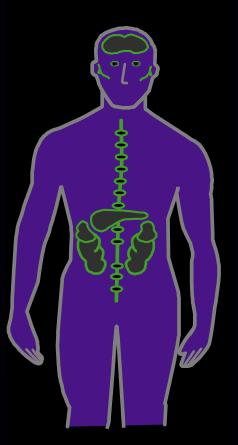
What Do My Genes Tell Me About Kidney Cancer?

Ramaprasad Srinivasan, M.D., Ph.D Investigator and Head, Molecular Cancer Section Urologic Oncology Branch, Center for Cancer Research National Cancer Institute

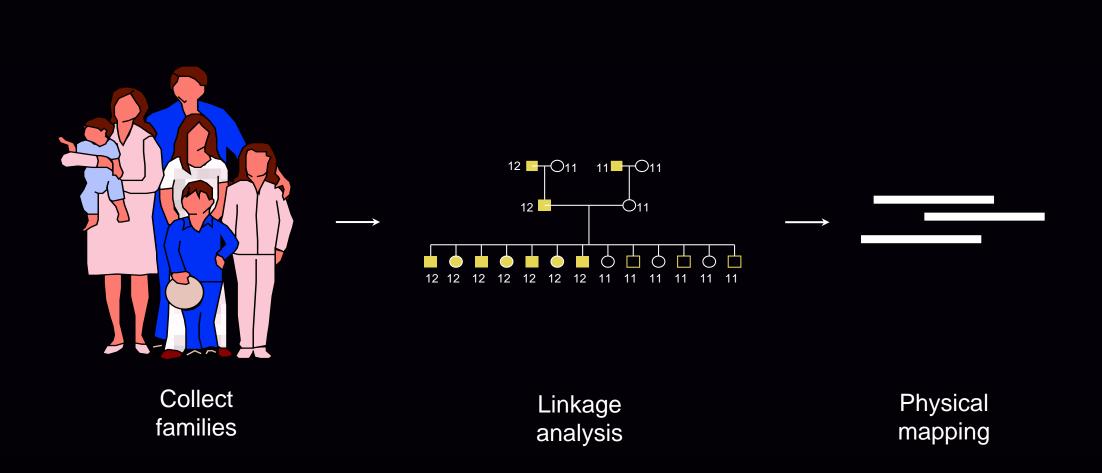
von Hippel Lindau Disease

- Tumors develop in:
 - Both Kidneys
 - Adrenal Glands
 - Pancreas
 - Brain or Spine
 - Eyes
 - Inner Ears

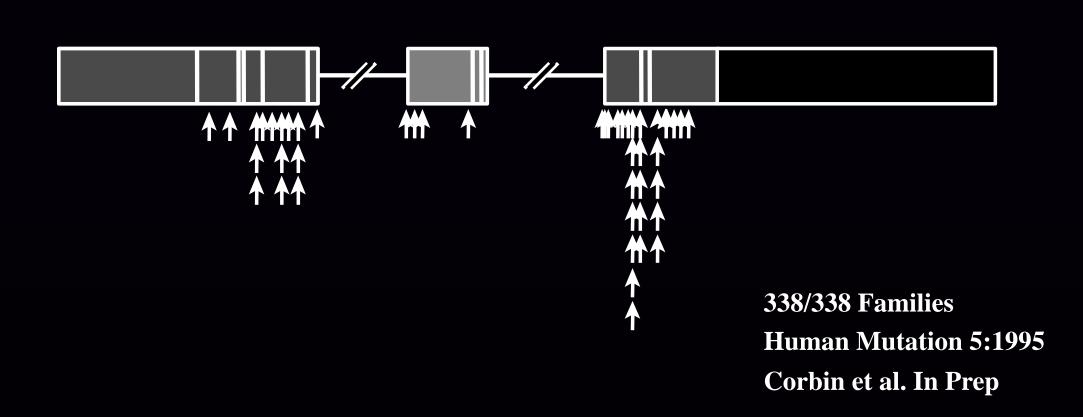


Identification of the VHL gene

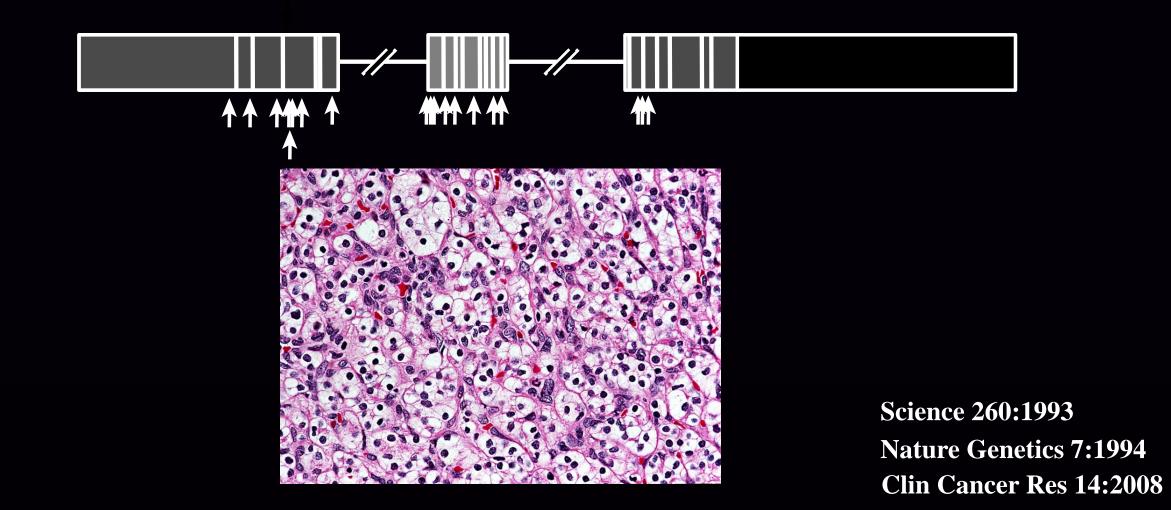
(W. Marston Linehan and Berton Zbar, NCI)



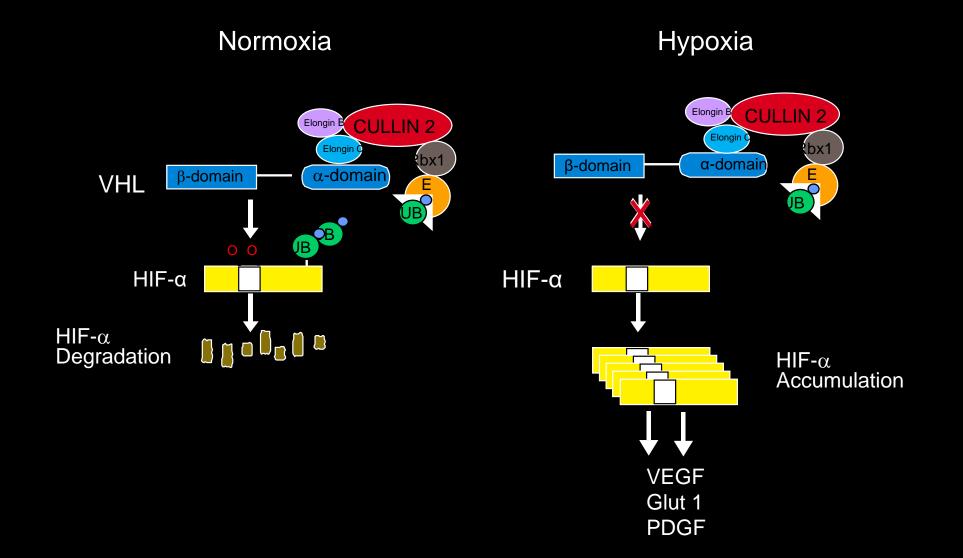
Germline VHL Mutations



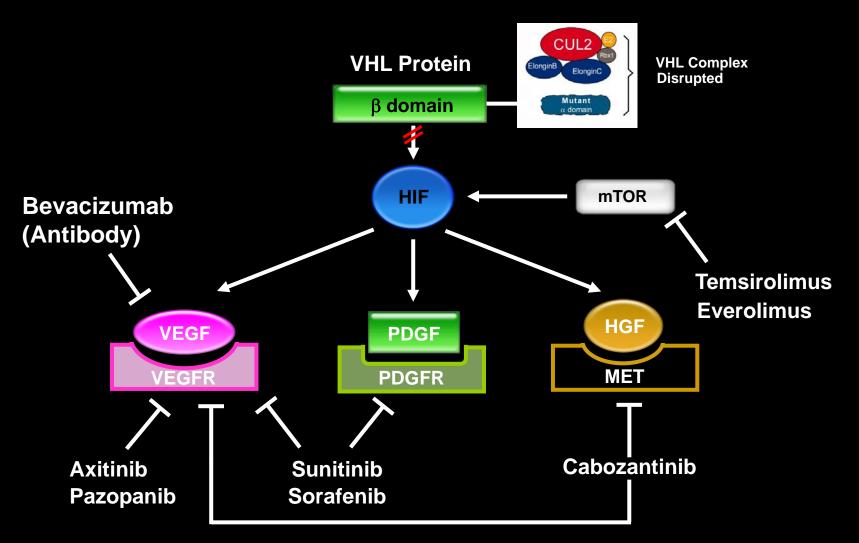
Sporadic Clear Cell RCC VHL Gene Mutations



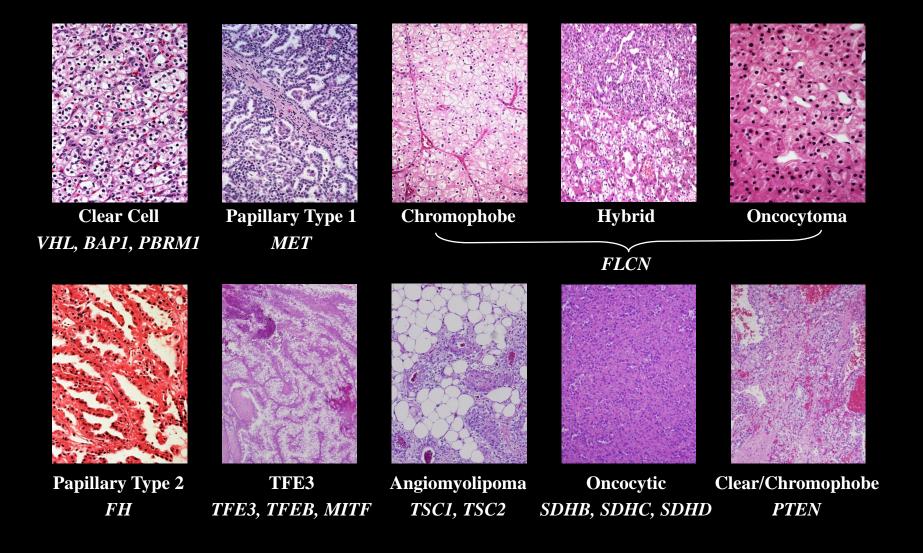
HIFα is targeted for degradation in normoxic, but not hypoxic cells



Targeting the VHL Pathway in Sporadic Clear Cell RCC



Histologic and Genetic Heterogeneity in Human Renal Epithelial Neoplasms



Genetic Alterations: Germline versus Somatic

<u>Germline Alteration</u>

- Identified by testing in blood cells, buccal mucosa etc
- Affects one allele in cells throughout the body
- Inherited (patients are born with these changes)
- Can be transmitted to offspring, usually in an autosomal dominant manner
- Cancer predisposition

Somatic Alteration

- Identified by testing tumor
- Alterations restricted to the tumor
- Not inherited

Who Should Undergo Germline Testing?

- Guidelines: e.g. American College of Medical Genetics and Genomics
- Factors to consider while referring for genetic evaluation
 - Presence of bilateral, multifocal tumors
 - Young age at diagnosis (generally < 50 years)
 - Histology- papillary type 1 and 2, collecting duct, tubulopapillary etc
 - Presence of other clinical features associated with herediatry RCC (e.g. multiple cutaneous leiomyomas)
 - Strong family history
- Patients should be assessed/counseled by a genetic counselor

I Have a Germline Mutation: Will I Get Kidney Cancer?

- Depends on the specific gene affected
- MET mutations highly penetrant
 - Almost all affected individuals will develop tumors by age 80
- Fumarate Hydratase Mutations
 - 15-30% of individuals with the mutation will develop kidney cancer
- No reliable way to predict who will develop cancer
 - Lifelong screening recommended for most patients

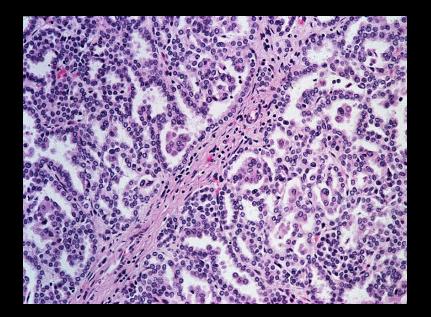
Identifying Genetic Alterations: How Does it Help?

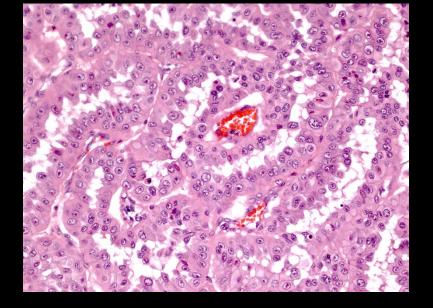
Identification of specific genetic alterations can

- Help establish a diagnosis
- Identify families at risk and enable appropriate screening
- Dictate choice of treatment
- ? Serve as a prognostic factor

Can my Genes Help Guide Therapy?

Papillary RCC: Histologic Subtypes

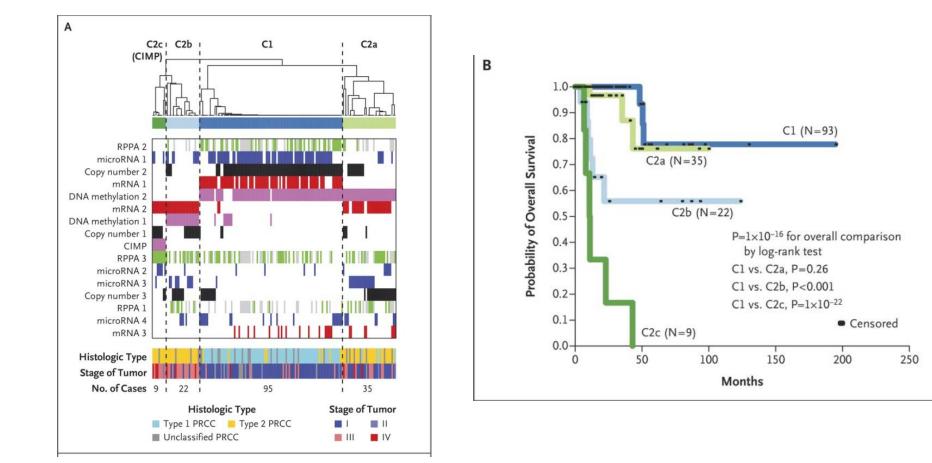




Type 1

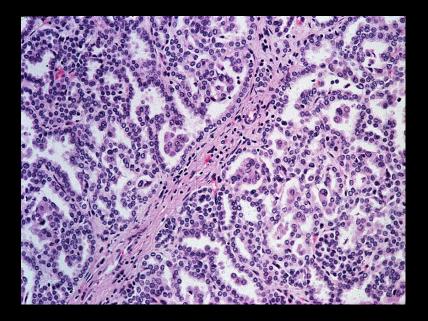
Type 2 (Non Type 1)

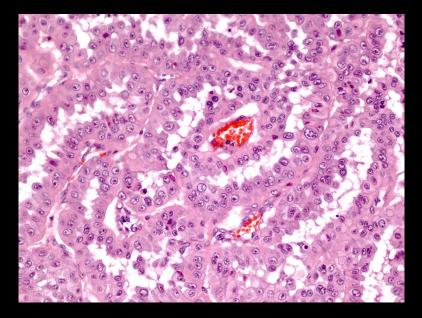
TCGA: Genetic Profiling of Papillary RCC: 4 Distinct Subgroups



NEJM 2016

Papillary RCC: Histologic Subtypes





Type 1

MET TERT Promoter EGFR CDKN2A/B

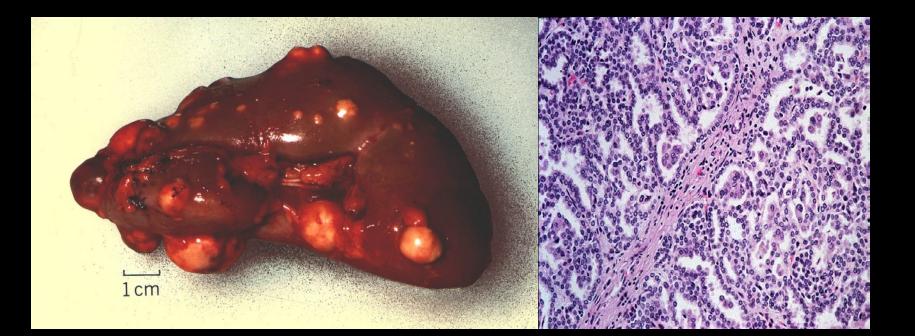
Type 2 (Non Type 1)

Fumarate Hydratase NRF2 Pathway CDKN2A/B SWI/SNF, Chromatin Remodeling

Hereditary Papillary Renal Cancer (HPRC)

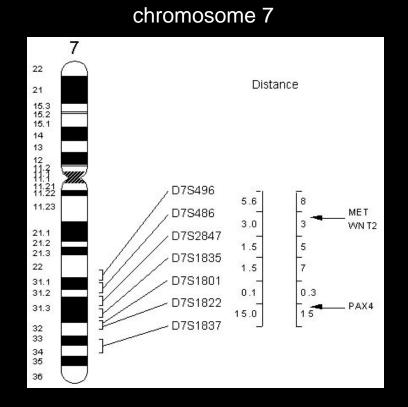
Familial form of type I papillary RCC

 Affected individuals present with bilateral multifocal papillary RCC



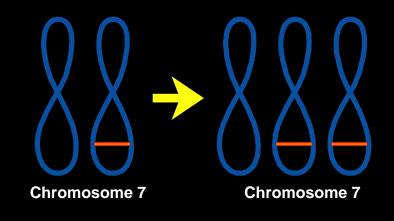
MET- The Gene for Hereditary Papillary Renal Cancer

Germline mutations in *MET* are the hallmark of HPRC



Location of *MET* on

Nonrandom duplication of chromosome bearing mutated *MET* allele



Schmidt et al., Nat Genetics, 1997

Met 'Activation' in Sporadic Papillary RCC

- Activating Mutations in MET
 - Somatic activating mutations seen in ~10% of sporadic papillary RCC
 - MET fusion or splice variants~ 5-7%
- Gain of chromosome 7
 - ~ 80% of type 1 papillary RCC
 - Both MET and its activating ligand HGF located on Ch 7
- Focal Amplification relatively rare
- MET and Ch7 alterations seen predominantly in type 1 papillary RCC (TCGA)

VOLUME 31 · NUMBER 2 · JANUARY 10 2013

JOURNAL OF CLINICAL ONCOLOGY

ORIGINAL REPORT

Phase II and Biomarker Study of the Dual MET/VEGFR2 Inhibitor Foretinib in Patients With Papillary Renal Cell Carcinoma

Toni K. Choueiri, Ulka Vaishampayan, Jonathan E. Rosenberg, Theodore F. Logan, Andrea L. Harzstark, Ronald M. Bukowski, Brian I. Rini, Sandy Srinivas, Mark N. Stein, Laurel M. Adams, Lone H. Ottesen, Kevin H. Laubscher, Laurie Sherman, David F. McDermott, Naomi B. Haas, Keith T. Flaherty, Robert Ross, Peter Eisenberg, Paul S. Meltzer, Maria J. Merino, Donald P. Bottaro, W. Marston Linehan, and Ramaprasad Srinivasan

Primary Endpoint: Overall Response Rate

| | Dosing Cohort A (n=37) | Dosing Cohort B (n=37) | TOTAL (N=74) |
|------------------------------|------------------------------|------------------------------|-----------------|
| Overall Response Rate | 5 (13.5%) | 5 (13.5%) | 10 (13.5%) |
| | | | |

- Duration of response: 18.5 months
- Median PFS: 9.3 months

Germline MET Mutations Associated with High Response Rate

<u>N=67 evaluable:</u>

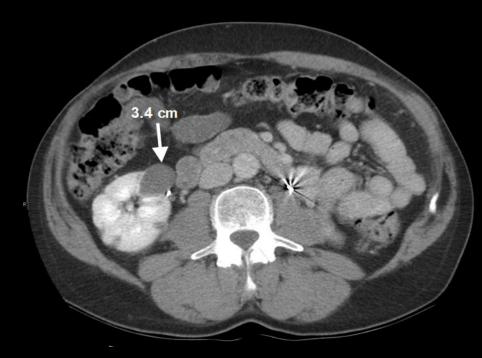
- Germline *MET* mutation (N=10)

- Mutated *MET:*
 - 5/10 PR (50%)
 - 5 SD (4 with >10% reduction in SLD of tumors)
- WT MET:
 - 5/57 (9%)

- Other *MET* alterations

- *MET* amplification (N=2): No responses
- Gain chromosome 7 (N=18): ORR 5%

Regression of a renal tumor in a patient with HPRC treated with Foretinib

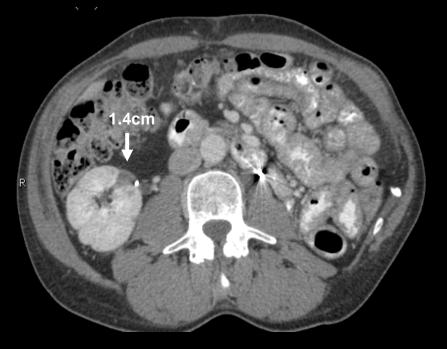


Pre-Treatment

Srinivasan, et al ASCO 2009

Regression of a Renal Tumor in a Patient with HPRC Treated with Foretinib



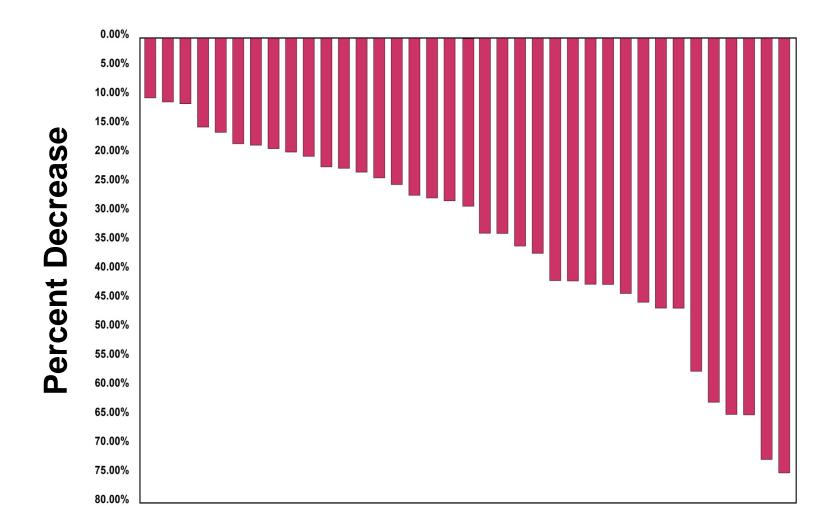


Pre-Treatment

Following 49 cycles of therapy

Srinivasan, et al ASCO 2009

Targeted Lesions in Patients with Germline *MET* Mutations



 $J_{\rm OURNAL \ OF \ CLINICAL \ ONCOLOGY}$

ORIGINAL REPORT

Biomarker-Based Phase II Trial of Savolitinib in Patients With Advanced Papillary Renal Cell Cancer

Toni K. Choueiri, Elizabeth Plimack, Hendrik-Tobias Arkenau, Eric Jonasch, Daniel Y.C. Heng, Thomas Powles, Melanie M. Frigault, Edwin A. Clark, Amir A. Handzel, Humphrey Gardner, Shethah Morgan, Laurence Albiges, and Sumanta Kumar Pal

European Journal of Cancer 87 (2017) 147-163



Original Research

Crizotinib achieves long-lasting disease control in advanced papillary renal-cell carcinoma type 1 patients with *MET* mutations or amplification. EORTC 90101 CREATE trial^{\Rightarrow}



Patrick Schöffski ^{a,b,*}, Agnieszka Wozniak ^b, Bernard Escudier ^c,

Impact of Met Status on Efficacy

| | Foretinib (N=74) | Crizotinib (N=23) | Savolitinib (N=109) |
|--|------------------|-------------------|---|
| Histology | All papillary | Type 1 Papillary | All Papillary |
| Stratification by Met Status (Post Hoc) | Yes | Yes | Yes |
| Stratification Criteria | Met Mutation | Met TK Mutation | Met TK Mutation Chromosome 7 Gain Focal Met amplification |
| Number MET + | 10 | 4 | 44 |
| Number MET- | 57 | 16 | 46 |
| ORR | | | |
| MET+ | 50% | 50% | 18% |
| MET- | 9% | 6% | 0% |
| PFS (Median) | | | |
| MET+ | NA | ? | 6.2 |
| MET- | | 3 | 1.4 |

Impact of Met Status on Efficacy

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| Number MET + Number MET- | 10 57 | 4 16 | 44 46 |
| ORR MET+ MET- | 50% 9% | 50% 6% | 18% 0% |
| PFS (Median) MET+ MET- | NA | ? 3 | 6.2 1.4 |

Impact of Met Status on Efficacy

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| Number MET + Number MET- | 10 57 | 4 16 | 44 46 |
| ORR MET+ MET- | 50% 9% | Comparable to | 18% 0% |
| PFS (Median) MET+ MET- | NA | (Historical) | 6.2 1.4 |

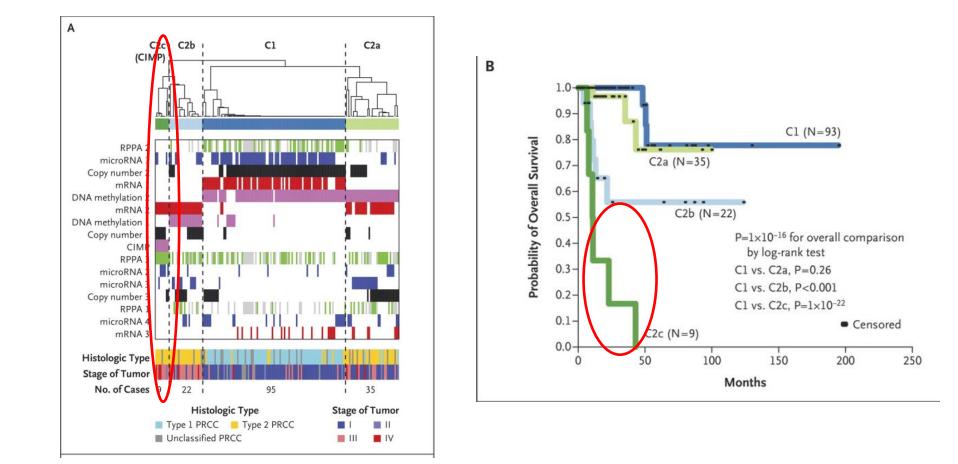
<u>Hereditary Leiomyomatosis</u> <u>Renal Cell Carcinoma: HLRCC</u>

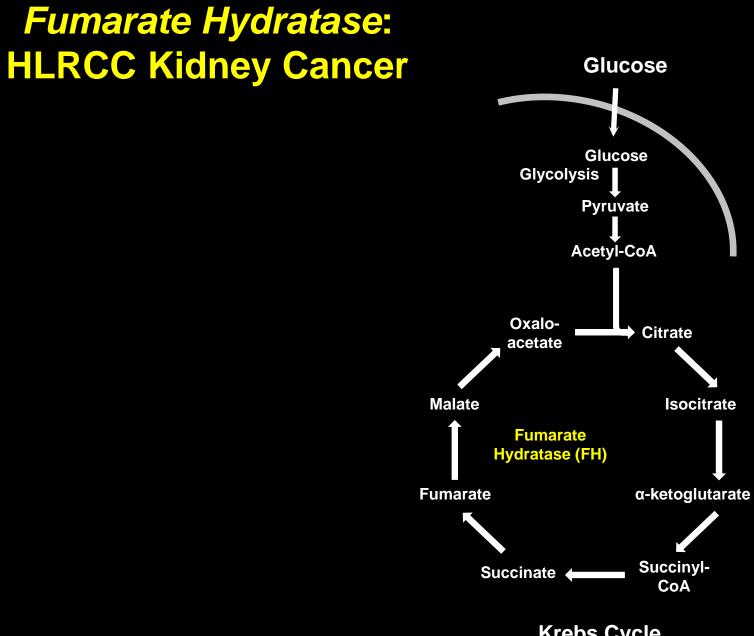
Cutaneous leiomyomas

• Uterine leiomyomas (fibroids)

• Renal cell carcinoma (Type 2 papillary RCC)

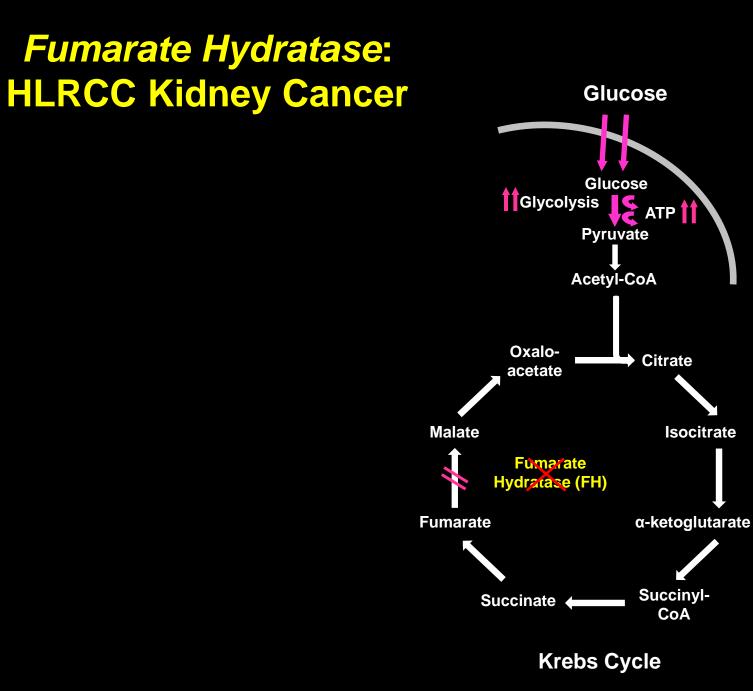
TCGA: Genetic Profiling of Papillary RCC: HLRCC Associated with Poor Prognosis



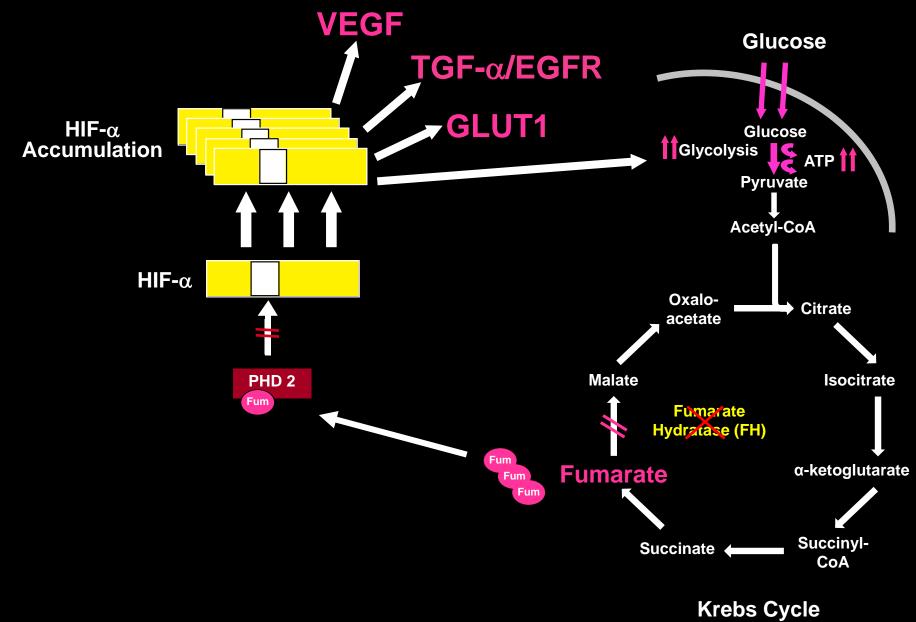


Isaacs, Cancer Cell, 2005 Tong, Cancer Cell, 2011

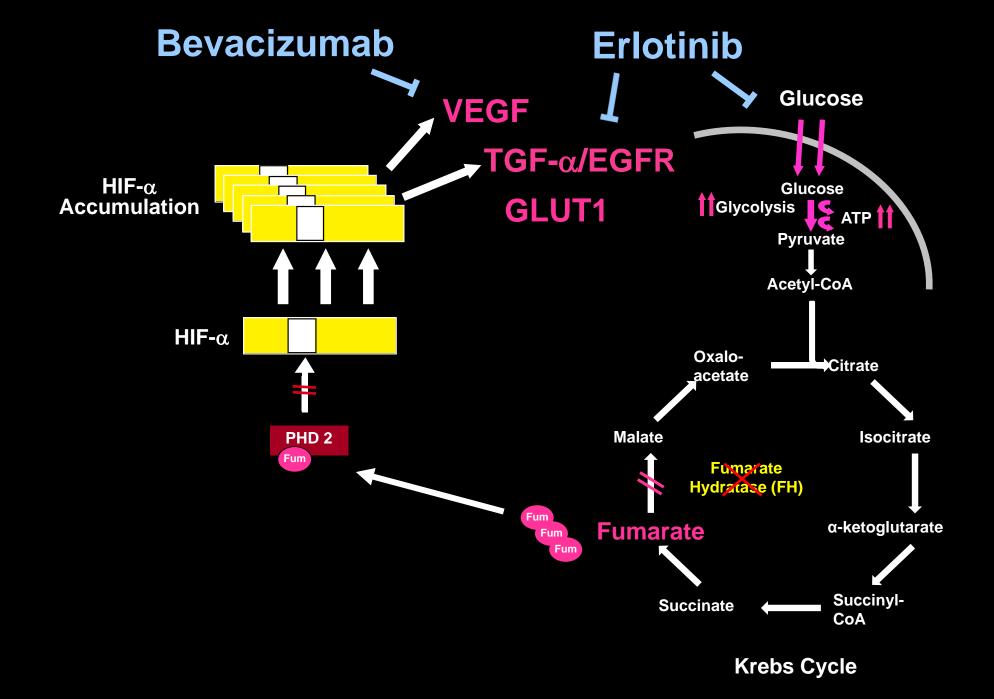
Krebs Cycle



Cancer Cell, 2005 Cancer Cell, 2011



Cancer Cell, 2005 Cancer Cell, 2011



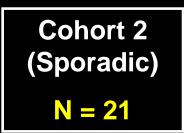
Study Design



N=41

Bevacizumab 10mg/kg IV q2 weeks plus Erlotinib150 mg PO daily

Cohort 1 (HLRCC) N = 20



Patient Demographics

| | Cohort 1 [HLRCC] | Cohort 2 [Sporadic] | Total |
|---------------------------|---------------------|------------------------|--------------|
| Number of Patients | 20 | 21 | 41 |
| Median Age (range), years | 46 (22 – 63) | 55 (35 – 73) | 52 (22 – 73) |
| Gender (%) | | | |
| Male | 11 (55%) | 15 (71%) | 26 (63%) |
| Female | 9 (45%) | 6 (29%) | 15 (37%) |
| MSKCC Risk Groups | | | |
| Favorable | 5 | 1 | 6 (15%) |
| Intermediate | 12 | 17 | 29 (70%) |
| Poor | 3 | 3 | 6 (15%) |
| Prior Systemic Therapy | | | |
| Νο | 14 | 9 | 23 (56%) |
| Yes | 6 | 12 | 18 (44%) |

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| No | 14 | 9 | 23 (56%) |
| Yes | 6 | 12 | 18 (44%) |

Bevacizumab plus Erlotinib in Papillary RCC - Efficacy

| Best Response by RECIST | Cohort 1 [HLRCC] (%) | Cohort 2 [Sporadic] (%) | Total (%) |
|------------------------------------|-------------------------|----------------------------|-----------|
| | N=20 | N=21 | |
| Confirmed Partial Response (PR) | 13 (65%) | 6 (29%) | 19 (46%) |
| Overall Response Rate | 65% | 29% | 46% |
| Stable disease (SD) | 7 (35%) | 13 (62%) | 20 (49%) |
| Disease Control Rate (SD+PR) | 100% | 91% | 95% |

36 Year Old Woman with HLRCC Associated Papillary RCC





 The term kidney cancer encompasses several different subtypes

 Individual subtypes are characterized by distinct genetic, clinical and histologic features

- Understanding genetic alterations is
 - Critical to the development of tailored treatment options
 - Increasingly becoming an integral part of the management of kidney cancer

Acknowledgements

<u>Urologic Oncology</u> Marston Linehan, M.D. Gennady Bratslavsky, M.D Mark Ball, M.D..

Clinical Team Julia Friend, PA-C Martha Ninos, RN Cheryl Royce, NP Erin Purcell, RN Geri Hawks, RN Andy Gillespie, RN Caitlin Drew, RN Debbie Nielsen, RN

<u>Berton Zbar, Ph.D</u> Len Neckers, Ph.D. Don Bottaro, Ph.D.

<u>UOB laboratories</u> Christopher Ricketts, Ph.D. Roma Pahwa, Ph.D. Cathy Vocke, Ph.D. Carole Sourbier, Ph.D. Youfeng Yang, M.S. Robert Worrell, Ph.D <u>Laboratory of Pathology</u> Maria Merino, M.D. Vanessa Moreno, M.D. Sara Gil Hernandez, M.D.

Ophthalmology Emily Chew, M.D Henry Wiley, M.D.

<u>Neurosurgery</u> Kareem Zaghloul, M.D. Prashant Chittiboina, M.D.

<u>Dermatology</u> Ed Cowan, M.D. Heidi Kong, M.D.

Radiology/Nuc Med Peter Choyke, M.D. Ashkan Malayeri, M.D. Clara Chen, M.D. Mark Ahlman, M.D. Brad Wood, M.D. Venkatesh Krishnaswamy, M.D. <u>UOB Staff</u> Rabindra Gautam Donna Drake Kristin Choo James Peterson Gabby Coello Cristiane Leite Janet Gichonge

<u>UOB Fellows</u> Eric Singer, M.D Brian Shuch, M.D. Abhinav Sidana, M.D. Mark Ball, M.D. Vladimir Valera, M.D.