

# Precision Medicine in RCC and Clinical Trial Design

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# Disclosures

- Research support: BMS, MSD, Novartis, Pfizer
- Consultancy (all non-remunerated): BMS, GSK, MSD, Pfizer, Novartis, Roche/Genentech

# Overview

- What is precision medicine?
- Are there any examples in kidney cancer?
- Why have advances in genomics not really impacted clinical practice?
- What are the clinical problems in this disease?
- How does this inform trial design?
- What does the future hold?

# What is precision medicine?

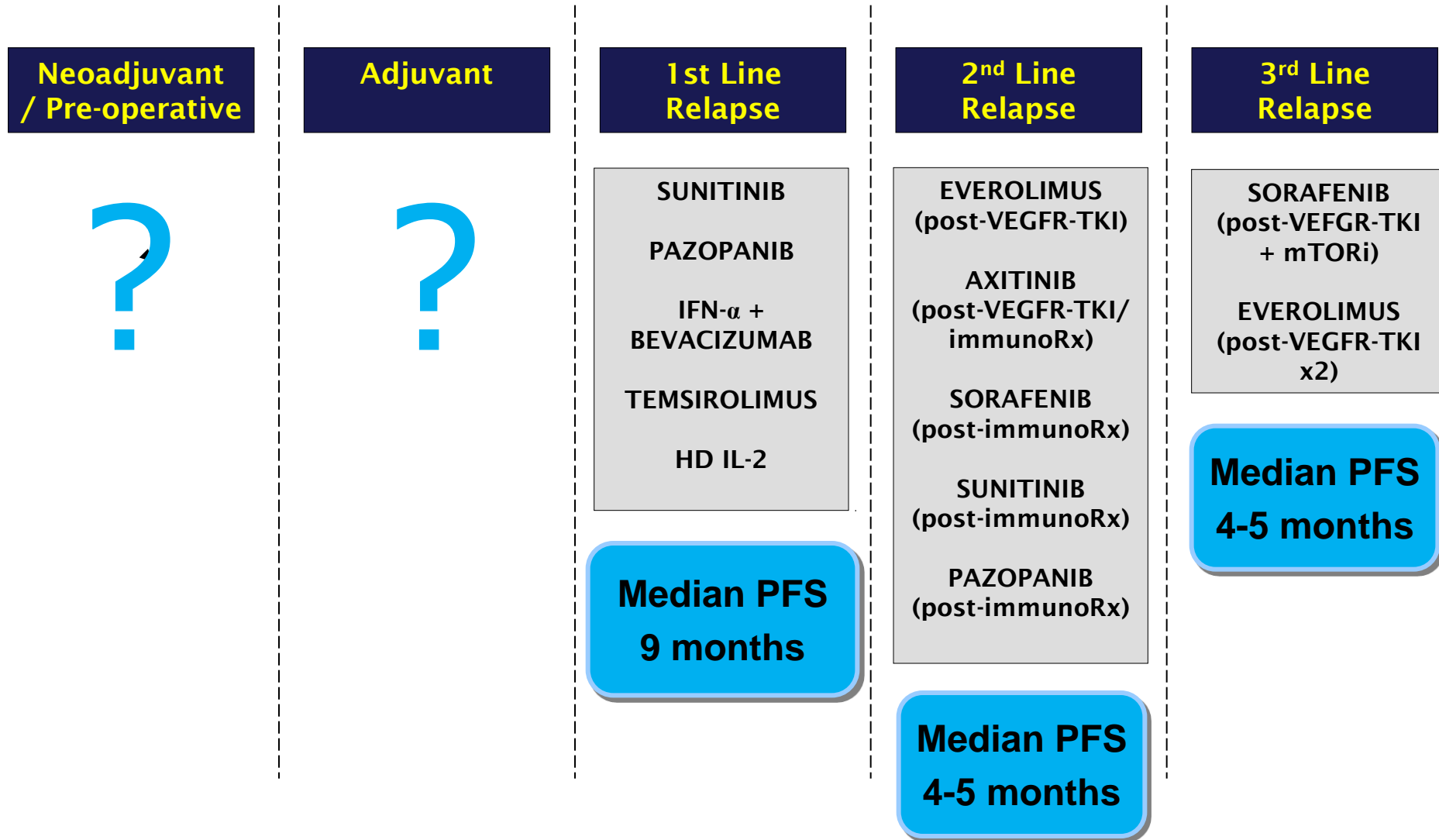
- 'Integration of molecular profiling with clinicopathological parameters to select optimal treatments for individual patients'
- Often refers to drug treatments and use of genomic information for patient care
- In RCC, we have lots of drugs...
- ...but all are cytokine, anti-VEGF or anti-mTOR



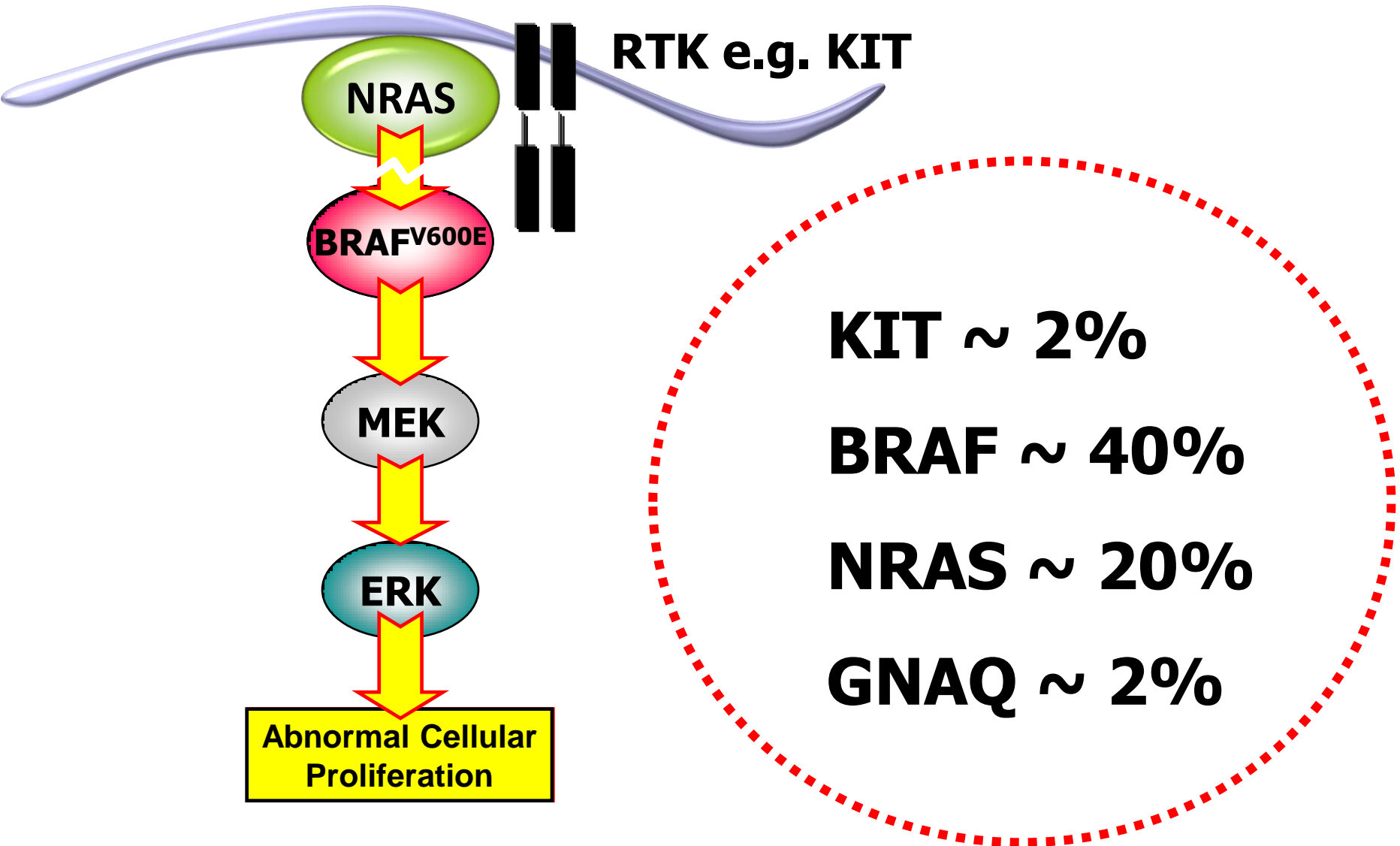
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- **Molecular profiling has no routine clinical role in 2014 in kidney cancer**

# Kidney Cancer Drug Therapy 2014



# Precision Medicine in Melanoma 2014



# Melanoma: Vemurafenib vs dacarbazine



# What about RCC?

- Most patients with ccRCC will get some tumour shrinkage from anti-VEGF therapy
- (Waterfall plot not as dramatic as targeting BRAF in melanoma though)
- Biggest problem in both diseases is acquired resistance to targeted therapy
- A lot is already known about this biologically **in patients** with melanoma (little in RCC)
- Clinically though we have a good idea in RCC which patients will not do well on therapy

# Why not more progress in RCC?

- RCC is **not** characterised by activating kinase mutations cf lung, melanoma, GIST etc
- Drug development for activating kinase mutations is tractable
- For tumour suppressor genes (which dominate RCC biology), targeting difficult
- Immunotherapy and anti-VEGF therapy act in the non-tumour compartment
- (NB No progress in developing predictive factors for these treatments in any tumour type)

# What about mTORi in RCC?

- If mTORi are predominantly targeting the tumour compartment, you would expect tumours with activation of this pathway to be sensitive to therapy...

# What about mTORi in RCC? RCTs:

- TEM vs IFN vs TEM + IFN in poor risk 1<sup>st</sup> line
- EVE vs sunitinib 1<sup>st</sup> line
- EVE + BEV vs IFN + BEV 1<sup>st</sup> line
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- EVE vs placebo post sunitinib/sorafenib
- TEM vs sorafenib post sunitinib **OS WORSE ON TEM**

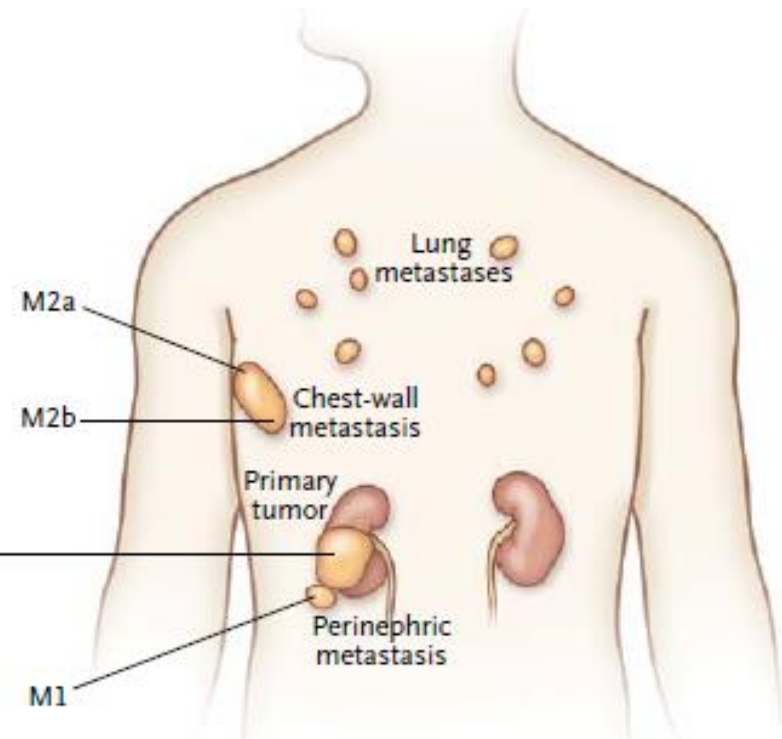
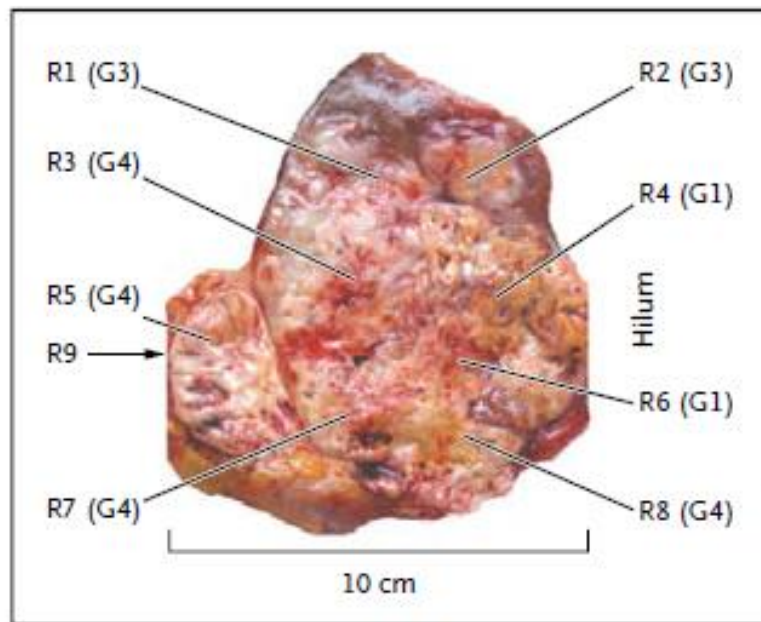
# What about mTORi in RCC? RCTs:

- TEM vs IFN vs TEM + IFN in poor risk 1<sup>st</sup> line
- *Choice of comparator arm controversial*
- *No other trials in poor risk group*
- *No yardstick for anti-VEGF activity*
- EVE vs placebo post sunitinib/sorafenib
- *Placebo control arm reasonable when trial recruited*

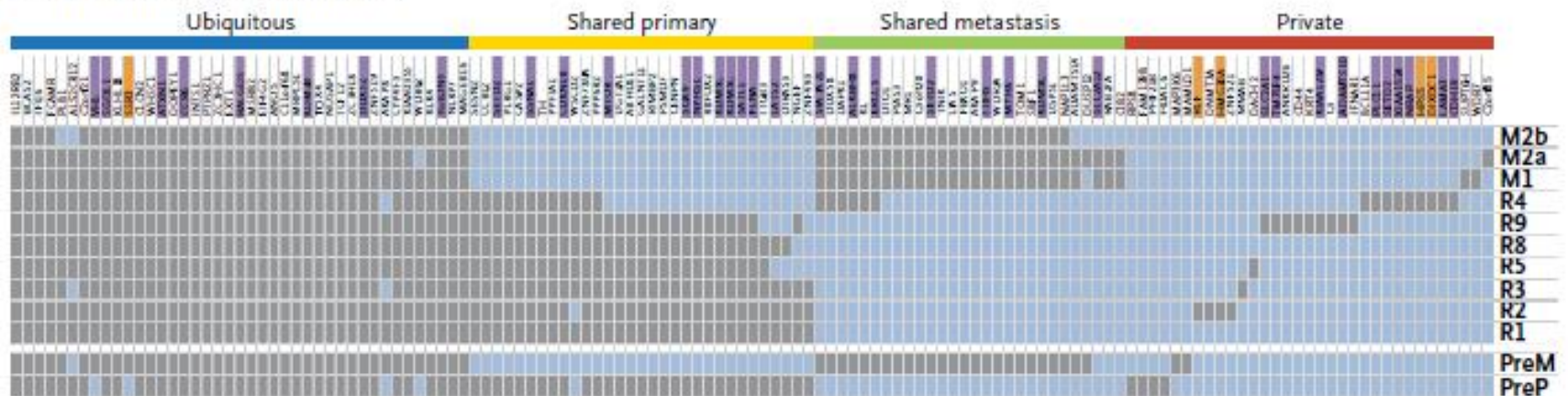
# What about mTORi in RCC?

- If mTORi are predominantly targeting the tumour compartment, you would expect tumours with activation of this pathway to be sensitive to therapy
- Clearly, on average, this is not the case in comparison with anti-VEGF therapy
- How could this be explained?
- Is there a subset of molecularly defined patients that might benefit?

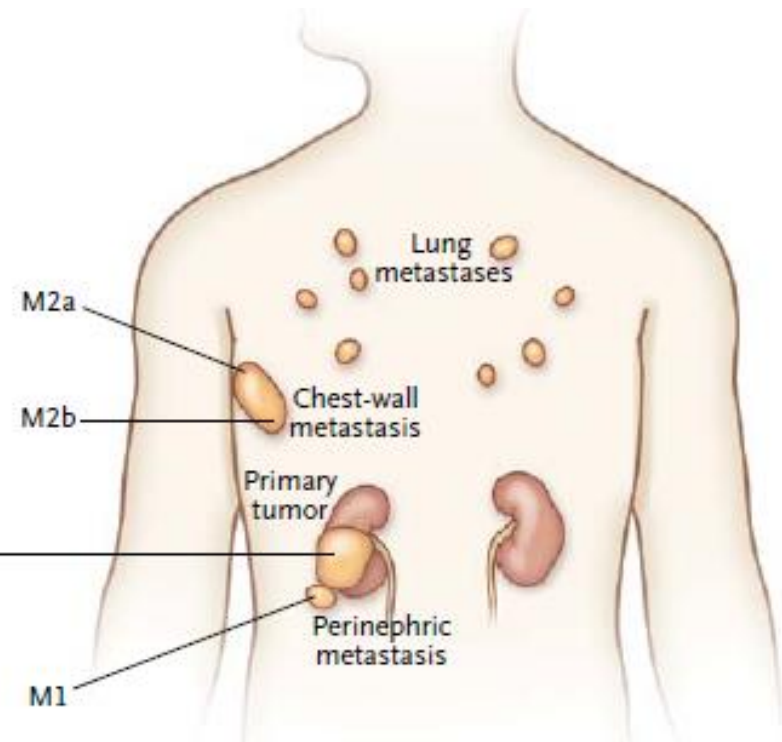
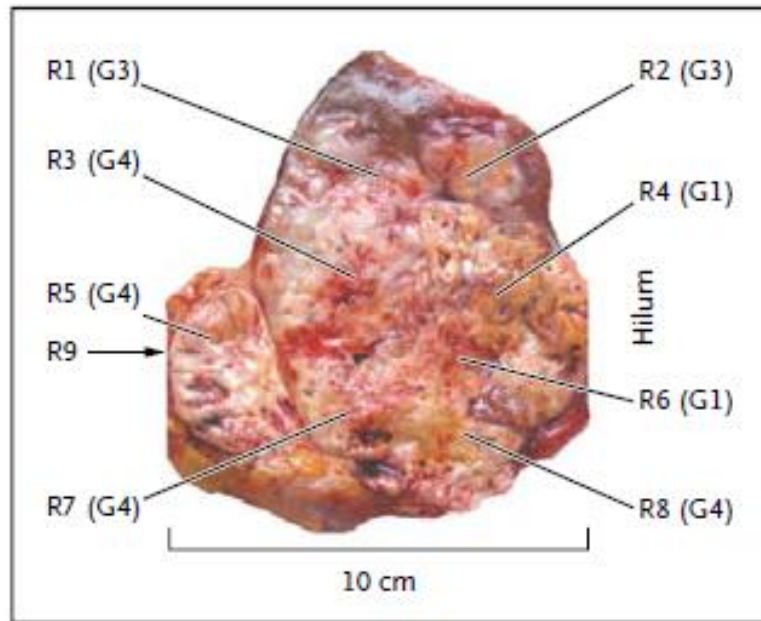
## A Biopsy Sites



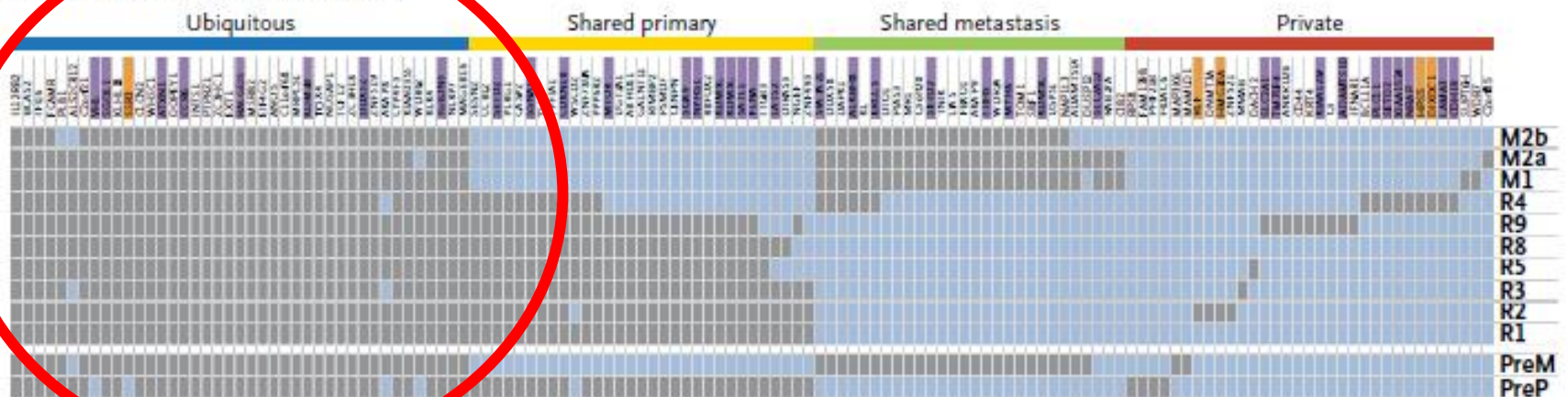
## B Regional Distribution of Mutations



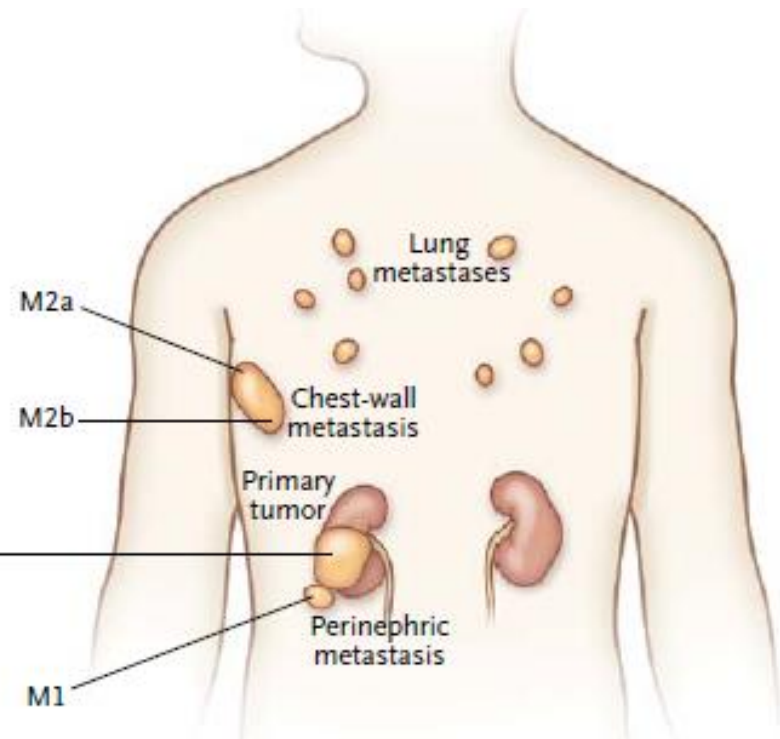
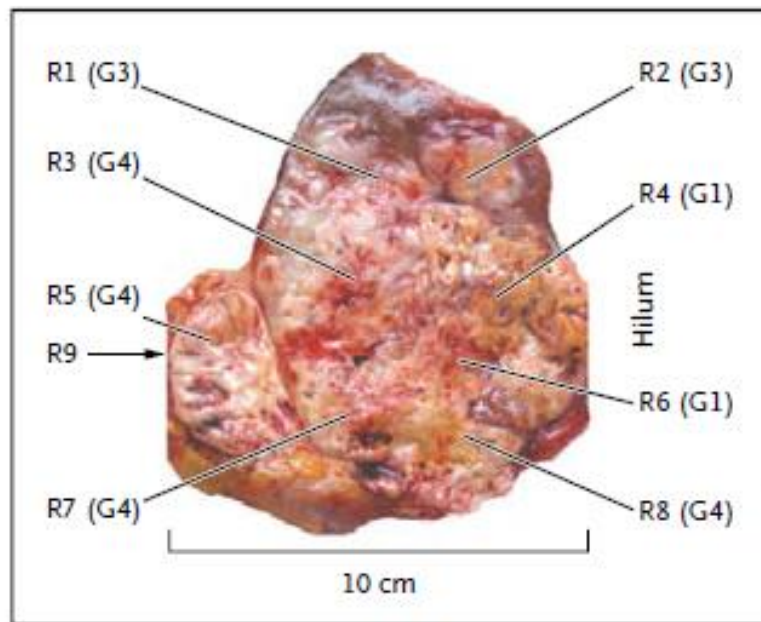
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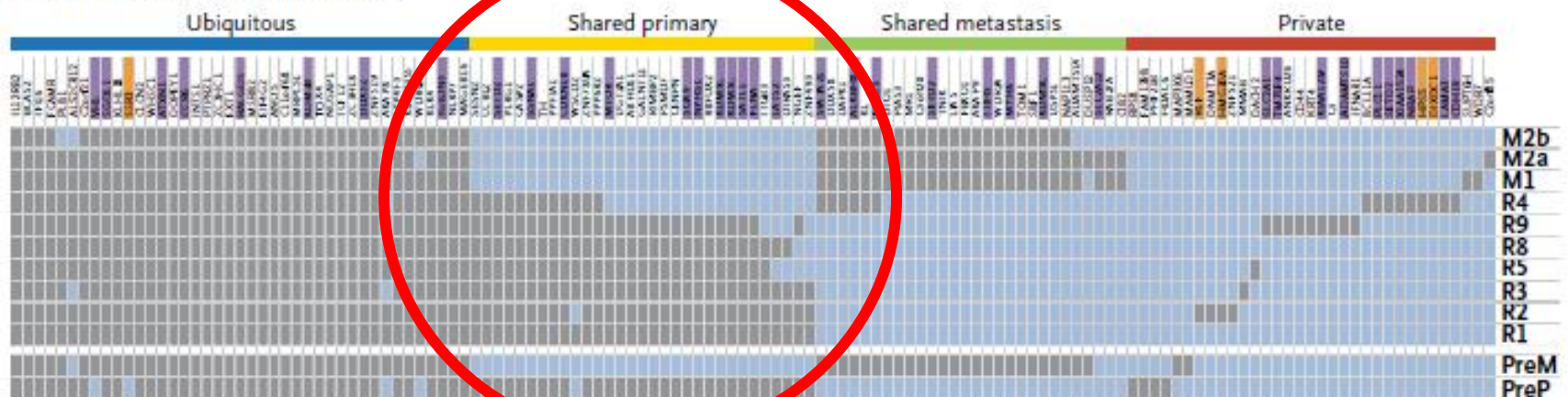
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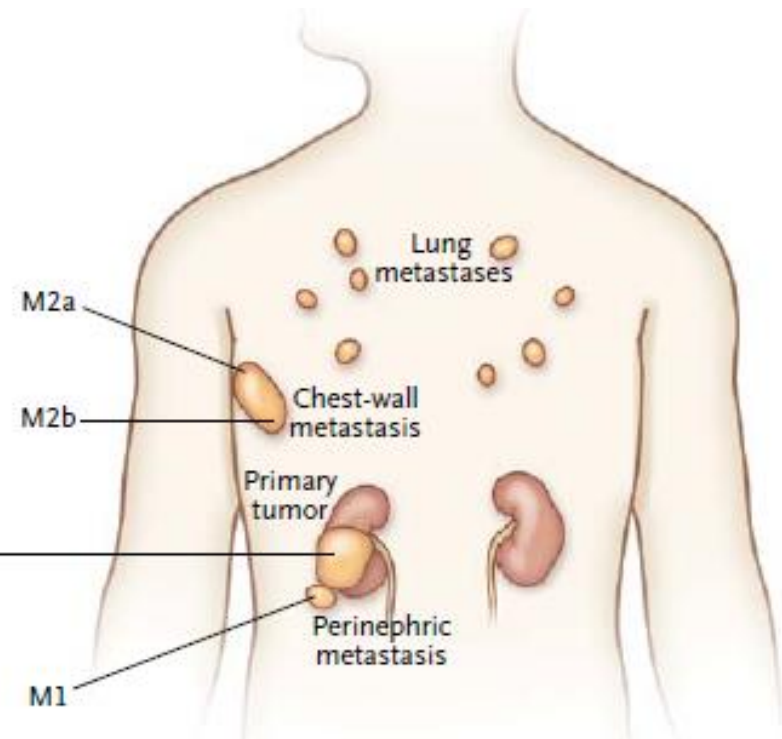
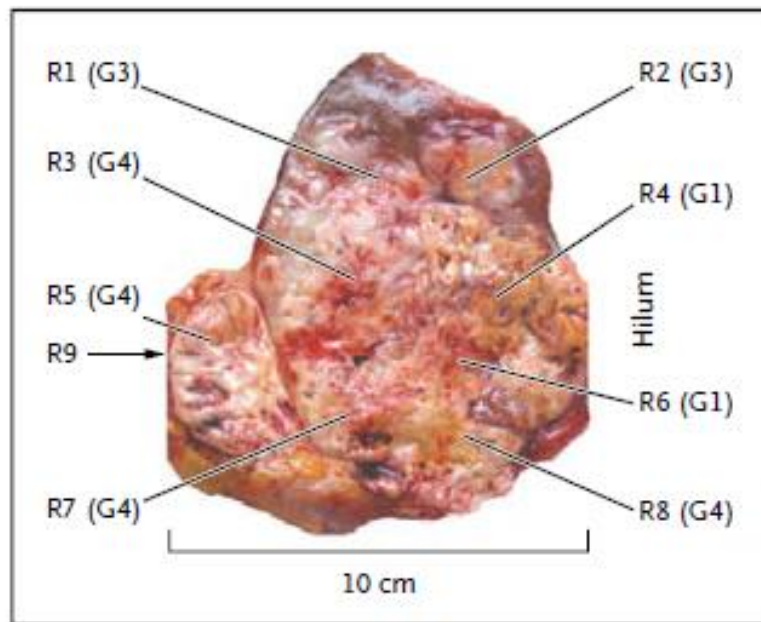
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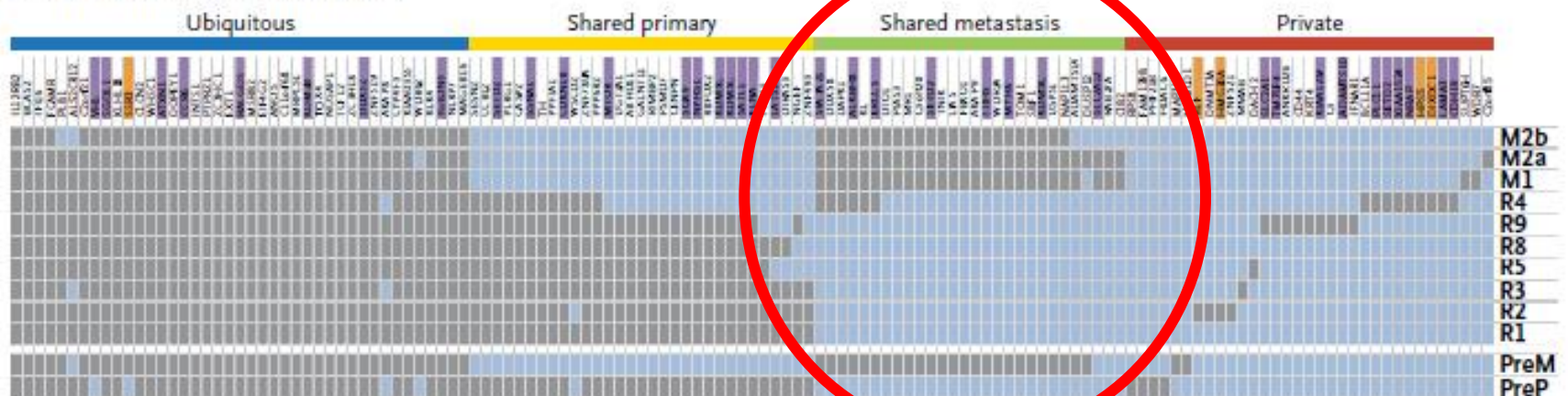
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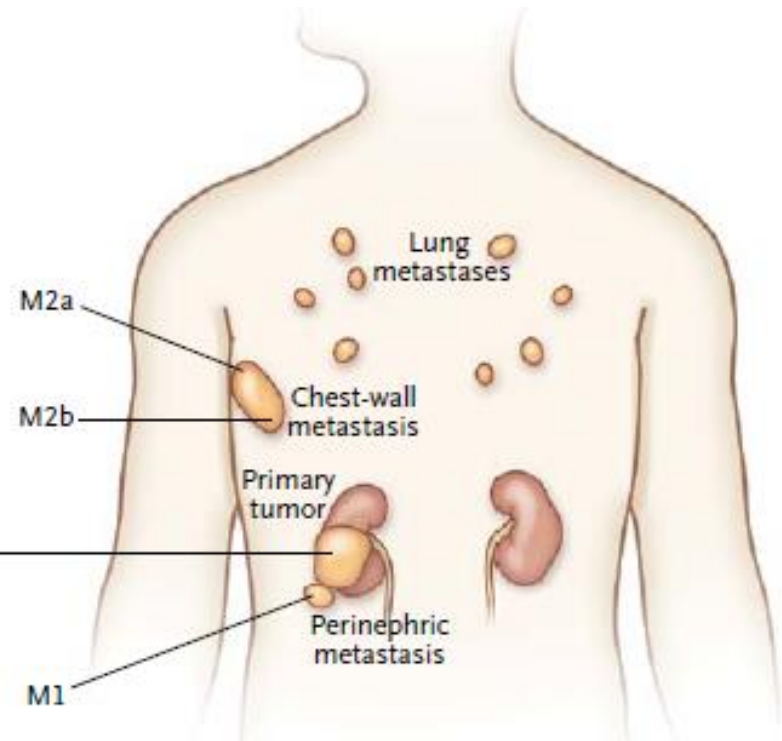
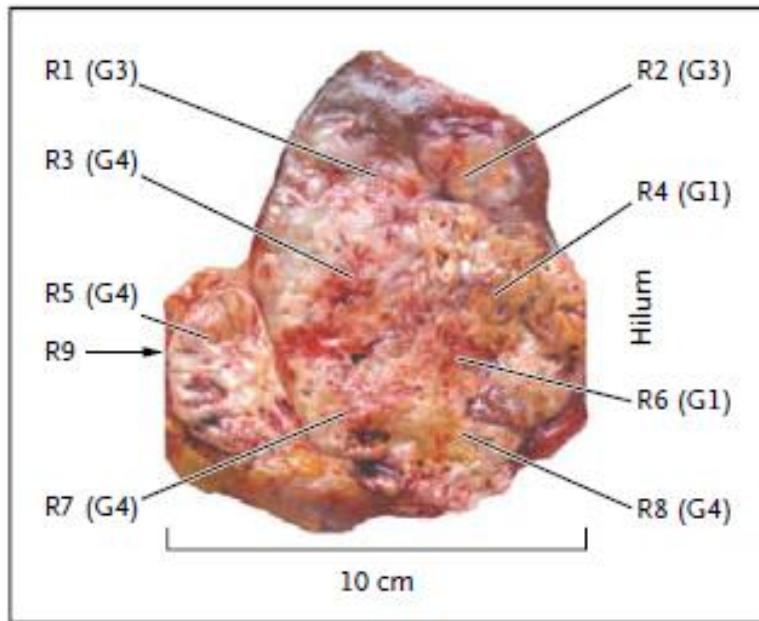
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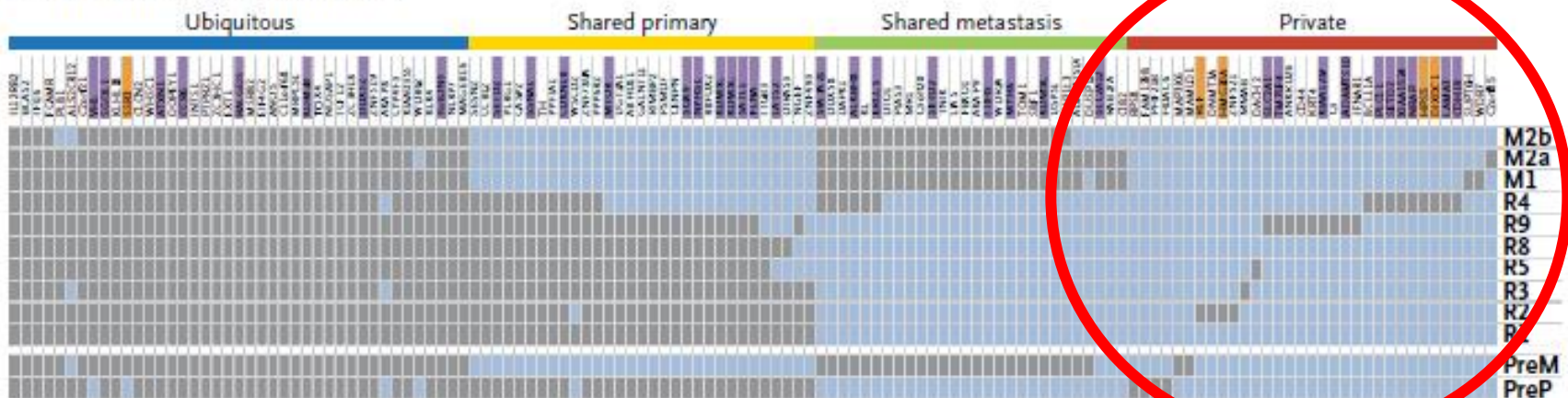
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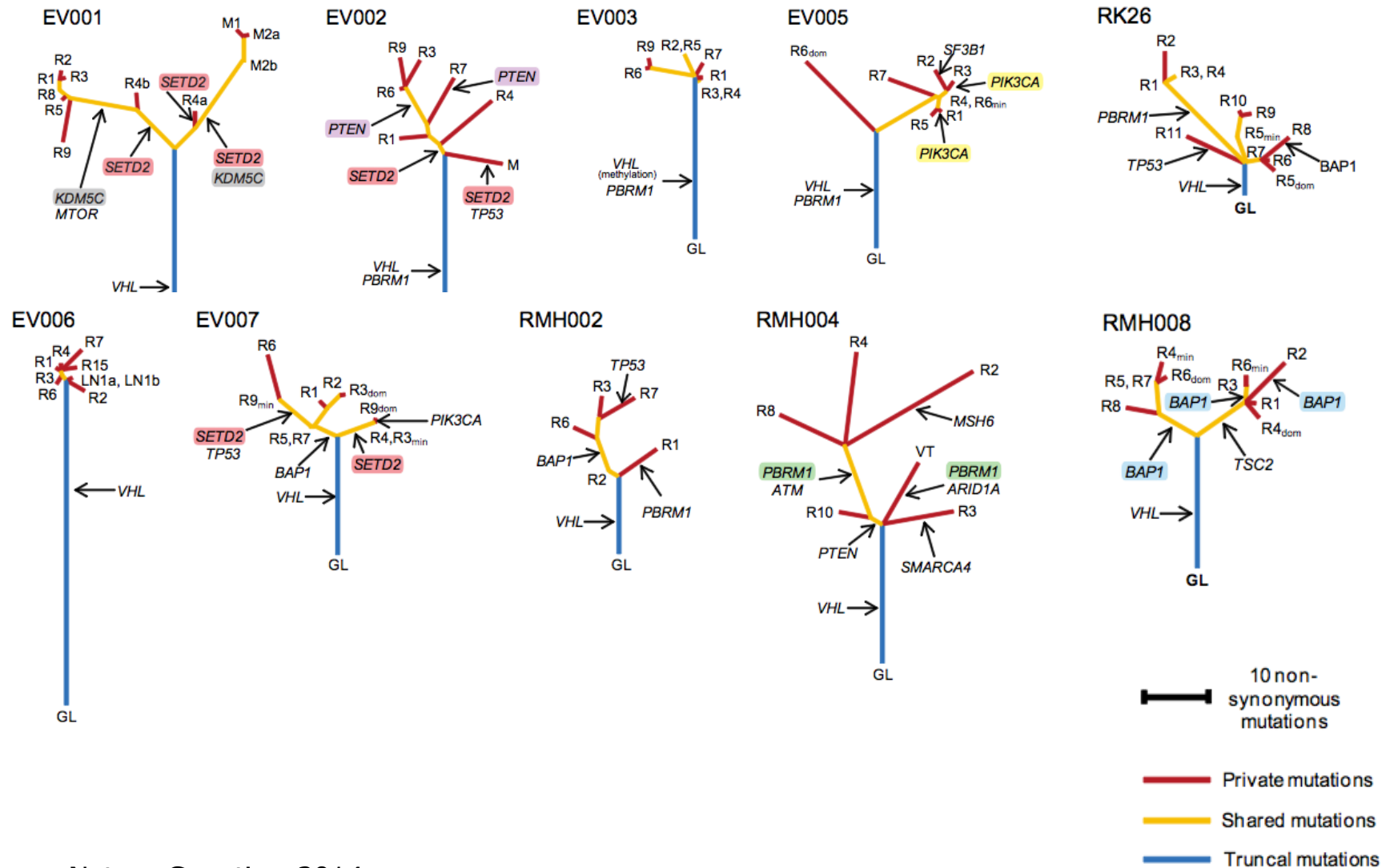


## B Regional Distribution of Mutations



**65% mutations are heterogeneous and not present in every biopsy**

# Branched Evolution in ccRCC



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**Table 1** Comparison of driver mutation prevalence in ccRCC samples

	Prevalence in TCGA samples ( <i>n</i> = 218 samples)	Prevalence in all M-seq samples ( <i>n</i> = 79 samples)	Prevalence in cases based on M-seq ( <i>n</i> = 10 cases)	Prevalence cases/prevalence M-seq samples
<i>PBRM1</i>	42%	39%	60%	1.5
<i>SETD2</i>	18%	27%	30%	1.1
<i>BAP1</i>	21%	24%	40%	1.7
<i>KDM5C</i>	7%	11%	10%	0.9
<i>TP53</i>	5%	5%	40%	8.0
<i>ATM</i>	3%	4%	10%	2.5
<i>ARID1A</i>	6%	1%	10%	10.0
<i>PTEN</i>	5%	10%	20%	2.0
<i>MTOR</i>	9%	8%	10%	1.3
<i>PIK3CA</i>	3%	4%	20%	5.0
<i>TSC2</i>	2%	4%	10%	2.5
PI3K-mTOR pathway	18%	28%	60%	2.2

# Target Trunks and Resolve Branches



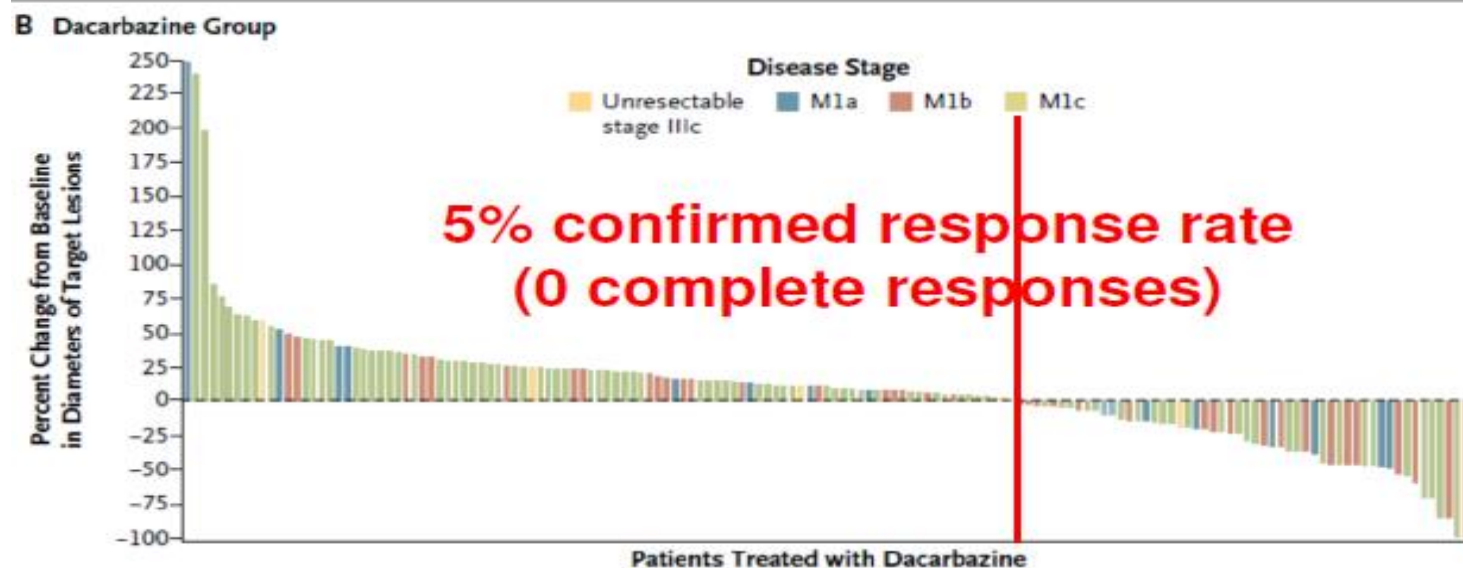
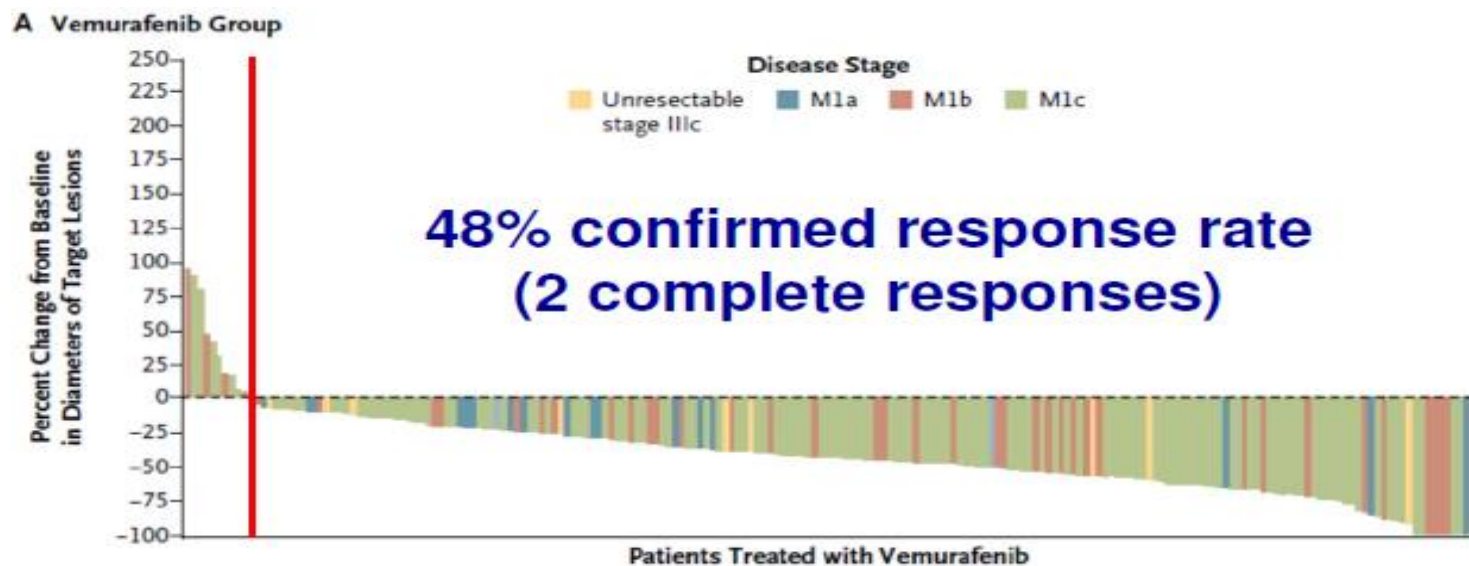
**Branched Genetic Events Present in  
Some Cancer Cells not others  
Dynamic during disease course**

**Monitor subclonal events to define  
drug resistance mechanisms**

**Trunk Genetic Events Present in  
Every Cancer Cell**

**DEFINE TRUNK DRIVERS**

# BRAF in Melanoma: 'Truncal Driver'



# mTORi in RCC summary

- mTORi active in RCC but less (on average in unselected patients) than anti-VEGF therapy
- mTOR pathway aberrations not infrequent
- Are they 'truncal drivers' though?
- Further study, particularly of metastatic sites / non-invasive technologies needed
- Understanding this could transform the use of mTORi in RCC

# What does this mean for clinical trial design?

- Relatively small 'tissue heavy' clinical trials are needed to understand these ideas better
- Need not involve drug therapy; disease evolution off therapy important too
- Our efforts to understand response and resistance to anti-VEGF and anti-mTOR therapy should continue
- We must also put the same efforts into investigating new drugs e.g. anti-PD1/PDL1

# What does this mean for clinical trial design?

- We need to frame our trials so that we are addressing important clinical issues e.g.
- Sarcomatoid histology does badly
- Patients presenting with mRCC do badly
- Patients presenting with mRCC unsuitable for cytoreductive nephrectomy do badly
- Patients with non-clear cell RCC do badly
- Patients with limited benefit from anti-VEGF do badly

# What does this mean for clinical trial design?

- We need to continue our engagement with industry to bring exciting drugs into RCC
- I would particularly highlight 2 trials
- Randomised phase 2: sunitinib vs MPDL3280A vs bevacizumab + MPDL3280A
- Well tolerated combination of drugs
- Cross-over to anti-PDL1 therapy is allowed
- Lack of cytoreductive nephrectomy allowed
- Sarcomatoid histology explicitly allowed

# What does this mean for clinical trial design?

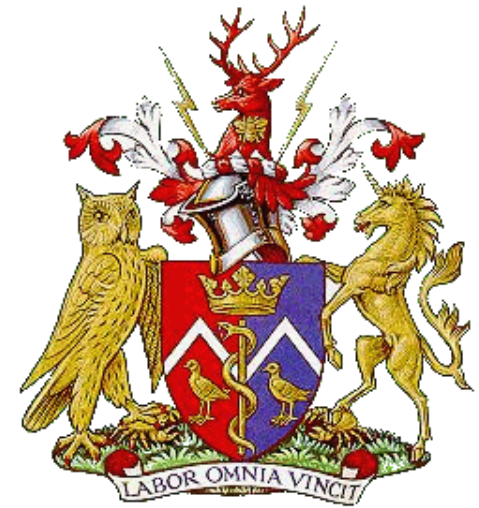
- The '214' trial: sunitinib vs ipilimumab + nivolumab ~1000 patient phase 3
- Melanoma clinicians familiar with ipilimumab + nivolumab and the '067' trial of ipi vs nivo vs the combination
- In melanoma, ipilimumab alone can durably control melanoma in 15-20%
- Early efficacy suggests activity in 40-50%
- Ipilimumab + nivolumab in RCC early efficacy ~ 40%

# Conclusions

- Major increase in understanding of RCC biology (especially genomic) last ~10 years
- This has yet to translate into molecular predictive factors for treatment in the clinic
- Inherently, there are some differences in RCC biology that may partly explain this
- Extensive efforts are being made and should continue with newer agents, particularly checkpoint inhibitors
- Clinical trial design must reflect clinical issues as well as molecular predictive factors

# Acknowledgements

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