

Cell Line Development for Multi-chain Biologics: *Leveraging the Leap In Transposase Platform*

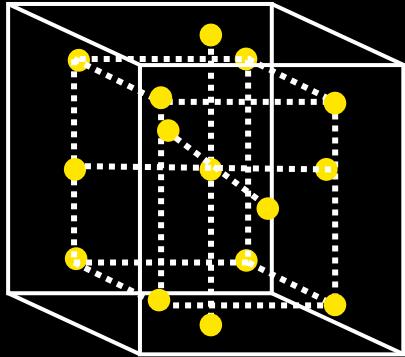
THE POWER OF THE POOL

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

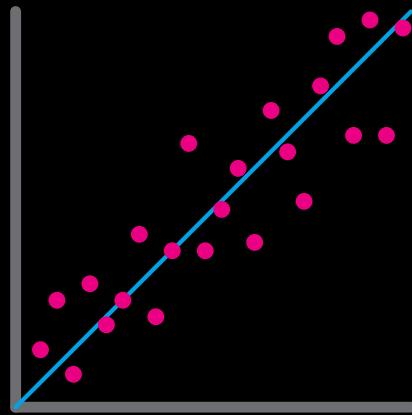


The 13th Annual
**BIOPROCESSING
SUMMIT** IN-PERSON • VIRTUAL
FLEXIBLE REGISTRATION

Design of Experiment



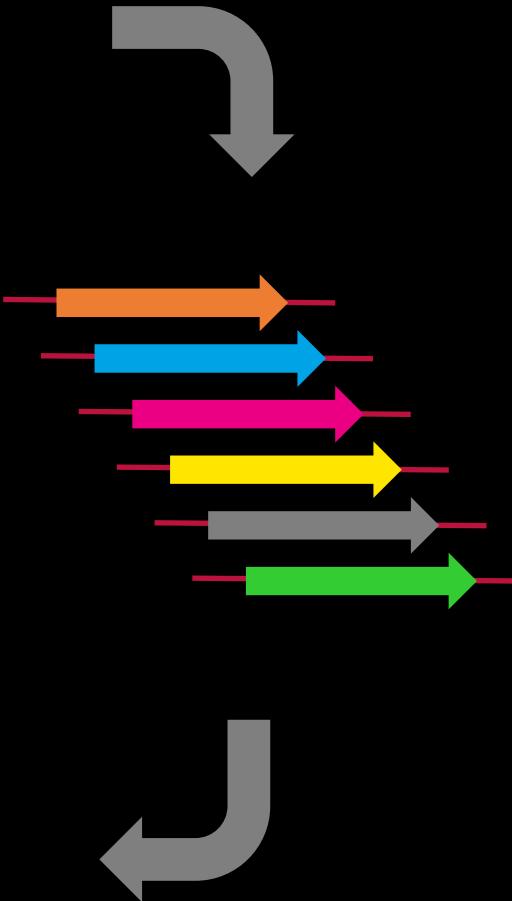
Machine
Learning



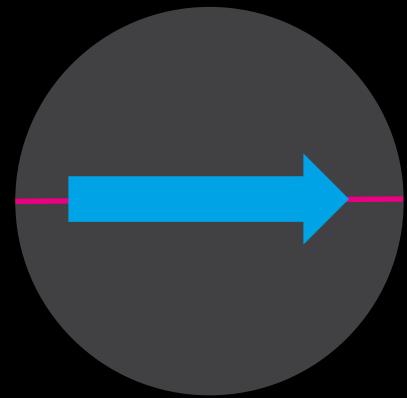
GPS
platform



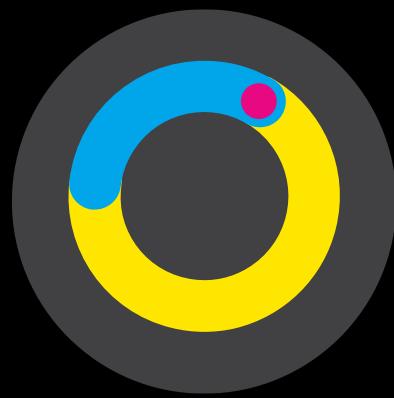
Test



The GPS Platform



gene GPS



vector GPS



protein GPS

ORF codon optimization

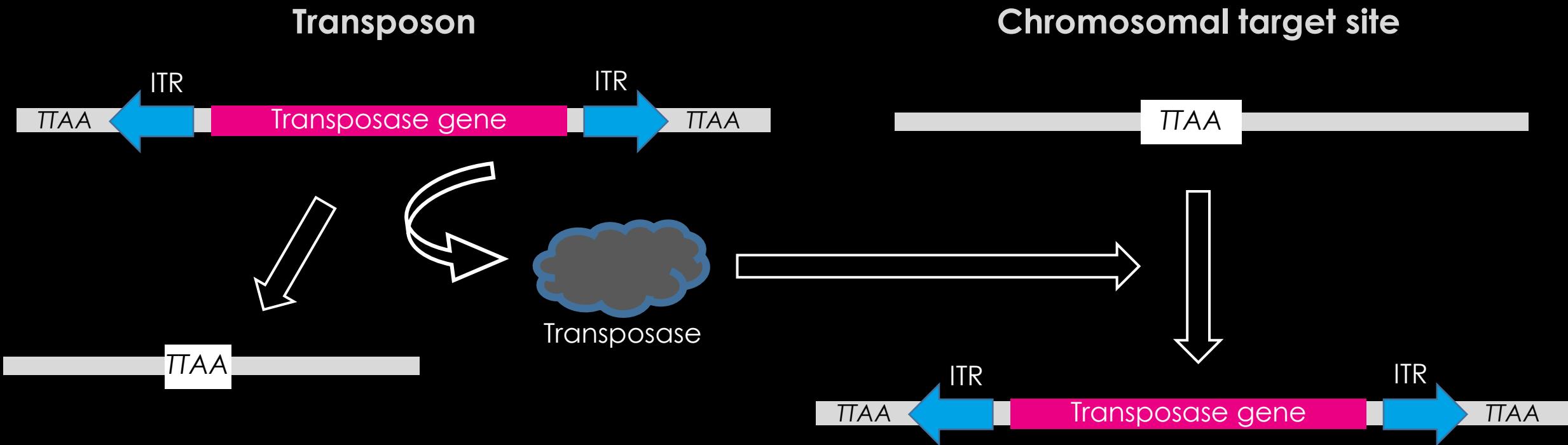
Expression vector element
optimization

Protein attribute
optimization

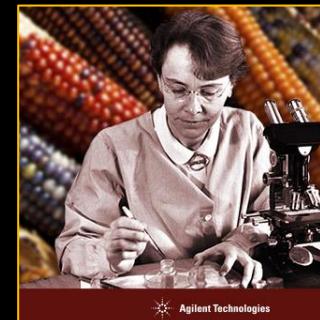


Leap-In Transposase® Platform

The life of a transposon-transposase pair

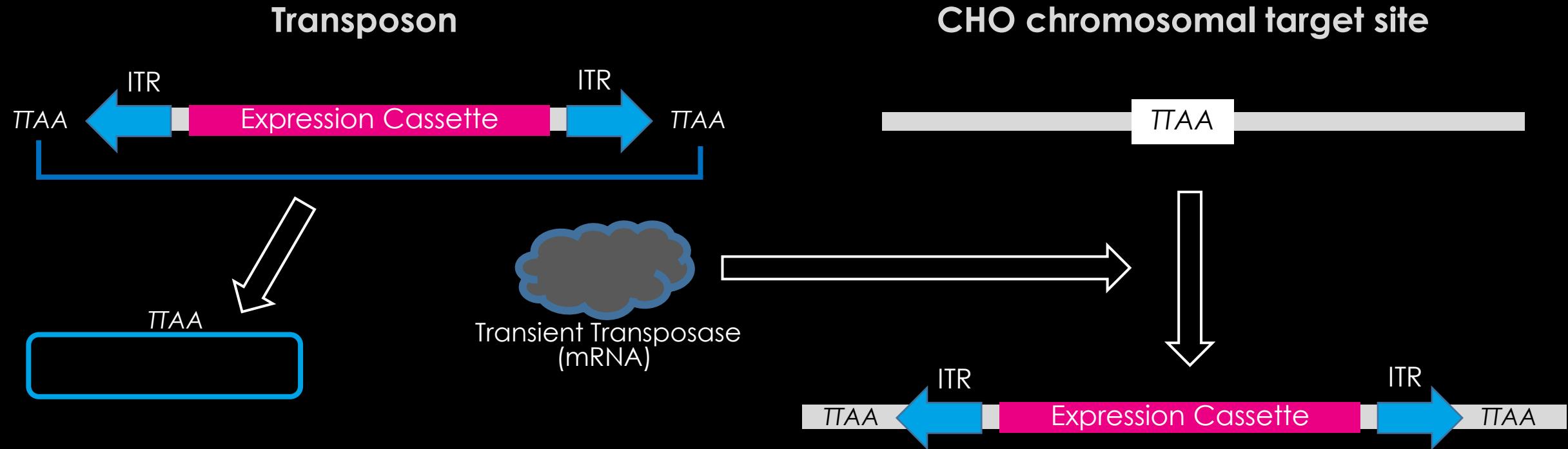


- Billions of years of evolutionary history
- Cut-paste mechanism
- Single copy integration at each site
- Perfect integration of elements between ITR's



1983 Nobel Prize in Physiology or Medicine

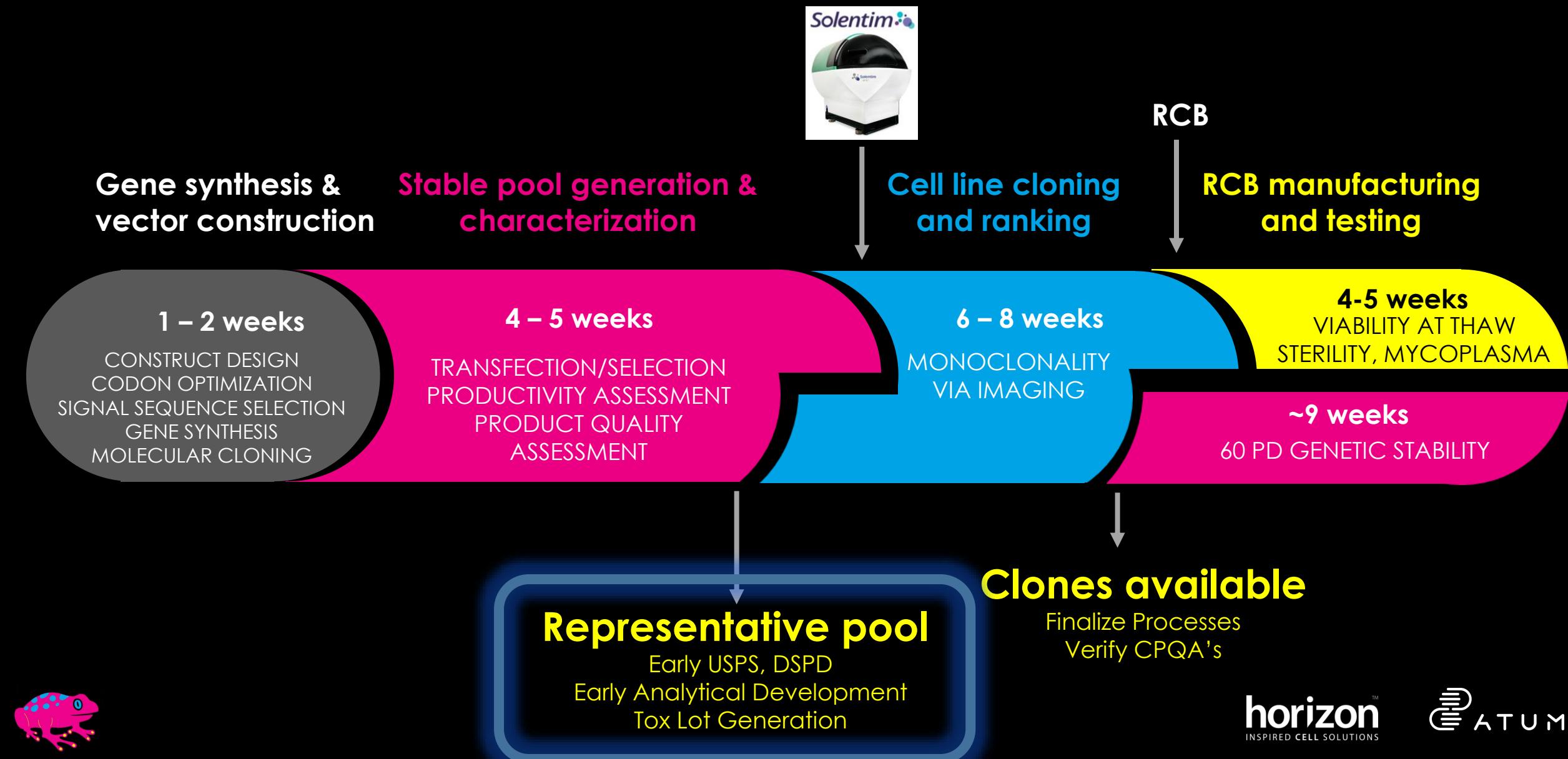
The life of a transposon-transposase pair



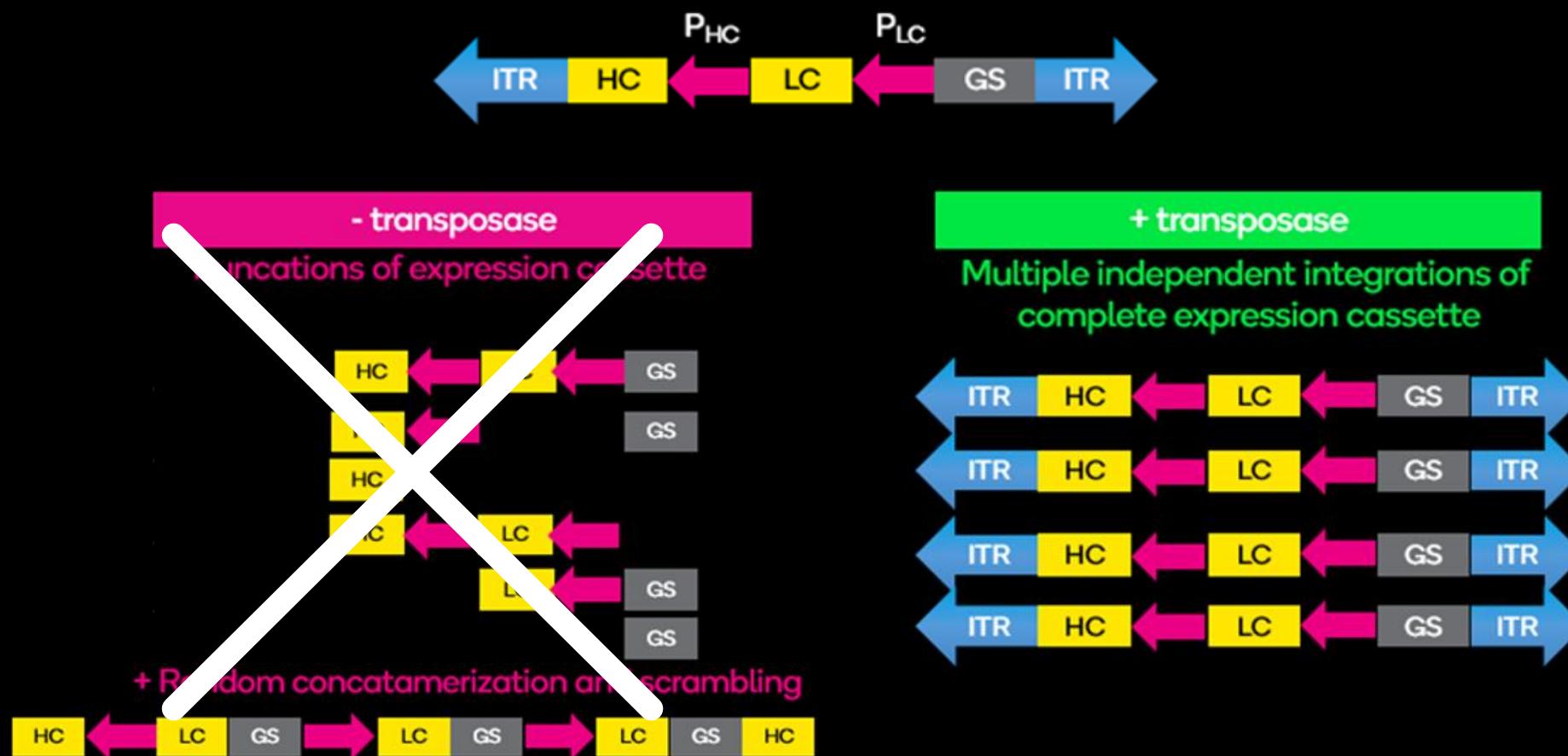
- Transient transposase = Stable insertion
- Single copy integrations at each site
- Multiple insertions (5 – 60+) across the genome
- Structural integrity maintained
- No size limitation



Transfection to RCB in ~12 weeks



Consistent, uniform presentation of Leap-In® transgenes



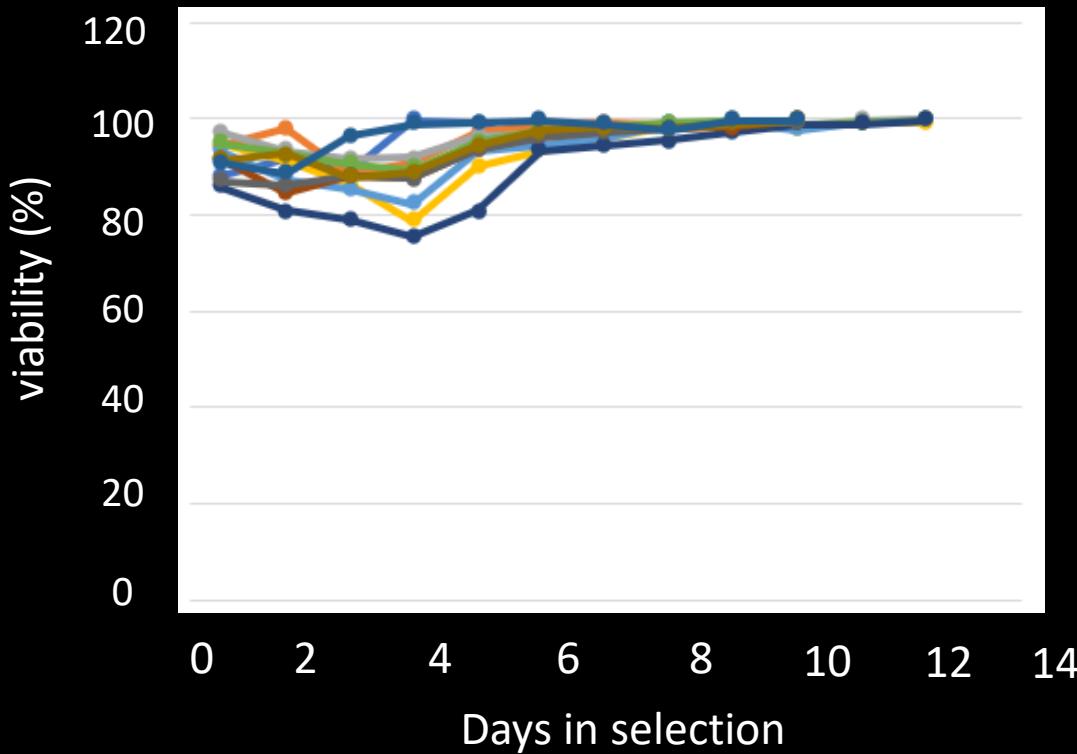
- In-silico designed expression construct maintained at every integration site
- On average, functionality of each integration is comparable
 - Expression and product quality



Rapid Stable CHO Pool Selection

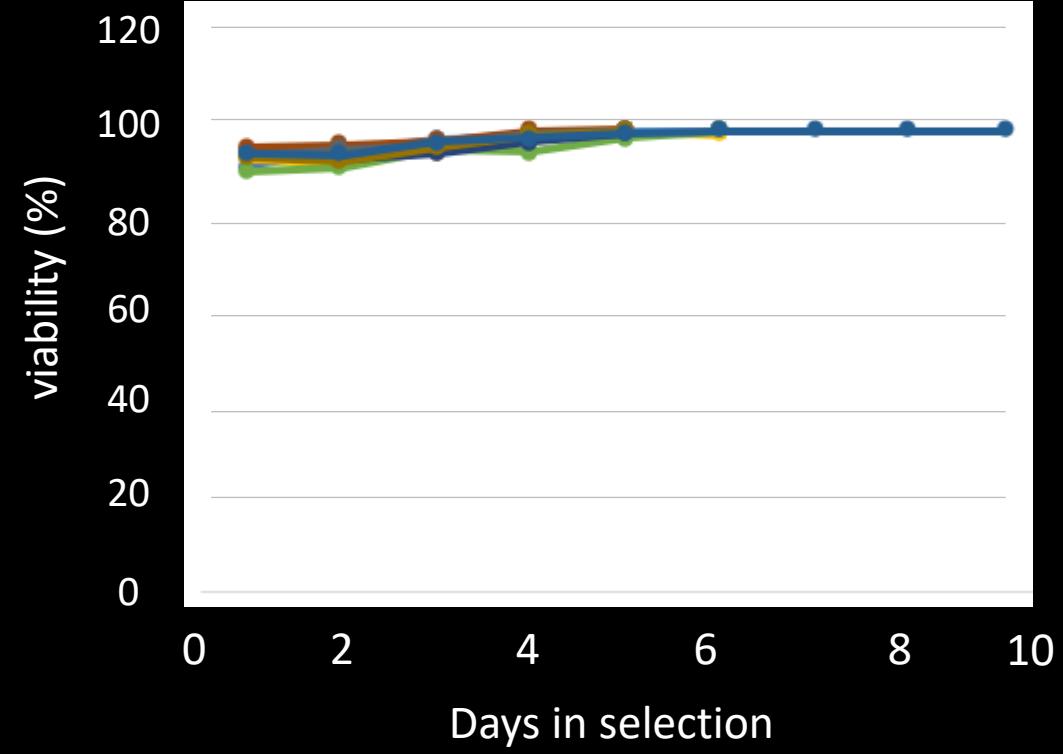


Vector-1



Selection in ~7-8 days

Vector-2



Selection in ~3-4 days

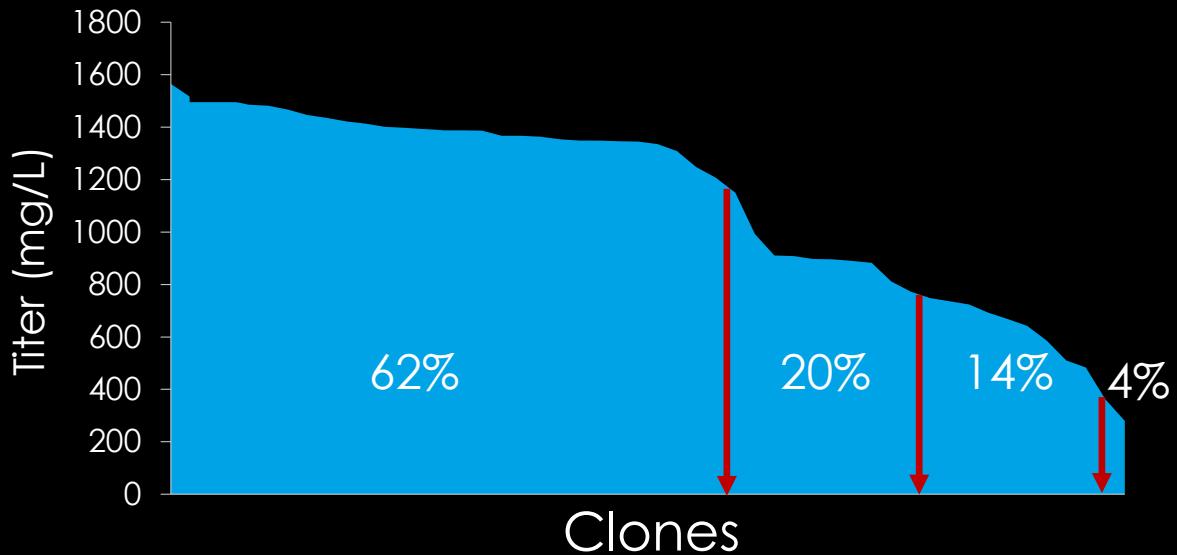


horizonTM
INSPIRED CELL SOLUTIONS

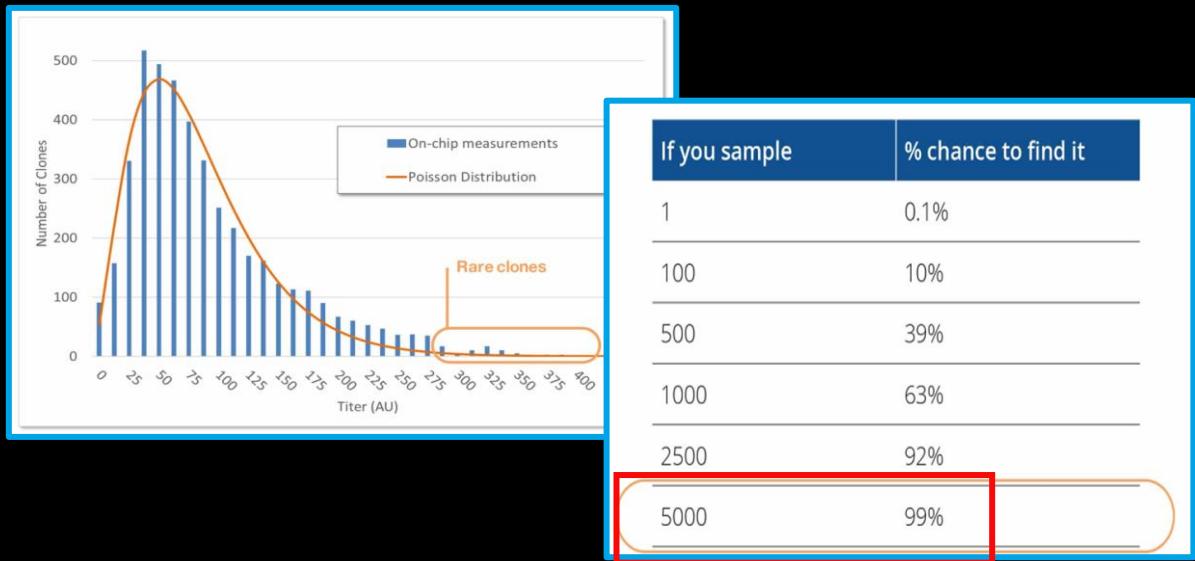
ATUM

CHO pools uniformity: Titer

Leap-In Transposase®



Random Integration



<https://www.berkeleylights.com/>

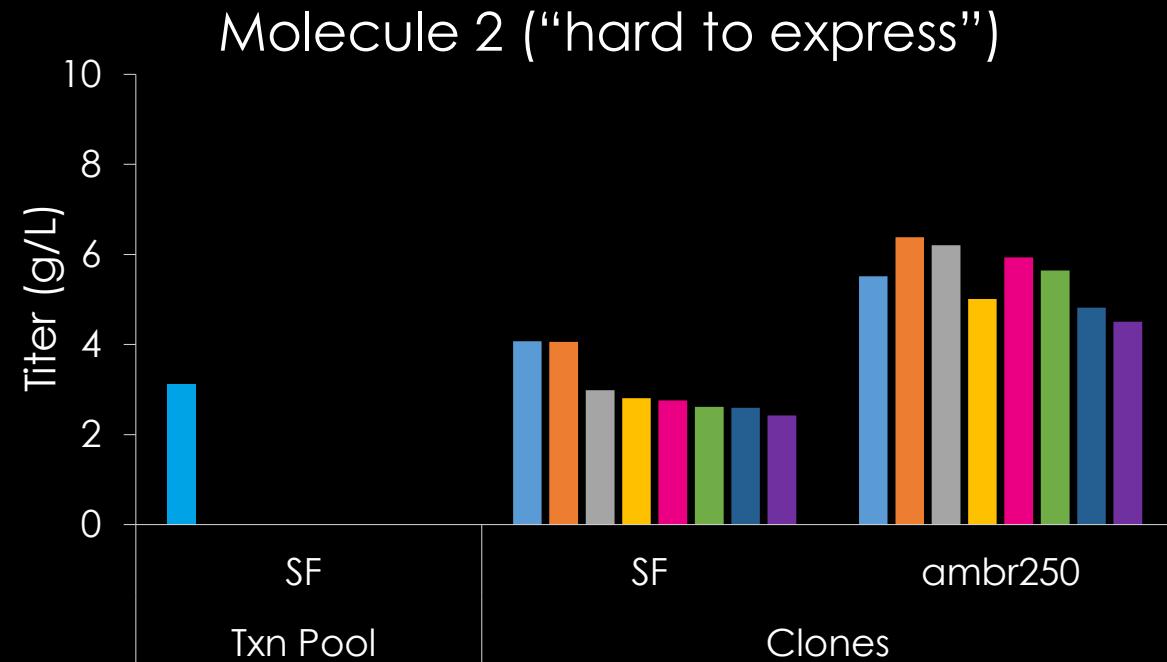
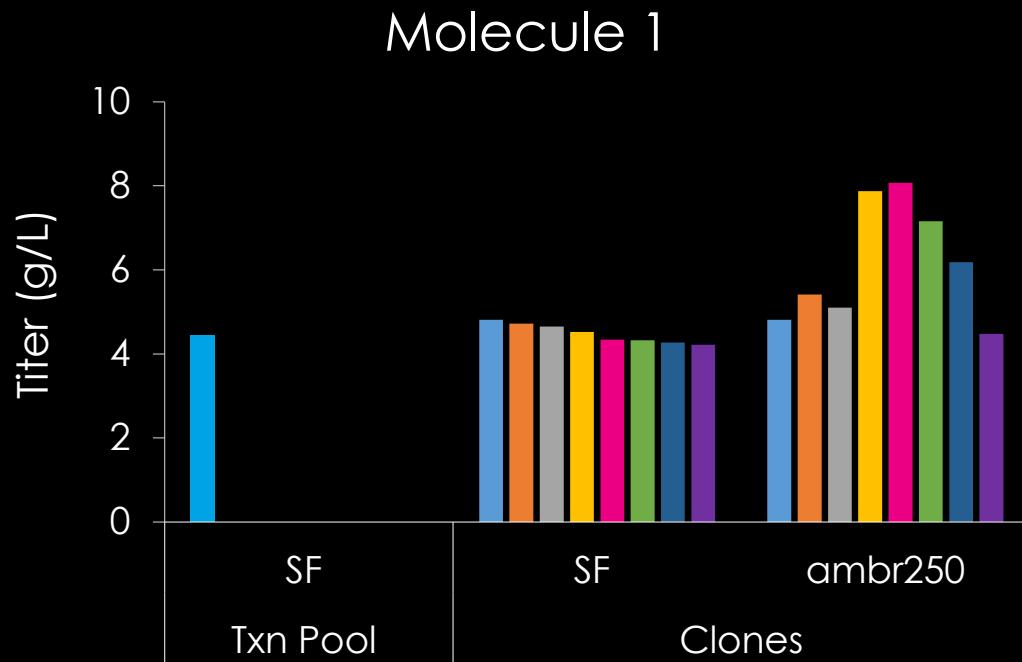
- 62% of clones in top quartile of expressers
- 82% of clones in top half of expressers
- **99% probability in under 200 clones**
- High producers rare

Robust High Titer Stable CHO Pools

Protein	Volumetric productivity
IgG1	7.7 g/L
IgG1	5.9 g/L
IgG1	5.3 g/L
IgG1	5.5 g/L
IgG1	5.3 g/L
IgG4	5.0 g/L
IgG4	5.0 g/L
Fc Fusion	3.5 g/L
3 ORF Bispecific	~7 g/L
3 ORF Bispecific	3 g/L

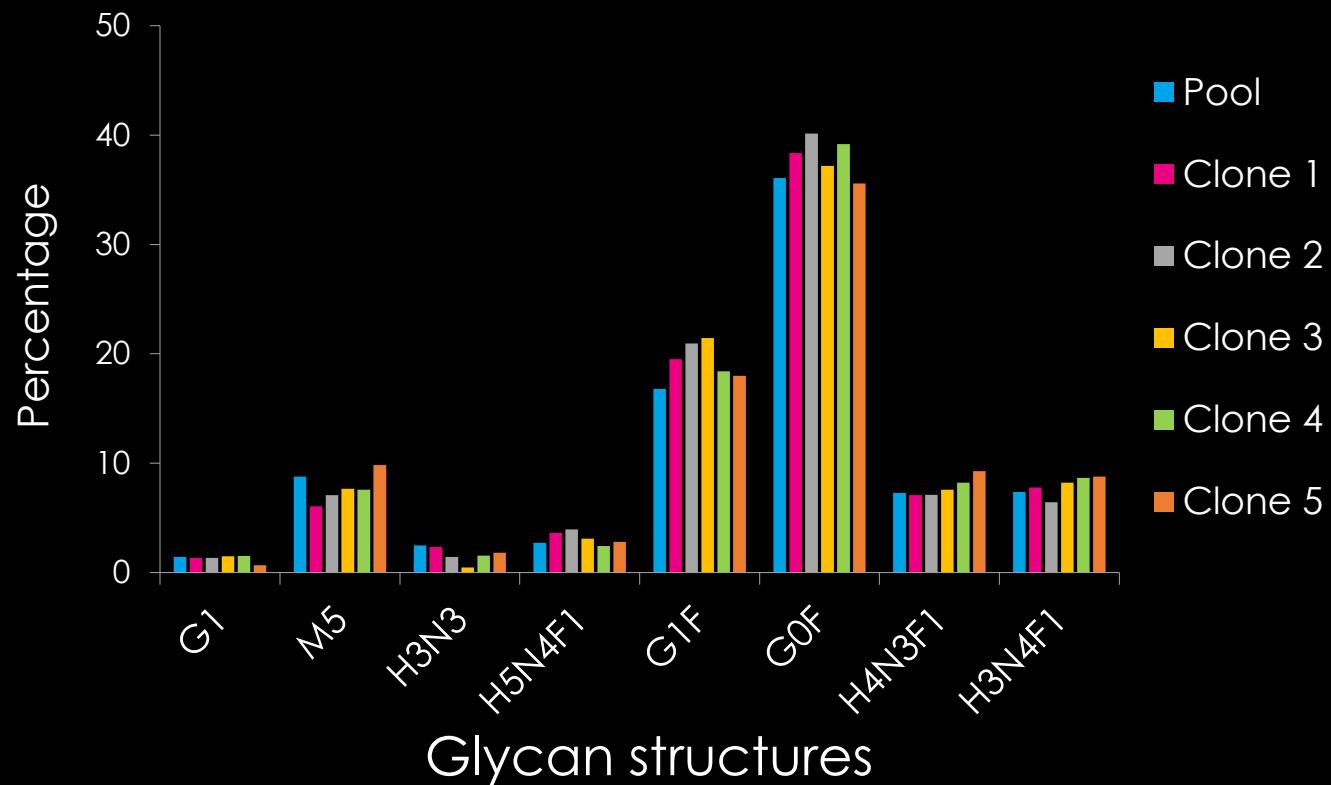


Pools predict clones: Titer

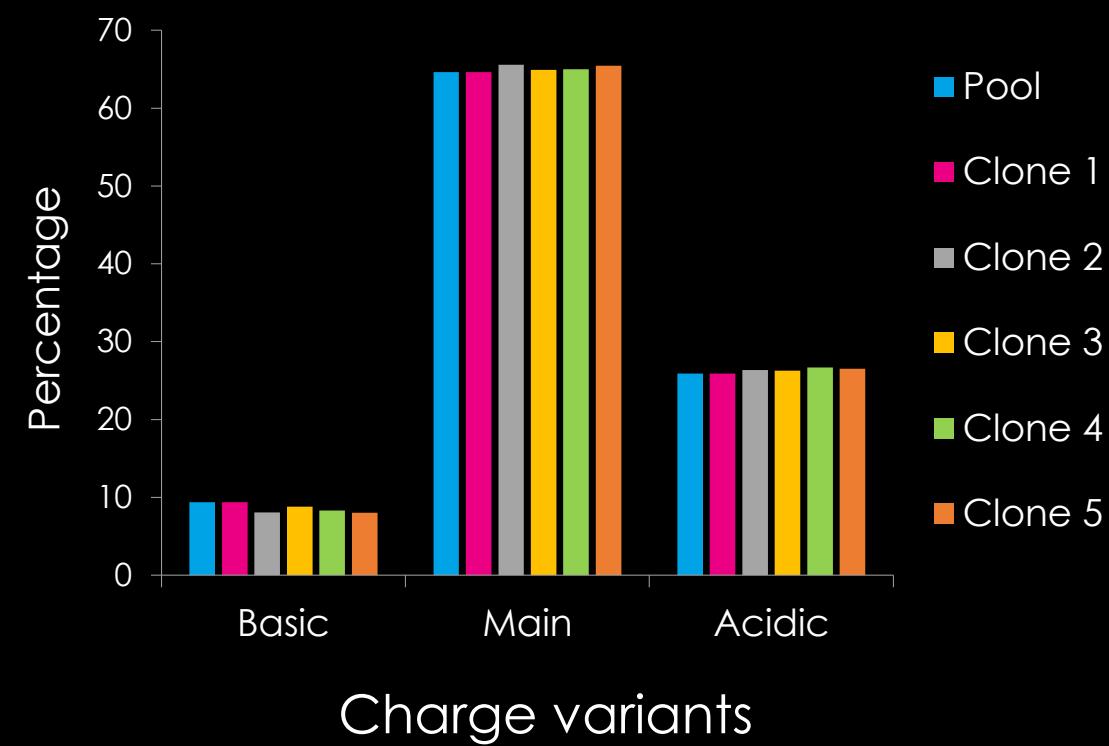


Pools predict clones: Product quality

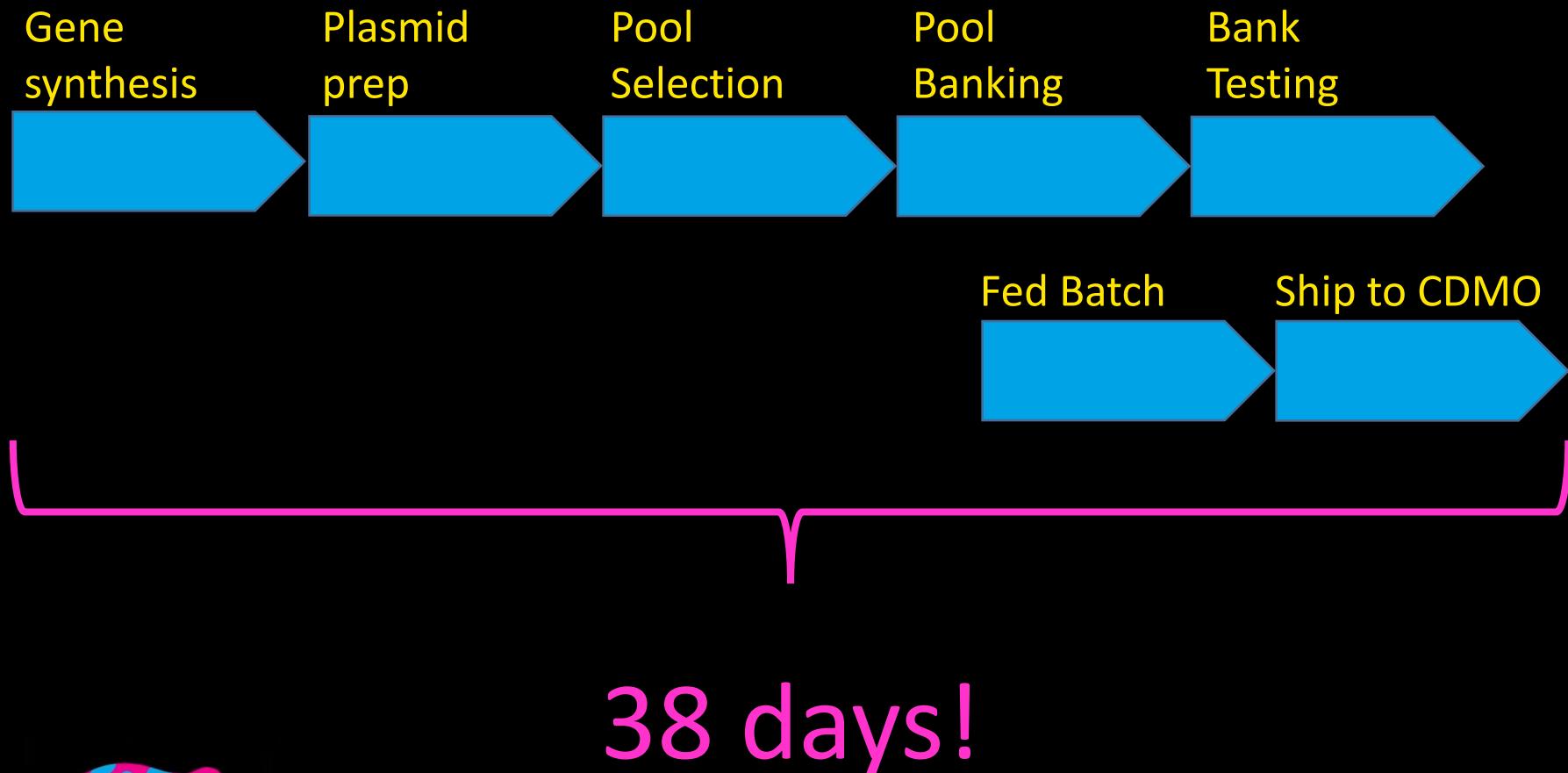
Glycans: Pool vs Clones



Charge variants: Pool vs Clones



COVID 19: ATUM Accelerated Timeline

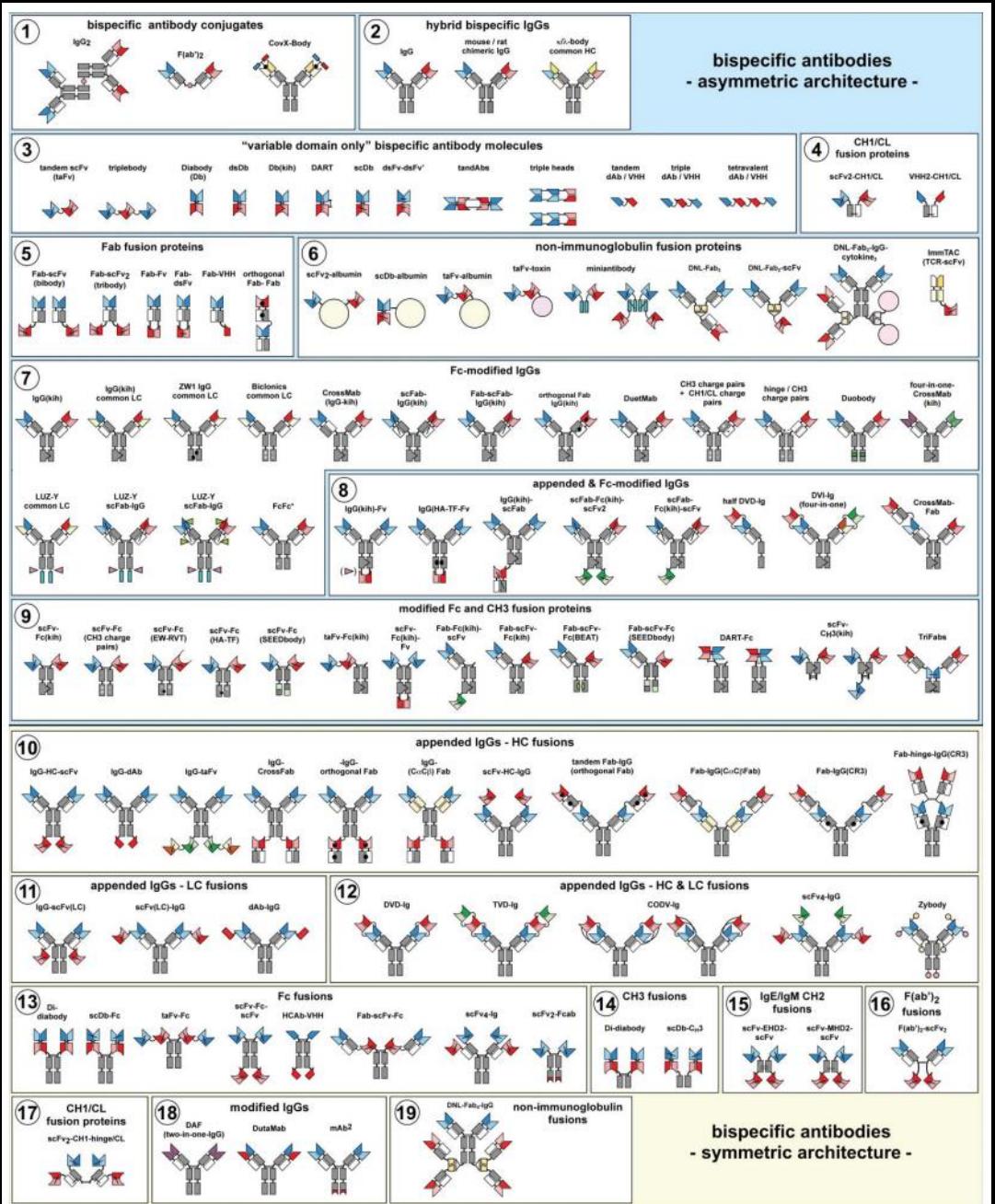


Leap-In transgenes structural feature summary

- Expression construct integrity maintained
- Each integration is equally functional
- Rapid, robust and predictive pool generation

THE POWER OF THE POOL





Beyond mAb's: 3 Chains and More

The “zoo” of bispecifics

Considerations for chain ratio balancing

Sequence

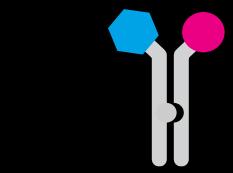
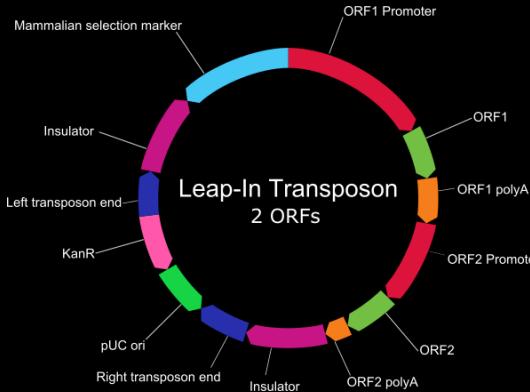
- Codon choice
- mRNA 2° structure
- Poly-A signal
- 5'/3' UTR choice
- mRNA stability
- Ribosomal entry/processivity
- Splice site donor/acceptor
- Signal sequences
- Etc.

Vector

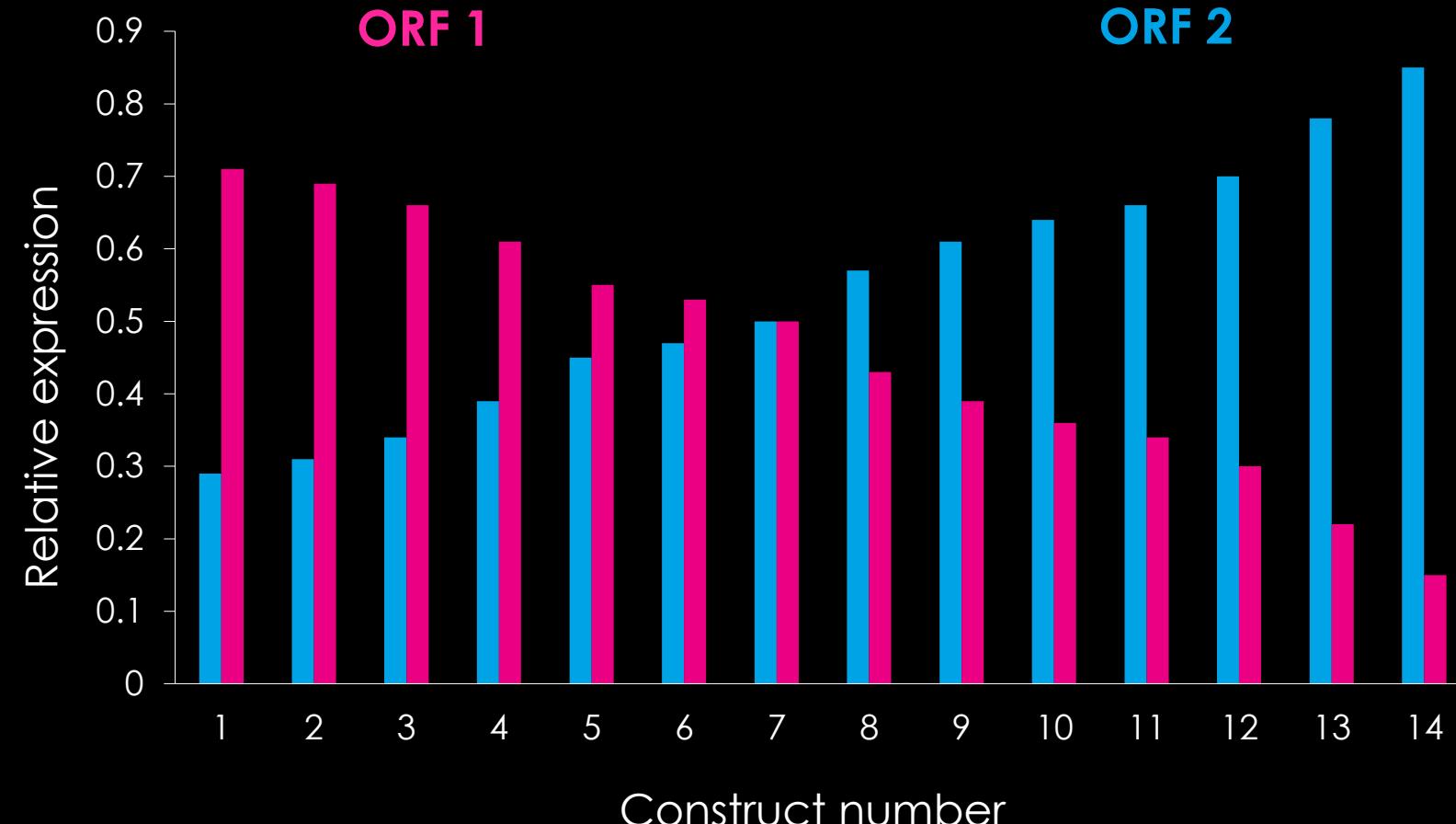
- Promoter choice
- Order of expression cassettes
- Number of expression cassettes
- Spacing of expression cassettes
- Directionality of expression cassettes
- Size of vectors
- Single vector or multiple vectors
- Choice of insulators
- Etc.



Controlling ratios with construct design: 2 ORFs



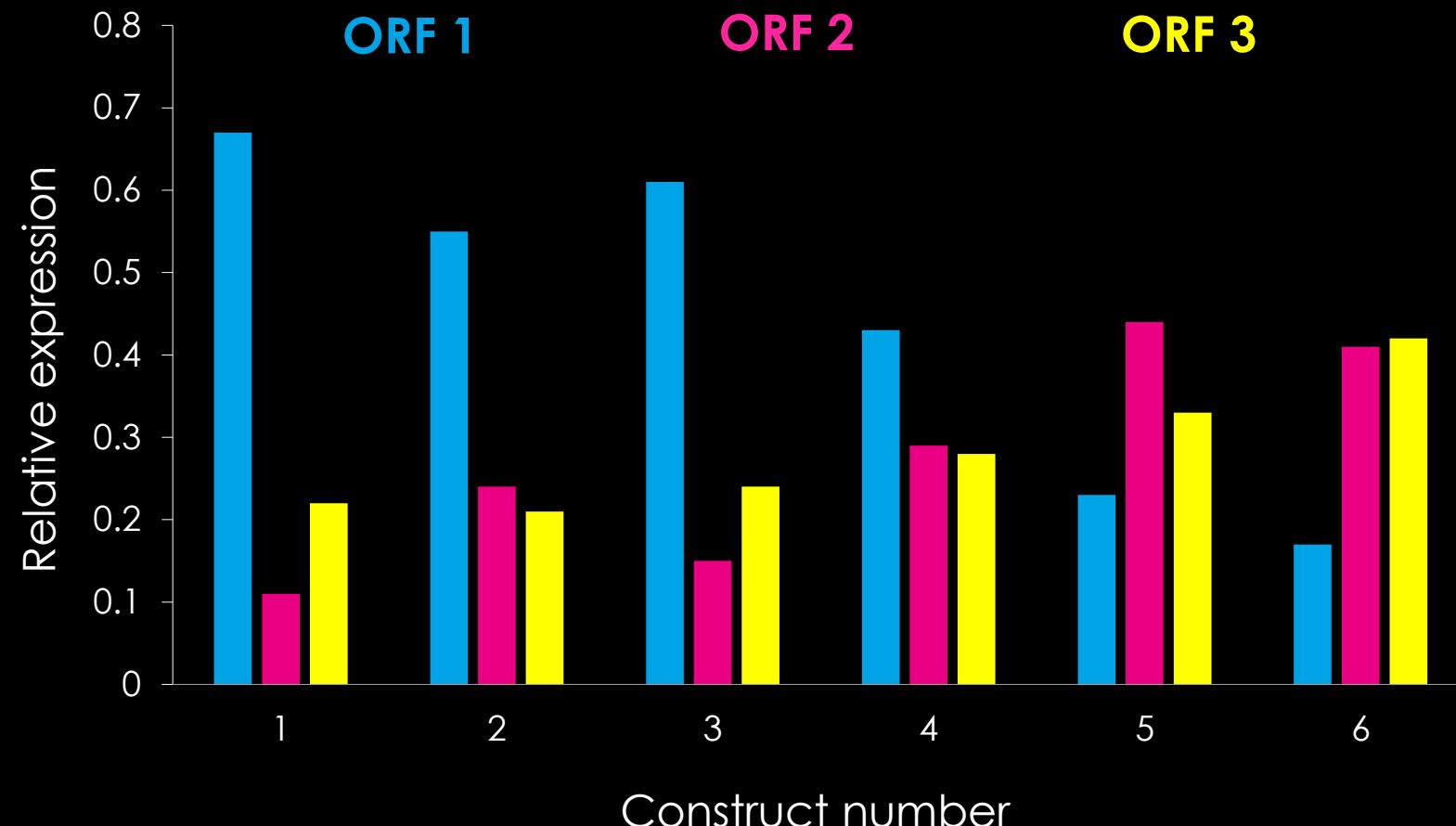
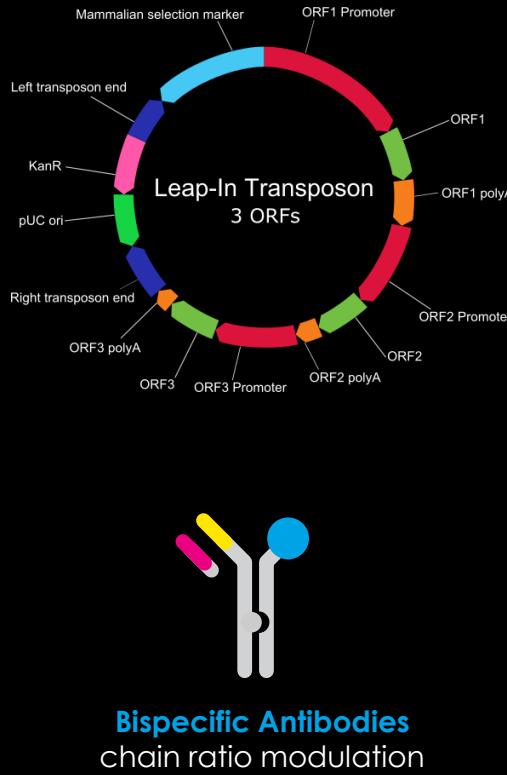
Bispecific Antibodies
chain ratio modulation



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INSPIRED CELL SOLUTIONS

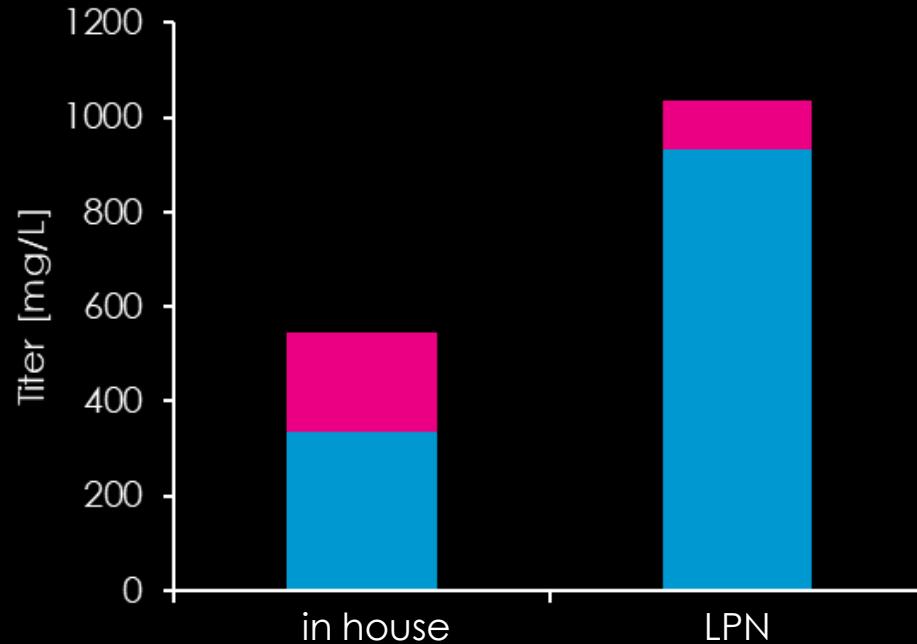
ATUM

Controlling ratios with construct design: 3 ORFs

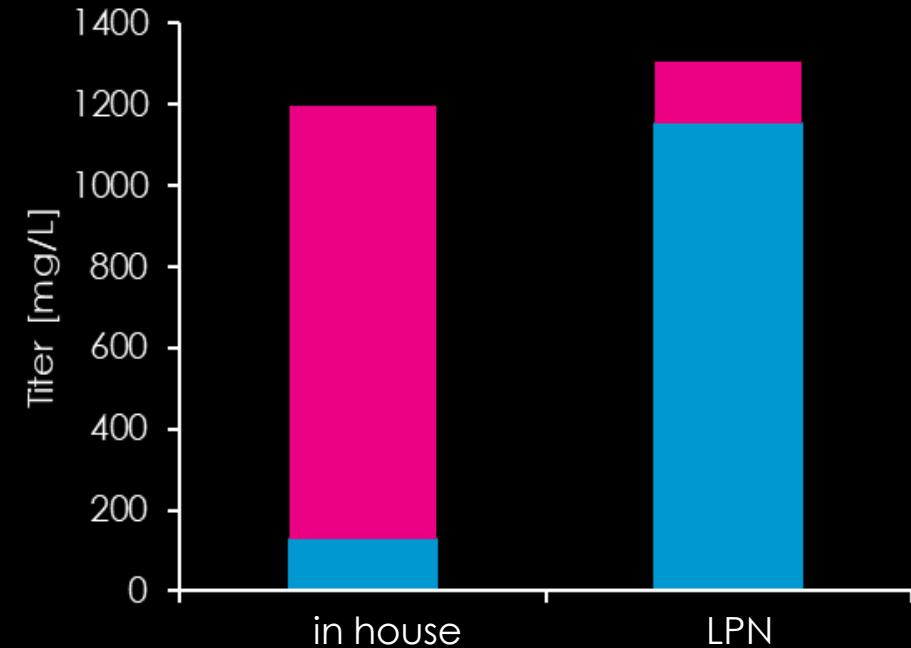


Controlling ratios with construct design: 3 ORFs

Desired product



Assembly variant



Leap-In enabled chain balancing = significantly improved product assembly



Case Study: 3-Chain Bispecific mAb

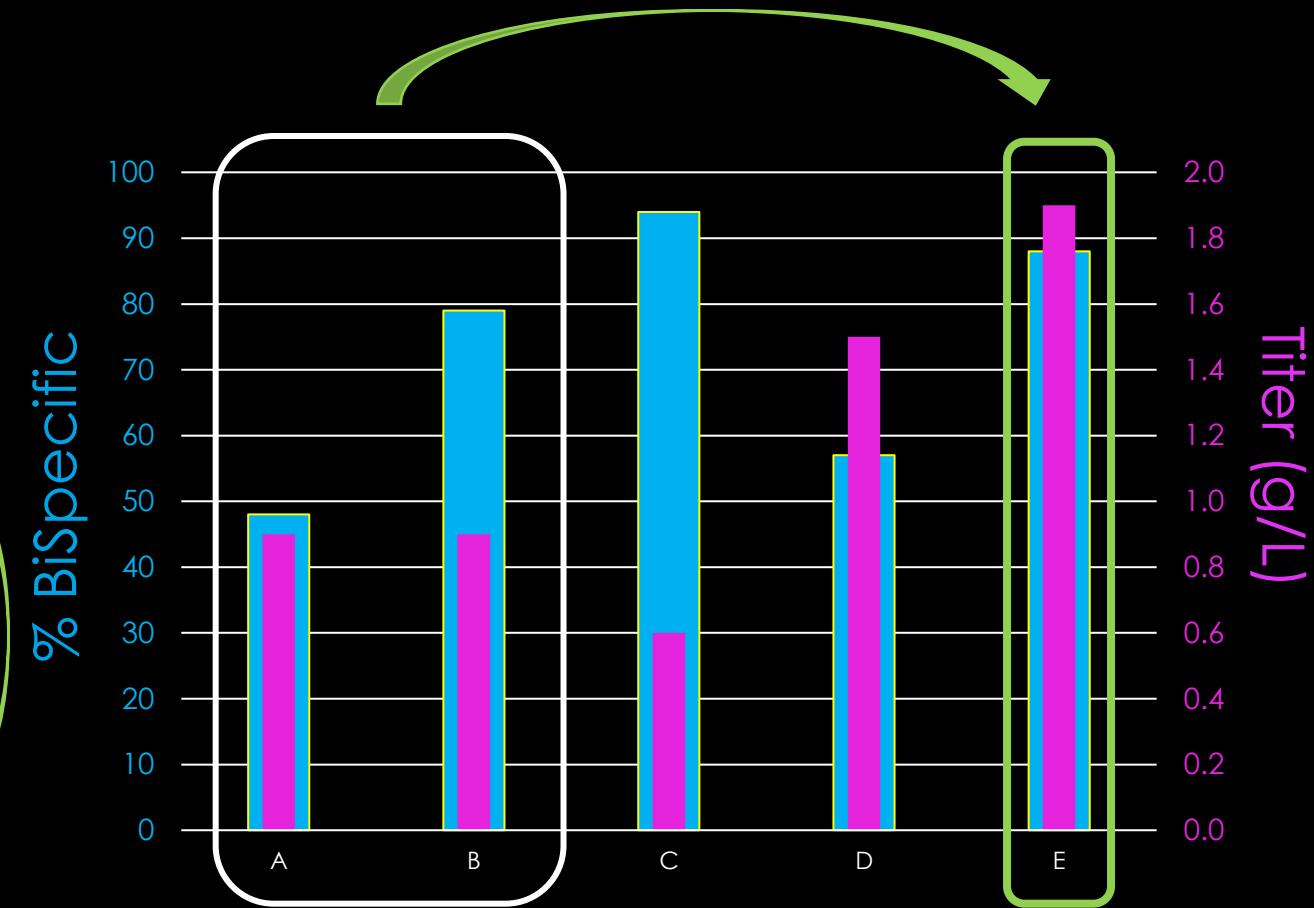
- 14 vector configurations
 - Varying expression levels
 - Varying expression ratios
- Leap In Transposase based pool selection
- Analytical assessment
 - Total titer
 - Chain expression: Relative and Amount
 - % Bispecific



Case Study: 3-Chain Bispecific mAb

Vector*	Expression Level [relative]		Expression Level
	LC	Sum of HC1+HC2 (normalized)	
A	comparable	1	med-low
B	comparable	1	low
C	significantly higher	1	low
D	moderately higher	1	high
E	comparable	1	high

* Subset of 14 vectors screened



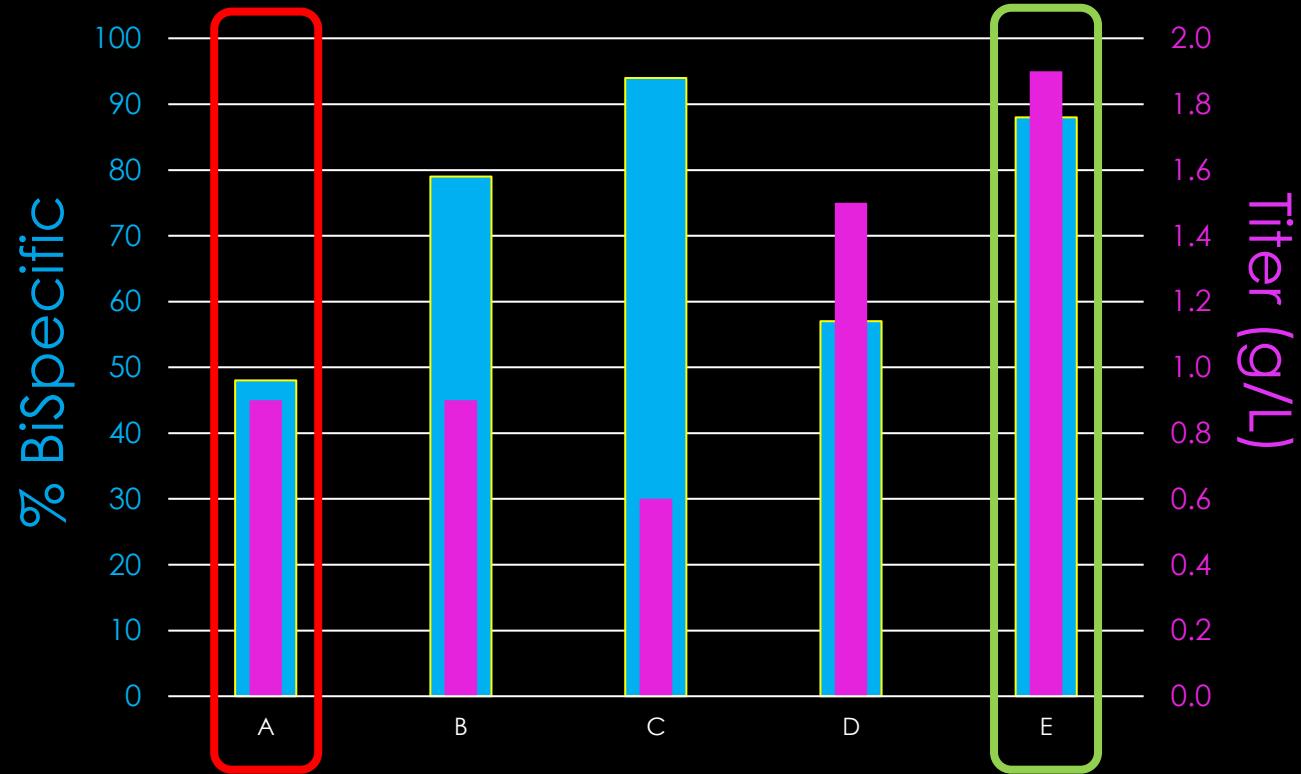
Screening vectors at pool stage
enables ID of high value pools



Case Study: 3-Chain Bispecific mAb

Vector*	Expression Level [relative]		Expression Level
	LC	Sum of HC1+HC2 (normalized)	
A	comparable	low	med-low
B	comparable	low	low
C	significantly higher	low	low
D	moderately higher	low	high
E	comparable	low	high

* Subset of 14 vectors screened

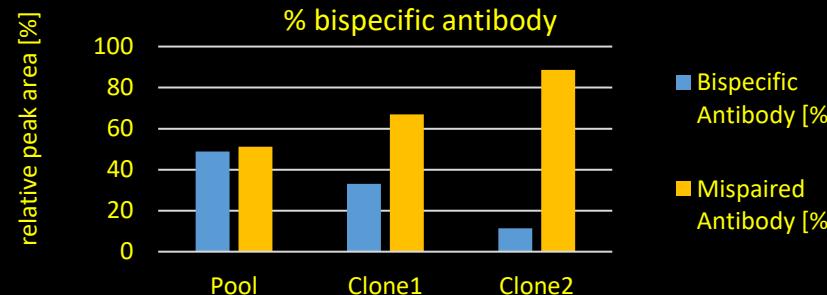
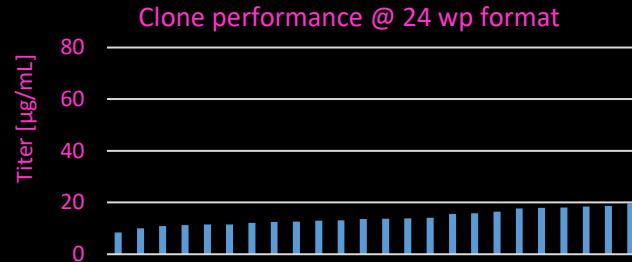


Screening vectors at pool stage
enables ID of high value pools



Case Study: 3-Chain BiSpecific mAb

Pool A

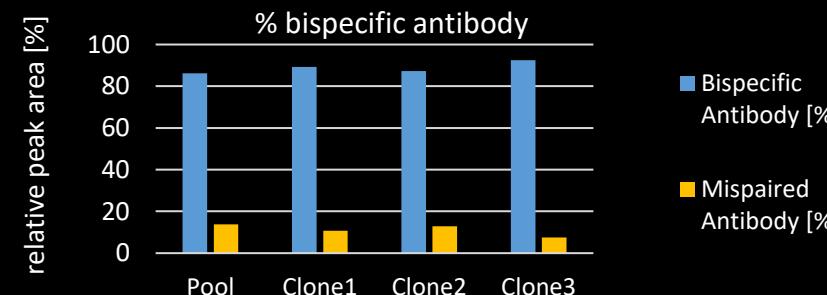
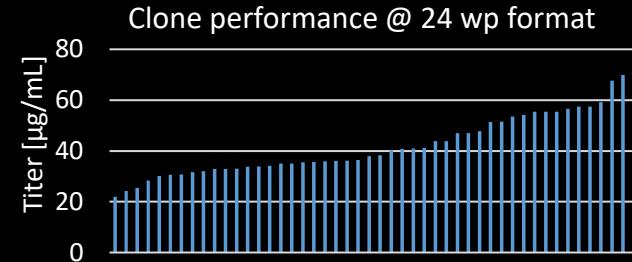


Pool and clone productivity

Pool*	derived clones*
0.9 [g/L]	up to 1.9 [g/L]

*Day 12 standard fed-batch

Pool E



Pool and clone productivity

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

*Day 12 standard fed-batch

Good pools predict good clones



The Power of the Pool

- Pools predict clones
 - Titer, glycan, charge variants, assembly
- Screen vector configurations at pool stage
- Uniquely enabling for complex molecules
- Enables rapid progress towards IND
 - Early transfer to CDMO
 - Process development in parallel with clone selection
 - Pools for tox lot and possibly Ph.I





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

