



Achilles Therapeutics AI-Powered Precision TIL Therapy

May 2022

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Precision targeting for solid tumors using clonal neoantigen-reactive T cells (cNeT)





Clonal neoantigens: a novel class of cancer target present on all tumor cells Developed a proprietary AI platform (PELEUS[®]) validated on real world patient data (TRACERx study)



Trackable precision T cell therapies

Able to target multiple tumor antigens with unprecedented precision and tracking in patients



Differentiated, scalable manufacturing

VELOS[™] manufacturing process designed to be closed and automated for commercial scale





Multiple near-term catalysts

Higher-dose cNeT monotherapy cohort and initial cNeT / PD-1 inhibitor combo cohort data in 2H 2022



Strong cash position

London (HQ) & Philadelphia with ~250 employees and cash runway into 2H 2024 (\$237M @ Mar 31, 2022)

Our Management team



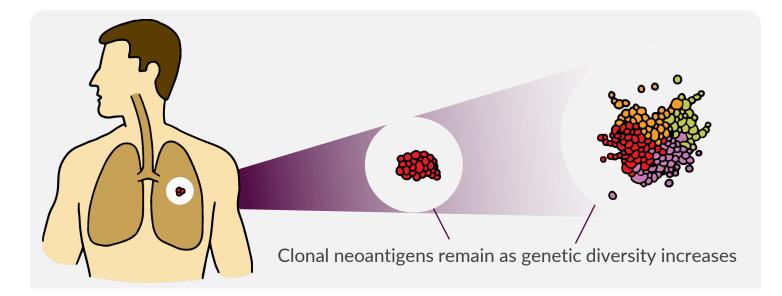




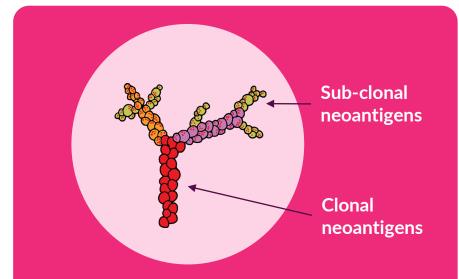


Ideal cancer targets: clonal neoantigens present on all tumor cells and absent from healthy tissue





- Tumors are constantly evolving and acquiring new mutations
- Genetic diversity enables tumors to develop resistance to standard therapies (e.g., loss of a target) which leads to disease relapse
- Despite increased genetic diversity the original mutations (clonal neoantigens) always remain¹
- Achilles' technology identifies the original (clonal) mutations and enables targeting of multiple antigens present only on tumor cells



Clonal neoantigens are

- Original mutations formed early in tumor evolution
- Expressed on all cancer cells¹ and absent from healthy tissue
- Found in multiple tumor types as tumor evolution principles apply across multiple solid cancers

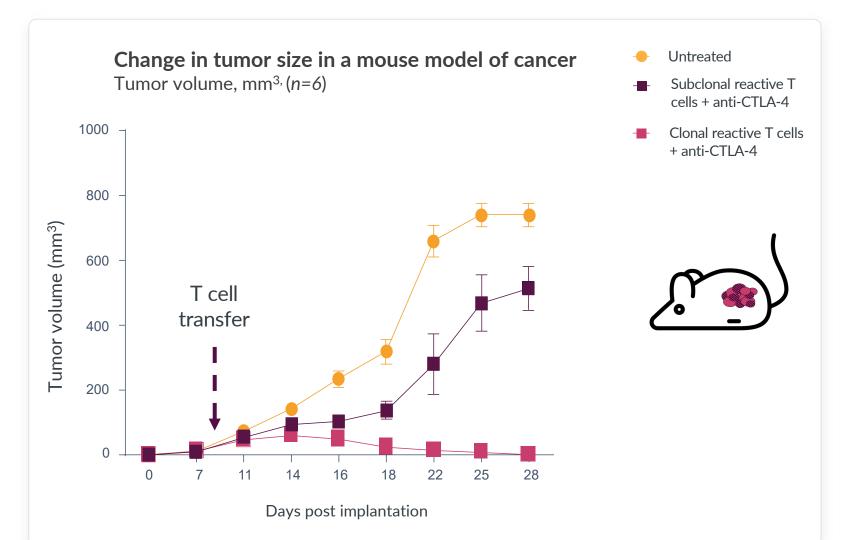
Clonality is key to the eradication of tumors

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Mice implanted with tumor cells expressing sub-clonal (on a sub-set of cells) and clonal (on all cells) antigens

Mice were then treated with murine T cells reactive to sub-clonal (■) and clonal antigens (■)

Targeting clonal antigens expressed on every tumor cell leads to complete tumor regression in this model





Clonal neoantigen identification is computationally complex

- Solving the heterogenicity problem requires processing of very large DNA data sets
- The process requires multiple steps and has no universally accepted protocol
- 22 groups using the same DNA sequencing data showed <20% overlap of neoantigen identification¹



Compares tumor DNA to healthy DNA to differentiate clonal and subclonal neoantigens

Achilles has developed an Alpowered solution

- Accurate neoantigen identification requires an advanced computational approach
- Al and machine learning have been developed to enable accurate and rapid processing of very large complex data sets
- Neoantigen predictions are then validated with real-world patient data



TRAOER

- Commercial rights to the largest longitudinal patient data set¹⁻⁴
- 3,200 tumor regions collected from 795 NSCLC patients over 5 years
- Multi-region data from primary & metastatic sites to confirm clonal status
- Unparalleled network of 15 NHS sites

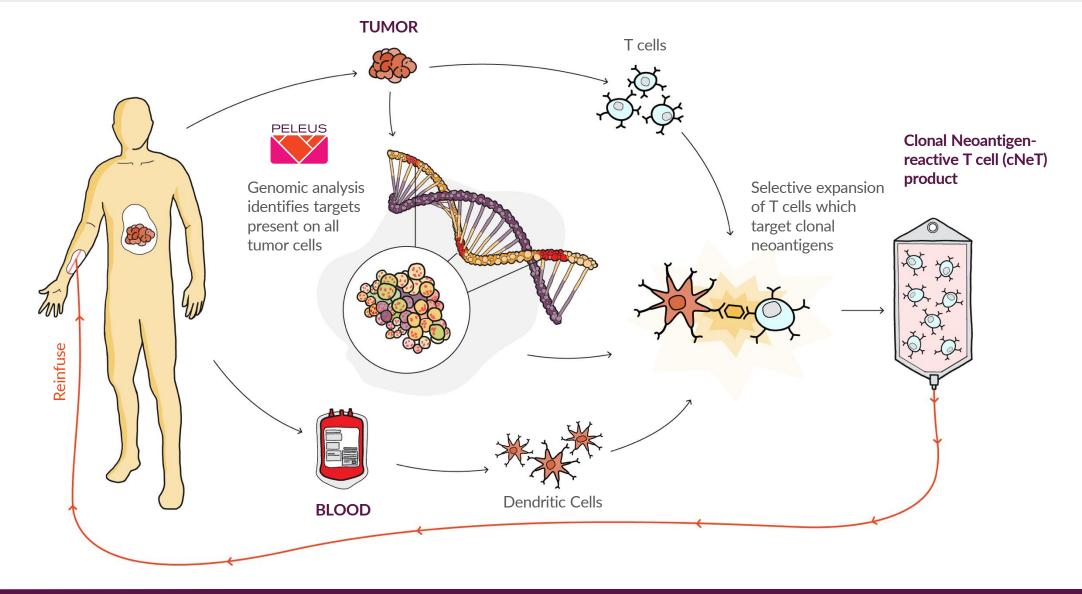


Reactive T cells are identified against clonal neoantigens

- Identification of T cells reactive to predicted clonal neoantigens in patient samples validates the PELEUS[®] platform
- Over 120 patients prospectively analyzed to date across multiple cancer types
- Patient tumor samples validate and train PELEUS[®] AI predictions

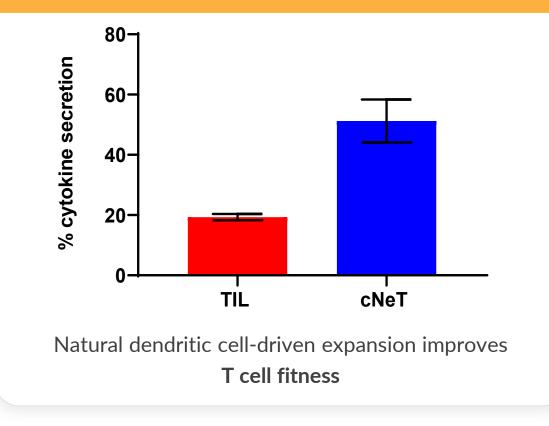
VELOS[™] process delivers precision clonal neoantigen targeting TIL therapy Cutting edge personalized genomics and machine learning enable targeting of all cancer cells









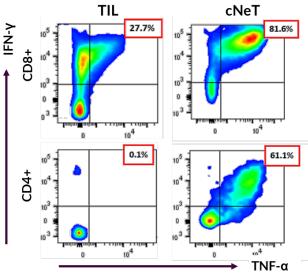


1. Achilles data measuring the ability of T cells to produce inflammatory cytokines in response to the same polyclonal stimulation

T cell specificity and function²

Cytokine secretion measured through flow cytometric analysis, n=1

VELOS[™] selectively expands tumor-derived clonal neoantigen reactive CD4+ and CD8+ T cells with improved **specificity and potency**



CD4+ and CD8+ T cells can work in concert to deliver robust and durable responses³⁻⁵

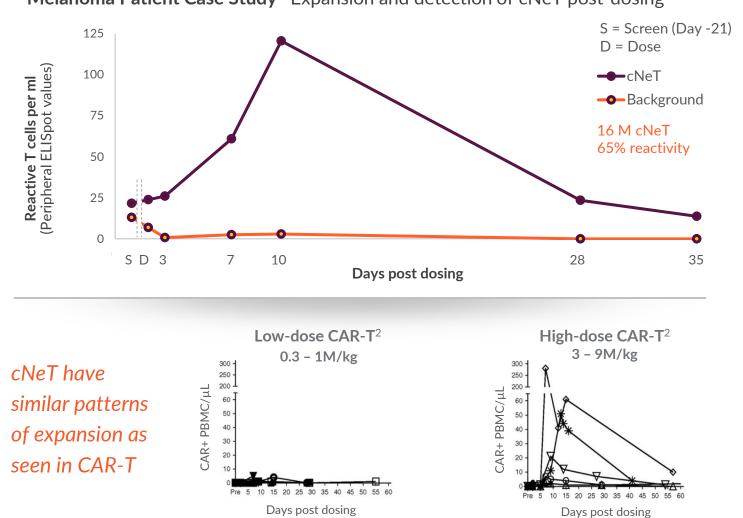
2. Achilles' data measuring the production of inflammatory cytokines in response to clonal neoantigens

- 3. Hunder et al., NEJM, 2008
- 4. Church et. al., Eur J Immunol, 2014
- 5. Antony et al. J Immunol, 2005

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The Achilles cNeT platform allows:

- Quantification of tumor reactivity and cNeT dose for each product (not possible with TIL therapy)
- Determination if increasing cNeT dose improves cNeT persistence and activity as seen in CAR-T therapy
- Correlation of cNeT dose and persistence with anti-tumor effect
- Product release on cNeT dose and basis for potency assays



Melanoma Patient Case Study¹: Expansion and detection of cNeT post-dosing

Aim to deliver clinical doses of >100M cNeT from our VELOS[™] manufacturing process



	Process 1	Process 2		VELOS Manufactured Cell Doses Compared to estimated lowest clinically active doses of
Pre-Expansion	T cells are harvested from the tumor	Cytokines boost tumor- reactive cell harvest		other T cell therapies
Expansion	Dendritic cells loaded with clonal peptides activate and drive cNeT expansion	Optimized DC-driven co- culture followed by short T cell boost increases final cNeT dose	T cell Dose	100 M 50 M 10 M 10 M Viral T cells ⁵
	2 • Maintains high for effector memory	Acturing timeline to Process 1 Anctional cell fitness and A phenotype France, Germany & Spain		Process 1 ¹ Clinical Trial Data

1. SITC 2021 Poster 543 2. ESMO IO 2021 Poster 58P

3. J Clin Invest 2021, Kristensen 4. Blood 2017, Mueller 5. Lancet 2003, Peggs

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Achilles has two ongoing Phase I/IIa clinical trials



Advanced NSCLC

CHIR

Melanoma

THETIS

Monotherapy

- Advanced unresectable or metastatic Stage III-Stage IV NSCLC
- Never-smokers and EGFR/ALK/Ros-1 mut excluded
- Open-label
- n = up to 40
- Option to open Cohort B in combination with a PD-1 inhibitor

Cohort A – Monotherapy

- Recurrent or metastatic malignant melanoma (n = up to 40); Open-label
- Acral, uveal and mucosal melanoma excluded

Cohort B – Combination with PD-1 inhibitor (nivolumab)

- n = up to 20 checkpoint refractory patients; Open-label
- CPI dosed 7-13 days prior to cNeT and restarted day 14 post-cNeT

Evaluating safety, tolerability and activity (RECIST), biomarkers of clinical activity and bespoke ctDNA assay

Ongoing in UK, EU and US

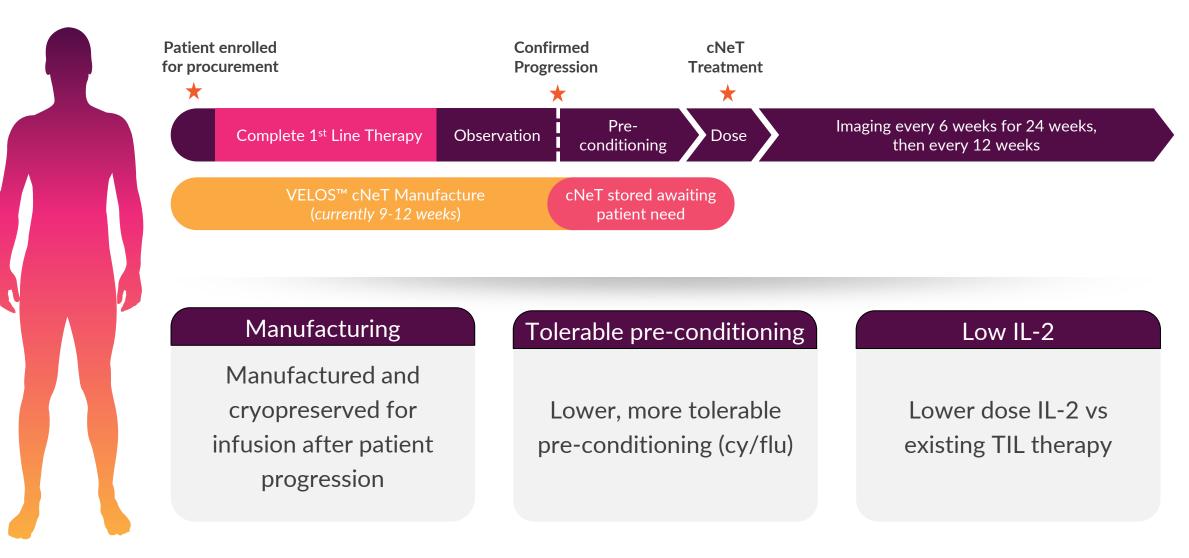
Evaluating safety, tolerability and activity (RECIST)

Ongoing in UK, expanding to EU & US

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cNeT therapies can be readily delivered within standard treatment pathways







Heavily pretreated patients with advanced cancer

- 2.5 median lines of prior therapy
- 14.2m median cNeT dose¹ in monotherapy
- All had progressive disease at time of lymphodepletion
- Five recurrent or metastatic malignant melanoma patients (THETIS)
- Three advanced unresectable or metastatic NSCLC patients (CHIRON)

cNeT tolerability profile

- Tolerability similar to standard TIL products not enriched for cNeT reactivities
- No higher-grade adverse events more commonly associated with the use of higher doses of IL-2
- Three SAEs deemed unlikely related and one ICANS event deemed to be possibly related to cNeT treatment



Activity 6-weeks post treatment

- Stable disease in 5 of 8 patients¹
- **Tumor reduction** in 2 of 4 lesions (55% and 90%) in patient with highest cell dose
- **Engraftment** in 5 of 7 patients, with dose associated response
- 2 to 28 reactivities observed in 7 of 8 cNeT
- Median dose of 14.2 M

Patient	cNeT Dose (M)	Reactivity	Engrafted ²	Best Response
C-03	0.1	0.20%		SD
T-12	2	13%	Y	SD
T-09	12	9%	Ν	SD
C-11	13	41%	Y	SD
T-05	16	65%	Υ	PD
C-10	21	3%	Ν	SD
T-11	42	5%	Y	PD
T-02 ³	287	77%	Y	PD

Median 2.5 lines of prior therapy

Next steps: Explore higher-dose cNeT monotherapy and combination with PD-1 inhibitor

- Data reported SITC Nov 12, 2021
- cNeT detected post infusion
- T-02 treated with steroids on day 11 to treat infectior

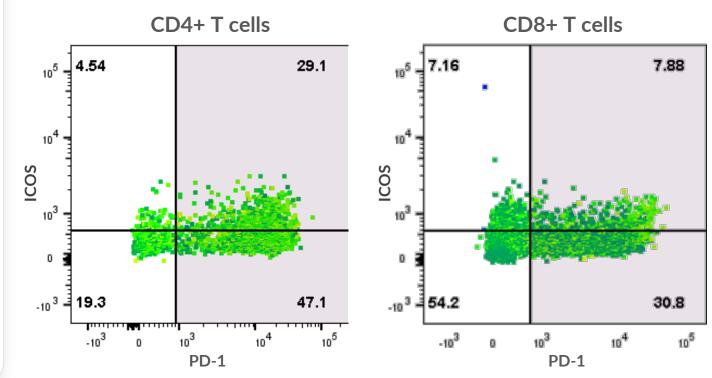
Potential for cNeT/PD-1 inhibitor combination in Melanoma (THETIS Cohort B)

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Checkpoint molecules in immune regulation

- Normal immune regulation uses immune checkpoints to inhibit T cells from attacking healthy tissue
- Our data show PD-1 checkpoint is upregulated on a fraction of cNeT which could partially reduce anti-tumor activity¹
- Targeting PD-1 with checkpoint inhibition could further increase cNeT activity

TIL-derived cNeT express elevated levels of PD-1 upon antigen encounter¹ Cell surface marker flow cytometric analysis (*n*=1)

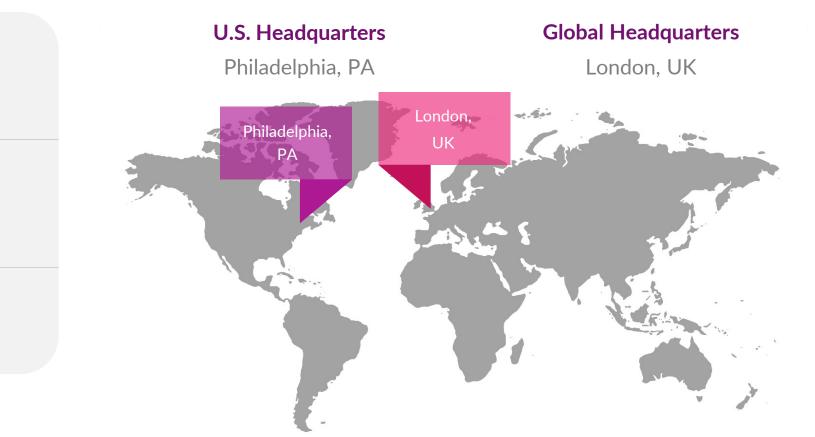


Scale-up of GMP manufacturing for late stage clinical trials and commercial launch









Global Operations

\$237M

CURRENT CASH BALANCE¹

2H 2024

CURRENT CASH RUNWAY

Key anticipated 2022 milestones and updates



	2022						
	Q1	Q2	Q3	Q4			
Higher-dose cNeT (Process 2)		First Patient Dose (Process 2)	Initial Process 2 monotherapy patient data	Initial Process 2 monotherapy patient data			
PD-1 + cNeT combo (Process 1 & 2)	Open CPI combo cohort (melanoma)		First Patient Dose (combo)	Initial combo patient data			
Facilities and Manufacturing	Establish US HQ and R&D	Catapult GMP License & US CDMO Partnership					
Tumor Archiving Program (TAP)		Initiate TAP program					

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NASDAQ:

ACHL





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