

# Dana-Farber/Harvard Cancer Center Kidney Cancer SPORE

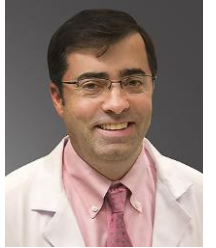
## A brief overview

2014-2019

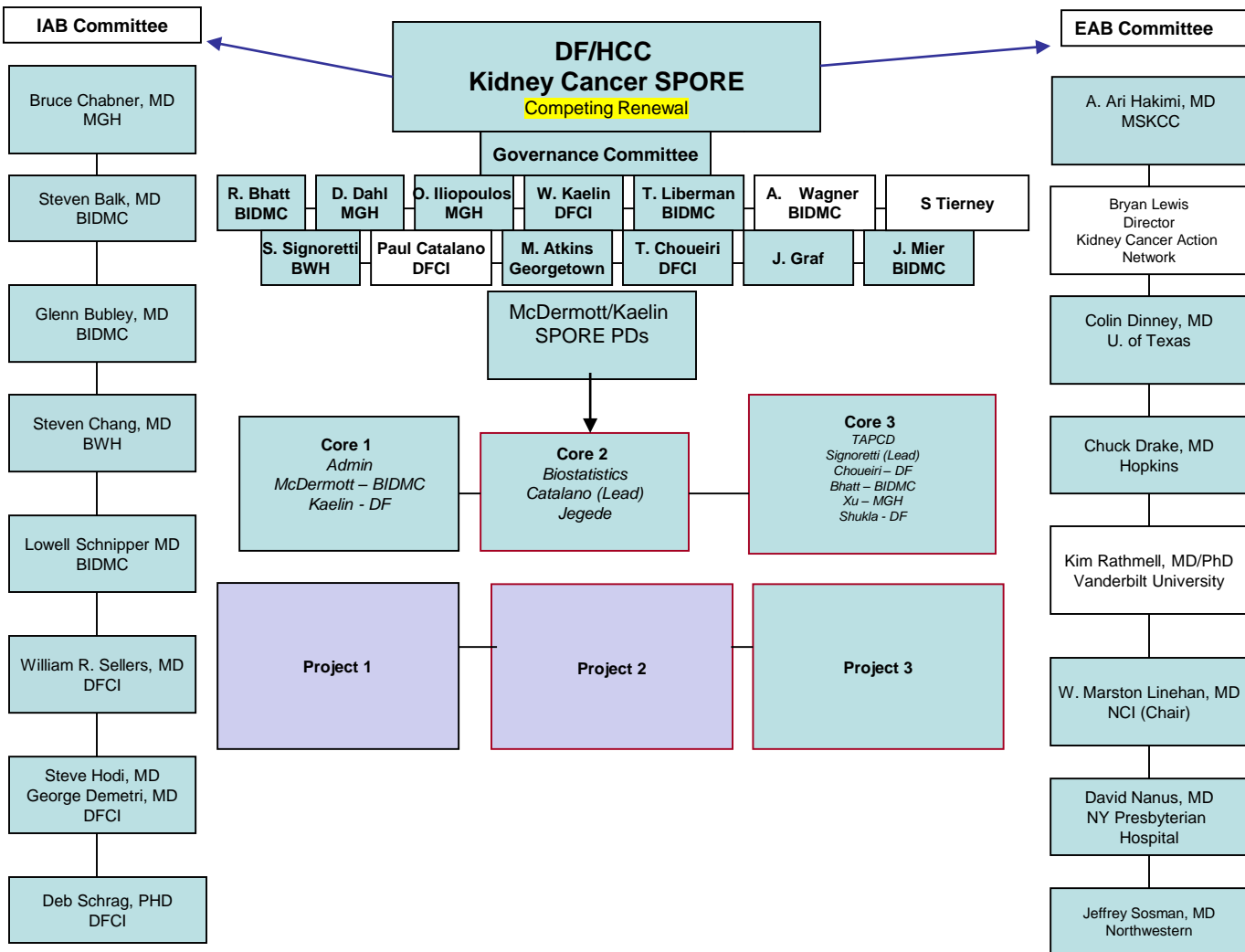
**Toni K. Choueiri, MD**

On Behalf of DF/HCC SPORE leadership

PI: McDermott and Kaelin



@DrChoueiri



# SPORE Related Clinical Research (2014-2019)

**Discovery**

**Translation**

**Impact**

Project	Agent	Target	Phase	Status
Angiogenic Escape	X4P-001	CXCR-4	1/2	Completed x 2
mTOR Targeting	TAK-228	mTOR kinase	2	Enrolling
PD-1 Targeting	Nivo then Ipi	PD-1, CTLA-4	2	Enrolling x 2
HIF2 Inhibition	PT-2385 PT-2977	HIF-2 $\alpha$	1/2	Completed P1 (x2) Enrolling P1/2 Combinations

# Beside Projects and Cores

- Career Development Awards (CDAs)
- Development Research Projects (DRPs)
- Director Choice Awards
- Patients and caregivers symposium (every year)
- E-Newsletters
- Integration within the DF/Harvard Kidney Cancer Program

# Efficacy of Novel HIF2a Inhibitor

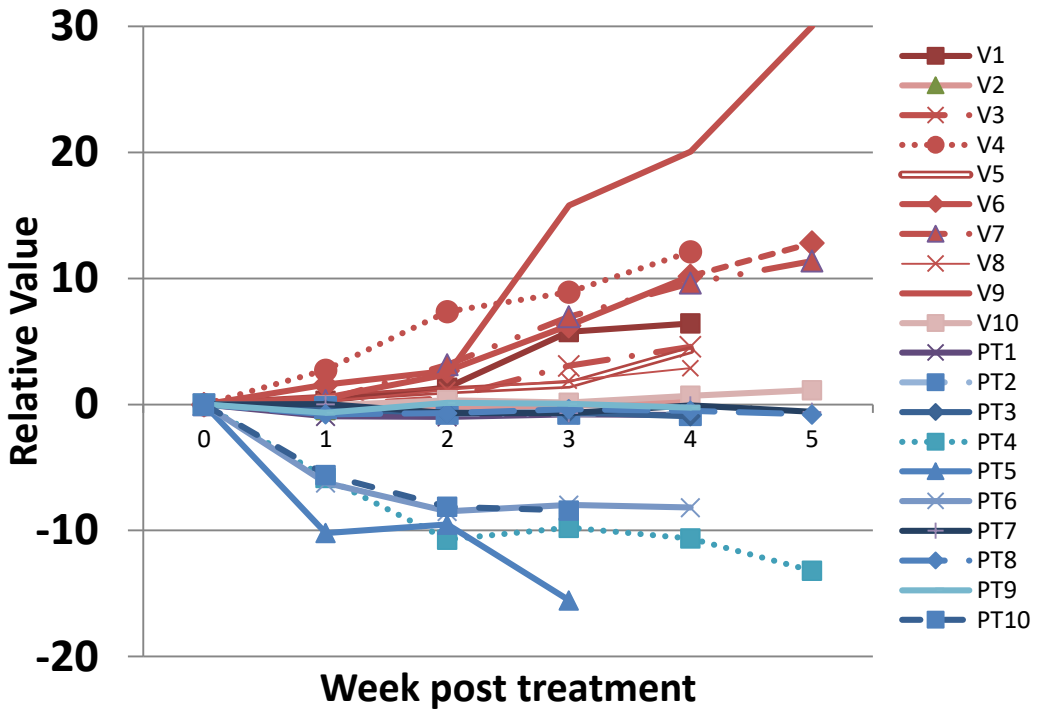
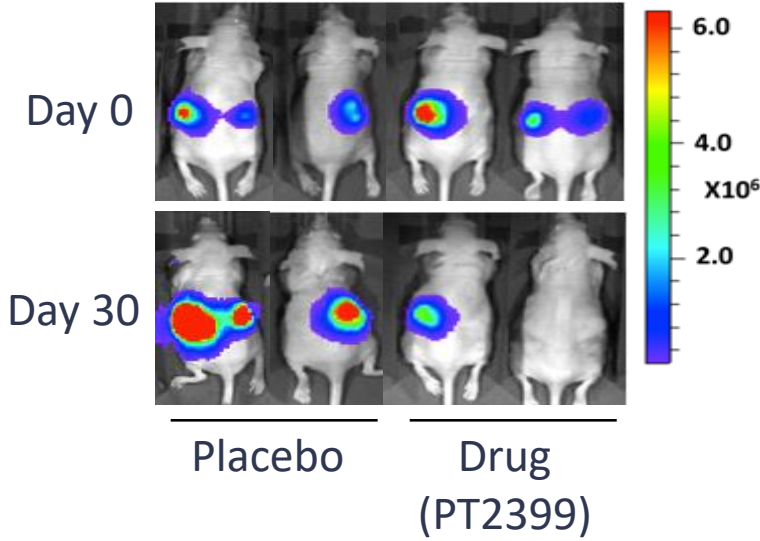
nature

2016

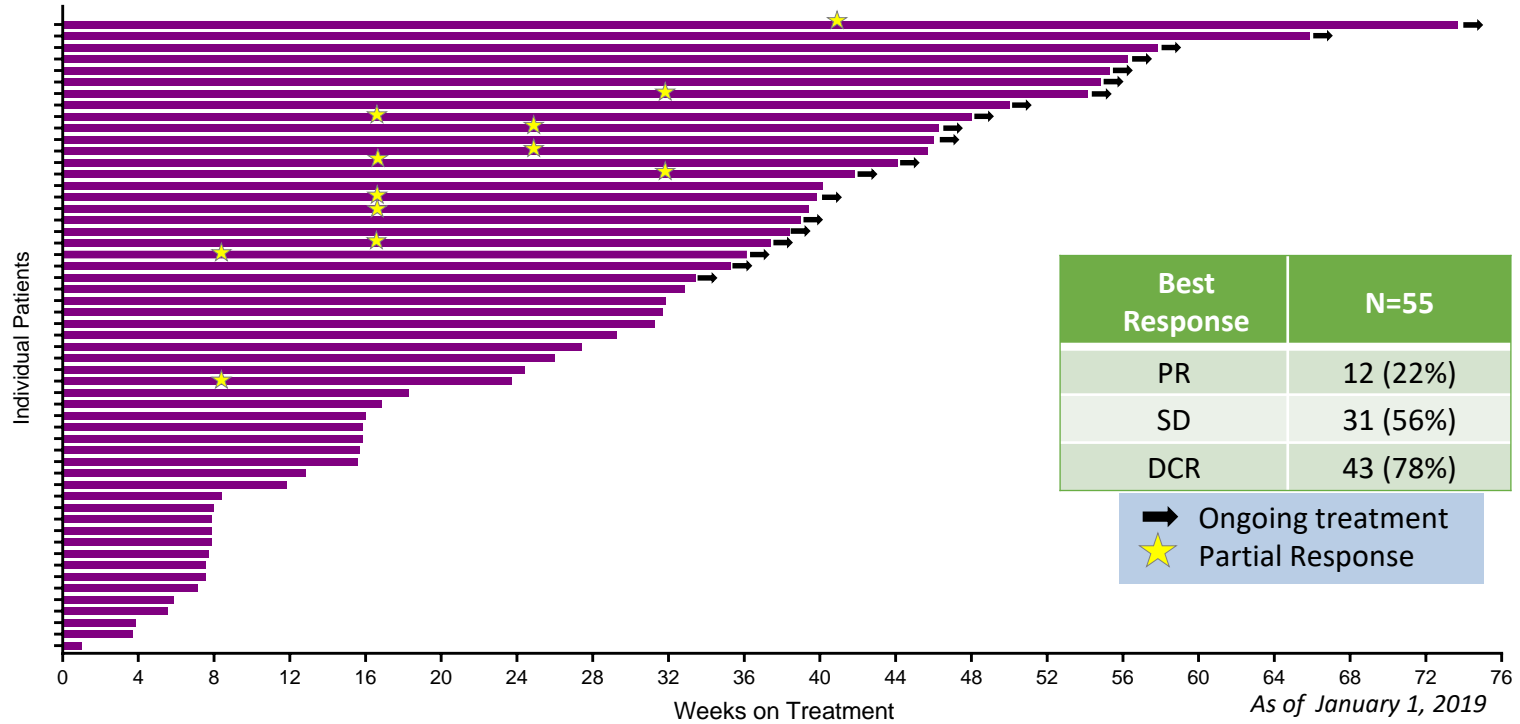


Kaelin<sup>DFCI</sup>

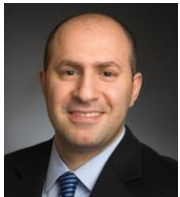
CMV-Luc 786-O Tumors



# Proof of Principle - HIF2a Inhibition: P1 PT-2977



Fourteenth European International Kidney Cancer Symposium  
 29-30 March 2019  
 Valamar Lacroma Hotel, Dubrovnik, Croatia  
 KidneyCancer.org  
 www.kidneycancersymposium.com

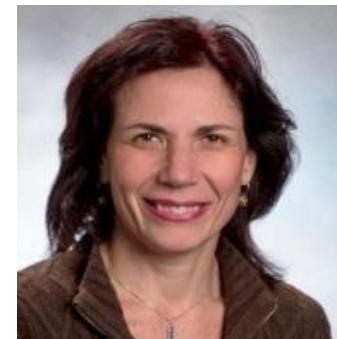
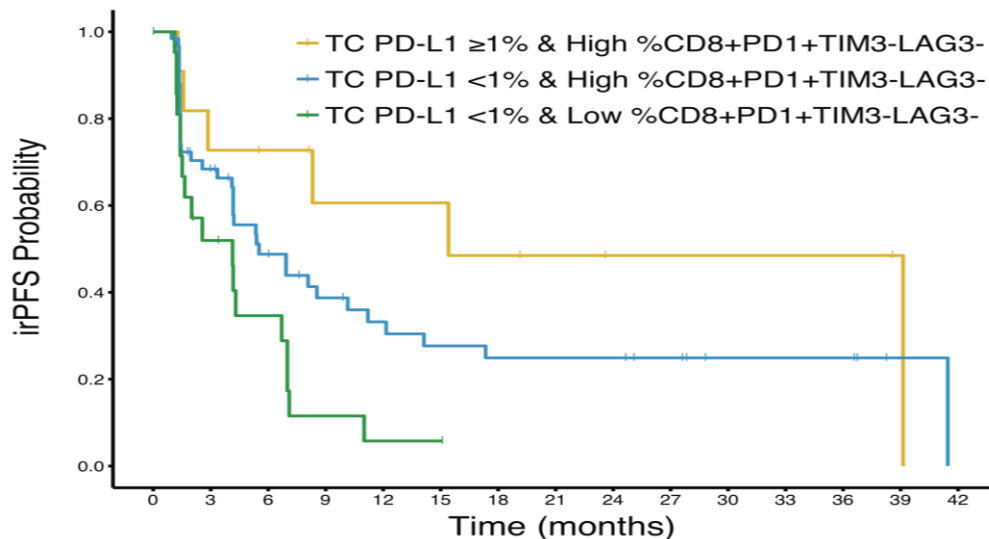


Choueiri<sup>DFCI</sup>

# Beyond PD-L1 expression

- Quantification of tumor infiltrating CD8-positive T cells
- Expression of multiple immune checkpoints on CD8-positive T cells

# Combined model (TC PD-L1 expression plus % of CD8+ cells that are PD-1+TIM-3-LAG-3-) identifies 3 groups of patients with distinct outcomes



Endpoints	TC PD-L1 expression $\geq 1\%$ and high % of CD8+ PD-1+TIM-3-LAG-3 <sup>a</sup> TIC (n = 11)		TC PD-L1 expression $< 1\%$ and high % of CD8+ PD-1+TIM-3-LAG-3 <sup>a</sup> TIC (n = 63 <sup>a</sup> )		TC PD-L1 expression $< 1\%$ and low % of CD8+ PD-1+TIM-3-LAG-3 <sup>a</sup> TIC (n = 23)		p-value
	n	%	n	%	n	%	
irORR	6	54.5	15	24.2	0	0.0	0.001
95% CI, %	23.4 - 83.3		14.2 - 36.7		0.0 - 15.4		
Median irPFS, months	15.4		5.5		4.1		0.013
95% CI	1.6 - 39.1		4.1 - 10.2		1.4 - 6.7		

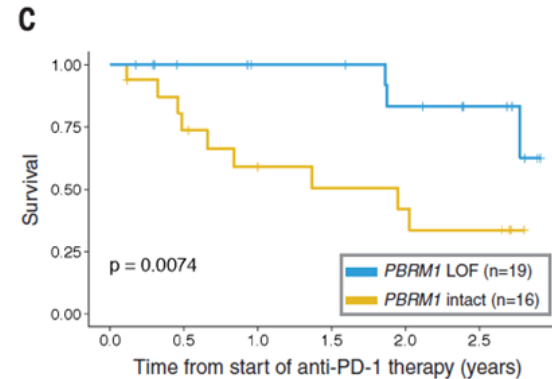
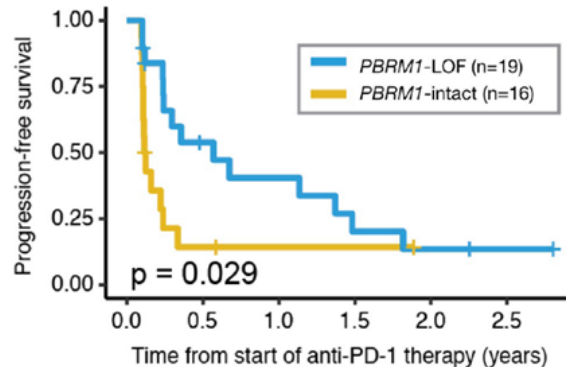
<sup>a</sup> irORR data for 2 patients were missing.



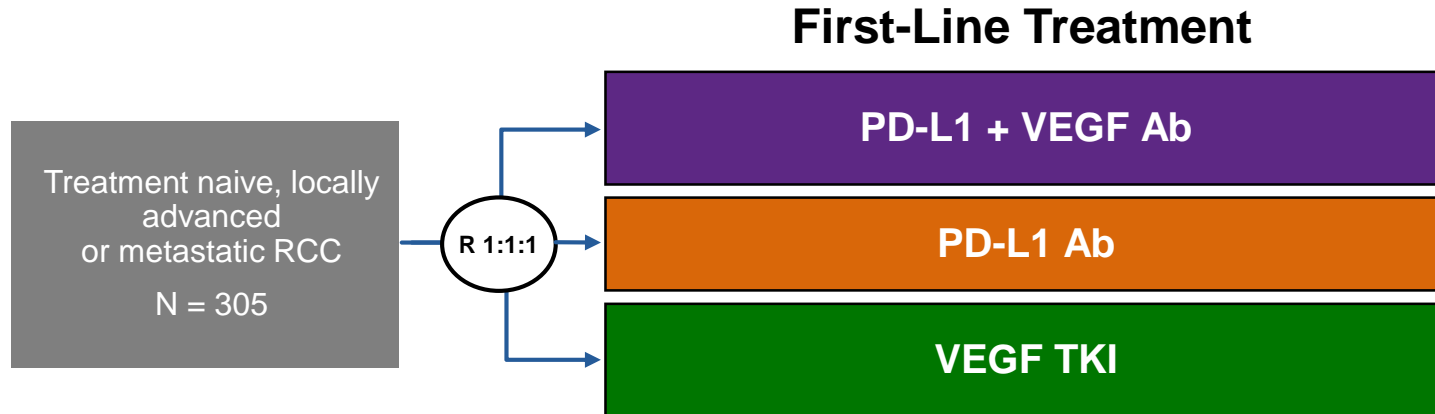
# Genomic correlates of response to immune checkpoint therapies in clear cell renal cell carcinoma

Diana Miao, Claire A. Margolis, Wenhua Gao, Martin H. Voss, Wei Li, Dylan J. Martini, Craig Norton, Dominick Bossé, Stephanie M. Wankowicz, Dana Cullen, Christine Horak, Megan Wind-Rotolo, Adam Tracy, Marios Giannakis, Frank Stephen Hodi, Charles G. Drake, Mark W. Ball, Mohamad E. Allaf, Alexandra Snyder, Matthew D. Hellmann, Thai Ho, Robert J. Motzer, Sabina Signoretti, William G. Kaelin Jr., Toni K. Choueiri, Eliezer M. Van Allen

**PBRM1 LOF enriched PFS and OS in Checkmate-009**



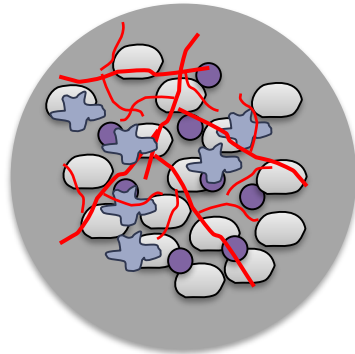
# IMmotion150 Trial Design: Randomized P2



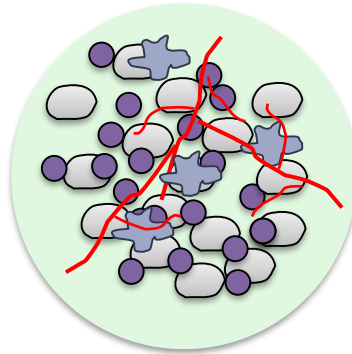
- IMmotion150 was designed to be **hypothesis generating** and inform the Phase III study IMmotion151
- **First Randomized Trial to:**
  - Explore **ICB** (atezo) + **Targeted Therapy** (bev)
  - Explore the **association between outcome** and **TME gene signatures**

TME, tumor microenvironment; ICB, immune checkpoint blockade

# Molecular Correlates of Differential Response to Atezolizumab ± Bevacizumab vs Sunitinib in mKC

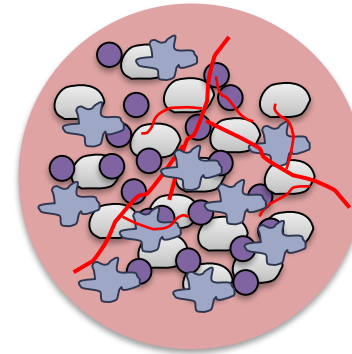


**Angiogenic**



**T-effector<sup>High</sup>**

**Myeloid Inflammation<sup>Low</sup>**

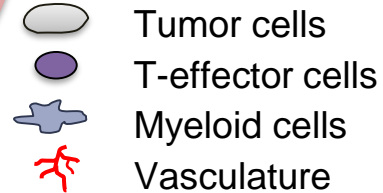


**T-effector<sup>High</sup>**

**Myeloid**

**Inflammation<sup>High</sup>**

***Immune Suppressed***



**Clinical Activity**

**VEGF TKI**

**PD-L1 Ab**

**PD-L1 + VEGF Ab**

# First-Line Phase 3 Trials in Advanced RCC

Control	Experimental Arm
Sunitinib	Axitinib + avelumab
Sunitinib	Bevacizumab + atezolizumab
Sunitinib	Nivolumab + cabozantinib
Sunitinib	Lenvatinib + everolimus or lenvatinib + pembrolizumab
Sunitinib	Axitinib + pembrolizumab
Sunitinib	Nivolumab + ipilimumab
Nivo/ipi	Nivolumab+ipilimumab
Adaptive	Nivolumab+ipilimumab followed by cabozantinib (Alliance)

All above with DF/HCC leadership or SC membership

**Should these approaches be applied to all patients?**

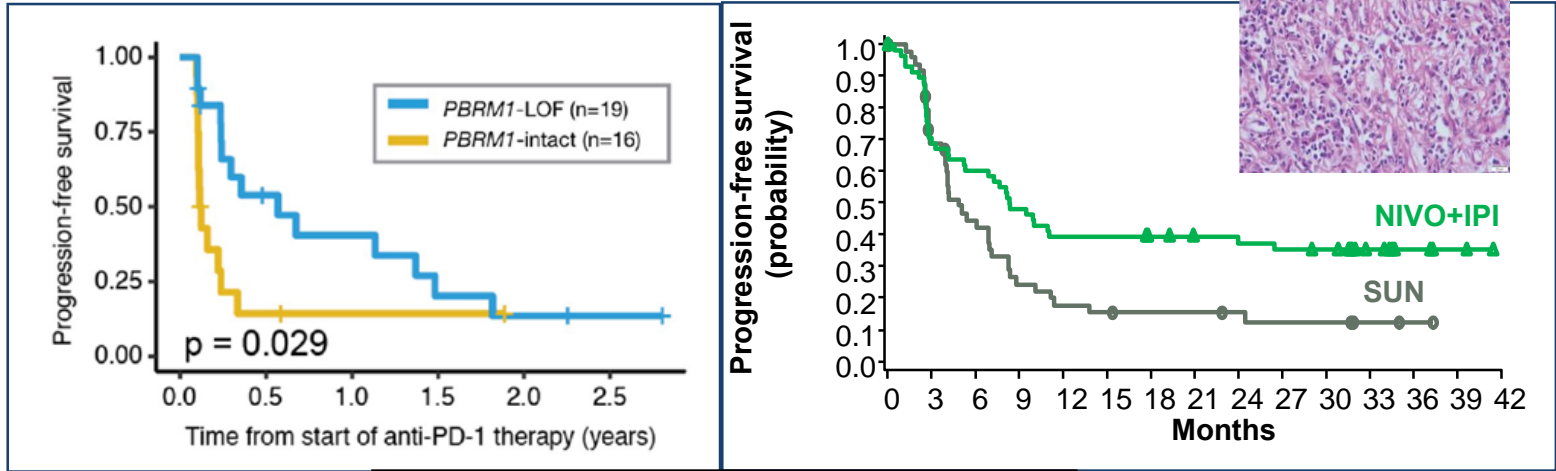
# DF/HCC KCP Immunotherapy Trials: Investigator Initiated

Trial	N	DF/HCC PI	Status
HCRN	200	McDermott	Enrolling
Omnivore	83	Harshman Choueiri	Enrolled
PROSPER	766	Harshman	Enrolling
Atezo/Bev ( <i>non-clear cell</i> )	60	Choueiri	Enrolled

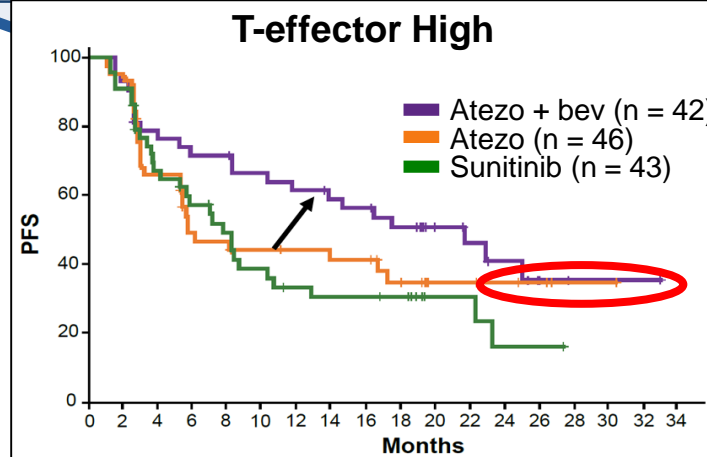
Close involvement with correlatives DF/HCC Lab- based colleagues:

- Signoretti Lab
- Wu lab
- Van Allen Lab
- Sharpe Lab
- Freeman Lab

# Biomarker Model



- All inter-related
- Some tumors may have a larger sweet spot



# Program Accomplishments 2016-present

## Awards

**Lasker Basic Medical  
Research Award 2016**

Bill Kaelin<sup>DFCI</sup>



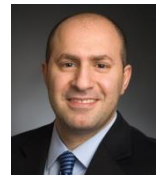
## SPORE Pathology Core Supports Collaborative Efforts



Sabina  
Signoretti<sup>BWH</sup>

Director

SPORE  
TAPCD Core



Toni  
Choueiri<sup>DFCI</sup>

Co-PI

International  
mRCC Data Base  
Consortium

## Large Grants

**UO1 Grant – 4/19**

**(PI: McDermott/Linehan)**

**Developing a Translational Pipeline  
for VHL Mutant Malignancies**

# Future Plans (DF/HCC Kidney SPORE)

- Competitive Renewal (2019-
- Promote the rationale application of therapies:
  - Integrative Biomarkers
  - Novel targets (e.g.: HIF-2, neoantigens)
  - Novel trials design (e.g. HCRN, OMNIVORE, PROSPER, Neovax)
- External funding (e.g. DOD)
- Continue to liaise with advocacy groups
- Mentor **and sponsor** the next-generation of Kidney Cancer researchers