

Efficacy of Cabozantinib Plus Nivolumab in Cluster 1/2 Metastatic Clear Cell Renal Cell Carcinoma

Results from OPTIC RCC, a phase II trial of a novel RNAseq-based biomarker

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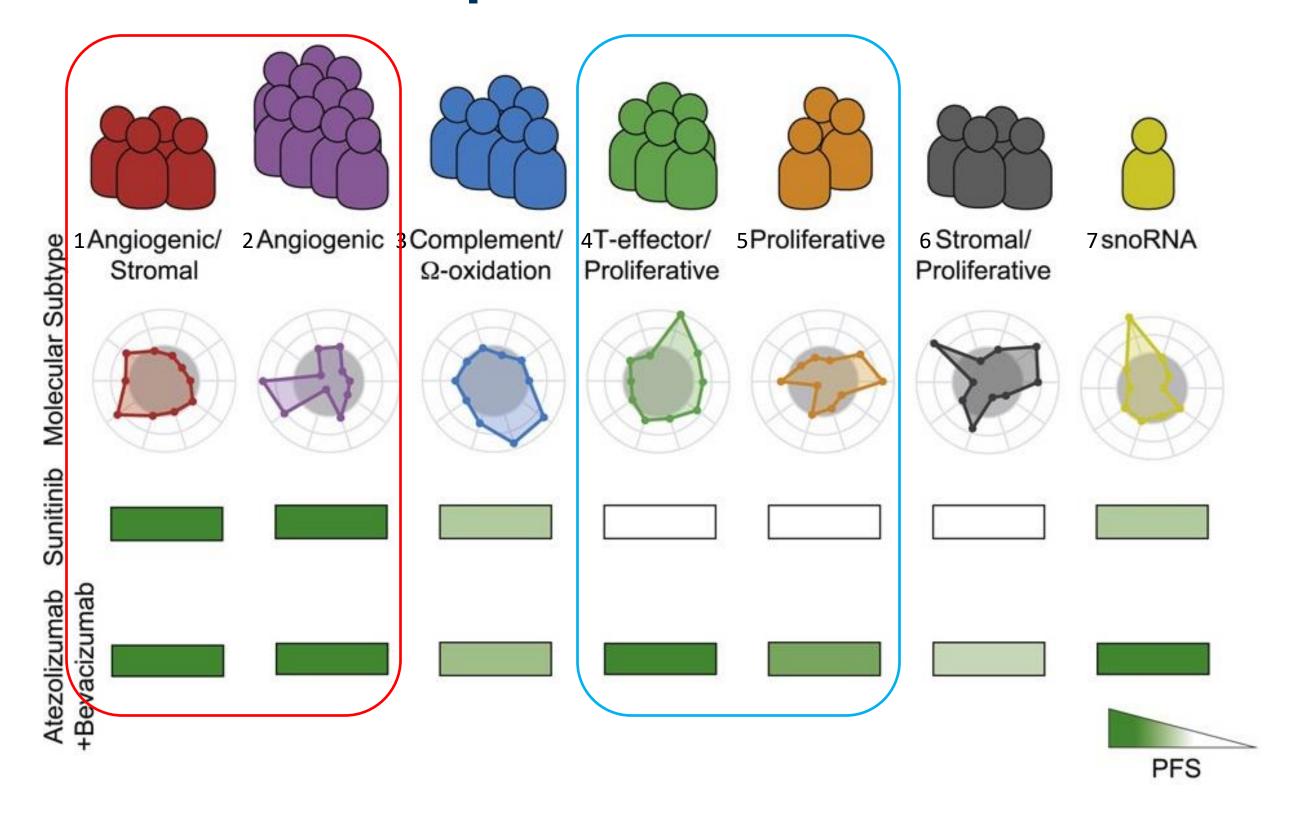


Declaration of Interests

- Invited speaker: Tempus
- Institutional research funding: BMS
- Ad board: Merck, Eisai, Exelixis



IMmotion 151 RNAseq-Defined Clusters



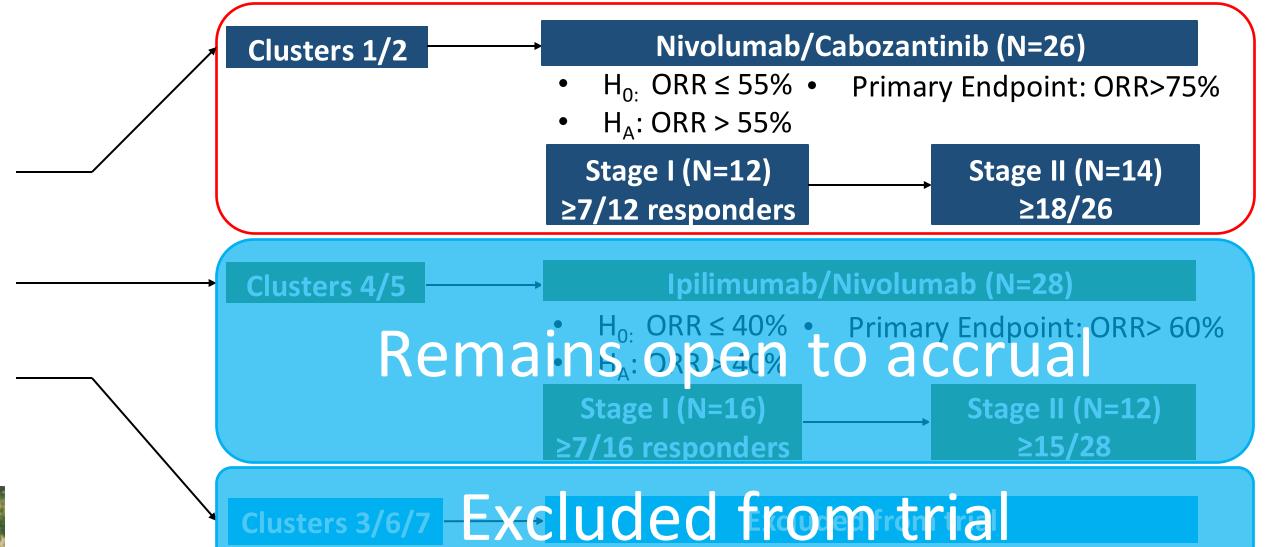


OPtimal Treatment by Invoking biologic Clusters in Renal Cell Carcinoma (OPTIC RCC)

Simon's Minimax Two-Stage Design

Key Eligibility Criteria

- ECOG 0 or 1
- Metastatic clear cell RCC without prior systemic therapy in any setting
- Available tumor tissue for RNA-sequencing/cluster prediction





Rini







Scott Haake, MD, PhD

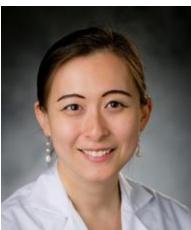
Haake

OPtimal Treatment by Invoking biologic Clusters in Renal Cell Carcinoma (OPTIC RCC)

Patient w/ stage IV ccRCC Isolate Sequence **mRNA** UACUCAGGUUCA Tempus xR Obtain metastatic +/primary tumor tissue **Automated** Al-Driven Vindhya **Analysis** Mar (UCI) Custom website



Ornstein (CCF) Barata (UH)



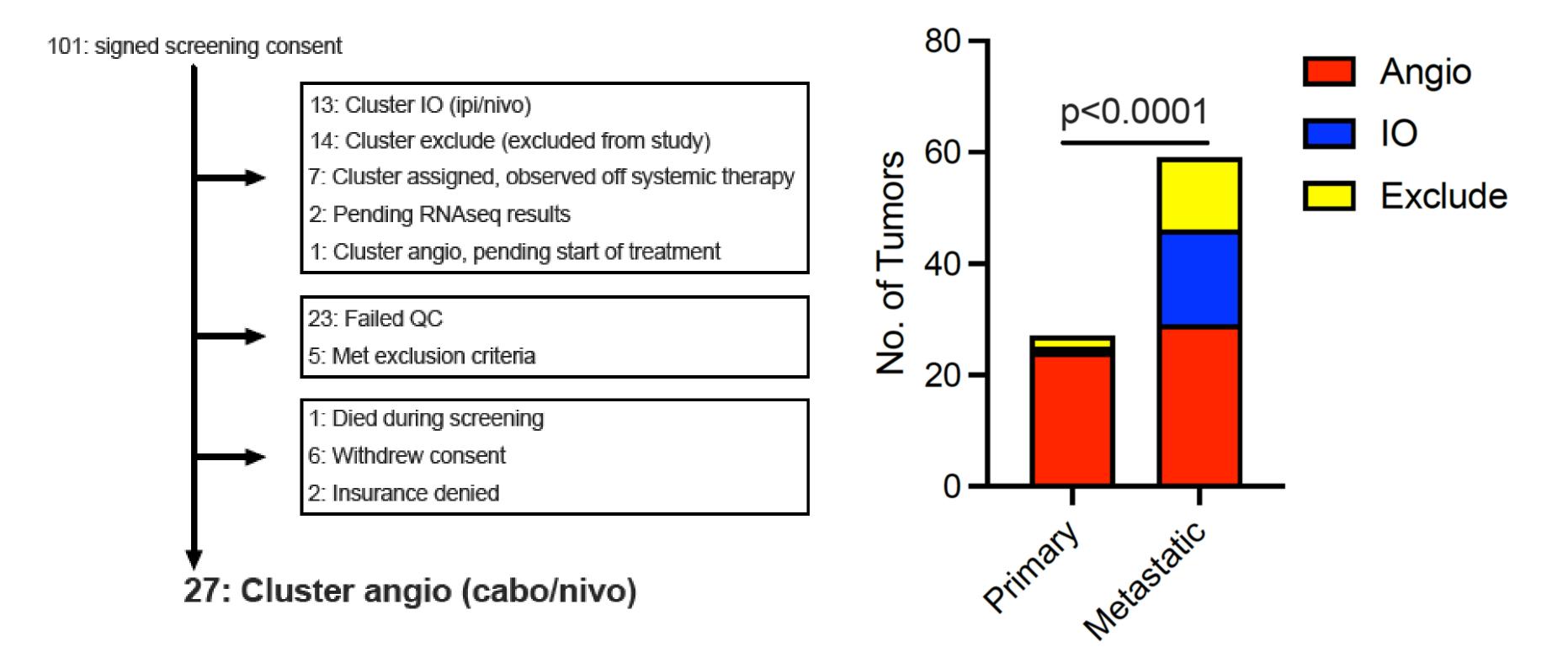
Zhang (UTSW)

Scott Haake, MD, PhD

disseminates results to

participating sites

CONSORT diagram & cluster distribution







Patient Characteristics

Characteristic (n=27)		
Age (year) – median (range)	68 (52-86)	
Gender – M:F (%)	56:44	
Race – White:Black:Asian (%)	89:7:4	
ECOG performance status – 0:1 (%)	70:30	
IMDC prognostic risk score – Favorable:Intermediate:Poor (%)	41:52:7	
Source of RNA – Primary:Metastatic (%)	22:78	
Sarcomatoid histology— no. (%)	0 (0)	
Nephrectomy – no. (%)	22 (81)	
No. metastatic sites – median (range)	2 (1-5)	
Most common sites of metastasis – no. (%)		
Lung	18 (67)	
Pancreas 8 (30		
Adrenal	7 (26)	
Lymph Node	7 (26)	
CNS	5 (19)	
Liver	4 (15)	
Soft Tissue 5 (19)		
Bone	2 (7)	
Other (peritoneum, cervix, spleen, thyroid)	5 (19)	

Objective Response

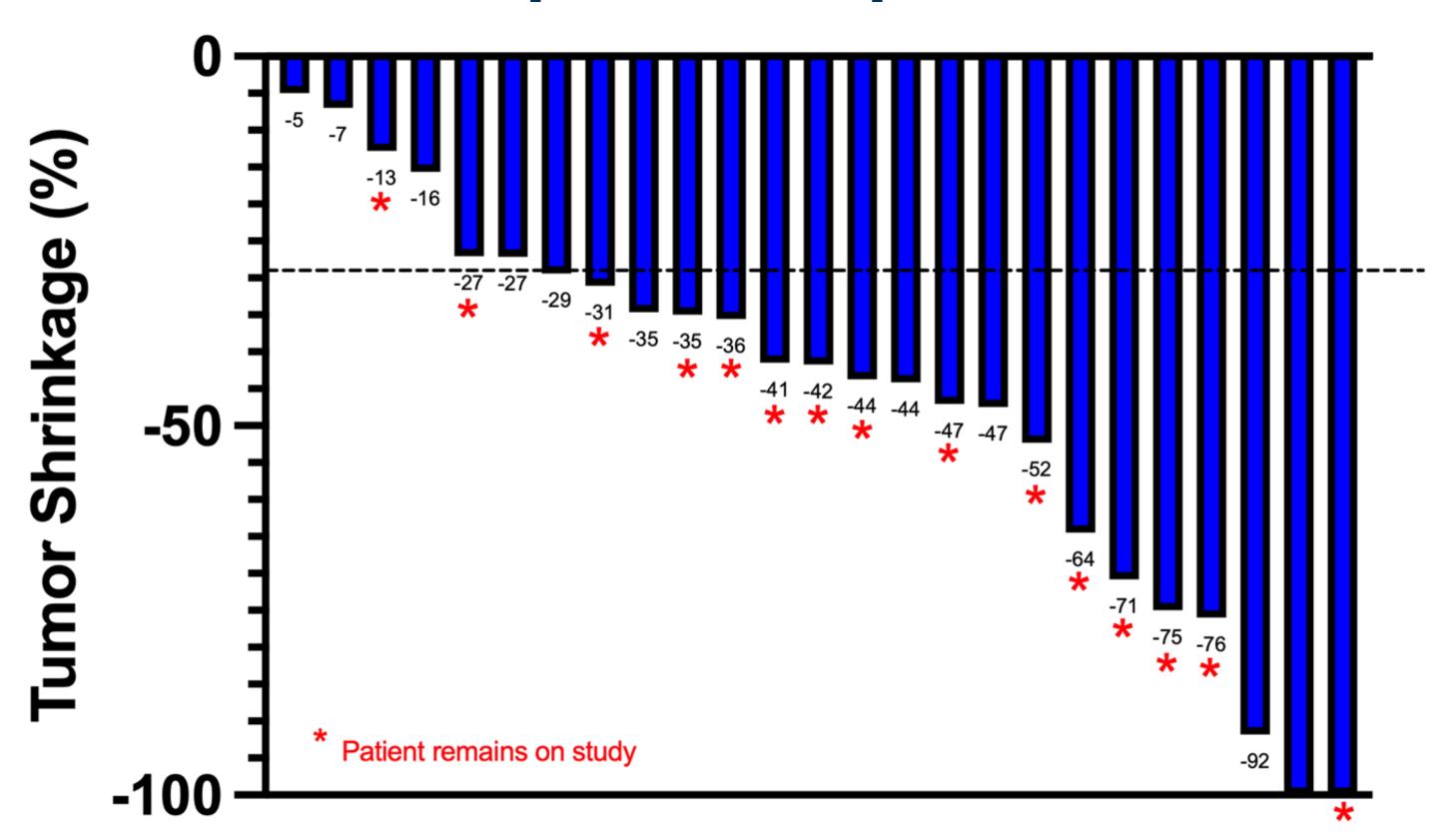
	Cluster 1/2 (n=25)*	
Objective response – no. (%)	18 (72)	
Best overall response – no. (%)		
Complete response	2 (8)	
Partial response	16 (64)	
Stable disease	8 (32)	
Progressive disease	0 (0)	
Patients with tumor shrinkage – no. (%)	25 (100)	
% tumor shrinkage – median (range)	42 (5-100)	

^{* 2} patients recently enrolled without a post-baseline scan to date

⁻ Median follow-up: 11.1 months (range 0.9 – 31.5); 17 of 27 patients remain on study



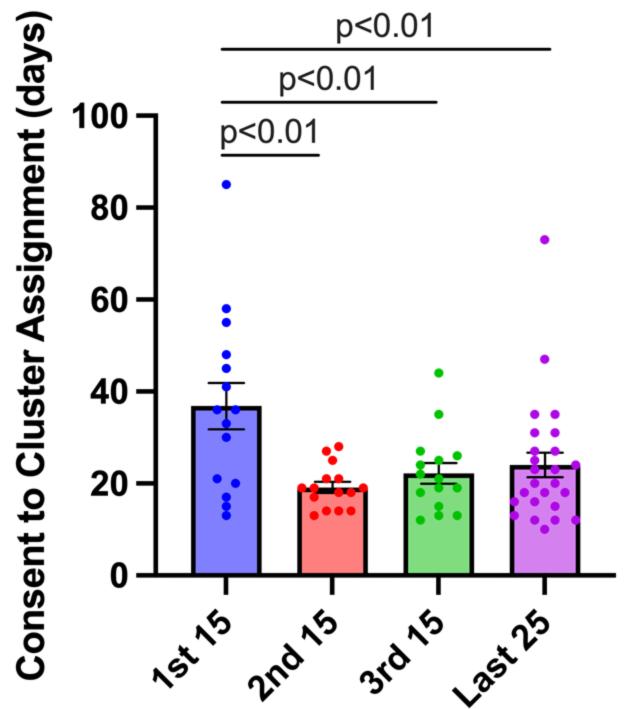
Depth of response







Consent-to-cluster turnaround time



			•	
Order o	of Pati	ient Ei	nrolli	ment

Order of patient enrollment	Consent-to-cluster (months) – median (range)
1 st 15	36 (13-85)
2 nd 15	19 (13-28)
3 rd 15	21 (12-44)
Last 25	20 (10-73)

X genes are enriched in best responding tumors

- Analysis from Anupama comparing pre-treatment gene expression in tumors with >50% shrinkage versus tumors with <50% shrinkage



Conclusions

- Prospective investigation of tissue-based RNAseq biomarkers to guide therapy is operationally feasible in a front-line metastatic ccRCC population
- Primary and metastatic tumors can exhibit different gene expression signatures, requiring metastatic tumor biopsies for a biomarker-driven study
- Selection of patients exhibiting an angiogenic gene expression signature enriches for clinical outcome to cabozantinib + nivolumab
 - High objective response rates, reduction of tumor burden in all patients and lack of primary progressive disease was observed.
- Time-to-event endpoints such as PFS and duration of response will require additional follow up



Acknowledgements

Patients

OPTIC RCC Leadership:

- Brian Rini
- Katy Beckermann

Site Pls:

- Pedro Barata
- Moshe Ornstein
- Tian Zhang
- Nataliya Mar

Data Scientists:

- Anupama Reddy
- Alex Nesta

Tempus:

- LaShantay Walls
- Rebecca Ramenaden

Others:

- Ellen Heimann-Nichols
- Yu-Wei Chen
- W. Kimryn Rathmell
- Carmen Gray
- Jason Brown
- Christopher Wee
- Eric Davis

Funding:

- US Department of Defense
- KidneyCANMany, many others!!







