

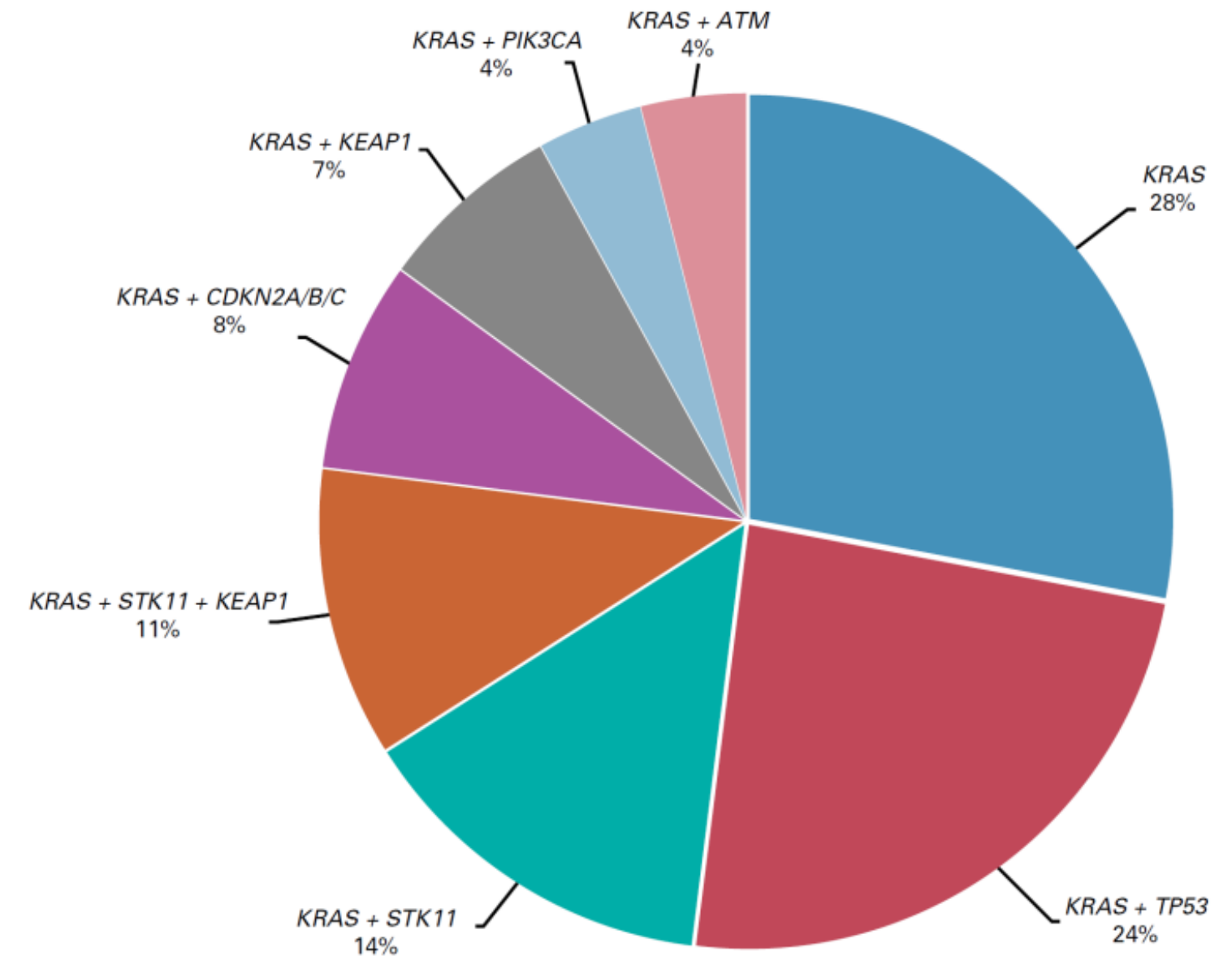
# Deciphering the Mechanism of Resistance to First Generation KRAS Inhibitors

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# Co-mutations as potentially predictive markers in *KRAS* mutated NSCLC

- Most common *TP53*, *STK11*, *KEAP1*
  - Arbour et al (n=330): *TP53* (42%), *STK11* (29%), *KEAP1* (24%)
  - Scheffler et al (n=1078): *TP53* (39%); (n=101 subset): *STK11* (20%), *KEAP1* (13%)
  - Aredo et al (Stanford; *KRAS*<sup>G12C</sup> mutation subtype): *TP53* 31% and *STK11* 27%
- Preclinical– differences in signaling, immune features, metabolic programming
- Prognostic
- Potentially predictive with treatment (e.g. immunotherapy)

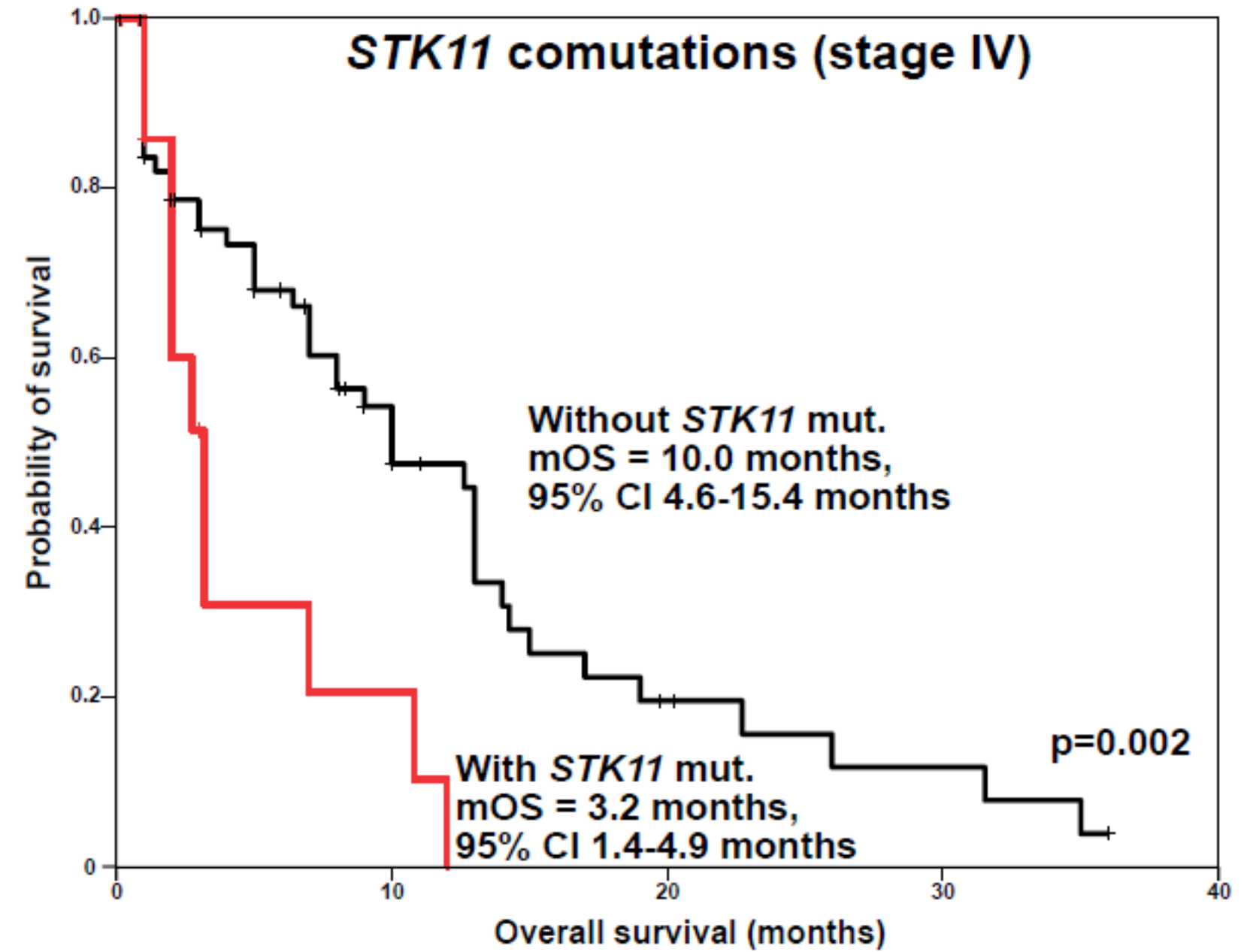
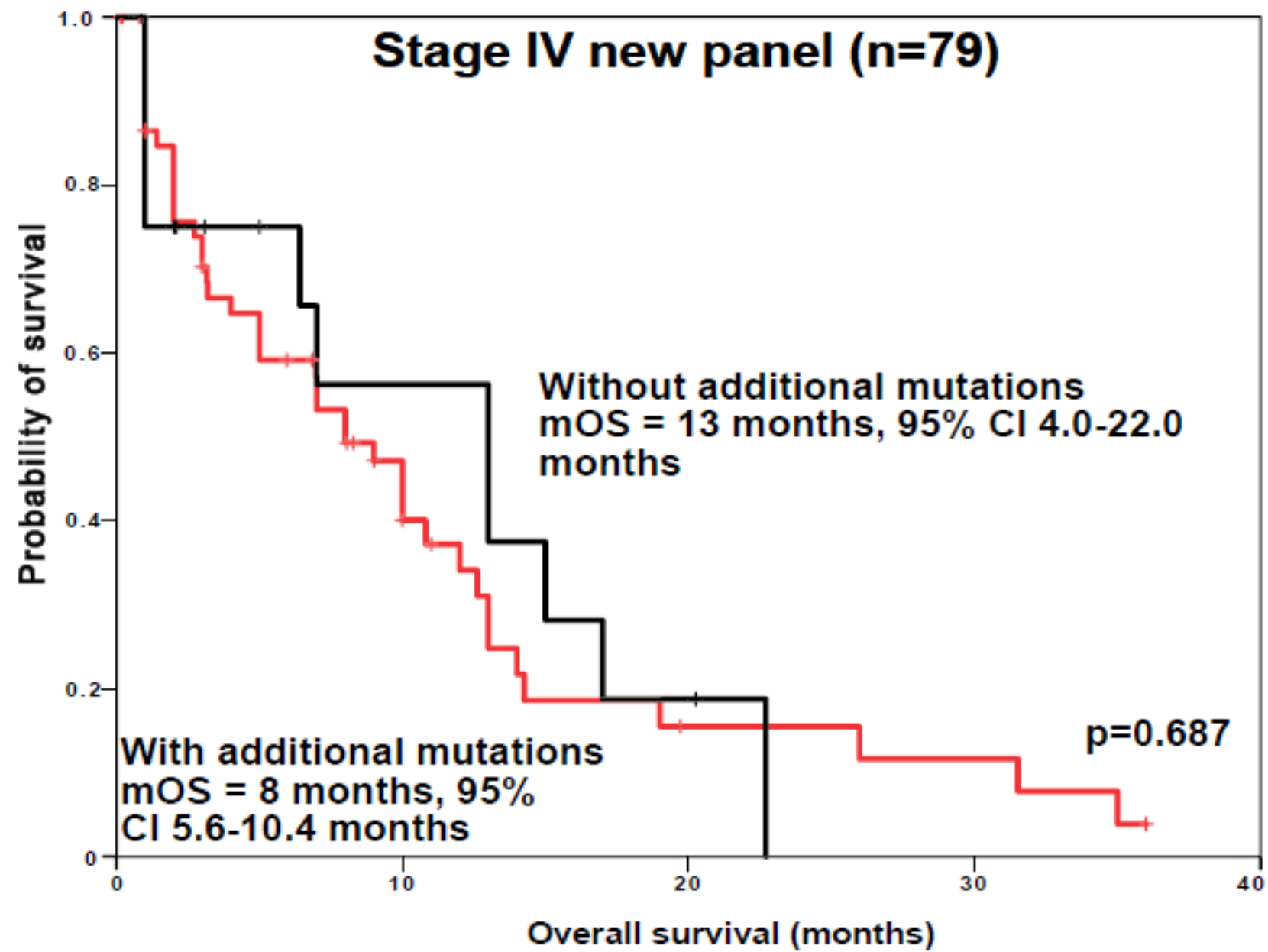


Clustering analysis of 776 cases of *KRAS* mutated NSCLC

Figure from Padda SK et al. JCO Precision Oncology 2021;5, 153-162

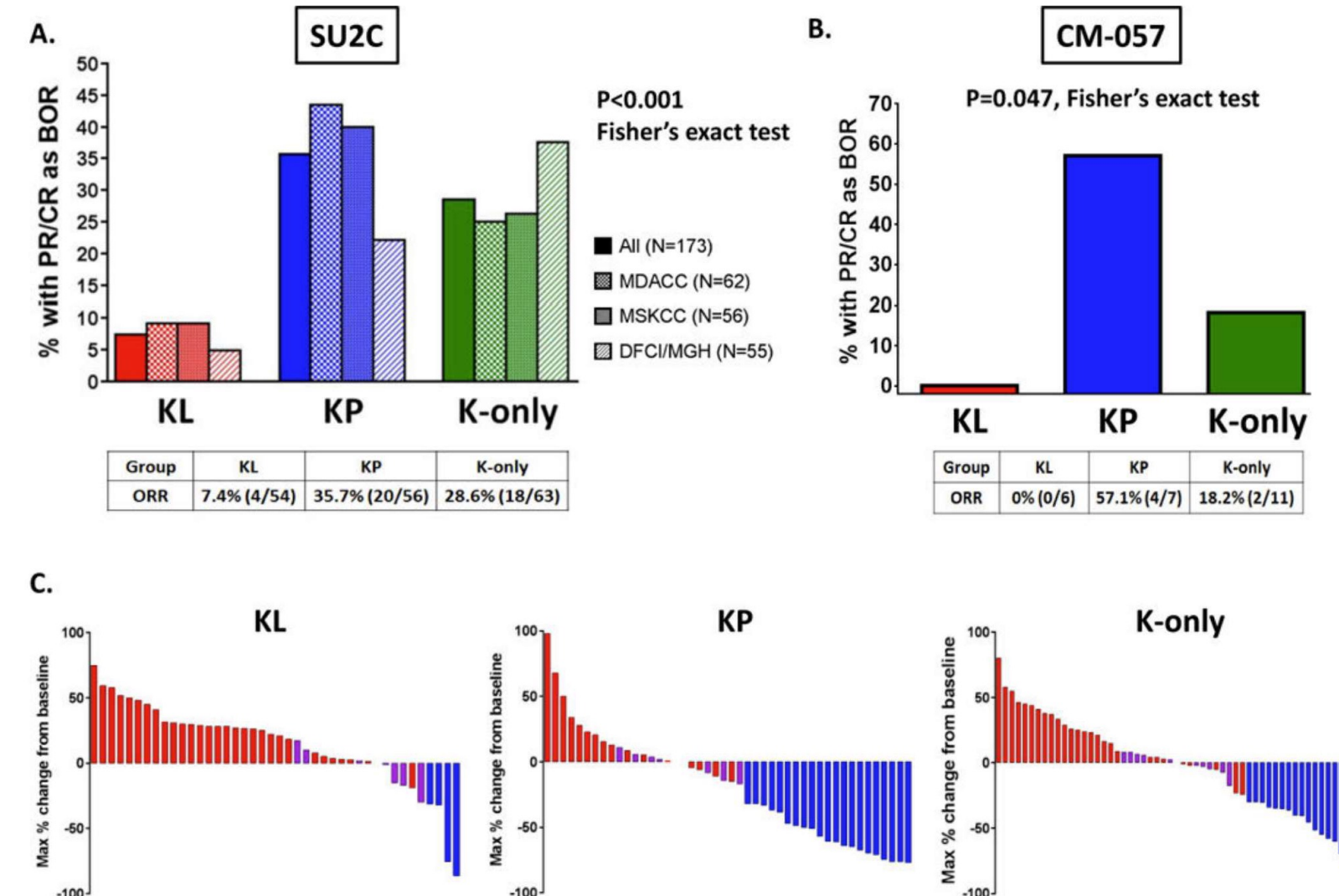
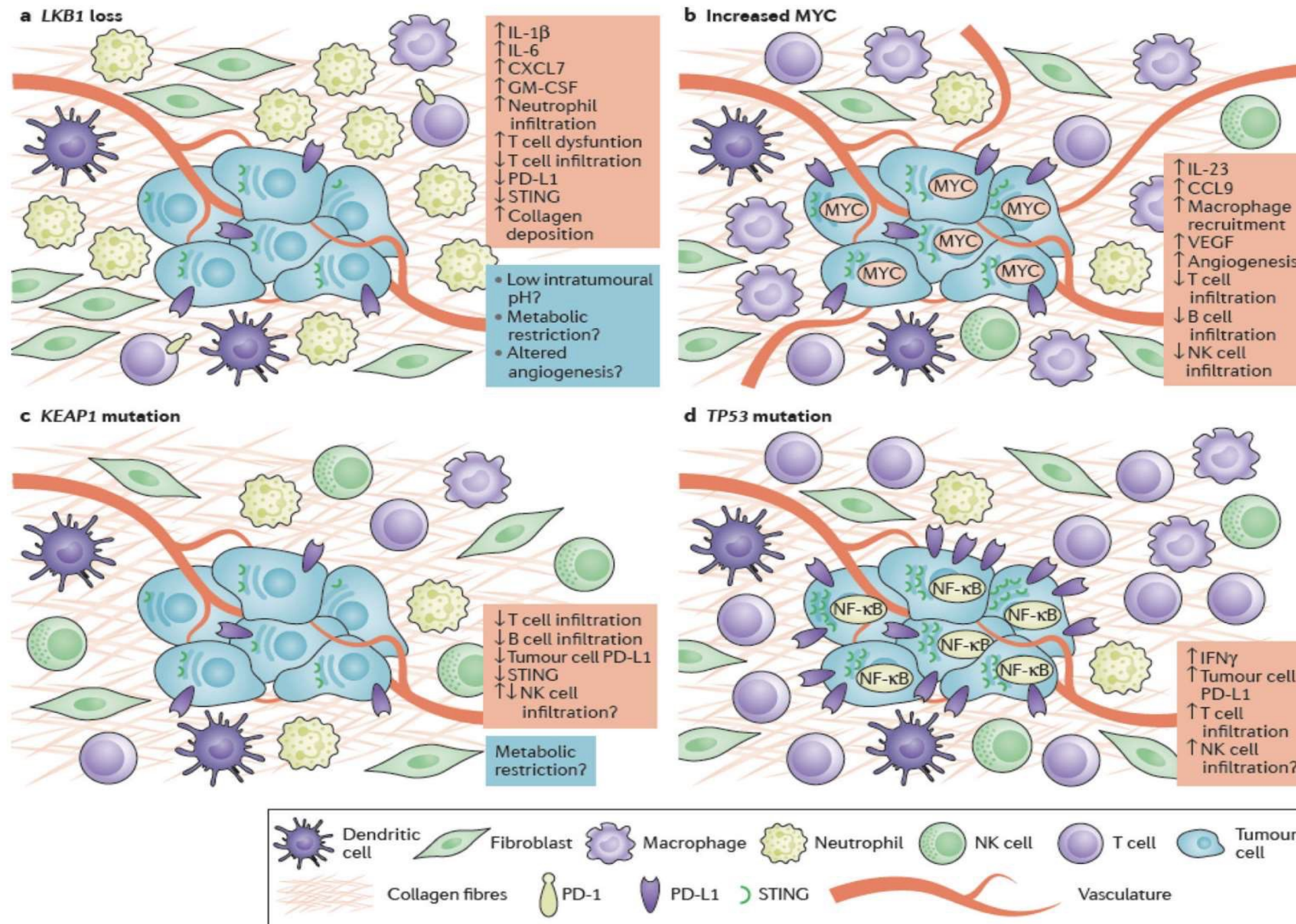


# Overall survival in *KRAS* mutant NSCLC by co-mutation





# Clinical Impact of KRAS co-alterations



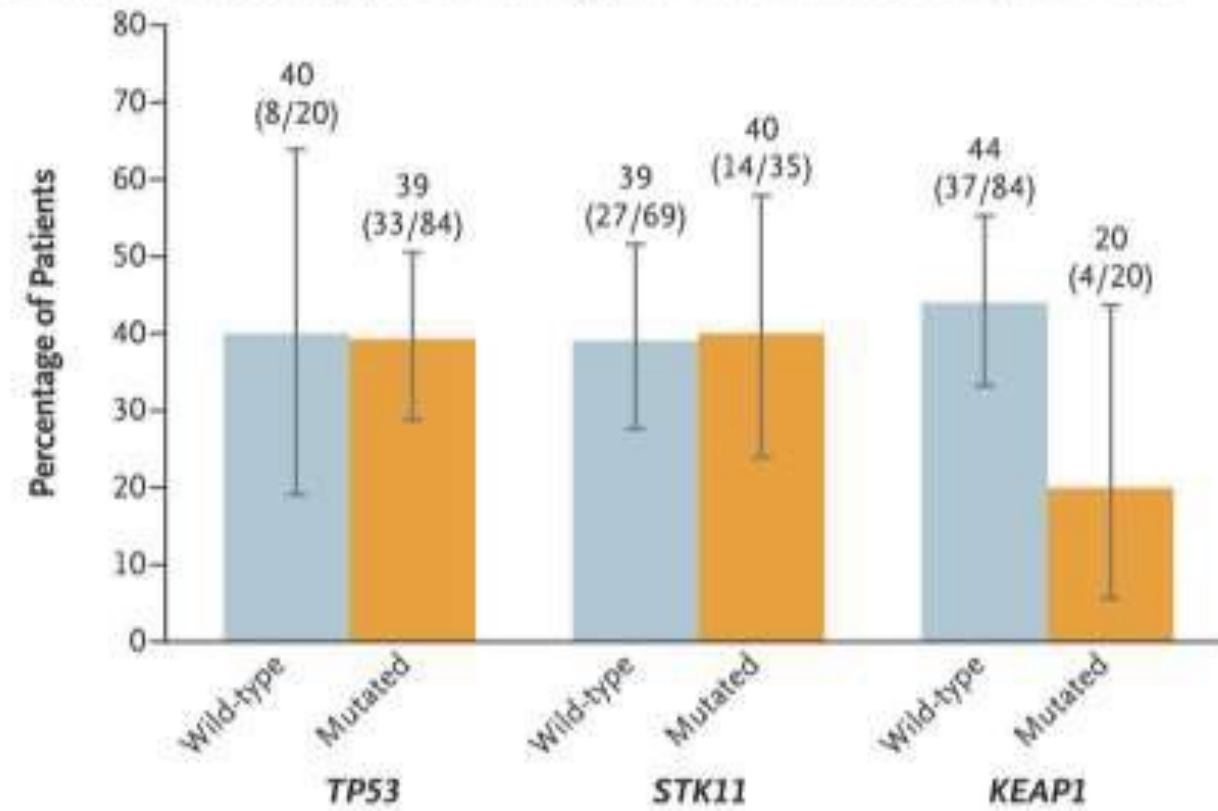
**STK11 loss is a major driver of primary resistance**



# Objective Response Rates by KRAS G12C Co-alterations TP53, STK11, KEAP1

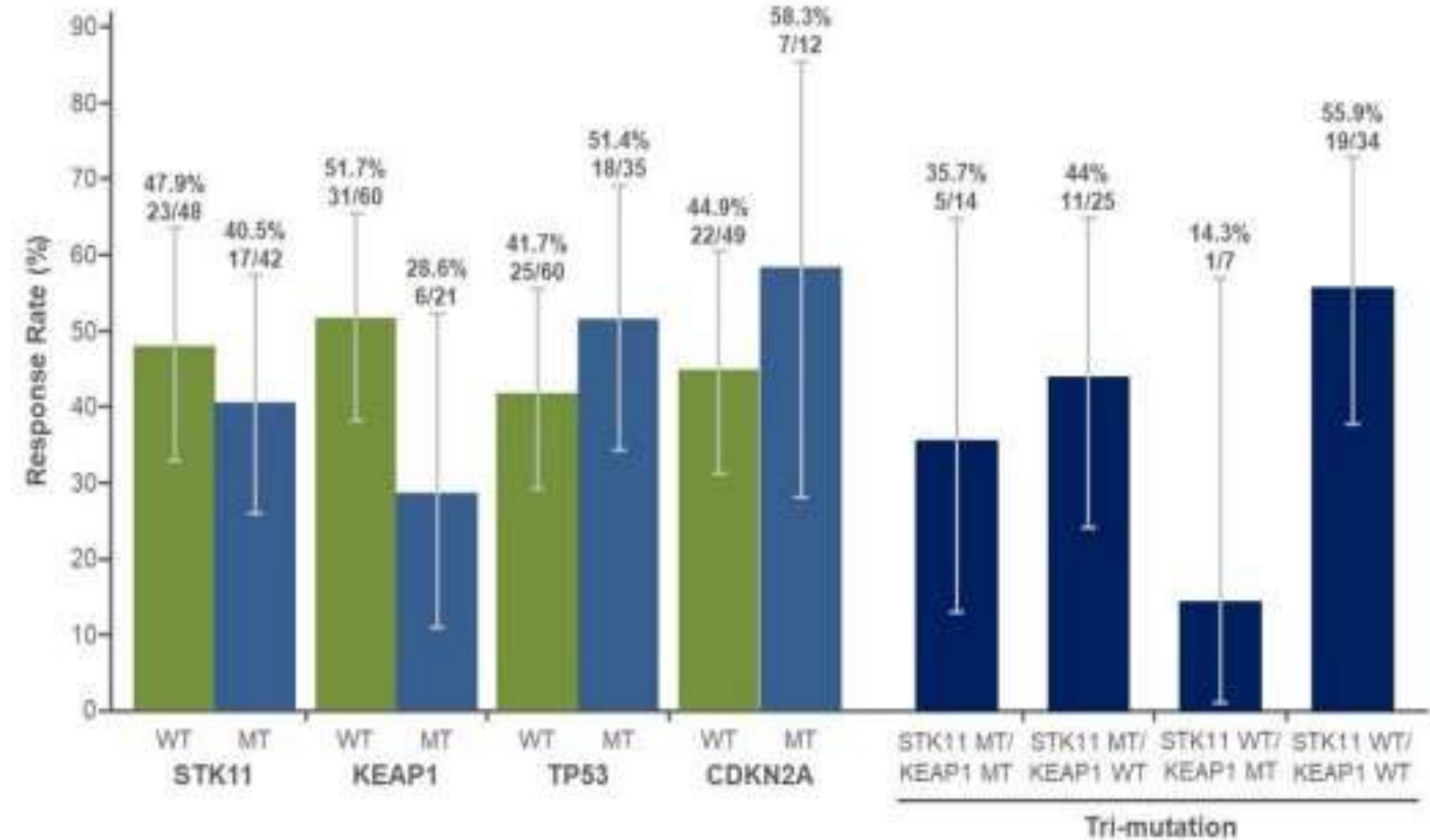
## Sotorasib

**B** Response According to Co-occurring Mutations in TP53, STK11, and KEAP1

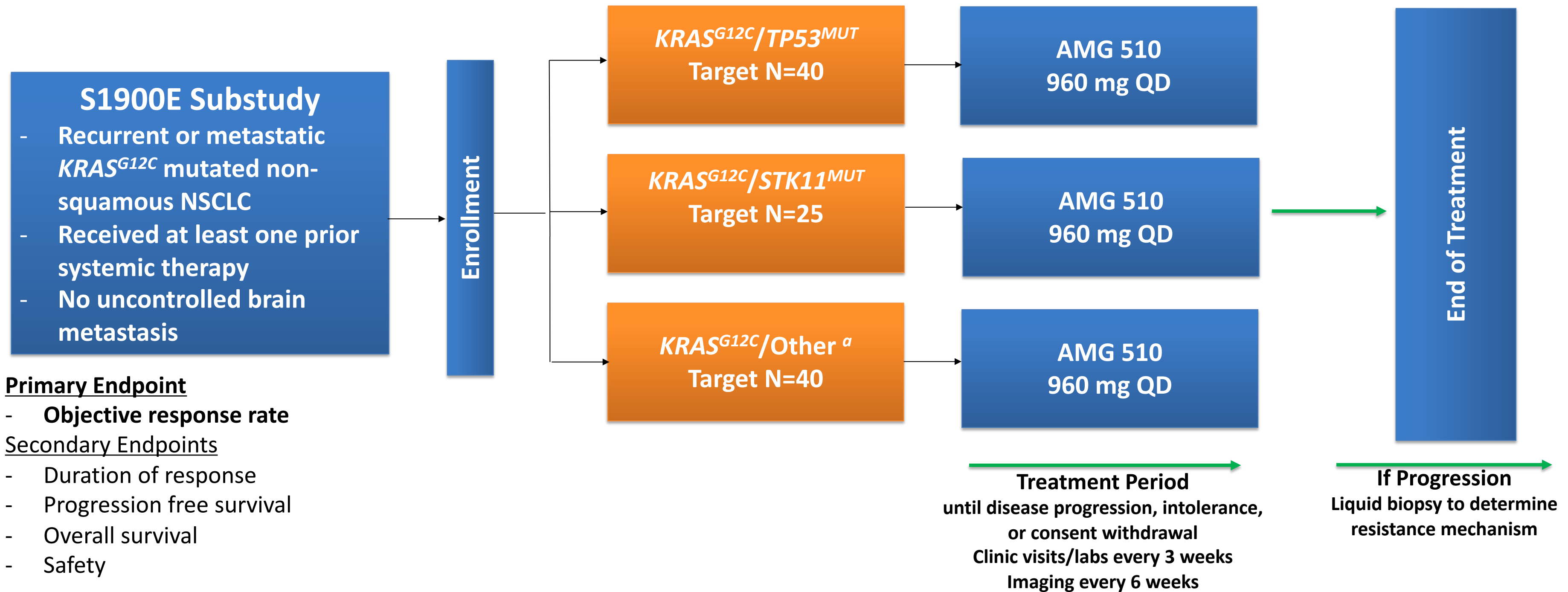


## Adagrasib

ORR in Patients Harboring KRAS<sup>G12C</sup> Co-mutations



# Lung-MAP S1900E—Evaluation of sotorasib in *KRAS* G12C mutant NSCLC with specific co-mutations



## Primary Endpoint

- **Objective response rate**

## Secondary Endpoints

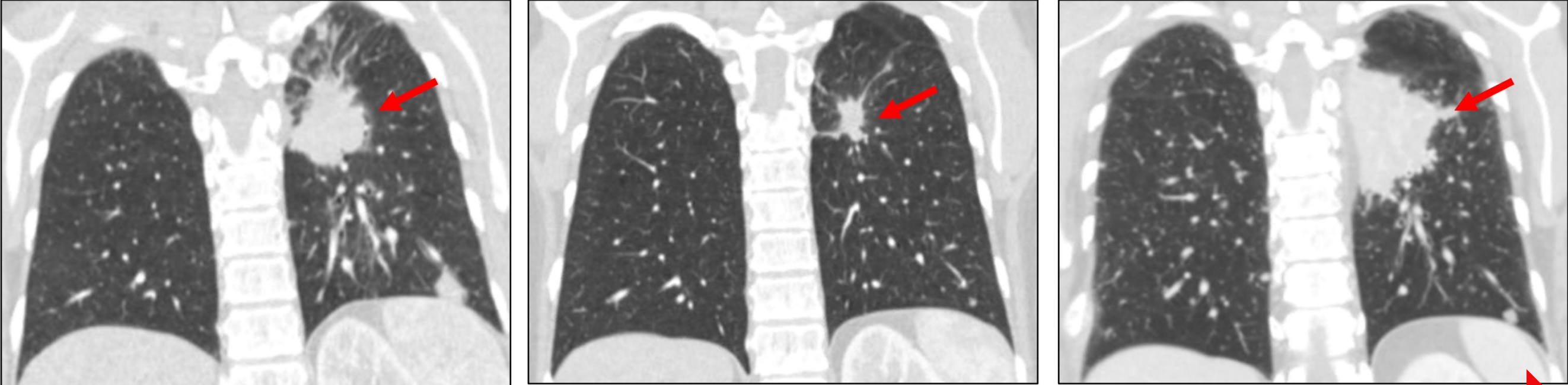
- Duration of response
- Progression free survival
- Overall survival
- Safety

<sup>a</sup>other co-mutations (e.g., *KEAP1*, *NFE2L2*, *CUL3*), double or triple co-mutations (e.g., *STK11/TP53*, *STK11/TP53/KEAP1*), or no co-mutations

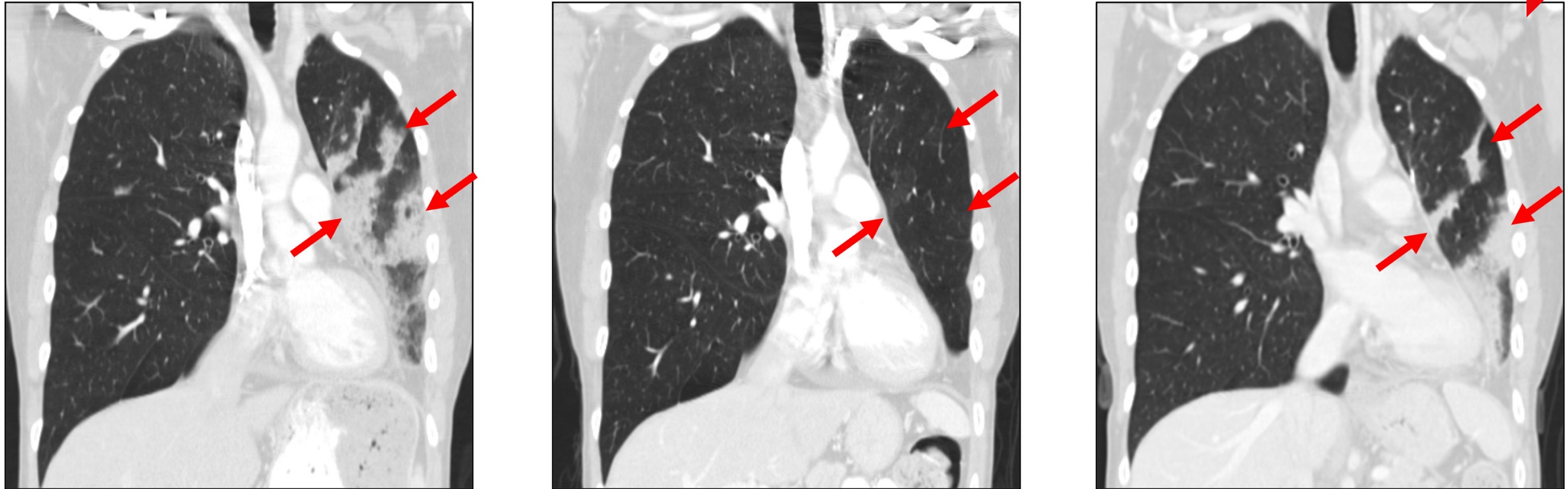


# Acquired Resistance

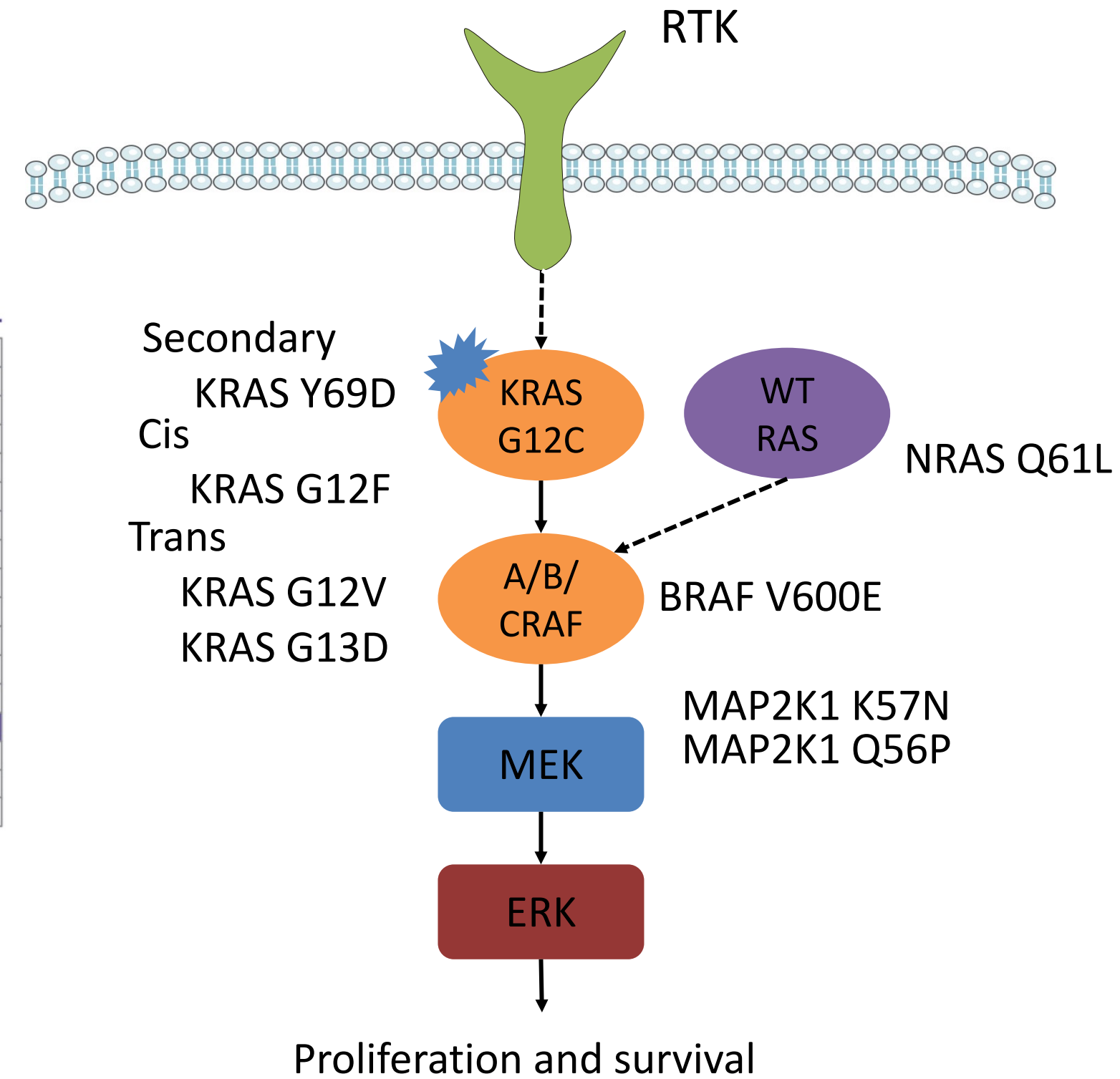
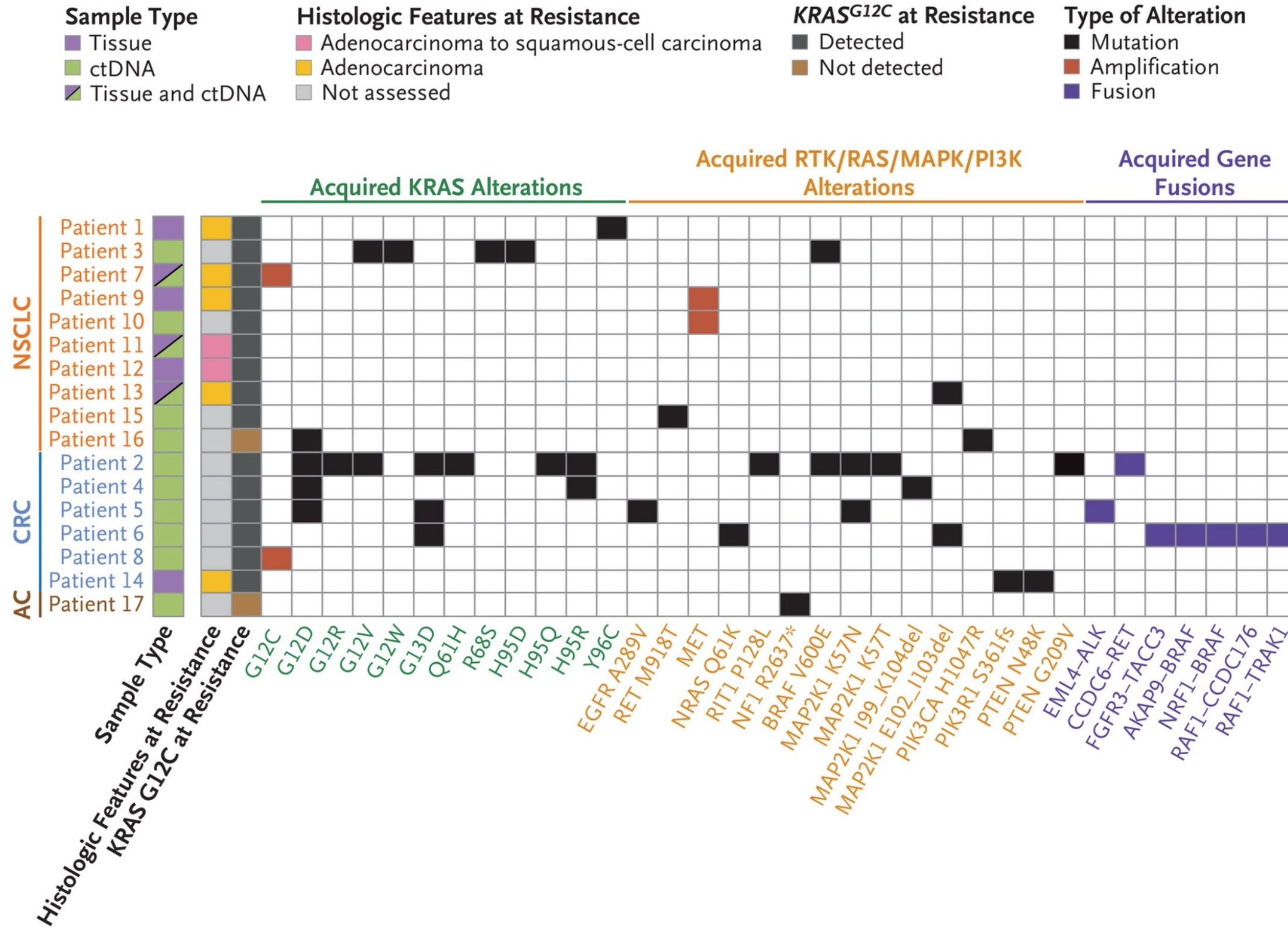
*KRAS G12C* mutant



Baseline Response Resistance



# Acquired Resistance to *KRAS* G12C Inhibitors

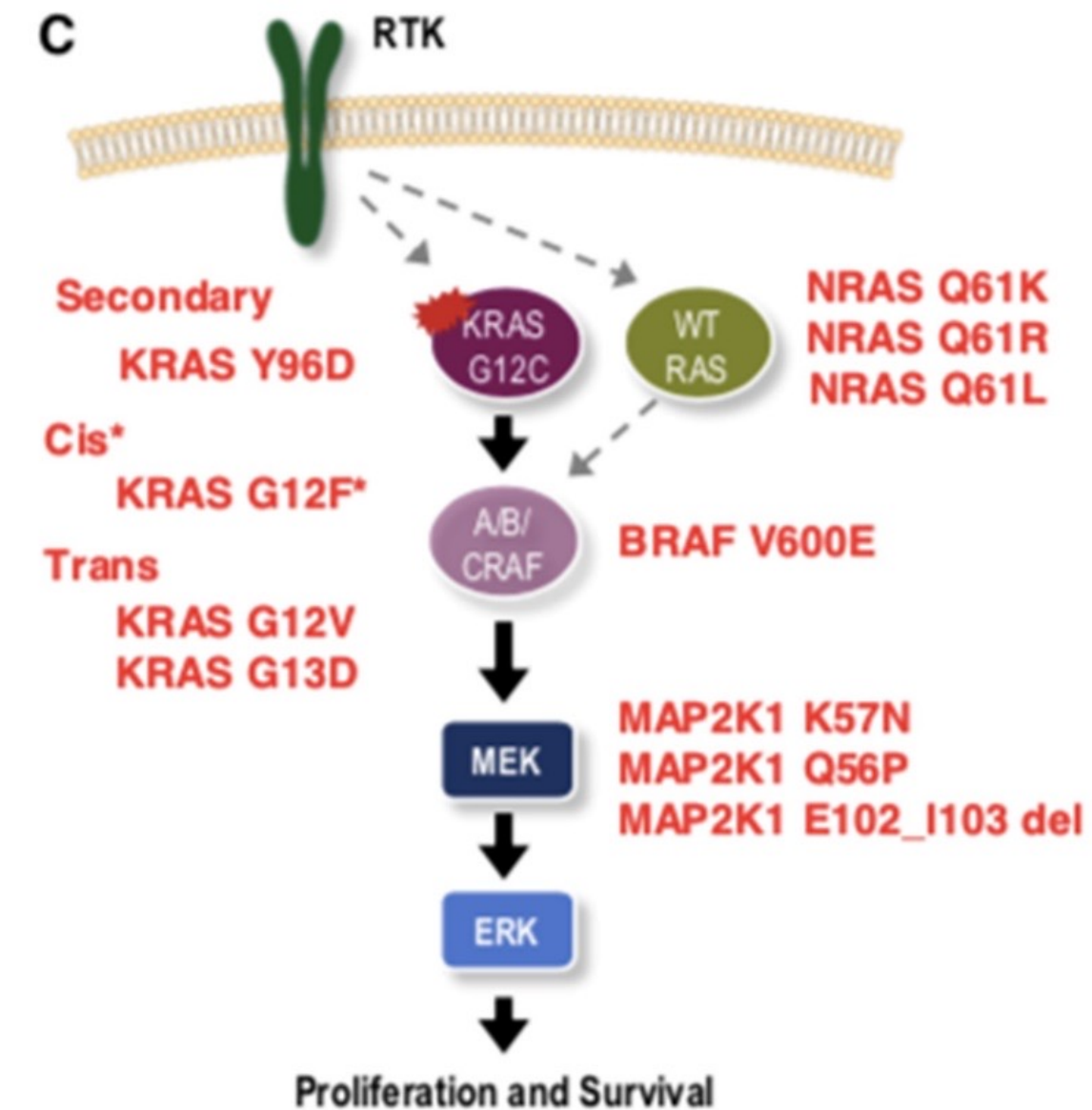




# Resistance to G12C Inhibitors

**B**

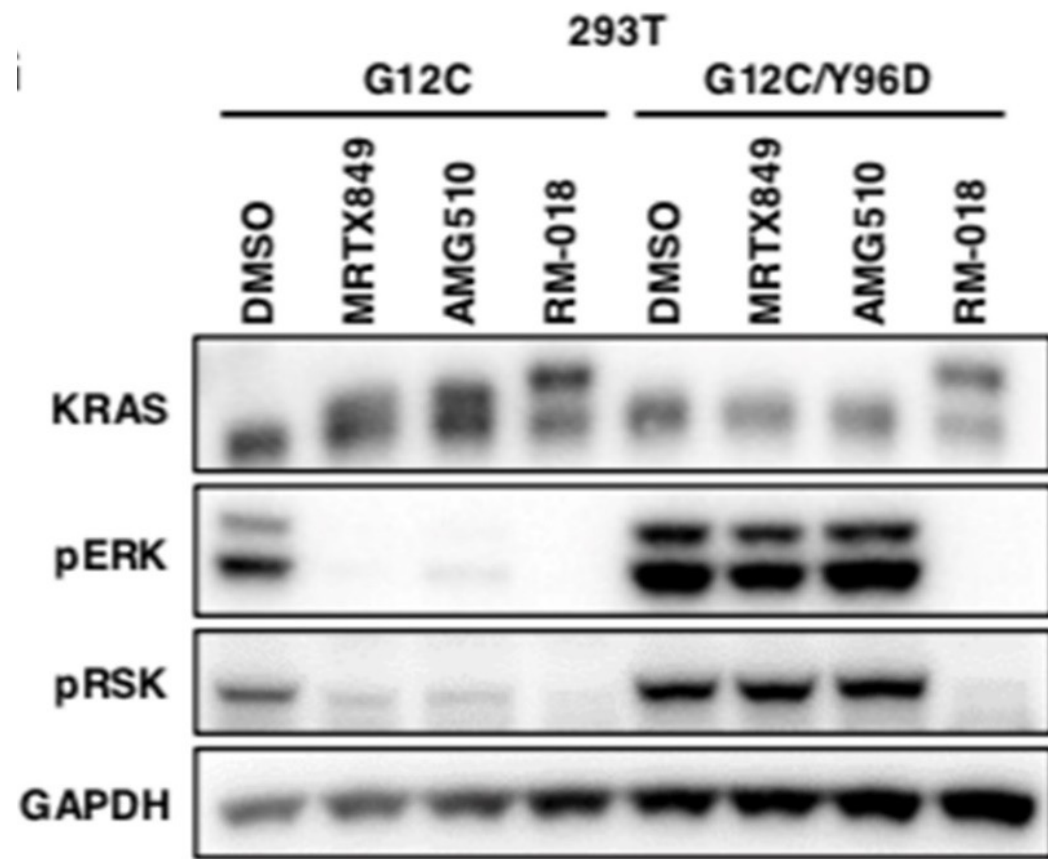
	Tumor		cfDNA		
	Pre-MRTX849	Pre-MRTX849	Days post-MRTX849 discontinuation:		
			0	9	51
TP53 F338fs	36.8%	0.22%	8.8%	10.1%	14.3%
KRAS G12C	21.3%	0.12%	31.7%	47.1%	24.9%
KRAS G12V	-	-	-	-	0.09%
KRAS G13D	-	-	-	0.13%†	0.04%
KRAS Y96D	-	-	0.4%	0.2%	-
NRAS Q61L	-	-	-	0.2%	-
NRAS Q61R	-	-	-	-	0.02%
NRAS Q61K	-	-	0.6%	0.6%	0.9%
BRAF V600E	-	-	0.1%	0.1%	0.5%
MAP2K1 K57N	-	-	0.05%†	-	0.3%
MAP2K1 Q56P	-	-	-	-	0.1%
MAP2K1 E102_I103del	-	-	-	0.12%†	0.2%



## Mechanisms of Resistance

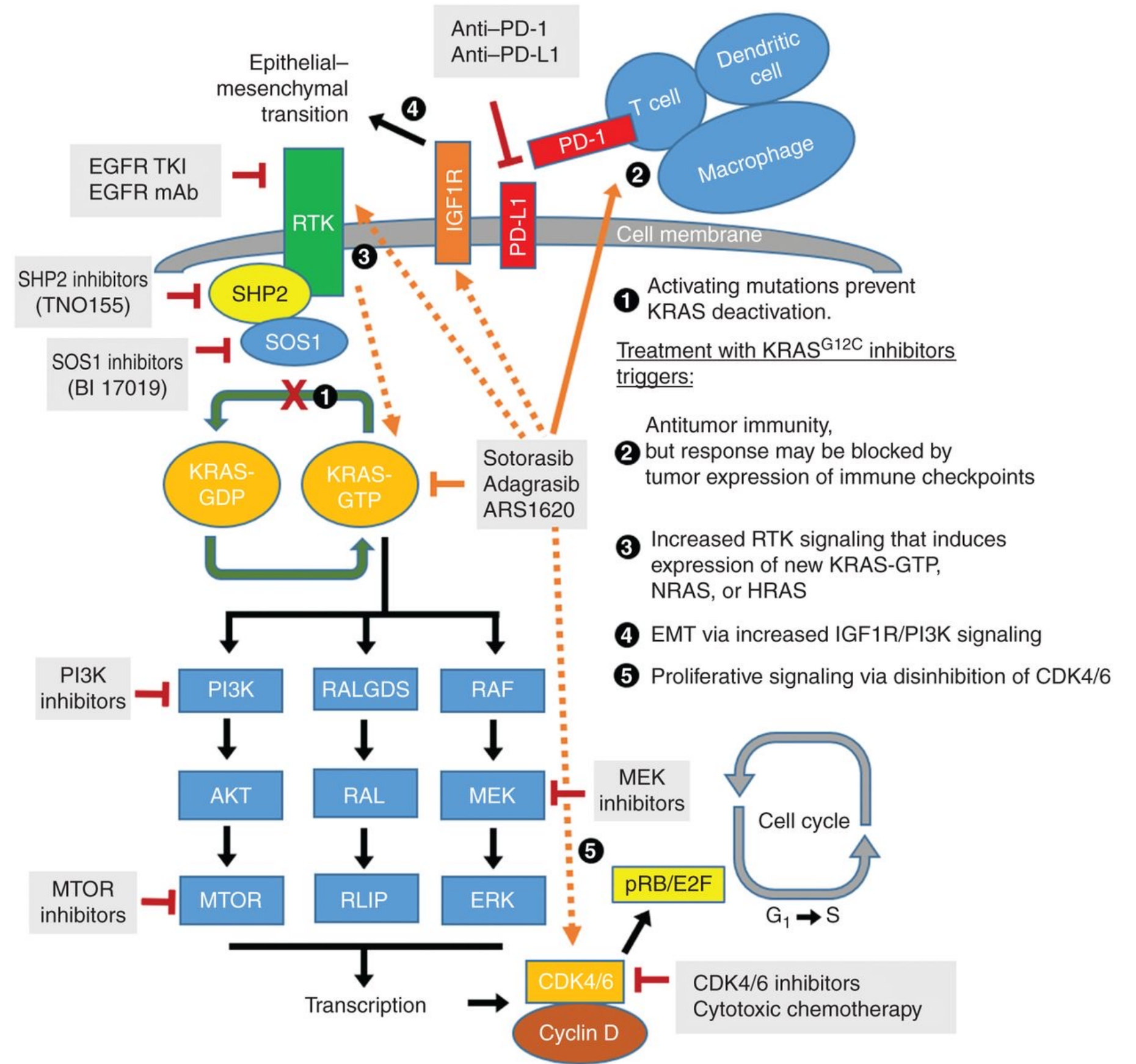
- 1) Activation of other RAS isoforms (NRAS)
- 2) Other KRAS activating mutations in trans (G13D, G12V)
- 3) Loss of G12C thru mutational switch to different KRAS mutation in cis
- 4) Novel KRAS Y96D alters drug binding

# Complexity of resistance to G12C inhibitors



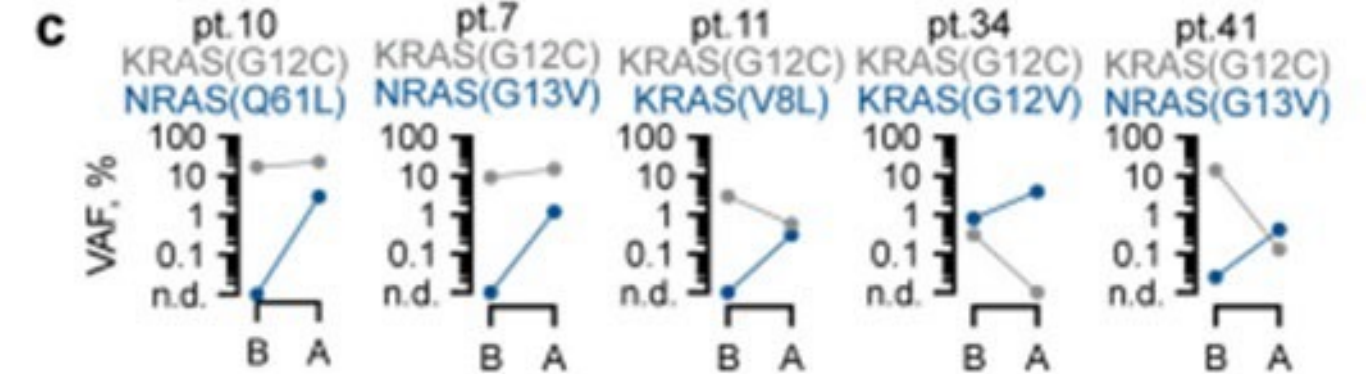
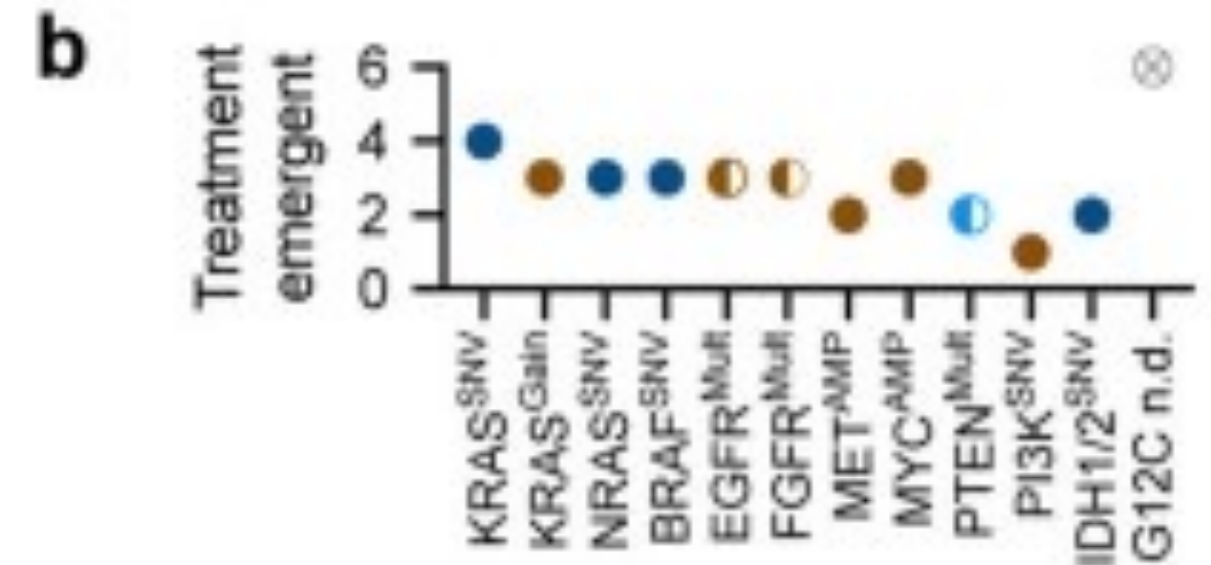
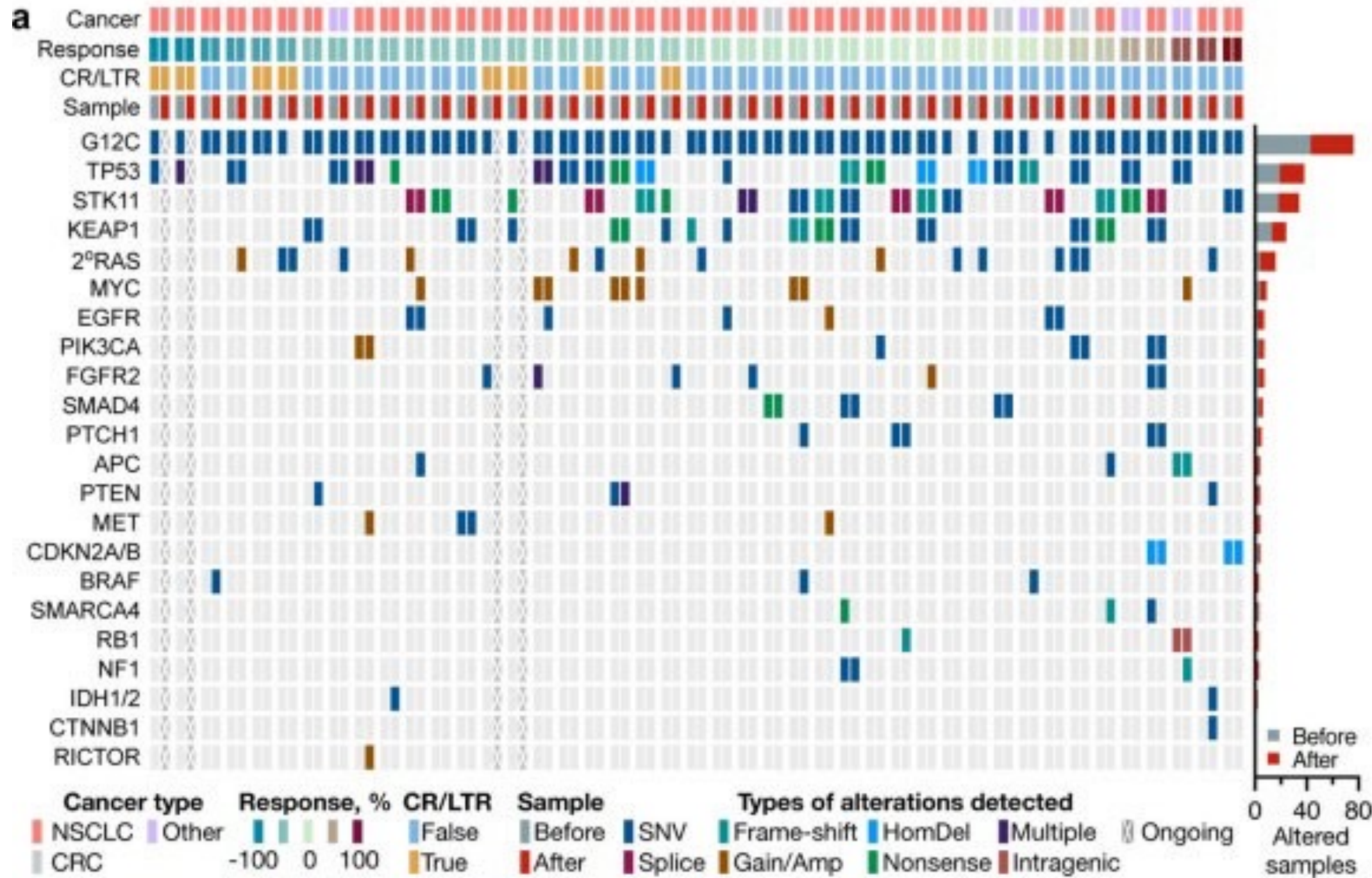
RM-018 blocks downstream signaling in KRAS G12C/Y96D

RM-018 = KRAS G12C (ON) inhibitor



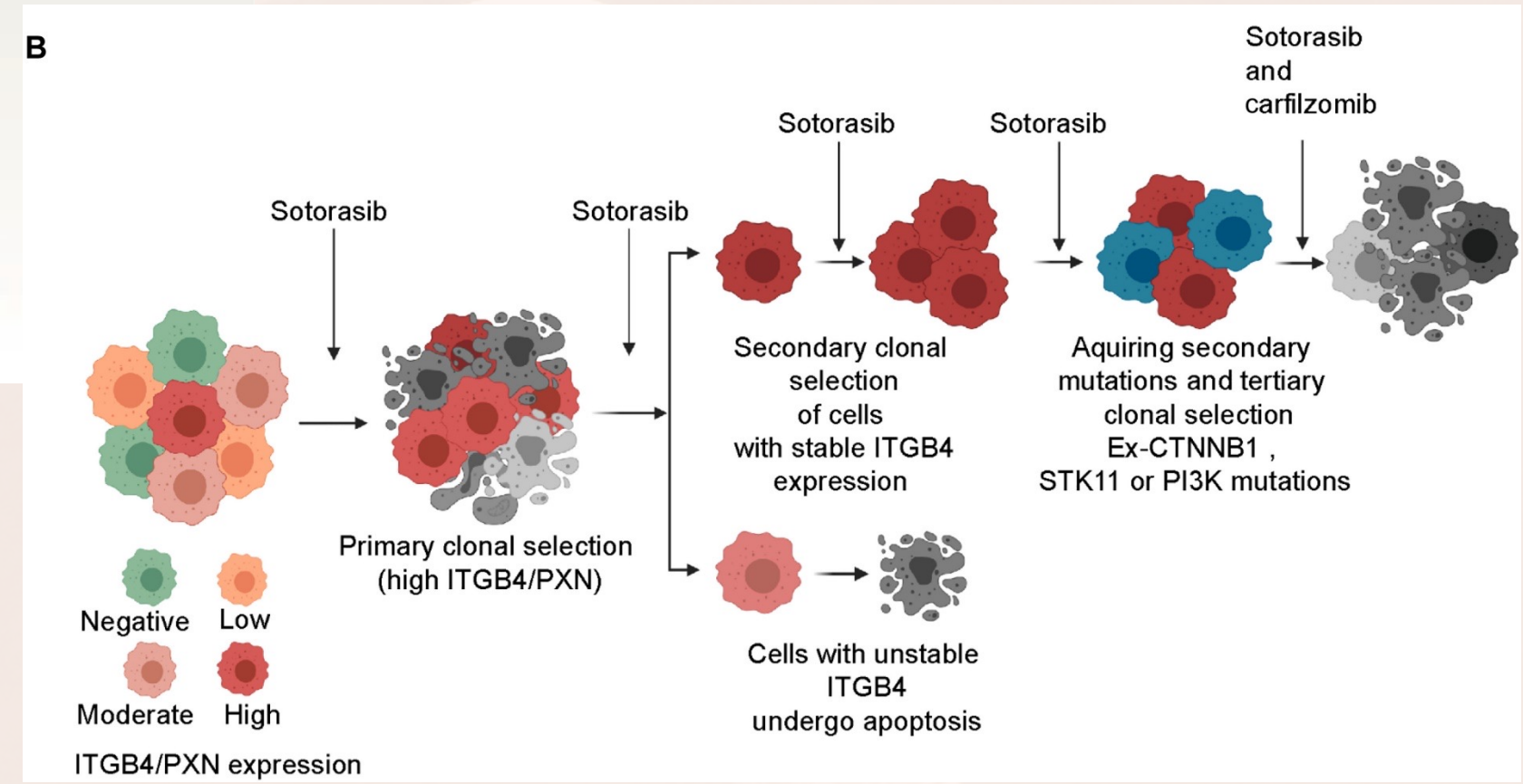
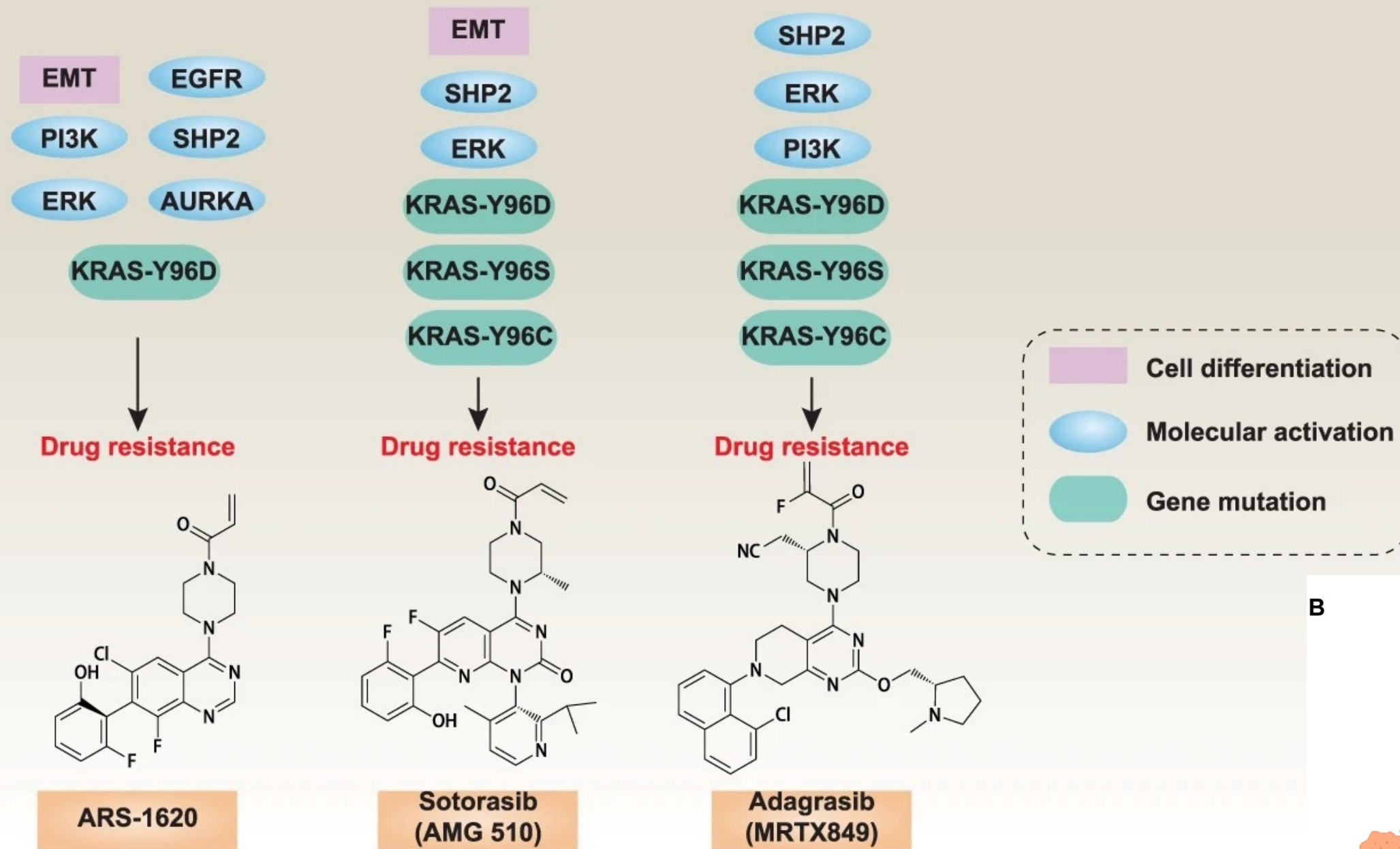


# Resistance to G12C inhibitors





# KRAS G12C resistance mechanisms

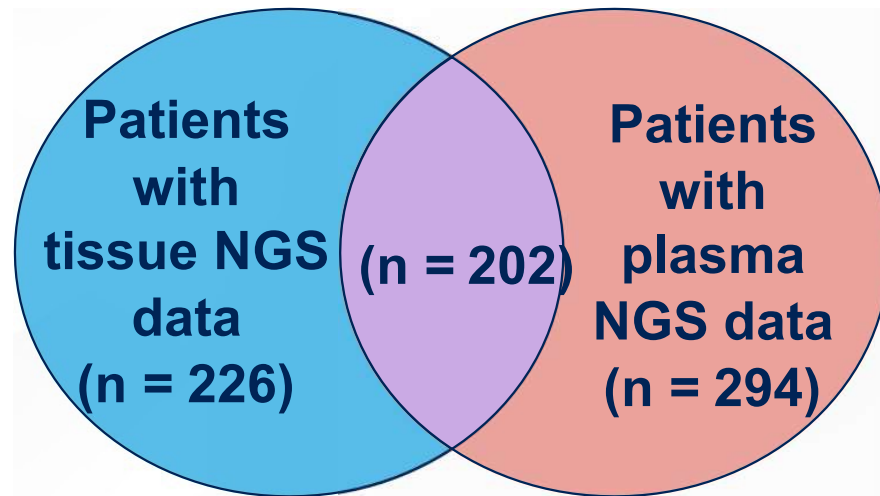


Liu J, Kang R, Tang D. Cancer Gene Ther 2022



# Biomarker Subgroup Analysis of Codebreak 200

ITT population  
(N = 345)



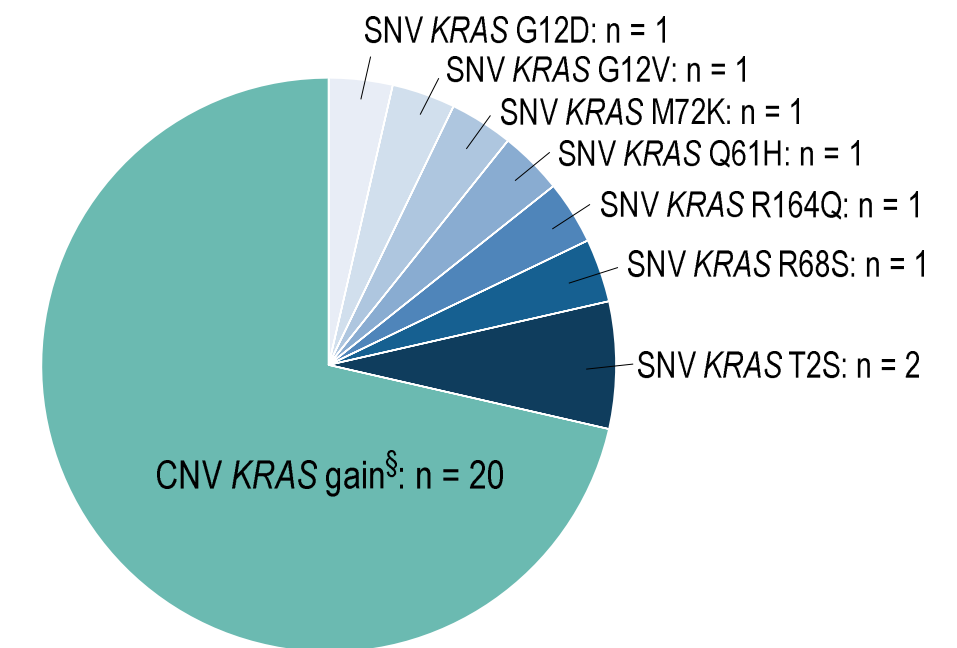
Biomarker evaluable population:  
patients with tumor and/or plasma NGS  
data  
(n = 318)

Baseline co-alterations were  
balanced in the Sotorasib  
and Docetaxel arms

## Sotorasib Retained PFS Benefit Versus Docetaxel Across Key Co-alteration Subgroups

- How to use this information in clinical practice?
  - STK11, KEAP1, TP53 ✓
  - Additional RAS (non-KRAS G12C) alterations ✗
  - ?NOTCH1 ✓ / ✗
- Limitations
  - Small numbers
  - Exploratory endpoints
- What about acquired resistance?

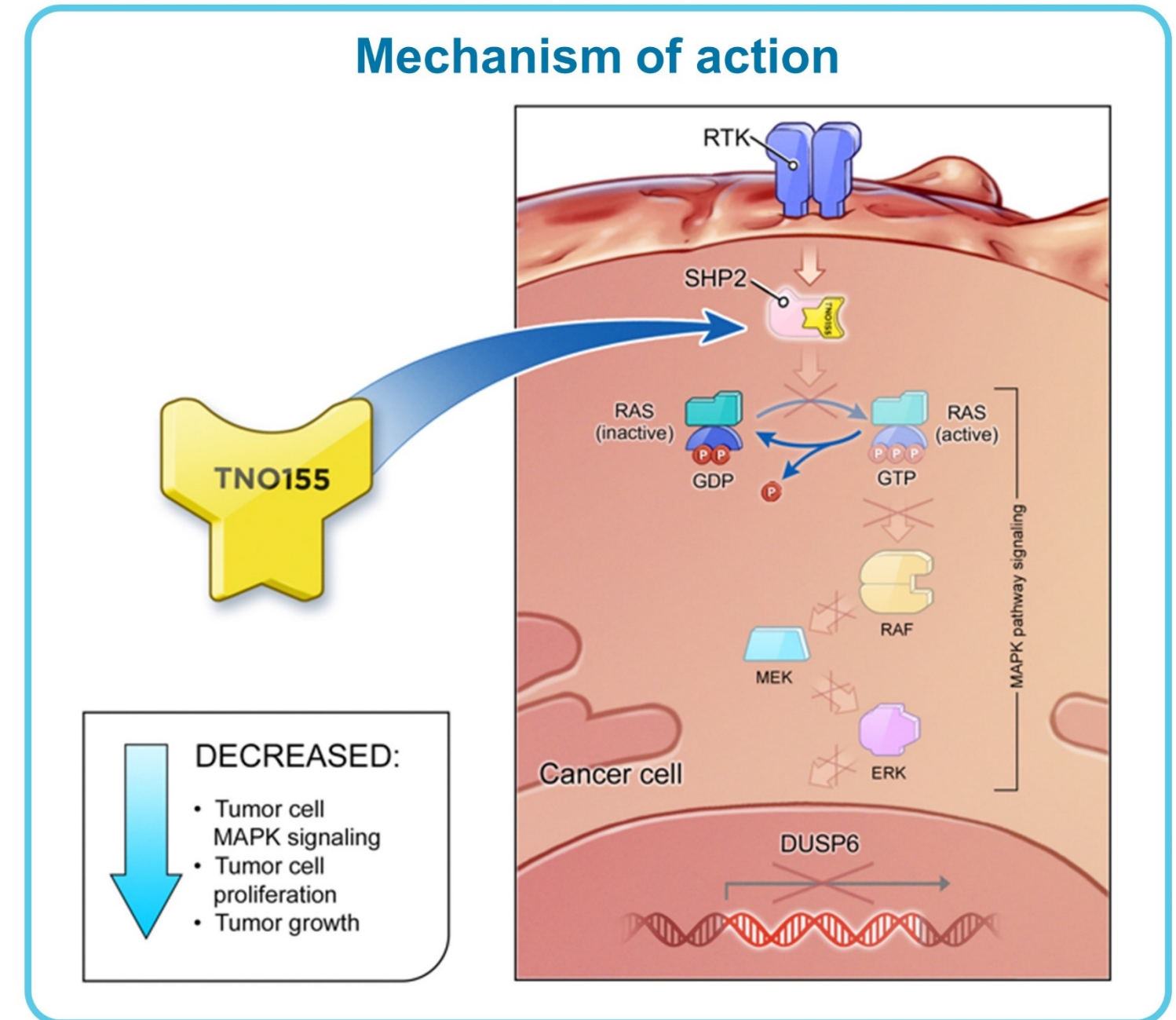
	Sotorasib (n = 164)	Docetaxel (n = 154)	Treatment Difference (P-value)
KRAS co-alteration*, n (%)	9 (5)	17 (11)	
ORR†, n (%)	0	0	–
Median PFS (m)	1.8	2.5	0.016‡
HR (95% CI)‡	1.74 (0.84, 3.58)		





# SHP2 in overcoming KRAS G12C inhibitor resistance

- SHP2 is a cytoplasmic phosphatase that transduces signaling from RTKs, such as EGFR, MET, and HER2<sup>1</sup>
- Upon activation of RTK, SHP2 activates signaling pathways, including the RAS/MAPK pathway<sup>1-3</sup>
- RTK and RAS signaling is frequently deregulated in many cancer types<sup>3</sup>
- Patients with RTK-driven cancers can develop resistance to RTK- and MAPK-targeted agents;<sup>4,5</sup> targeting SHP2 may potentially offer an alternative treatment option for these patients
- TNO155 is an orally bioavailable, selective, first-in-class allosteric inhibitor of wild-type SHP2<sup>6</sup>
- CTNO155X2101 (NCT03114319) is an ongoing first-in-human, open-label dose escalation/expansion trial of TNO155 in adults with select advanced solid tumors; here we describe data from the TNO155 single-agent dose-escalation part



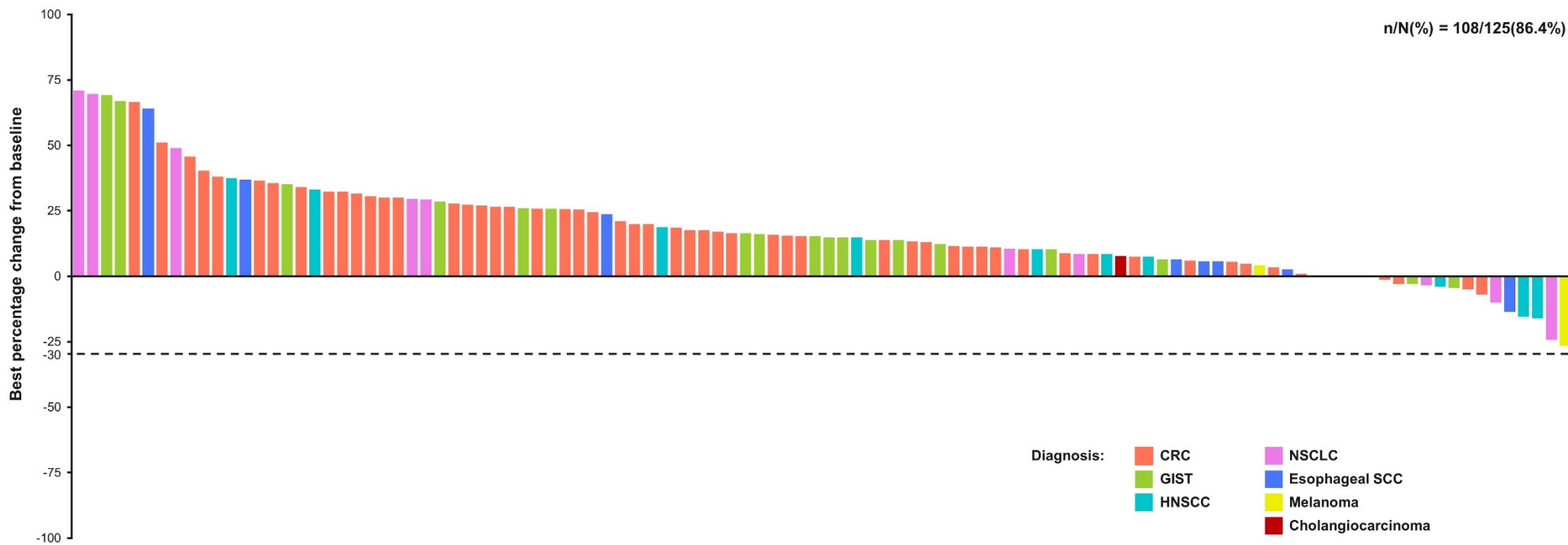
EGFR: epidermal growth factor receptor; HER2: human epidermal growth factor receptor 2; MAPK: mitogen-activated protein kinase; RTK: receptor tyrosine kinase; SHP2: Src homology region 2 domain-containing phosphatase-2.

1. Grossmann KS *et al. Adv Cancer Res.* 2010;106:53–89; 2. Matozaki T *et al. Cancer Sci.* 2009;100:1786–1793; 3. Sanchez-Vega F *et al. Cell.* 2018;173:321–337; 4. van der Wekken AJ *et al. Crit Rev Oncol Hematol.* 2016;100:107–116; 5. Wang WL *et al. Cancer Chemother Pharmacol.* 2011;67(suppl 1):S15–S24; 6. LaMarche MJ *et al. J Med Chem.* 2020;63:13578–13594.



# TNO155—Best percentage change in target lesions and duration of response in dose escalation patients

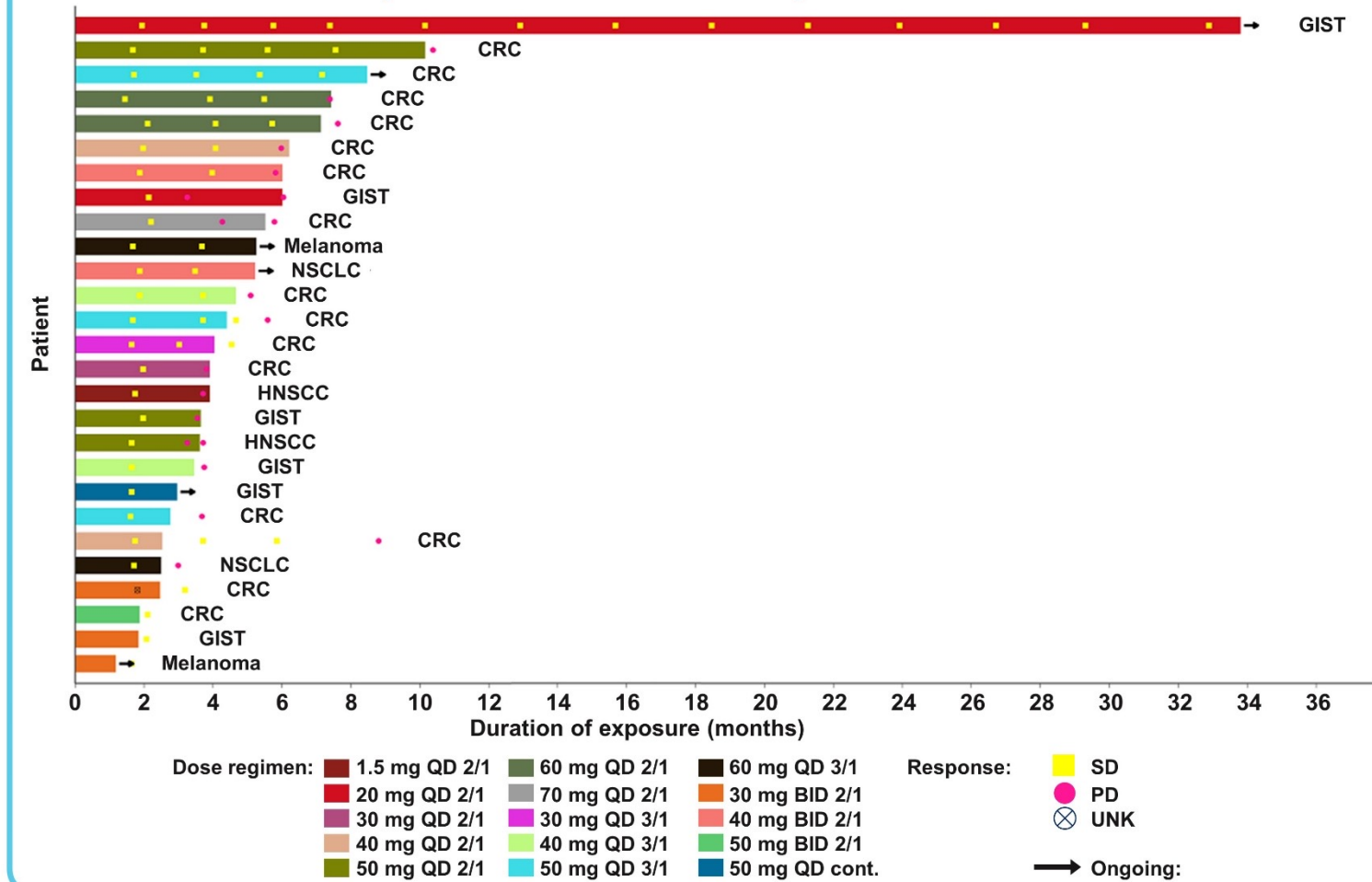
Best percentage change from baseline in sum of target lesion diameters



CRC: colorectal cancer; GIST: gastrointestinal stromal tumor; HNSCC: head and neck squamous cell cancer; NSCLC: non-small cell lung cancer; SCC: squamous cell cancer.

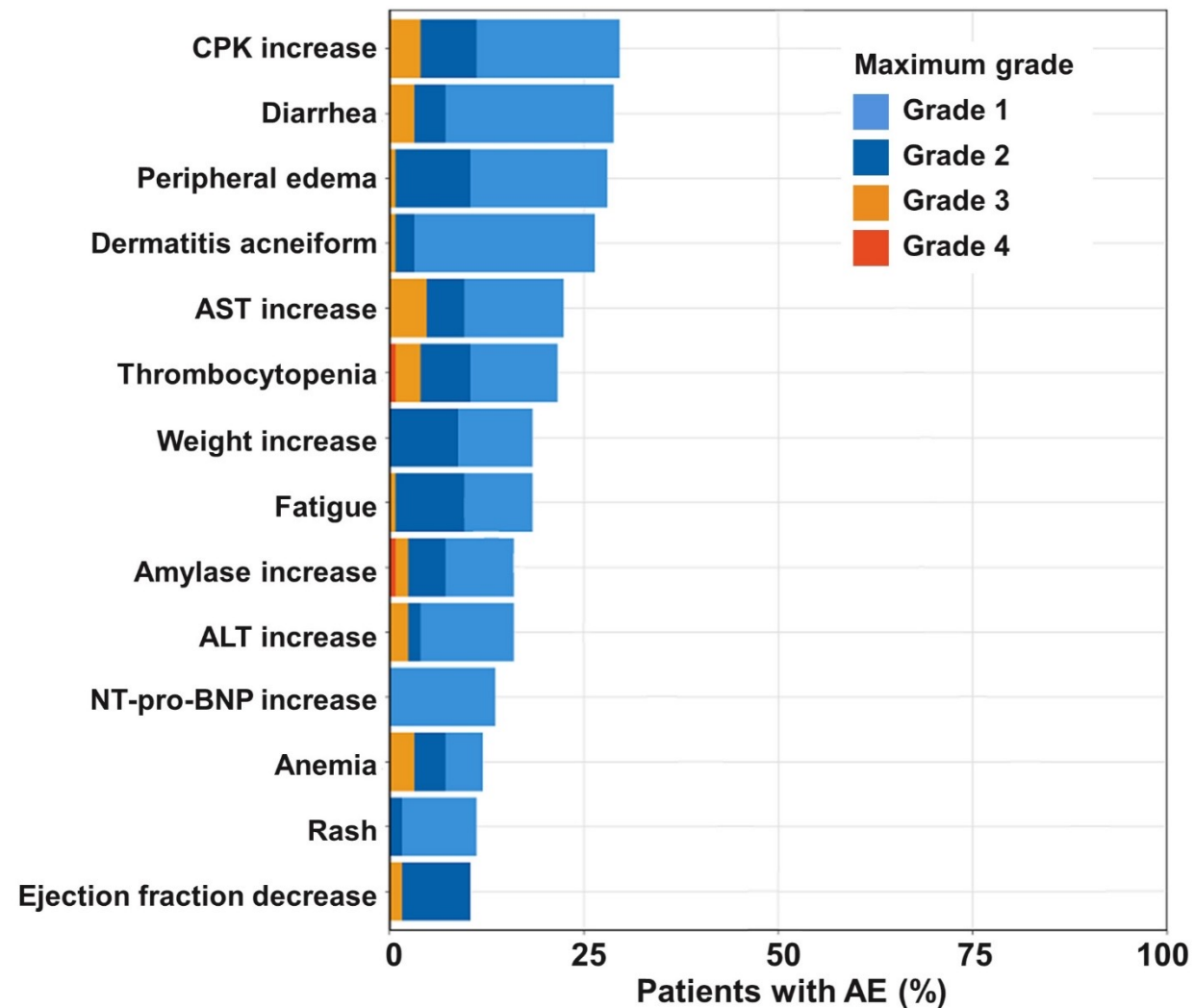
- Best response was SD
- Median duration of SD 5.6 months (95% CI: 1.6, 32.9)
- Only 7 *KRAS* mutant patients

Duration of exposure to TNO155 in patients with stable disease



# Treatment-related AEs to TNO155, SHP2 inhibition

## Treatment-related AEs, ≥10% of patients



- Most AEs were Grade 1 or 2 in severity
- There were no suspected-related Grade 5 AEs
- Ejection fraction decreases/left ventricular dysfunction of any grade were reported in 13/125 (10%) patients
  - Five patients (4%) had LVEF decreases of ≥10% from baseline to a value below 50%, with only one below 40%
    - Four resolved within 7–9 days and one in 23 days
  - The majority of ejection fraction decreases were mild (n=11 Grade 2) and were identified as a result of frequent monitoring\*

AE terms which are equivalent are grouped for reporting: rash and maculopapular rash are reported as rash; platelet count decrease and thrombocytopenia are reported as thrombocytopenia; neutrophil count decrease and neutropenia are reported as neutropenia; ejection fraction decrease and left ventricular dysfunction are reported as ejection fraction decrease.

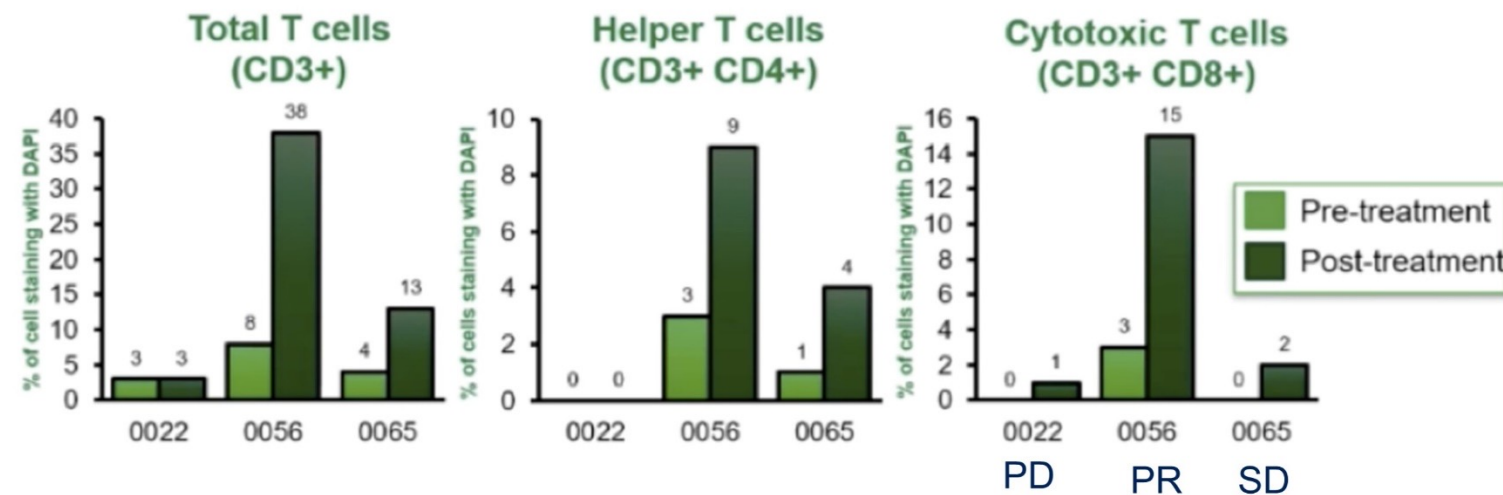
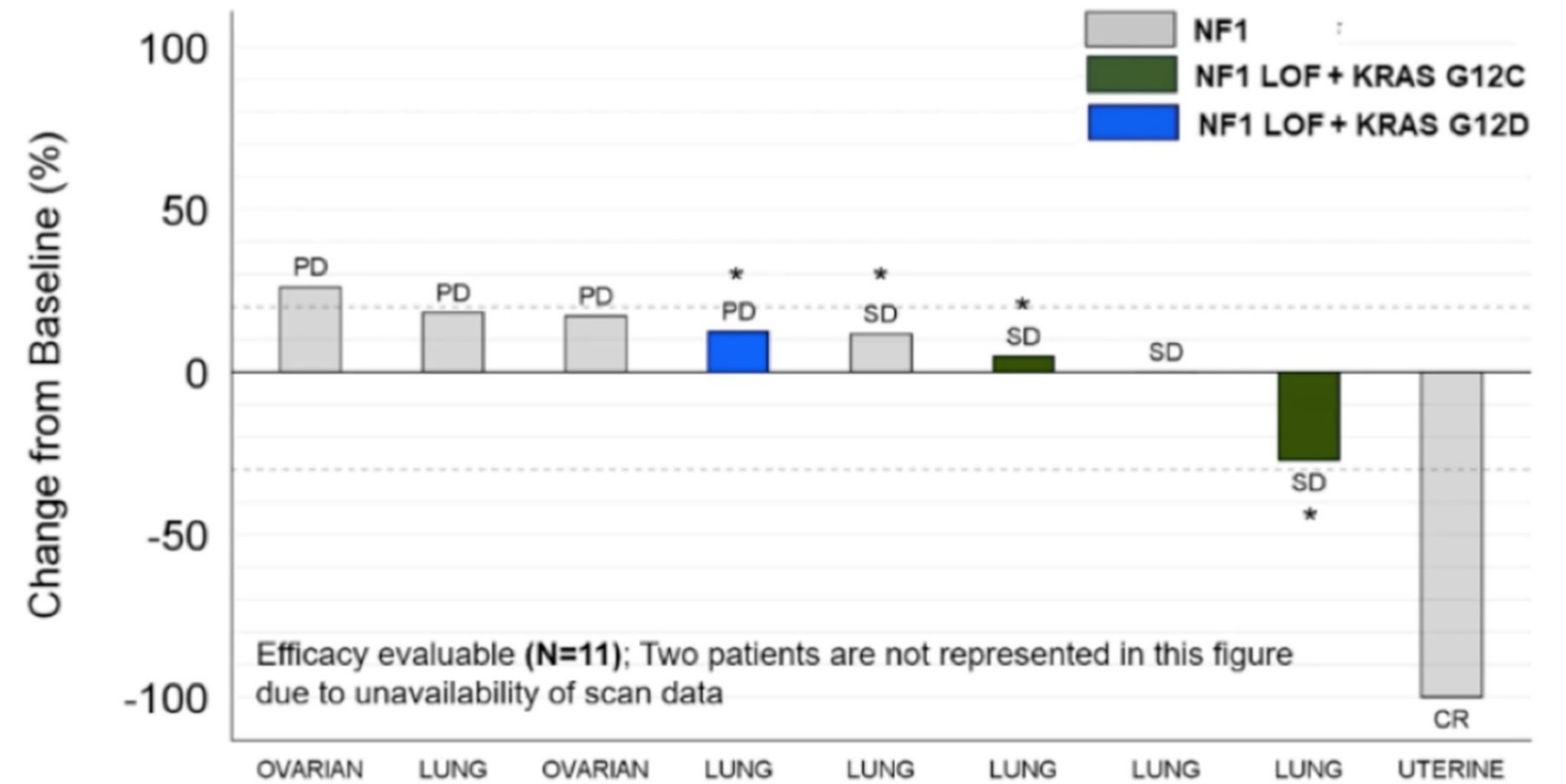
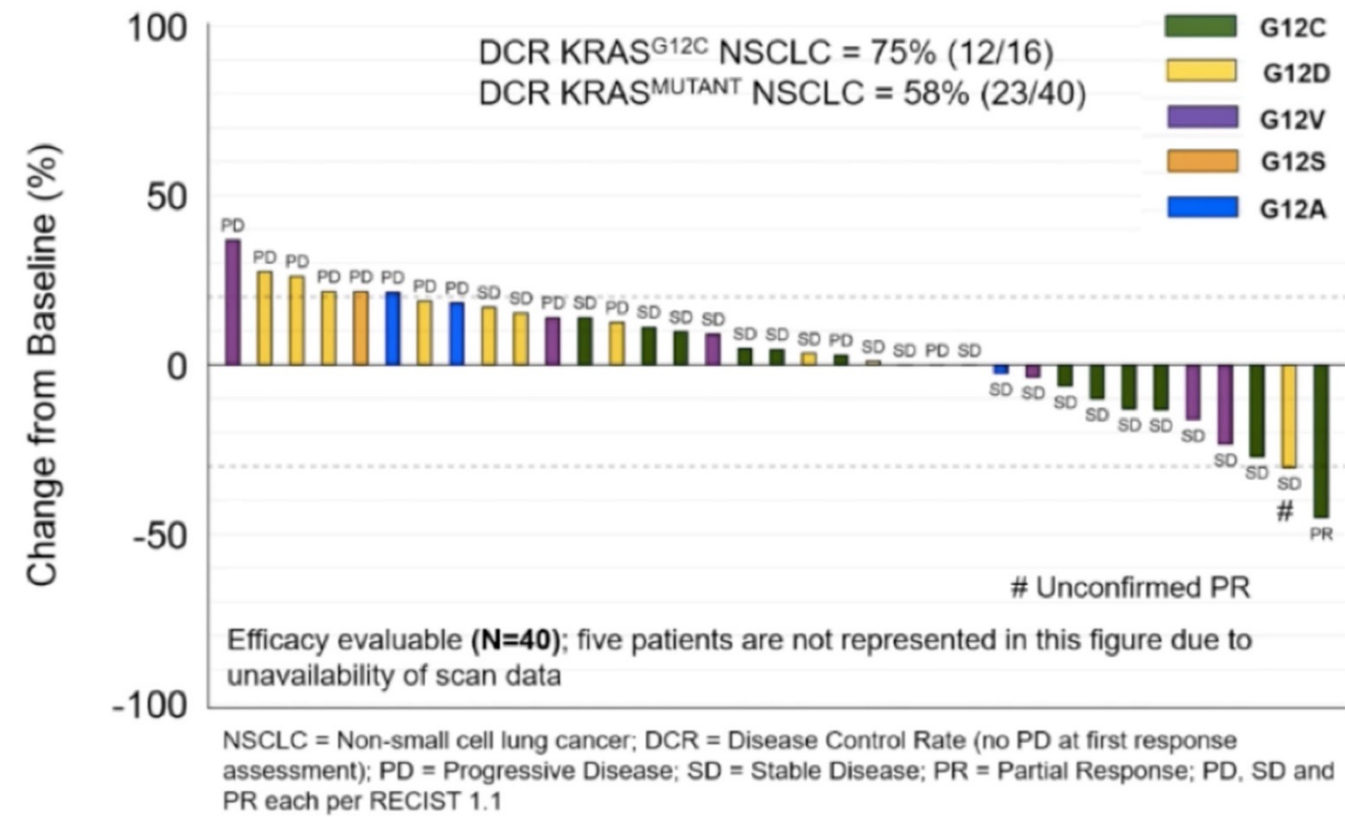
\*Cardiac imaging by echocardiogram or MUGA scan was required at baseline, C1D14 (2/1 schedule) or C1D21 (3/1 or continuous schedules), C2D1, C2D14 or C2D21, C3D1, then D1 of every even cycle through C8, then D1 of every third cycle.

AE: adverse event; ALT: alanine amino transferase; AST: aspartate amino transferase; C: cycle; CPK: blood creatine phosphokinase; D: day; LVEF: left ventricular ejection fraction; MUGA: multiple gated acquisition;

NT-pro-BNP: N-terminal prohormone brain natriuretic peptide.



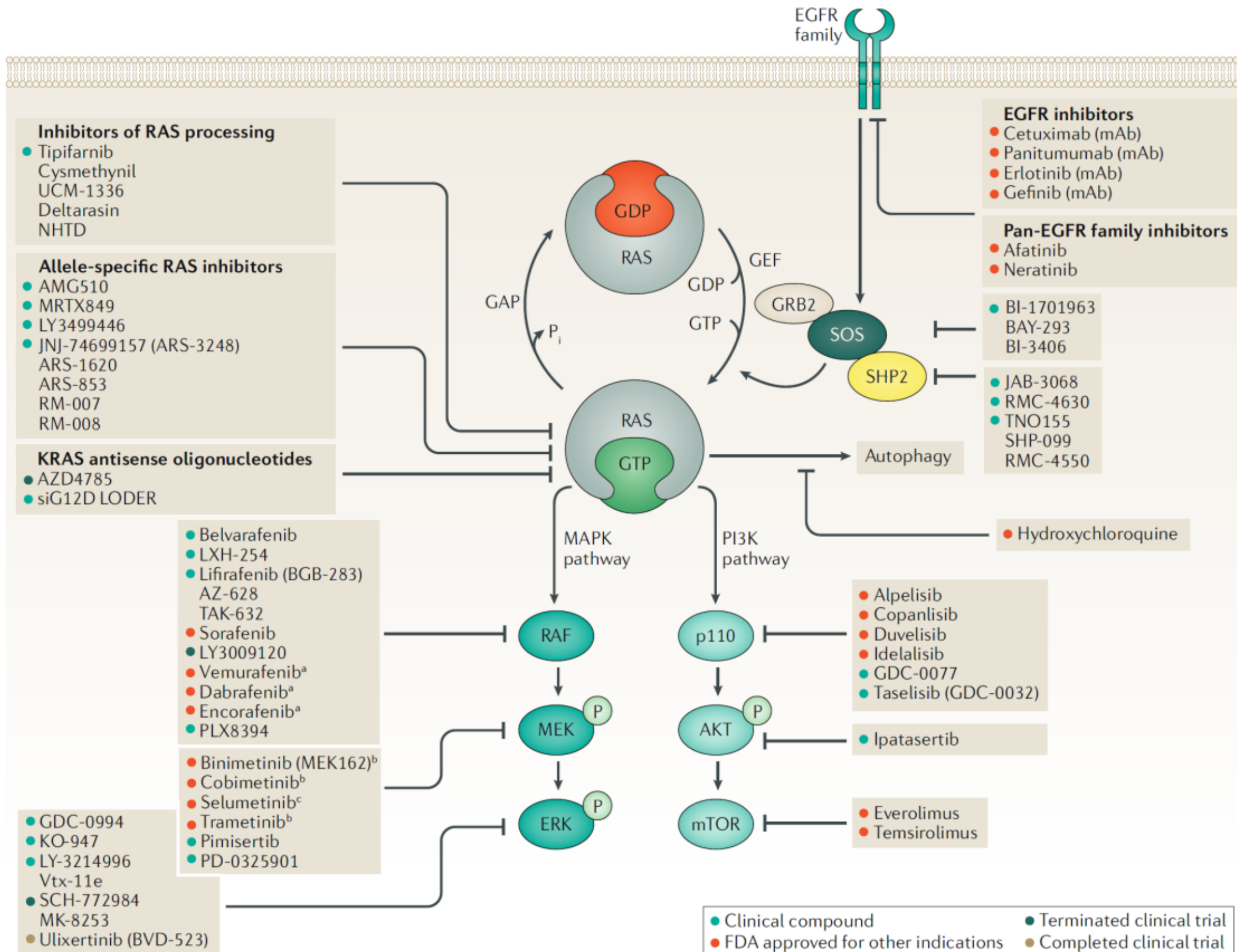
# RMC-4630 (SHP2i) single agent activity—enriched for *KRAS* G12C and *NF1* LOF



Preliminary evidence of increased T cell infiltration

Koczywas et al. AACR 2021 LBA 001

# Emerging Treatment Combinations in *KRAS*-Mutant NSCLC



KRAS G12C inhibitor



PD-1 inhibitor

Pan-EGFR TKI

SHP2 inhibitor

CDK4/6 inhibitor

FGFR inhibitor



Thank you!

QUESTIONS?