

# Leap In Transposase Platform

*The power of the pool*

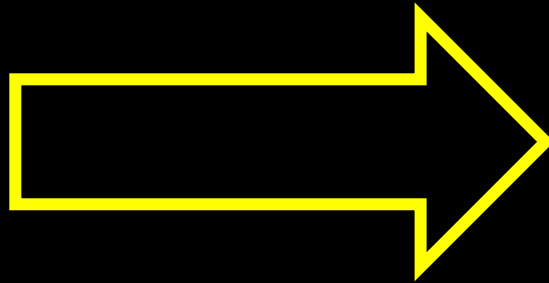


Oren Beske, Ph.D.  
obeske@atum.bio



# Transposase – Transposon: Mechanistic detail

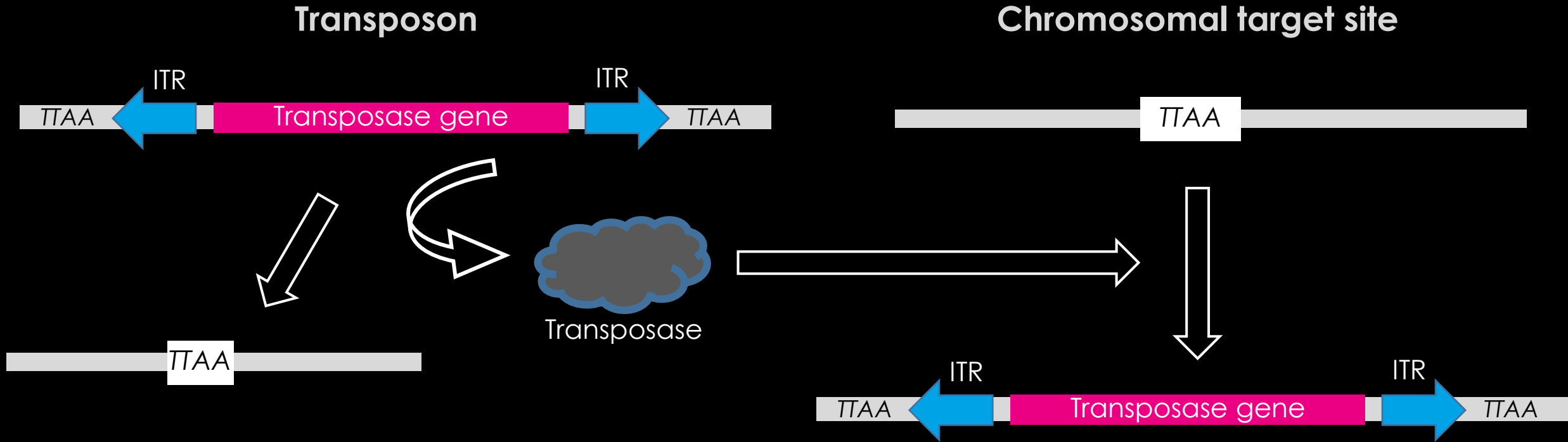
Cut



Paste



# The life of a transposon-transposase pair

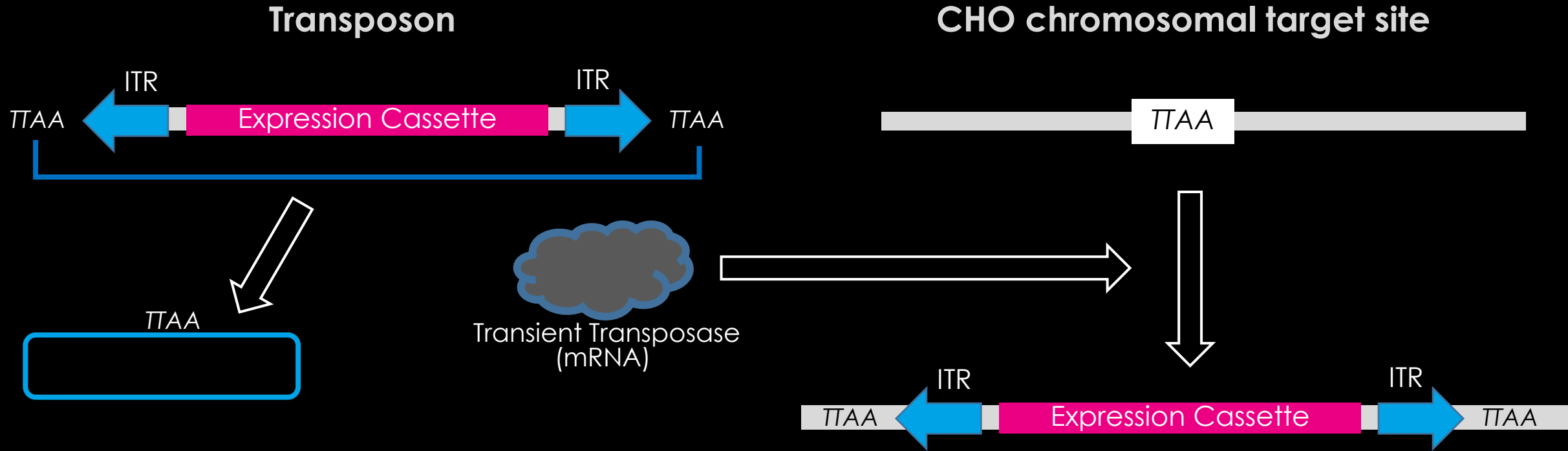


- Billions of years of evolutionary history
- Cut-paste mechanism
- Single copy integration at each site
- Perfect integration of elements between ITR's



1983 Nobel Prize in Physiology or Medicine

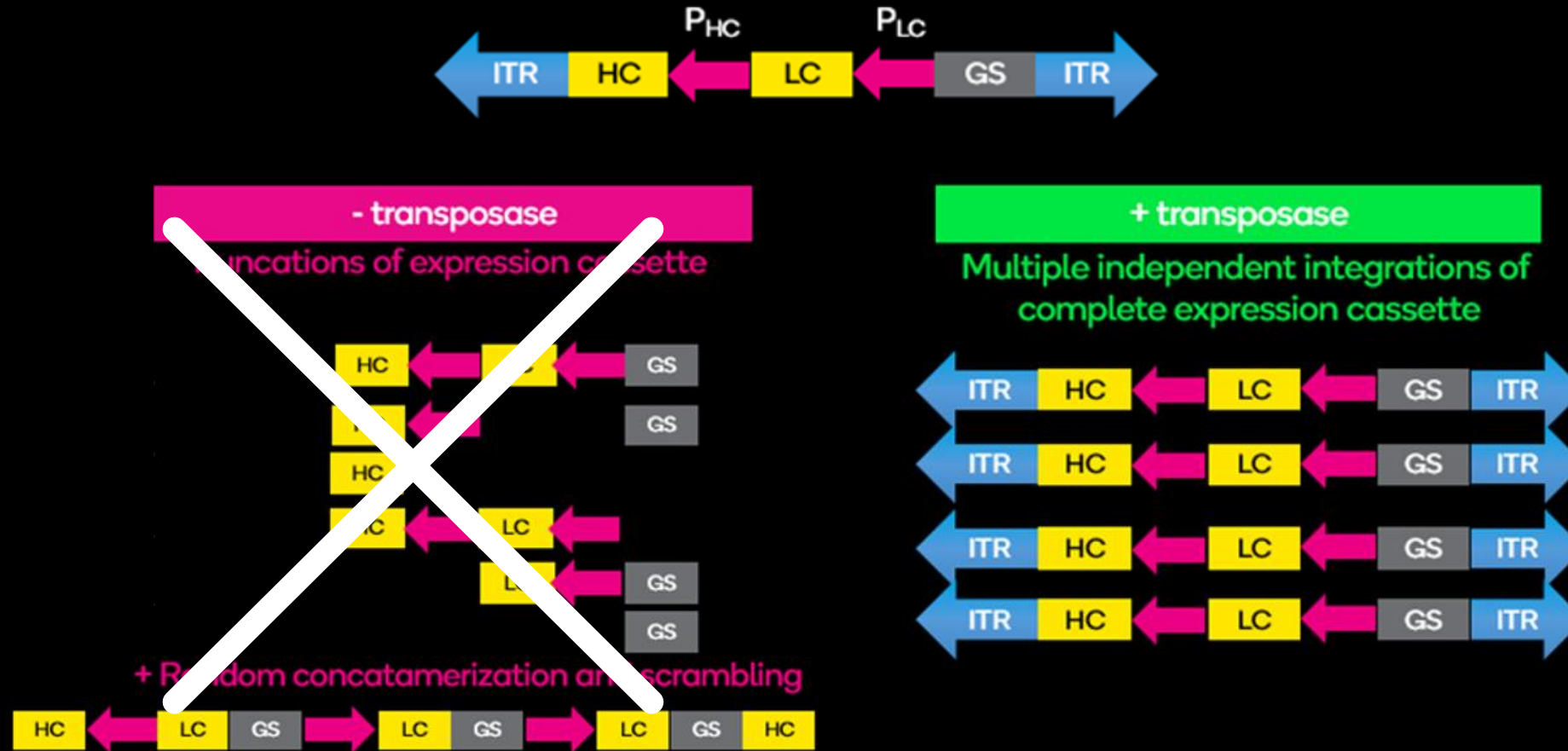
# The life of a transposon-transposase pair



- Transient transposase = Stable insertion
- Single copy integrations at each site
- Multiple insertions (5 – 60+) across the genome
- Structural integrity maintained
- No size limitation

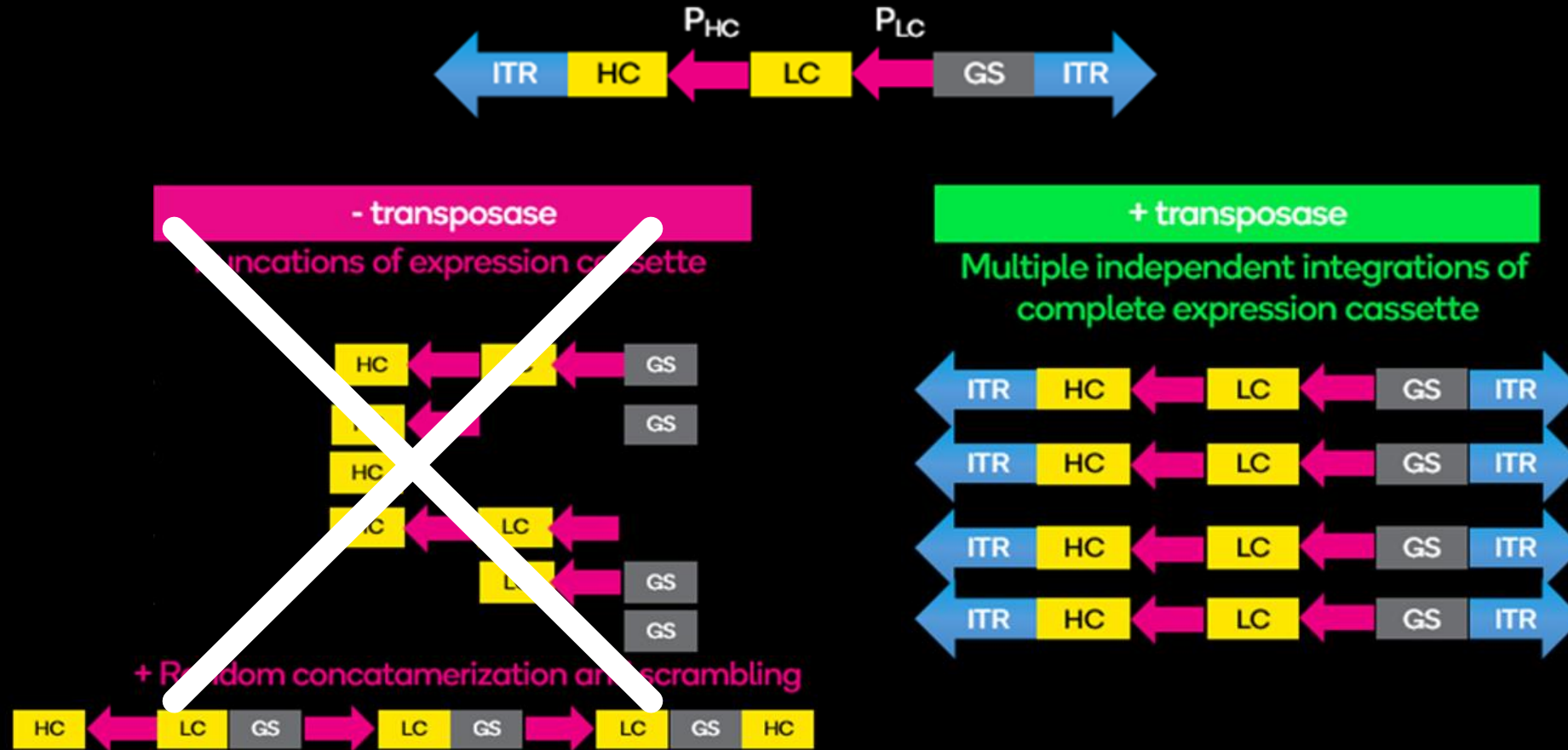


# representation of Leap-In<sup>®</sup> transgenes



- In-silico designed expression construct maintained at every integration site
- On average, functionality of each integration is comparable
  - Expression and product quality

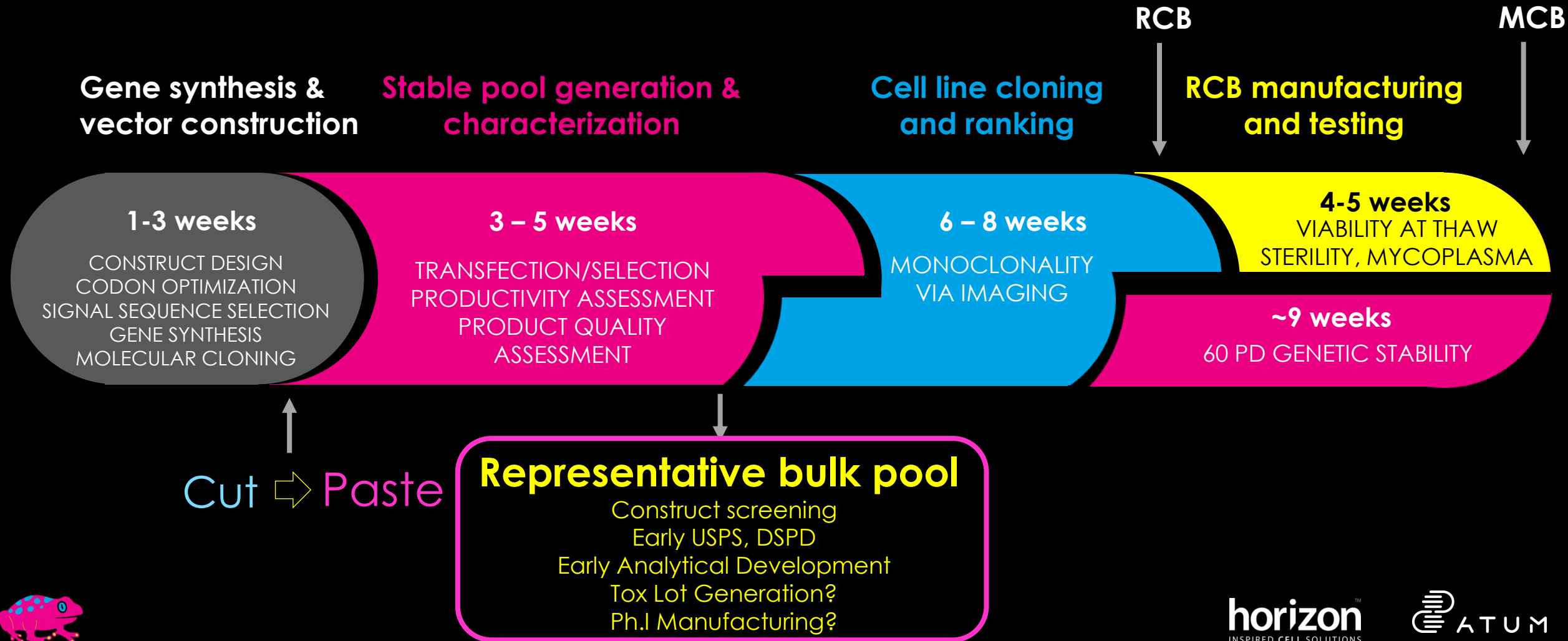
# representation of Leap-In<sup>®</sup> transgenes



Highly uniform and predictive bulk pools



# rapid and robust: sequence to MCB



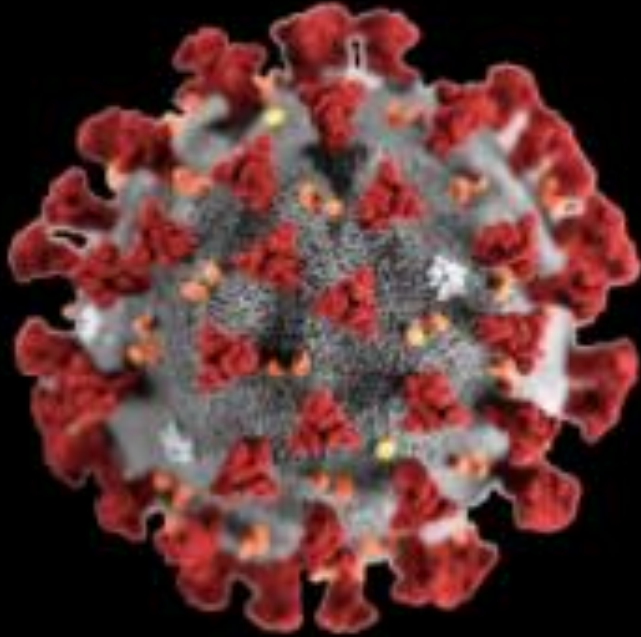
# Leap-In Transposase CLD platform

- Expression construct integrity maintained
- Rapid, robust and representative bulk pool generation
- Robust, high titer and extremely stable clones





# antibodies For COVID 19 treatment



- Eleven candidate therapeutic mAb's
- Desire to initiate human trials ASAP
- Rapid progress: sequence to Ph.I
- Use bulk cell pools for GMP manufacturing

# rapid cell line development: bulk pools



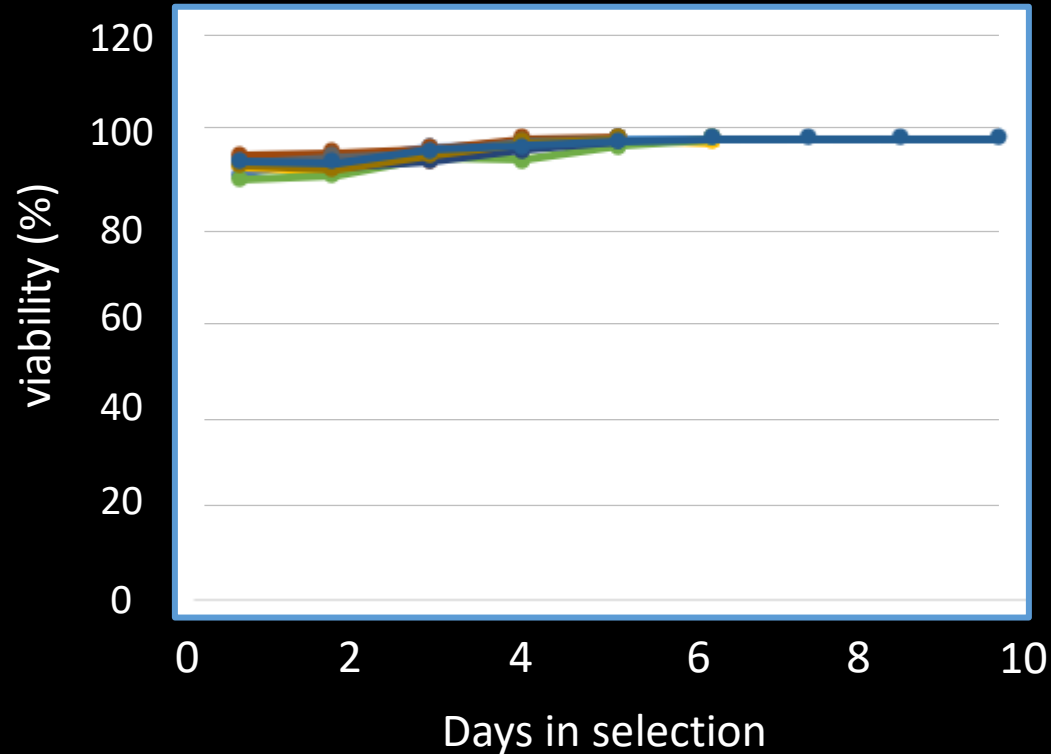
- Two vector sets for each of 11 mAb's
- Create Leap-In Transposase<sup>®</sup> derived pools
- Test expression in platform fed-batch format
- Freeze RCB's for transfer to CDMO



# COVID 19: ATUM accelerated timeline: 1

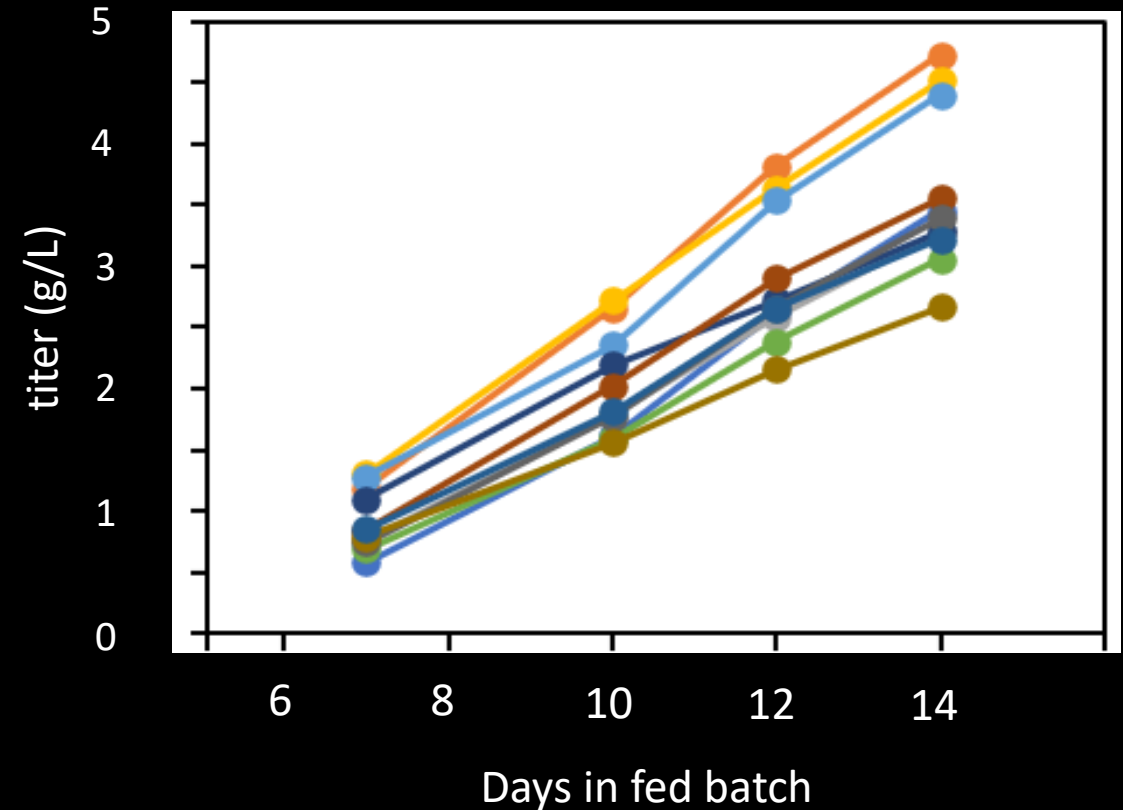


Vector-2



Selection in ~3 days

Vector-2



Titer: 2.5 – 4.8 g/L



# COVID 19: ATUM Accelerated Timeline: 1



Gene  
synthesis

Plasmid  
prep

Pool  
Selection

Pool  
Banking

Bank  
Testing

Fed Batch

Ship to CDMO

*IND filed*

CDMO: Intensified  
fed batch process

>12 g/L

38 days



# COVID 19: ATUM accelerated timeline: 2



Rapid cGMP Manufacturing of COVID-19 monoclonal antibody using stable CHO cell pools

Rita Agostinetto<sup>1</sup>, Jessica Dawson<sup>2</sup>, Angela Lim<sup>2</sup>, Mirva Hejjaoui-simoneau<sup>3</sup>, Cyril Boucher<sup>3</sup>, Bernhard Valldorf<sup>4</sup>, Adin Ross-gillespie<sup>3</sup>, Joseph Jardine<sup>5</sup>, Devin Sok<sup>5</sup>, Dennis Burton<sup>5</sup>, Thomas Hassell<sup>6</sup>, Hervé Broly<sup>7</sup>, Wolf Palinsky<sup>3</sup>, Philippe Dupraz<sup>3</sup>, Mark Feinberg<sup>6</sup>, and Antu Dey<sup>8</sup>

<sup>1</sup>Merck Serono SpA

<sup>2</sup>EMD Serono Biotech Center Inc

<sup>3</sup>Ares Trading SA

<sup>4</sup>Merck KGaA

<sup>5</sup>The Scripps Research Institute

<sup>6</sup>International Aids Vaccine Initiative

<sup>7</sup>Merck Serono SA-Corsier-sur-Vevey

<sup>8</sup>Greenlight Biosciences Inc

Pools 6.0 g/L



200L Preclinical Safety



2000L Phase I

Preprint on Authorea.com

“.. Enabled manufacturing of early clinical trial material within 4.5 months ...”



# COVID 19: ATUM accelerated timeline: 3



**Towards Maximum Acceleration of Monoclonal Antibody Development:  
Leveraging Transposase-Mediated Cell Line Generation to Enable GMP  
Manufacturing within 3 Months using a Stable Pool**

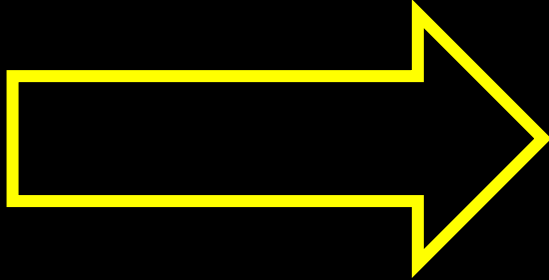
Valerie Schmieder<sup>1</sup>, Juergen Fieder<sup>1</sup>, Raphael Drerup<sup>2</sup>, Erik Arango Gutierrez<sup>2</sup>, Carina Guelch<sup>3</sup>, Jessica Stolzenberger<sup>4</sup>, Mihaela Stumbaum<sup>5</sup>, Volker Steffen Mueller<sup>6</sup>, Fabian Higel<sup>6</sup>, Martin Bergbauer<sup>7</sup>, Kim Bornhoeft<sup>8</sup>, Manuel Wittner<sup>9</sup>, Petra Gronemeyer<sup>10</sup>, Christian Braig<sup>11</sup>, Michaela Huber<sup>12</sup>, Anita Reisenauer-Schaupp<sup>13</sup>, Markus Michael Mueller<sup>14</sup>, Mark Schuette<sup>15</sup>, Sebastian Puengel<sup>1</sup>, Benjamin Lindner<sup>1</sup>, Moritz Schmidt<sup>1</sup>, Patrick Schulz<sup>1</sup> and Simon Fischer<sup>1,\*</sup>

1: Cell Line Development, Bioprocess Development Biologicals, Boehringer Ingelheim Pharma GmbH & Co. KG, Biberach an der Riss, Germany

Journal of Biotechnology, 2022



fundamental mechanism

Cut  Paste

... what you paste matters ...









# considerations for chain ratio balancing

## Sequence

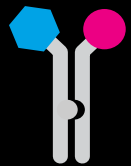
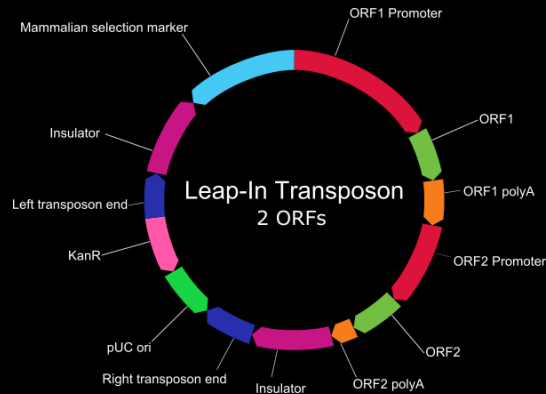
- Codon choice
- mRNA 2° structure
- Poly-A signal
- 5'/3' UTR choice
- mRNA stability
- Ribosomal entry/processivity
- Splice site donor/acceptor
- Signal sequences
- Etc.

## Vector

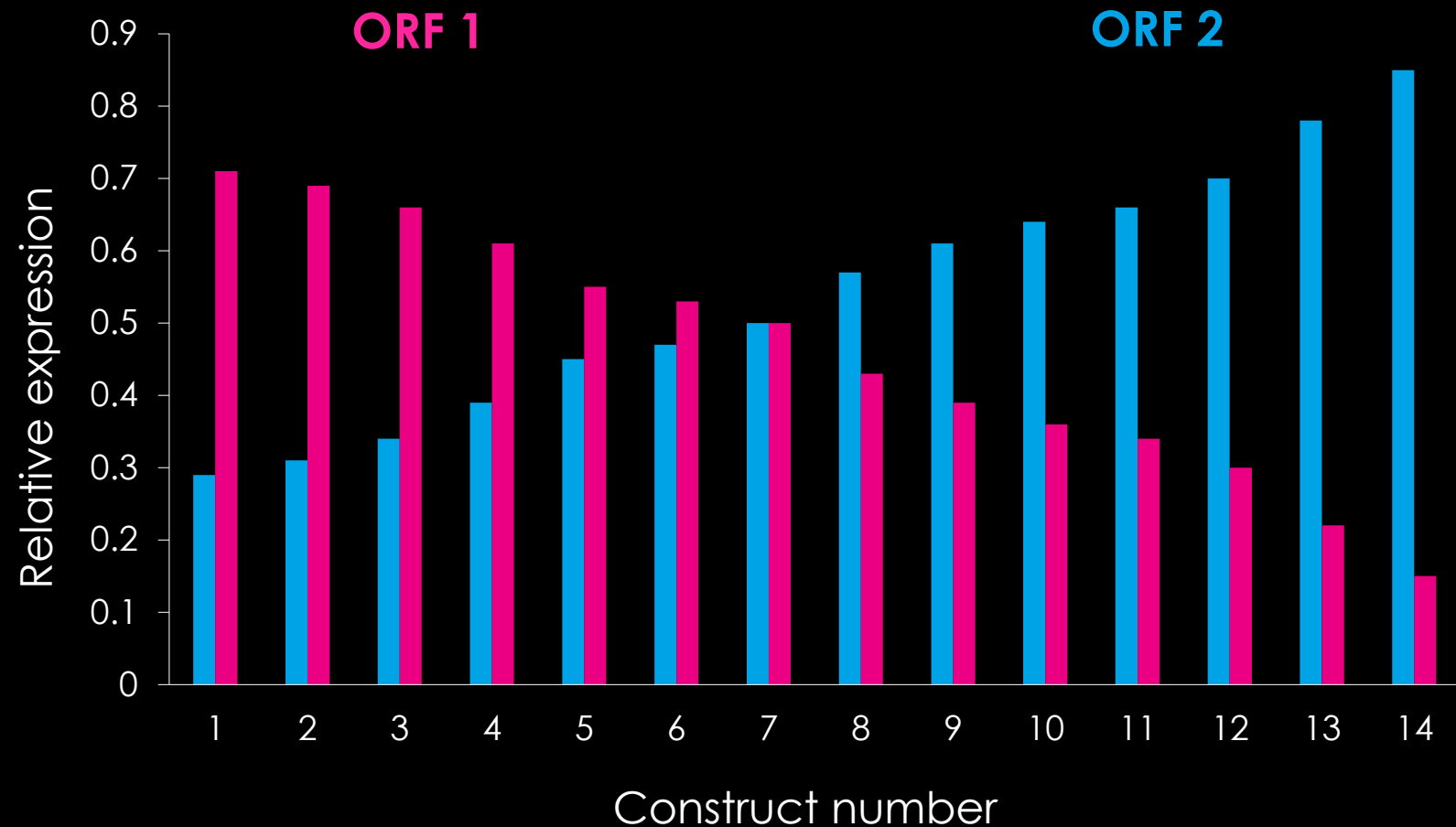
- Promoter choice
- Order of expression cassettes
- Number of expression cassettes
- Spacing of expression cassettes
- Directionality of expression cassettes
- Size of vectors
- Single vector or multiple vectors
- Choice of insulators
- Etc.



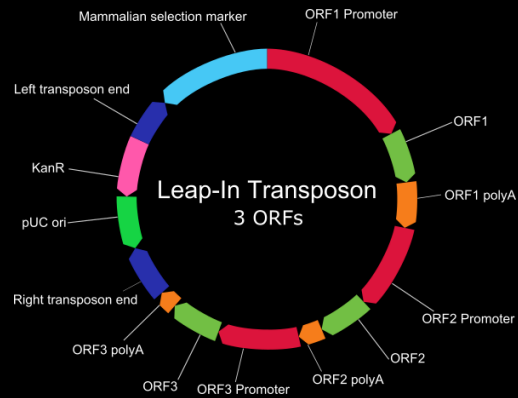
# controlling ratios with construct design: 2 ORFs



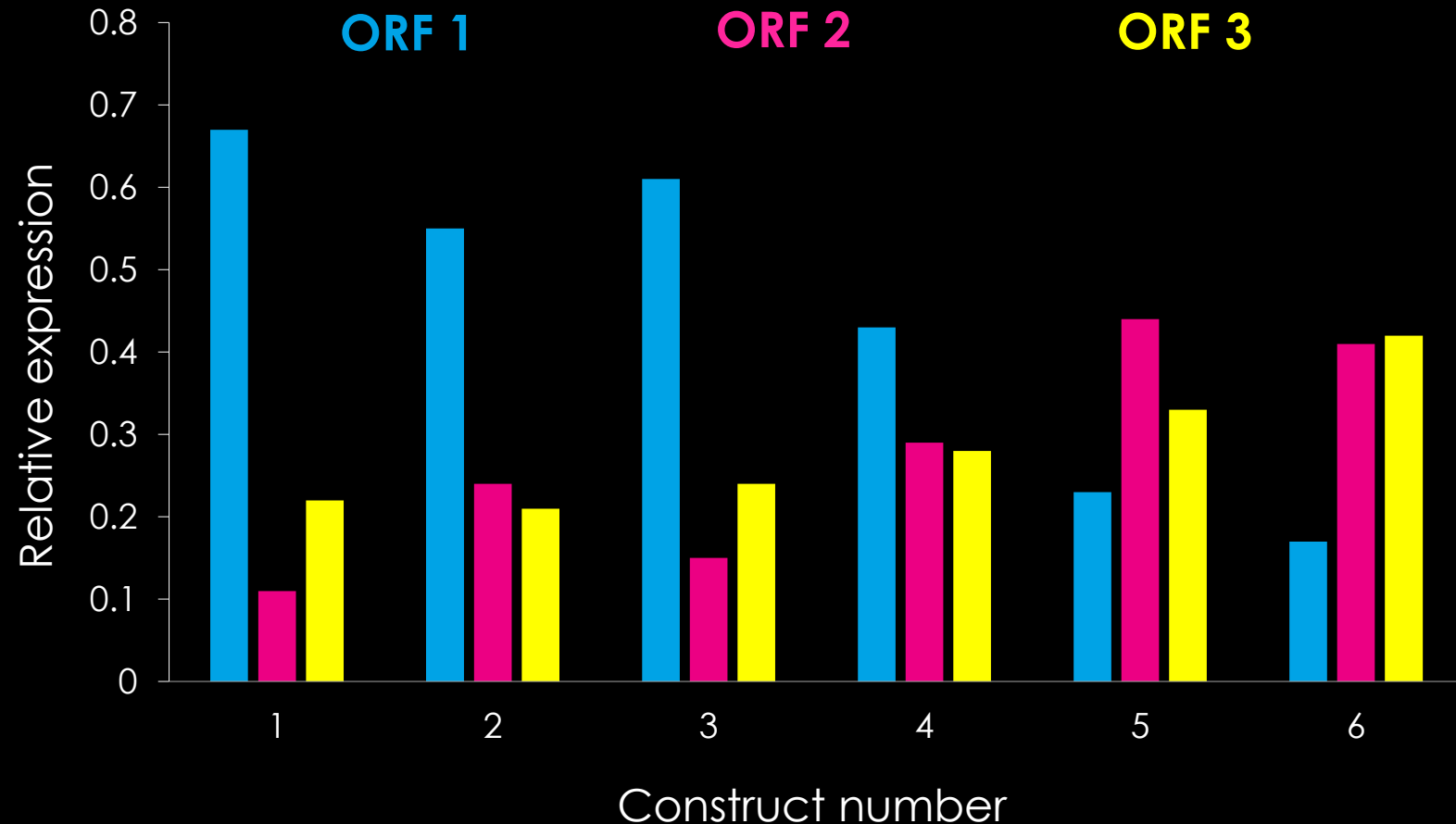
**Bispecific Antibodies**  
chain ratio modulation



# controlling ratios with construct design: 3 ORFs



**Bispecific Antibodies**  
chain ratio modulation



# case study

- Medium sized biotech
- Leap In Transposase licensee
- Two and four chain bispecifics
- Strong internal biologics capabilities

*( ... heavily redacted data set ... )*

# case study-1: two chain bispecific

## Random Integration Pool

Fed-batch Expression d14: Normalized = 1

61.4% Proper Assembly

## Leap In Transposase Pool

Fed-batch Expression d16: Normalized =1.9

90% Proper Assembly

## Leap In Transposase Clones

| Clone | Titer<br>(Normalized) | % Assembly |
|-------|-----------------------|------------|
| A     | 2.5                   | 98.4       |
| B     | 2.5                   | 98.3       |
| C     | 2.6                   | 98.2       |
| D     | 2.5                   | 98.0       |
| E     | 2.4                   | 94.9       |

Good pools predict good clones

# case study-2: four chain bispecific

## Random Integration Pool

Fed-batch Expression d14: Normalized = 1

10.6% Proper Assembly

## Random Integration Clones

| Clone | Titer<br>(Normalized) | % Assembly |
|-------|-----------------------|------------|
| A     | 0.82                  | 87.5       |
| B     | 0.71                  | 86.0       |
| C     | 0.44                  | 72.1       |
| D     | 0.55                  | 77.1       |
| E     | 0.52                  | 60.5       |

Random integration clones lost expression and  
few clones with reasonable product quality

# case study-2: four chain bispecific

## Random Integration Pool

Fed-batch Expression d14: Normalized = 1  
10.6% Proper Assembly

## Leap In Transposase Pool

Fed-batch Expression d16: Normalized = 1.1  
85.6% Proper Assembly

## Leap In Transposase Clones

| Clone | Titer<br>(Normalized) | % Assembly |
|-------|-----------------------|------------|
| A     | 1.0                   | 92.9       |
| B     | 1.1                   | 93.5       |
| C     | 0.96                  | 91.8       |
| D     | 0.95                  | 92.8       |
| E     | 1.2                   | 88.8       |



# case study-2: four chain bispecific

## Random Integration Pool

Fed-batch Expression d14: Normalized = 1  
10.6% Proper Assembly

## Leap In Transposase Pool

Fed-batch Expression d16: Normalized = 1.1  
85.6% Proper Assembly

## Leap In Transposase Clones

| Clone | Titer<br>(Normalized) | % Assembly |
|-------|-----------------------|------------|
| A     | 1.0                   | 92.9       |
| B     | 1.1                   | 93.5       |
| C     | 0.96                  | 91.8       |
| D     | 0.95                  | 92.8       |
| E     | 1.2                   | 88.8       |

Leap In derived clones retained expression and  
all top clones had excellent product quality



# licensee comments

## The Leap-In transposase platform enables:

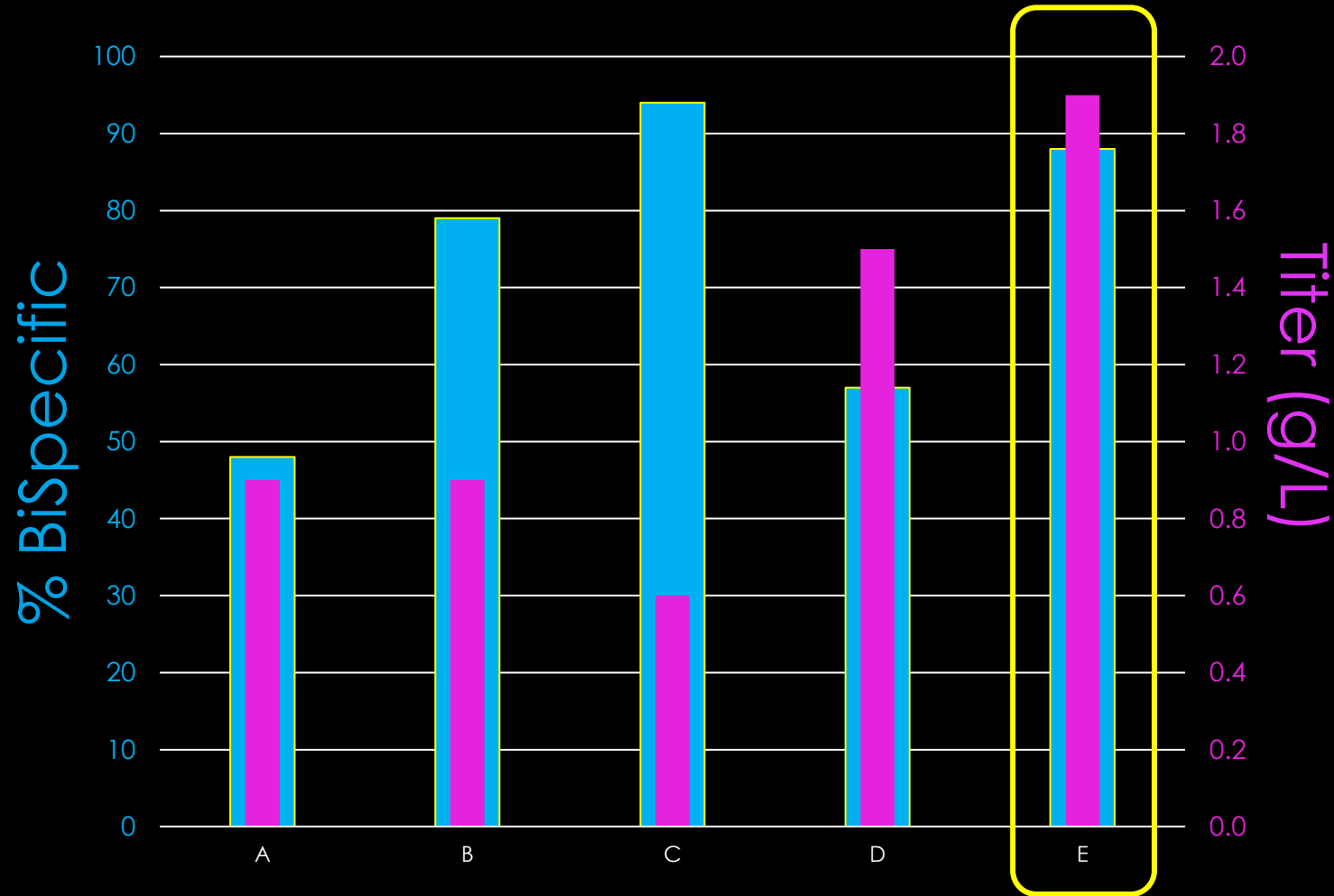
- Higher titers and better protein quality in stable pools.
- More homogenous cell populations in stable pools.
- Low clonal variability = fewer clones need to be screened.
- Timelines can be shortened by ~8 weeks.

# case study-3: 3-chain bispecific mAb

- Known to be difficult
  - Low titer
  - Poor assembly
- 14 vector configurations
  - Varying expression ratios
  - Varying expression levels
- Leap In Transposase based pool selection
- Analytical assessment
  - Total titer
  - Chain expression: Relative and Amount
  - % Bispecific



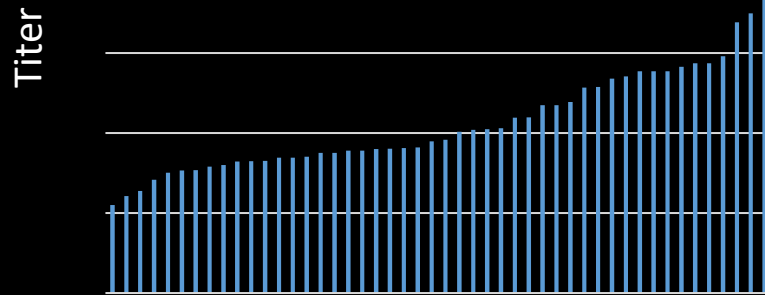
# case study: 3-chain bispecific mAb - bulk pools



# case study: 3-chain bispecific mAb - clones

## Pool E

Clone performance @ 24 wp format

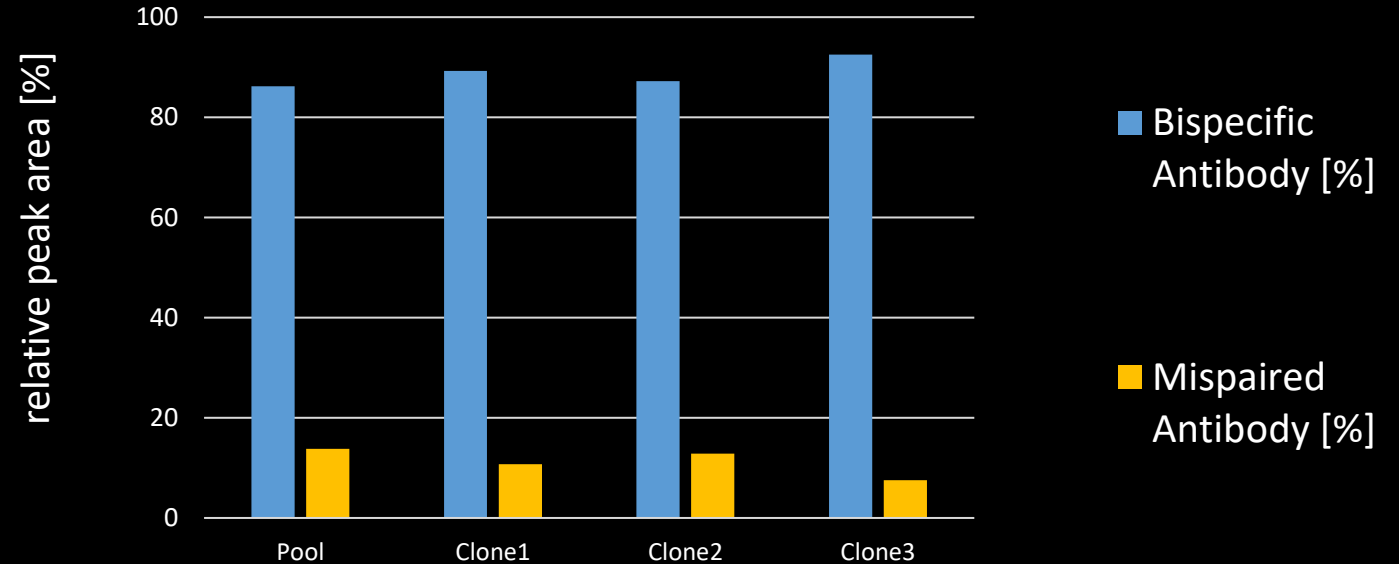


Pool and clone productivity

| Pool*     | derived clones* |
|-----------|-----------------|
| 1.9 [g/L] | up to 5.5 [g/L] |

\*Day 12 standard fed-batch

% bispecific antibody

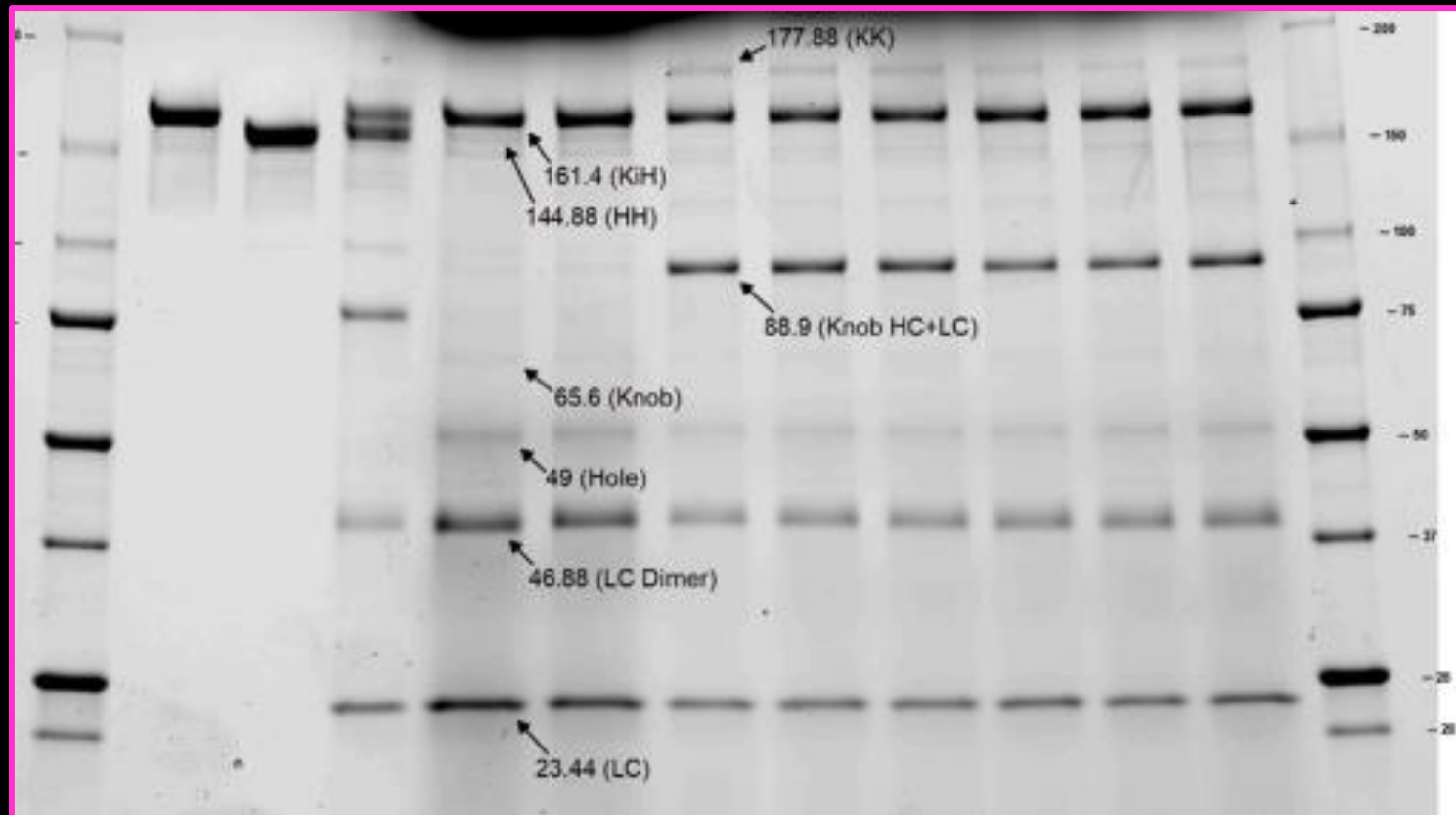


Good pools predict good clones



... and there is more ...

supernatant SDS PAGE



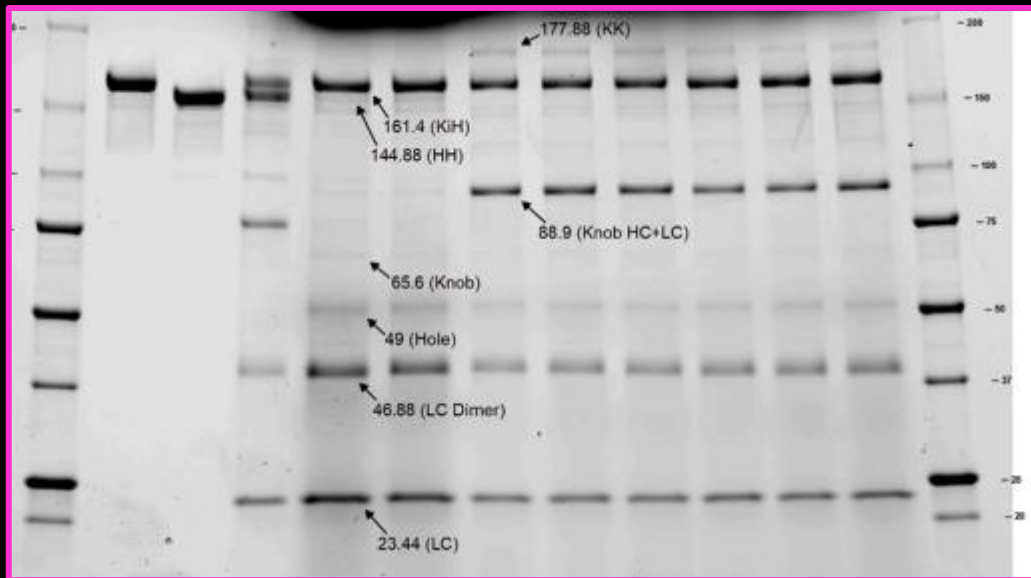
# ... and there is more ...

An innovative platform to improve asymmetric bispecific antibody assembly, purity, and expression level in stable pool and cell line development☆☆☆

Yanling Wang<sup>a,\*1</sup>, Haoran Qiu<sup>a,1</sup>, Jeremy Minshull<sup>b</sup>, Wilburt Tam<sup>a</sup>, Xichan Hu<sup>a</sup>, Carl Mieczkowski<sup>a</sup>, Weibin Zheng<sup>a</sup>, Chun Chu<sup>a</sup>, Wenqiang Liu<sup>a</sup>, Ferenc Boldog<sup>b</sup>, Claes Gustafsson<sup>b</sup>, Jean-Michel Gries<sup>a</sup>, Wenfeng Xu<sup>a</sup>

<sup>a</sup> Hengenix Biotech Inc., 430 N McCarthy Blvd, Milpitas, CA, USA

<sup>b</sup> ATUM, 37950 Central Court, Newark, CA, USA



“ ... efficiently generate clones with titer above 6 g/L within 3 months from vector to top 10 clones ... ”



... and more ...

Leap In Licensee with a 3-chain platform:

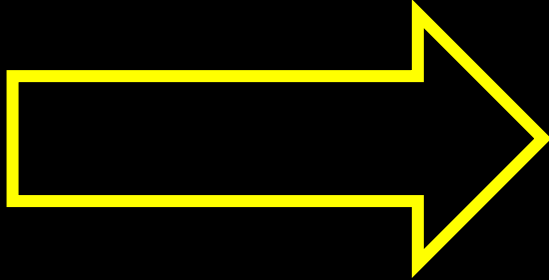
“ ... 5-7 g/L with >97% correct assembly ... ”

CLD customer with a novel 4-chain platform:

“ ... 6-11 g/L with >85% correct assembly \* ... ”



fundamental mechanism

Cut  Paste

... what you paste matters ...





# robust market adoption

- Launched ~4.5 years ago
- Offered as a service by ATUM: >175 projects delivered
- >40 active licensees: 5 of top 7 pharma
- >20 regulatory filings in ~3 years: Including Ph.II/III
- Tech transfer to ≥ 13 CDMO's

IT NETWORKS



# ATUM

- Gene synthesis, vectors
  - Large, complex, routine
  - Host optimized
- Protein production
  - 96-well to 100's of grams
  - mAbs to others
  - Mammalian, e. coli, other
- Protein analytics
  - MS, HPLC, CE, other
  - Developability
- Cell based assays
  - FACS, signaling, other
  - Primary immune cells
- **Protein Engineering**





# Thank You

Oren Beske  
obeske@atum.bio



## Partners:

Horizon Discovery  
Rentschler Biopharma  
Our Customers

Technology presented is protected by issued US patents 10435696, 10344285, 10287590, 10253321, 10233454, 10041077, 9771402, 9580697, 9574209, 9534234, 9493521, 9428767, 9290552, 9206433, 9102944, 8975042, 8825411, 8635029, 8412461, 8401798, 8323930, 8158391, 8126653, 8005620, 7805252, 7561973, 7561972 and pending applications

