



Achilles Therapeutics Al-Powered Precision Cell Therapy Targeting All Tumor Cells

March 2023

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Transforming the treatment of solid tumors with precision T cell therapy



Company founded 2016

Nasdaq IPO: ACHL 2021 Early clinical proof of concept 2022

Clinical update 2023



Global Headquarters London, UK



Two active clinical programs

~200 employees

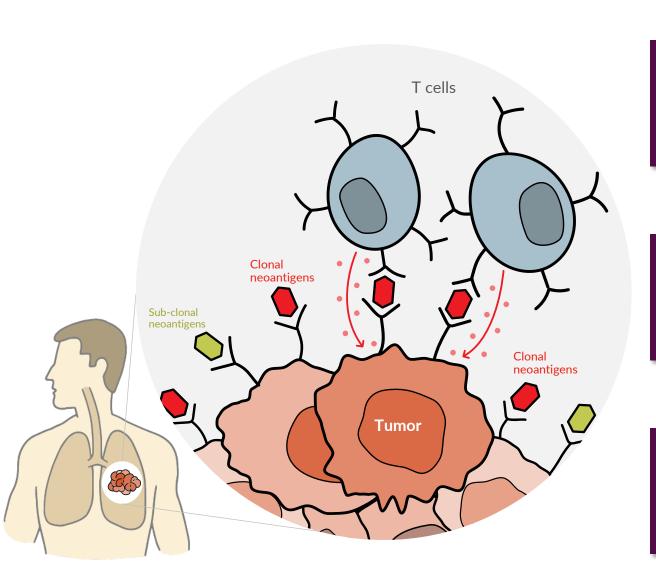
\$173M cash

U.S. Headquarters Philadelphia, PA



Targeting clonal neoantigens with patented technology, linking mechanism and potency





Clonal neoantigens: a novel and ideal cancer target Only target present on all cancer cells, absent from healthy tissue

Unique and world leading capability to identify clonal neoantigens

Only neoantigen platform validated on real world data

World class scientific platform

Demonstrated target engagement supporting mechanism of action

Clinical-stage precision targeting for solid tumors using clonal neoantigen-reactive T cells (cNeT)





Targeting clonal neoantigens: a novel class of cancer target present on all tumor cells

We have developed a proprietary patent protected AI platform (PELEUS®) that is validated on real world patient data (TRACERx) and which can be used to identify personal clonal neoantigens



Controlled precision therapy

Scientific platform that can quantify, characterize and track tumor reactive T cells, target engagement and mechanism of action



Emerging PoC for cNeT in NSCLC

Durable disease control achieved with cNeT monotherapy, 71% (5/7) NSCLC patients (including 1 PR and 4 SDs) with encouraging safety and tolerability



Near-term clinical milestones

Clinical and translational updates in 2023: 15-20 new patients across NSCLC (CHIRON) monotherapy and melanoma (THETIS) monotherapy and in combination with check-point inhibitor (anti-PD-1)



Strong cash position supports all planned operations into mid-2025

Cash runway of \$173M (£143M) as of December 31, 2022

Experienced leadership with decades in cell therapy drug development









Robert Coutts CFO



Syncona



CSO

Iraj Ali CEO

Syncona McKinsey&Company



Daniel Hood General Counsel







Shree Patel EVP, Patient Supply Operations

Cell Medica



Jim Taylor CBO



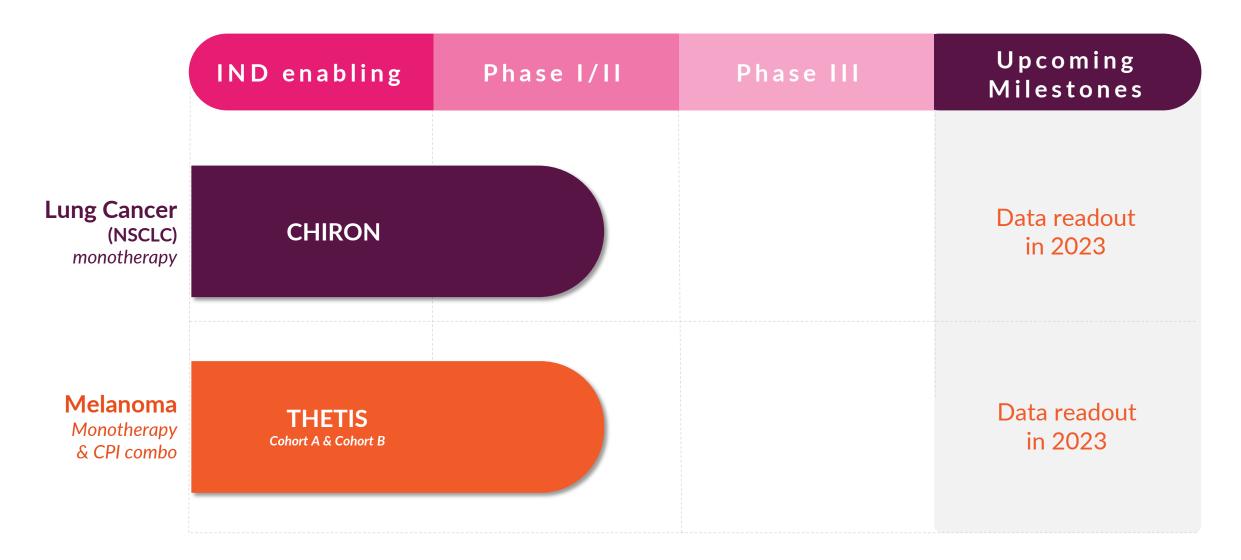
Ed Samuel EVP, Technical **Operations**





Differentiated pipeline of precision T cell therapies across multiple solid tumors





Cancer is driven by mutations to DNA which create targets for the immune system

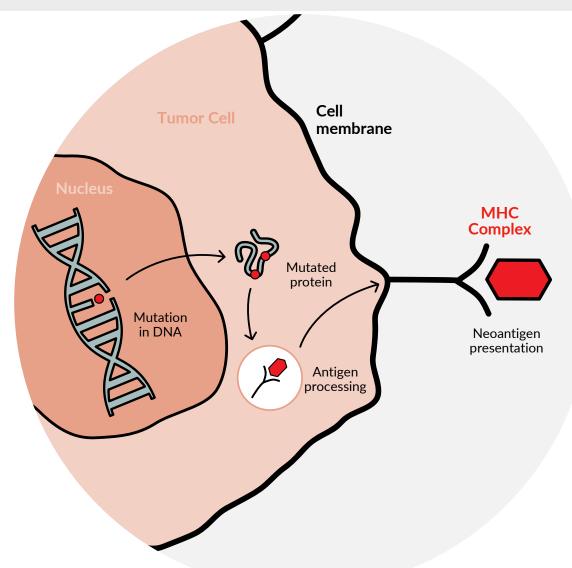


DNA damage causes genetic alterations which can lead to cancer

Mutated proteins from these alterations create antigens for the immune system

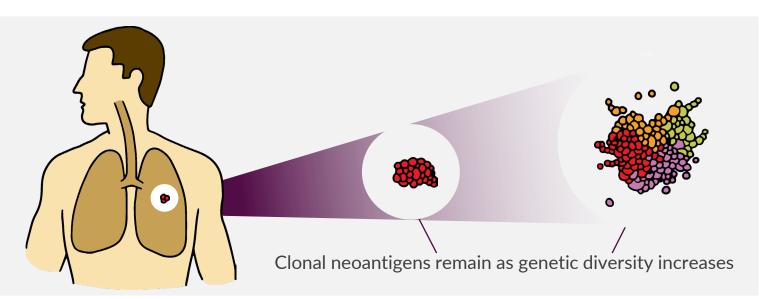
Neoantigens presented on cell surface via MHC molecules recognized by T cells

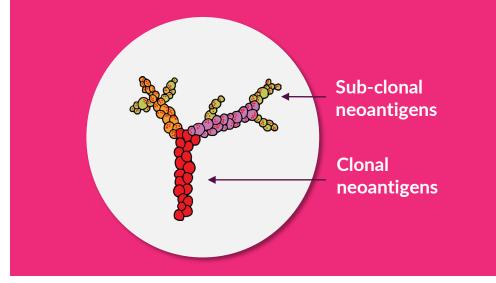
T cells will recognize neoantigens as foreign and destroy the tumor cell



A landmark study (TRACERx) demonstrated that a specific group of neoantigens (clonal neoantigens) are present on all tumor cells in cancers





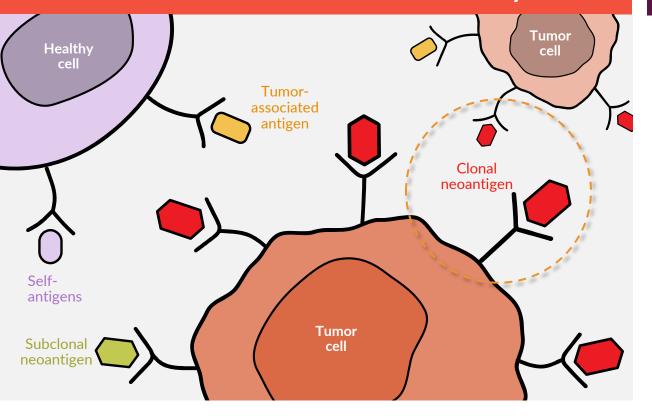


- Tumors are constantly evolving and acquiring new mutations
- This genetic diversity enables tumors to develop resistance to standard therapies
- However, original mutations (clonal neoantigens) are passed down and remain present in all tumor cells¹⁻⁴
- We can identify the original clonal mutations for each patient and so can target multiple antigens present only on tumor cells²⁻⁴

Multiple lines of clinical evidence support neoantigens as relevant cancer targets and clonal neoantigens as the predominant predictive biomarker for positive clinical outcome



Clonal neoantigens: the only known target present on all tumor cells and absent from healthy tissue



Multiple lines of evidence link neoantigens with positive clinical outcomes

- Neoantigen-reactive T cells are correlated with improved outcomes for CPI and TIL therapy¹⁻³
- Disease-free survival is driven by the presence of clonal neoantigens in untreated NSCLC⁴
- Evidence that clonal neoantigens are the driver of overall survival in checkpoint (CPI) therapy⁵⁻⁷
- mRNA vaccines targeting neoantigens now clinically validated showing recurrence-free survival benefit vs anti-PD-1 alone⁸

Litchfield et al. Cell 2021

Lauss et al. Nat Commun. 2017 Nov 23:8(1):1738

Kristensen et al. J Clin Invest. 2022 Jan 18;132(2):e150535

Rosenthal et al. Nature. 2019. 567 479–485

Rizvi et al. 2015 Cancer Immuno 348(6230):124-8

^{6.} McGranahan et al. 2016 Science 351:1463-1469

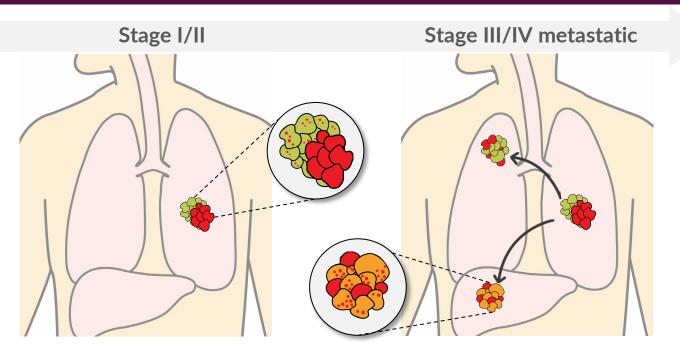
^{7.} Litchfield et al. Cell 2021

https://clinicaltrials.gov/ct2/show/NCT03897881

TRACERx is a unique asset that enables Achilles' neoantigen identification capability



TRACERx: patients (815) enrolled with early stage to advanced NSCLC and followed over several years with extensive tumor sampling, DNA sequencing and neoantigen analysis



- Biopsies taken over five years tracking disease progression
- DNA sequencing and bioinformatic analysis confirm which neoantigens (clonal) are conserved at all tumor sites (red)

TRAGER

- TRACERx is the largest longitudinal real-world patient data set of its kind¹⁻⁴
- Unique asset with a network of 15 NHS hospital sites using a harmonised protocol
- Extensive sequencing data (>4,000 biopsy samples across 815 patients) identify clonal neoantigens at primary and metastatic sites¹⁻⁴
- Clonals neoantigens proven to be present in all tumor sites, at all time points⁴
- Clonal neoantigens can be identified by their specific sequence "signatures" using our patent protected PELEUS platform

PELEUS: Patented Al-driven neoantigen prediction model built and validated on real-world data



Al-powered neoantigen prediction

- Neoantigen identification requires an advanced computational approach
- Al and machine learning developed to enable reliable and rapid processing of complex patient DNA data
- Our neoantigen prediction method is patented and validated with realworld patient data

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



Method for identification of clonal neoantigens can be applied to multiple tumor types

PELEUS: An industry leading neoantigen platform

- Uniquely positioned to validate our clonal neoantigen predictions (>10,000 predicted clonal neoantigens validated with human patient data)
- Our AI-platform can predict neoantigen immunogenicity based on real world T cell reactivity data
- Our extensive real-world patient datasets can be mined for targetable mutations and associated TCRs

Clonal neoantigens can be targeted with a range of therapeutic modalities

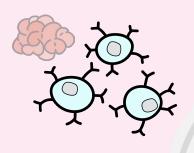


Current Achilles approach

Alternative modalities

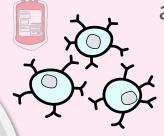
TIL-based cNeT

Sourcing cNeT from TIL - a clinically validated approach across multiple solid tumor settings



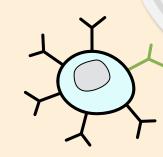
Blood-based cNeT (cNeT)

Investigational approach exploring use of blood as source of cNeT, without the need for surgery

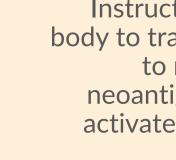


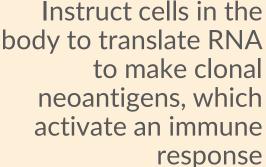
TCR-therapy for shared neoantigens

Engineered TCR therapy equips activated T cells with specific receptors that can target shared neoantigens



Clonal neoantigen vaccines (mRNA)

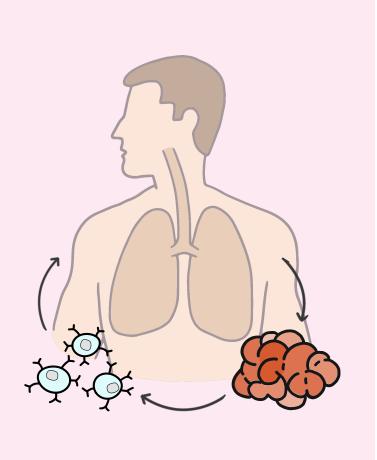




Compelling clinical data support TIL based approaches, but a significant opportunity remains



TIL: impressive clinical responses seen in multiple late-stage settings





31% ORR TIL monoTx in PD-1 refractory melanoma (n=153)¹



21% ORR TIL monoTx in PD-1 refractory NSCLC (n=28)²



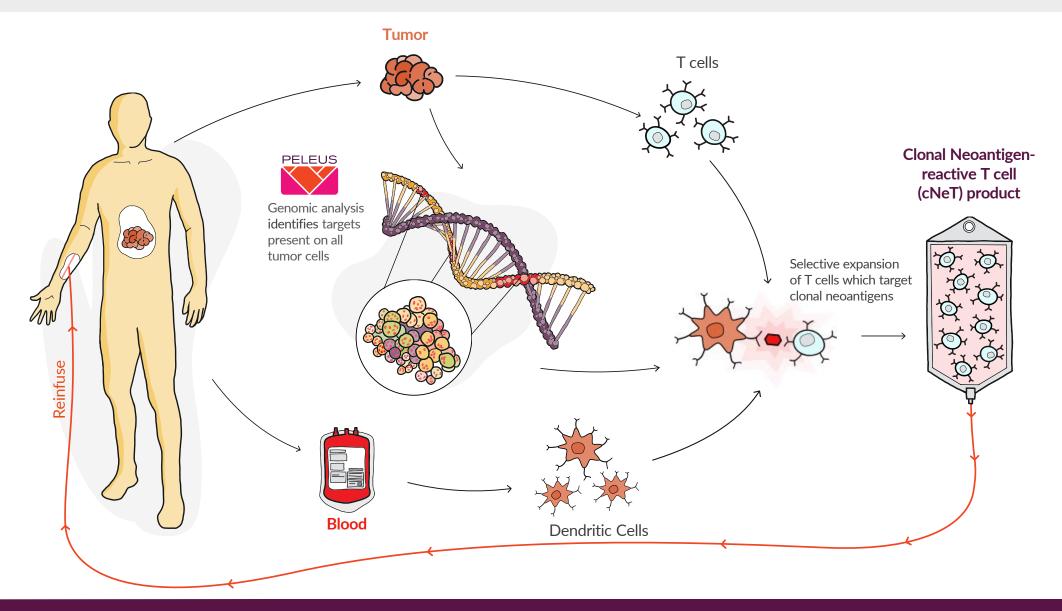
44% ORR TIL monoTx in pre-treated cervical cancer (n=27)³

- TIL therapy uses uncontrolled expansion that will enrich both subclonal and clonal neoantigen reactive T cells and can result in T cell exhaustion
- Opportunity to build on and improve traditional TIL therapy: by prospectively targeting clonal neoantigens we aim to enrich the tumor reactive component and deliver a more potent product

Achilles process delivers precision clonal neoantigen targeting T cell therapy (cNeT) Cutting edge personalized genomics and machine learning enable targeting of all cancer cells



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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

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

Ongoing in UK, Europe and US

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)

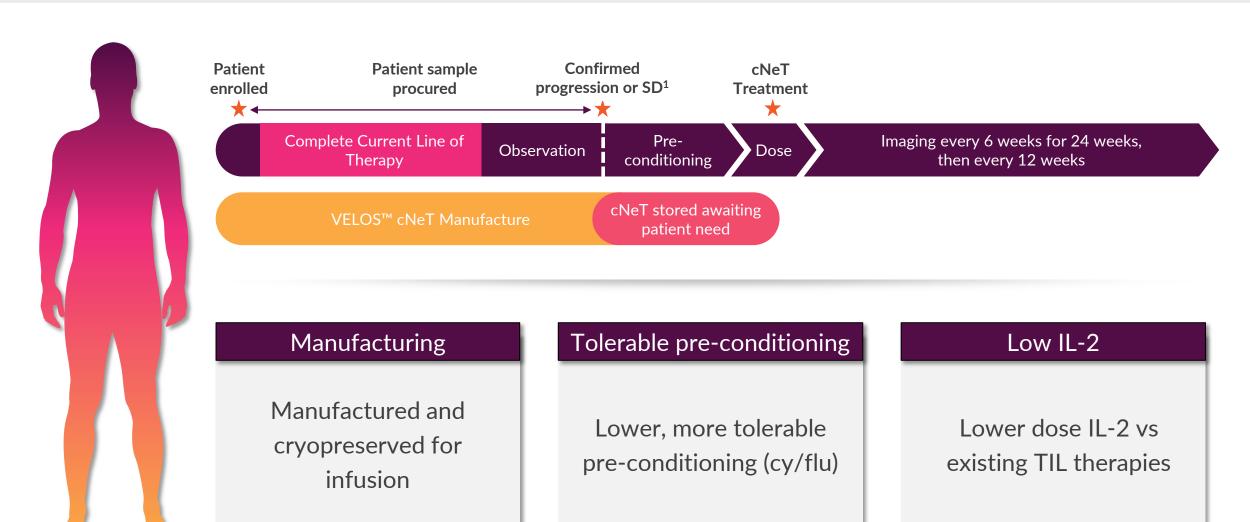
- n = up to 20 checkpoint refractory patients; Open-label
- CPI dosed 7-13 days prior to cNeT and restarted day 14 post-cNeT

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

Ongoing in UK and Europe, expanding to US

cNeT therapies will be readily delivered within standard treatment pathways





cNeT were generally well tolerated in the fourteen patients treated in CHIRON & THETIS

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Heavily pretreated patients with advanced cancer

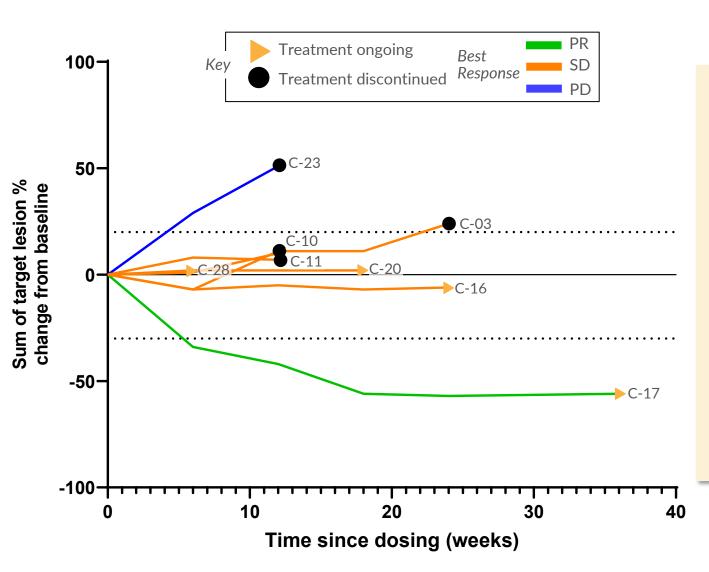
- Eight advanced unresectable or metastatic NSCLC patients (CHIRON)
- Six relapsed/refractory melanoma patients (THETIS)
- Two median lines of prior therapy, all patients refractory to checkpoint inhibitor (CPI)
- All patients had progressive disease at time of lymphodepletion
- Process improvements delivering median cNeT dose of 78M (n=3 dosed patients)

cNeT tolerability profile¹

- Tolerability similar to standard TIL
- No new cNeT-related SAEs or dose-limiting toxicities since last report (ESMO 2022)
- Lower dose lymphodepletion and lower dose IL-2 well tolerated
 - 124/130 (95%) scheduled IL-2 doses delivered
- Lymphopenia and neutropenia the most common AFs

8 CHIRON (NSCLC) patients dosed with Best Response of PR and SD¹





Early proof-of-concept demonstrated in NSCLC

- Disease control at >12 weeks observed in 5 of 7 evaluable patients (71%), including one PR (>36 weeks)
- -4 of 7 (57%) out to >18 weeks
- PR/SD with lower dose lymphodepletion and IL-2
 - Supports potential for wider applicability of cNeT, including in an ambulatory setting

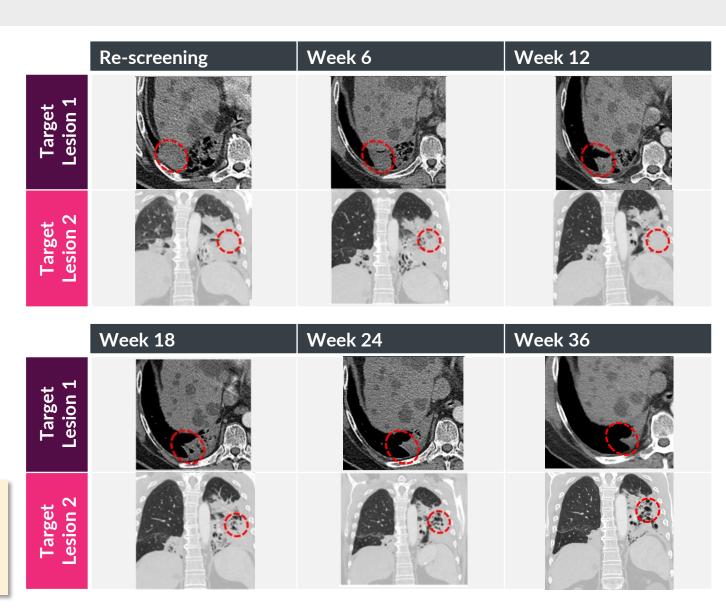
Patient C-17: 56% reduction in total target lesion size vs. baseline at week 36



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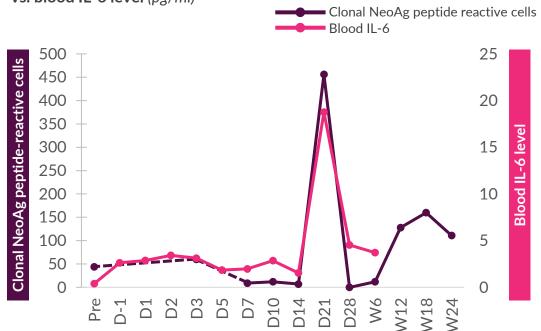
Total target lesion reduction of 56% at wk 36, with a 64% reduction in Target Lesion 2



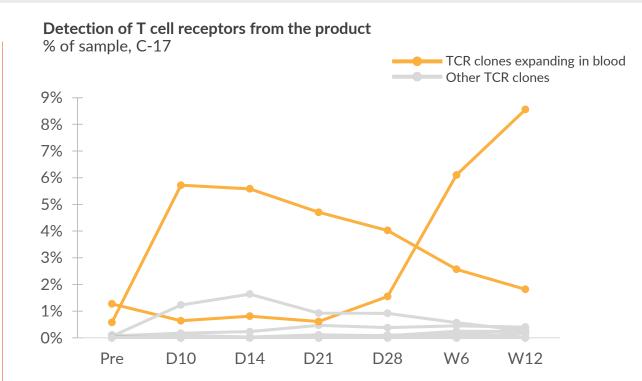
Patient C-17: cNeTs expand and persist beyond week 12 coincident with tumor regression



Clonal neoantigen peptide-reactive cells in blood (normalized spot count) vs. blood IL-6 level (pg/ml)



Clonal neoantigen reactive T cells detected in blood-post dosing, with peak at Day 21 – coincident with peak in serum cytokine associated with T cell activity (IL-6)



T cell clones that are clonal neoantigen-specific are identified expanding in the patient beyond 12 weeks and to a greater extent than other patients

Durable clinical benefit and encouraging safety and tolerability data with cNeT therapy



Lymphodepletion & IL-2 well tolerated



Lower dose lymphodepletion and IL-2 and 95% of IL-2 doses well tolerated

 Supports potential for wider applicability of cNeT, including in an ambulatory setting

Early PoC in NSCLC



- Disease control >12 weeks in 71% patients, including one PR (>36 weeks¹)
- Potential for deep, durable clinical responses with reduced lymphodepletion and II -2

cNeT Driving
Anti-tumor
Activity



- Engraftment & cytokine profiles supportive of cNeT driving anti-tumor activity
- Active cNeT peak expansion at day 21 coincides with peak in IL-6 (marker of activity)

Efficient scale-up of GMP manufacturing to align with clinical and commercial need



- Flexible manufacturing allows efficient alignment of scale-up
- GMP facilities at Royal Free Hospital in London and Catapult site in Stevenage, UK currently support global clinical trial manufacturing
- Identified and initiated tech transfer to CDMO in Philadelphia, USA, in preparation for expansion
- Design work complete for GMP modular facility to support late stage clinical and commercial supply



Focus for CHIRON and THETIS data readouts in 2023



Clinical

Dose & deliver data from 15-20 additional patients with cNeT monotherapy (lung & melanoma) and CPI combo (melanoma)

Drive additional confirmed responses in CHIRON and THETIS patients with higher cNeT doses

Translational



Leverage world-class translational science platform to define actionable cNeT features of response

Process



Continue PELEUS™ and process development to optimize dose and identify new sources of clonal neoantigens





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