Initial Data From a Phase 1, First In Human Clinical Trial For T-Plex, A Multiplexed, Enhanced T Cell Receptor-Engineered T Cell (TCR-T) Therapy For Solid Tumors

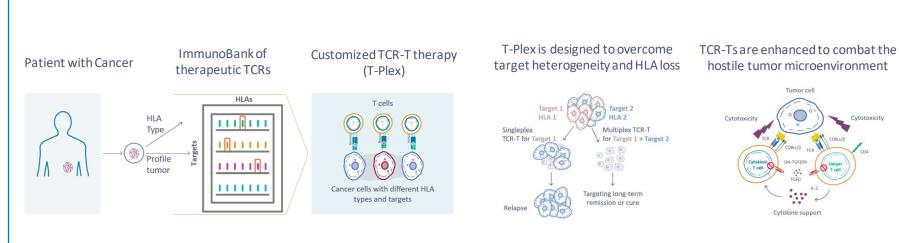
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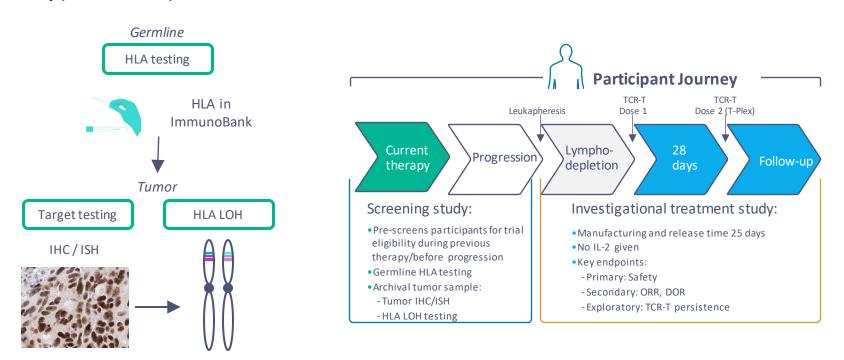
Background and Rationale

- Advanced solid tumor malignancies are difficult to cure due to complex biology.
- Lack of sufficient endogenous anti-tumor T cells hinders an immune response, even in the presence of checkpoint
- T cells engineered with exogenous T cell receptors (TCR-Ts) can target tumor specific antigens presented on human leukocyte antigens (HLAs) to kill tumor cells. However, solid tumors are notoriously heterogenous with variable target antigen expression and may also have HLA loss of heterozygosity (LOH) in up to 40% of tumors, a common resistance mechanism to avoid immune detection.
- First-generation TCR-Ts targeting single antigens had limited response rates (30-50%) and short durations of response (3-4 months), possibly due to tumor target heterogeneity and HLA LOH, an immuno-suppressive tumor microenvironment, poor TCR-T persistence and/or TCR-T exhaustion.
- Our solution is to administer multiple enhanced TCR-Ts targeting different shared antigens, commonly expressed in solid tumors, presented on diverse and frequent HLA types (T-Plex).
- The TCR-Ts are enhanced with co-delivery of CD8 α/β to engage helper T cells and a dominant negative TGF β receptor (DN-TGFβRII) to enhance T cell expansion/persistence which should enable potent tumor killing and long-term persistence.
- Product and study design details were presented at ASCO 2023 (Abstract #2554).
- Initial data from the screening study are reported here.



Prospective Identification of Target and HLA Expression is Designed to Maximize Speed Within the Participant Journey

- The screening study (NCT05812027) evaluates participants with solid tumors to determine germline HLA types, target expression and tumor HLA/LOH status.
- Pre-identification of participants allows for potential rapid enrollment into the investigational treatment study (NCT05973487).



302 participants were enrolled into the screening study across 14 sites since Sept 2023.

Pa	rticipant Demographi	cs (n=287)
Age	Median	63
	Range	21-85
Gender	Male	132 (46%)
	Female	156 (54%)
Race	White or Caucasian	230 (80%)
	Black or African American	12 (4%)
	Asian	6 (2%)
	Hispanic or Latino	3 (1%)
	American Indian or	2 (1%)
	Alaskan Native	
	Other or Unknown	34 (12%)

Demographics and Sample Testing Schema

Participants enrolled into the screening study with a variety of solid tumors.

cancer, esophageal cancer, osteosarcoma, uterine cancers and meningioma

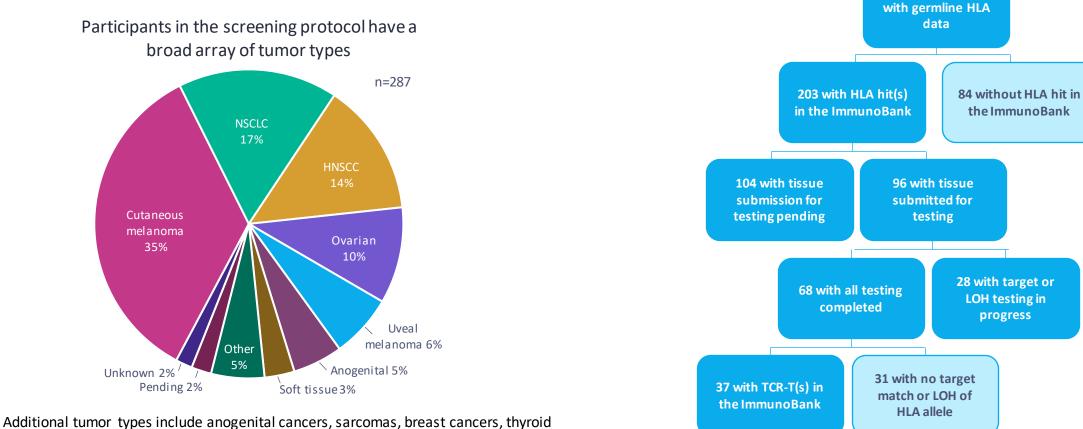
80% of participants exhibit intact HLA haplotypes

■ LOH negative ■ LOH positive

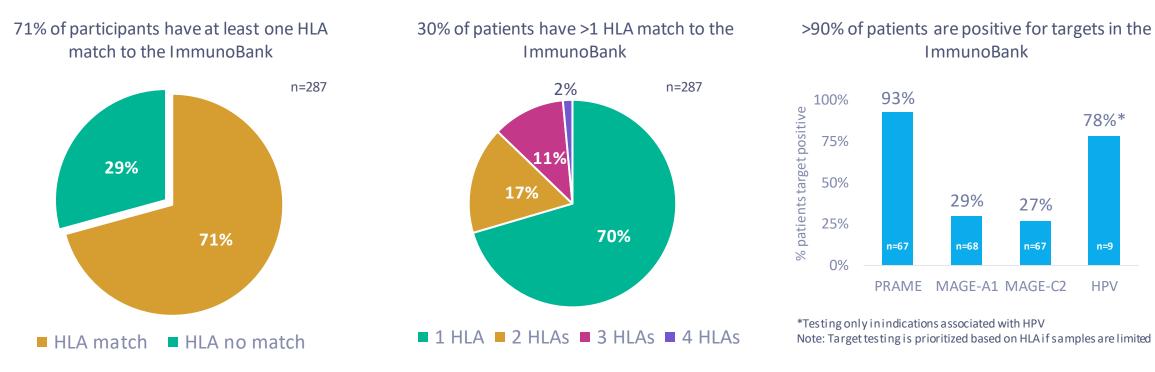
for the targetable HLA allele are not submitted for LOH testing

*Samples that are negative for target/HLA match or are homozygous

- Germline HLA data were available for 287 participants as of the 8 May 2024 data cut.
- Melanoma, non-small cell lung cancer, head and neck cancer, and ovarian cancer account for >75% of tumor types.



The Majority of Participants Have At Least One TCR-T Match in the ImmunoBank, and Many Have Multiple TCR-T Matches that Could **Qualify for T-Plex**



>50% of participants with all testing completed

All testing Any TCR-T

complete match

in cancer cells have at least one TCR-T in the ImmunoBank n=55* 80%

qualify for T-Plex n=37 22% 73% Singleplex ■ 2-T-Plex Combo ■ 3-T-Plex Combo

27% of participants with TCR-T match that could

287 participants

Key Inclusion/Exclusion Criteria and Specifications for Study NCT05973487

Inclusion Criteria

- Age ≥ 18, all genders
- Diagnosed advanced melanoma, non-small cell lung cancer, head and neck cancer, ovarian cancer, cervical or anogenital cancers, or other solid tumors that may express relevant targets
- Failure of standard of care, including any relevant targeted therapy or checkpoint inhibitor
- ≥ 1 HLA with a TCR-T match and no LOH of that HLA in the tumor
- ≥1 target match
- ECOG PS 0-1 with adequate organ function
- ≥ 1 measurable lesion per modified RECIST v1.1

Exclusion Criteria

- · CNS metastases that are symptomatic or in need of
- treatment; carcinomatous meningitis
- months of enrollment
- Systemic corticosteroids within seven days of enrollment
- Concurrent therapy that has not yet met washout

Major cardiac pathology or stroke/TIA within 12

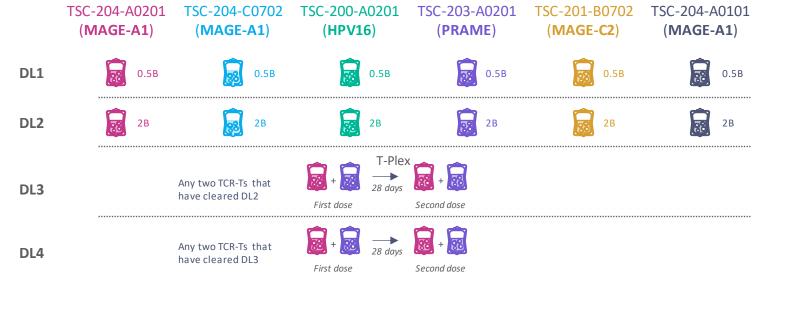
- requirements
- Hypersensitivity to fludarabine, cyclophosphamide, or study drug components
- Presence of an HLA type that may interfere with the HLA type being targeted

Protocol Specifications

- Lymphodepletion: Cyclophosphamide (60 mg/kg on Days -7 and -6); fludarabine (30 mg/m2 on Days -7 to -3)
- Hospitalization of first three singleplex and first three T-Plex patients for three to seven days post cell infusion

Dose Escalation Scheme Provides a Path to Multiplex TCR-T in the Phase 1 Study

- TCR-Ts currently in the master protocol target PRAME on HLA-A*02:01; HPV16 on HLA-A*02:01; MAGE-A1 on HLA-A*02:01, HLA-A*01:01, or HLA-C*07:02; or MAGE-C2 on HLA-B*07:02.
- All TCR-Ts in the ImmunoBank and master protocol are first tested as singleplex therapies in dose levels 1 and 2 before becoming available for multiplexing in dose levels 3 and 4.



Conclusions and Acknowledgements

- The screening study is underway and widely adopted, with over 300 participants enrolled.
- >50% of participants with all testing completed (n=68) have at least one TCR-T in the ImmunoBank and 27% of those could qualify for T-Plex.
- Potential participants have been pre-identified across all six singleplex cohorts, as well as some T-Plex cohorts (n=37 and n=10, respectively).
- Eligibility for T-Plex is expected to increase as the ImmunoBank grows via additional IND applications.
- The **first participant** (with metastatic cutaneous melanoma) was dosed with TSC-203-A0201 targeting PRAME; no acute toxicities were observed.
- Manufacturing for three additional participants is underway.

We extend a warm thank you to the participants, families and treatment teams associated with the screening and investigational treatment studies.

For more information, please contact: clinicaltrials@tscan.com