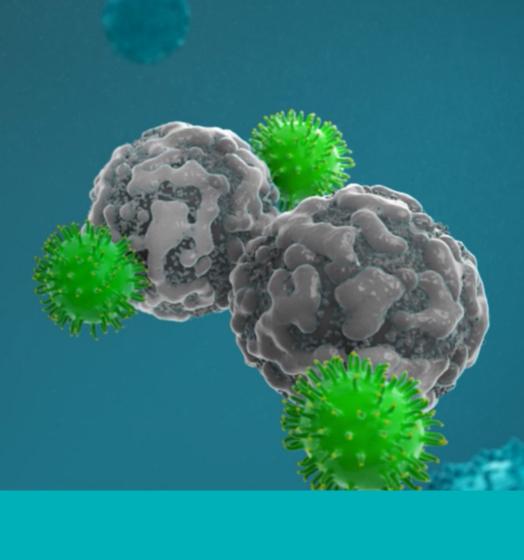
Prospective BASECAMP-1 experience in patients with gastrointestinal (GI) cancer: Identifying patients with human leukocyte antigen (HLA) loss of heterozygosity (LOH) for a future therapeutic trial exploiting LOH as a tumor vulnerability



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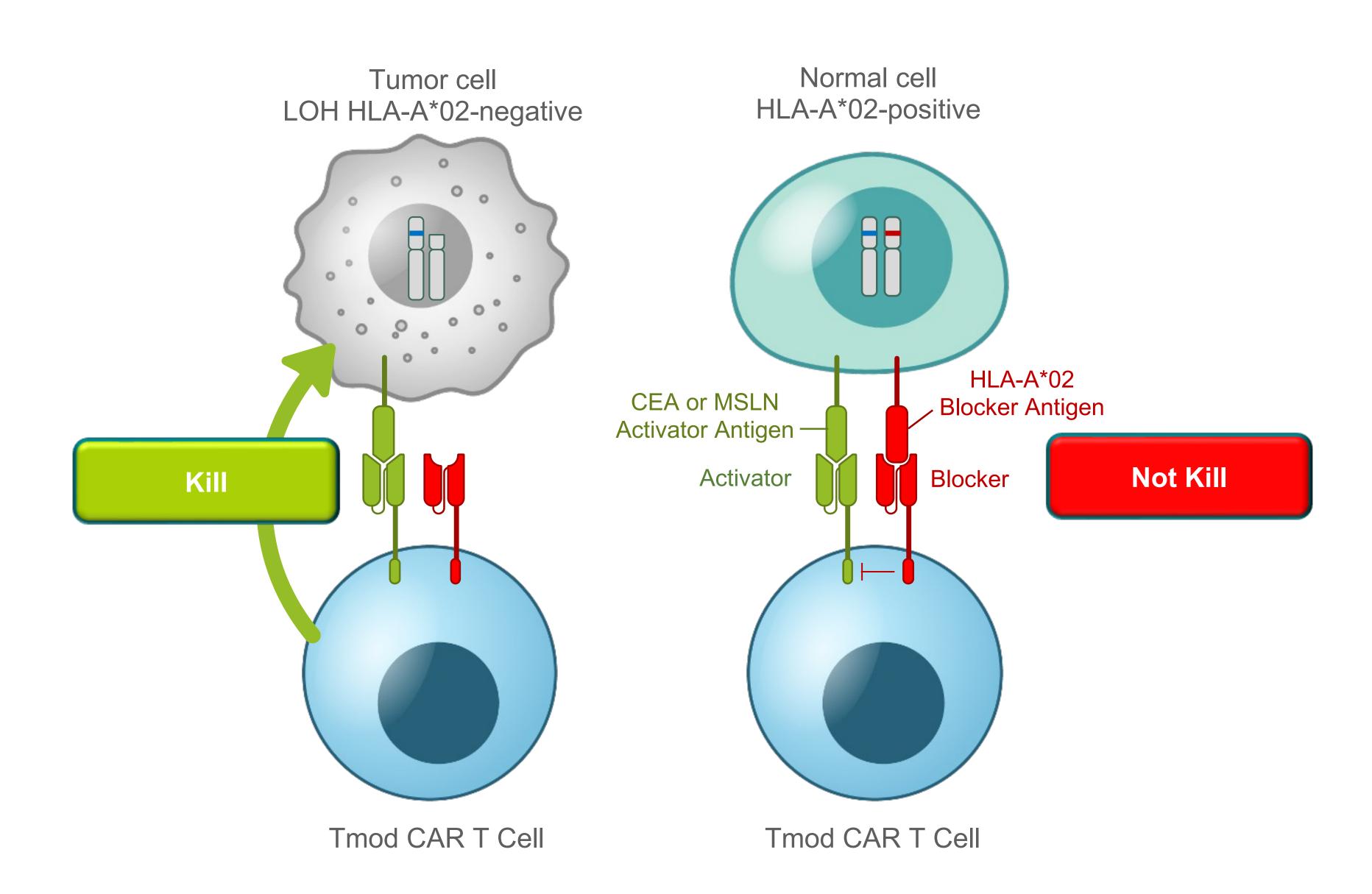
BACKGROUND AND STUDY OBJECTIVES

- Metastatic colorectal, pancreatic (PANC), gastric, and esophageal cancers are the leading causes of gastrointestinal (GI) cancerrelated mortality (metastatic 5-year survival rate of 15%, 3%, 6% and 5% respectively) [1]
- Chimeric antigen receptor (CAR) T-cell therapy has demonstrated clinical outcomes in hematologic malignancies [2,3]. However, translating engineered T-cell therapies to solid tumors proves difficult due to a lack of tumor-specific targets that distinguish cancer cells from normal cells. In previous studies, the use of carcinoembryonic antigen 5 (CEA) T-cell receptors and mesothelin (MSLN) CARs both resulted in dose-limiting on-target, off-tumor toxicities [4-6]
- Tmod™ CAR T-cell is a logic-gated cell therapy which addresses these challenges by leveraging dual receptors capable of killing tumor cells, while leaving healthy cells intact [7]. Tmod platform technology is a versatile system that may be applied to T cells and natural killer cells in autologous and allogeneic settings
- A2B530 is a CEA-directed and A2B694 is a MSLN-directed Tmod construct utilizing a LIR-1-based inhibitory receptor (blocker) targeting human leukocyte antigen (HLA)-A*02
- HLA loss of heterozygosity (LOH) may provide a means to distinguish between tumor from normal tissue in a definitive manner due to this irreversible, clonal loss within tumor cells [7,8]. The 2 receptors of the Tmod CAR T-cell platform comprise an activator that recognizes an antigen present on the surface of normal and tumor cells and a blocker that recognizes a second surface antigen from an HLA allele lost only in tumor cells
- In the Tempus real-world database, LOH occurs in 12.2% to 26.0% of advanced solid tumors with an average of 16.3% in 10,867
- HLA-A*02 LOH can only be therapeutically exploited if patients are identifiable through a feasible and timely clinical workflow. The Tempus xT is a next-generation sequencing (NGS) platform commonly used for patients with GI cancer which can readily
- BASECAMP-1 (NCT04981119) is an ongoing study with key objectives: 1) To determine and identify patients with somatic HLA LOH eligible for Tmod CAR T-cell therapy; and 2) Subsequent leukapheresis and manufacturing feasibility for future Tmod CAR
- Eligible patients identified in BASECAMP-1 will be referred to the EVEREST A2B530 CEA Tmod or A2B694 MSLN Tmod interventional studies

STUDY RATIONALE

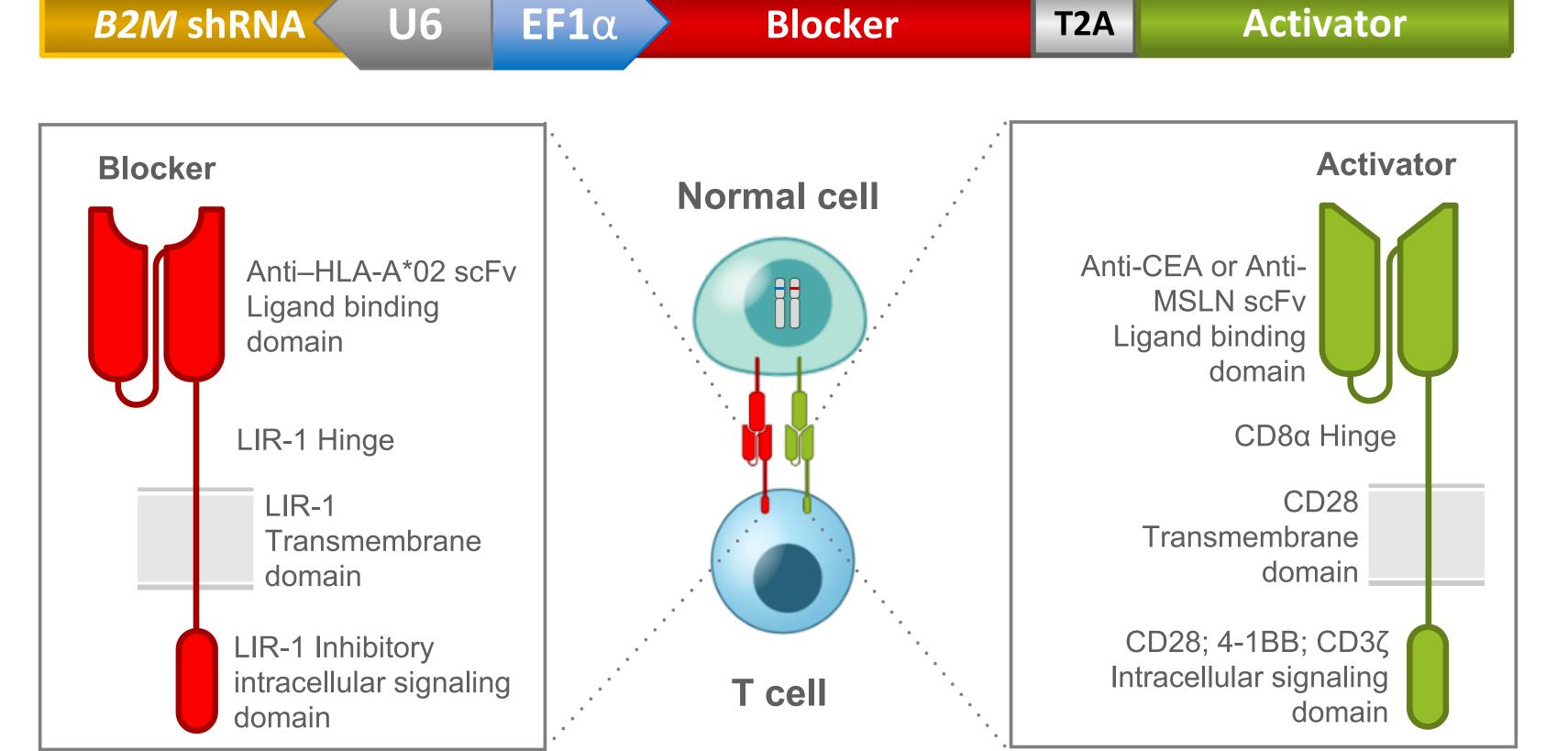
identify HLA LOH

Figure 1. Logic-gated CAR T with the goal of reducing toxicity: CEA or MSLN (activators) and HLA-A*02 (blocker)



- CAR, chimeric antigen receptor; CEA, carcinoembryonic antigen 5; HLA, human leukocyte antigen; LOH, loss of heterozygosity; MSLN, mesothelin.
- A2 Bio's Tmod CAR T HLA LOH approach has been published by Hamburger et al 2020 (Figure 1) [7]
- HLA was selected as blocker target; first blocker HLA-A*02 is the most prevalent allele in the US population Activators include CEA and MSLN, which are both well-studied targets but have dose-limiting toxicities in previous studies
- CAR T HLA-A LOH approach is independently validated by Vogelstein/Kinzler, 2021 [8]

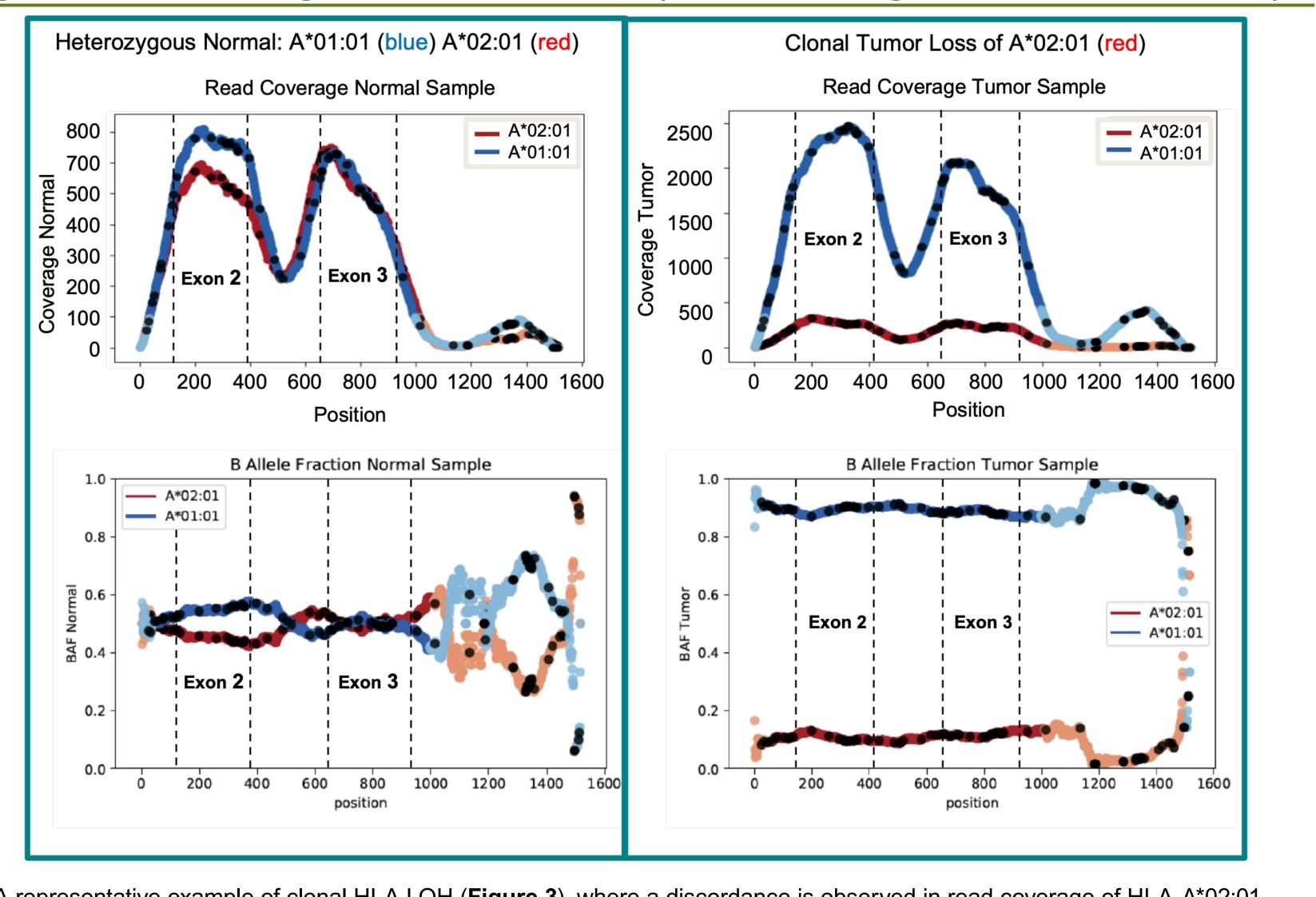
STUDY RATIONALE (cont.)



B2M shRNA. β2-microglobulin short-hairpin RNA; CEA, carcinoembryonic antigen 5; EF1α, elongation factor-1α; HLA, human leukocyte antigen; LIR, leukocyte immunoglobulin-like recepto MSLN, mesothelin; scFv, single-chain variable fragment; T2A, thosea asigna virus 2A.

- CAR activator: 3rd-generation CAR T with both signal 1 (CD3ζ) and signal 2 activation domains (CD28 & 4-1BB) CAR blocker: LIR-1 is a member of the immune inhibitory receptor family and contains 4 immunoreceptor tyrosine-based inhibition motifs in its signaling domain [10]
- Replicant incompetent single lentivirus transgene: The activator and blocker receptors are co-expressed in a single construct containing a cleavable T2A linker (Figure 2)

Figure 3. Read coverage and B allele fraction (ratio of coverage for allele 1 and allele 2)



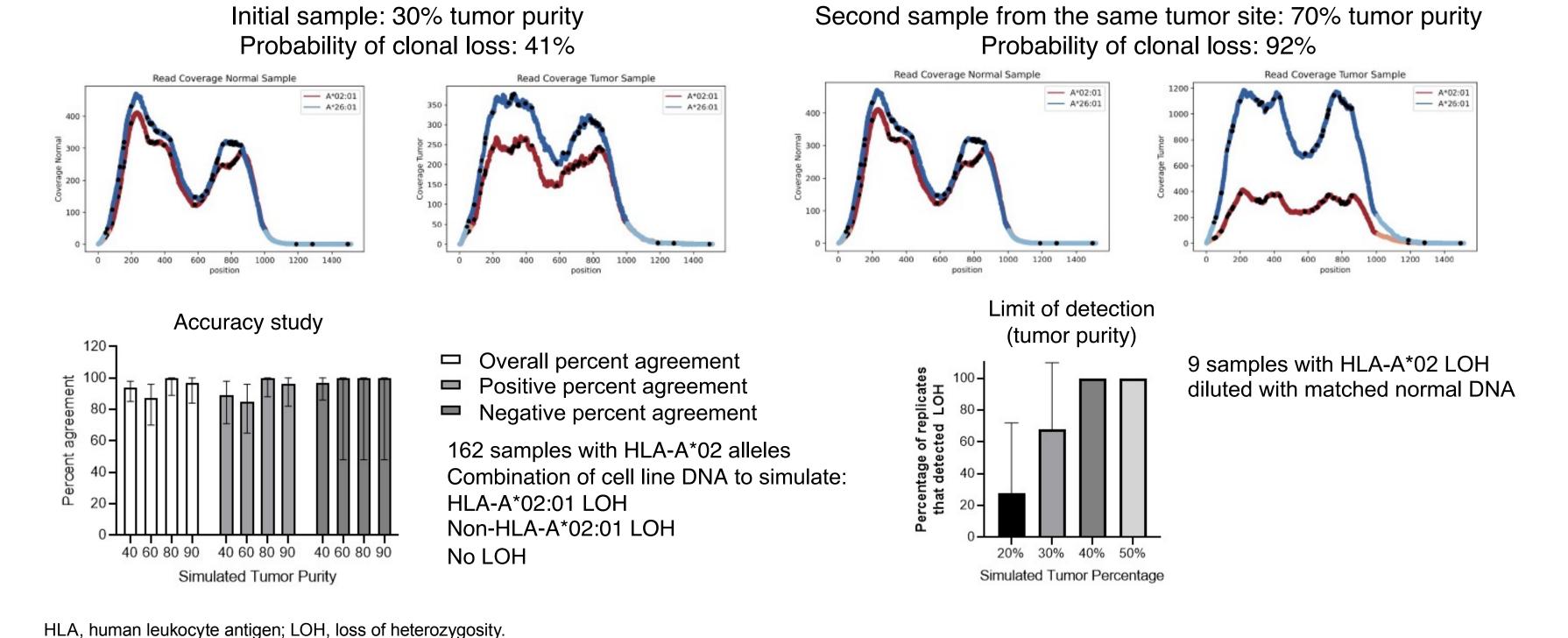
- A representative example of clonal HLA LOH (Figure 3), where a discordance is observed in read coverage of HLA-A*02:01 between the tumor and matched-normal samples [9, 11]
- HLA-A*02 LOH can be reliably detected using the Tempus xT next-generation sequencing (NGS) assay (Table 1)

Table 1. Frequency of HLA-A LOH in advanced GI tumors^a

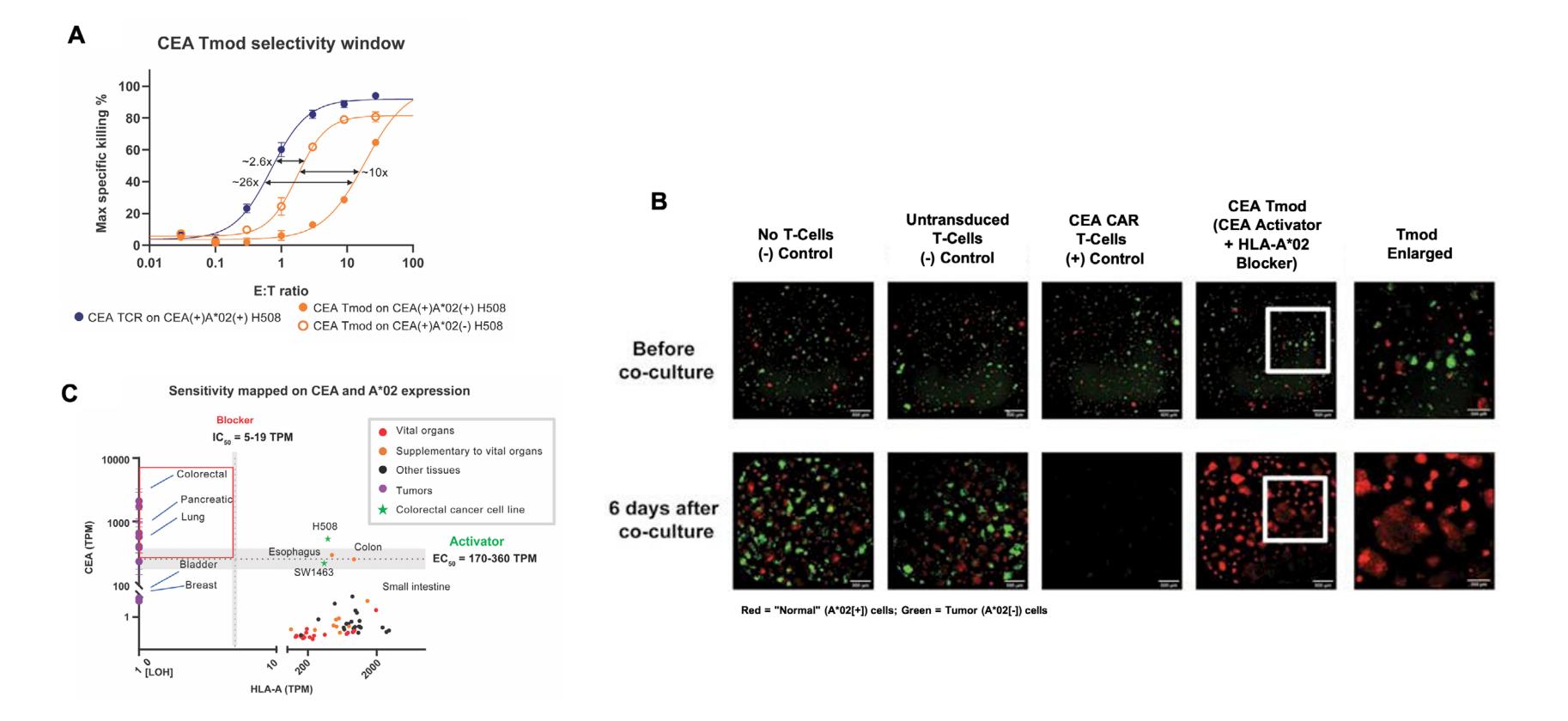
	Tempus HLA-A LOH advanced disease real-world [9]	TCGA HLA-A LOH primary tumors [12]
Average	16.3% (n=10,867)	12.6% (n=10,844)
Range in GI cancers	15.6%-20.8% (n=3,035)	9.6%-33.1% (n=1,424)
Colorectal	15.6% (n=1,854)	9.6% (n=615)
Pancreatic	19.6% (n=675)	33.1% (n=184)
Gastroesophageal	20.8% (n=506)	16.2% (n=625)

HLA, human leukocyte antigen; LOH, loss of heterozygosity; TCGA, The Cancer Genome Atlas. ^aTempus data contain more advanced disease, and TCGA data have more primary tumors.

Figure 4. Higher tumor purity allows for more accurate prediction of HLA-A*02 LOH



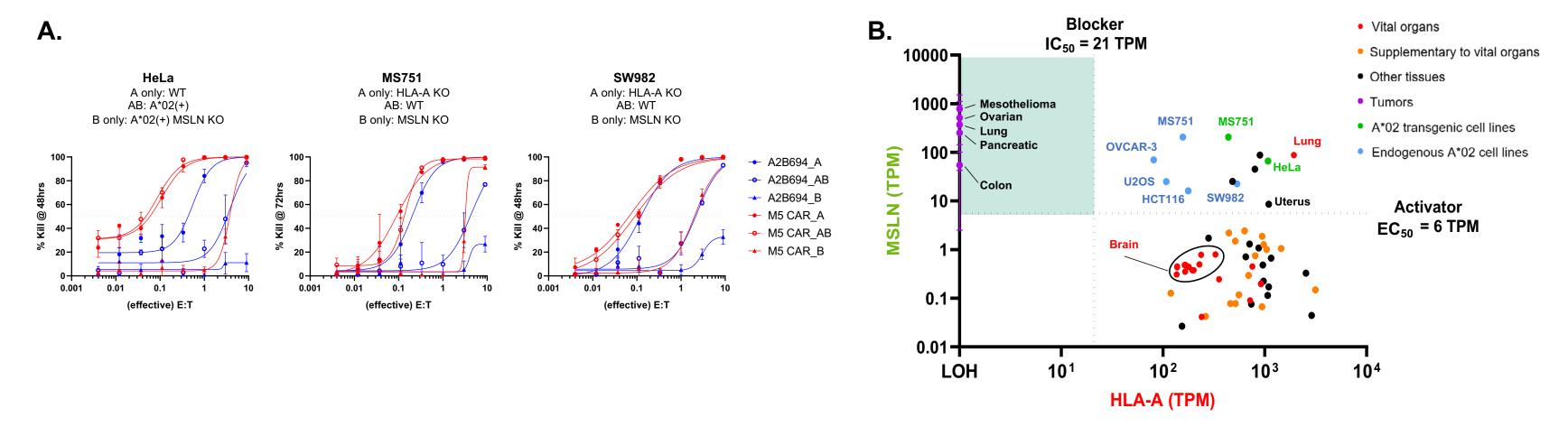
igure 5. CEA Tmod (A2B530) in vitro study provides a therapeutic safety window comparable to NCI benchmark CEA TCR-T [4,13]



CAR. chimeric antigen receptor: CEA. carcinoembryonic antigen 5; EC₅₀, half maximal effective concentration; E:T, effector-to-target; HLA, human leukocyte antigen; IC₅₀, half maximal inhibitor concentration; NCI, National Cancer Institute; TCR, T-cell receptor; TPM, total particulate matter. Tmod provided selectivity at varying effector-to-target (E:T) ratios with "normal" CEA(+)A*02(+) cells and tumor CEA(+)A*02(-)

- colon cancer cell lines (Figure 5A) Mixed A*02(+) and A*02(-) cell cultures show Tmod's ability to discriminate between "normal" (A*02[+]) and tumor (A*02[-]) cells
- (Figure 5B) CEA and HLA-A*02 standard plots were generated using CEA surface expression data from mRNA data (Figure 5C)
- CEA Tmod Jurkat or T-cell effective concentration and inhibitory concentration were graphed with the tumor and normal expression values for the CEA and A*02 antigens, along with multiple cell lines

igure 6. MSLN Tmod (A2B694) in vitro study provides a therapeutic safety window comparable to M5 benchmark MSLN CAR-T [14]

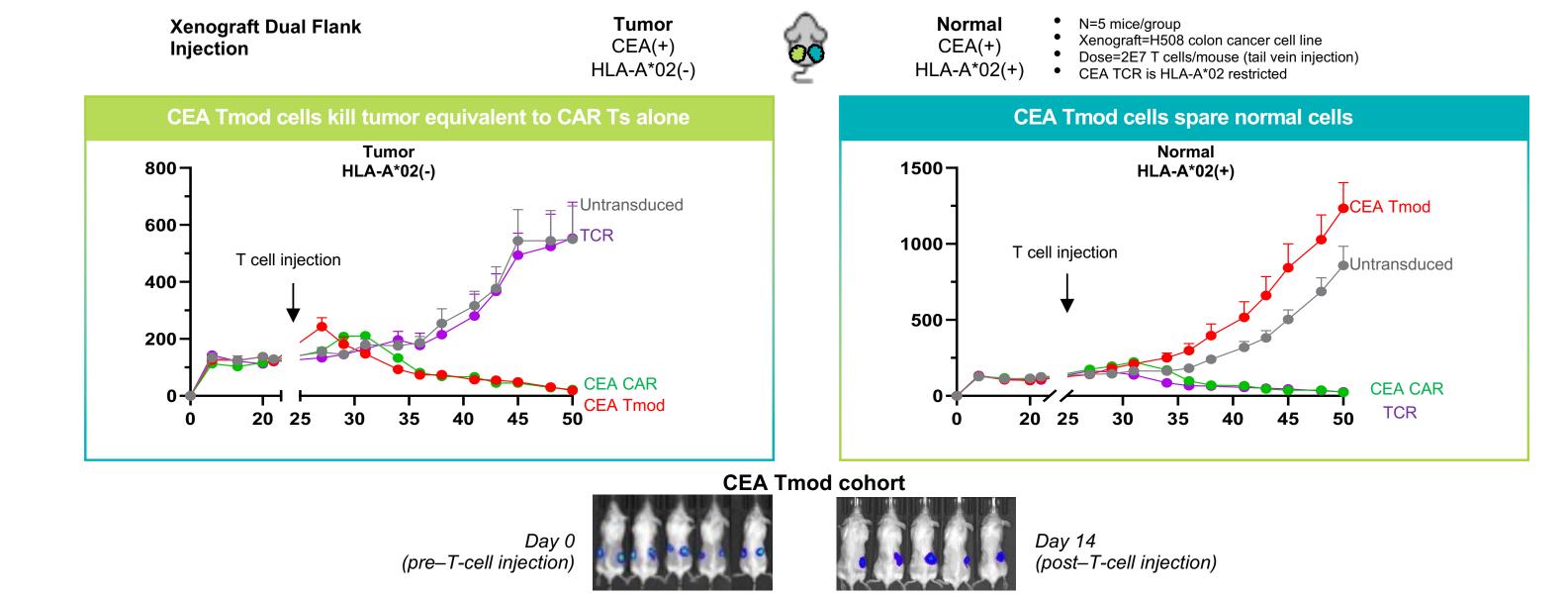


(A) Cytotoxicity data of A2B694 constructs from a representative HLA-A*02(+) donor against three endogenous MSLN(+) cell lines with native or engineered expression of HLA-A*02. M5 CAR expressing cells are shown for comparison.

(B) MSLN Tmod are predicted to kill tumors while protecting normal tissues. Data representation requires standard curves (not shown) that relate surface protein level to RNA-Seg value for cell lines and tissues. Purple points represent tumor types with HLA-A expression set at 0 TPM to account for selection of A*02(-) tumors by LOH. Tumor data from TCGA database: normal tissue data from the GTEx database.

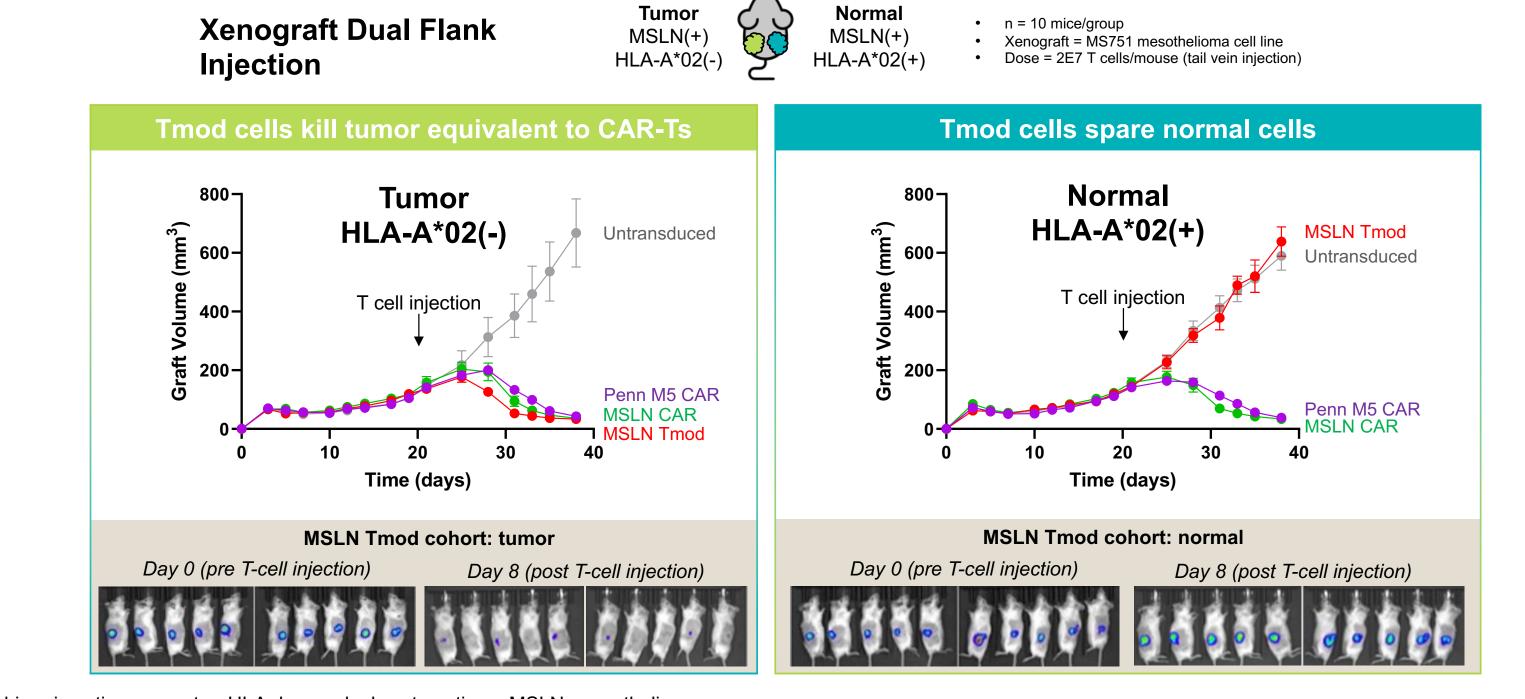
CAR, chimeric antigen receptor; EC₅₀, half maximal effective concentration; GTEx, Genotype-Tissue Expression; HLA, human leukocyte antigen; IC₅₀, half maximal inhibitory concentration MSLN, mesothelin; TCGA, the Cancer Genome Atlas; TPM, total particulate matter.

Figure 7. CEA Tmod (A2B530) in vivo study demonstrates potency comparable to NCI Figure 10. Tempus clinical workflow benchmark CEA TCR-T [4,13]



CAR, chimeric antigen receptor; CEA, carcinoembryonic antigen; HLA, human leukocyte antigen; NCI, National Cancer Institute; TCR, T-cell receptor.

Figure 8. MSLN Tmod (A2B694) in vivo study demonstrates potency comparable to M5 benchmark MSLN CAR-T [14]



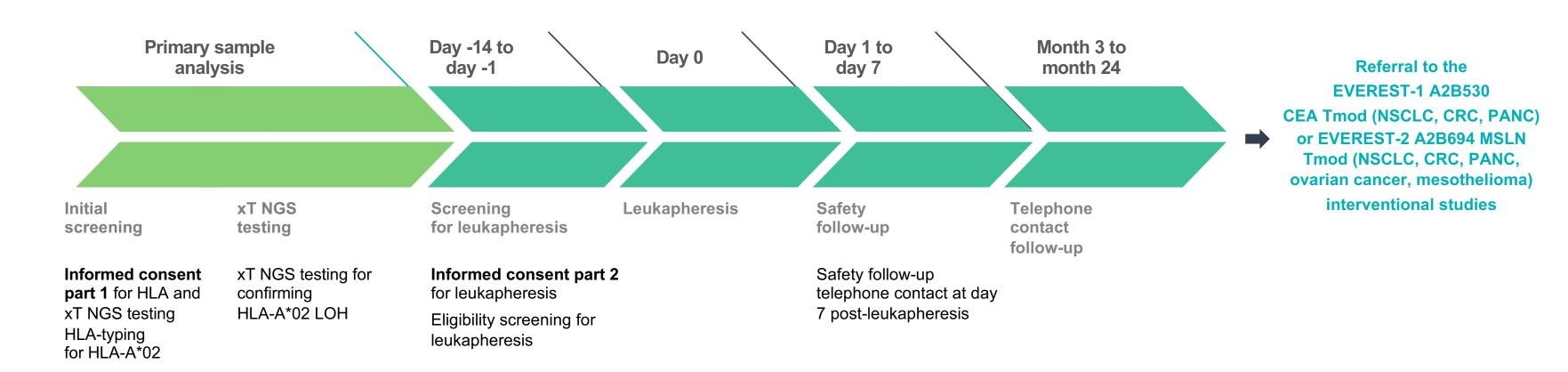
In vivo studies show that Tmod maintains selectivity

for relapsed status

- Tumor (HLA-A*02[-]) and "normal" (HLA-A*02[+]) cells were implanted subcutaneously in NOD scid gamma (NSG) mice CAR T cells or Tmod CAR T cells were administered via tail veins when tumor reached 100-150 mm³ Approximately 2 weeks following cell infusion, Tmod CAR T-cell treated mice (shown in red) experienced selective regression of tumor
- grafts while "normal" tumor grafts continued to grow. Mice treated with CEA or MSLN CAR T cells (shown in green) experienced regressions of both tumor and "normal" tumor grafts (Figures 7 and 8)

STUDY DESIGN AND METHODS

Figure 9. Study Schema for BASECAMP-1 (NCT04981119)



CEA, carcinoembryonic antigen 5; CRC, colorectal cancer; HLA, human leukocyte antigen; LOH, loss of heterozygosity; MSLN, mesothelin; NGS, next-generation sequencing; NSCLC, non-small cell lung cancer; PANC, pancreatic cancer.

- Participants will be initially screened to identify germline HLA-A*02 heterozygosity by central NGS. If HLA-A*02 heterozygosity is confirmed, tumor tissue will be analyzed by xT NGS testing to determine if somatic tumor HLA-A*02 LOH is present (Figures 9 and 10)
- If the tumor demonstrates HLA-A*02 LOH and the participant screens eligible, the participant will undergo leukapheresis Participants enrolled in the study who undergo leukapheresis will be evaluated for safety 7 days after leukapheresis and followed

Banked T cells will be available for subsequent autologous Tmod CAR T-cell therapy at the time of relapse

STUDY DESIGN AND METHODS (cont.)

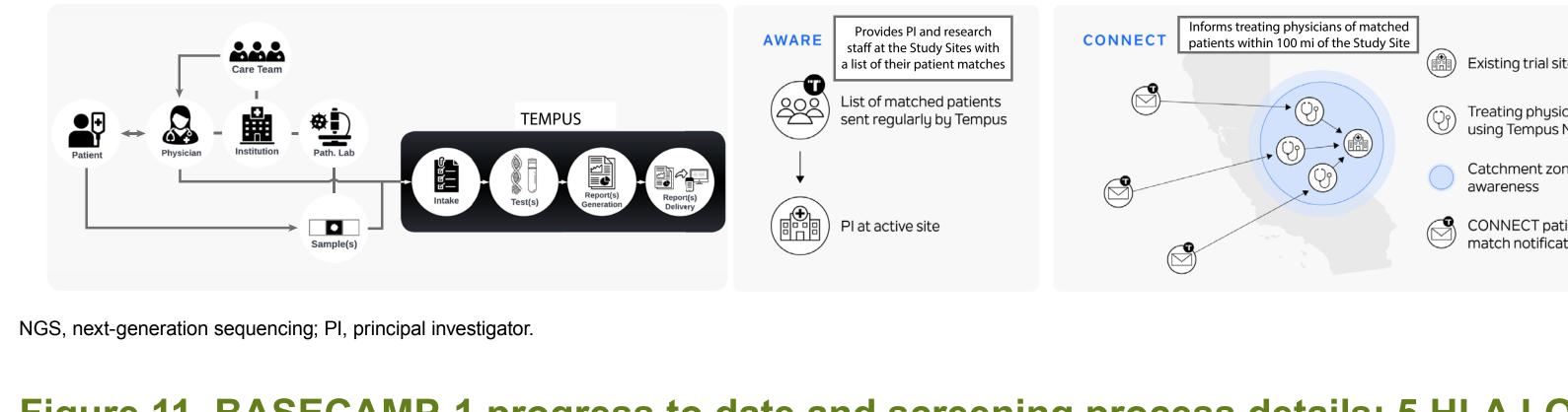
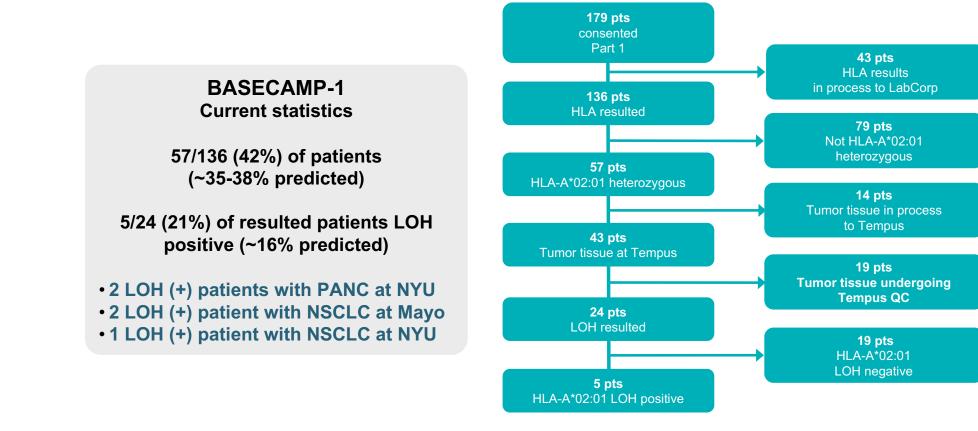


Figure 11. BASECAMP-1 progress to date and screening process details: 5 HLA LOH patients identified (Updated data cut on December 4, 2022)



HLA, human leukocyte antigen; LOH, loss of heterozygosity; NSCLC, non-small cell lung cancer; PANC, pancreatic cancer; QC, quality control.

CONCLUSIONS

- BASECAMP-1 prospective identification of HLA-A*02 LOH is feasible in the real-world setting
- Clonal HLA LOH is an irreversible discriminator between tumor vs normal cells that can be exploited for AND-NOT logic-gated Tmod CAR T to reduce on-target off-tumor toxicity [7,8]
- The frequency of HLA LOH among advanced GI solid tumor cancers in Tempus real-world dataset is 16.3%, with a range of 15.6%-20.8% between colorectal, pancreatic, and gastroesophageal tumors [9]
- Tempus xT platform is able to identify patients with clonal HLA LOF
- LOH results can be obtained within a clinically feasible workflow and timeframe, although samples with a <40% tumor purity have a reduced sensitivity for LOH detection, an issue recurrently observed in patients with PANC
- BASECAMP-1 (NCT04981119) study is currently enrolling patients to identify HLA-A*02 LOH subjects with colorectal, pancreatic,
- non-small cell lung cancer, mesothelioma or ovarian cancer and then to bank their T cells for future EVEREST A2B530 CEA Tmod or A2B694 MSLN Tmod interventional studies

SITE LIST

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- Mavo Clinic. Rochester Principal Investigator: Julian Molina, MD, PhD Sub-Investigator: Yi Lin. MD. PhD Sub-Investigator: Caleb Smith, MD
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Locke F, et al. N Engl J Med. 2022;386(7):640-654

Maude S, et al. N Engl J Med. 2018;378(5):439-446

0. Borges L, et al. *J Immunol*. 1997;159(11):5192-5196.

13. Sandberg M, et al. *Sci Transl Med.* 2022;14(634).

Hamburger A. et al. *Mol Immunol*, 2020:128:298-310.

Hecht J, et al. *J Clin Oncol.* 2022; 40(4 suppl):190-190.

14. Tokatlian T, et al. J Immunother Cancer. 2022;10:e003826.

Perera J, et al. *J Immunother Cancer.* 2019;7(suppl 1):P103.

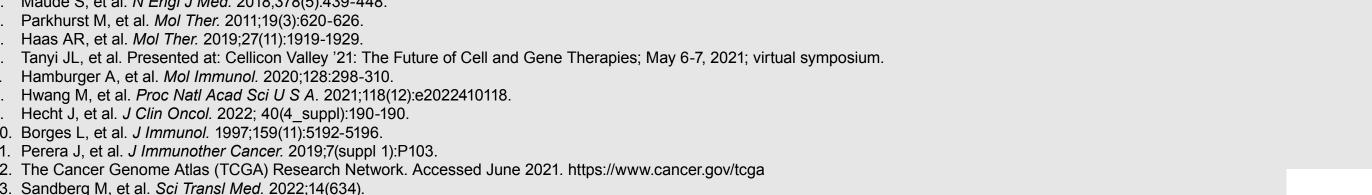
Parkhurst M, et al. Mol Ther. 2011;19(3):620-626

Haas AR, et al. *Mol Ther.* 2019;27(11):1919-1929.

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