Translation of a complex, 3x CRISPR edited, lentiviral vector engineered T cell process onto the Cellares Automated Cell Therapy Manufacturing Platform



<u>Mercy Gohil</u>, Margaret Karwoski, Irina Kulikovskaya, Fang Chen, Vanessa E. Gonzalez, Joseph A. Fraietta, Julie Jadlowsky, Gabriela Plesa, Megan M. Davis [1] Dmitry Malkov, Anna Sahakyan, Henry Nguyen, Victor Rocha, Julie Fang [2]

[1] Perelman School of Medicine at the University of Pennsylvania, Center for Cellular Immunotherapies, Philadelphia, PA USA [2] Cellares, Inc. South San Francisco, CA USA

Background

The University of Pennsylvania (UPENN) has a longstanding history of administering autologous investigational cell therapy products into humans to demonstrate clinical proof of concept. Investigational cell therapy products implementing gene editing combined with viral integration of transgenes (TCR and CAR constructs) offer potential application in development of universal "off-the-shelf" cell therapy products. The NY-ESO-1-transduced CRISPR 3Xedited cells (NYCE), developed at UPENN, have demonstrated enhanced tumor specificity, improved persistence, and a favorable safety profile in clinical trials (Stadtmauer et al., 2020). This is achieved by knocking out endogenous TRAC, TRBC, and PDCD1 genes while integrating a lentiviral vector encoding for the cancer-specific transgene NY-ESO-1 TCR (8F-TCR). Incorporating multiplex genome editing into a fully-closed clinical cell manufacturing process presents challenges for scalability and automation. The Cellares Cell Shuttle platform was evaluated as a potential solution to address these challenges. Initial feasibility studies generated 3X CRISPR edited transduced cells (Edited 8F-TCR) using the gene editing unit of the platform. Edited 8F-TCR cells generated on the Cell Shuttle instrumentation, demonstrated TCR-mediated, antigenspecific cytotoxicity. These preliminary results suggest potential for the complex manufacturing process of adoptive T cell therapies to be adapted onto a scalable, automated manufacturing platform.



CUSTOMIZABLE ELECTROPORATION PROCESS PARAMETERS

- Cell concentration in electroporation buffer
- Voltage, pulse width, number of pulses
- Electroporation volume per batch
- Number of electroporation batches
- · Post-electroporation recovery temperature and duration

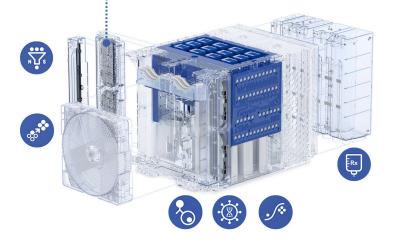


Figure 1. Schematic of the manufacturing process on the Cellares Cell Shuttle technology.

Blue: Automated process steps

Gray: Manual process steps

DAY

-2

0

8



REAGENT PREPARATION

Prepare and store reagents in automationcompatible bottles on the Cell Shuttle for automatic retrieval based on process demand.

ENRICHMENT

Automated enrichment of white blood cells via Centrifugal Counterflow Elutriator.



LABEL WASH & MAGNETIC SELECTION

Automated incubation and removal of unbound selection reagents, followed by magnetic selection.

ACTIVATION

Automated transfer of selected cells and activation reagent to bioreactor in the CC and incubation for 48h at 37 °C.

ELECTROPORATION

Automated buffer exchange and concentration, followed by execution of customized electroporation protocol.

TRANSDUCTION

Automated transfer of transduction reagent and incubation for 24h at 37°C.

EXPANSION

Automated perfusion expansion to reach target dose of Edited 8F-TCR cells

HARVEST WASH

Automated harvest and concentration, followed by CC removal from the Cell Shuttle

FILL & FINISH

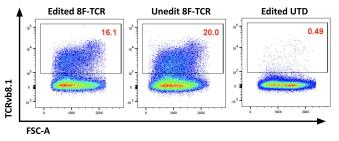
Drug substance removal from CC, followed by formulation and cryopreservation using a controlled-rate freezer

Results

f 1 High efficiency CRISPR-Cas9-mediated modifications and successful detection of the engineered TCR

The frequencies of TRAC, TRBC, and PDCD1 gene disruptions were quantified using Inference of CRISPR Edits (ICE) from Sanger sequencing data. Edited 8F-TCR sample achieved high frequency of gene disruption (>84% across all genes), while no gene disruption was detected in the unedited control (Unedited 8F-TCR). The frequency of transgenic TCR (Vb8.1) was assessed via flow cytometry, achieving 16% in the thawed final product.

Sample	Description	Editing Efficiency (%)			Transduction
		TRAC	TRBC	PDCD1	Efficiency (%)
Edited 8F-TCR	Transduced (8F-TCR) and Electroporated (CRISPR 3X)	93	84	91	16.1
Unedited 8F-TCR	No Electroporation Control	0	0	0	20.0
Edited UTD	No Transduction Control	93	83	91	0.5

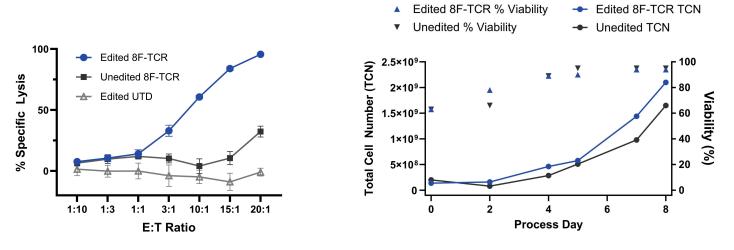


Antigen-specific cytotoxicity against NY-ESO-1-expressing cancer cells

Edited 8F-TCR effector cells (E) exhibited TCR-mediated dose-dependent lysis of NY-ESO-1 expressing Nalm6 target cancer cells (T). In contrast, unedited 8F-TCR cells exhibited less than 10% lysis of target cells with E:T ratios up to 15:1, while Edited UTD cells showed no lysis across all tested ratios.

3 In-process analytics demonstrate automation feasibility

Large-scale expansion of Edited 8F-TCR cells demonstrated high viability (>94%) and fold-change (15x) by day 8. Higher cell number is expected with further expansion. Memory phenotype of Edited 8F-TCR cells was comparable to smallscale controls (data not shown).



Conclusion

We successfully translated an engineered T cell process, incorporating multiplex genome editing (Edited 8F-TCR), onto the Cellares Cell Shuttle instrument, producing cellular material that exceeded potency, efficacy, and safety requirements.

The endogenous TCR was effectively replaced with the cancer antigen-targeting 8F-TCR, achieving stable expression, as confirmed by flow cytometry. The engineered T cells exhibited strong cancer-specific cytolytic activity and IFN-gamma release in a co-culture assay, highlighting their potential for targeted immunotherapy.

These results demonstrate the capability of the Cellares Cell Shuttle technology to manufacture engineered T cells while streamlining complex workflows to meet the growing patient demand for these life-saving cell therapies.