



Memorial Sloan Kettering  
Cancer Center

# PD39-05: Prevalence and Landscape of Actionable Genomic Alterations in Renal Cell Carcinoma

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# Introduction

- Actionable genomic alterations include somatic mutations and structural alterations that predict response to targeted therapy
- Defining the landscape of actionable alterations in RCC may identify therapeutic targets and inform targeted therapy trials
- **OBJECTIVE:** describe the prevalence and landscape of actionable alterations and the corresponding evidence supporting the alteration as predictive of response to targeted therapy in RCC



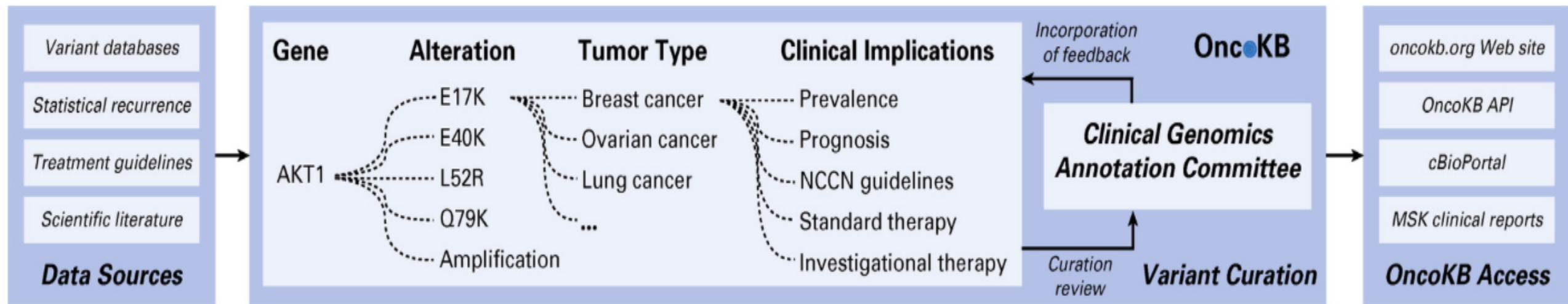
## Methods

- Institutional clinical sequencing database queried to include tumor samples sequenced across all cancers
- Actionable alterations with clinical/biologic evidence supporting an association with response to targeted therapy stratified by level of evidence using an oncology knowledge database (OncoKB)<sup>1</sup>

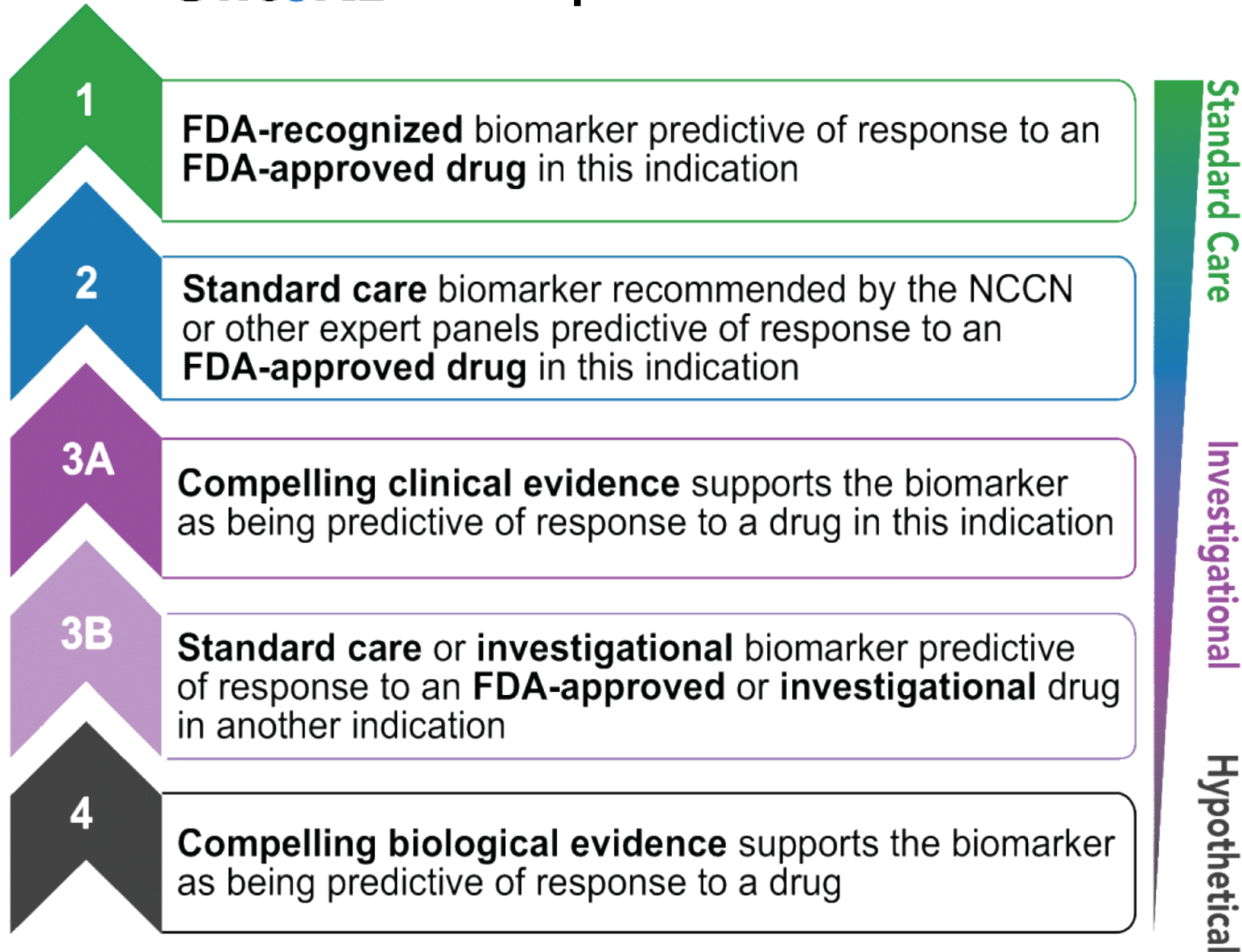
<sup>1</sup>Chakravarty et al. *JCO PO*. 2017

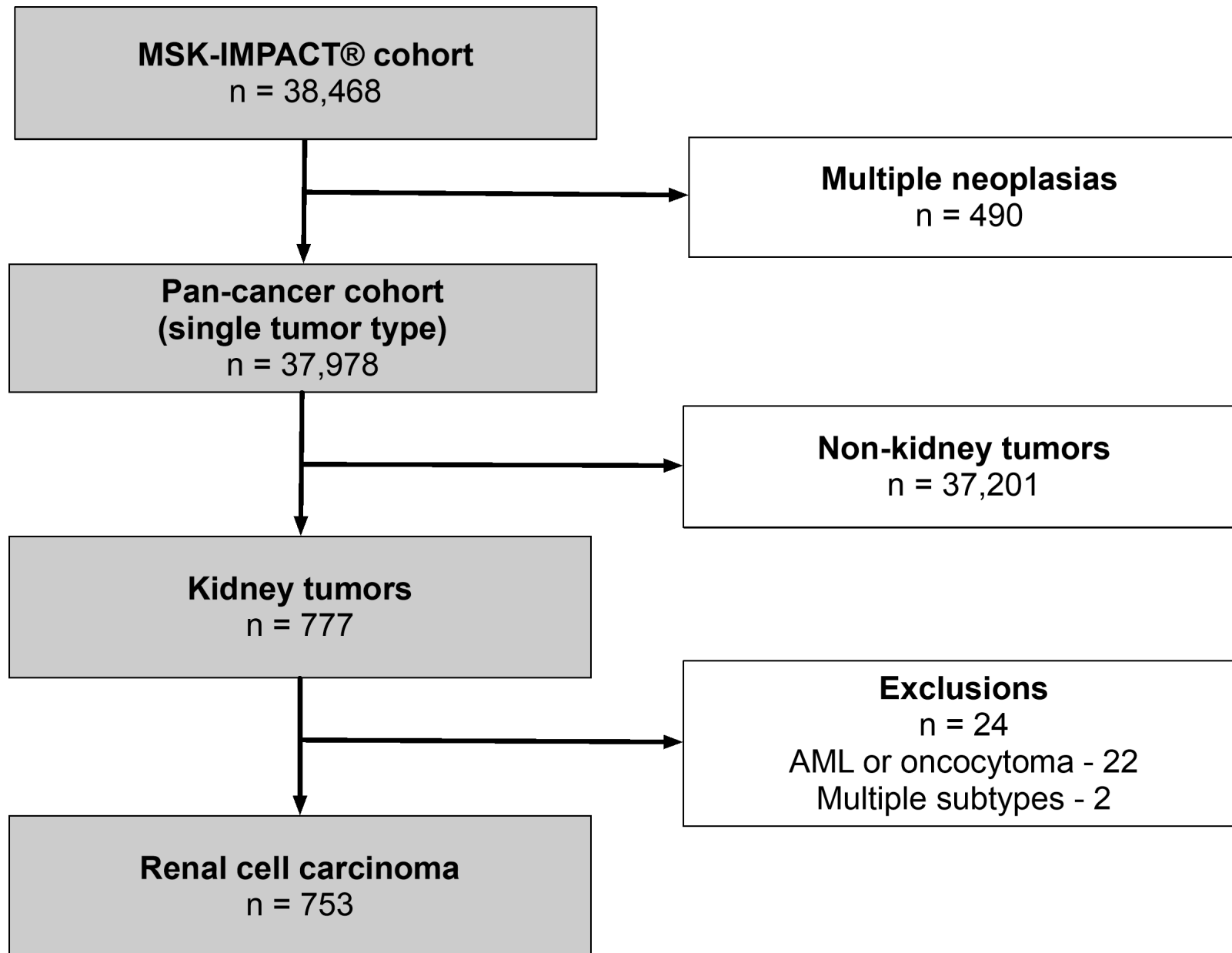


# OncoKB Workflow



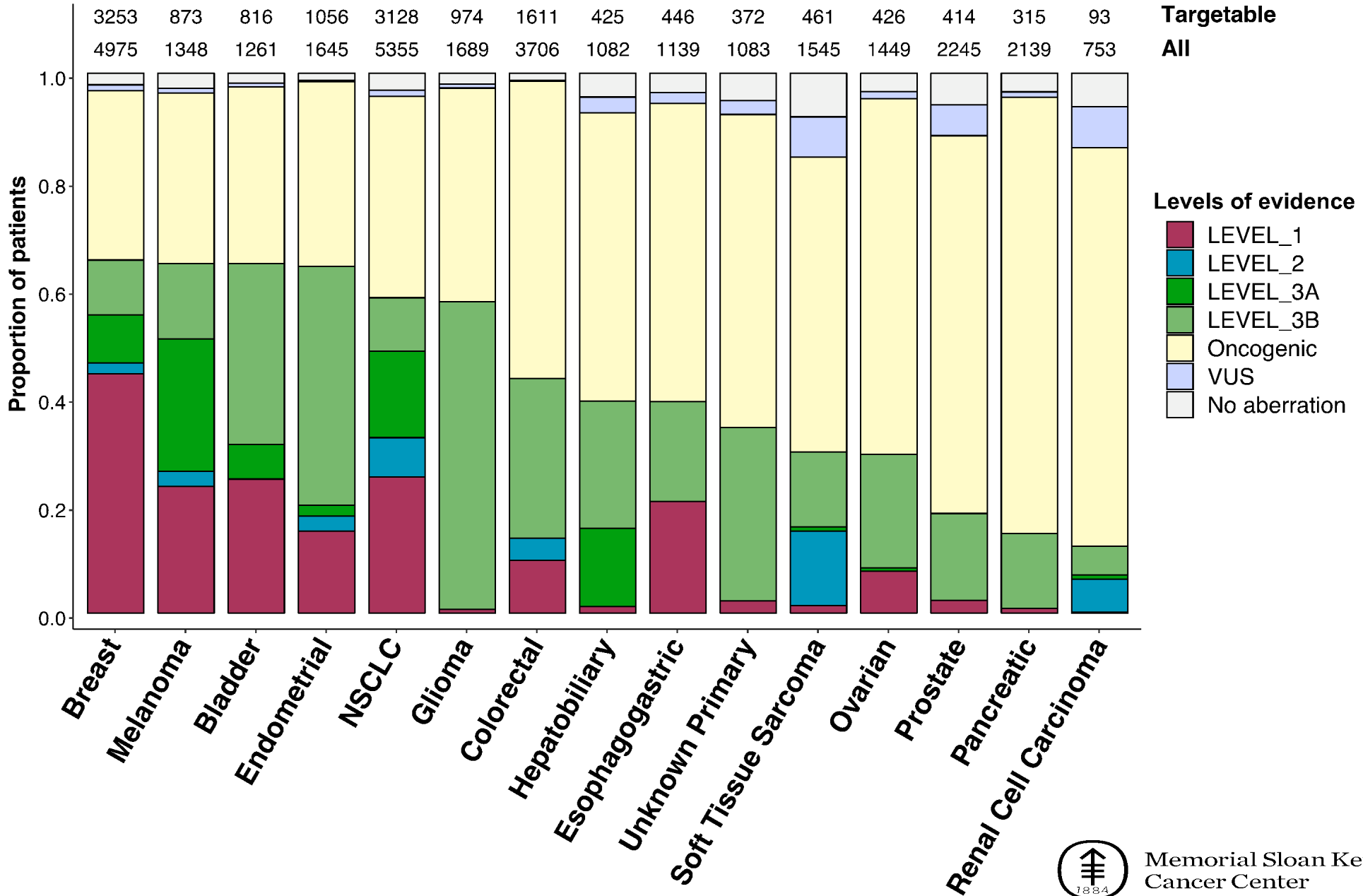
# OncKB Therapeutic Levels of Evidence





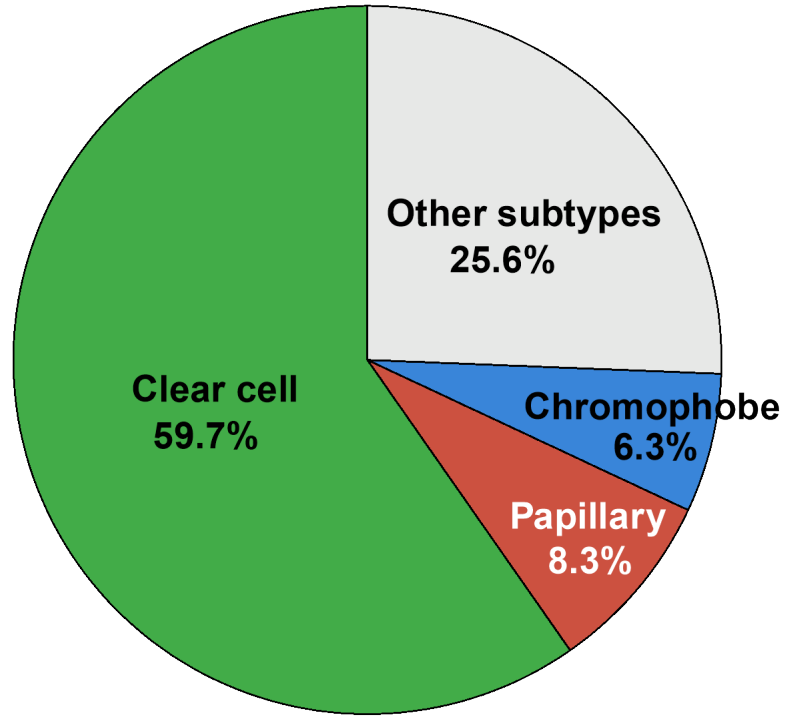
# Pan-cancer MSK-IMPACT cohort

## Targetable alterations by cancer type (15 most common)

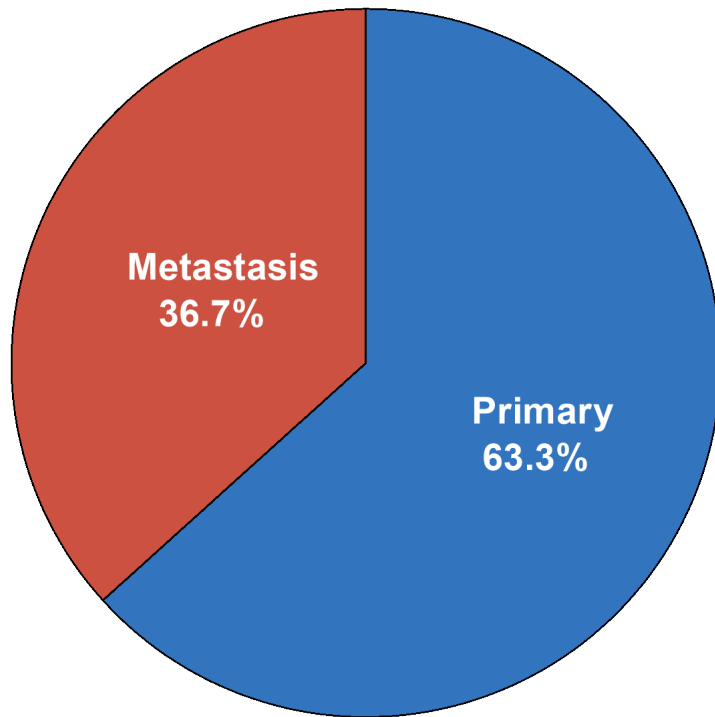




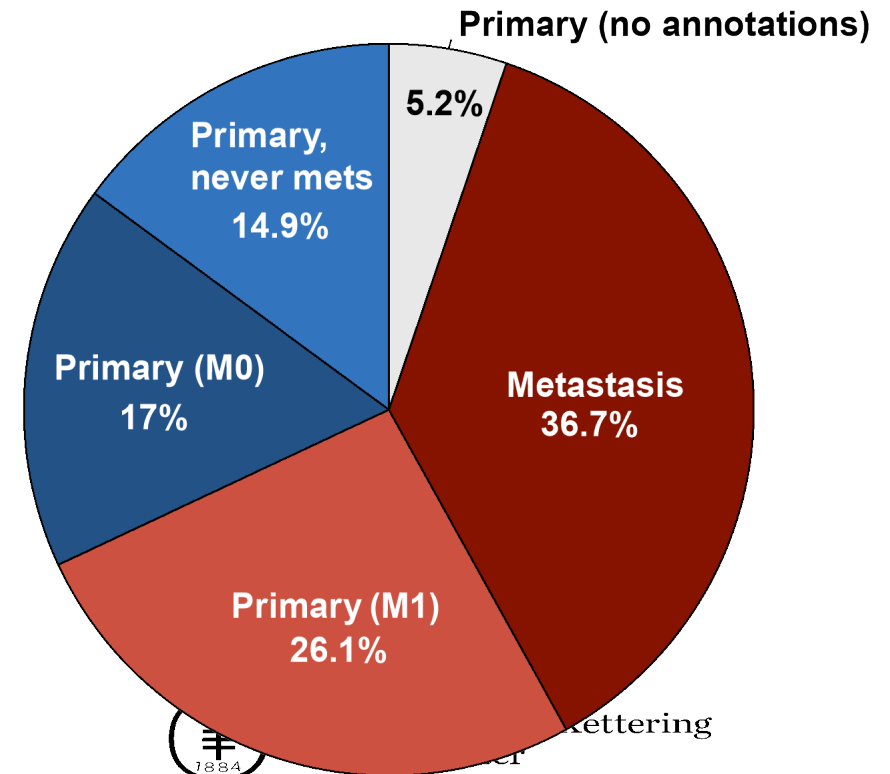
RCC histologic subtypes



Sample type

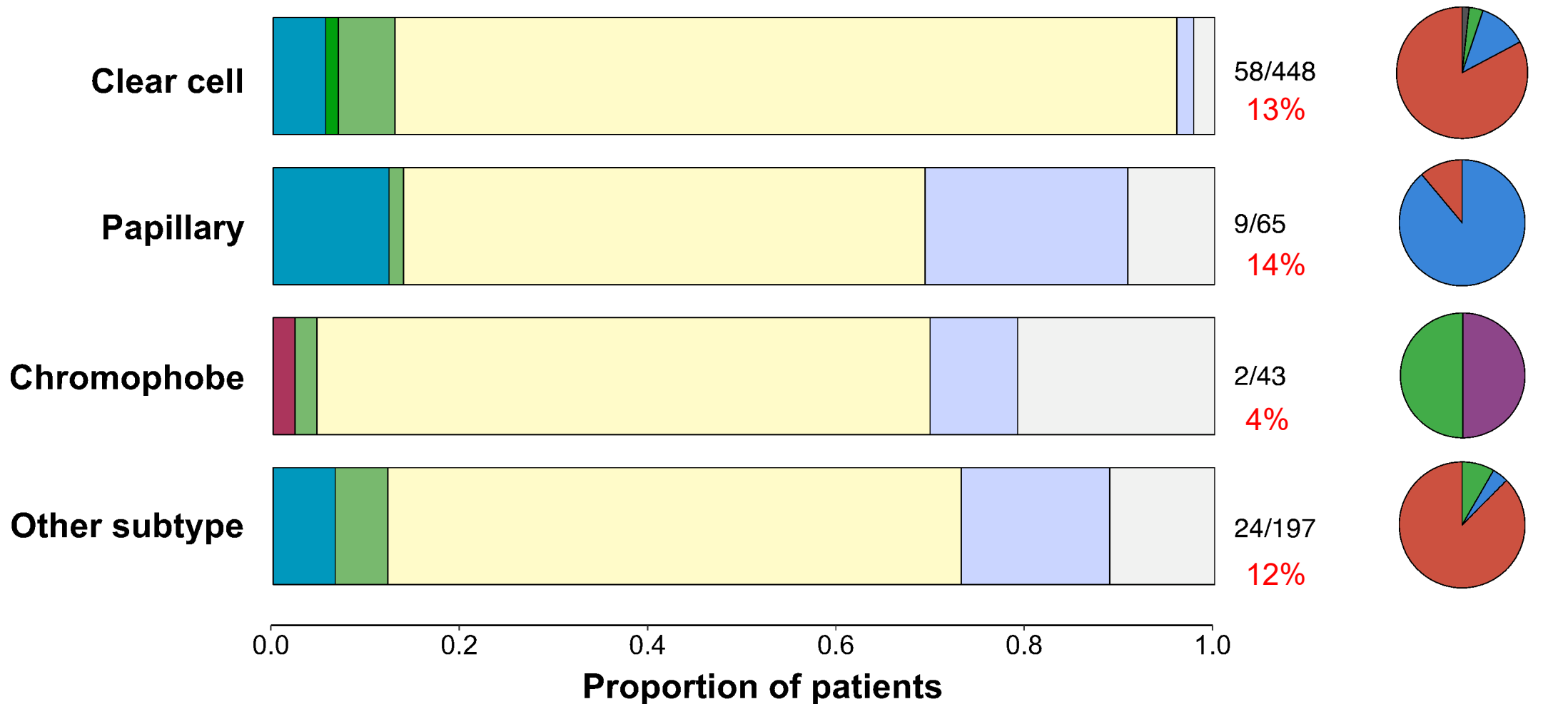


Clinical annotations

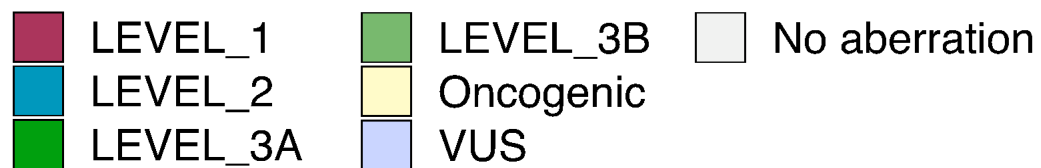




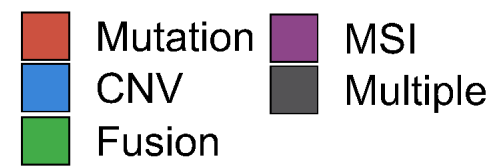
# Targetable alterations in Renal Cell Carcinoma (n=753)



## Levels of evidence

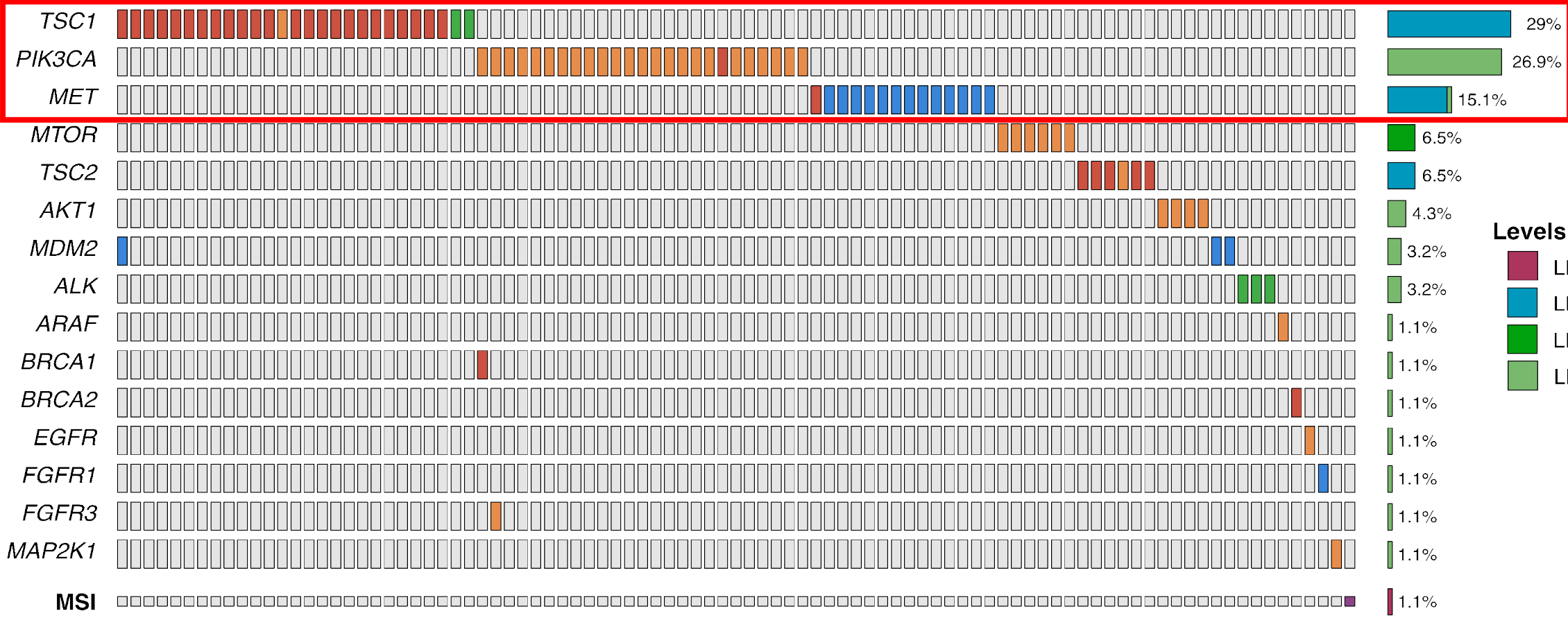


## Type of alteration



# Targetable alterations in RCC

RCC subtype 



**Levels of evidence**

- LEVEL\_1
- LEVEL\_2
- LEVEL\_3A
- LEVEL\_3B

**RCC subtype**

- Clear Cell
- Papillary
- Chromophobe
- Unclassified
- FH-Deficient
- Collecting Duct

**Type of somatic alteration**

- WT
- Truncating Mutation
- Fusion
- Missense Mutation
- Copy-number
- MSI



- SUMMARY

- The prevalence of actionable alterations in RCC is 12%
- Type of alteration varies by histologic subtype
  - Clear cell RCC → acquired somatic mutations
  - Papillary RCC → copy number variations
  - Chromophobe RCC → MSI/fusion (rare)

- NEXT STEPS

- Validation of results (TCGA, TRACERx)
- Clonality of alterations



# Acknowledgments

## Mentors/Co-authors

- **A. Ari Hakimi, MD**
- **Timothy Chan, MD**
- Robert Motzer, MD
- Martin Voss, MD
- Paul Russo, MD
- Jonathan Coleman, MD
- Maria Carlo, MD
- Chung-Han Lee, MD
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*We wish to thank our patients, their families, and their caregivers*