

# *Applying basic immunology to treat cancer*

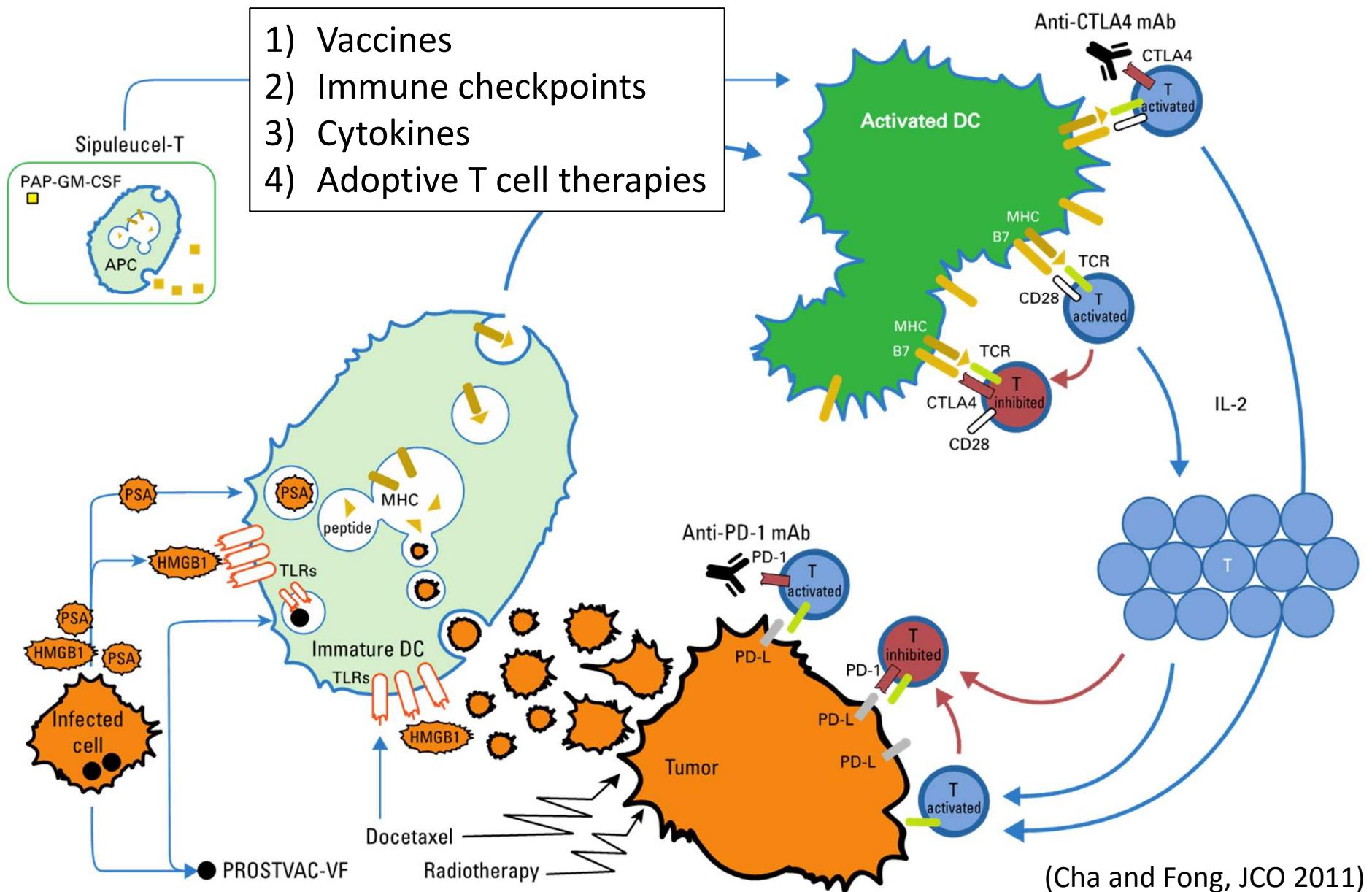
Lawrence Fong, MD



# COI/disclosures

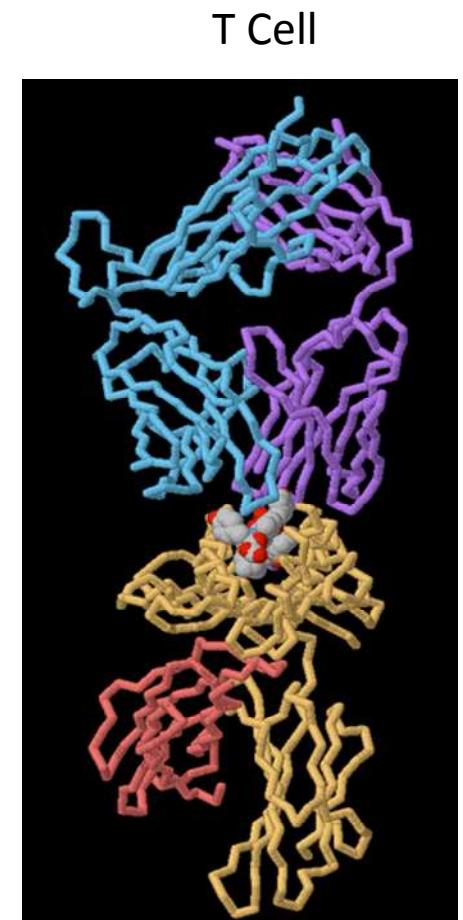
- We have received research support from Dendreon and Bristol Myer Squibb.
- Sipuleucel-T and Ipilimumab will be discussed in disease setting that are off-label and are investigational.

# Approaches to cancer immunotherapy



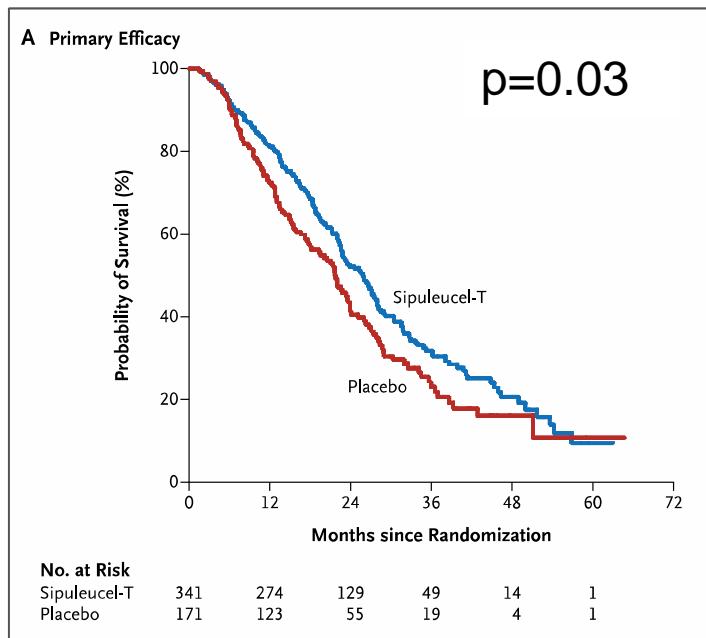
# Candidate tumor antigens

Class	Antigen
Developmental: (cancer/testis)	Mage, Gage, Lage families
	NY-ESO-1
	telomerase
Viral:	EBV, HPV
Overexpressed:	Her2neu, CEA, Muc-1
Differentiation antigens:	Prostate Acid Phos, PSA, PSMA
	tyrosinase, gp100, Mart-1/Melan-A
Tumor specific:	Id, bcr-abl
	p53, ras
	Tumor specific mutations

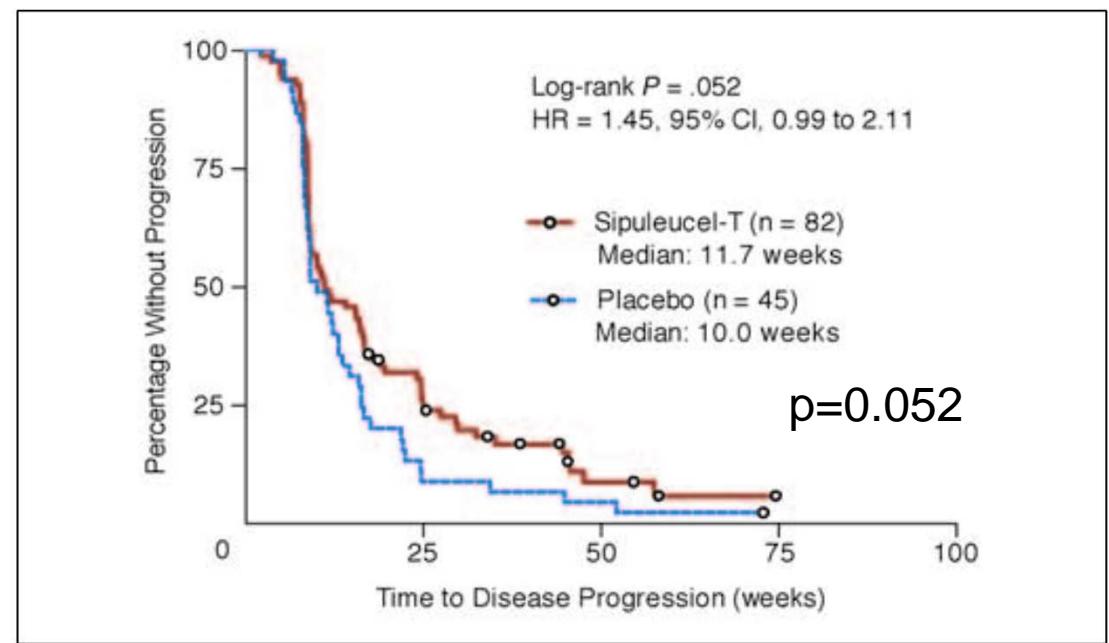


# Sipuleucel-T improves overall survival in metastatic castration resistant prostate cancer (CRPC)

Overall Survival

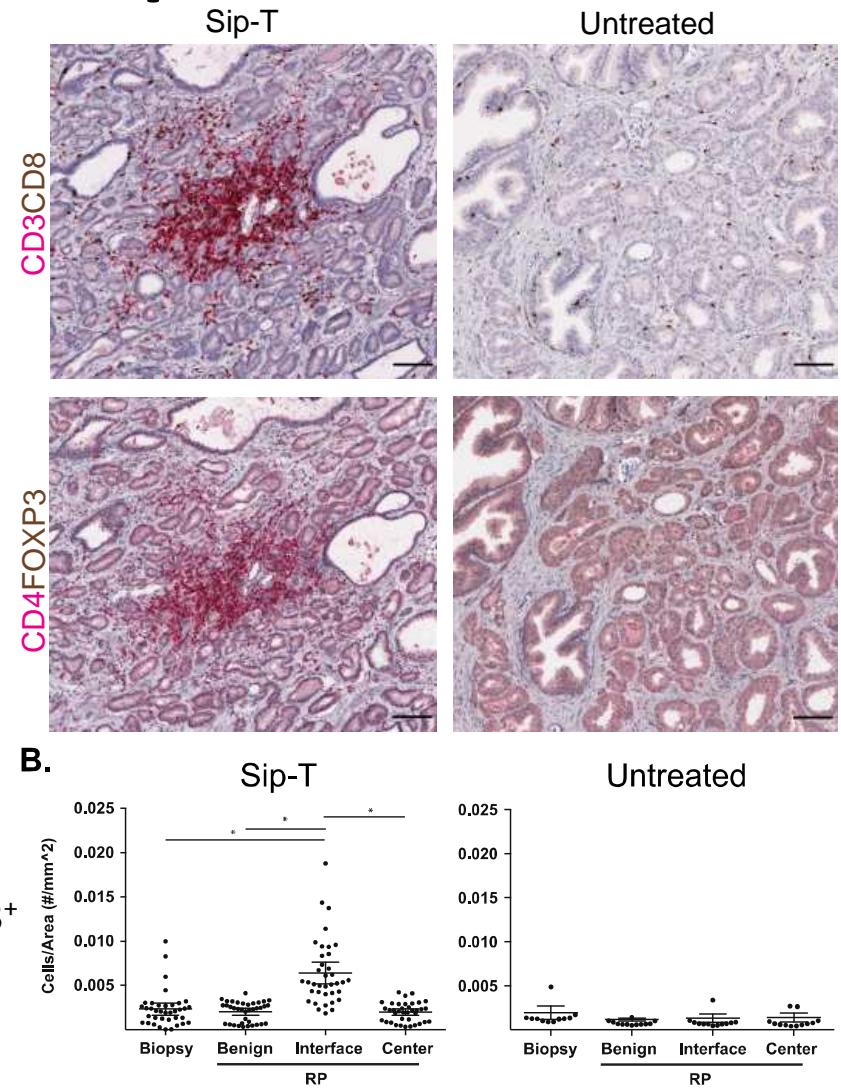
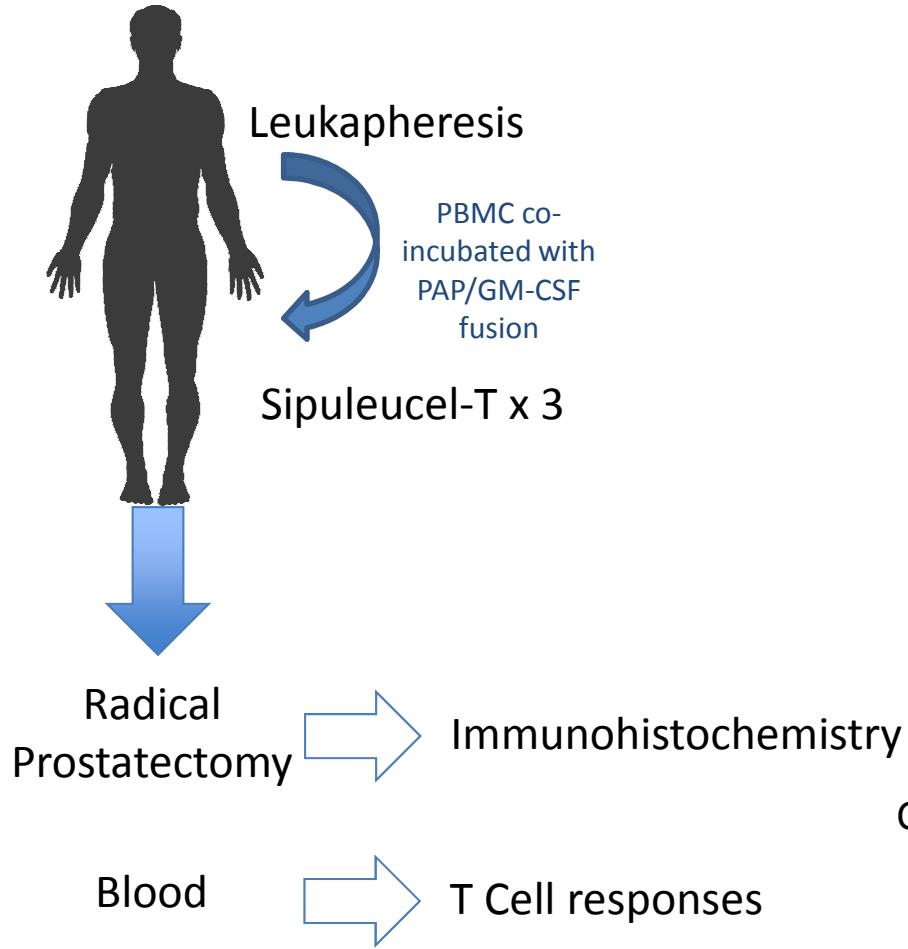


PFS



(Small JCO 2006, Kantoff NEJM 2010)

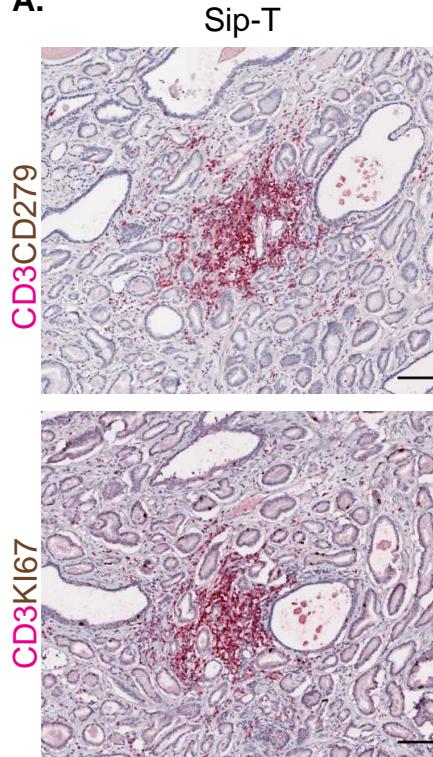
# Sipuleucel-T treatment generates T cell responses in the prostate



(Fong et al, JNCI 2014)

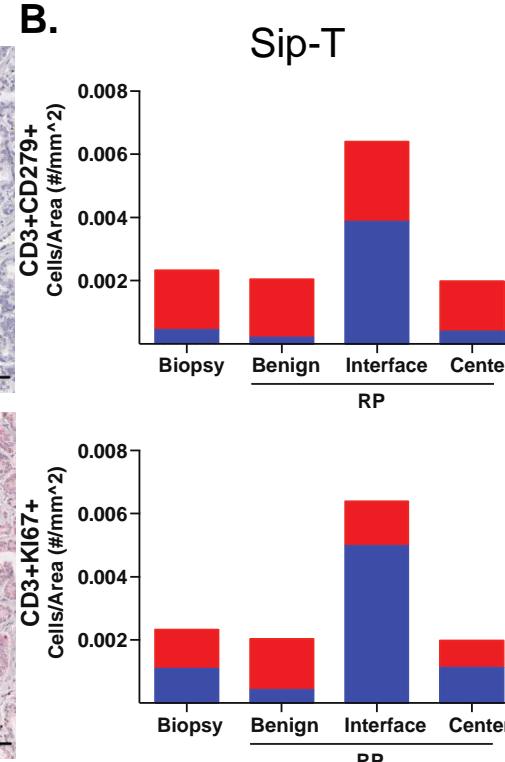
# T cells at the interface express PD-1 and are proliferating

A.



Untreated

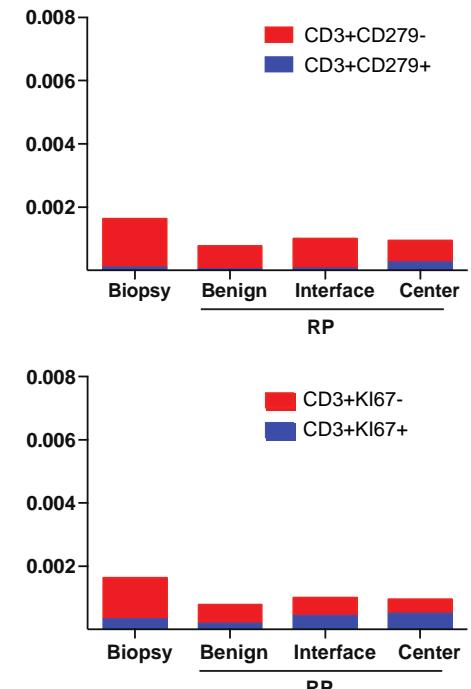
B.



Sip-T

Untreated

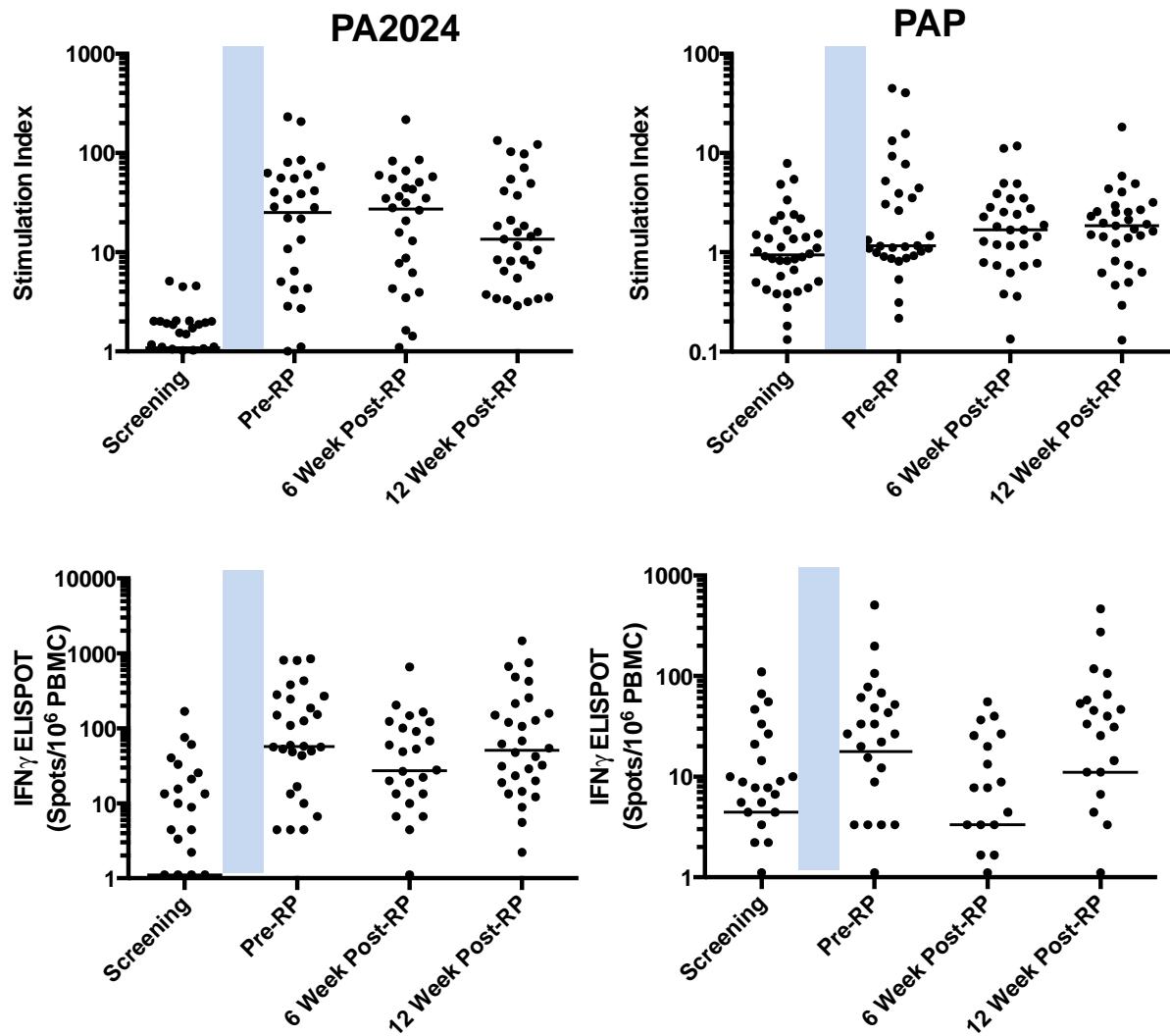
CD3+CD279-  
CD3+CD279+



CD3+KI67-  
CD3+KI67+

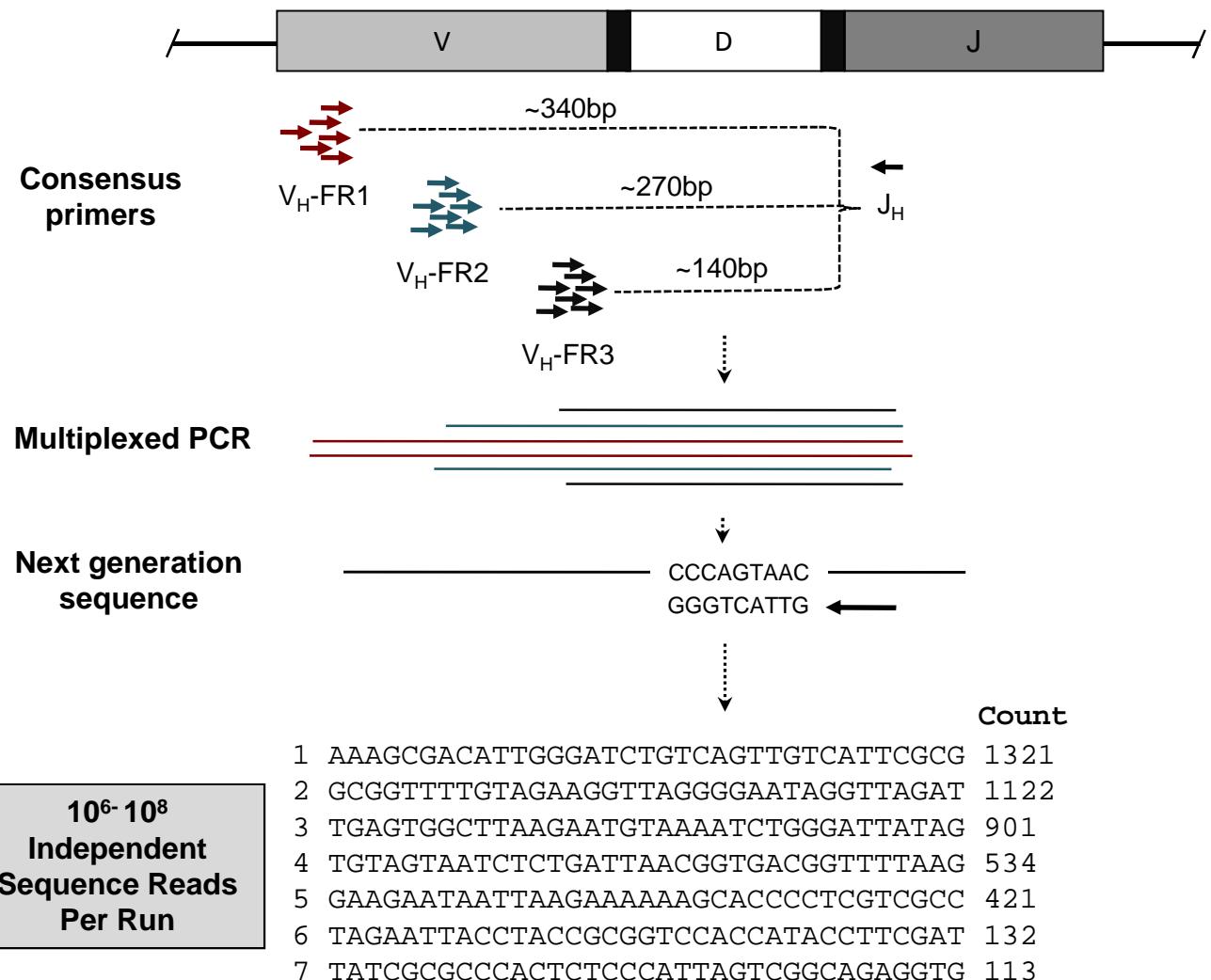
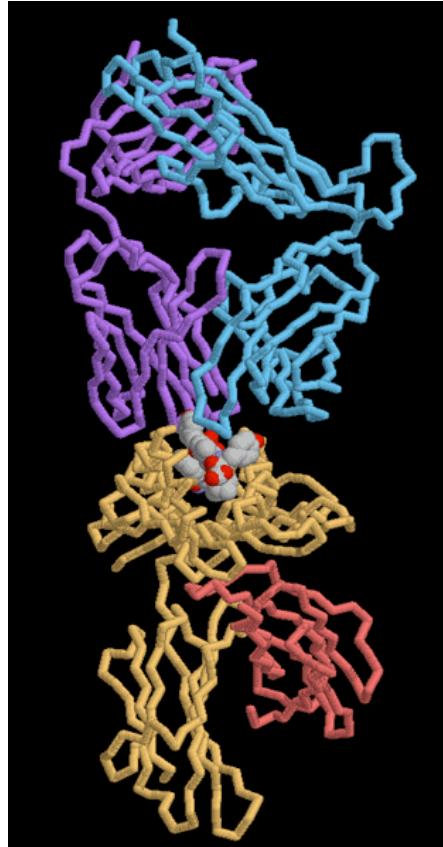
(Fong et al, JNCI 2014)

# Neoadjuvant Sip-T induces circulating T cell responses



(Fong et al., JNCI 2014)

# T cell clonotype tracking by TCR $\beta$ sequencing



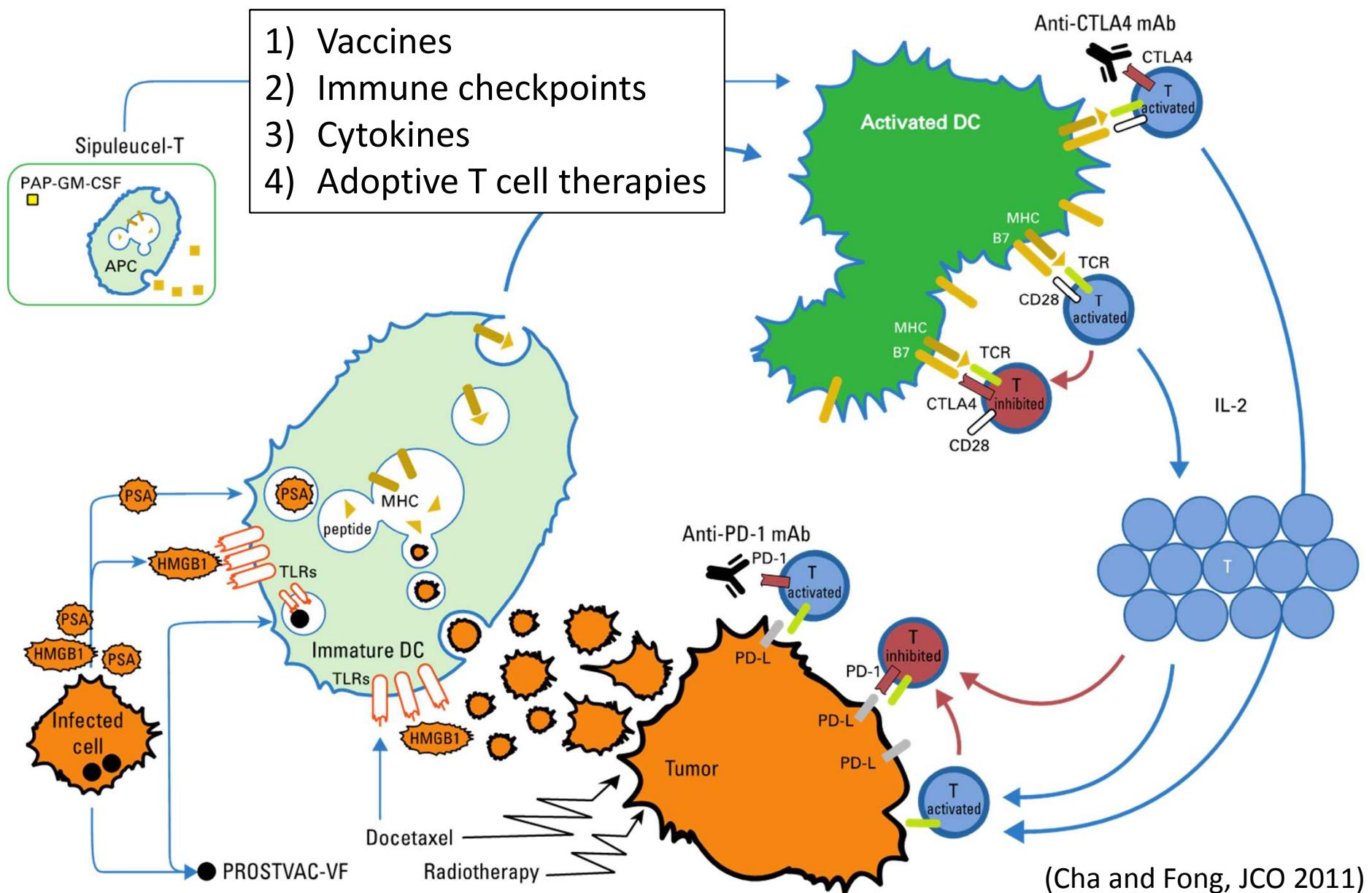
$10^6$  -  $10^8$   
Independent  
Sequence Reads  
Per Run

(adapted from Aaron Logan)

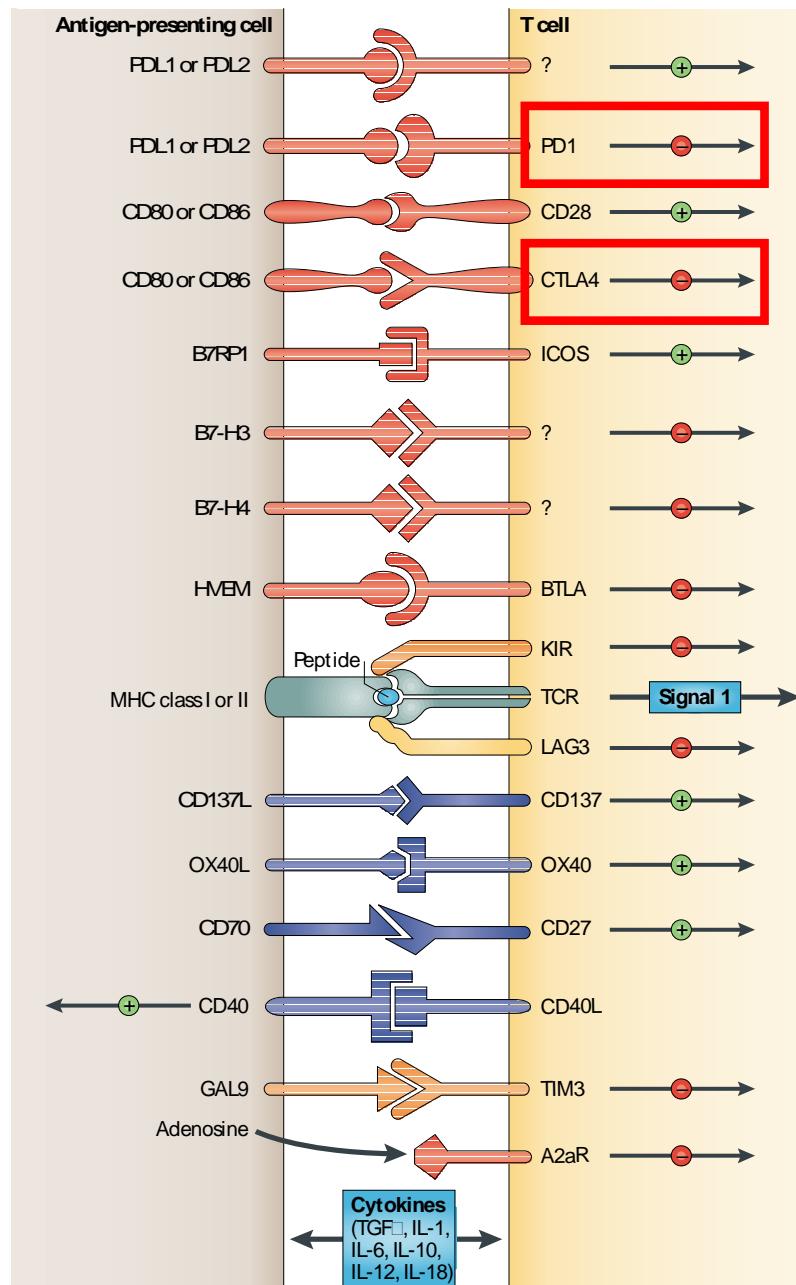
# Conclusions

- Neoadjuvant Sip-T recruits CD3 T cells to the tumor interface.
- Sip-T induces narrowing of the circulating T cell repertoire.
- Many of the circulating T cells induced by treatment can be found in the tumor tissue.
- Sip-T leads to a diversification of the T cells in the prostate.

# Approaches to cancer immunotherapy

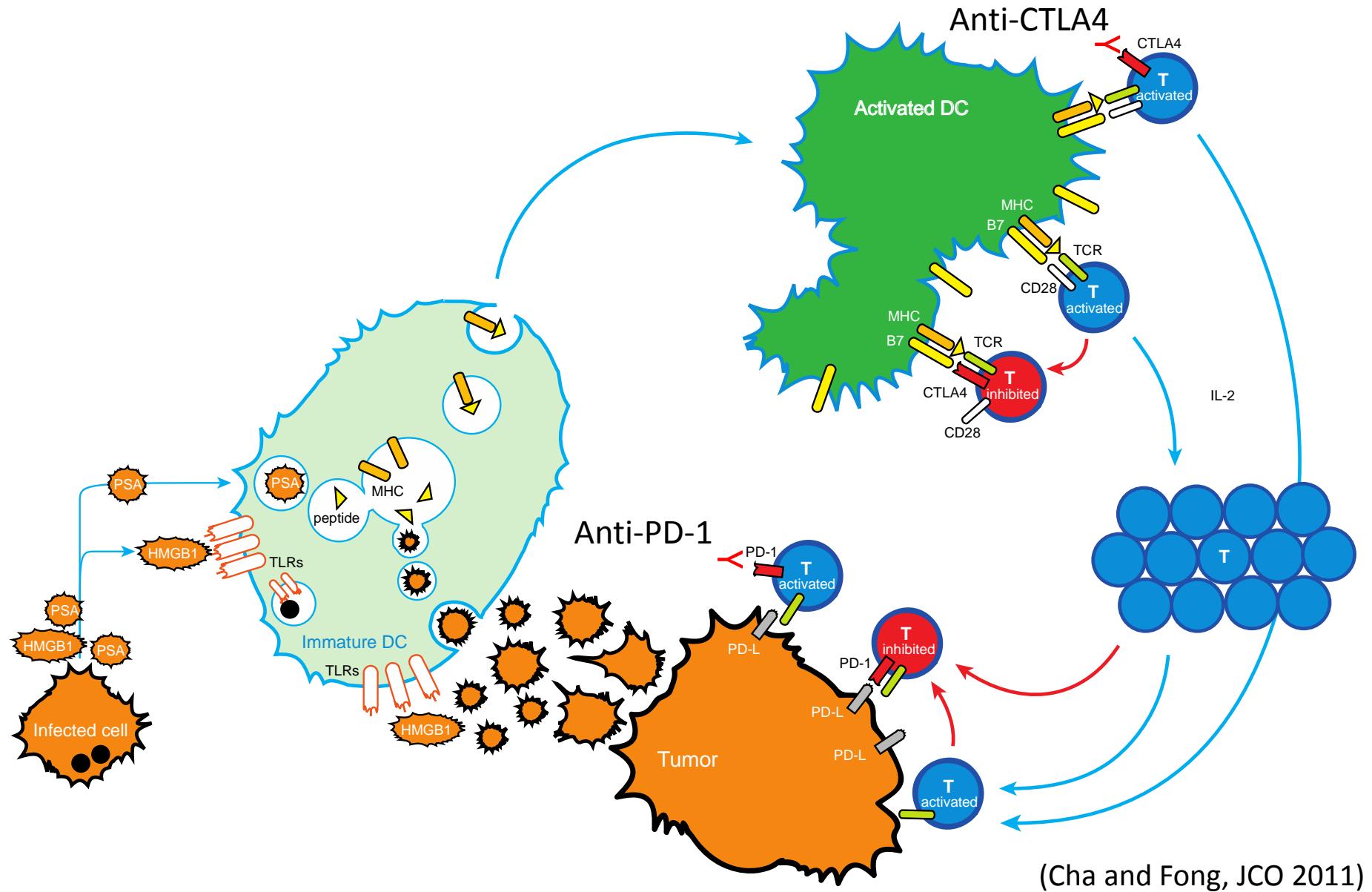


# Co-stimulation and co-inhibition

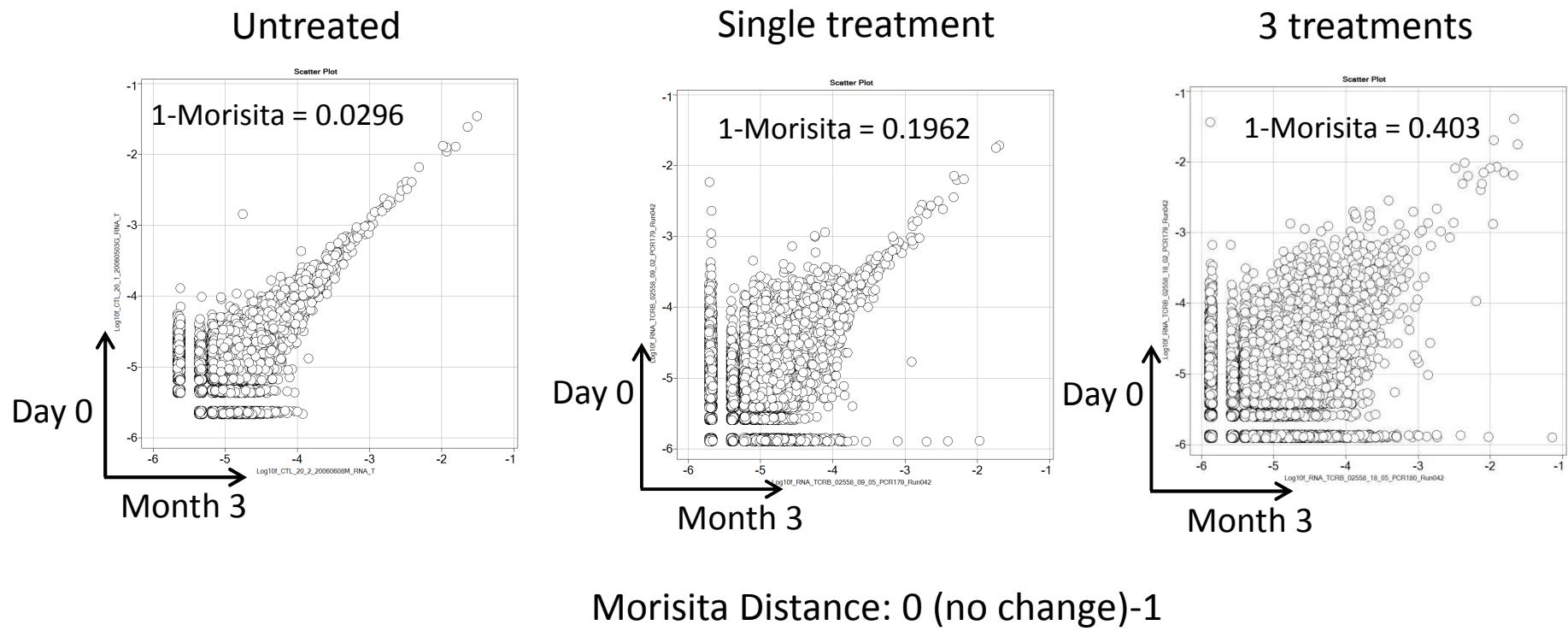


(Pardoll. NRC 2012)

# Checkpoint inhibitors rely on immune priming to endogenous tumor antigen



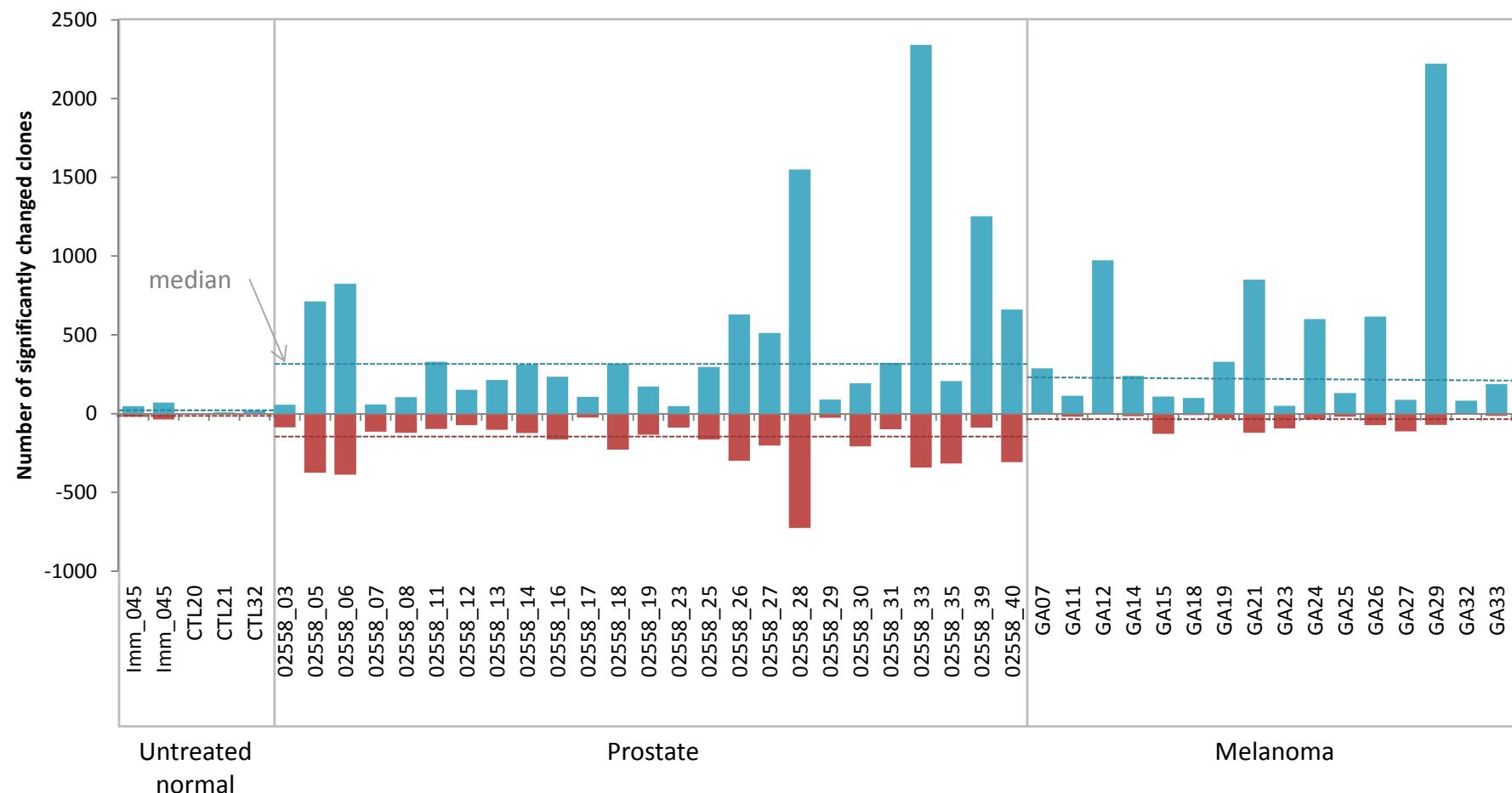
# CTLA-4 blockade induces remodeling of the T cell repertoire



(Cha et al., Science TM 2014)

# CTLA-4 blockade increases TCR diversity overall

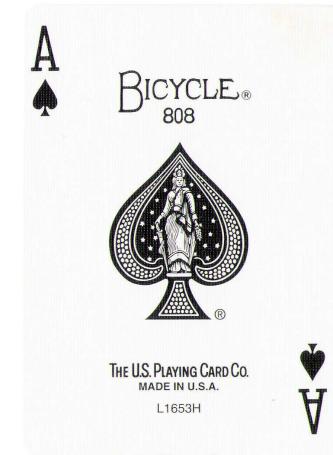
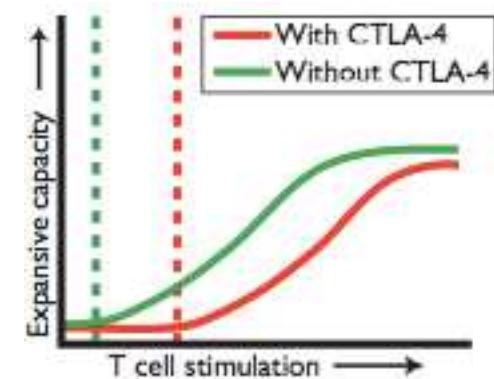
Number of clones changing in abundance after CTLA-4 blockade (one month)



(Cha et al, Science TM 2014)

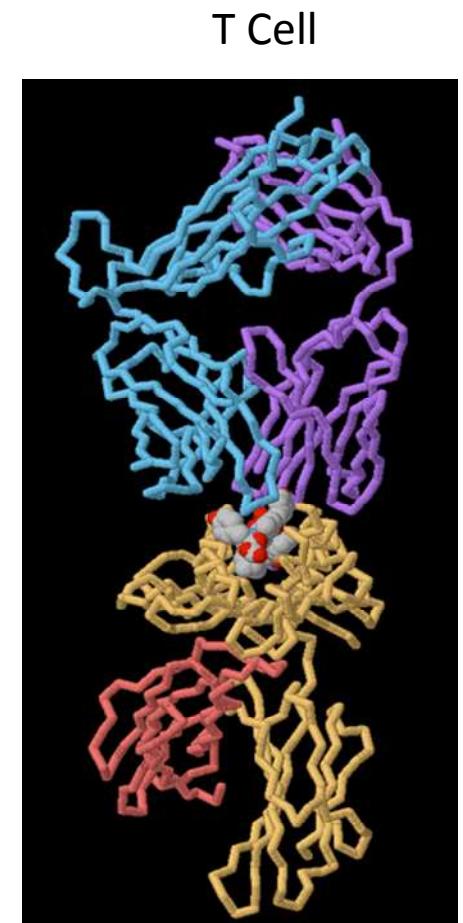
# Model

- Threshold model: CTLA-4 helps set the threshold for T cell activation  
Prediction: Low avidity T cells would be allowed to proliferate leading to increased TCR diversity.



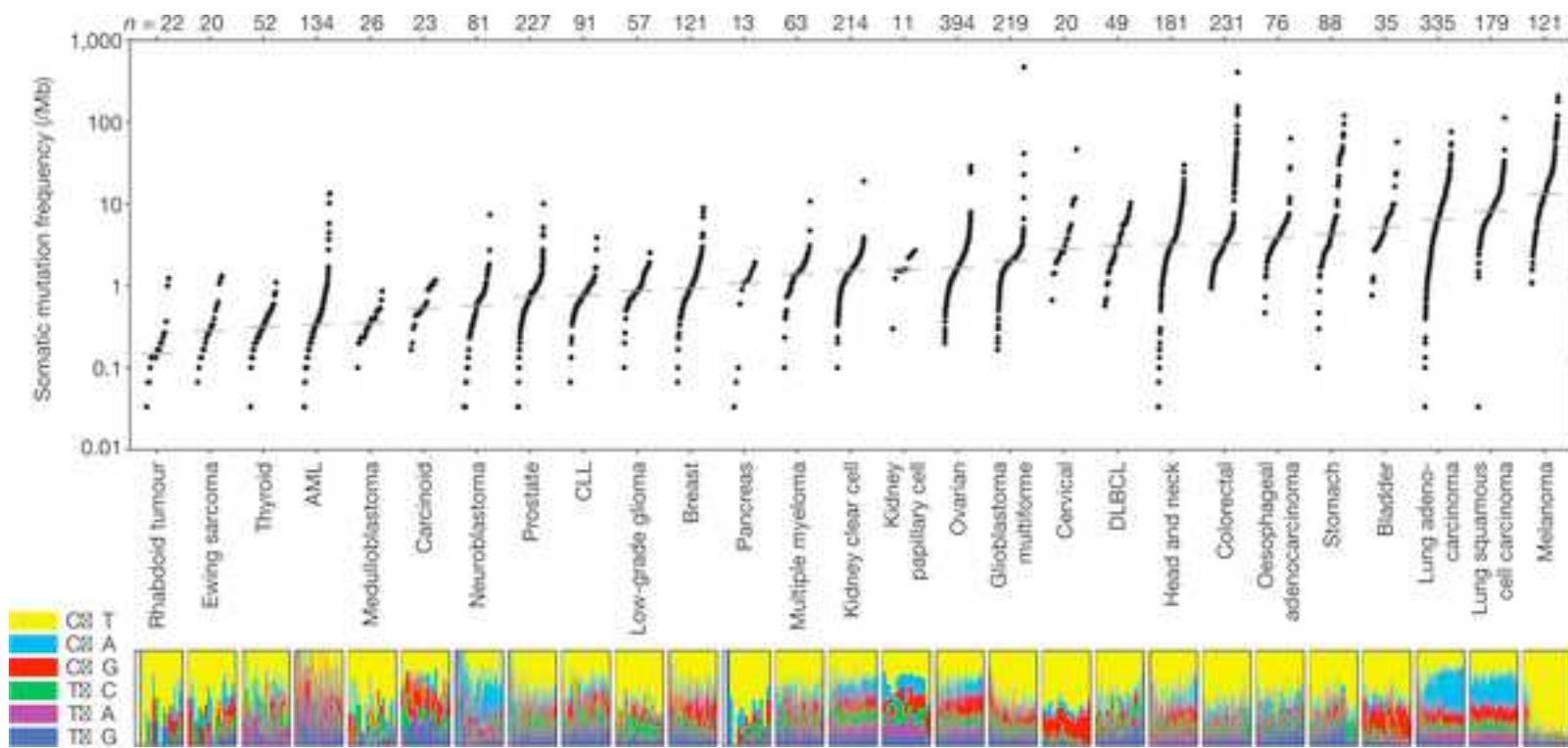
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Tumor specific:	Id, bcr-abl
	p53
	Tumor specific mutations



Tumor

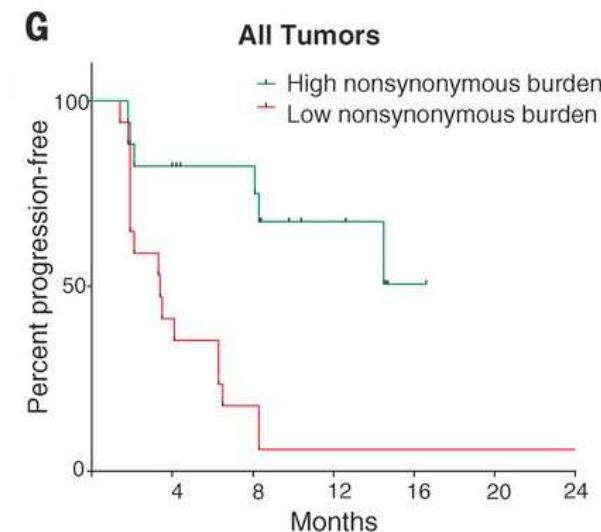
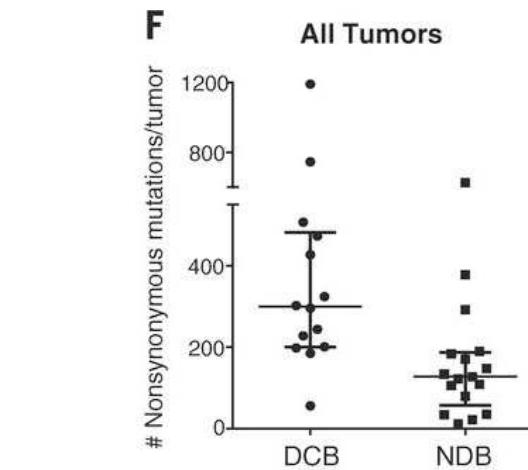
# Why are these different cancers responding?



(Lawrence et al, Nat 2014)

# Immune response to tumor mutations

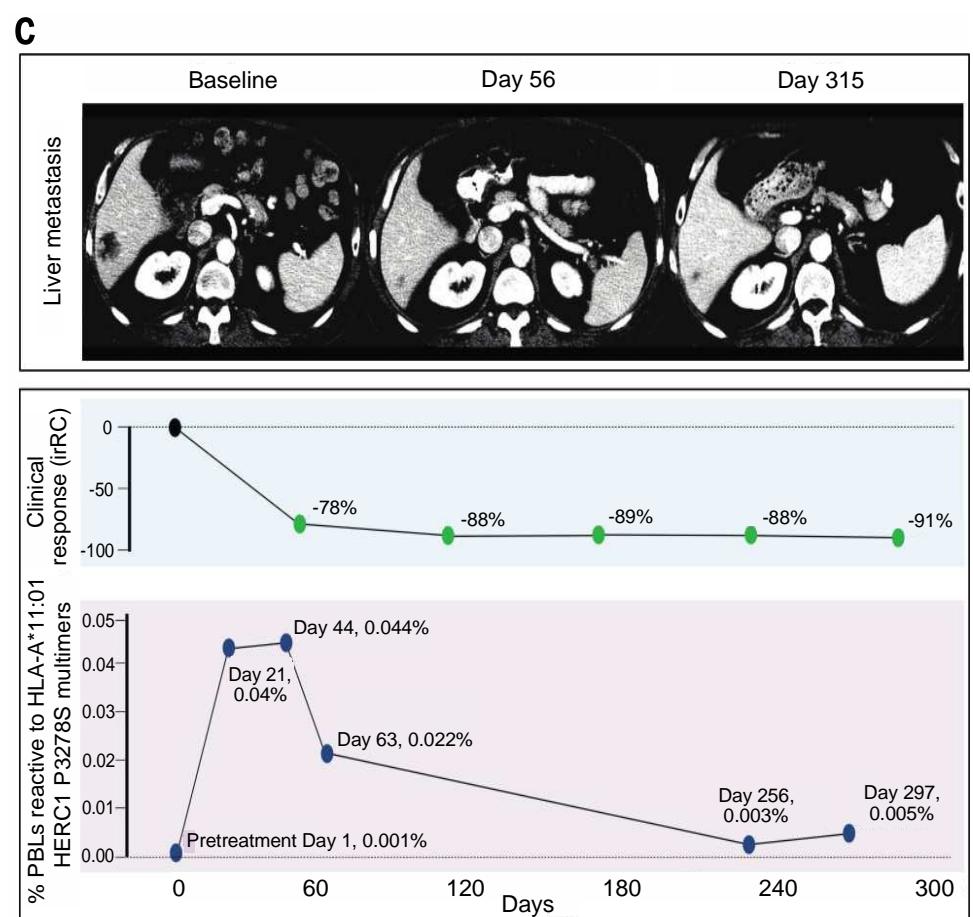
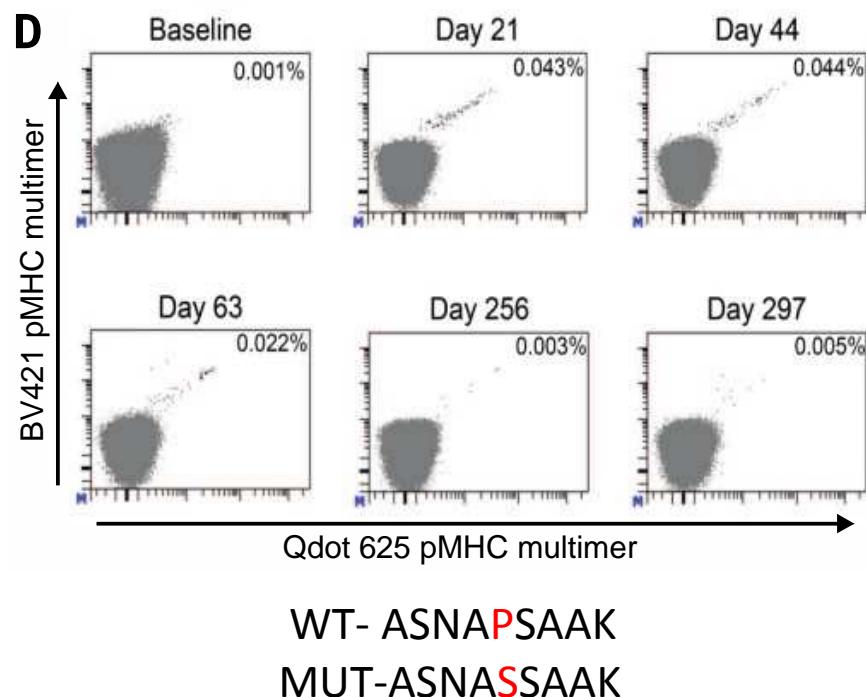
- Non-small lung cancer patients treated with anti-PD-1
  - Durable clinical benefit (PR, SD at 6mos) or not
- Whole-exome sequence tumor
  - Enumerate the non-synonymous mutations



(Rizvi et al. Science 2015)

# T cells recognizing neoantigens are expanded with treatment

- Identify candidate “neoantigens” by seeing which mutation will enhance binding to MHC
- Stain for these reactive T cells



(Rizvi et al. Science 2015)

# Conclusions

- Cancer patients possess a more diverse T cell repertoire.
- CTLA-4 blockade induces remodeling of the T cell repertoire leading to greater T cell diversity.
- Tumor mutations can provide a pool of endogenous antigens against which T cell responses can respond.

# Acknowledgements

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Peter Carroll

Jeff Simko

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UCLA

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Dendreon

Nadeem Sheikh

James Trager

Todd Devries

Mark Frohlich

Sequenta

Malek Faham

Mark Klinger

Adaptive

David Hamm



Patients and their families.



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NCI R01 CA194511



# Question 1

1) Cancer can stimulate:

- A. CD4 T cells
- B. Regulatory CD4 T cells
- C. CD8 T cells
- D. B cells
- E. All of the above

# Question 2

2) Sources of tumor associated antigens can include

- A. Differentiation antigens
- B. Cancer/testis antigens
- C. Overexpressed self antigens
- D. Neoantigens derived from tumor associated mutations
- E. All of the above