

Targeting High-Risk Leukemias with Redirected WT1-Specific T- Cells

Aude Chapuis, MD

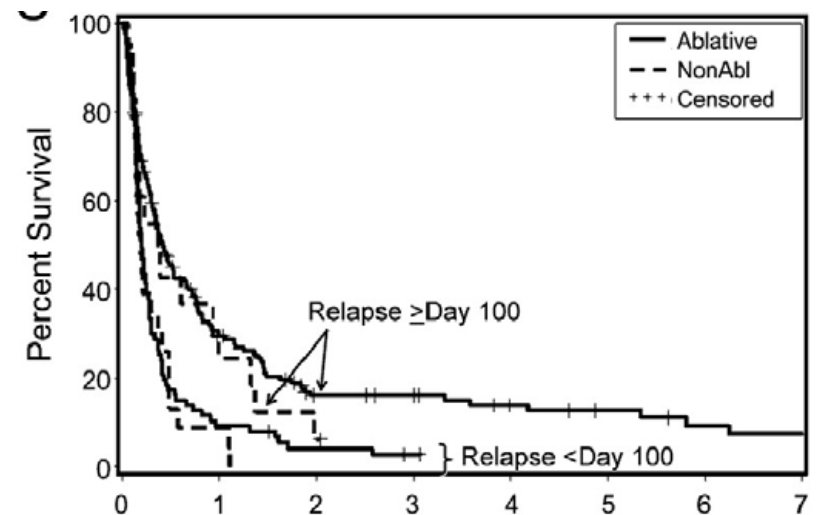
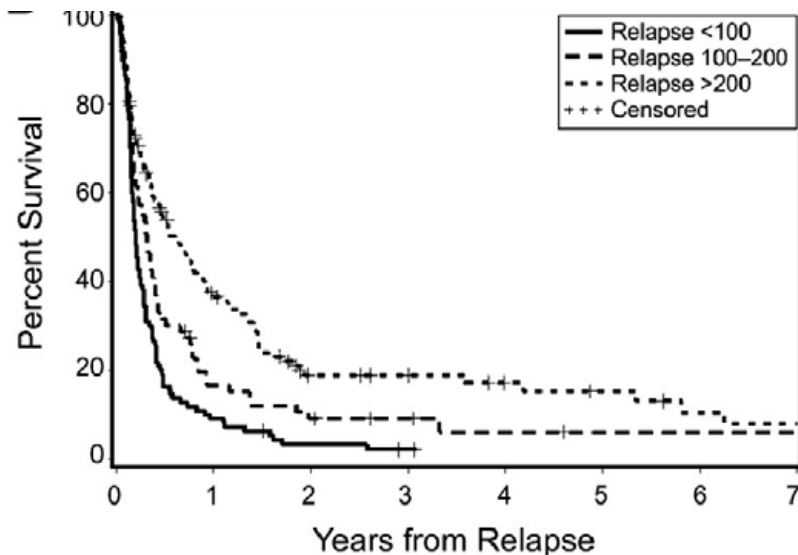
Fred Hutchinson Cancer Research Center
Seattle

Background: High-Risk Acute Myeloid Leukemia (AML) and Myelodysplastic Syndrome (MDS)

- Definition of high-risk AML:
 - Unfavorable Cytogenetics/FISH/molecular abnormalities:
 - Del 5q; del 7q; abnormalities involving 3q, 9q, 11q, 20q, 21q, 17p; t(6;11); t(15;17); complex karyotype; FLT3-ITD mutation.
 - Refractory / beyond 1st remission.
 - AML caused by previous chemotherapy (therapy-related).
 - AML arising for a previous hematologic disorder.
- Definition of high-risk MDS:
 - MDS with IPSS score >1.5 and/or
 - Unfavorable cytogenetics.
- **For these patients, the probability of post-HCT relapse (>95% fatal) and/or death is >50% within 2 years** (Gyurcokza B., JCO 2010; Deeg J., Blood 2002; Gratwohl A., BMT 1996; Radich J., Semin. Hematol. 2010).

Leukemias that relapse post-HCT have a very poor prognosis

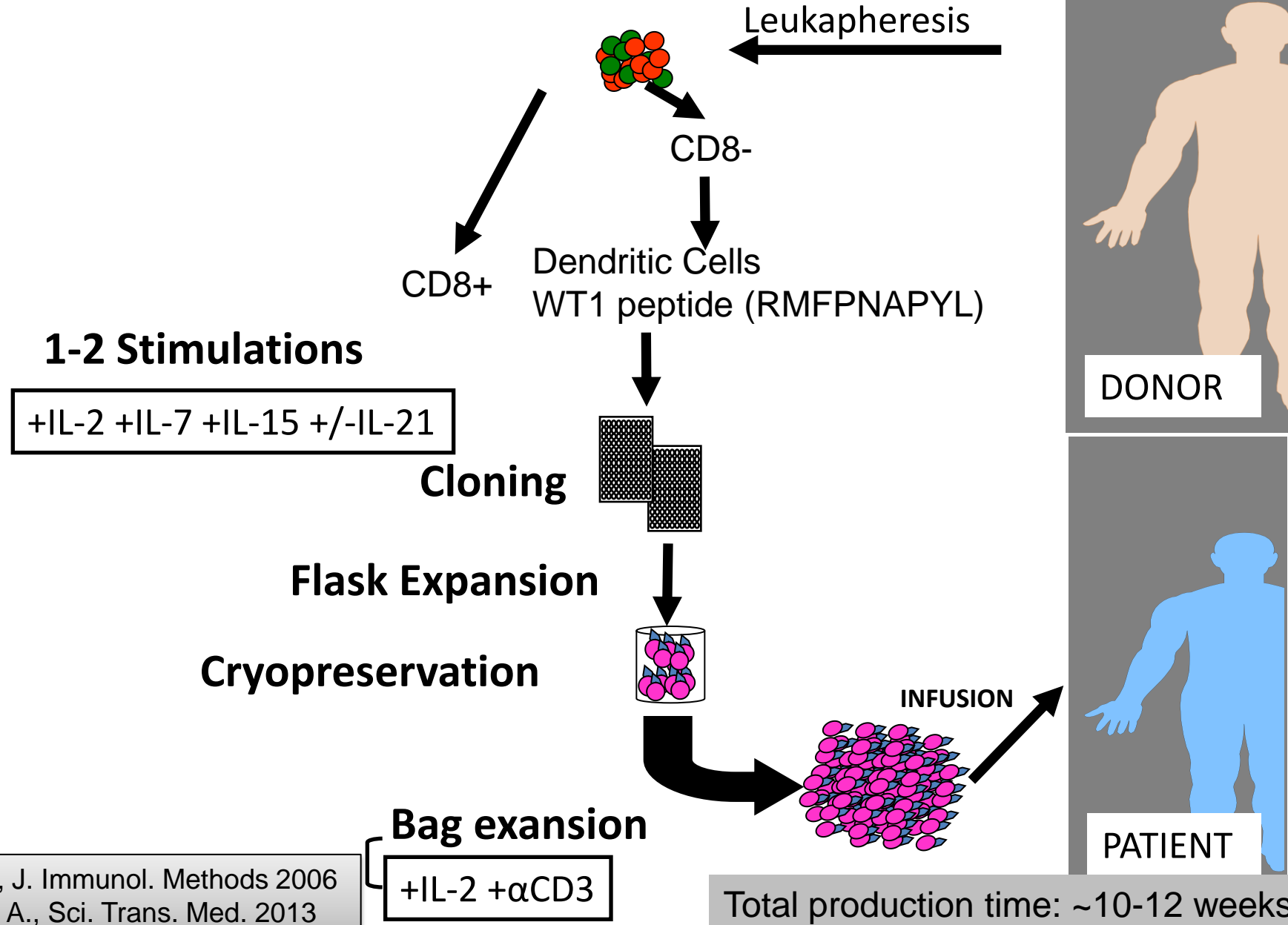
- Current post-HCT treatment options include withdrawal of immunosuppression, re-induction chemotherapy, donor lymphocyte infusion from the original HCT donor, second non-myeloablative transplant.
- Patients with early post-HCT relapse have a worse prognosis with shortened survival compared to patients who relapse later.
- **Trials designed to prevent relapse in patients with high-risk leukemias constitute a high priority, but no predictably effective therapy exists.**



Rationale for targeting WT1 in patients with high-risk leukemias

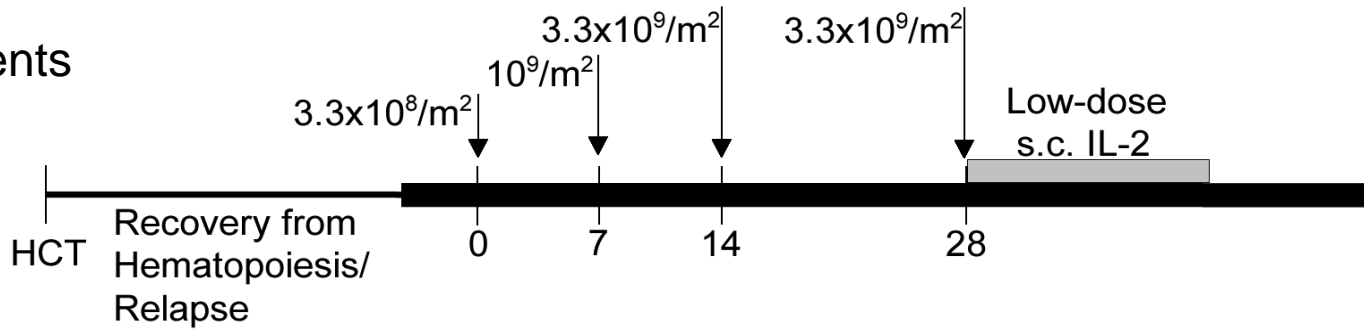
- WT1, a zinc finger protein that regulates gene expression
 - originally characterized as a gene associated with Wilm's tumor (Pritchard-Jones K., *Nature* 1990).
- Over-expressed (10->1000 fold) in AML, ALL, CML, MDS blasts
 - higher levels correlated with worse prognosis (Bergman L., *Blood* 1997).
- Low-level expression in adult kidney podocytes, testis sertoli cells, mesothelial lung cells (mesodermal origin) and CD34⁺ progenitor cells (Inoue K., *Blood* 1997).
- CD8⁺ T-cells can distinguish difference in protein expression between physiologic WT1 expression levels and leukemic cells (Gao L et al., 2000).
- WT1 vaccine studies have resulted in anti-tumor responses, including some long-term complete remissions in solid tumors and leukemias (Oka Y., *Immunotherapy* 2010; Van Tendeloo *PNAS* 2010; Ochsenreither S., *J Immunother* 2010).
- No toxicities to normal tissues expressing physiologic levels of WT1 have ever been reported.

Generation of WT1-specific CD8⁺ T cell clones from donor PBMC: Methods

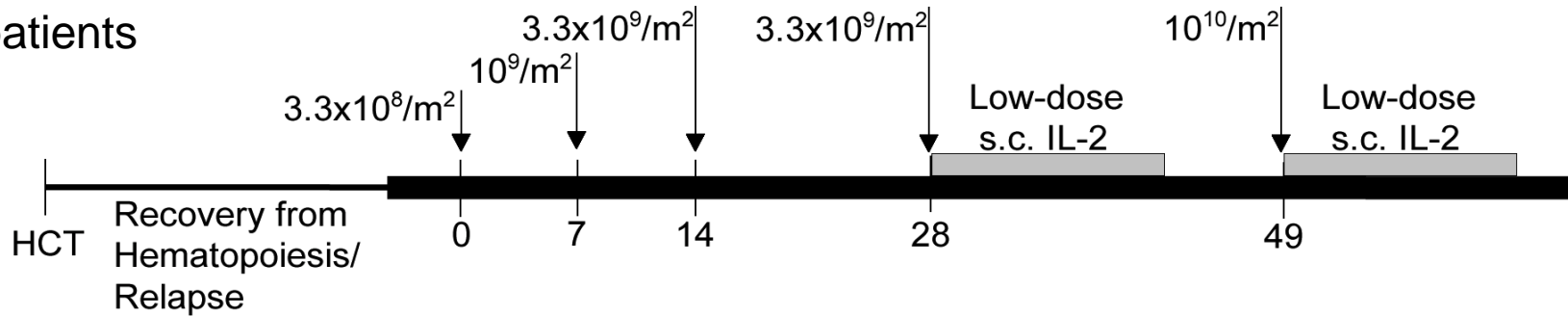


Infusion of donor-derived WT1-specific CD8⁺ T cell clones: Treatment plan

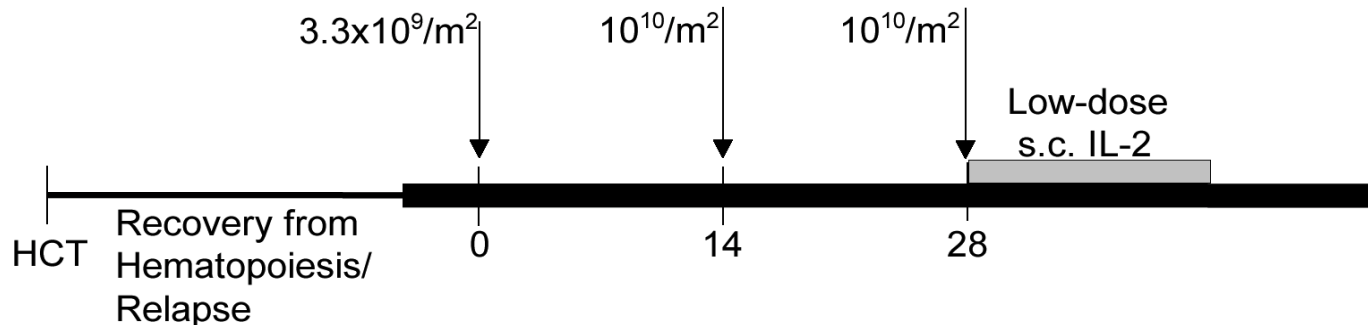
First 2 patients



Second 2 patients



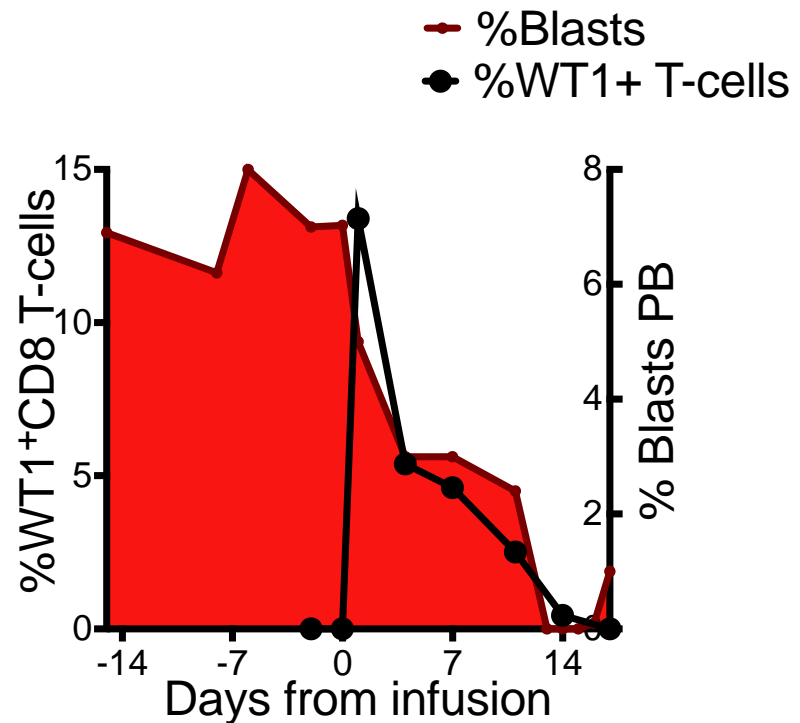
Last 7 patients



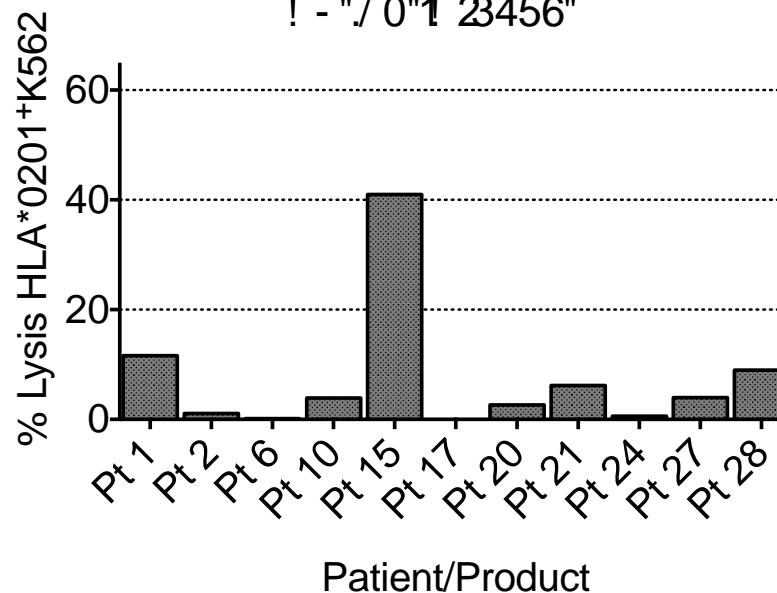
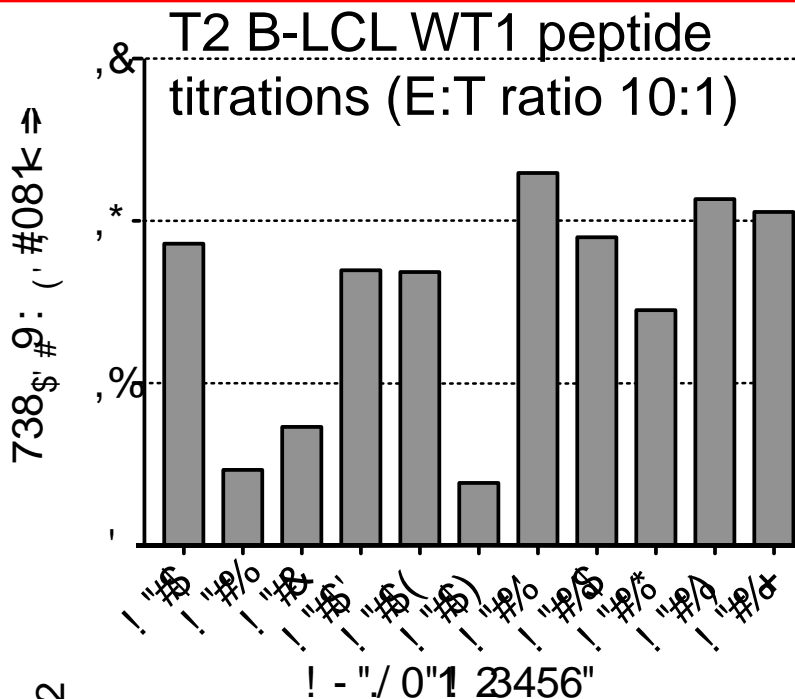
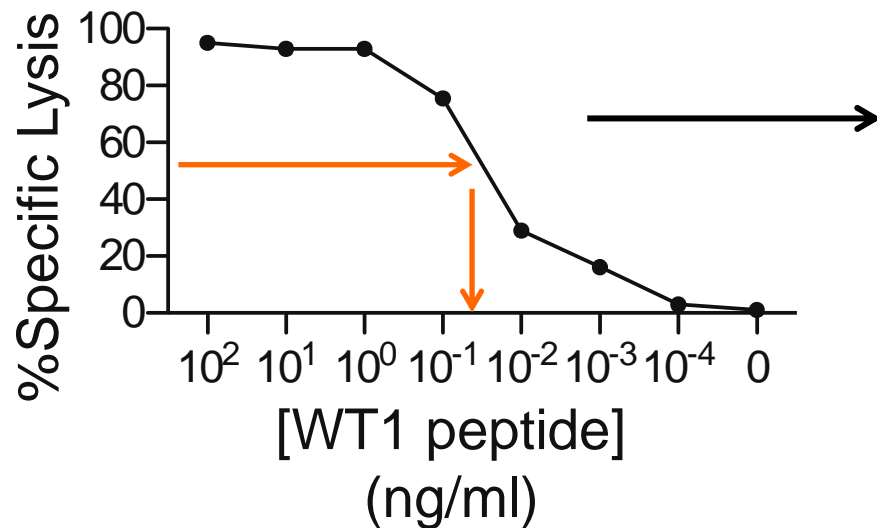
Infusion of donor-derived WT1-specific CD8⁺ T cell clones: Results

- 11 patients with high-risk or relapsed leukemias received escalating doses of WT1-specific T cells for a maximum dose of 10^{10} cells/m².
- No injuries were observed to tissues expressing physiologic levels of WT1.
- Evidence of direct albeit transient anti-leukemic activity.

Patient 15



Variability in the avidity of infused clones



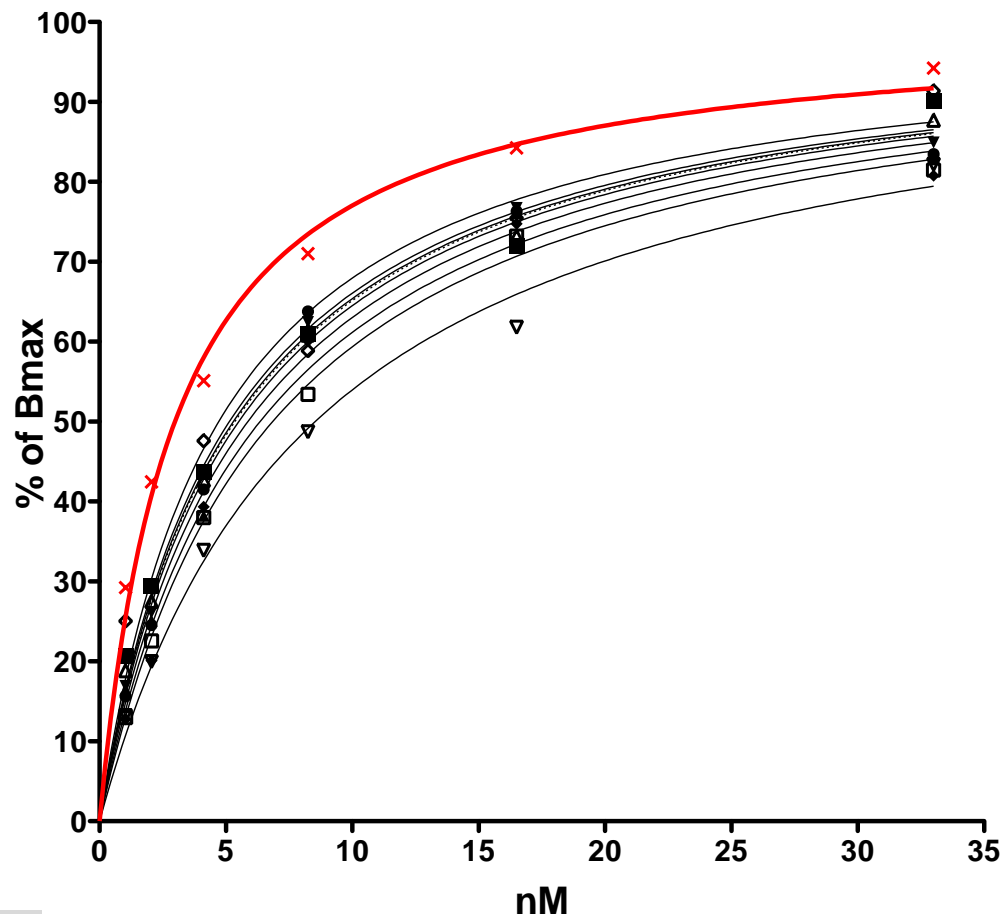
Rationale for targeting WT1 with a naturally occurring TCR of higher affinity

- Although the most avid clone (based on WT1 peptide titrations) from each patient/donor pair was selected for infusion, **the avidities obtained were variable.**
- T cells expressing **higher affinity TCRs** exhibited improved recognition of WT1⁺ target cells.

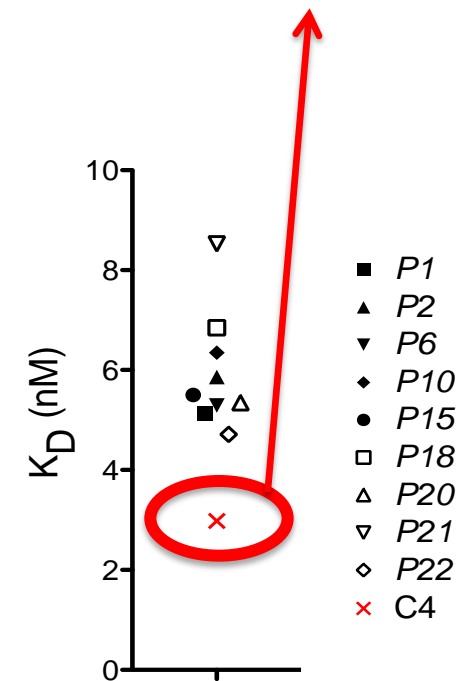
As TCR affinity is a major determinant of T-cell avidity, expressing a higher affinity TCR is associated with better recognition of leukemic cells

Selection of a naturally occurring high-affinity HLA A*0201-restricted WT1-specific TCR

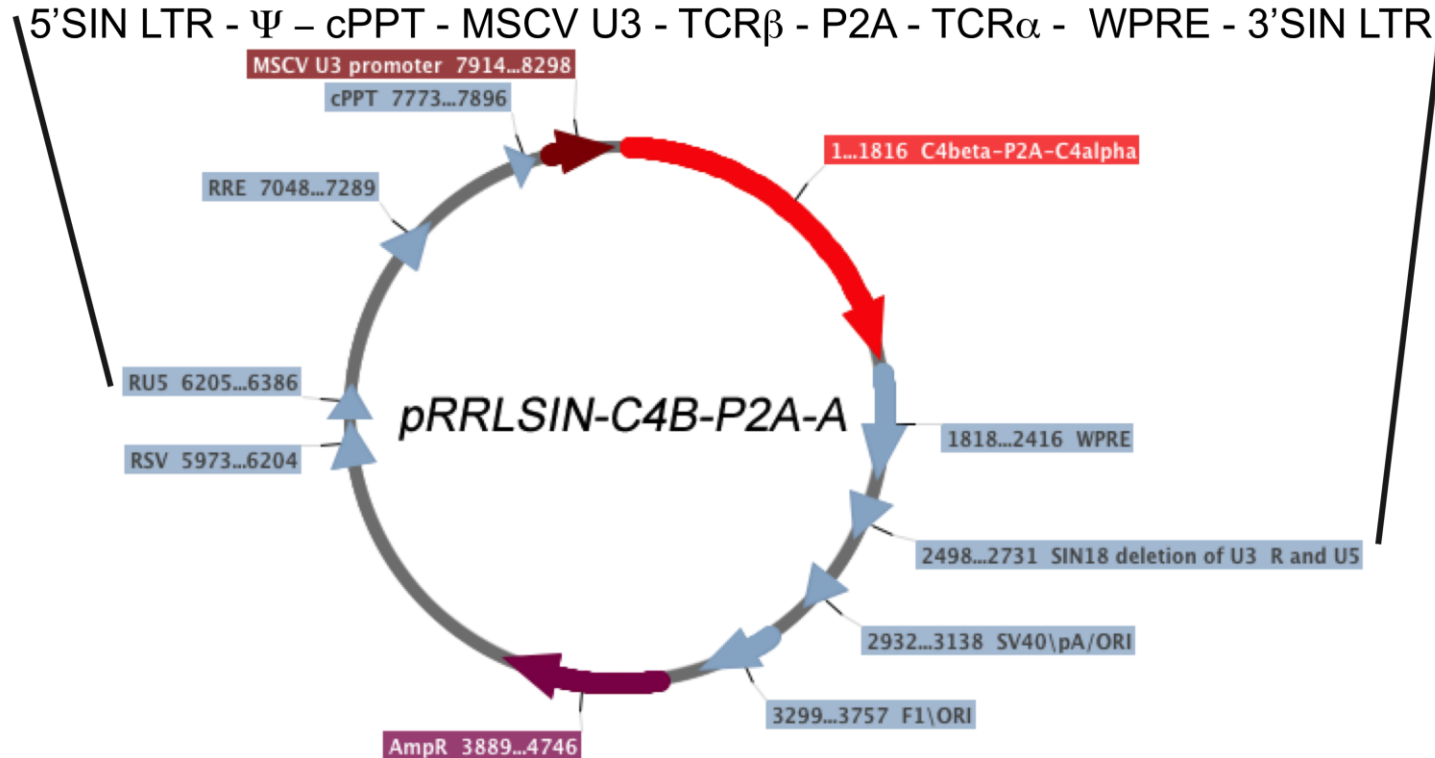
Relative affinity of the 'parent C4 clone' compared to clones administered to patients:



The K_D of C4 is 2.98 nM



Insertion of the TCR_{C4} in a lentiviral SIN vector with a strong internal promoter



SIN LTR = Self-inactivating long terminal repeat

Ψ = Packaging signal

cPPT = Central polypurine tract

MSCV = MSCV U3 promoter

TCRβ = Beta chain of the WT1-specific TCR

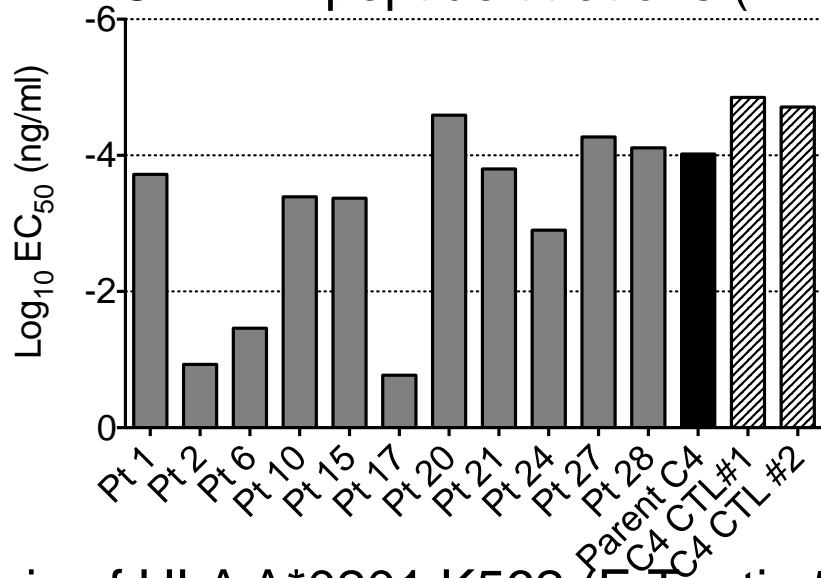
P2A= 2A element from the porcine teschovirus

TCRα = Alpha chain of the WT1-specific TCR

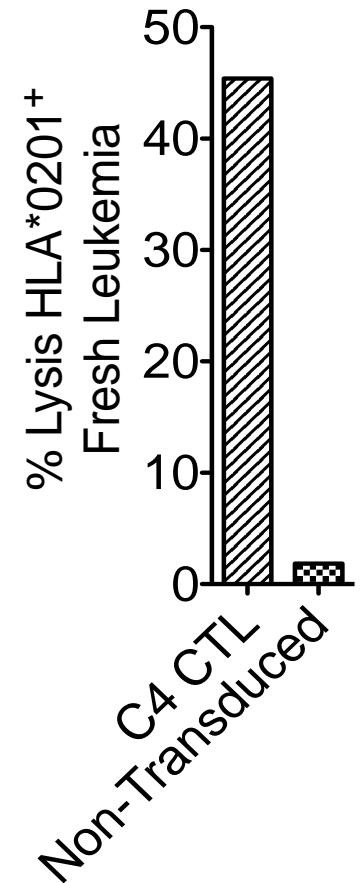
WPRE = Woodchuck Hepatitis Virus Posttranscriptional Regulatory Element

Relative avidities of the parent C4 clone and TCR_{C4} CTL

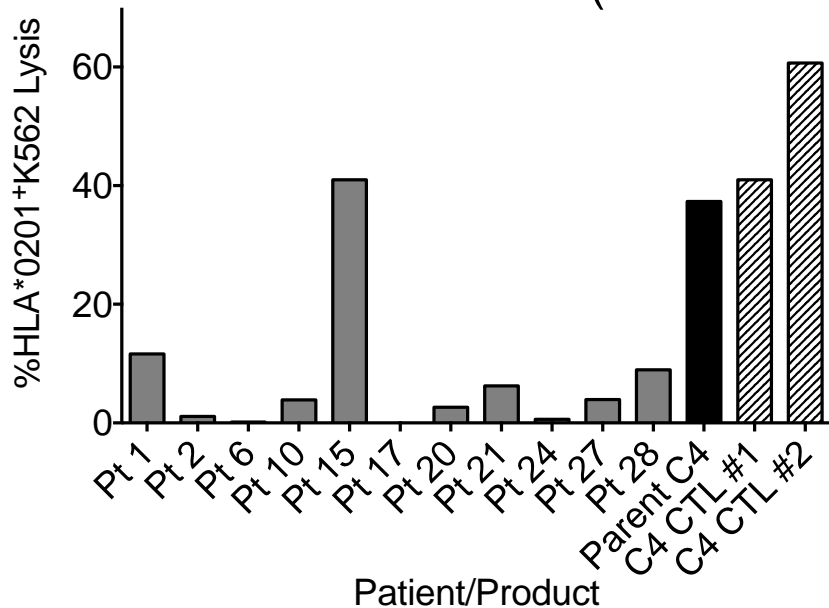
T2 B-LCL WT1 peptide titrations (E:T ratio 10:1)



Lysis of fresh HLA A*0201⁺ Leukemia cells



Lysis of HLA A*0201 K562 (E:T ratio 5:1)



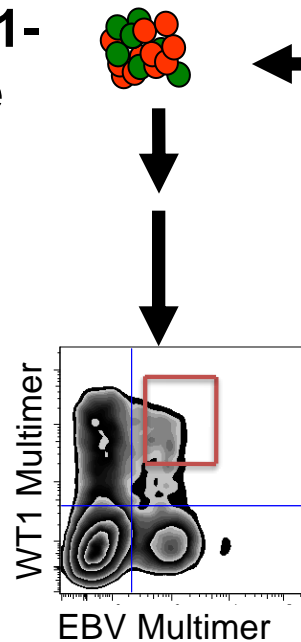
Generation of TCR_{C4}-transduced products for infusion

1. **Stimulation with HLA A*0201-restricted **EBV/CMV** peptide**

2. **Transduce with TCR_{C4} Lentiviral supernatant**

+IL-2 +IL-7 +IL-15

3. **Sort on virus- and WT1-multimer⁺ cells**



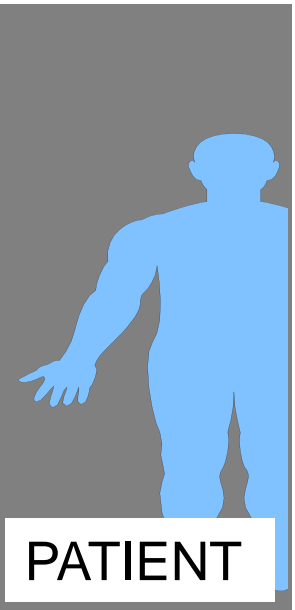
4. **Flask expansion**

+IL-2+αCD3

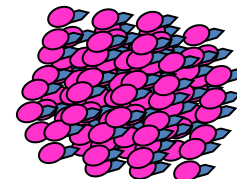
5. **Bioreactor expansion**

+IL-2+αCD3

G-mobilized
Leukapheresis



INFUSION



Total production time: ~5 weeks

Leukemia trial: Patient characteristics -update

Pt#	M/F	Age	EBV/ CMV	Disease	Leukemia WT1 expression*	Disease Characteristics at 1st CTL infusion
1	M	56	EBV	AML, chloroma 5y after 1st myeloablative HCT	ND	PET+ Chloroma
2	F	50	CMV	AML, 2nd HCT for relapse	YES	MRD by flow (0.03%)
3	M	48	EBV	High-risk AML	ND	NED
4	M	25	EBV	High-risk AML	YES	NED
5	M	48	EBV	High-risk AML	YES	70% peripheral blasts
6	F	20	CMV	High-risk AML 2nd HCT for relapse	ND	NED
7	F	33	EBV	High-risk AML	YES	MRD by flow (0.08%)
8	F	62	EBV	High-risk AML	ND	MRD by flow (0.26%)
9	F	67	CMV	High-risk AML	YES	MRD by FISH (0.3%)
10	M	74	EBV	High-risk AML	ND	NED
11	M	59	EBV	High-risk AML	ND	NED
12	F	55	EBV	High-risk AML	ND	NED
13	F	59	EBV	High-risk AML	ND	NED
14	M	17	EBV	High-risk AML, relapse after 1st HCT	ND	MRD by flow (0.02%)

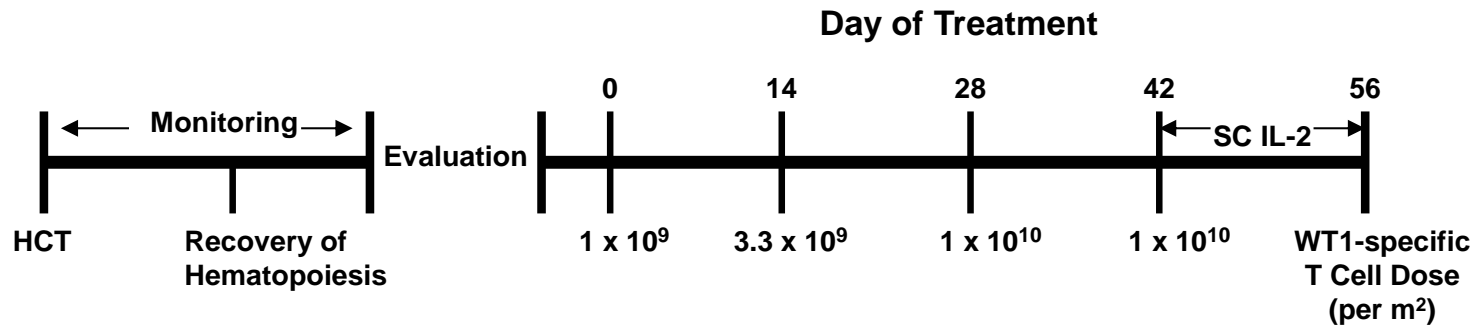
*>250-fold expression compared to normal donor marrow

TCR_{C4}-transduced EBV/CMV-specific CD8⁺ T-cells – plan of treatment

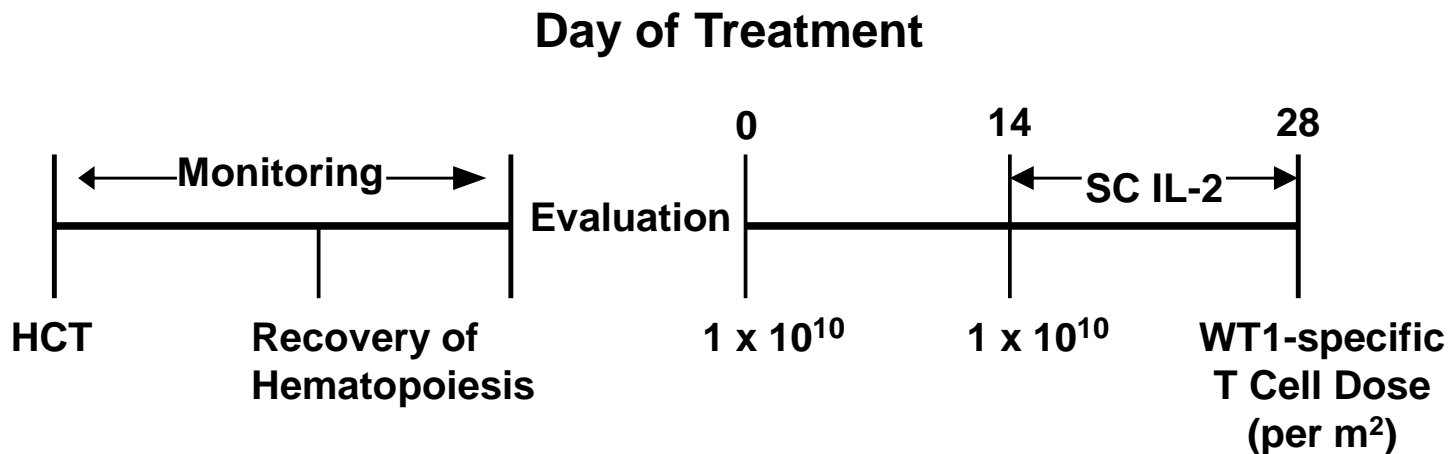
1st Phase: Dose escalation (5 patients completed 4 infusions):

Arm 1: Preventative treatment after HCT.

Arm 2: Treatment of detectable disease.



2nd Phase (7 patients):



TCR_{C4}-transduced EBV/CMV-CTL do not injure normal tissues expressing physiologic levels of WT1

NCI CTCAE v4.0	Grade 3	Grade 4
Fever (within <24 hours of infusion)	2	0
Chills (within <24 hours of infusion)	1	0
Hypotension (within <24 hours of infusion)	1	0
Lymphopenia (<10 days after infusion)	8	1
Thrombocytopenia	3	0
Transient maculo-papular rash	1	0
GVHD	0	0

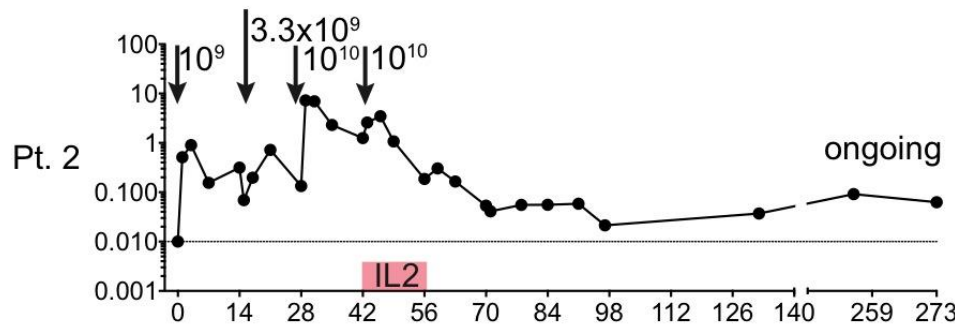
Preliminary Clinical Outcomes

- 7 pts received TCR_{C4}-transduced T cells with **NED post-HCT** either on the prevention arm or after salvage chemotherapy for relapse:
 - All are alive with NED average **240 days** (range 93-525 days) after infusions, **328 days** (range 161-605) days after HCT.
- 3 pts received TCR_{C4}-transduced T cells with **MRD post-HCT**:
 - 1 **cleared MRD** and is alive without additional therapy **557** days after infusions.
 - 2 pts continue to have detectable but decreased MRD 183 and 86 days after infusions.
- 4 pts **progressed** during the time they were receiving the TCR_{C4}-transduced T cells:
 - 1 had a resistant extramedullary chloroma.
 - 1 had 70% peripheral blasts (increasing rapidly).
 - 2 had received salvage therapy for early relapse post-HCT (1 pt received CMV-specific cells, no persistence).

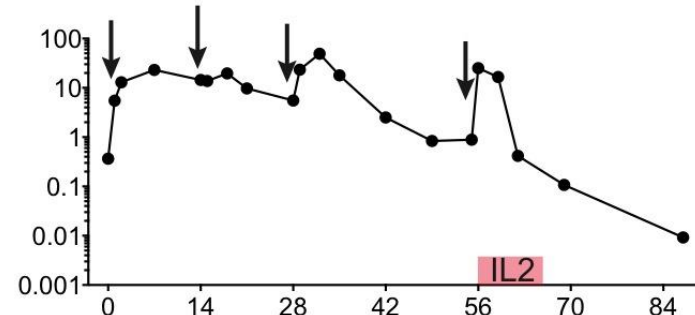
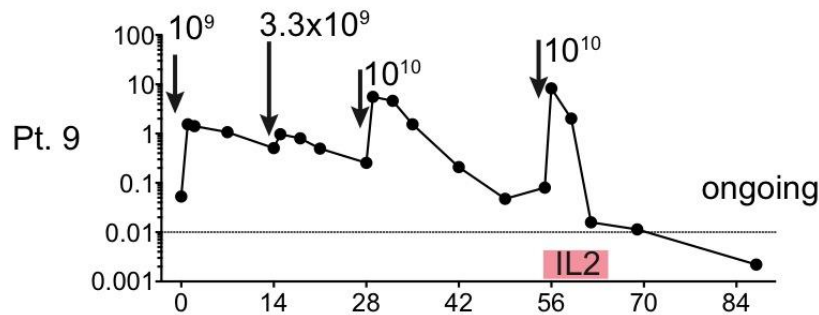
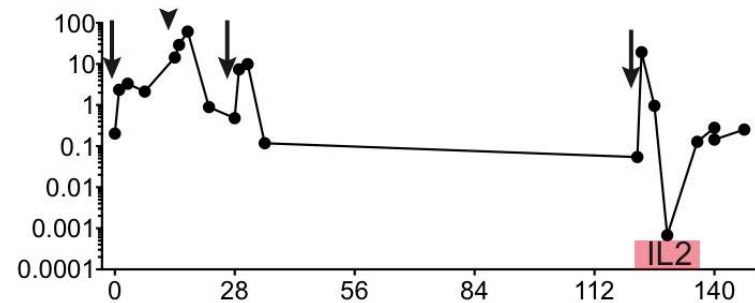
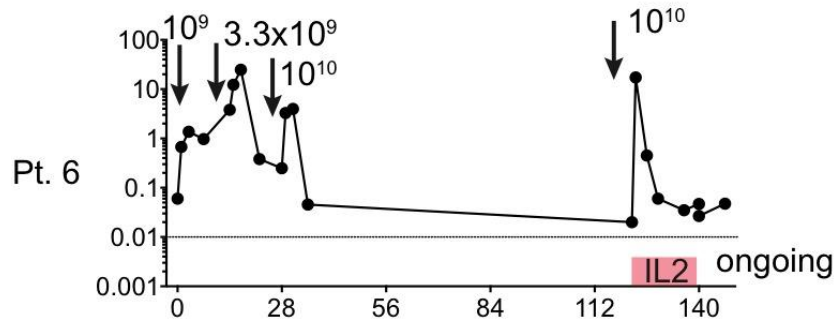
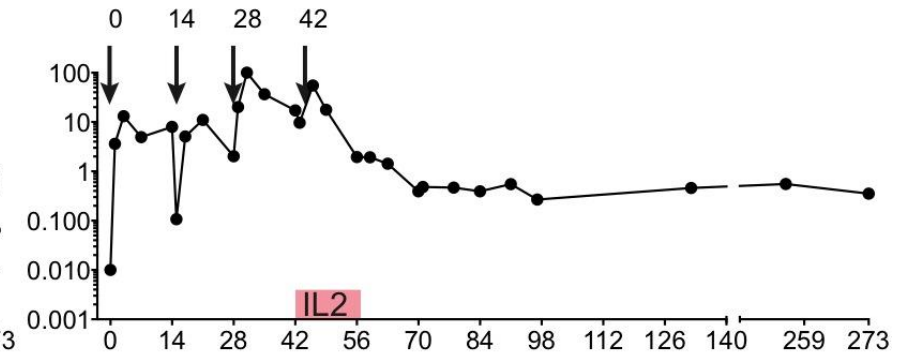
Assessment of persistence after infusions: CMV-Specific CTL

3 of 3 patients:

% WT1⁺ CD8⁺ T cells:



WT1⁺ CD8⁺ T cells / μ l

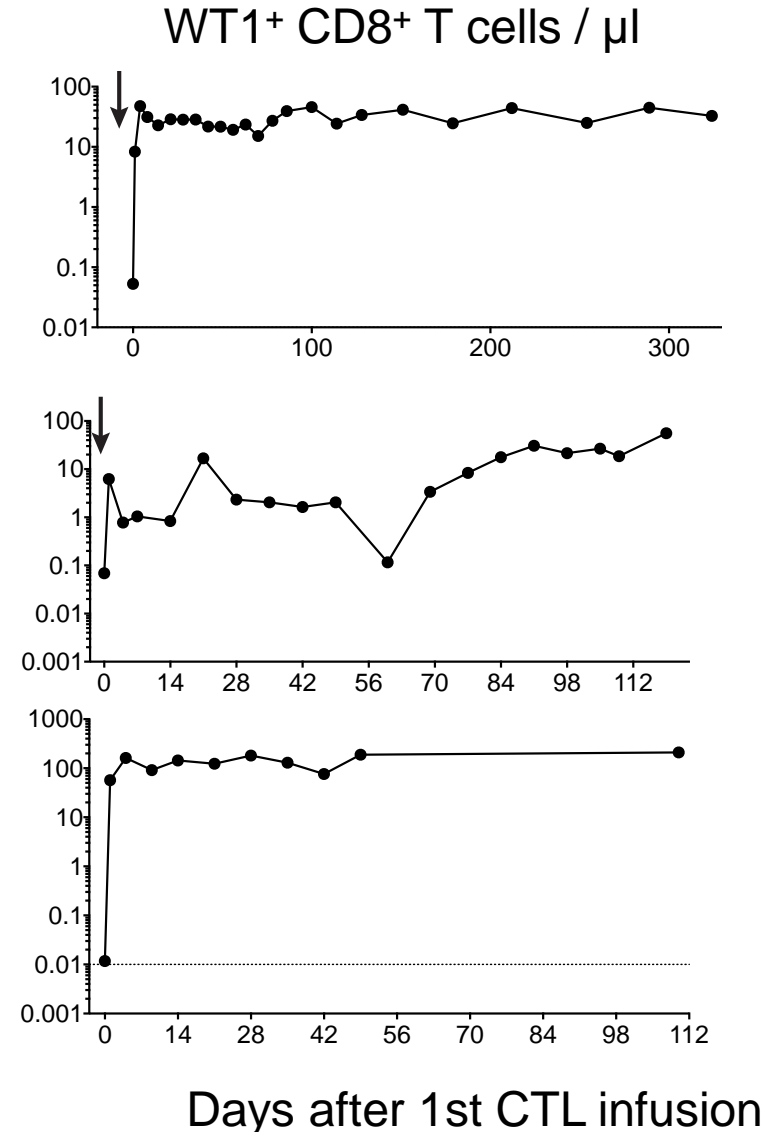
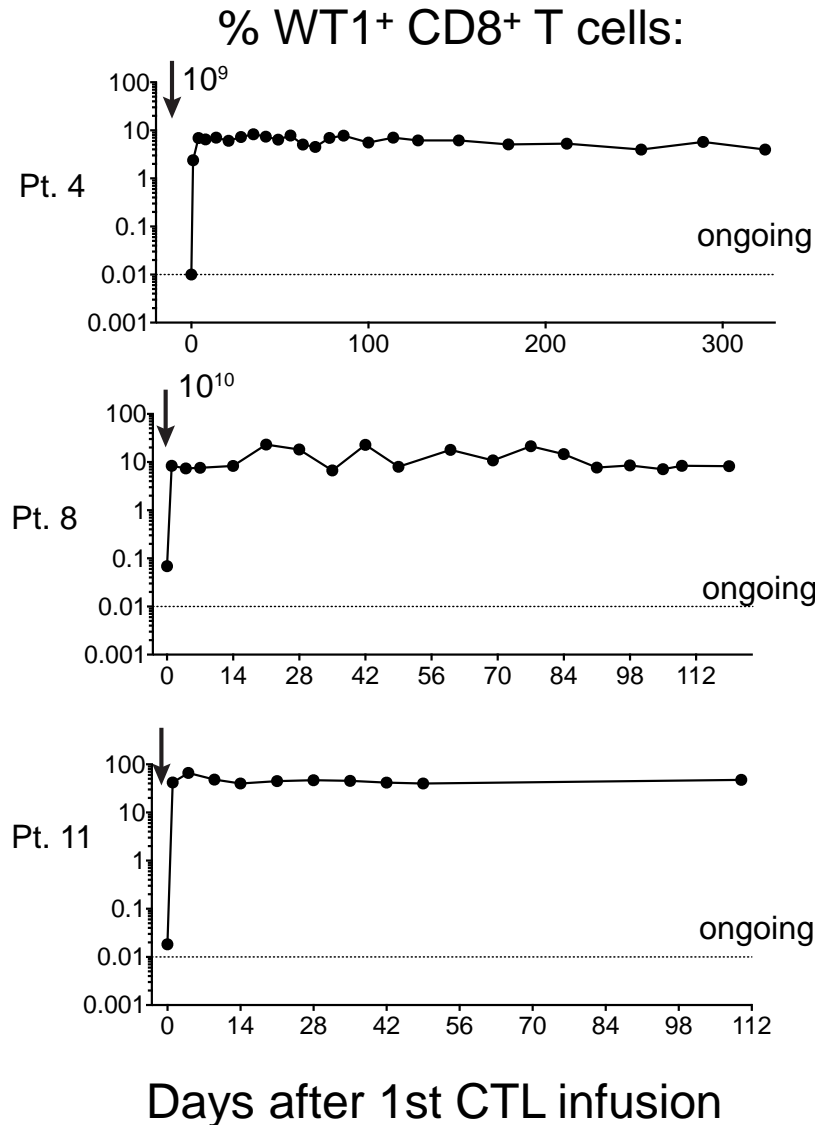


Days after 1st CTL infusion

Days after 1st CTL infusion

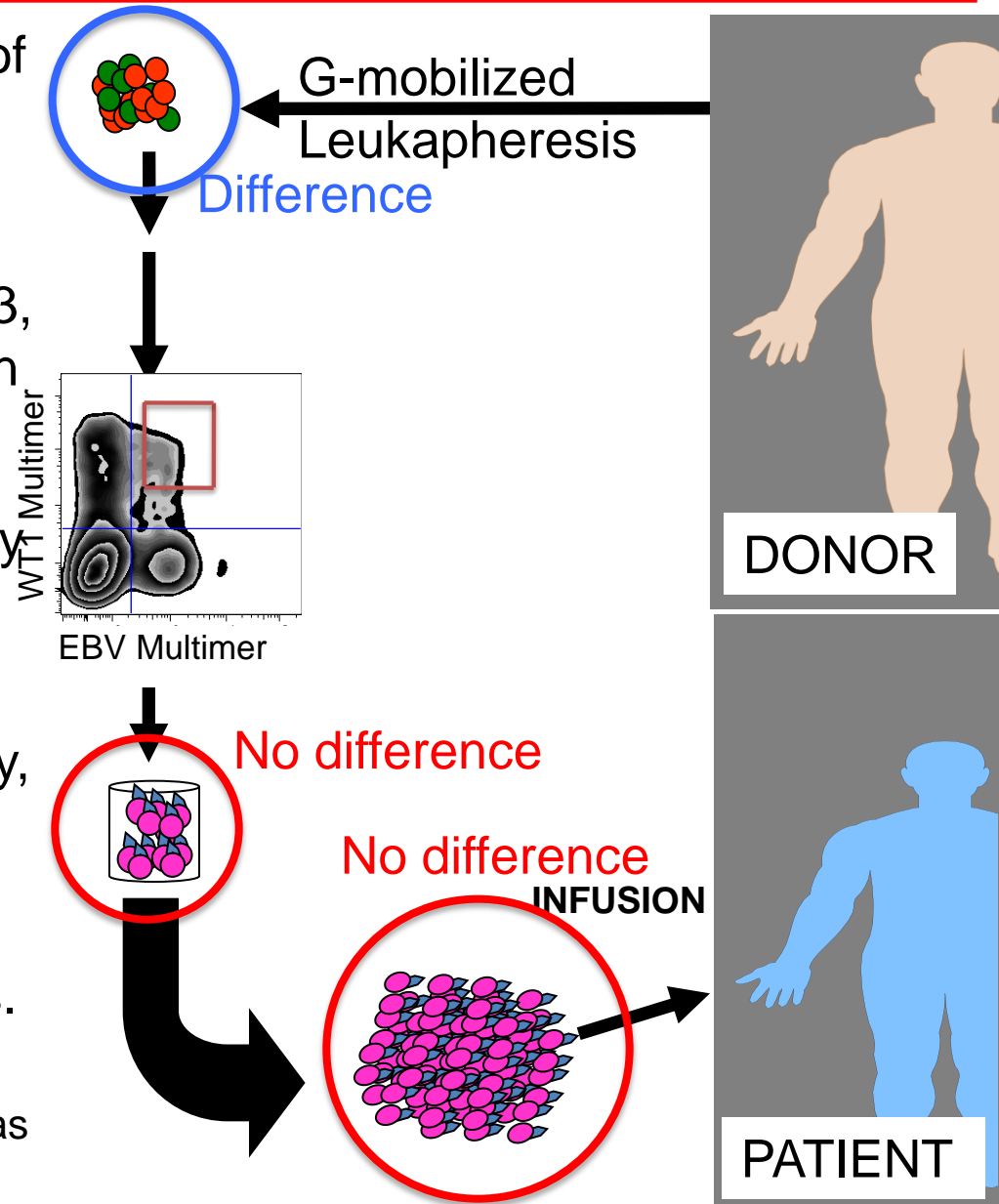
Assessment of persistence after infusions: EBV-Specific CTL

6 of 7 patients who could be followed > 42 days after infusions:



Why are EBV-specific CTL persisting longer with higher frequencies *in vivo* than CMV-specific CTL?

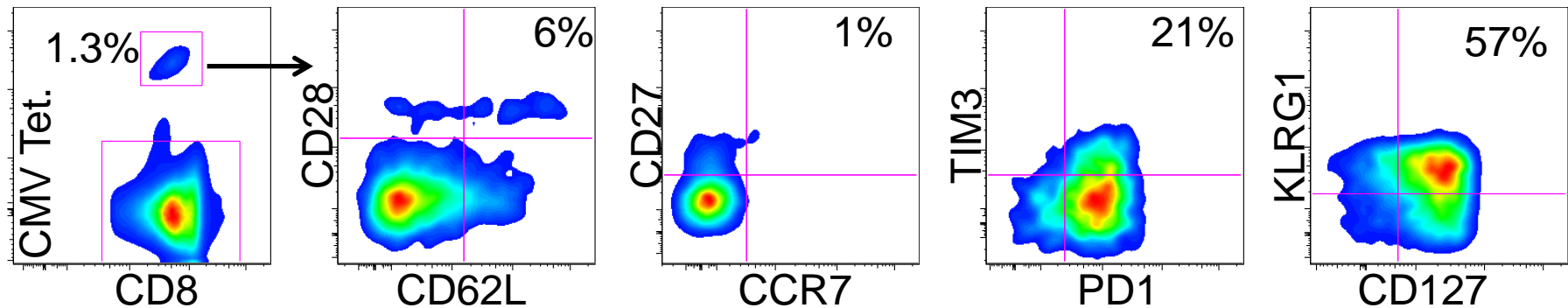
- No difference in the expression of surface markers associated with memory (CD28, CD27, CD127, CD62L, CCR7) or activation/exhaustion (PD1, TIM3, 2B4, KLRG1, CD160) on infusion products.
- CMV infection is characterized by continued low-level replication (Sylwester AW, JEM 2005).
- EBV infection establishes latency, infrequent reactivations.
- EBV-specific cells are present in a 'resting state' in normal donors. Secrete IL-2, less perforin upon cognate Ag recognition (Makedonas G, PLOS pathogens 2010).



Why are EBV-specific CTL persisting longer with higher frequencies *in vivo* than CMV-specific CTL?

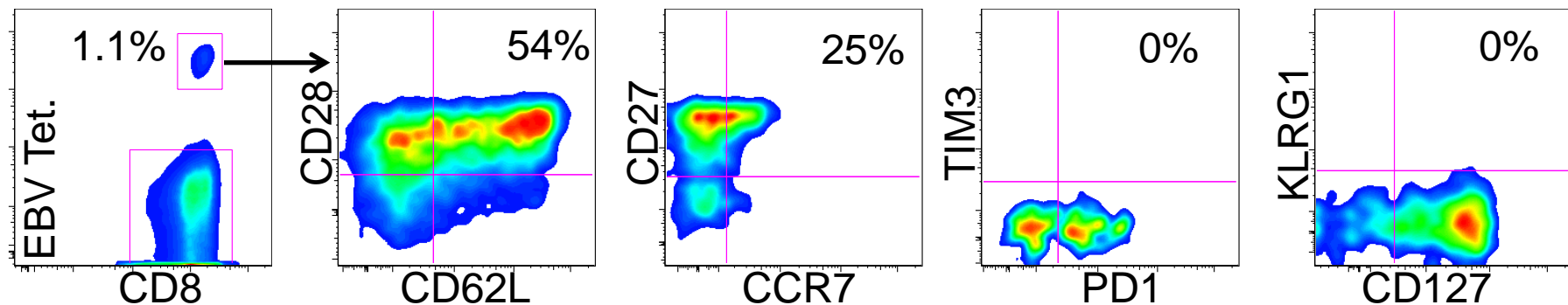
Donor Pt. 3: **CMV-specific substrate**

Gated on Tet.⁺ CD8⁺ T-cells

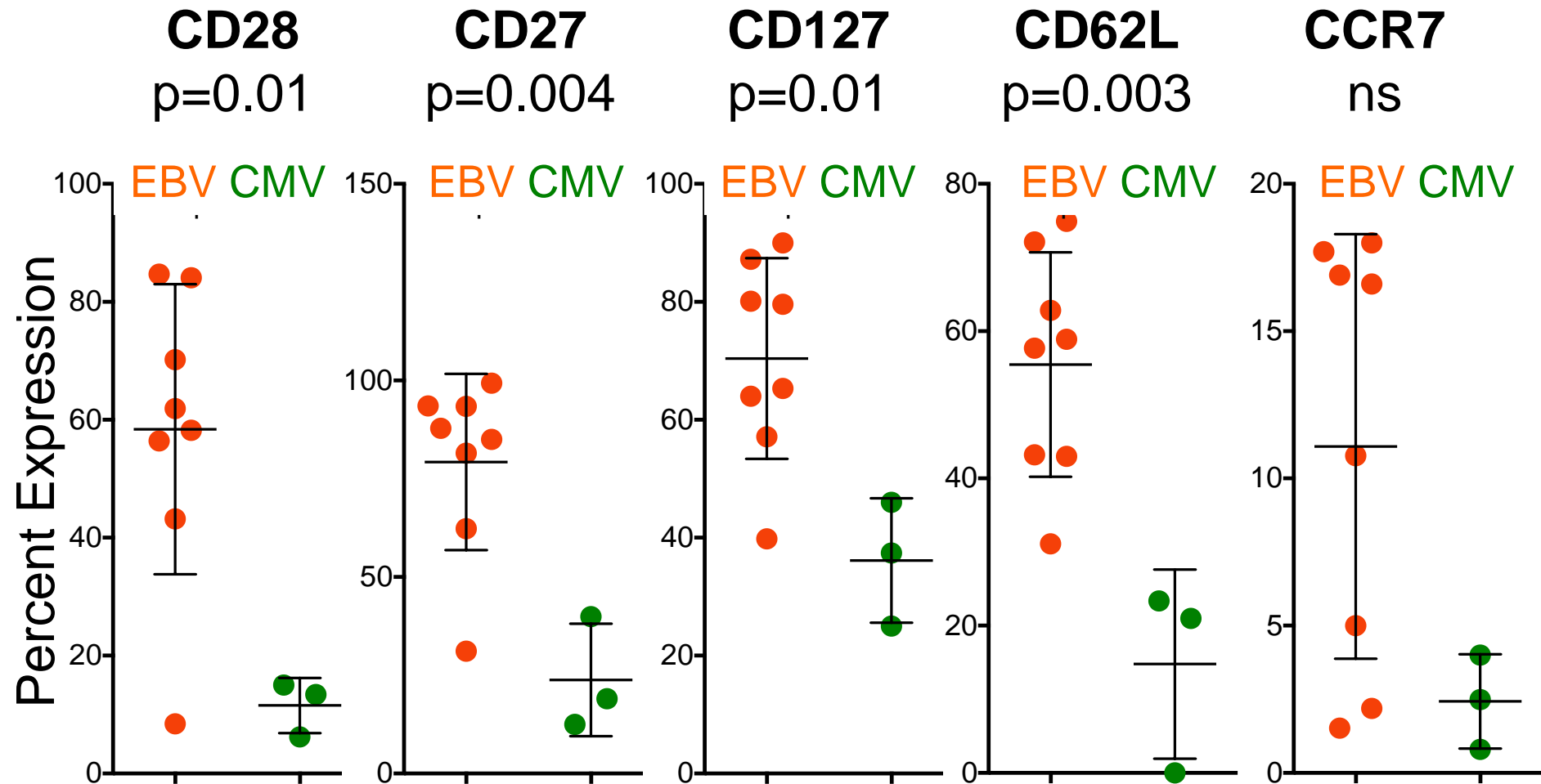


Donor Pt. 4: **EBV-specific substrate**

Gated on Tet.⁺ CD8⁺ T-cells



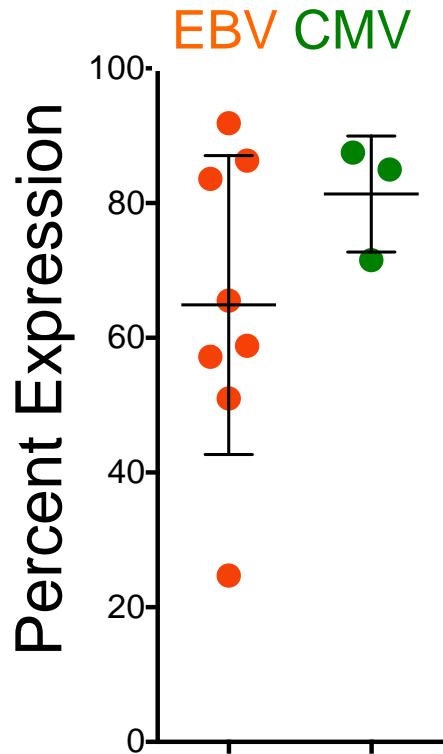
Differential expression of markers associated with memory on donor EBV and CMV-specific T-cells



Differential expression of activation/exhaustion markers on donor EBV and CMV-specific T-cells

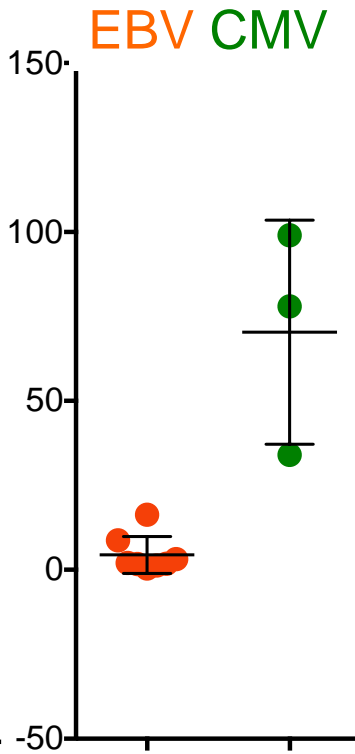
PD1

ns



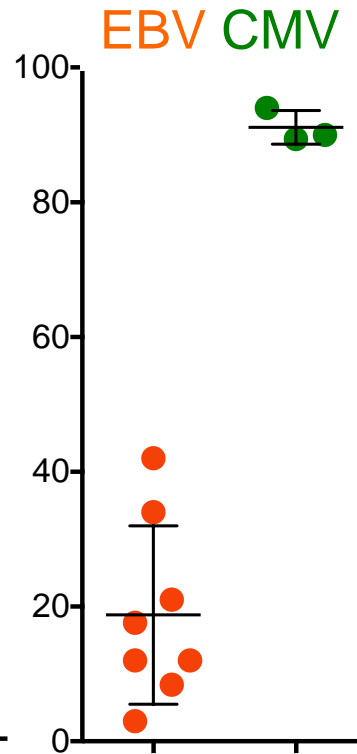
TIM3

$p=0.0002$



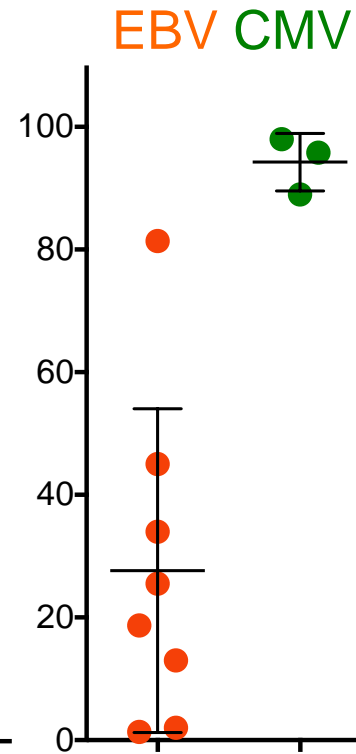
2B4

$p<0.0001$



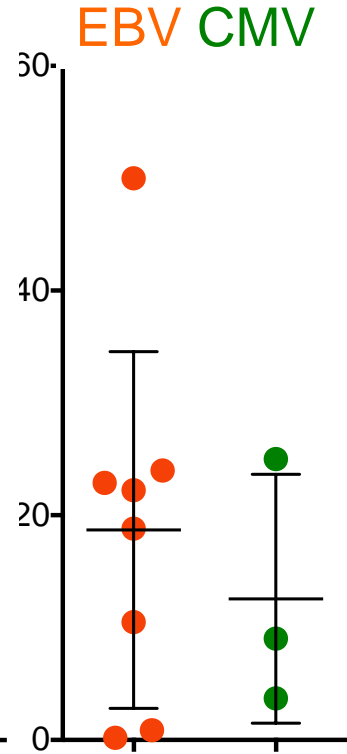
KLRG1

$p=0.003$

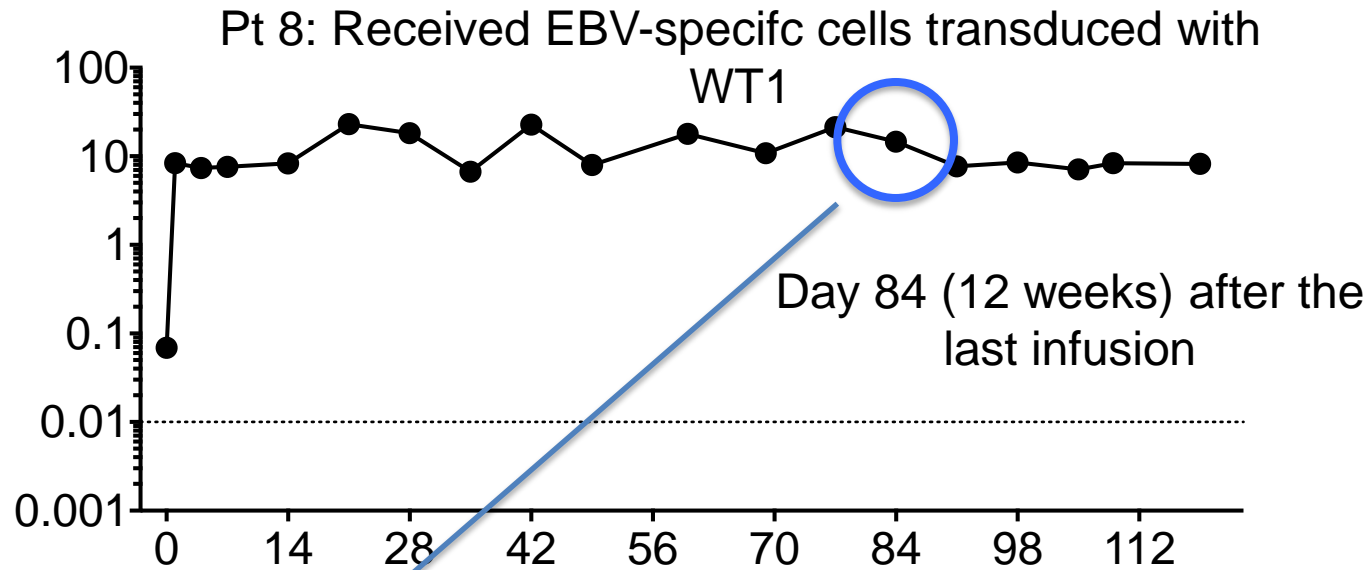


CD160

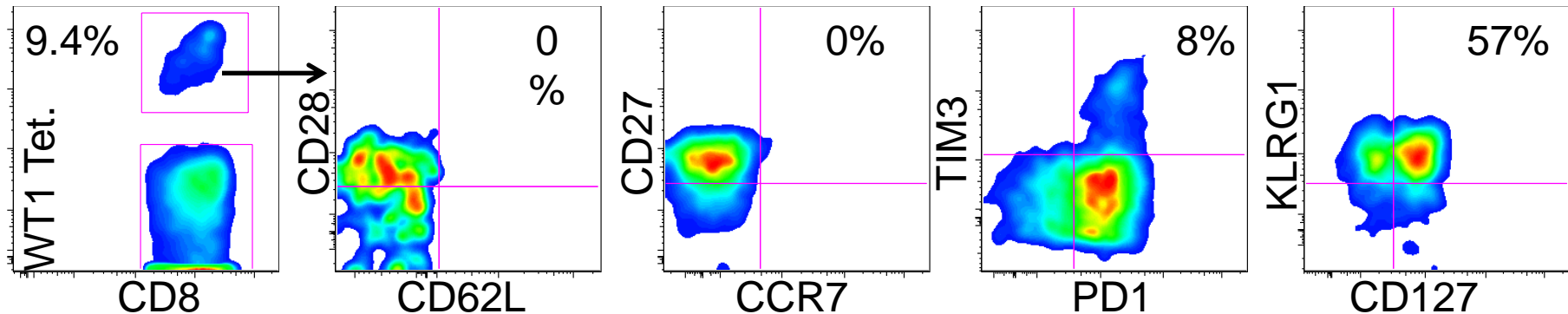
ns



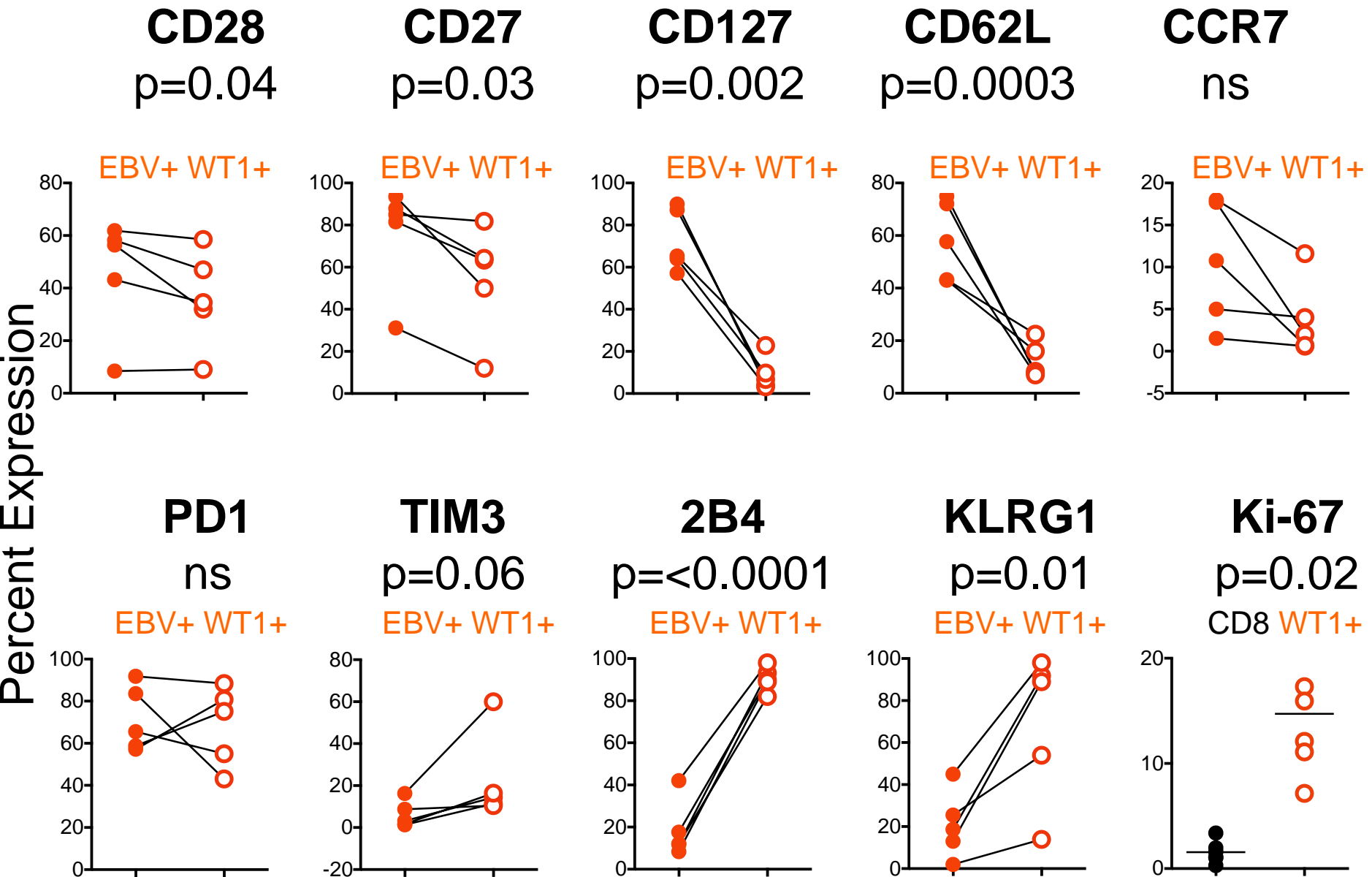
Phenotype of persisting WT1-specific T-cells ~12 weeks after transfer



Gated on Tet.⁺ CD8⁺ T-cells



EBV-transduced WT1-specific T-cells ~12 weeks after transfer have a effector phenotype



Summary and future directions

- Infusion of up to 10^{10} TCR_{C4} transduced virus-specific cells +/- low-dose s.c. IL-2 is safe and **not toxic to tissues expressing low physiologic levels of WT1**.
- Transducing a population of substrate cells which contain cells in a resting state with a '**central memory**' (T_{CM}) phenotype results in prolonged *in vivo* persistence after transfer.

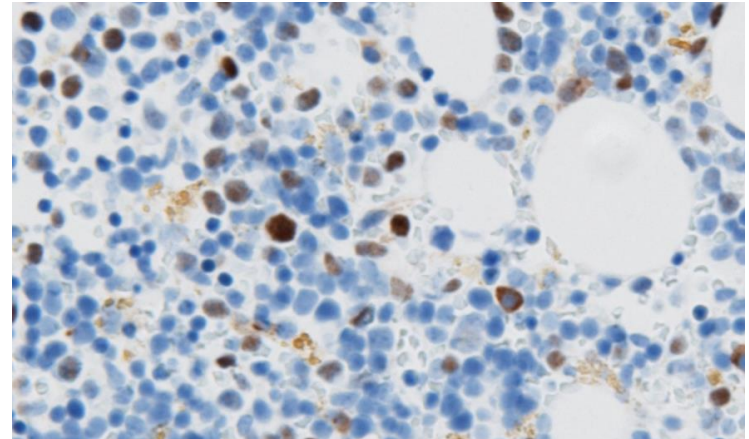
Rationale for targeting Non Small Cell Lung Ca (NSCLC) with WT1-specific cells:

- NSCLC is potentially an immunologically targetable disease (Brahmer H, NEJM 2012; Topalian S, NEJM 2012).
- >96% of NSCLC express WT1 by RT-PCR (Oji K, 2002; Menssen 1995)
- Autologous setting allowing less constraint in the selection of the substrate cells.
- **Prospective comparison between cells generated from the naïve pool compared to cells generated from long-lived memory.**

Cytoplasmic expression of WT1 in Non Small Cell Lung Cancer

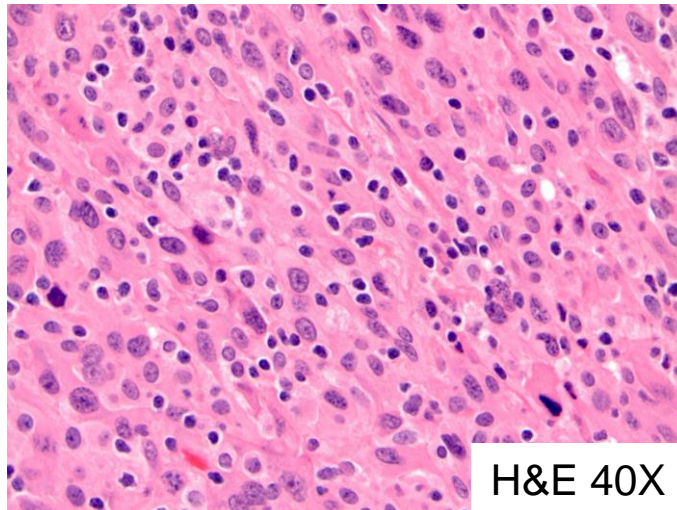
High-risk leukemia: 24%
abnormal CD34+ cells in the
bone marrow:

Anti-WT1 antibody 6F-H2 (Dako) 40X

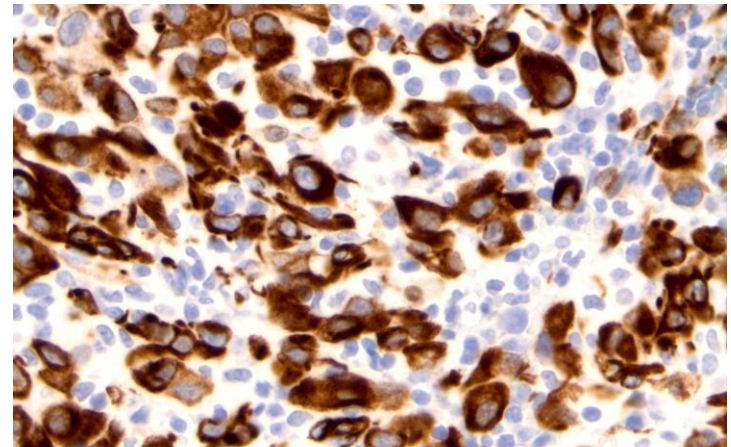


Anti-WT1 antibody 6F-H2 (Dako) 40X

NSCLC:



H&E 40X



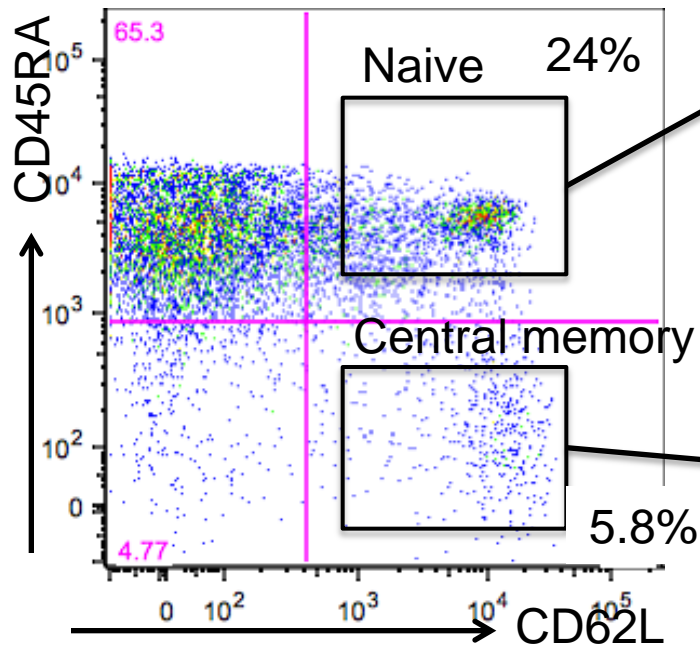
Anti-WT1 antibody 6F-H2 (Dako) 40X

Generation of polyclonal T_{CM} -TCR $_{C4}$ and T_N -TCR $_{C4}$

Non Small Cell Lung Patient 1:

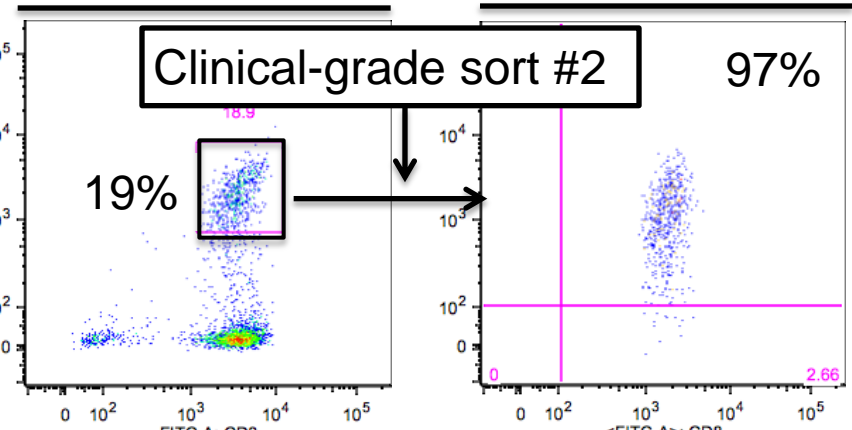
-> Clinical-grade sort #1
-> TCR $_{C4}$ transduction
-> Rapid Expansion Protocol (REP)

Gated on CD8 $^{+}$ T Cells

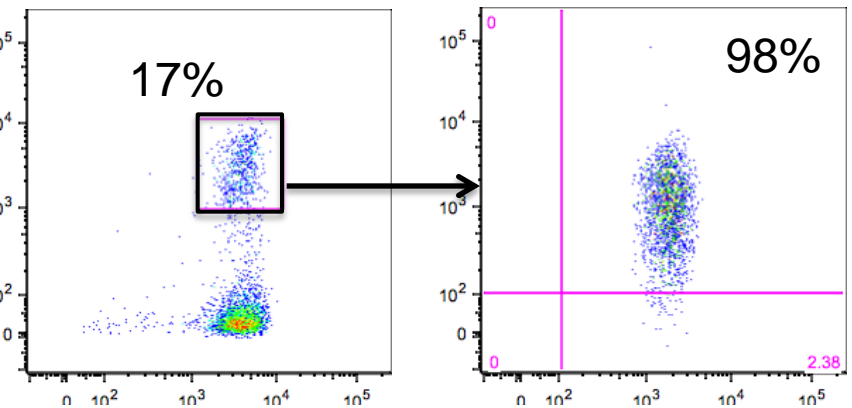


WT1 tetramer

After stimulation and transduction



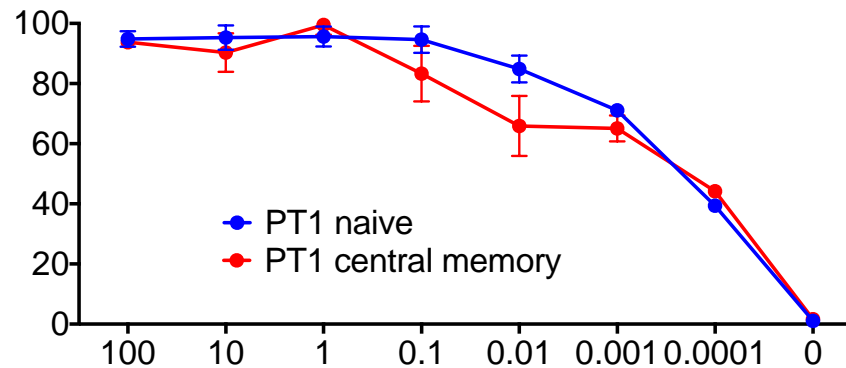
After 2nd sort and REP



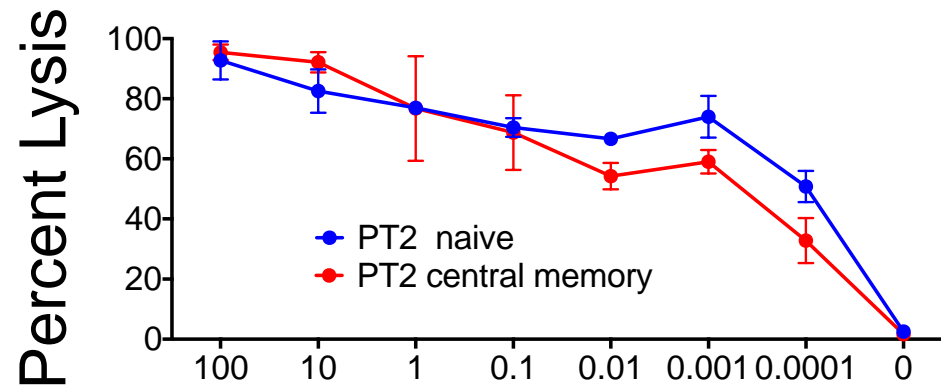
CD8

T_N and T_{CM} TCR_{C4} -transduced substrate cells lyse target cells with identical avidity

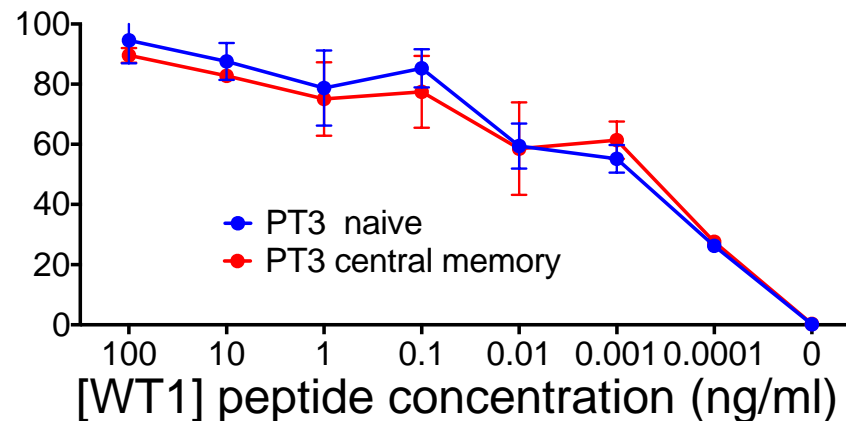
NSCLS Pt1



NSCLS Pt2

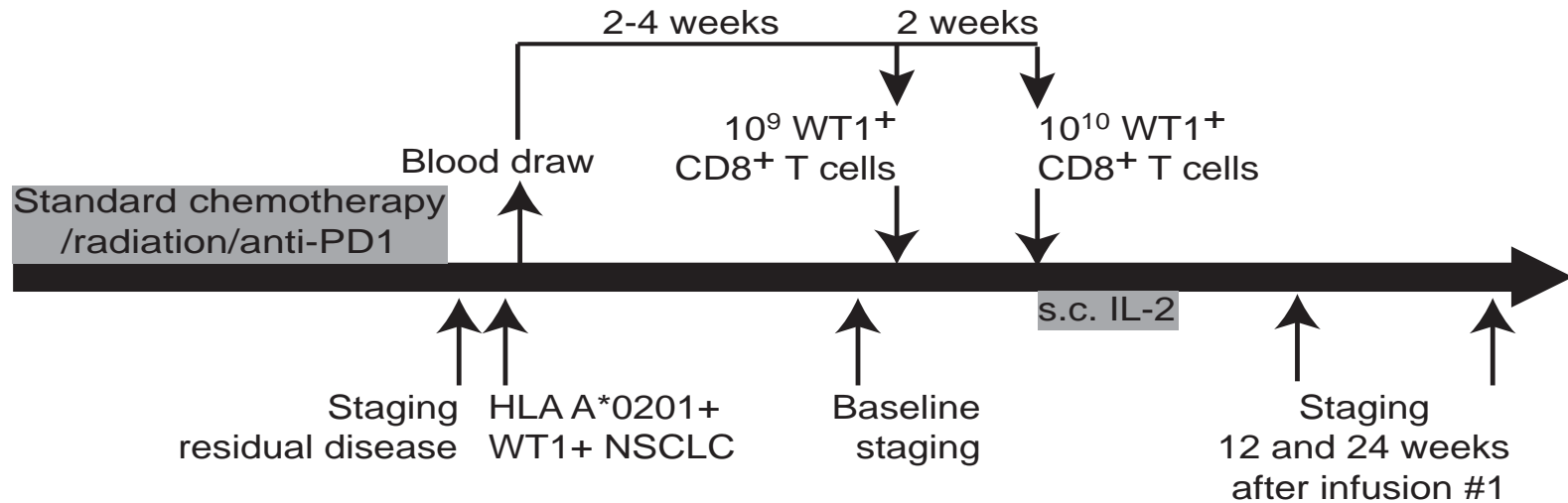


NSCLS Pt3

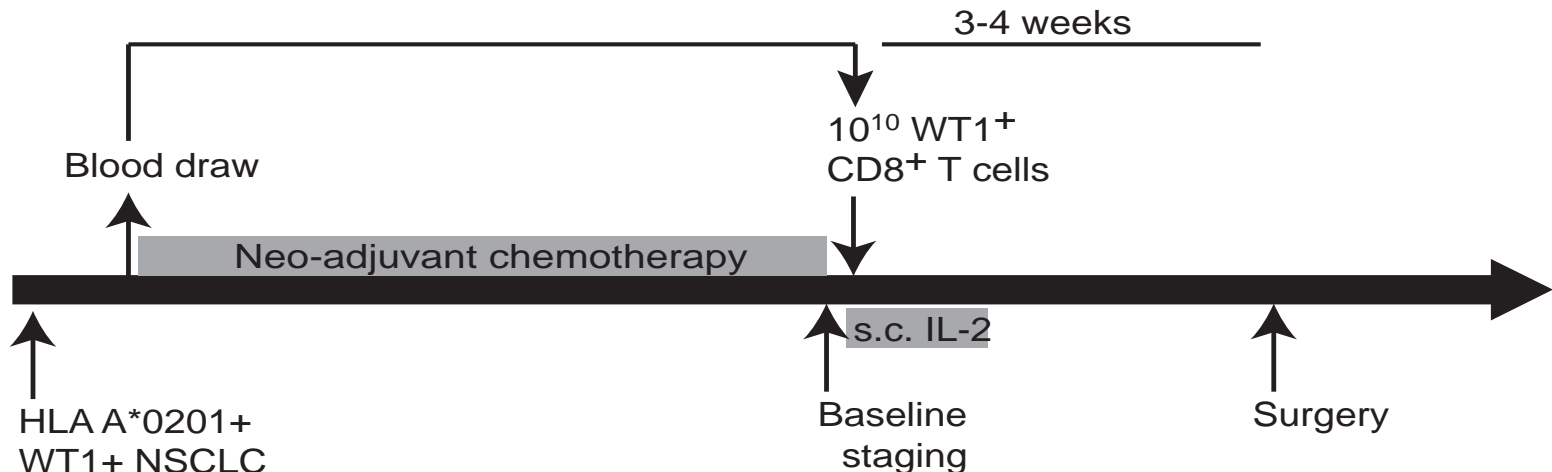


Infusion of TCR_{C4}-transduced T_N and T_{CM} in patients with NSCLC

Arm 1



Arm 2



Acknowledgements

Philip Greenberg

Tom Schmitt

Felecia Wagener

Hieu Nguyen

Gunar Ragnarsson

Colette Chaney

Kieu-Thu Bui

Heather Sloan

Ilana Roberts

Merav Bar

Daniel Egan

Rachel Perret

Natalie Duerkopp

Kendall Shibuya

Jianhong Cao

Daniel Hunter

Sylvia M. Lee

Sheila Gelder

Anna Marie Kane

Ted Gooley

Funding:

NIH NCI P01

NIH NCI K08

Bezos Family
Foundation

Guillot Family
Foundation

Juno Therapeutics