The Development of Cellular Immunotherapy for Patients with Cancer

Steven A. Rosenberg, M.D., PhD. Surgery Branch, National Cancer Institute

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GOAL

Develop effective immunotherapies for patients with metastatic cancer based on the adoptive transfer of immune cells with anti-cancer activity

Cancer Deaths in the U.S.

	New Cases	<u>Deaths</u>
Total	1,688,780	600,920
Solid cancers	1,515,870	542,620
Hematologic	172,910	58,300

1 in 2-3 Americans will develop an invasive cancer

1 in 5 Americans will die of cancer

The development of effective immunotherapies for patients with metastatic epithelial solid cancers that cannot be cured by any available treatment and result in 90% of cancer deaths.

Summary of Cell Transfer Protocols for the Treatment of Patients with Metastatic Melanoma* (median f/u over 8 years)

Total				PR				CR	OR_
			num	ber	of p	atie	nts (d	luration in months)	
194			e	60 (3	1%)			46 (24%)	106 (55%)
84,	71+	·,70+	·,63+	·,58+	·,55⊦	⊦,51+	⊦,37,	152+,142+,137+,136+,136+,	
36,	28,	25,	22,	21,	19,	19+	,14,	134+,132+,123+,121+,118+,	
14,	14,	14,	13,	12+	,11,	11,	11,	114+,113+,113+,112+,110+,	
10,	10,	9,	9,	9,	9,	8,	8,	110+,108+, 64+, 63+, 63+,	
7,	7,	7,	7,	7,	6,	6,	6,	63+, 62+, 62+, 62+, 62+,	
6,	6,	6,	5,	5,	5,	5,	5,	61+, 61+, 60+, 60+, 60+,	
4,	4,	4,	4,	4,	4,	3,	3,	60+, 59+, 58+, 57+, 54+,	
3,	3,	3,	2					53+, 53+, 50+, 45+, 45+,	
								43+, 39+, 38+, 27, 19,	
								14+	

*from four trials (5 groups) using different lymphodepleting regimens

(44 of 46 Complete Responders ongoing from 14 to 152 months) (44 of 46 Complete Responders received a single treatment)



⁽Median followup: 8.6 years)

Question

What are the phenotypic characteristics of the cells that mediate cancer regression in vivo?

Approach:

High dimensional single cell transcriptome analysis of up to 10,000 cells

(tSNE analysis: t-distributed stochastic neighbor embedding)

Pilot study of 16 melanoma Rx1 I.P samples by CyTOF



Gated on CD3+ CD45+ 22 Clusters were defined

(Sri Krishna, Frank Lowery)

CyTOF analyses of 16 pilot Rx1 I.P. samples indicated that Cluster 1 is significantly higher in CR Rx1 relative to NR Rx1





Cluster 1 was highly enriched in CD39neg,CD69neg cells.

(Sri Krishna, Frank Lowery)

CD39neg,CD69neg "stem-like" lymphocytes appear to be the effector cells responsible for melanoma treatment



(Sr Krishna, Frank Lowery



CD8+ DN T-cells are curative in murine tumor ACT models

12

Question

What do TIL recognize that enables the in vivo destruction of the last melanoma cell?

Specific cancer regression in the absence of offtumor on-target, toxicities in patients led us to explore the role of specific cancer mutations as the targets of TIL.

Mining the Cancer Exome to Identify Immunogenic Cancer Mutations

For a mutation product to be a cancer antigen it has to:

- 1) be processed intracellularly into a 9-11 amino acid peptide
- 2) the peptide must fit and be presented in the groove of on one of the patient's surface MHC molecules

Thus, only rare mutations will be antigenic.

Screen all Cancer Mutations to Test their Immunogenicity

Whole exome sequencing and RNA-Seq to identify all cancer mutations

Introduce every cancer mutation into the patient's antigen presenting cells (APCs) as a 25mer with the mutated amino acid in the middle

Coculture APCs with patient TIL that mediated cancer regression

If TIL recognize the mutation interferon-gamma is secreted thus identifying the cancer antigen

<u>Advantages</u> No need to predict peptide binding to MHC. All candidate peptides and all MHC loci are included in the screen. No tumor cell line necessary.

Immunogenic Mutations in Patients with Metastatic Melanoma

Patients evaluate d	Median	Total	Screened	Immunogenic Epitopes
(#)		(# of mutations)	(of total)	(#)
76	318	44381	13400 (30%)	180

Patients with mutation-reactive T cells in TIL: 55/76 = 72%

Immunogenic mutations (of # screened): 180/13400 = 1.3%

8% CD4 92% CD8

All neoantigens were unique, none shared

Immunogenic Mutations in Patients with Gastrointestinal Cancers

Patients Evaluated	Patients with Neoantigens	Median	Total	Screened	Immunogenic
(number)		(number o	f mutations)		
130	104/80%	123	20046	15256 (76%)	210

Immunogenic mutations of number screened: 210/15256 = 1.3%

47% CD8 53% CD4

All neoantigens were unique except 2 patients shared the same KRAS mutation restricted by Cw*0802

Updated 10/8/20

Mutated Antigens Recognized by TIL from 195 Patients with Epithelial Cancers

Cancer		# of patients screened	# of patients with neoantigen reactivity	Total # of neoantigens recognized
Colorectal		95	80 (84%)	166
Anal		1	1 (100%)	1
Cholangiocarcinoma		14	10 (71%)	17
Pancreatic		11	7 (64%)	11
Esophageal		5	4 (80%)	7
Endometrial		3	3 (100%)	4
Breast		43	29 (67%)	100
NSCLC		11	8 (73%)	34
Ovarian		7	6 (86%)	16
Stomach		4	2 (50%)	6
Prostate		1	1 (100%)	1
	Total	195	151 (77%)	363

All neoantigens were unique except for 2 KRAS antigens.

An advantage of targeting mutations is its applicability to target multiple cancer types.

Responses in Patients with Chemorefractory Metastatic Solid Epithelial Cancers

TIL0/200Selected TIL3/2512 %Selected TIL + Pembro6/2623 %

45 y.o. female with metastatic cholangiocarcinoma

- 12/2009 Right hepatectomy for cholangiocarcinoma
 - 4/2010 Multiple lung and liver metastases Received cisplatin and gemcitabine: PD
 - 5/2011 Taxotere chemotherapy: PD in lung and liver
 - 3/2012 Unselected TIL from resected lung lesion infused; PD
- 10/2013 TMG approach with cells selected to target a unique ERBB2IP cancer mutation (of 26 mutations)

(Tran et al, Science 344:641-5, 2014)



J.A. 51 year old female with metastatic breast cancer

2003	Localized Ductal Carcinoma in Situ; underwent mastectomy		
Aug. 2013	ER+, PR+ invasive breast cancer metastatic to mu groups, chest wall, bone	ultiple nodal	
Sept. 2013	Pacitaxel chemotherapy	Progressed	
Feb. 2014	Arimidex	Progressed	
Sept. 2014	Xeloda chemotherapy	Progressed	
Oct. 2014	Navelbine chemotherapy	Progressed	
Nov. 2014	Taxotere, Adriamycin, Cytoxan chemotherapy	Progressed	
Jan. 2015	Lucitanib (TKI inhibitor)	Progressed	
Sept. 2015	Everolimus (mTOR inhibitor)	Progressed	
Dec. 2015	NCI for cell transfer immunotherapy targeting mu by her cancer (62 mutations) Received 80e9 cells	itations expressed	
	She is now in an ongoing complete response of n wall, and liver metastases 58 months after treat	nultiple nodal, chest tment	

J.A.: ACT using autologous lymphocytes targeting somatic mutations



Pre-Treatment

14 Months

Apparently Random Somatic Mutations wereTargeted in this Patient with Metastatic Breast Cancer

SL3A2: 4F2 cell-surface heavy chain

Function: Required for the function of light chain amino-acid transporters

KIA0368: Proteasome-associated protein ECM29 homolog Function: Adapter/scaffolding protein that binds to the 265 proteasome

CADPS2: Calcium-dependent secretion aviator 2 Calcium-binding protein Function: Involved in exocytosis of vesicles filled with neurotransmitters and neuropeptides

CTSB: Cathepsin B. Thiol protease Function: Which is believed to participate in intracellular degradation and turnover of proteins

23% of infused cells contained neoantigen reactivity. All 8 neoantigen TCR present in PBL at 6 weeks.

Patient C.R. Treated with anti-Kras Reactive Lymphocytes

49 y.o. female with metastatic colon cancer

- 9/5/13 Sigmoid colectomy, partial cystectomy Multiple lung metastases
- 5/14/14 Radiotherapy to bladder suture line
- 9/13/14 FOLFOX chemotherapy: PD
- 3/29/15 Two lung metastases resected for TIL
- 7/1/15 TMG approach to target unique cancer mutations (61 somatic mutations)

Response after infusion with KRAS^{G12D}-reactive TIL



- 6/7 lesions regressed at 9 months post ACT
- 1 lesion (#3) progressed at 9 months; excised; patient NED over 5 years after treatment²⁷

RNA analysis of freshly isolated TIL from metastatic cancer



⁽Frank Lowery, Sri Krishna)

Recognition of random somatic mutations is the <u>"final common pathway"</u> explaining cancer regression from most immunotherapies for solid cancers.

IL-2 anti-CTLA4 anti-PD1 anti-CD40 Tumor infiltrating lymphocytes Any intracellular protein can potentially be a "cancer antigen" if mutated and processed intracellularly to a peptide that can bind to the autologous MHC.

(About 1 in 70 mutated neoepitopes are neoantigens.)

Bad news: Treatment will be highly individualized and thus complex. Good news: Virtually <u>all</u> cancer patients are potentially eligible. **Tumor suppressor gene**

50% of human cancers contain a p53 mutation

Most frequently mutated gene in human cancers

Mutations occur throughout the gene but there are up to 10 major "hot spots"

TCRs available for the treatment of patients with p53 mutations

usi	ng "o	ff-the	shelf" ar	nti p53	TCRs
TCR source	p53 mutation	mutation frequency (%) *	HLA restriction	HLA frequency (%)†	Potentially treatable patient (%)
4141; 4196	R175H	4.406	A*02:01	47.40	2.088
4259	Y220C	1.521	A*02:01	47.40	0.721
4273	R248W	2.527	DPB1*02:01	27.30	0.690
4149; 4343	Y220C	1.521	DRB3*02:02	32.80	0.499
4127	G245S	1.494	DRB3*02:02	32.80	0.490
4285	R175H	4.406	DRB1*13:01	10.00	0.441
4259	Y220C	1.521	DRB1*04:01	17.30	0.263
4304	M237I	0.411	DRB1*01:01	20.00	0.206
4266	R248W	2.527	A*68:01	2.80	0.071
4324	T211I	0.076	C*06:02	11.20	0.009
4293	Y236S	0.018	DRB3*02:01	0.33	0.006
4350	L111R	0.02	DRB1* 08:03	7-20% in Chinese populations	0.004
4350	L111R	0.02	A*11:01	14.00	0.003
4114	C135R	0.08	Class II	na	na
Total					5.5%

(P. Kim)

iTCR THERAPY OF BREAST CANCER: PATIENT M.K.

48 yo female with breast cancer

Feb. 2012	Lumpectomy, I.n., dissection, local radiotherap ER ⁻ , PR ⁻ , Her2 ⁺ & breast cancer	y & tamoxifen for
Sept. 2014	Local recurrence Doxataxel, Carboplatin, Herceptin	Progressed
March 2015	Navelbine, Herceptin, Pertuzumeb	Progressed
Aug. 2015	Kodcyla	Progressed
July 2016	Xeloda, Herceptin	Progressed
Feb. 2017	Lapatimid, Herceptin	Progressed
Dec. 2017	Ibrance, Anastrazole, Herceptin	Progressed
Dec. 2018	Doxil	
May 2019	Taxol, Herceptin	
Sept. 2019	NCI Pericardial window for tamponade ACT with 5.3x10 ¹⁰ (p53 cells)	

90% response: Recurred at 6 months

Breast cancer: Treatment with anti-p53 (R175H) TCR



PRE

Breast cancer: Treatment with allogeneic anti-p53 (R175H) TCR

9-18-19 Pre

1-6-20 Day 60



Blueprint for Cancer Immunotherapy Directed Against the Common Epithelial Cancers

Target the immunogenic somatic mutations unique to the autologous patient's cancer.

Raise a library of T-cell receptors against shared cancer mutations (e.g. Kras, p53)

Cell transfer therapy can mediate durable regressions in patients with metastatic cancer refractory to other treatments.

T-cells that recognize unique somatic mutations can be found in TIL and PBL in patients with common epithelial cancers.

Identification and targeting of mutations unique to each cancer or shared mutations such as KRAS or p53 has the potential to extend cell therapy to patients with common epithelial cancers.

- 1. Unique somatic mutations in an intracellular protein
- 2. Mutations in driver oncogenes or tumor suppressor genes that can be shared among patients.

e.g.: Kras p53 PIK3CA

3. Non-mutated proteins on the surface of cells that are not essential for survival and can be recognized by antibodies or specific ligands

e.g.: CD19 (CAR: αCD19 antibody)

Kras protein is a GTPase essential for normal tissue signaling.

Activating mutations are essential steps in the development of many cancers.

Eight "hotspot" mutations in the KRAS gene at amino acid positions 12, 13,

and 61 account for greater than 80% of all KRAS mutations.

Library of TCRs Available for Treatment of Patients with KRAS Mutations

			The stable	Potentially
Mutation	Frequency (%)	HLA HLA Restriction (%)	Frequency (%)	Patients (%)
G12D	35 %	C*0802 A*1101 DRB3*02	8 – 14 % 14 – 30 % 32 %	2.8 % 4.9 – 10.5 % 11.2 %
G12V	24 %	A*1101 C*0102 DRB3*02 DRB1*0701 DRB1*0301 DRB1*0101	14 – 30 % 1.3 – 7.2 % 32 % 26 % 20 % 18 %	3.4 – 10.5 % 1.3 – 7.2 % 11.2 % 6.1 % 4.8 % 4.3 %
G12R	3 %	DQA1*0505 DRB5*01	35 % 35 %	1 % 1 %
TOTAL				29 %

(Noam Levin, 2020)

Immunogenic Mutations in Patients with Metastatic Breast Cancer

Patients evaluate d	Median	Total	Screened	Immunogenic Epitopes
(#)		(# of mutations)	(of total)	(#)
43	119	7936	4722 (60%)	100

Patients with mutation-reactive T cells in TIL: 29/43 = 67%

Immunogenic mutations (of # screened): 100/4722 = 2.1% 74% CD4 26% CD8

All neoantigens were unique, none shared

(updated 10/8/20)

CAR-T cells for the Treatment of Solid Epithelial Cancers

CAR-T cells require the use of monoclonal antibodies (MoAb) that recognize molecules on the cell surface.

MoAb were first described by Kohler and Milstein 45 years ago. (Nature 256:495-7, 1975)).

Two major problems:

- 1) No MoAb have been identified that recognize cell surface molecules <u>unique</u> to epithelial cancers.
- 2) Normal cells expressing the target of MoAb are highly sensitive to destruction.

Blueprint for the identification of mutation-reactive T-cells in common epithelial cancers



Blueprint for the generation of mutation-reactive T-cells in common cancers



Advantages of this approach:

No need to predict peptide binding to MHC.

All candidate peptides and all MHC loci are included in the screen.

No tumor cell line necessary.

Two Critical Issues in the Development of Cancer Immunotherapies

Identify the antigenic targets on the cancer cell

Identify the characteristics of the immune cells that recognize and destroy cancer cells

Potential improvements in targeting of somatic mutations in epithelial cancers

Increase the frequency and number of mutation reactive cells in the infusion product

Enrich the mutation-reactive cells using markers in the fresh cell suspension or upregulated after activation (41BB)

Transduce mutation-reactive TCRs into naïve or CM cells (FDA approval to use a GMP 293GP line to produce transient vectors with minimal testing)

Add anti-PD-1 (reexpressed by infused cells in vivo) or other CPM

Knockout CISH or PD-1 (or other inhibitory molecules) on transferred cells

Vaccinate with mutations recognized by transferred cells

Obtain mutation-reactive TCRs from circulating lymphocytes and mutations from paraffin sections or liquid DNA) – eliminate need for tumor resection

Improve methods to identify multiple mutation targets expressed by tumor (robotics)

ADVANTAGES OF CELL TRANSFER THERAPY

- 1. Administer large numbers of highly selected cells with high avidity for tumor antigens.
- 2. Administer cells activated ex-vivo to exhibit anti-tumor effector function.
- 3. Potentially identify exact cell subpopulations and effector functions required for cancer regression in vivo.
- 4. Manipulate host prior to cell transfer to provide altered environment for transferred cells.

Isolation of ERBB2IP reactive cells



Use enriched ERBB2IP autologous lymphocytes for treatment

Objective response of lung and liver metastases ongoing for over 7years

(Tran et al, Science 344:641-5, 2014)

M.B. Selected TIL



Tumor suppressor gene

50% of human cancers contain a p53 mutation

Most frequently mutated gene in human cancers

Mutations occur throughout the gene but there are up to 10 major "hot spots"

Pt. [4355] Breast cancer course following ACT



CAR-T cells for the Treatment of Solid Epithelial Cancers

CAR-T cells require the use of monoclonal antibodies (MoAb) that recognize molecules on the cell surface.

MoAb were first described by Kohler and Milstein 45 years ago. (Nature 256:495-7, 1975)).

Two major problems:

1) Very few MoAb exist that recognize cell surface molecules <u>unique</u> to cancer (e.g. EGFRvIII).

2) Normal cells expressing the target of MoAb are highly sensitive to destruction.

(Most MoAb in clinical use target growth factors, such as EGFR or Her2, that are expressed on the surface of normal cells.)

Gene signature of CD39- CD69- cells by single cell transcriptome analysis shows enrichment of memory markers

CD8+ CD39+ CD69+ N = 1016 Cells S1PR1 KI67 15 15 CCL3 Upper 2 Quantiles KLF2 IFNG CD69 (TPM) 10 Log₁₀ FDR CDK1 CD8+ CD39- CD69-CD8+ CD39+ CD69+ 0 I AMAVCR2 10 15 5 -1.0 -0.5 0.0 0.5 1.0 ower 2 Quantiles ENTPD1 (TPM) Log Fold Change DN DP CD8+ CD39- CD69-

N = 4420 Cells

- 1. Unique somatic mutations in an intracellular protein
- 2. Mutations in driver oncogenes or tumor suppressor genes that can be shared among patients.

e.g.: Kras p53 PIK3CA

3. Non-mutated proteins on the surface of cells that are not essential for survival and can be recognized by antibodies or specific ligands

e.g.: CD19 (CAR: αCD19 antibody)

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G12R	3 %	DQA1*0505 DRB5*01	35 % 35 %	1 % 1 %
TOTAL				29 %

(Noam Levin, 2020)

A novel method to screen for T cell responses to p53 "hotspot" mutations



Synthesize one TMG (ten 25mers) encoding the top ten p53 "hotspot" mutations.

Synthesize the top 10 25mer peptides.

Coculture patient TIL with TMGs and peptides.

ADVANTAGES OF CELL TRANSFER THERAPY

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- 2. Potentially identify exact cell subpopulations and effector functions required for cancer regression in vivo.
- 3. Manipulate host prior to cell transfer to provide altered environment for transferred cells.

Antigen Recognition by TCRs and CARs



Patient E.K. Treated with Autologous Anti-CD19 CAR T-Cells

48 year old male with follicular non-Hodgkin lymphoma

Aug. 2002	diagnosed with stage IV lymphoma 7 cycles PACE chemotherapy (cisplatin, doxorubicin, cyclophosphamide, etoposide)
April 2004	idiotypic/KLH vaccine (5 doses)
Sept. 2007	ipilimumab
Nov. 2007	6 cycles EPOCH-R chemotherapy (etoposide, predisone, vincristine, cyclophosphamine, rituximab)
May 2009	To NCI for treatment with autologous anti-CD19 CAR transduced T cells

In ongoing progression-free regression over 10 years later.

(Blood 116:3875-86, 2010; 119:2709-20, 2012)



Bone marrow biopsies showed extensive CLL before treatment and nearly absent B-lineage cells after treatment

Before treatment



3 months after treatment





In Patient 8, normal blood B cells were eliminated after CAR-transduced T cell infusion



In contrast, T and NK cell counts rapidly recovered after treatment





Treatment of Patients with Refractory Diffuse Large B-cell Lymphoma

	Objective	Complete Response		
	Response (%)	Total	Öngoing	
Surgery Branch	73%	47%	42%	
Kite Pharma	82%	54%	40%	

In 2012 SB/NCI signed a CRADA to transfer our technology to Kite Pharma. In 2017 Kite received FDA approval. In October 2017 Kite was sold to Gilead Sciences, Inc. for \$11.9 billion. 36 y.o. female with metastatic cervical cancer

- 10/2011 Presented with fungating cervical mass, lung and intraperitoneal metastases
- 11/29/11 Radiation therapy and cisplatin chemotherapy
- 10/06/12 Cancer progressed. She underwent hysterectomy and excision of both ovaries
- 11/2012 toDeveloped liver, lymph node, intra-abdominal1/2013metastases and urinary tract obstruction requiring a stent
- 3/15/13 At NCI/Surgery Branch treated with cell transfer immunotherapy (75 billion of her own tumor infiltrating lymphocytes and IL-2)

Experienced complete regression of all disease including relief of urinary obstruction and remains disease-free over 7 years later.

(Stevanovic et al, Science 356:200, 2017)

Patient S.S. with metastatic cervical cancer treated with cell transfer immunotherapy



NeoTCR signature projection onto T-cell transcriptome maps



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Genes in NeoTCR signature						
CXCL13	LRRN3	ASB2	ENTPD1	GALNT2	TIGIT	=
NSMCE1	HMOX1	GZMB	CLIC3	NELLZ	CXCR6	
AC092580.4	DUSP4	LAG3	NDFIP2	LAYN	PTMS	
GPR25	ACP5	CARS	IL4R	CD9	тох	