



AFNT-111, a preclinically safe and effective
TCR-engineered T cell therapy targeting the
oncogenic driver KRAS G12V mutation
(Poster #244)

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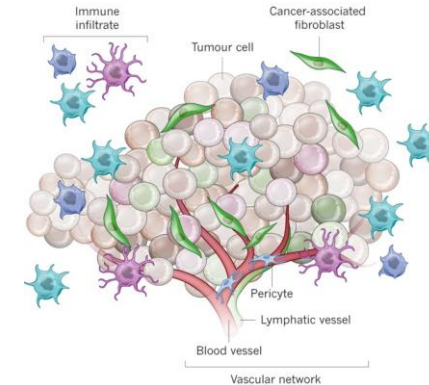
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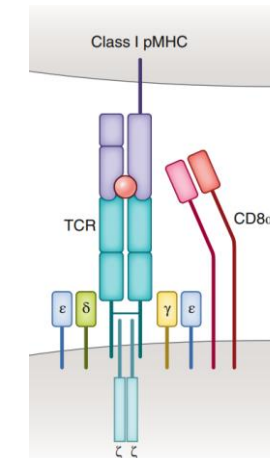
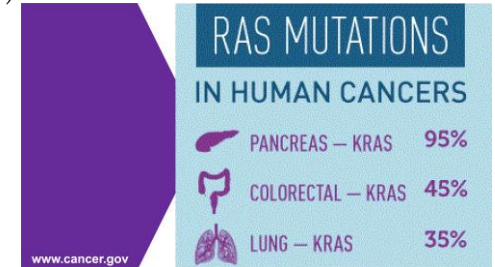
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- Oncogenic driver mutations responsible for the initiation and maintenance of cancer are ideal targets
 - Addresses tumor heterogeneity (cancer dependency on oncodrivers)
 - Neoantigens are unique to cancer and limit on-target/off-tumor tox
- KRAS is the most frequently mutated oncogene in solid tumors
 - KRAS mutations (mKRAS) promote cell growth and suppress apoptosis
 - RAS family of genes are responsible for up to 30% of all human cancers
- TCR-T cell therapies are an optimal modality to attack mKRAS
 - Able to target intracellular KRAS presented on the cancer cell surface
 - Clinical proof of concept for targeting KRAS demonstrated in multiple T cell therapy clinical studies^{1,2}



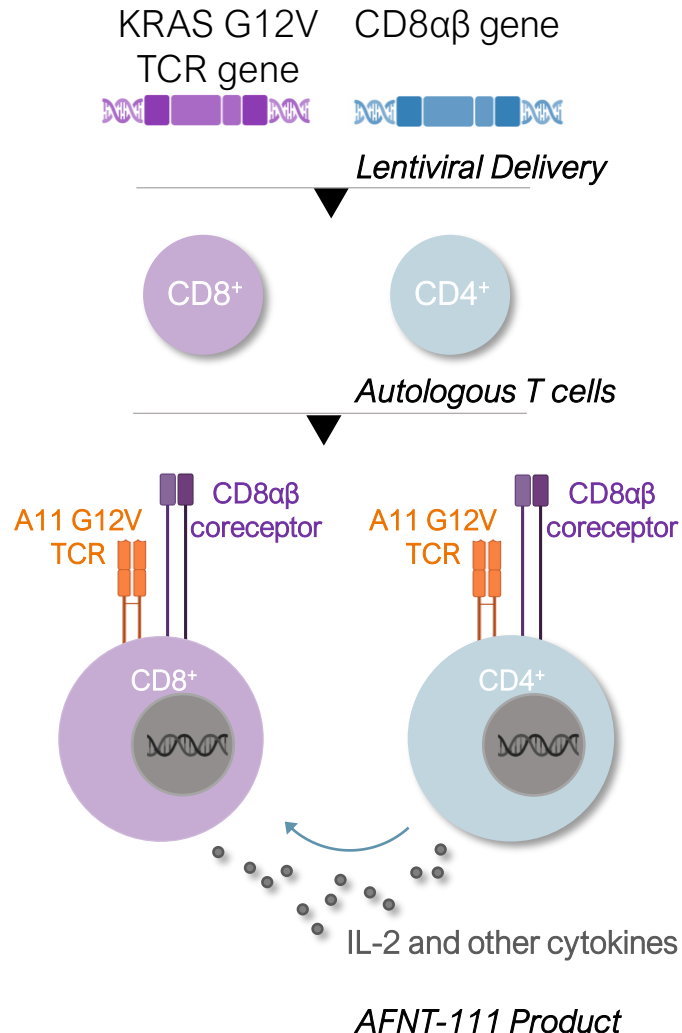
From Joglekar AV and Li G, Nat Methods (2021)

Tackle a high unmet medical need



Utilize a therapeutic modality with clinical PoC

¹Tran et al. NEJM (2016), ²Leidner et al. NEJM (2022)

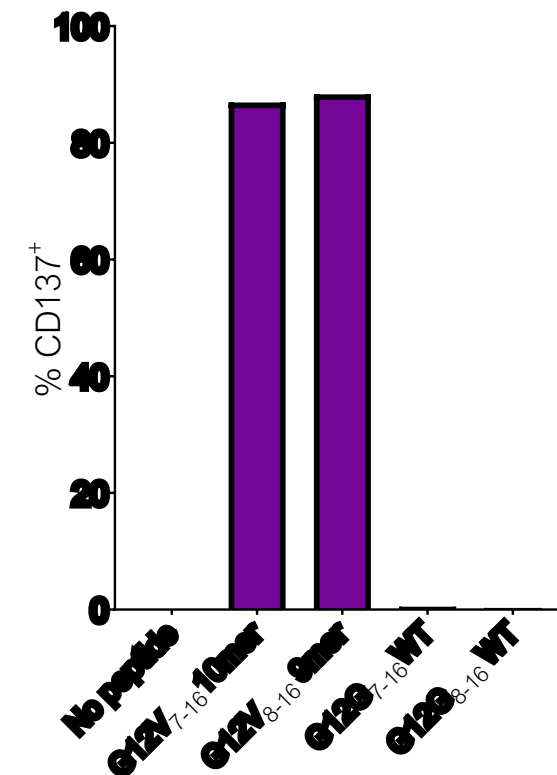
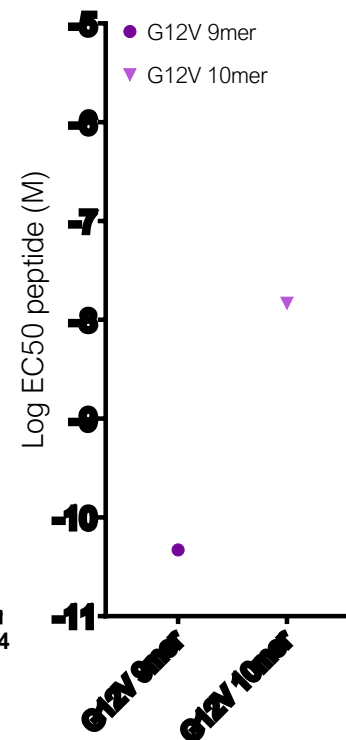
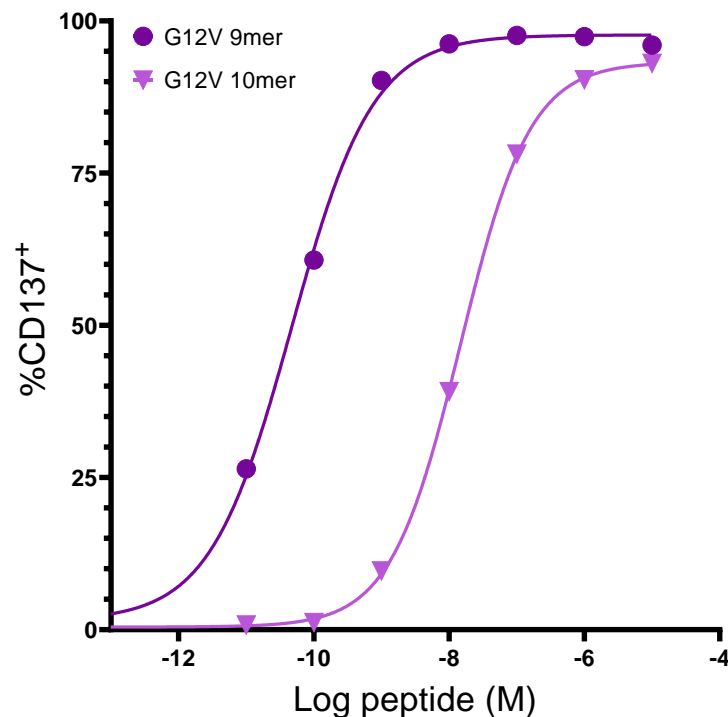
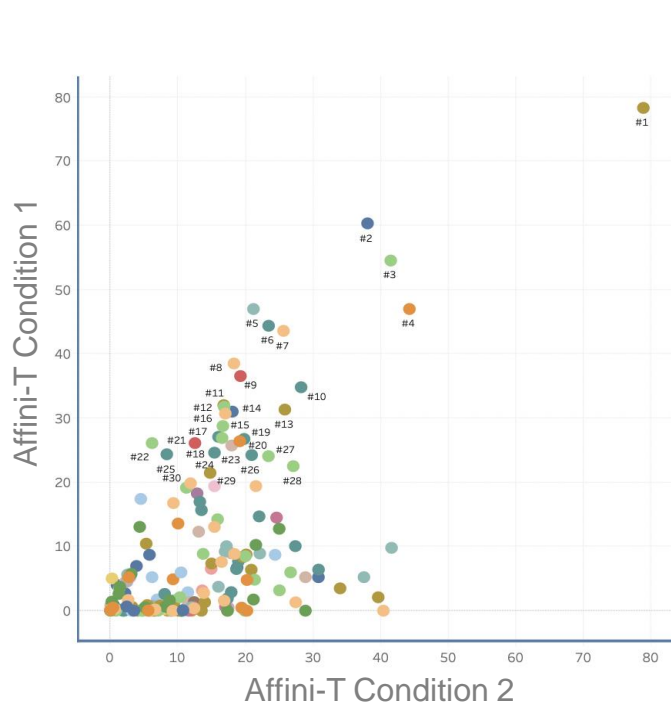


I. TCR Therapy

- Robust TCR Discovery platform identifies high affinity, high functional avidity TCRs from the natural TCR repertoire of healthy donors (no affinity maturation)
- AFNT-111: Autologous CD4⁺ and CD8⁺ T cells lentivirally-transduced with CD8αβ co-receptor and high affinity TCR to KRAS G12V presented by HLA-A*11:01

II. Coordinated CD8⁺ and CD4⁺ T Cell Response

- CD8α/β co-receptor expression in CD4⁺ T cells enables MHC I target recognition and a coordinated CD8⁺/CD4⁺ T cell response
- CD4⁺ T cells retain a helper phenotype, engage CD8⁺ T cells in the TME, promote effector activation, and prevent T cell exhaustion leading to enhanced CD8⁺ T cell persistence



- High affinity KRAS G12V TCR candidates identified from healthy donors that have cleared thymic selection

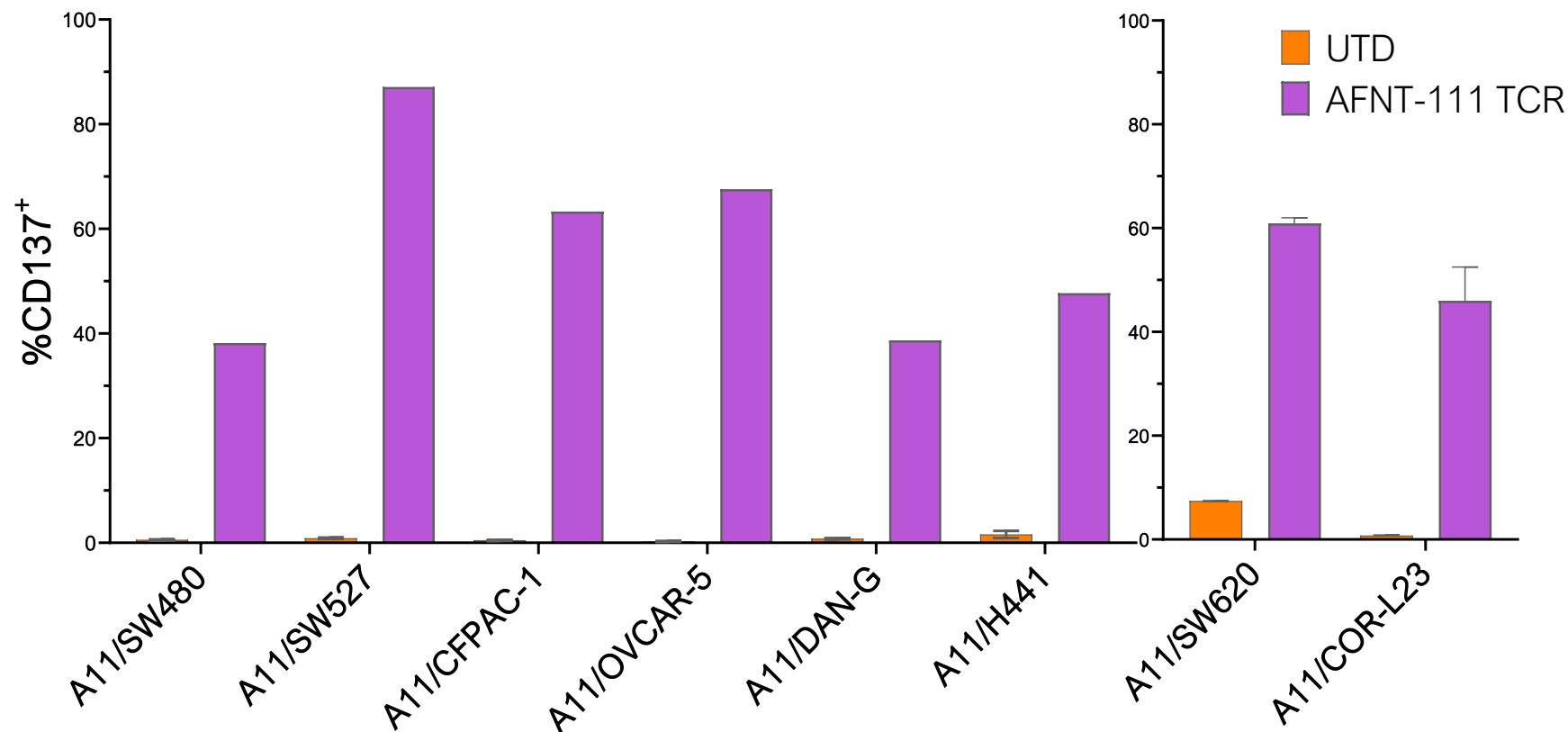
- Lead AFNT-111 TCR exhibits high functional avidity: EC50 in low pM
- Recognition of both 9mer and 10mer KRAS G12V epitopes with no reactivity to WT KRAS

**Peptide dose response CD137 T cell activation assay*

AFNT-111 TCR is activated by endogenously presented KRAS G12V across diverse tumor cell lines

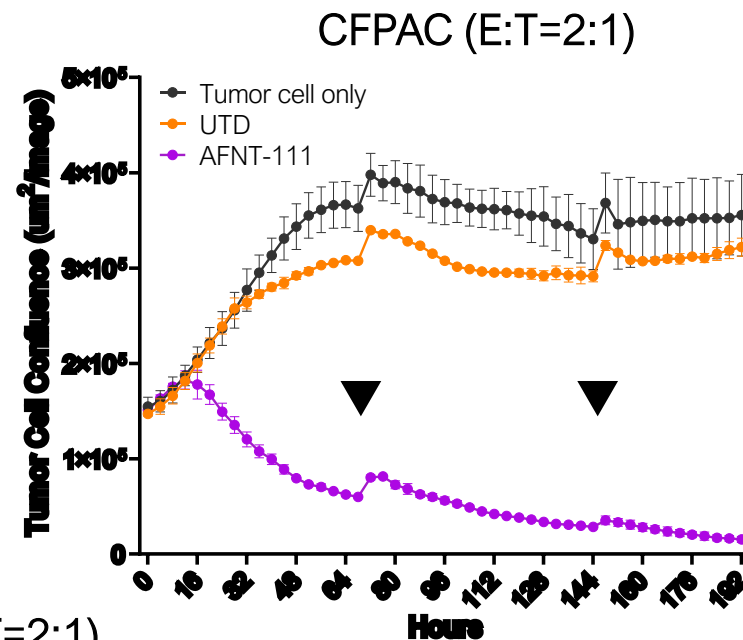
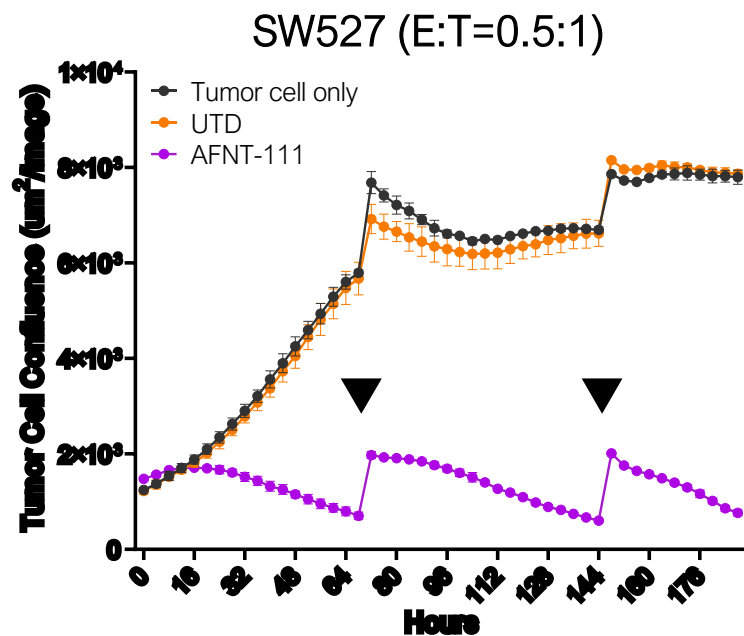


Cell lines used	Cancer derivation
CFPAC-1	Pancreatic
DAN-G	Pancreatic
SW620	Colon
SW480	Colon
COR-L23	Lung
NCI H441	Lung
SW527	Breast
OVCAR-5	Ovarian

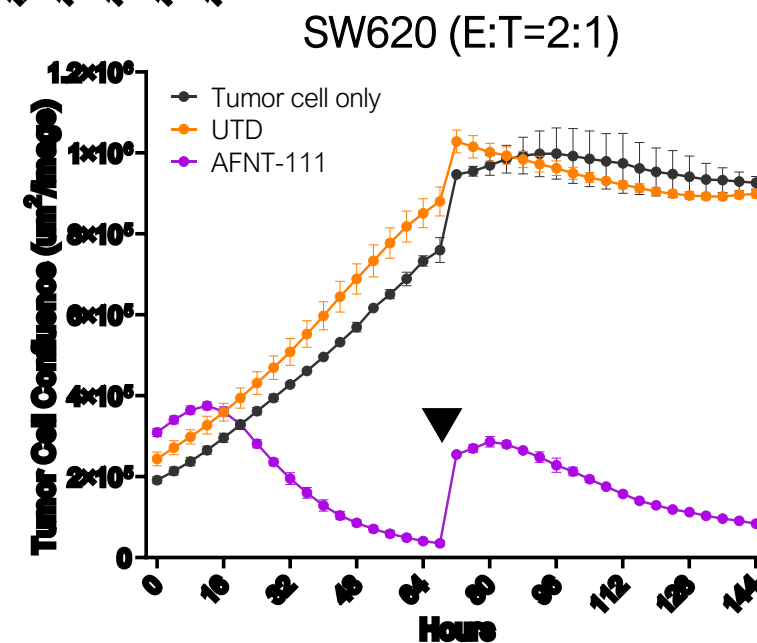
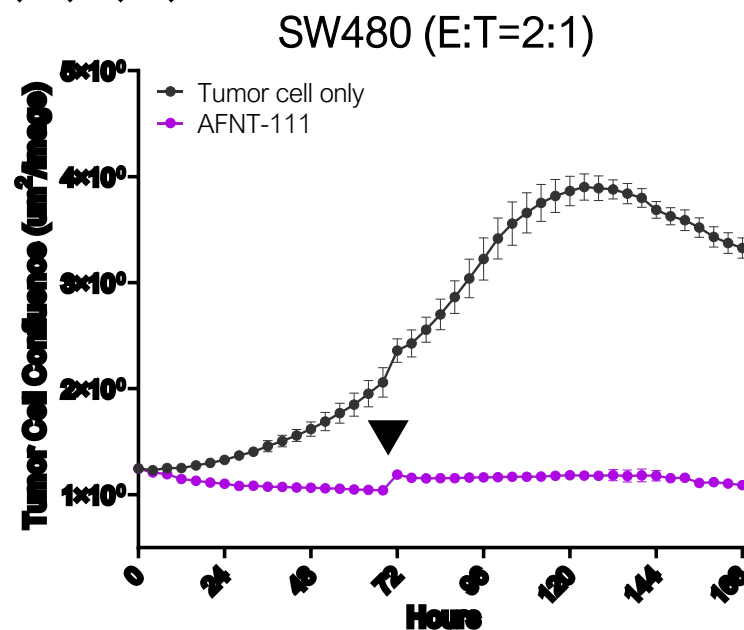


*CD137 T cell activation assay after co-culture with tumor cell lines

AFNT-111 T cells exhibit robust *in vitro* cancer killing in repeat tumor challenge assay

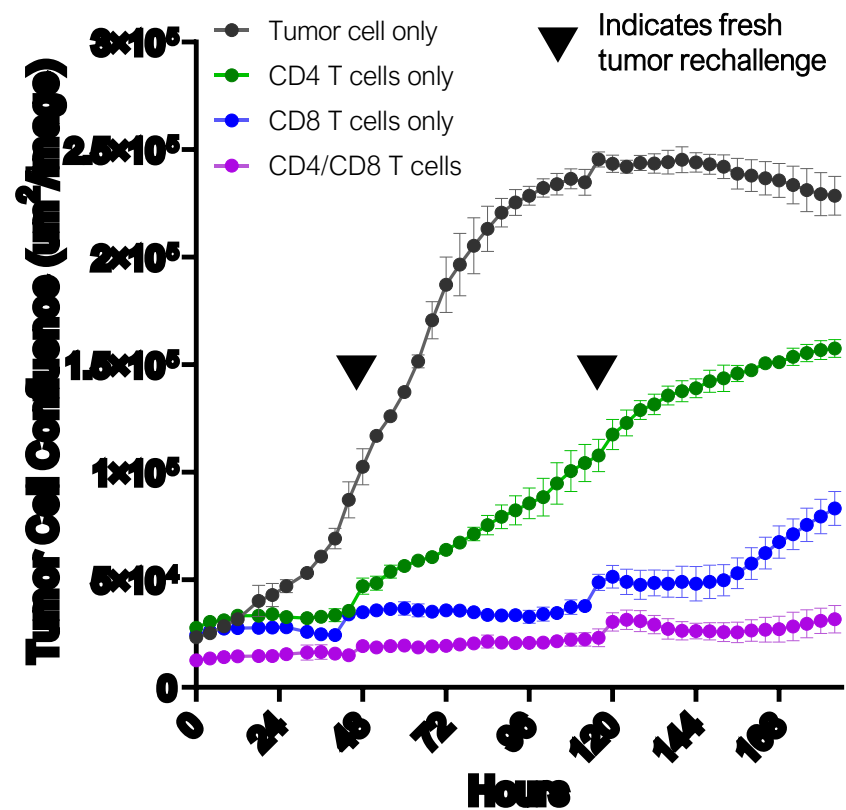


▼ Indicates fresh tumor rechallenge

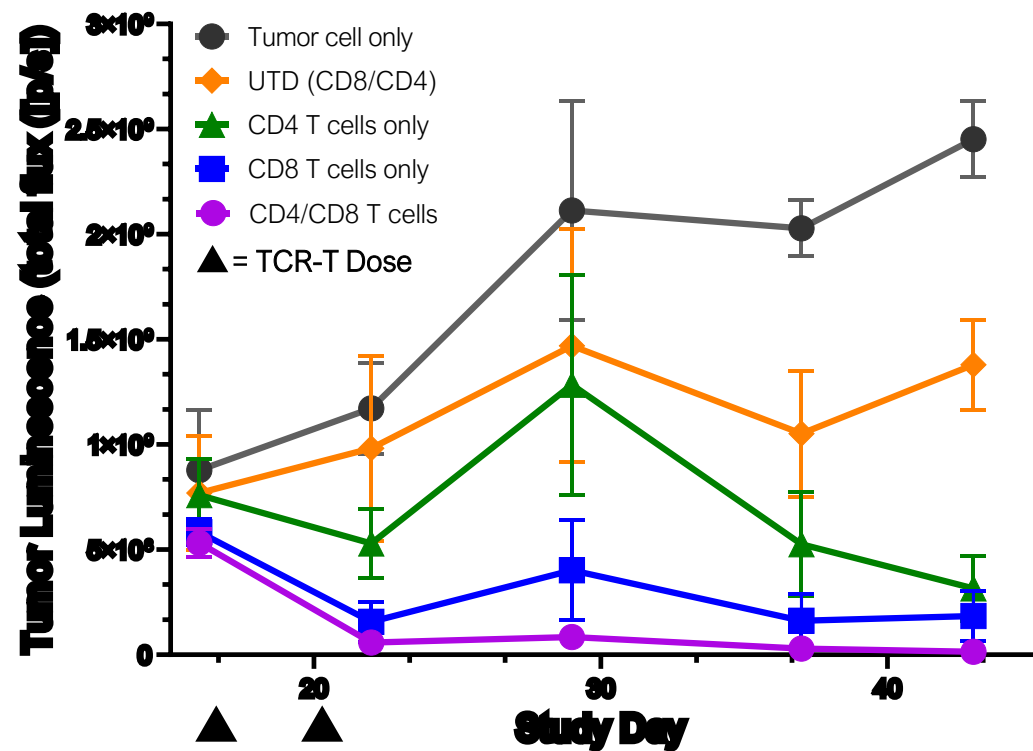


*Incucyte live tumor cell visualization assay

CFPAC (*In vitro* tumor challenge)



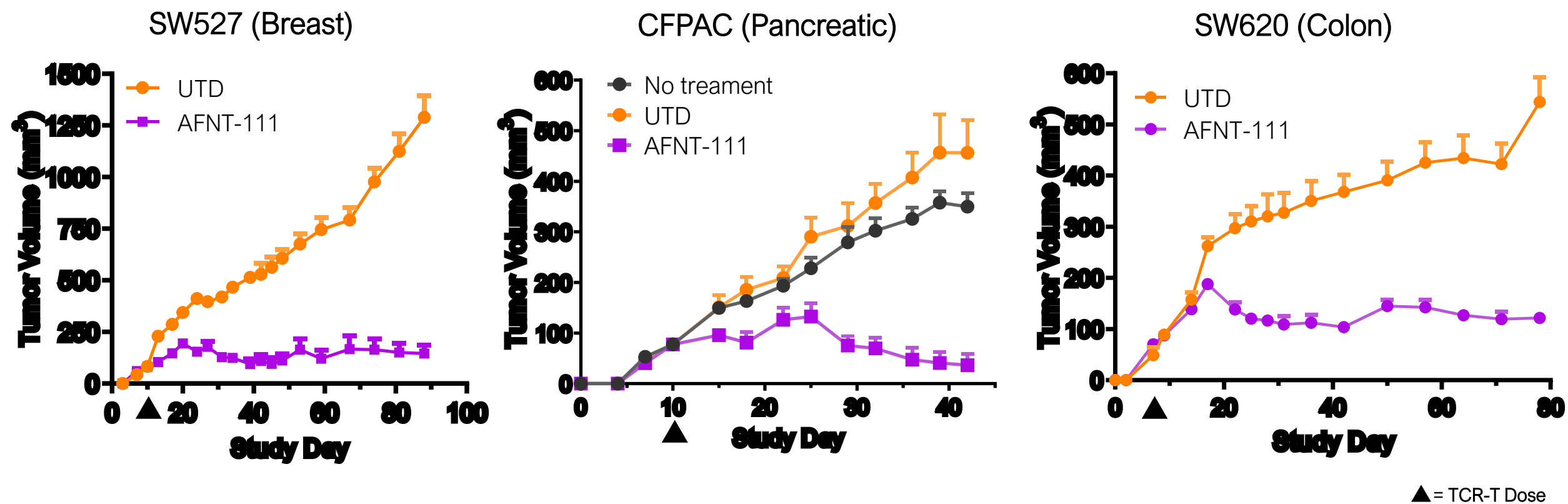
CFPAC-Luc (*In vivo* tumor challenge)



CD8⁺ T cell killing is enhanced by the presence of CD4⁺ T cells both *in vitro* and *in vivo*

*Incucyte live tumor cell visualization assay; Total T cells kept constant in CD4 only, CD8 only and CD4/CD8 groups

NSG mice randomized after IP tumor implantation (5 mouse/group)
Dose: two administrations of 7 × 10⁶ (D8) and 8 × 10⁶ (D21) AFNT-111 T cells



- NSG mice randomized after SC tumor implantation (5 mouse/group)
 - Dose: single IV administration of 3×10^6 AFNT-111 T cells on D07
- NSG mice randomized after SC tumor implantation (5 mouse/group)
 - Dose: single IV administration of 1×10^7 AFNT-111 T cells on D10
- NSG mice randomized after SC tumor implantation (5 mouse/group)
 - Dose: single IV administration of 3×10^6 AFNT-111 T cells on D09

XScan mutagenesis assay

p1 p2 p3 p4 p5 p6 p7 p8 p9

XVGAVGVGK

V**X**GAVGVGK

VV**X**AVGVGK

VVG**X**VGVGK

VVGAV**X**GVGK

VVGAVGV**X**GK

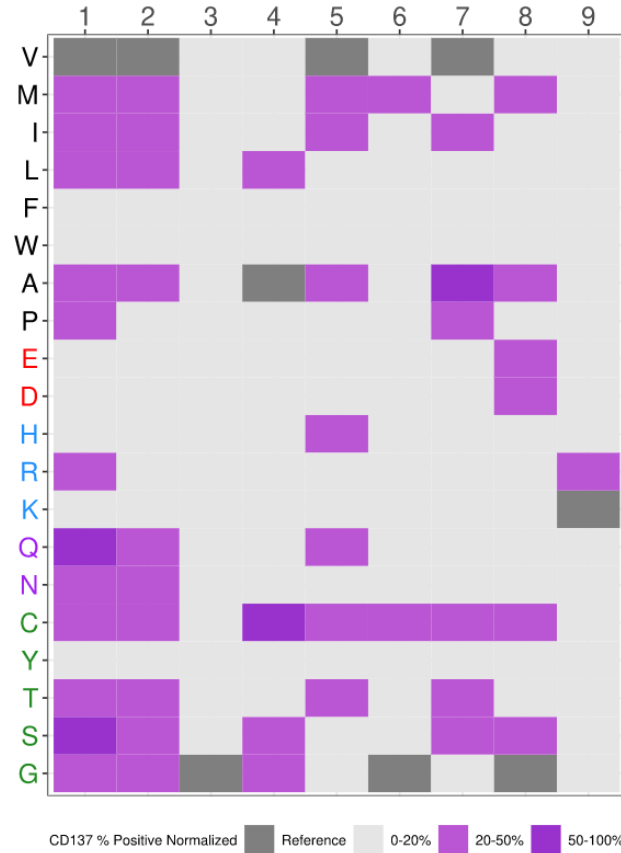
VVGAVGVGX

VVGAVGVGX

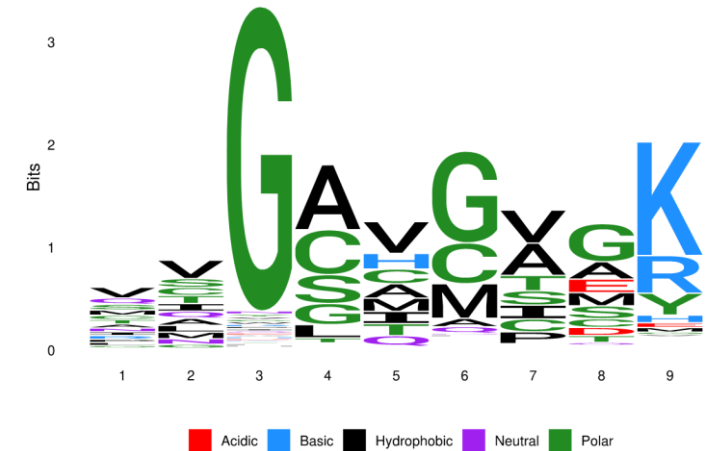
VVGAVGVGX

X= All other 19 amino acids

Data summary



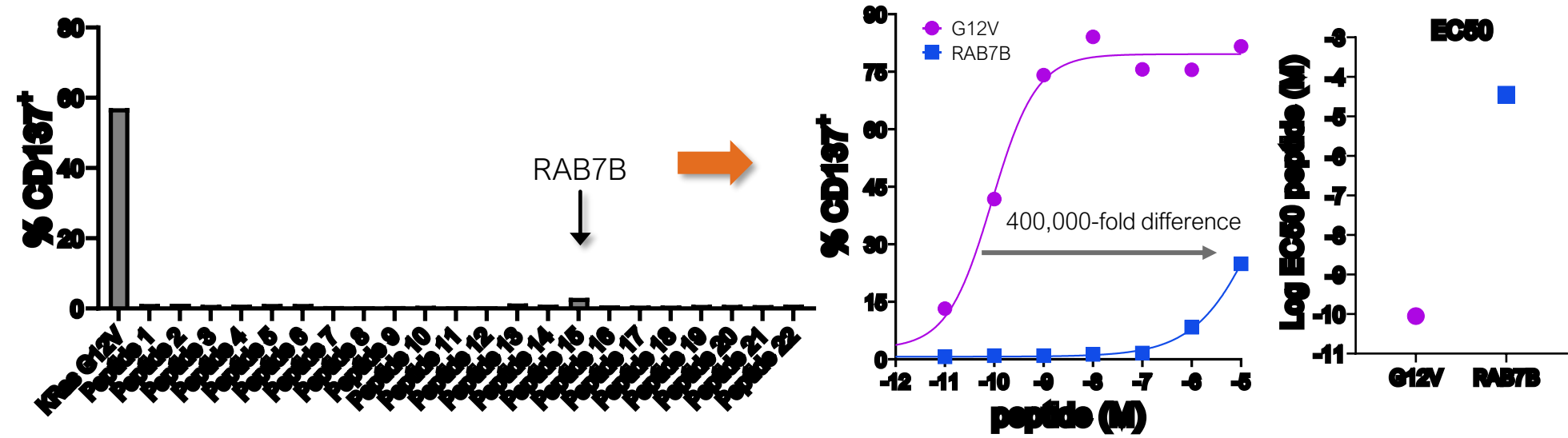
Recognition motif of AFNT-111 TCR



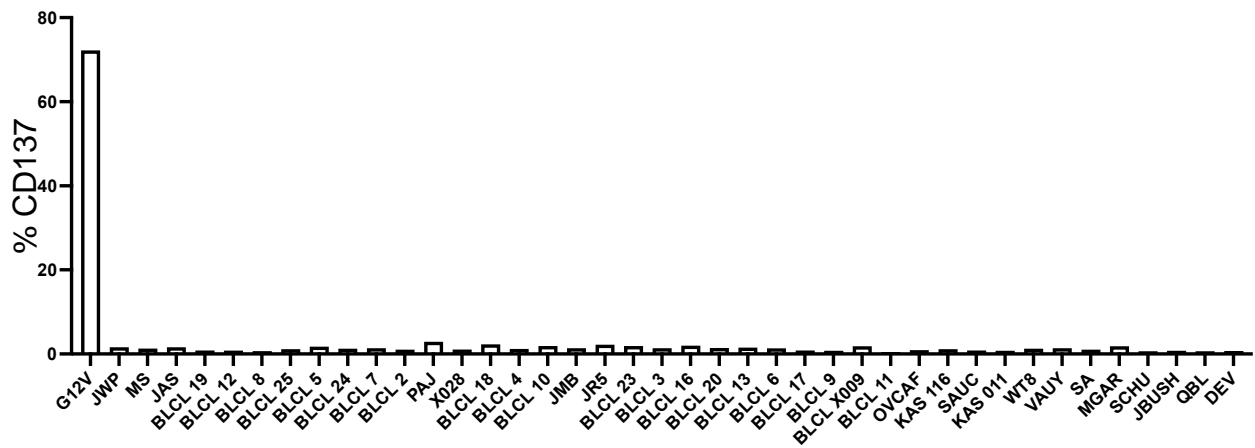
- XScan assay used to define the peptide recognition motif of AFNT-111
- In silico* assessment across human proteome reveals only 22 peptides of potential concern (next slide)

*AFNT-111 T cells incubated with XScan peptides and assessed for CD137⁺ T cell activation

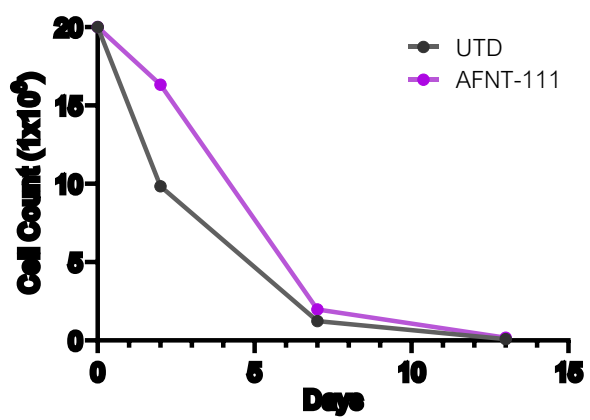
- No cross-reactive self-peptides of concern identified
- Peptide dose response studies confirm that TCR does not respond to physiological concentrations of RAB7B



- No allo-reactive responses detected
- B-LCL library covers >95% of the most common HLA alleles



- No cytokine-independent growth



AFNT-111, a preclinically safe and effective TCR-engineered T cell therapy targeting the oncogenic driver KRAS G12V mutation (Poster #244)

- AFNT-111 is a potent and specific preclinical TCR-engineered T cell therapy that is cytotoxic to KRAS G12V-expressing tumor cells both *in vitro* and *in vivo*
- Crossreactivity and alloreactivity assessments established a strong AFNT-111 preclinical safety profile
- AFNT-111 IND anticipated in 1H'23
- First-in-human clinical studies in collaboration with Fred Hutch Cancer Research Center
 - Phase I Objectives: Safety, tolerability, maximum tolerated dose (MTD), and preliminary efficacy
 - Clinical Indications: Advanced or metastatic Pancreatic ductal adenocarcinoma (PDAC), Colorectal adenocarcinoma (CRC), Non-small cell lung cancer (NSCLC)
 - Target population: HLA-A*11:01⁺; KRAS G12V⁺ patients

KRAS G12V T cell receptor-engineered T cells expressing the durability FAS-41BB switch receptor exhibit a potent, persistent, coordinated CD4/CD8 anti-tumor response in vitro and in vivo (Poster #342)

- Affini-T therapeutics is also developing a next generation TCR-T cell therapy with novel FAS-41BB switch receptor armoring to increase fitness, persistence, and anti-tumor activity



- Xingyue He
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