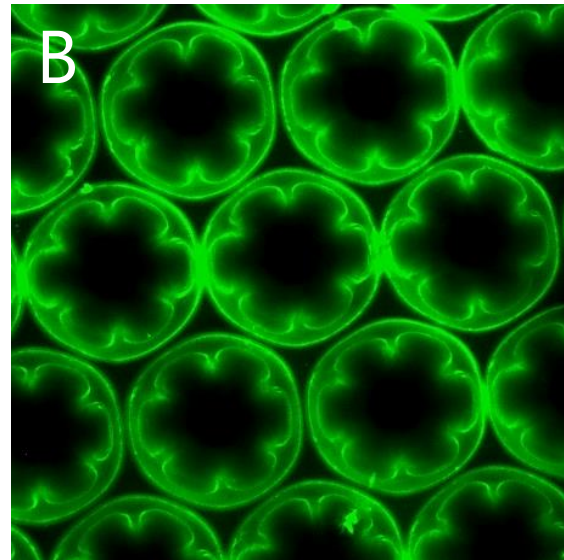
# Versatile On-chip Assay Format for Single Cell Analysis

**Top view of Nanoculture wells (A-D).** Wells are embedded with antigen as indicated, followed by detection with fluorescence labeled secondary mAb

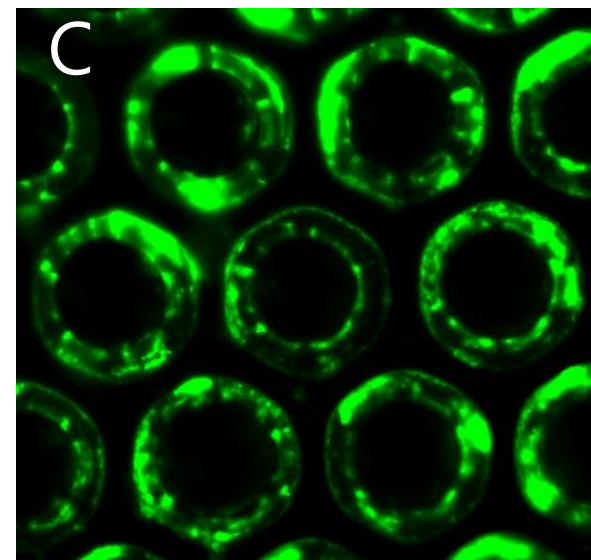
Anti-IgG



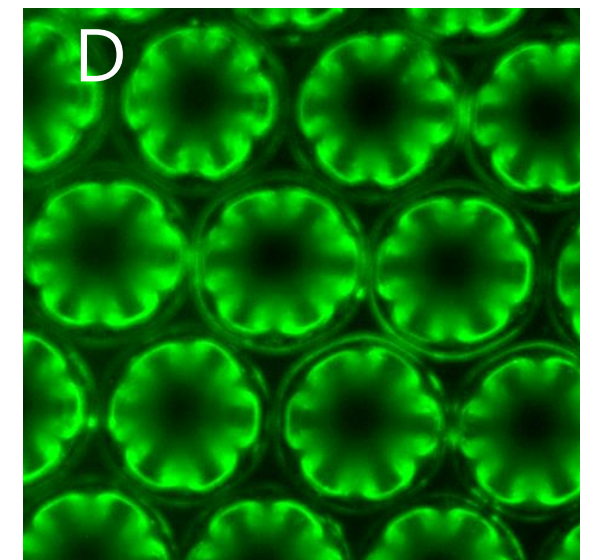
Live B-cell secreting Anti-RSV F-protein IgG (=antigen)



Antigen= Human AR-105 mAb

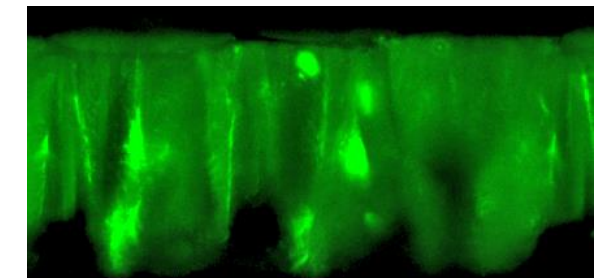
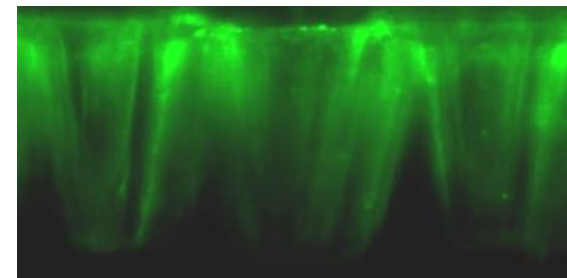


Antigen= adsorbed whole cell *P. aeruginosa* bacteria expressing alginate (detected w/ anti-alginate mAb)



Antigen= Adsorbed intact rotavirus expressing G1 protein (detected with anti-G1 mAb)

Sideview of Nanoculture wells

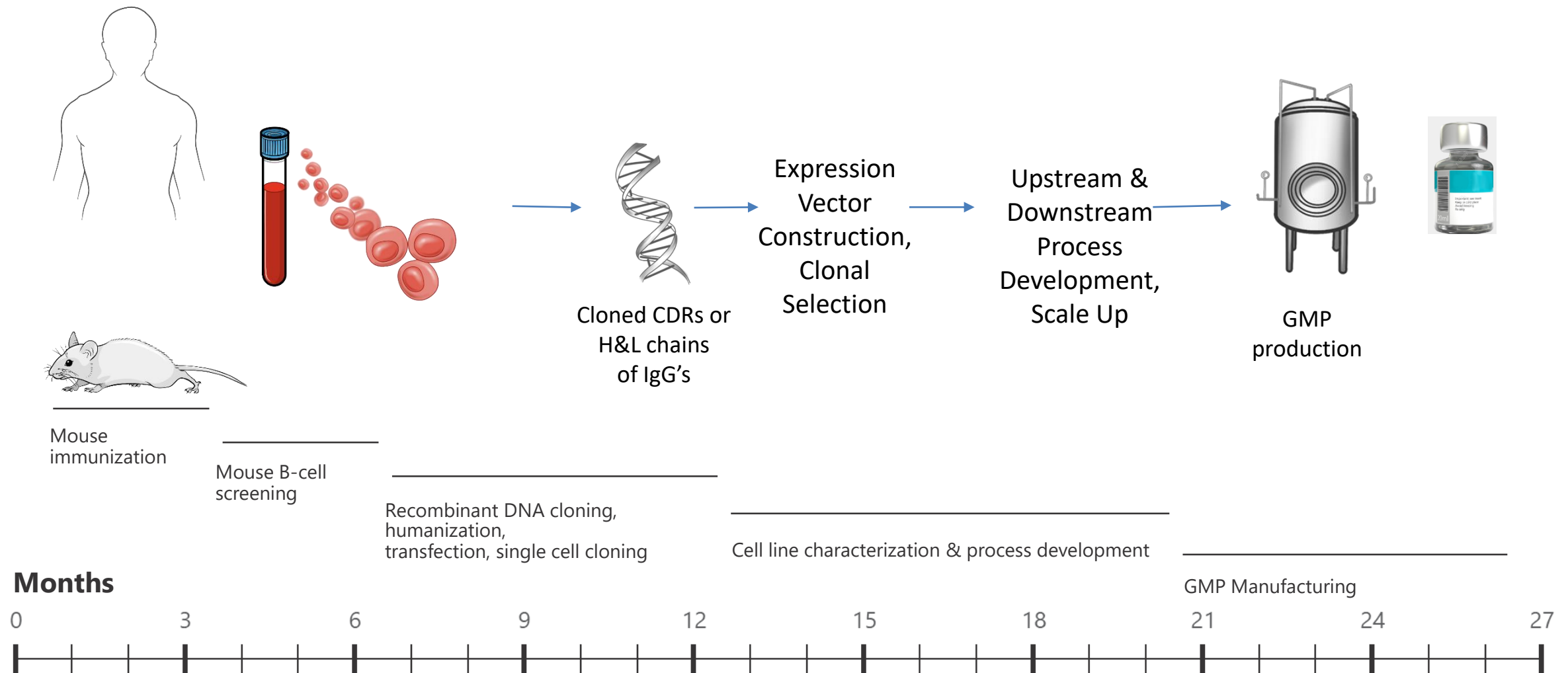




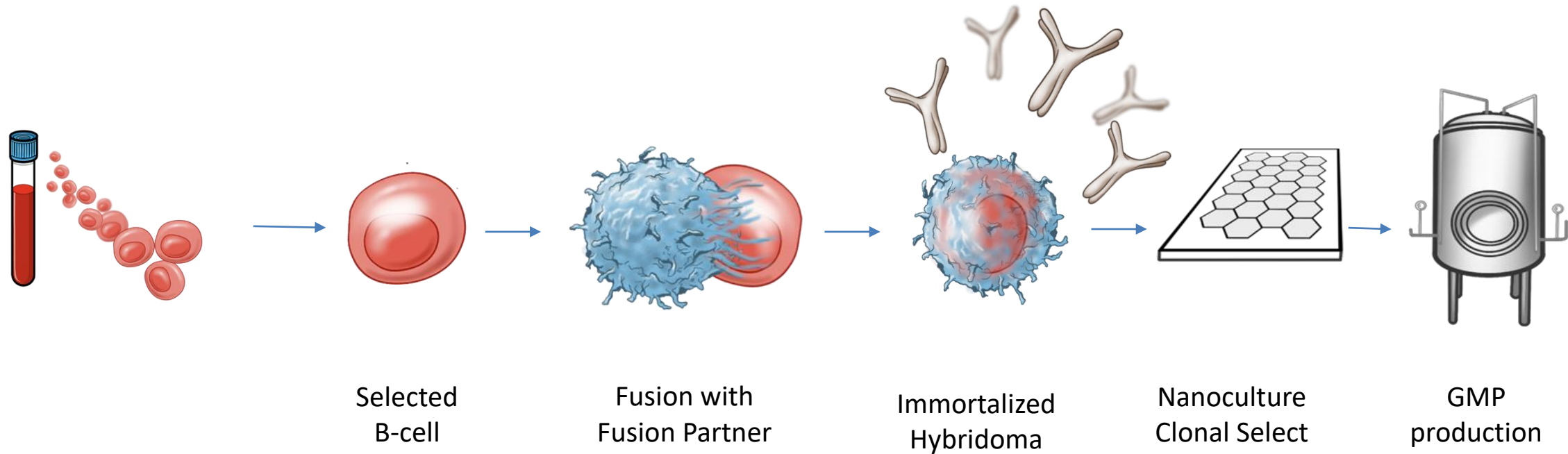
# APEX™ Technology Suite

- Single cell cloning & mAb discovery
- **B-cell immortalization using cell fusion**
- Enhanced mAb production of established cell lines
  - CRISPR-enabled activation of transcription regulatory genes
- mAb production CHO cell line technology

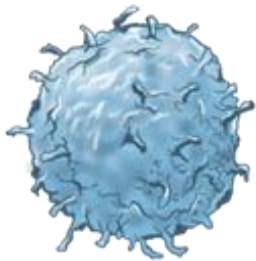
# Current Practice: Recombinant approaches to mAb discovery & production requires 1.5 - 2 years from discovery to drug in vials



# Bypassing Recombinant Approaches Using B-cell Fusion & Immobilization



# Engineering of the Fusion Partner Cell Line



Genetically engineered, non-secreting IgG mouse heteromyeloma cell line developed with high cell fusion efficiency

Genetic modifications introduced to **activate master transcription factors** associated with increased transcription and translation of H & L chain genes of monoclonal antibodies. Exhibits

- Immortality
- CHO-level productivity
- Stability

## Key Feature

Contains stable integrated CRISPR machinery

- Cas enzyme, accessory proteins, and guided RNAs necessary to activate endogenous master transcription factors

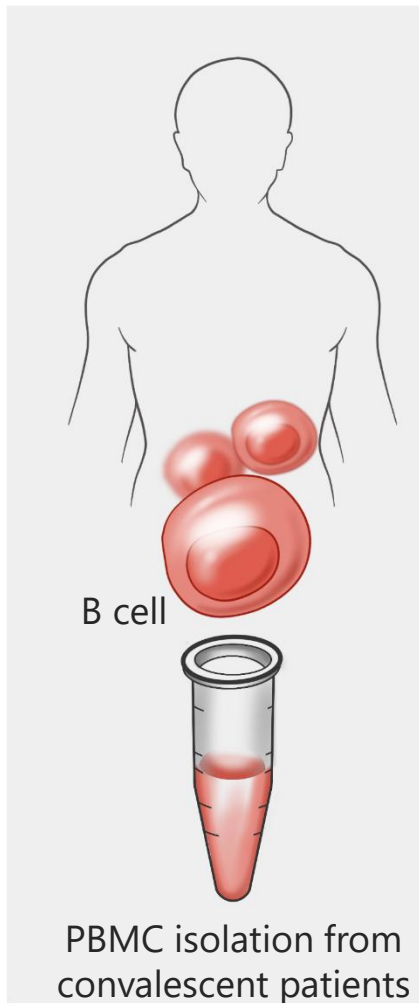
# **B-cell Screening & Cell Fusion Proof-of-Concept Study: RSV mAb Discovery**

RSV-F antigen used for PoC study

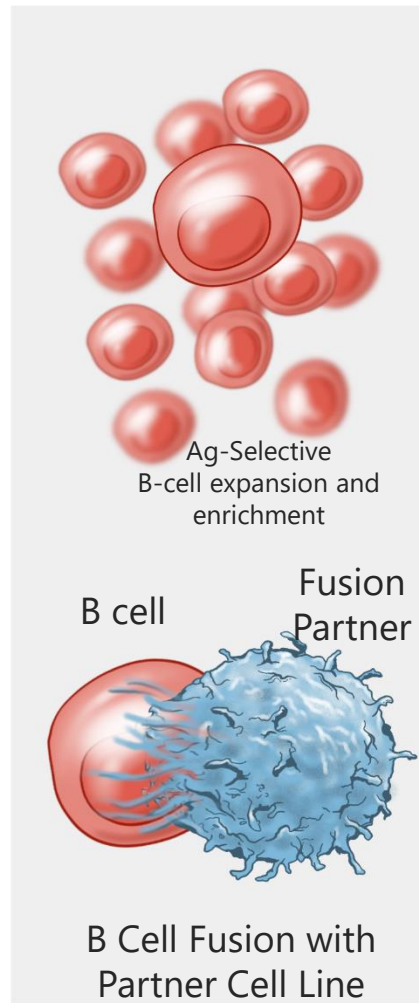
- Availability of primary RSV-F+ B-cells from biobanks
- Well described structure of RSV-F protein
- Antigen specific tools, assays and reagents are available

# Repertoire Screening & Cell Fusion Work Flow

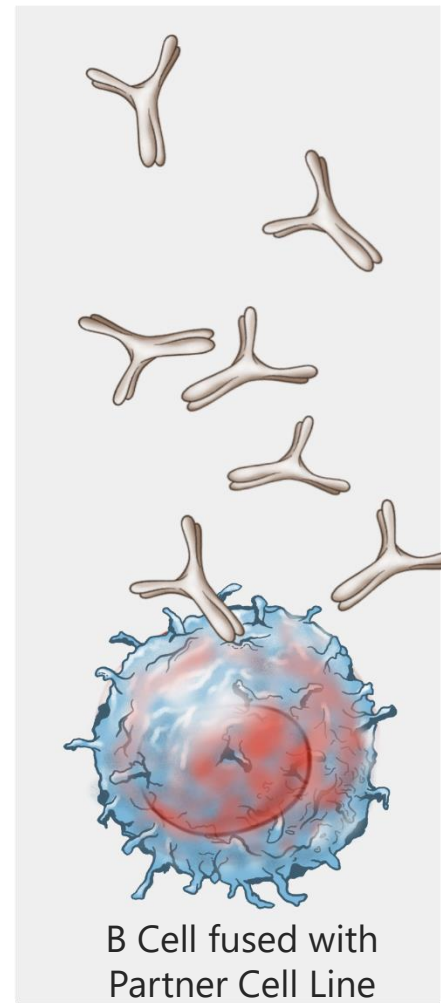
## ISOLATE PBMCs



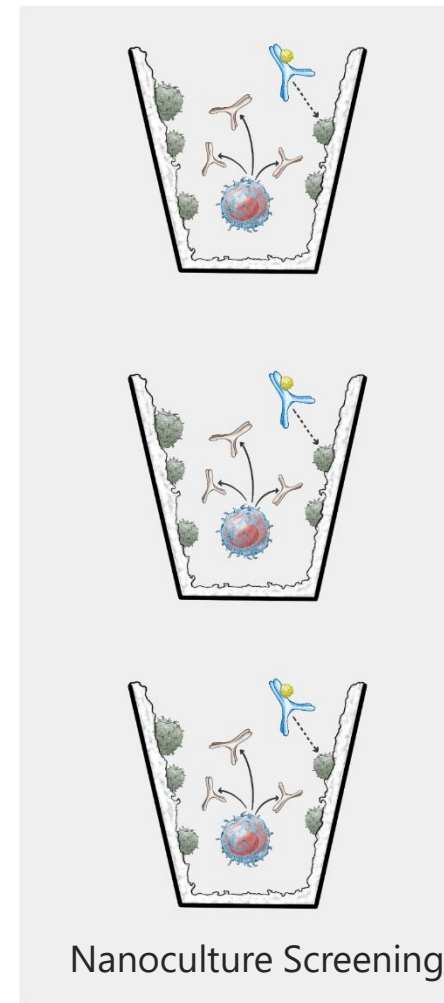
## B-CELL EXPANSION



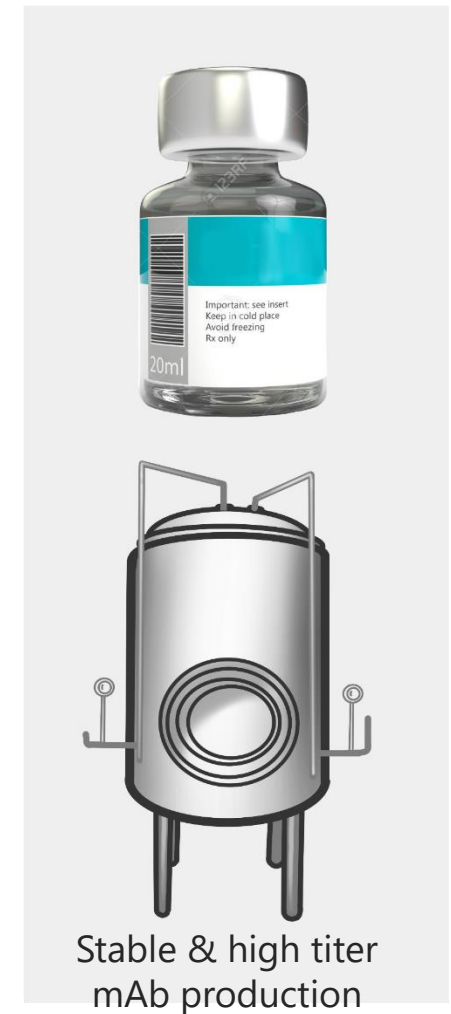
## IMMORTALIZED MAB SECRETING PROGENY



## CLONAL SELECTION



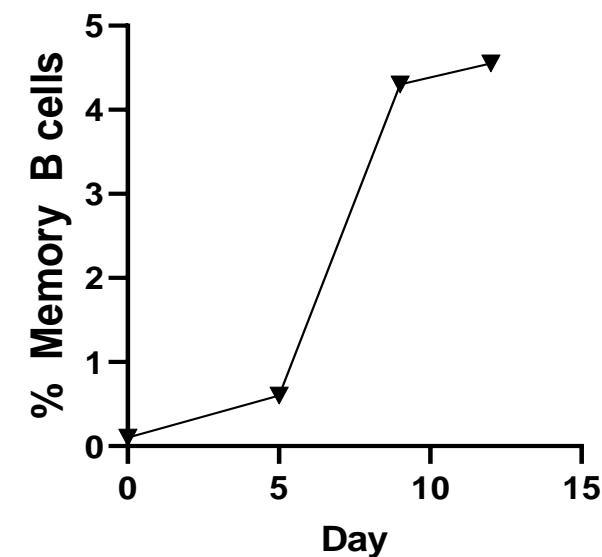
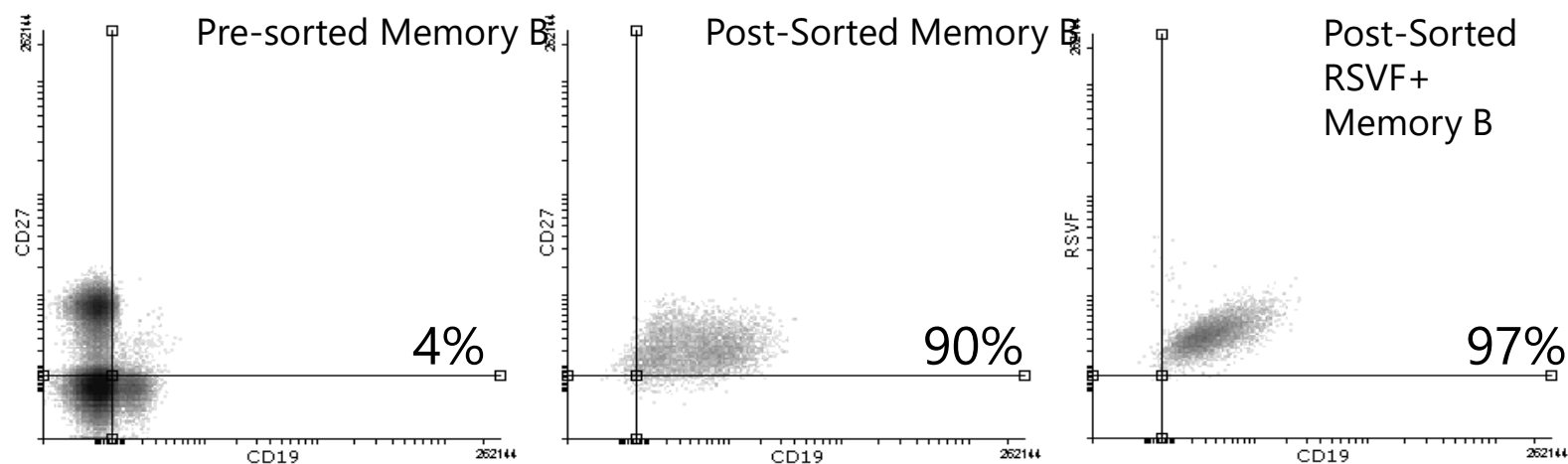
## PRODUCTION





# RSV-F Ag-specific Human B Cell Activation

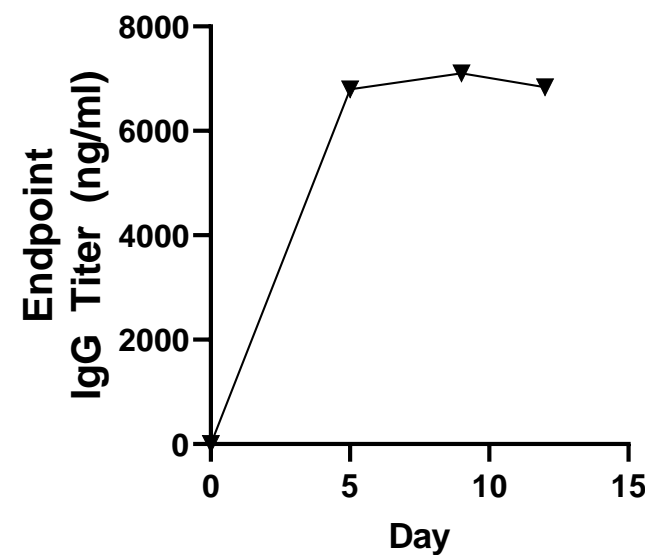
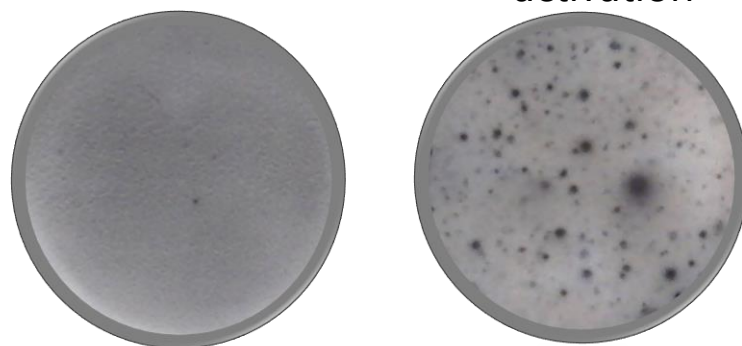
## Day 7 Ag-specific Human B-cell Activation



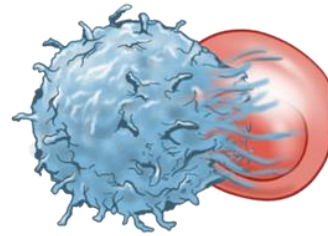
## RSV-F memory B-cell secretion

Day 7 PBMCs

Day 7 RSVF B-cell activation

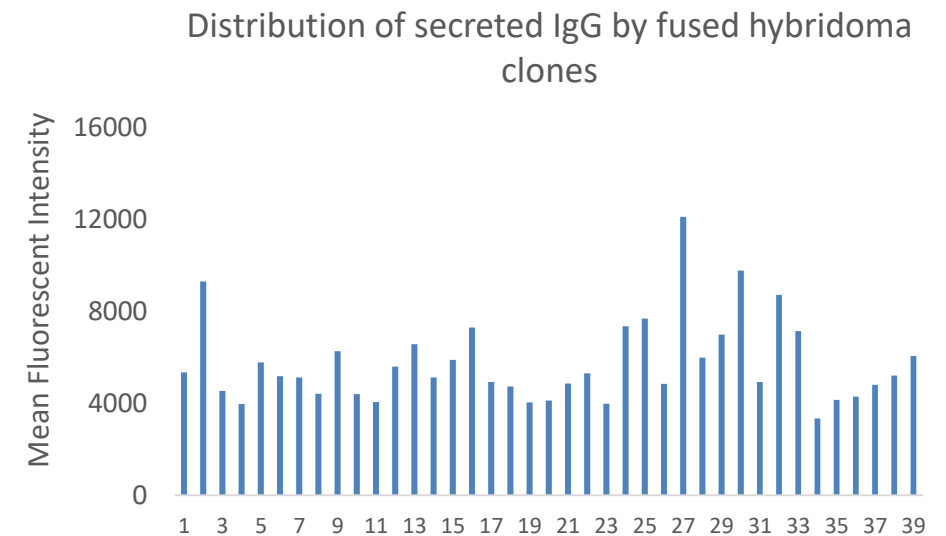
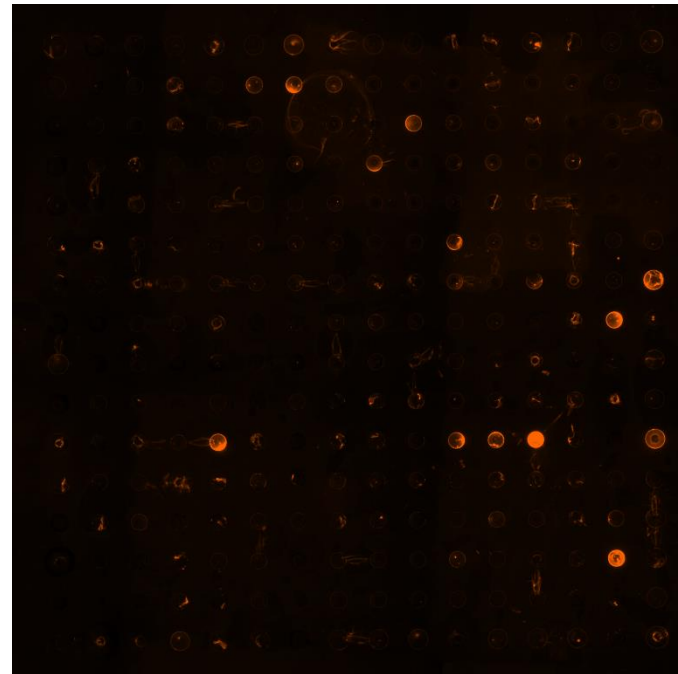
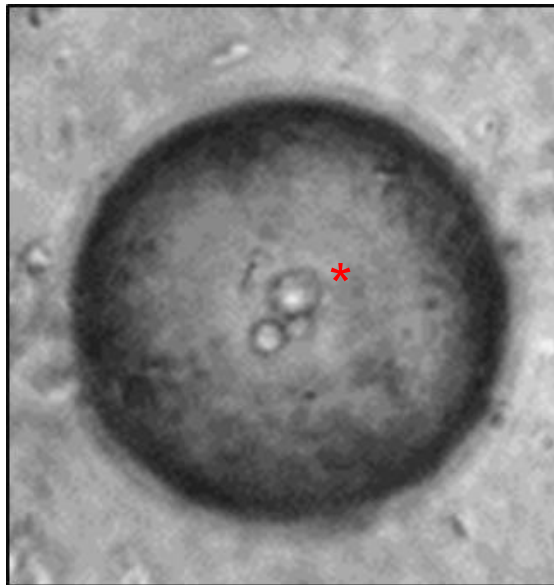


# Engineered Fusion Partner Exhibits High Fusion Efficiency



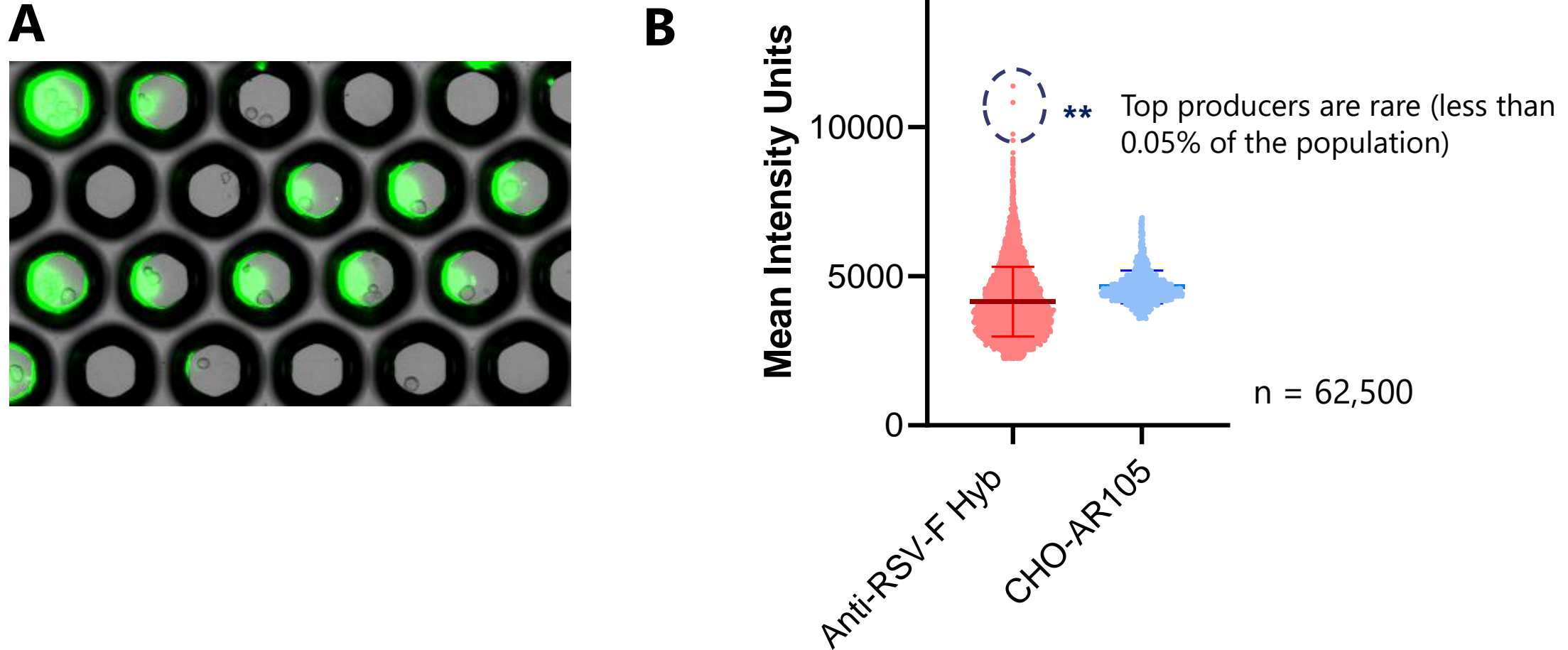
**On-chip Single Cell Fusion: 10%-20% Fusion Efficiency [i.e. 10-20% of progenies that fused, survived, and are producing mAb]**

*After Fusion*



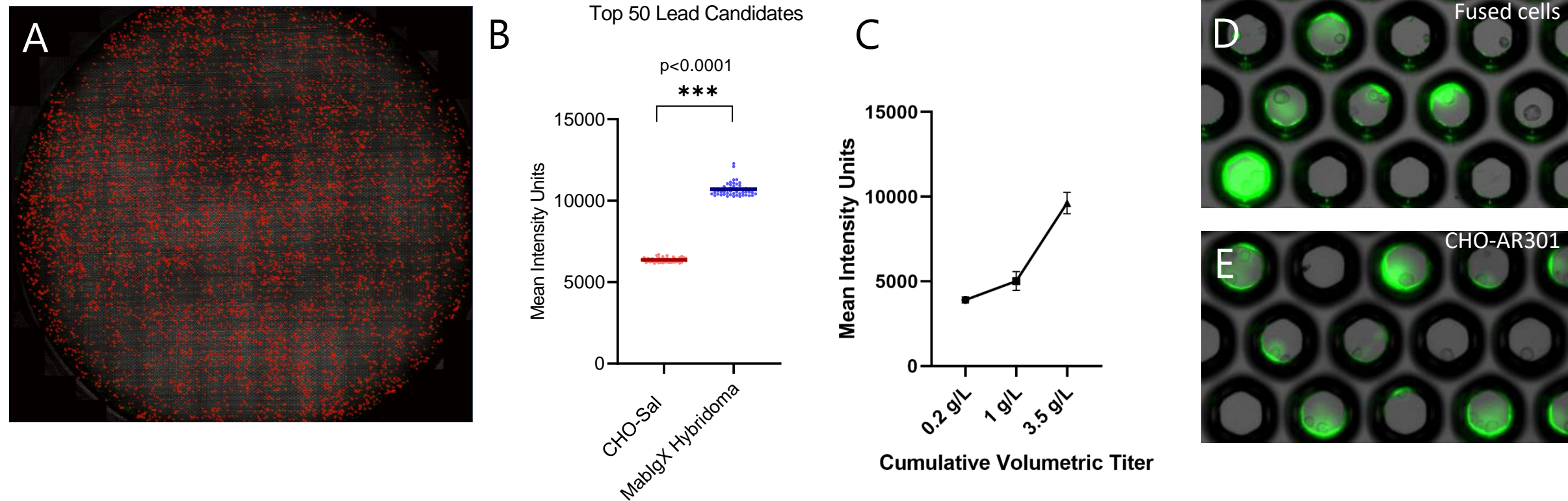
Left: 16x16 array of fused hybridoma at Day 27 post-fusion; Right: distribution of hybridoma clones that are producing IgG. Fluorescence intensity is used as a metric of IgG secretion level

# Rapid identification of high-producing RSV-F specific hybridoma



**A.** Panel of nanowells of hybridoma producing Human anti-RSVF mAb. **B.** Distribution of fluorescent ring intensity of anti-RSVF production compared to CHO-Aer. \*\* Denotes production levels higher than CHO-Aer producing at 3.5 g/L.

# High content screening allows for rapid identification of rare high-producing hybridoma clones

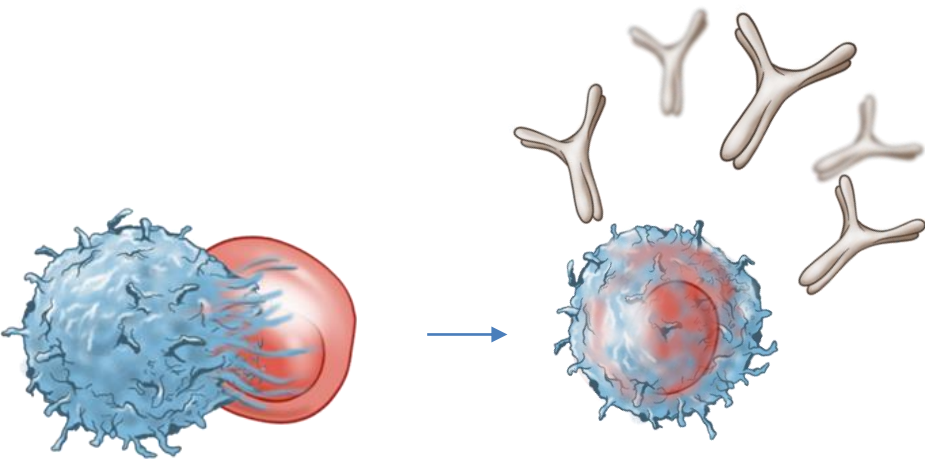
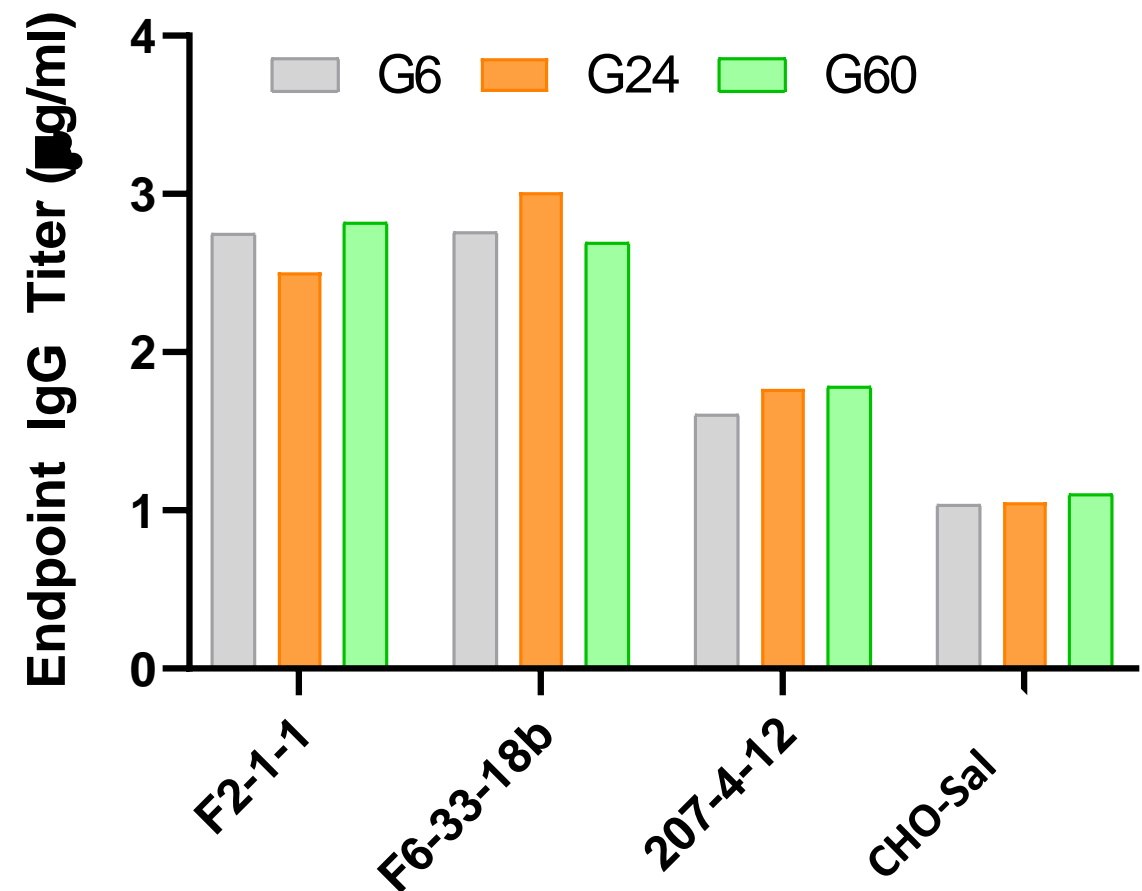


**(A)** Hybridomas generated from cell fusion compared to CHO-Sal were screened for mAb secretion on nanocultures. ~24,000 positives identified for MabIgX<sup>®</sup> hybridomas/ ~ 33,000 positives identified for CHO-Sal.

**(B)** Mean fluorescence intensity of the (B) 50 highest producing clones and **(C)** Correlation of mean intensity units to cumulative volumetric titer of cell lines with experimentally-determined productivity ( $*** p < 0.0001$  by Student's t-test,  $n = 50$ ).

**(D-E)** Representative panel showing single cell IgG production of **(D)** Hybridoma and **(E)** CHO-Sal (1 g/L volumetric titer)

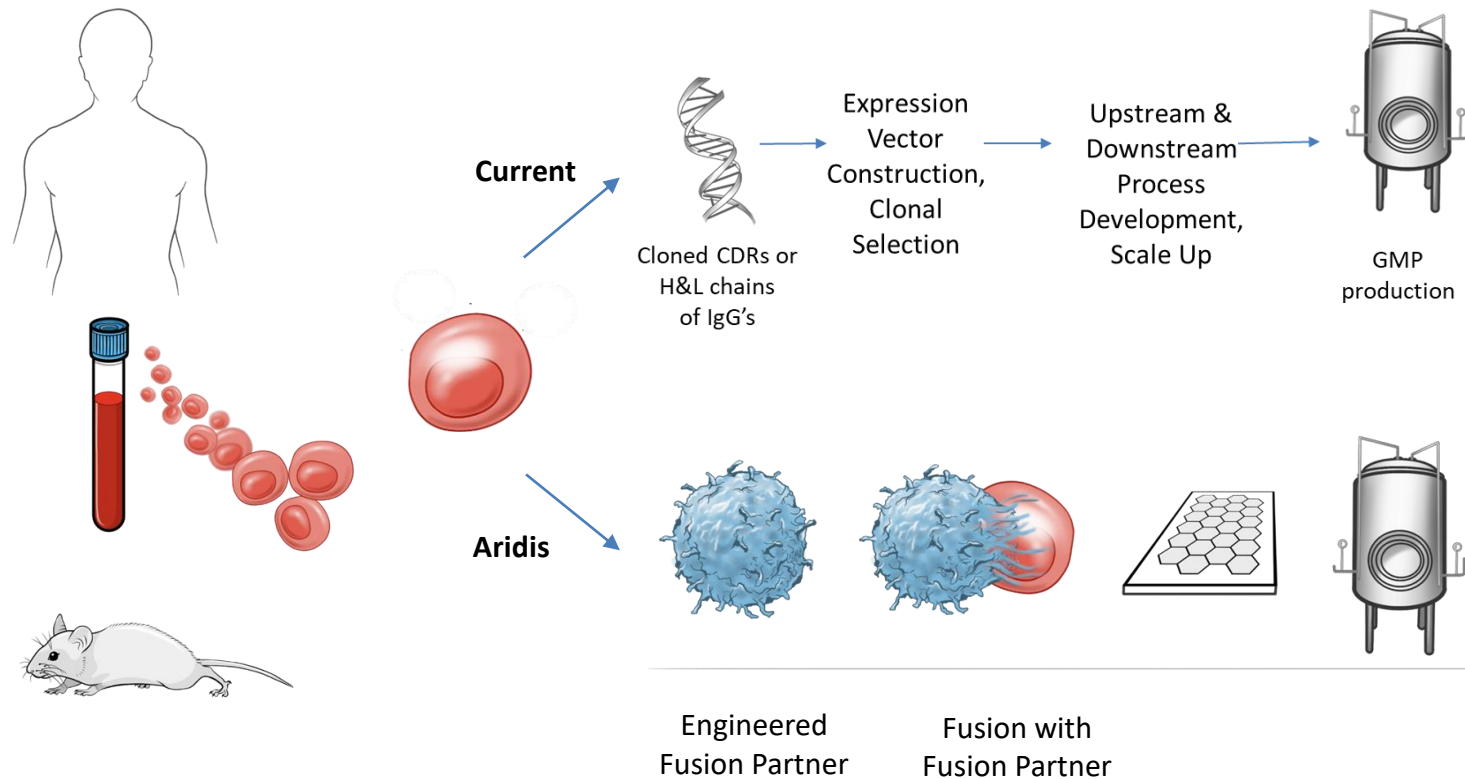
# Stable Hybridoma Productivity Comparable to CHO over 60-Generations



- Endpoint titer of 3 hybridoma clones producing human IgG demonstrates consistent productivity at each time point
- Titer at generations 6, 24, and 60 shown
- CHO-Sal producing at 1 g/L was used as a comparator for all stability studies



# Cell Fusion to Create Hybridoma: Advantages & Disadvantages



## Advantages:

- Well established process

## Disadvantages:

- Up to ~1 year longer than the hybridoma approach

## Advantages:

- Convenient B-cell starting point
- Bypasses recombinant steps
- Little to no process development needed
- Fastest to clinical manufacturing

## Disadvantages:

- Involves hybridoma\*

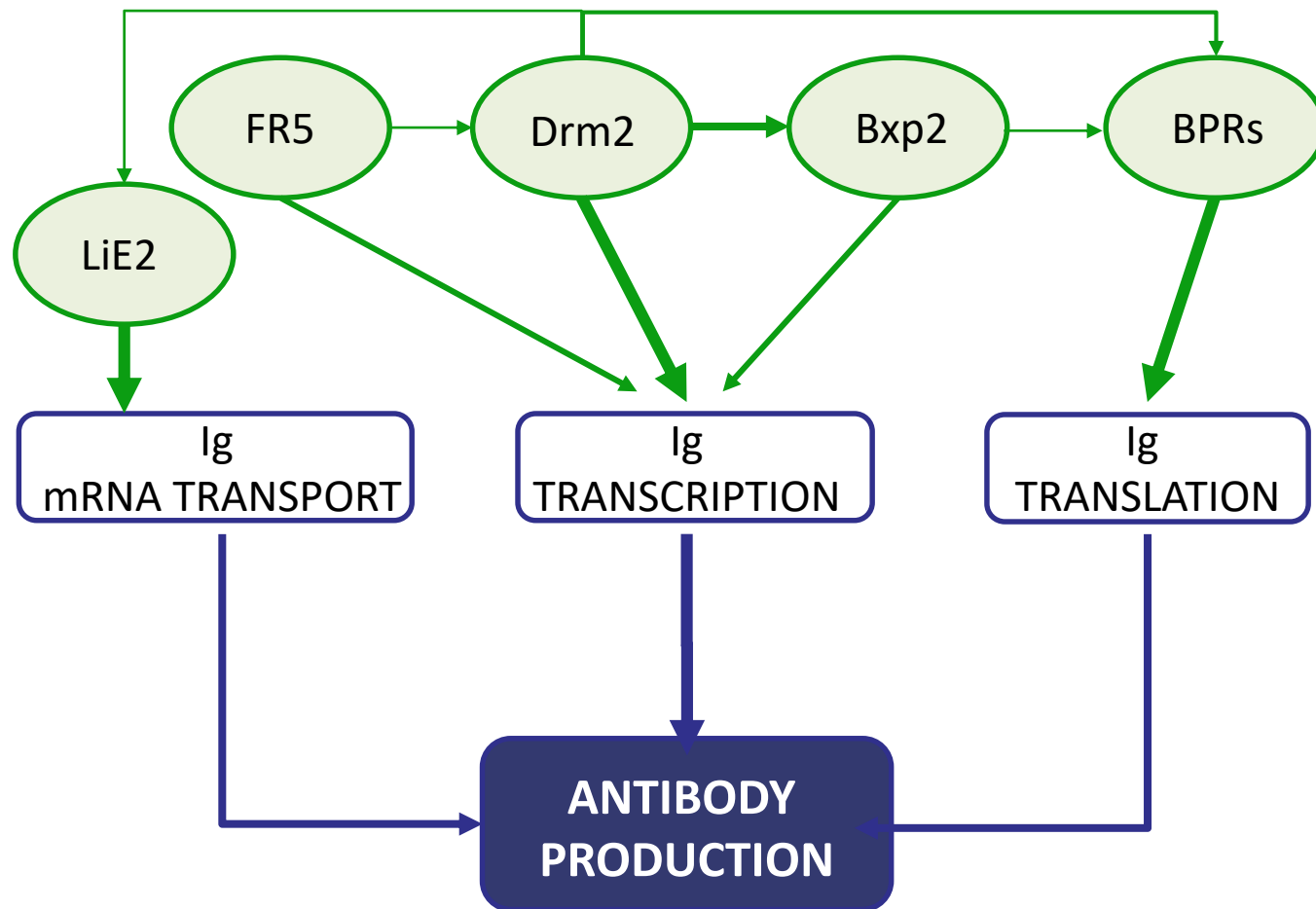
\*Can derive CHO while in Phase 1-2 clinical studies



# Outline

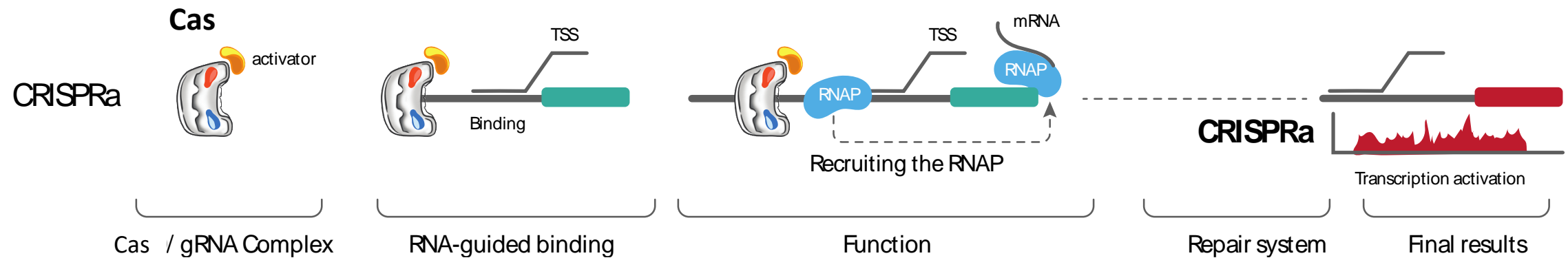
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# Key Master Transcription Factor Genes of Interests



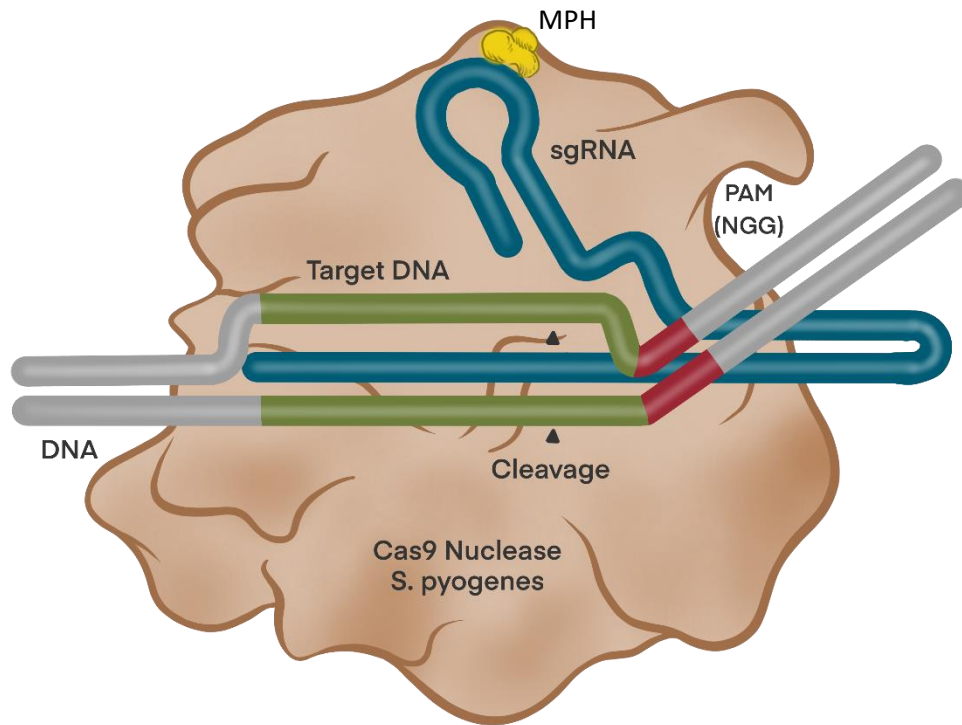
- **Transcription factor genes selected for CRISPR mediated addition & activation**
  - Drm2
  - Fr5
  - Bxp2
- **Promotor activation is universal and can be applied to any host cell irrespective of B cell lineage**

# CRISPR-Cas Targeted Gene Activation



\*Transcriptional state site (TSS), homology-directed repair (HDR), non-homologous end joining (NHEJ)

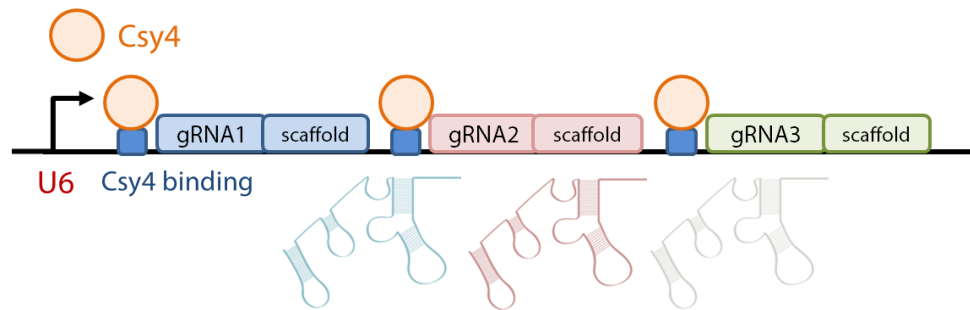
# Multiplex CRISPR Guided Activation (CRISPRa)



1. In traditional activation, open reading frames (ORF) of genes need to be cloned under a ubiquitous promoter. The size of ORF is a rate-limiting step as only 1-2 genes can be activated
2. Short gRNA (20 nt) for transcriptional activation allows for multiple concomitant activation of genes
  - Potent activation of genes from 20-100 fold
3. Stable expression of dCas-VP64 recognizes canonical protospacer motif (PAM) on gRNA targeting regions within 200bp of the transcriptional start site (TSS) of the plasma cell master regulators: Drm2, Fr5 and Bxp2.
4. Activation of the three transcription factors: Drm2, Fr5 and Bxp2 trigger a cascade of events to enhance antibody production:

# Multiplex TF Activation by CRISPRa

Single Module



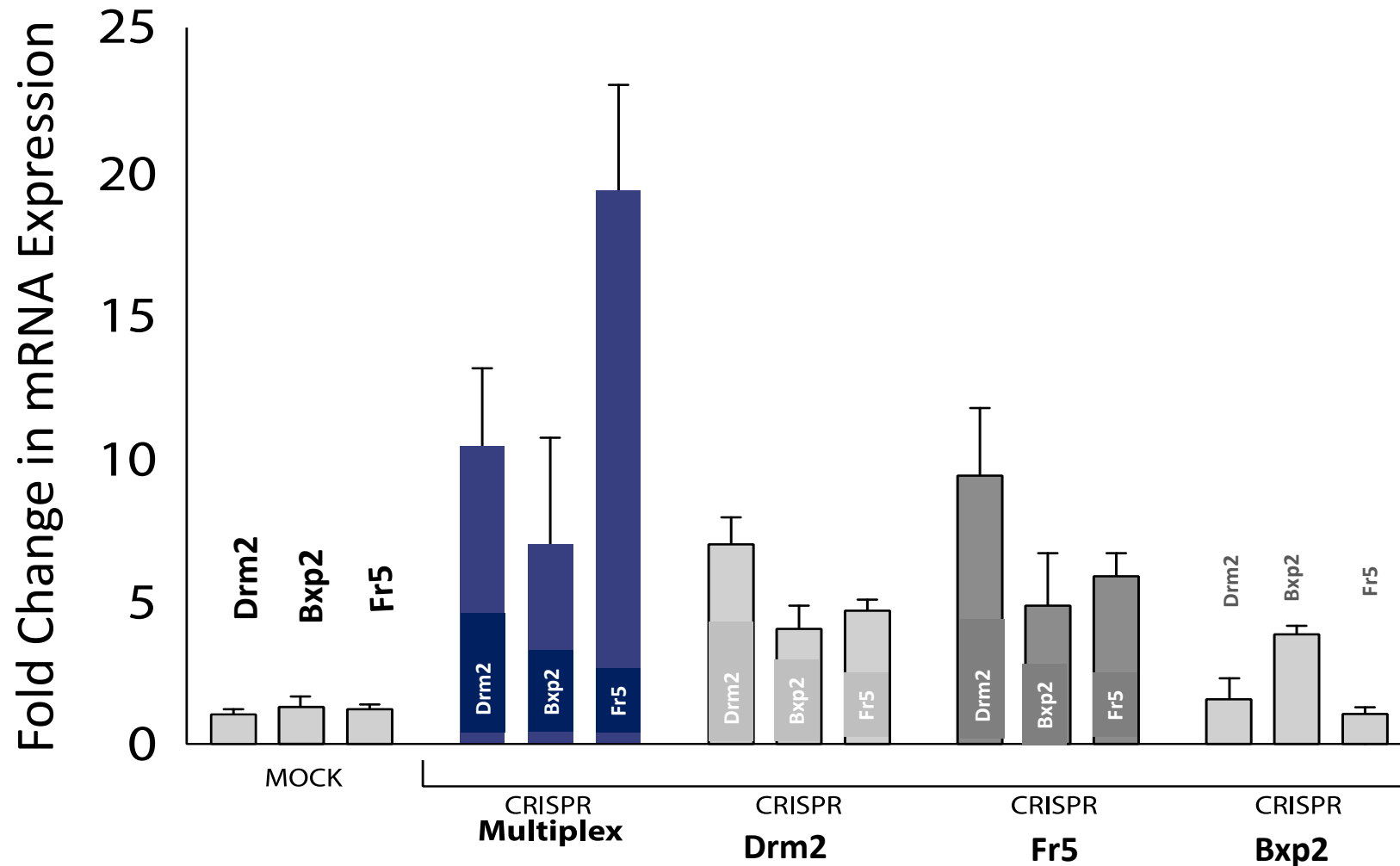
## Current Challenges

- gRNA that allow for potent transcriptional activation needs to be empirically determined
- Not all gRNAs will activate gene transcription
- Multiple transfection and transduction are required for a single gene
- Current CRISPRa technologies at best can only activate two gRNAs at a time

## Aridis Multiplex CRISPRa

- Utilizes a unique and modular gRNA scaffold design for multiple processing of gRNAs in a single vector
- Synergistic activation of multiple genes in the sample molecular pathway superior to a single gRNA activation

# Potent synergistic activation of transcription factors in Fusion Partner by multiplex guided RNAs



## Synergistic activation of master transcription factor genes

- Predicted synergistic activation of targeted transcription factors with single gRNA
- Multiplex approach showed higher transcriptional activation transcription factors compared to single gRNA activation

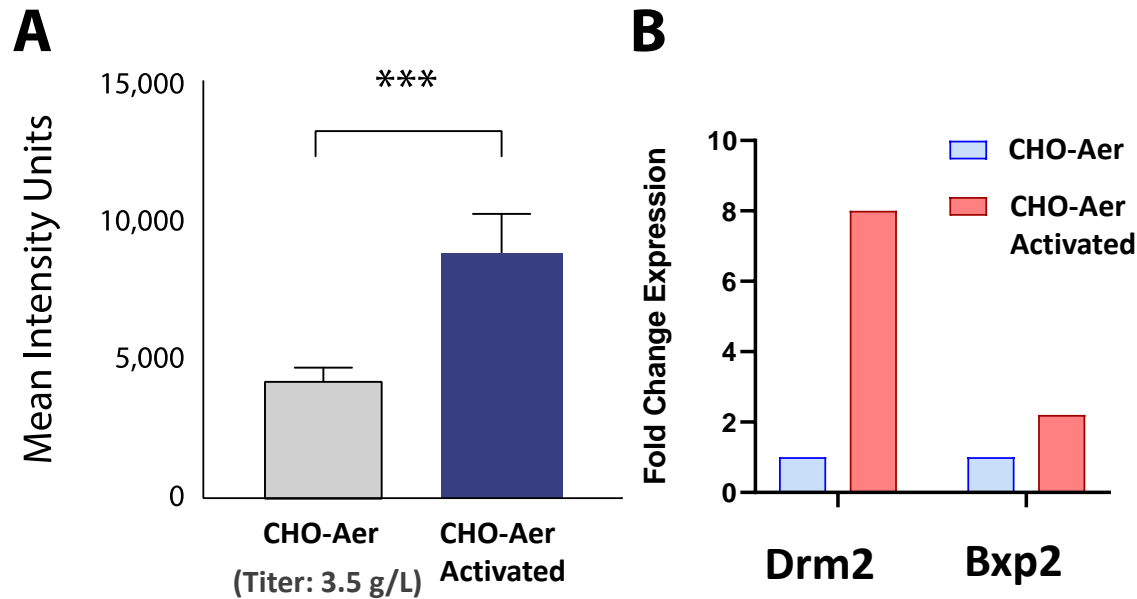


**Hypothesis: CHO production cell lines have similar master transcription regulatory elements, but are not comparably activated or utilized as in B-cells**

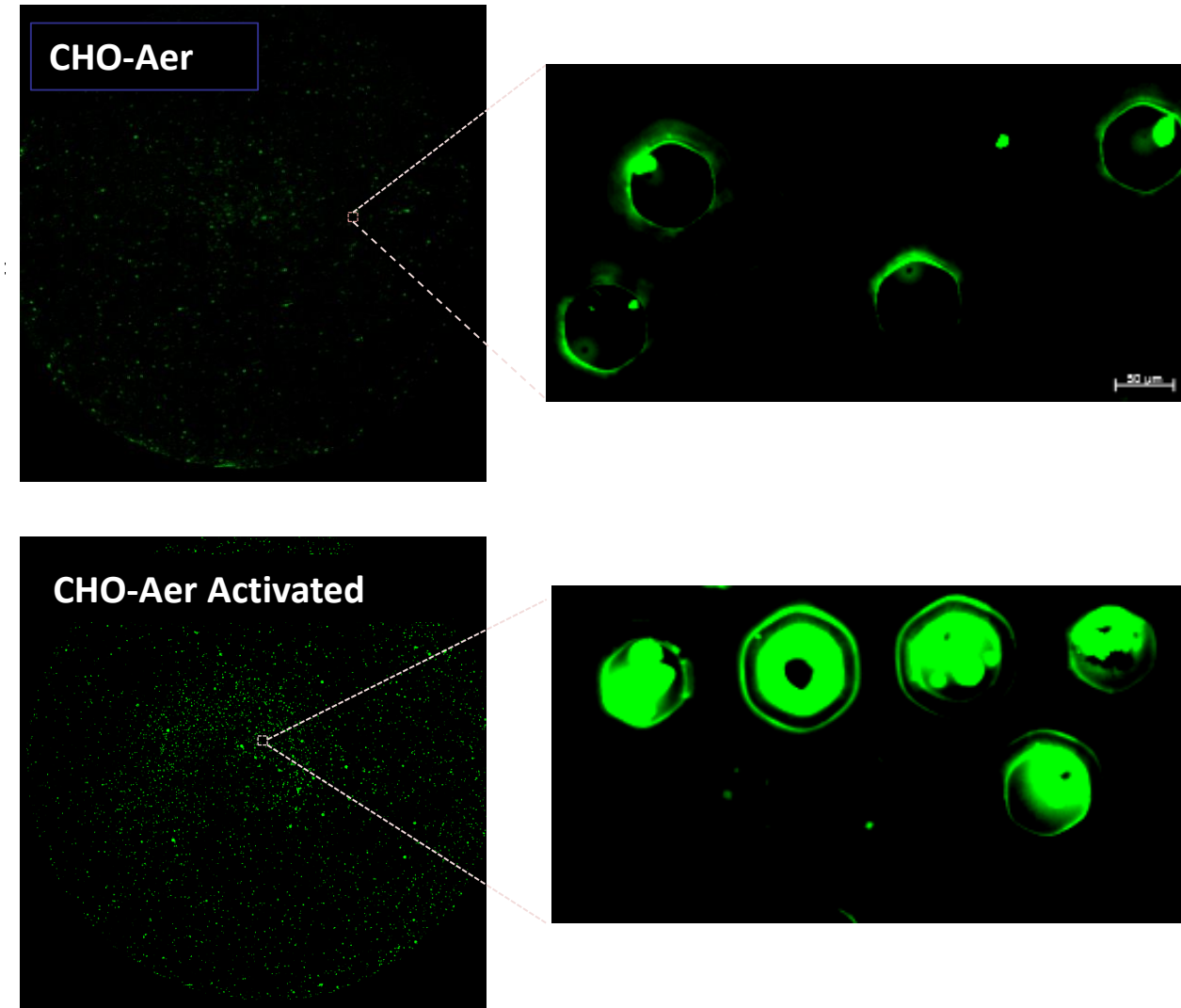
**Question: Can B-cell MTREs Enhance Productivity of Existing mAb Production CHO Cell Lines?**



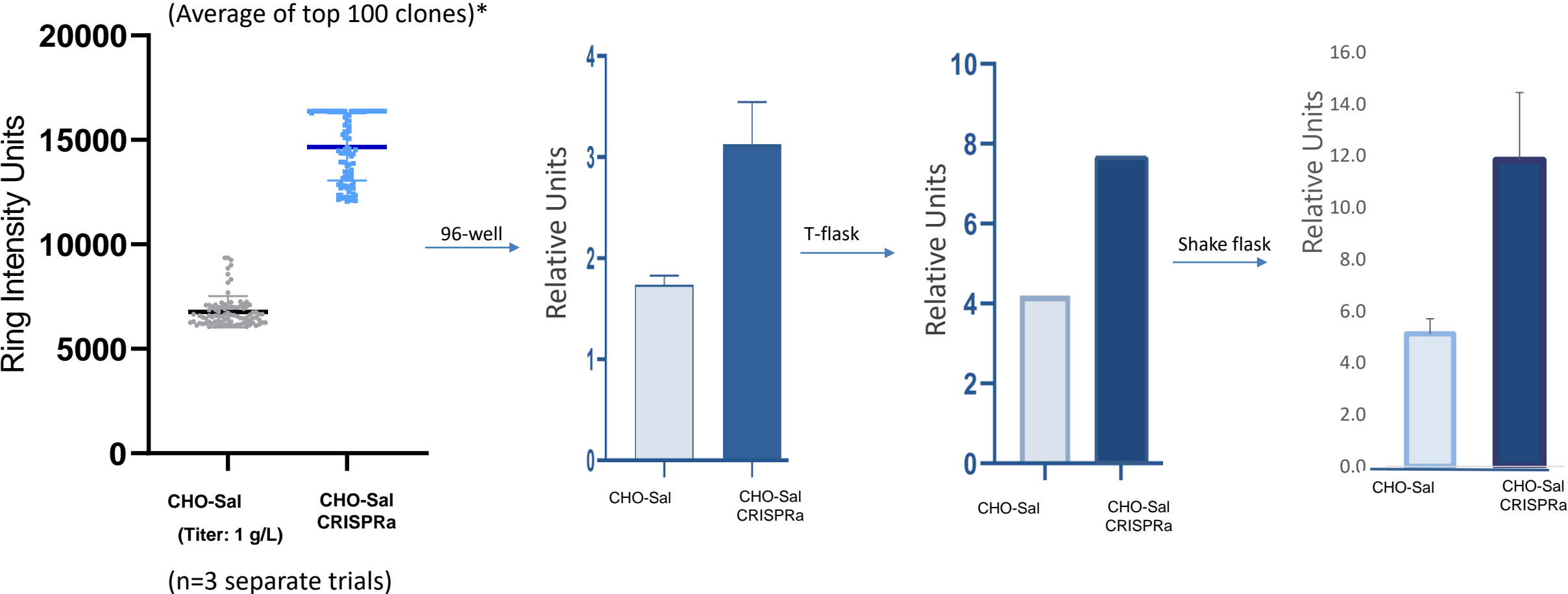
# CRISPR-Activation of Endogenous Master TF Regulatory Elements Increases Productivity a CHO Production Cell Line



**A-B.** Significant increase in single cell IgG production measured using nanoculture arrays. **B.** Expression of Drm2 & Bxp2 in CRISPRa CHO **C.** Representative panel. Right panels show an inset of single well containing 100,000 nanowells. Data represents average titer over 100,000 wells. \*\*\*  $p < 0.0001$  by Student's t-test,  $n = 100,000$ .



# Case Study #2: CRISPR-Activation of Endogenous Master Transcription Regulatory Elements of CHO-Sal Cell Line

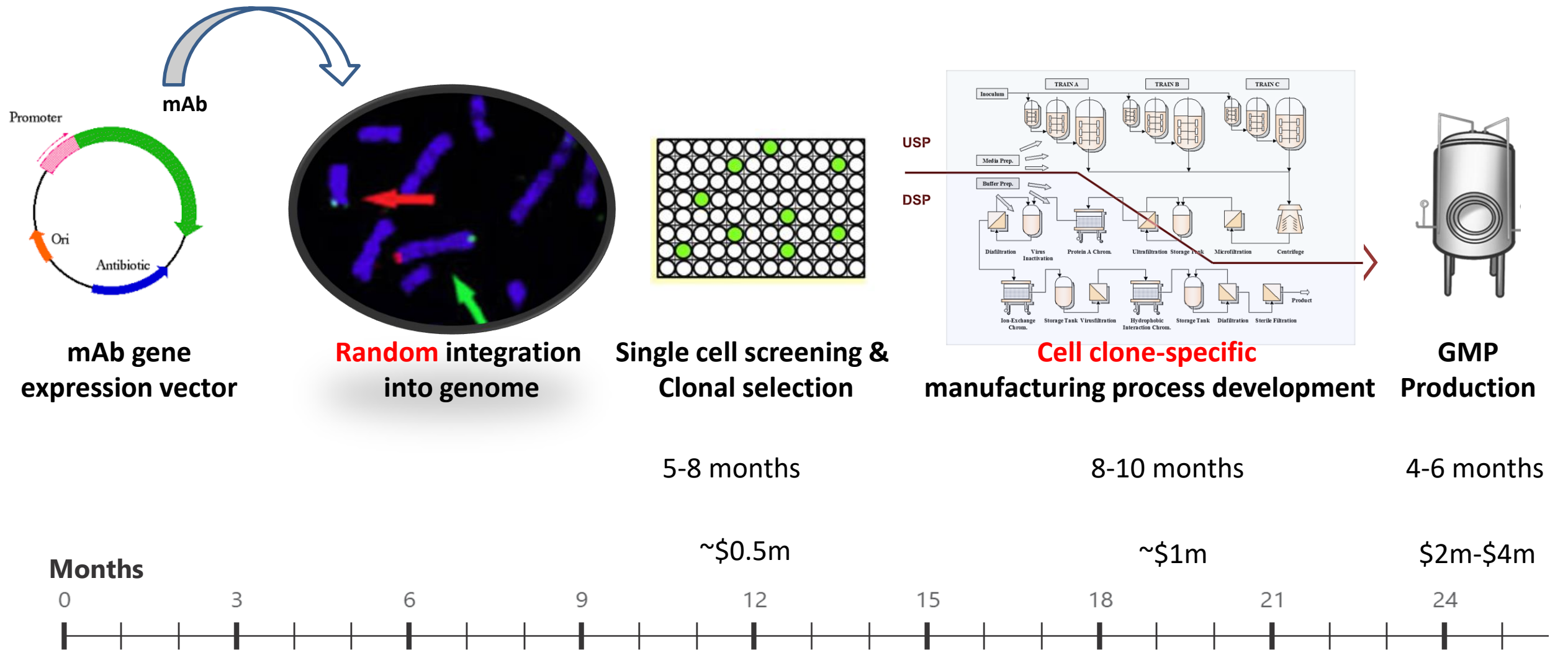


\*Pools of APEX CHO-Sal after 3 weeks selection in antibiotics were seeded on nanowells and top 50 clones were isolated using micromanipulator. E2 clone was further subcloned and expanded for 3 weeks in CHO CD medium. Following expansion, CHO-Sal and APEX CHO-Sal (E2) clone was plated on nanowells for IgG Diffusion Assay. Figure on left shows top 100 lead clones ring intensity distribution after 1 day for original CHO-Sal (volumetric titer: 1 g/L) and stable APEX CHO-Sal (E2 clone) with stable MTRE

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# Current State of the Art in mAb Manufacturing



## Product Concept:

- *Engineered CHO cell lines with CRISPR-mediated transcription activation harboring a full length mAb H/L genes that is designed to be swapped out for the CDR or H/L of interest*
  - *Integration sites and H/L gene copy numbers are known and unchanged*
- *CDR or H/L can be switched from the cell line without significant modification of established upstream & downstream manufacturing processes*



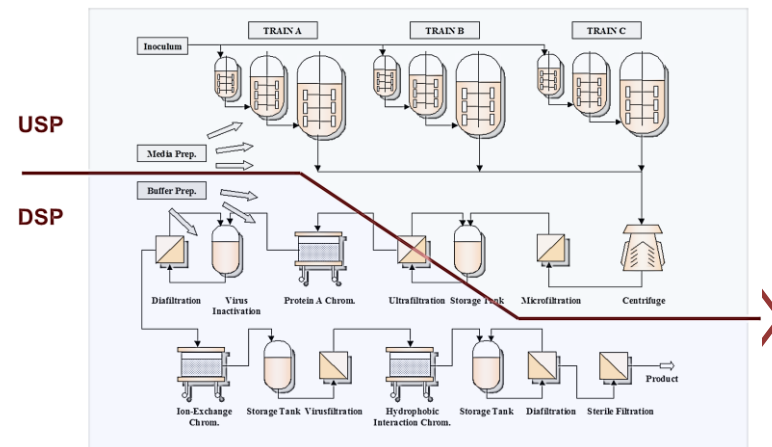
# APEX<sup>s</sup> BREATH<sup>TM</sup> CHO Master Cell Line:

## Development of a Manufacturing Process Template

Master Cell Line  
Harboring IgG<sub>1</sub>

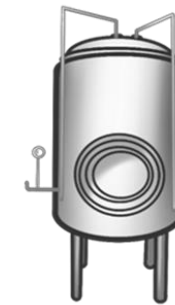


✓ MTRE activated  
By CRISPR



✓ Manufacturing Process  
to be Developed

✓ Scalability to  
GMP Manufacturing to be  
demonstrated

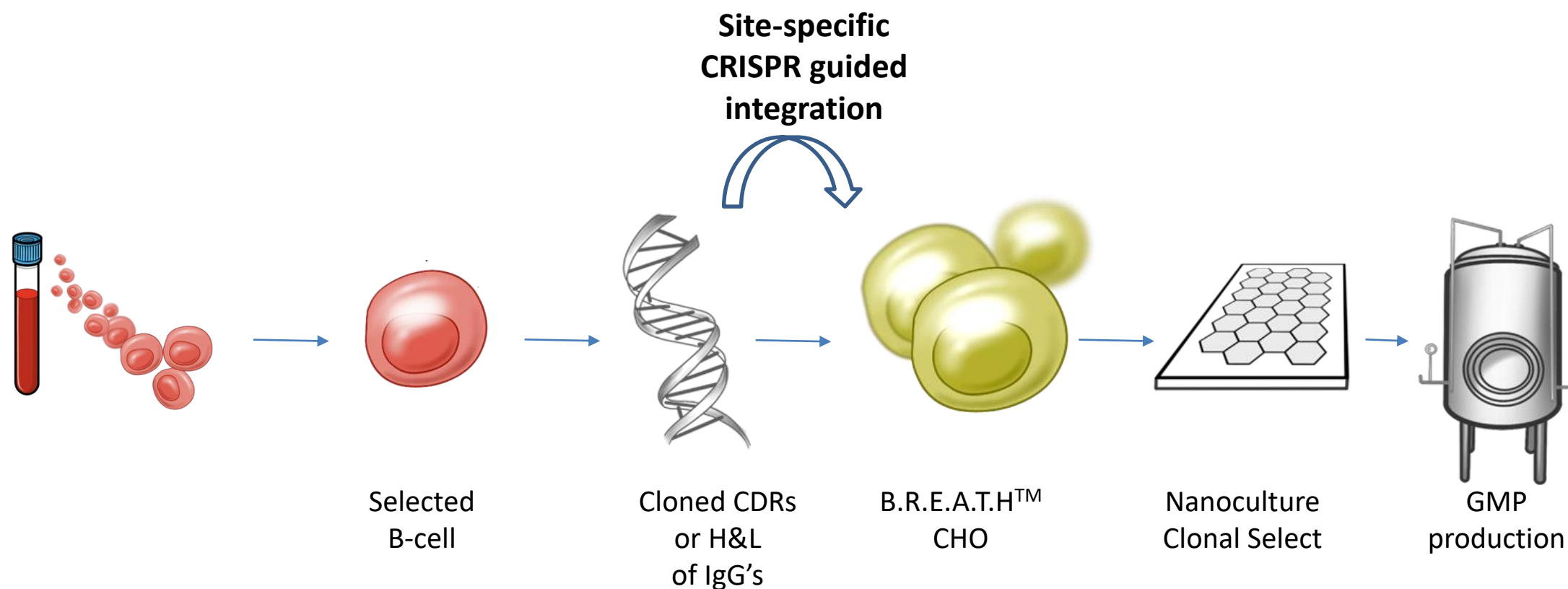


Ready for IgG<sub>1</sub>  
Gene Swap

B.R.E.A.T.H<sup>TM</sup> B-cell Regulatory Elements Assisted Transcription Host

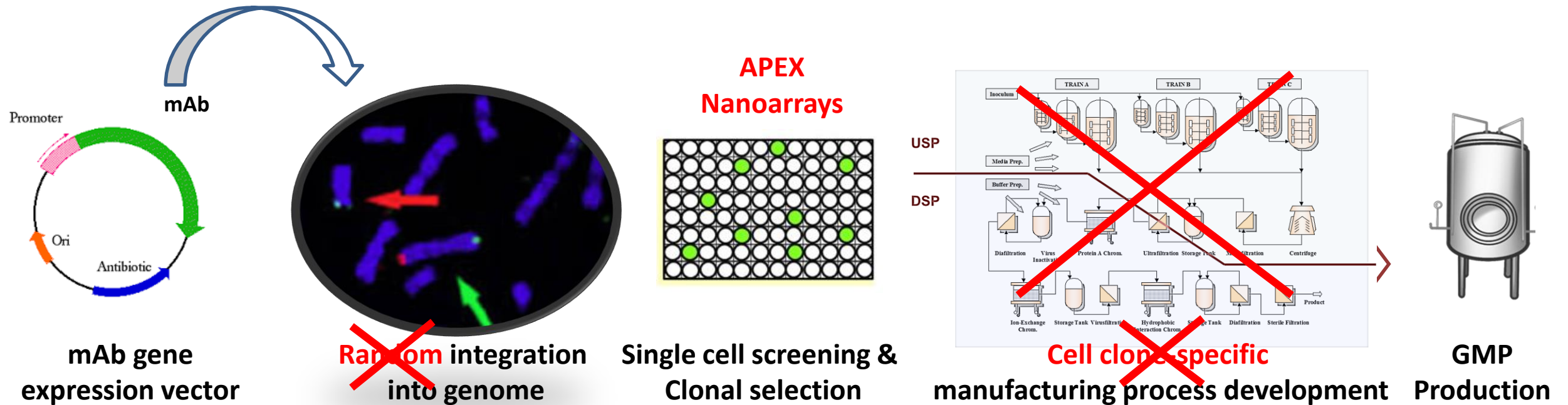
# APEX™ CHO Cell Line: CDR or H/L Chain Swappable

## CHO Host



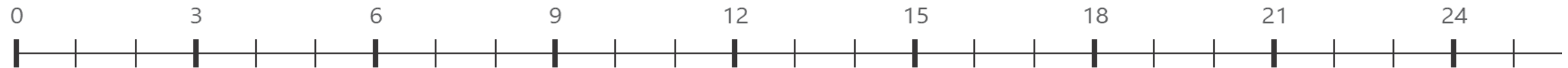
B.R.E.A.T.H™ B-cell Regulatory Elements Assisted Transcription Host

# APEX™ Technology Solutions



APEX: Discovery, Development, and Manuf.

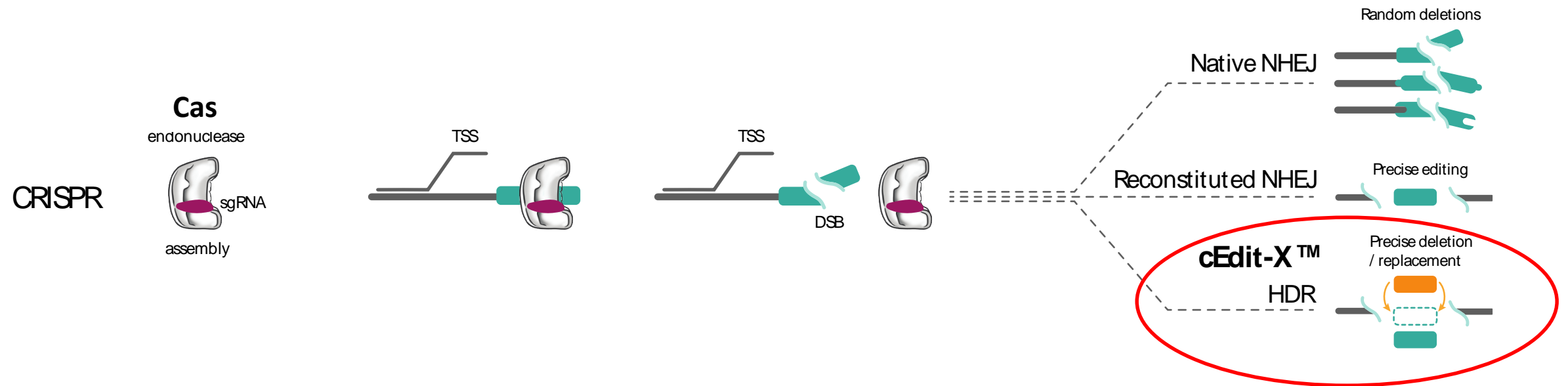
Months



~12-15 months time saving

~\$0.8m - \$1.2m cost saving

# CRISPR-Cas Targeted Gene Replacement



\*Transcriptional state site (TSS), homology-directed repair (HDR), non-homologous end joining (NHEJ)

# Engineering of CHO with B-cell Master Transcription Factors (B.R.E.A.T.H<sup>TM</sup> CHO)



Genetically engineered transfection-ready production CHO cell line harboring H & L chains of a human IgG1 and B-cell master transcription factor regulatory elements

## **Proprietary CRISPR-Mediated Activation**

Genetic modifications introduced to activate transcription factors to promote productivity and stability

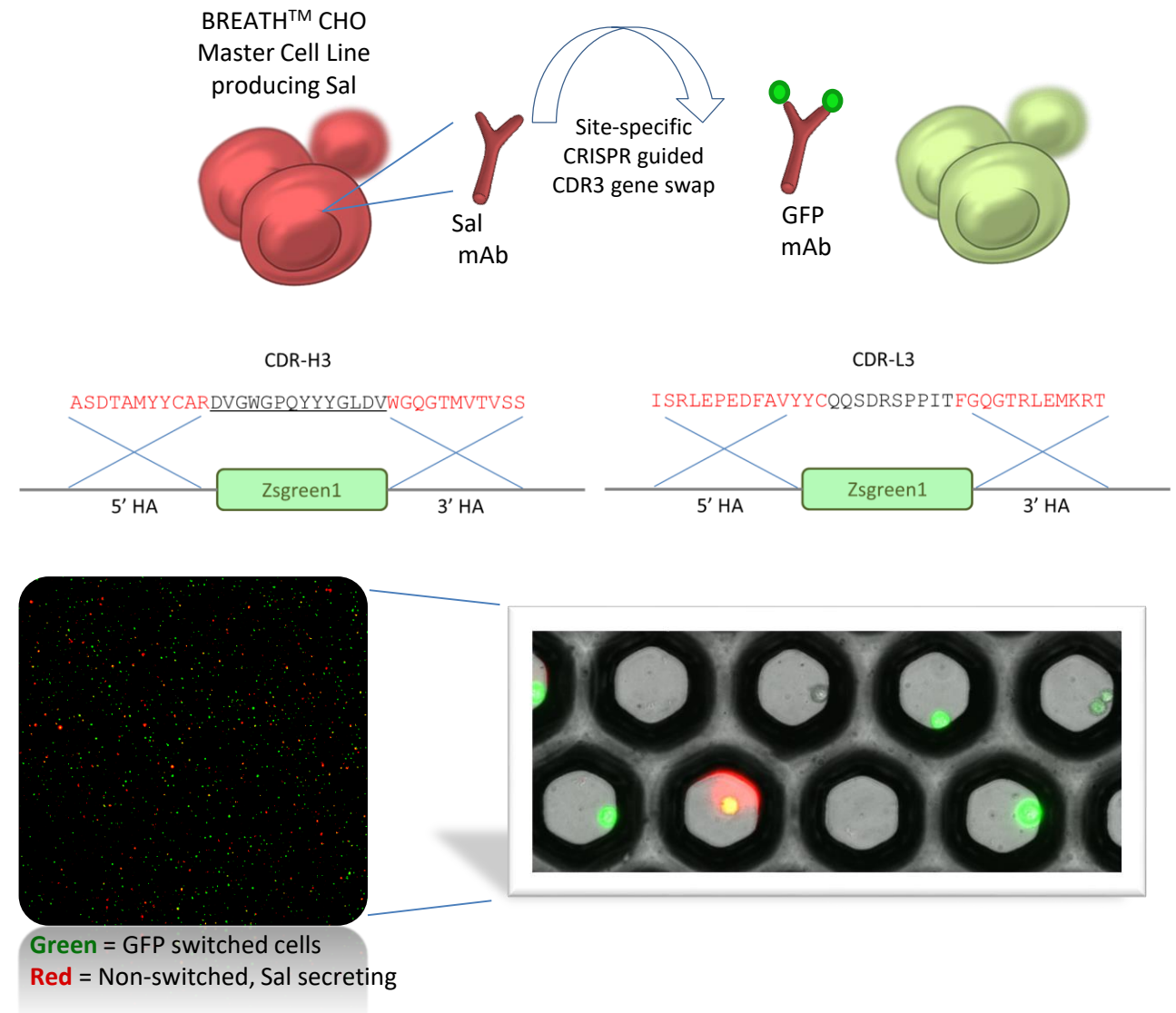
- High productivity
- Stability

## **Engineered for Productivity**

- Engineered as a host cell line to accommodate cloning of human H & L chain CDRs or full-length H & L chains by homologous recombination
- Proprietary vector system allows for dual promoter, high-copy amplification of plasmid DNA

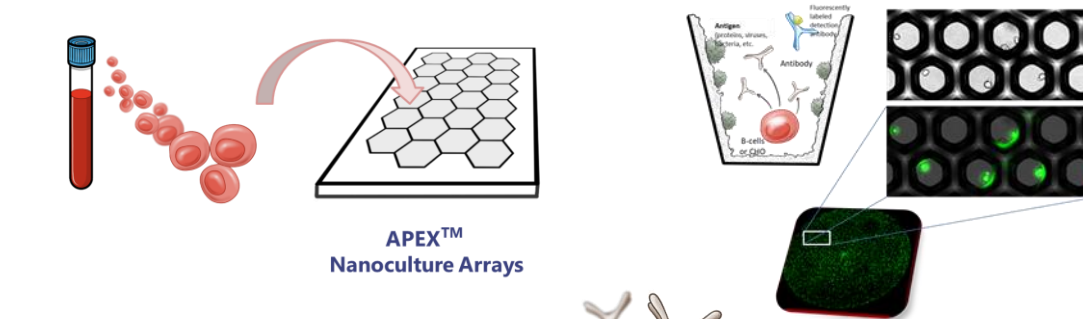
# Target Specific Insertion of Heavy and Light Chain CDR3 Genes

Aridis' approach technology utilizes CRISPR/Cas to achieve site specific integration of antibody heavy and light chain genes for maximum productivity

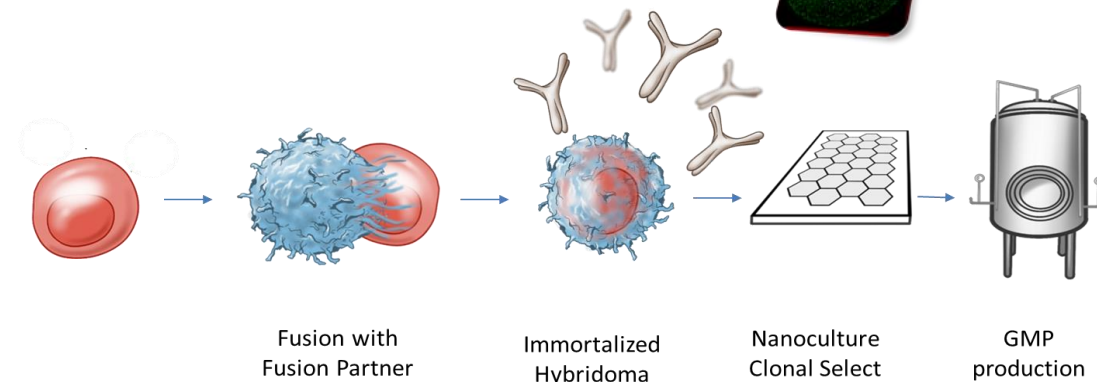


# APEX™ Technology Suite

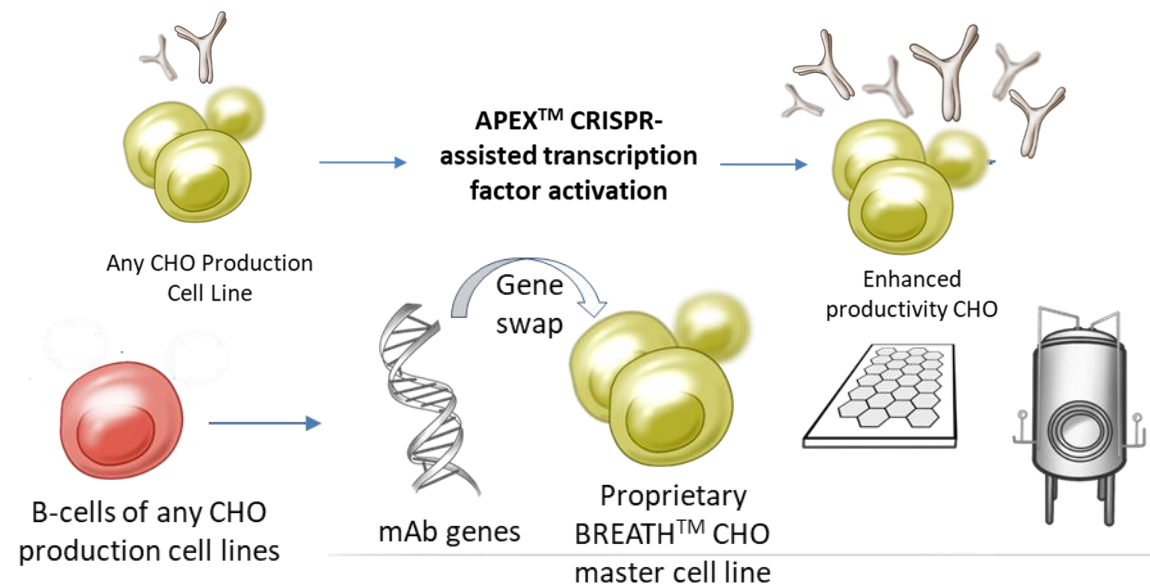
## B-cell Repertoire Screening: APEX™ NanoArrays



## B-cell immortalization using cell fusion



## mAb Productivity Enhancement: APEX™ CRISPR-assisted TF activation



## BREATH™ CHO Master cell line designed for CDR swapping