



Single-cell Multi-omics for Accelerated Therapeutic Characterization and Release

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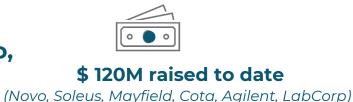
CEO Mission Bio

Moving precision medicine FORWARD.

Our mission is to help researchers and clinicians unlock single-cell biology to accelerate discovery, development, and delivery of advanced therapeutics

Market Leader in Single-cell Multi-omics









>30
Peer-reviewed publications

>60%
NCI Cancer
Centers

1000+ End users worldwide ~100
Assays built for cancer

50M+Genomes sequenced

Integrated DNA & protein analysis from thousands of single cells

Expertise in microfluidics, molecular biology, and informatics

Translational Applications Across Market Segments

THE LEADER IN SINGLE-CELL MULTI-OMICS ANALYSIS



TRANSLATIONAL CANCER CENTERS

Resolve Clonal Heterogeneity



BIOPHARMA

Enhance Biomarker Development



CELL AND GENE THERAPY

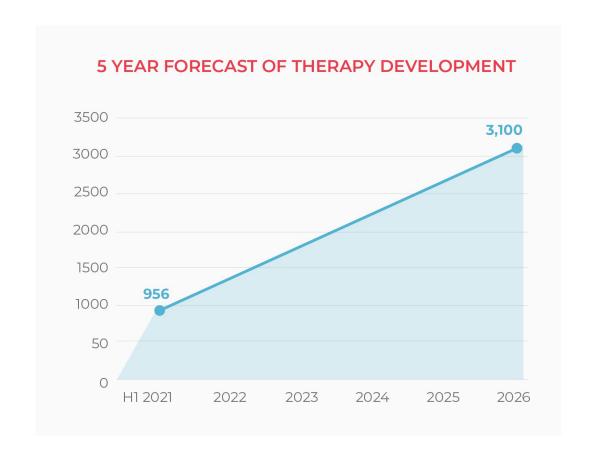
Accelerate Approval And Expand Product Labeling



Rapid Development of Cell & Gene Therapies



- H1 2021 strongest financing half on record
- Increasing # developers around world
- Forecasts predict rapid development of therapies by 2026



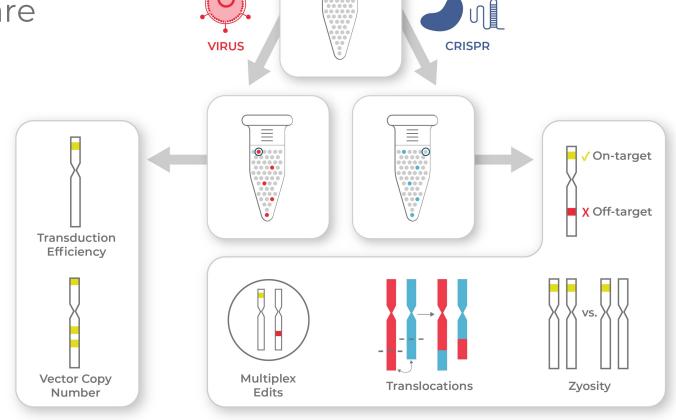


Cell & Gene Therapies Are Complex

Gene-modified cell therapies are heterogeneous

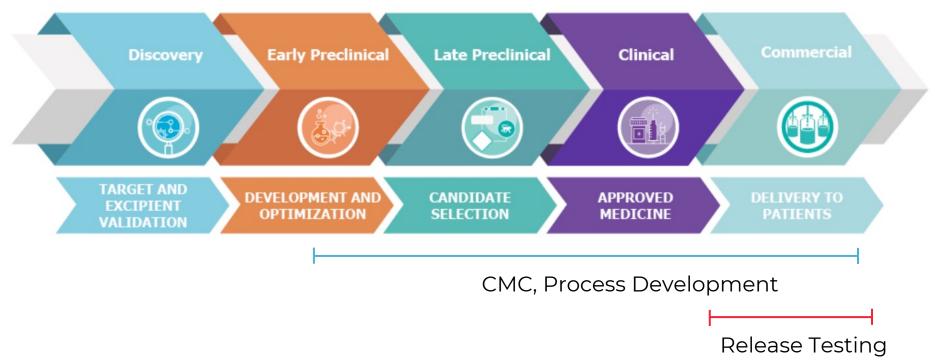
Cell-to-cell variation in:

- Genome engineering alterations by virus or gene editing
- Cell state / type (immunophenotype)





Expedited Pipeline for Therapy Development



- Accelerated development & reduced time to market
- Challenging to develop optimized assays in short timelines
- Regulators require precise, specific, & sensitive analytics for approval & release



Thorough Testing is an Unmet Need in Development

Cell and Gene Therapies Necessitate New Lot Release Test Methods

October 31, 2019

Cynthia A. Challener

BioPharm International, BioPharm International-11-01-2019, Volume 32, Issue 11

Convergence: Inadequate testing of cell and gene therapies draws FDA concern

Posted 15 September 2021 | By Joanne S. Eglovitch

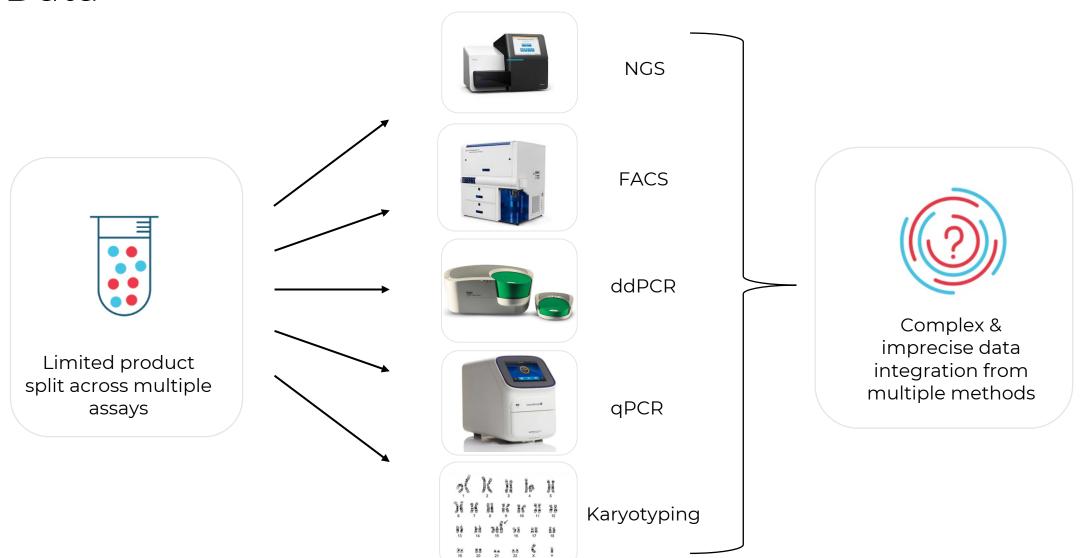
FDA seeking more consistency from cell, gene therapy developers, top official says

Published May 19, 2021

For Cell and Gene Therapy Products, Early Characterization is Critical

- Analytical characterization is often inadequate
- Multiple assays are needed to measure multiple attributes
- Robust release tests needed to ensure batch consistency

Conventional Multi-assay Workflows Yield Complex Data





Mission Bio Can Help

Who we Are

Leaders in high-throughput single-cell analysis with the goal of enabling advanced therapeutic development

Only solution for genotype + phenotype multi-omics at single-cell level

How we Help

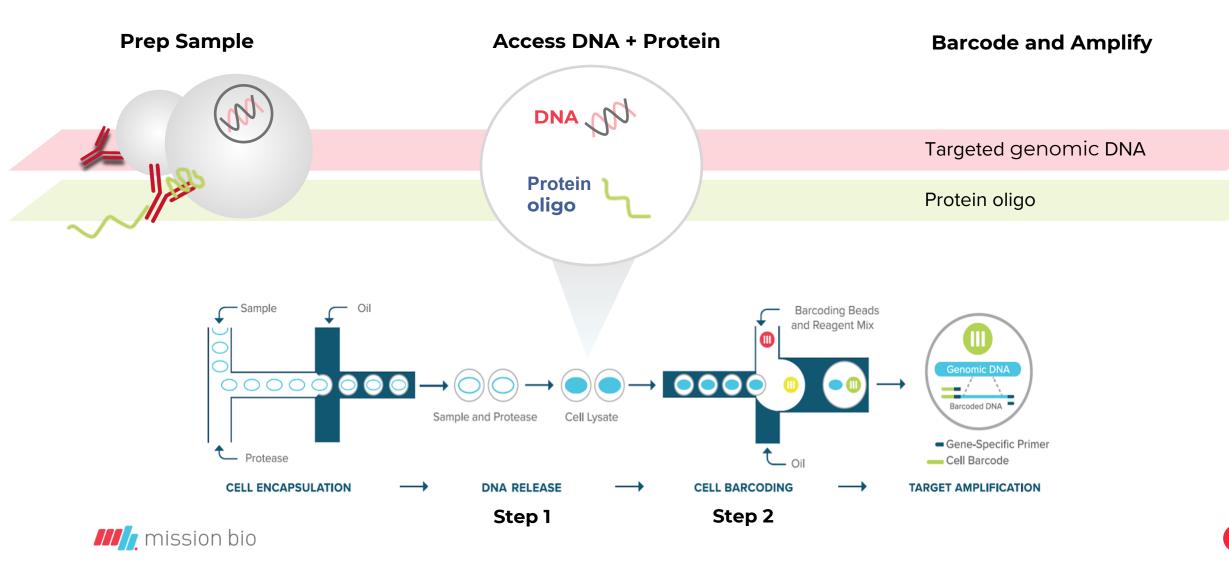
- ✓ Strategic partnership
- √ Consultation
- ✓ Single-cell assay design
- ✓ Wet lab processing
- ✓ Data analysis
- ✓ Assay transfer







Tapestri Enables Single-cell DNA + Protein Multi-omics



Tapestri Cell and Gene Therapy Solutions

Transgene Delivery M. Transduction efficiency **CRISPR** Vector Copy Number (VCN) √ On-target X Off-target Transduction Efficiency Multiplex **Vector Copy Translocations** Zyosity

Number

Edits

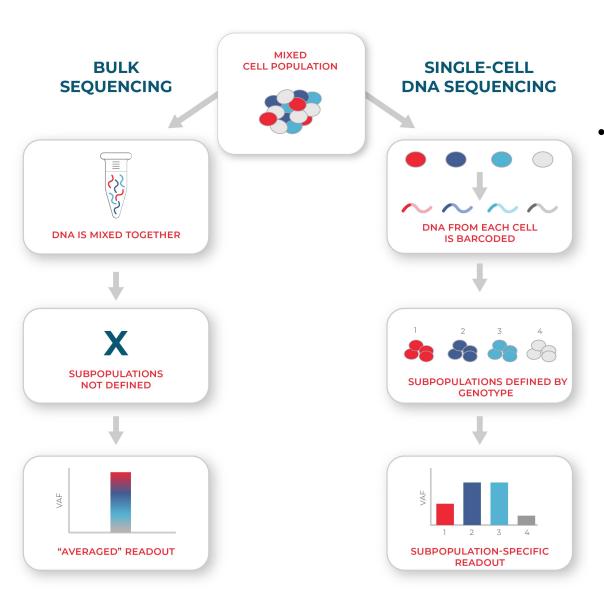
Gene Editing

- On-/ off-target edits
- Multiplex edits
- Chromosome aberrations
- Zygosity of edits



Single-cell Advantages Over Bulk Assays

- Bulk assays mix DNA across cells
- Reports average readouts

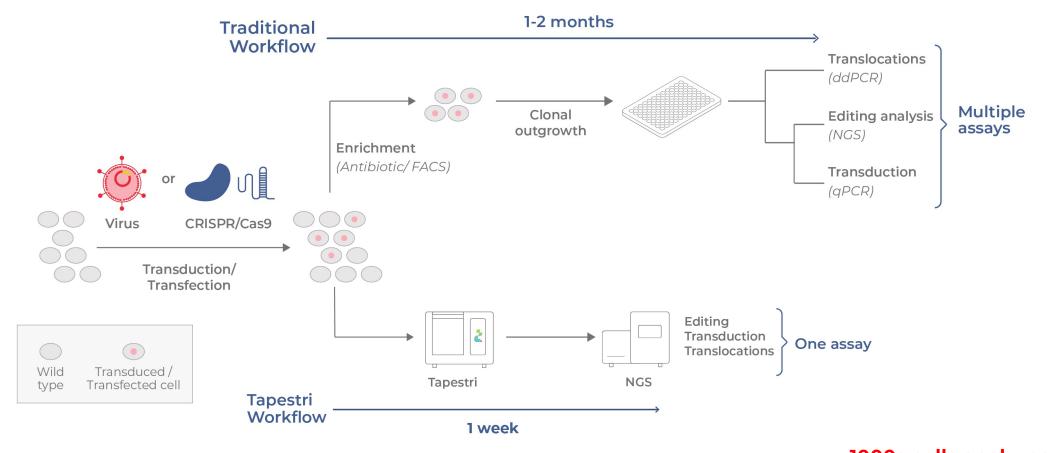


Single-cell DNA sequencing identifies genetic alterations in individual cells

- ✓ Co-occurrence
- ✓ Zygosity
- ✓ Rare events (0.1% of cells)



Single Assay Measures Multiple Parameters in Less Time



- 1000s cells analyzed
- 1-week workflow
- 1 multi-omics assay

- Bypass cell enrichment
- Bypass clonal outgrowth



Analysis of Multiplexed Gene Editing

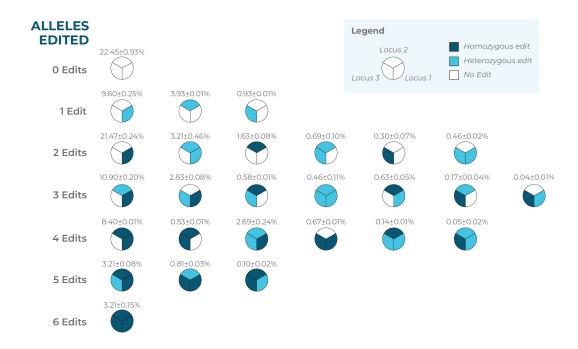
Experiment: Multiplex CRISPR Editing to Obtain Cells with a Triple Knockout

BULK DATA

Edit site 3 Edit s

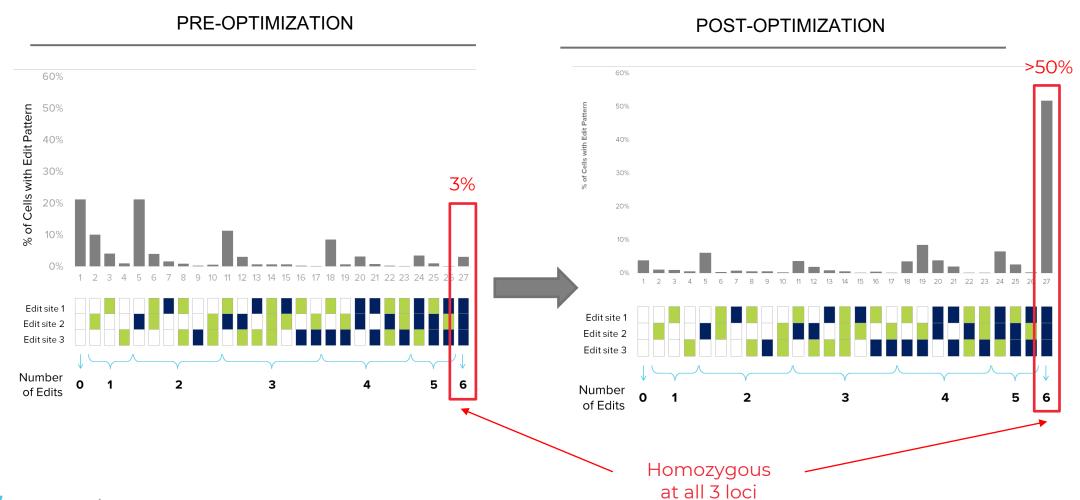
- Bulk shows editing efficiency for each locus
- No zygosity information
- No knowledge of how edits co-occur

SINGLE-CELL DATA



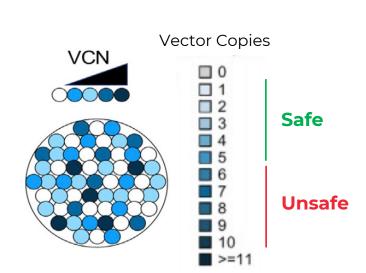
- All 27 different edit combinations detected
- zygosity & co-occurrence of edits

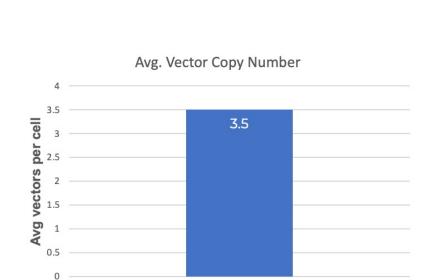
CRISPR Editing Optimization with Single Cell



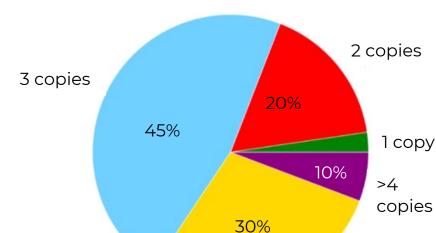


Single-cell Analysis of Vector Copy Number





BULK DATA



SINGLE-CELL DATA

 Cell-to-cell variation in VCN

- Fails to report distribution of cells with different vector copies
- Potentially unsafe cells missed

 % of cells with different numbers of vector copies

4 copies

Potentially unsafe cells detected



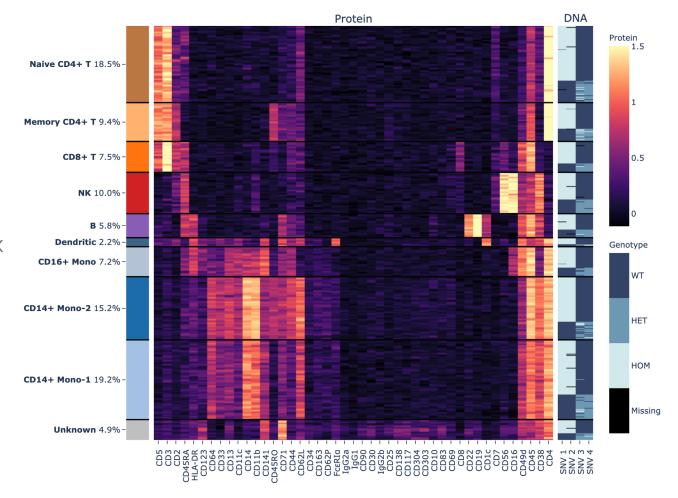
Genotype and Phenotype Correlations at Single-cell Level

Ready to use 45-plex TotalSeq[™]-D Human Heme Oncology Cocktail

A menu of >120 individual TotalSeq-D antibodies by BioLegend available now.

Custom AOCs can be spiked into the 45-plex TotalSeq™–D Human Heme Oncology Cocktail

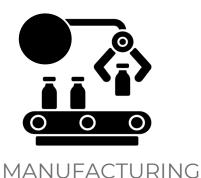
Protein Cluster Signature vs Analyte and Barcode, Subsorted by DNA





Mission Bio's PAD Team Dedicated to Seamlessly Integrate Tapestri into your CGT Workflow





R&D/ Pre-Clinical

Clinical Trials

Commercial

- IND-enabling studies
- CMC, analytics
- Protocol optimization



- Reverse translational studies
- Product & process development



- Release testing
- Patient follow-up

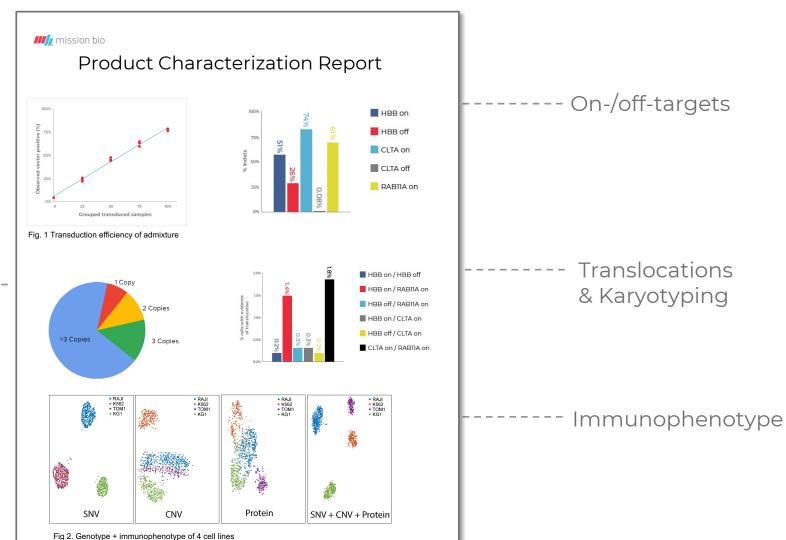




Detailed Analytical Report with Each PAD Project

Transduction -----efficiency

Viral vector copy ---





Single-cell Transduction Assay for Lot Release

Mission Bio Transfers First Tapestri GMP-Ready CGT Assay to Avance Biosciences for Cell-Based Therapies

GMP-compliant test for transduction efficiency marks a key milestone for Tapestri's ability to power single-cell analysis in clinical trial settings for cell & gene therapies

SOUTH SAN FRANCISCO, February 2, 2021 — Mission Bio, the pioneer in high-throughput single-cell DNA and multi-omics analysis, announced today the first tech transfer of a clinical trial-ready cell and gene therapy (CGT) assay from its Tapestri Platform to Avance Biosciences, Inc., a leading-edge CRO based in Houston, Texas, to assess transduction efficiency for an autologous cell therapy currently under development. This collaboration marks the first time the Tapestri Platform will undergo qualification and validation within a good manufacturing practice (GMP) setting, paving the way for its routine use in clinical trials on CGT materials to support the next wave of life-saving CGT treatments.

The Tapestri Platform will allow Avance to analyze cells that have been transduced with lentivirus, identifying transgene integration within individual cells without the need to grow clonal populations. Tapestri analysis not only reduces the timeframe of conventional workflows, but also enables large sample sizes (at the scale of thousands of individual cells) to be rapidly assessed — yielding highly accurate and precise measurements of transduction efficiency.

Assessing the safety and efficacy of CGT products is more challenging than conventional therapies because of the many potential variations between genetically altered cells. Cells modified with a viral vector will differ in transduction efficiency and the number of copies incorporated into the genome. Cells modified with tools like CRISPR can vary in on- and off-target edits, zygosity, or aberrant translocations. Before Tapestri, workflows required synchronizing data from multiple genotypic and phenotypic assays over several weeks, an approach with reduced specificity and limited ability to characterize cells with simultaneous edits.



Single-cell Transduction Assay for Lot Release

Project: Single-cell transduction assay for lot release for gene-modified cell therapies.

1. Lentiviral transduction assay & analytics developed though PAD for proof of concept.



2. Final assay developed & tested across different lots for two products



3. Assay is transferred & validated at Avance (CRO)







4. Use for lot release across multiple drug candidates in clinical trials





Single-cell Analysis is Shaping the Future

- Accommodate accelerated development timelines
 - Multiple assays combined in 1 workflow
 - Assess 1000s of cells in each run
- Improved resolution- single cell level
 - Protocol optimization
 - Safety: detect rare events
- Partnering with Mission Bio
 - Assay design
 - Development
 - Bioinformatics
 - Tech transfer









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