

Evolving Targeting Therapy for BRAF-mutated Colorectal Cancer

Scott Kopetz, MD, PhD.

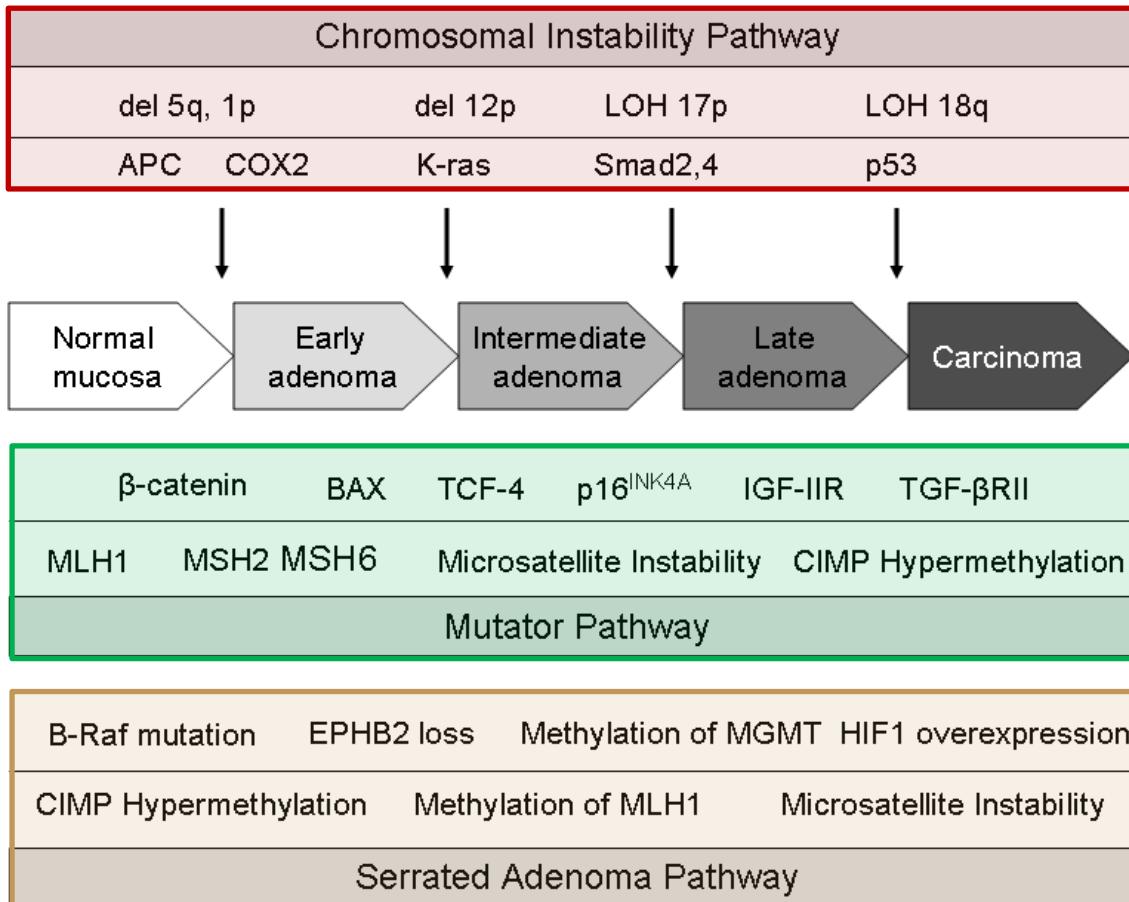
Department of GI Medical Oncology

MD Anderson Cancer Center

Disclosures

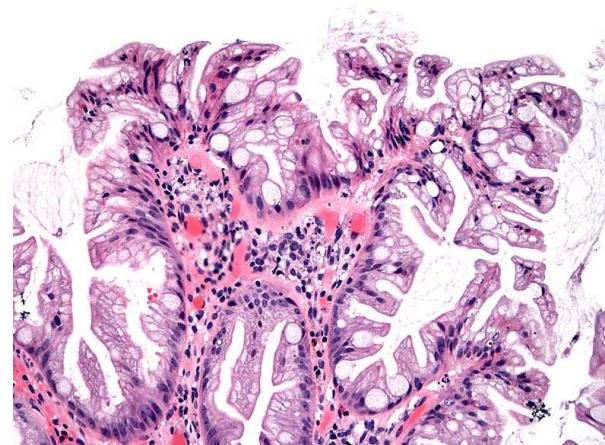
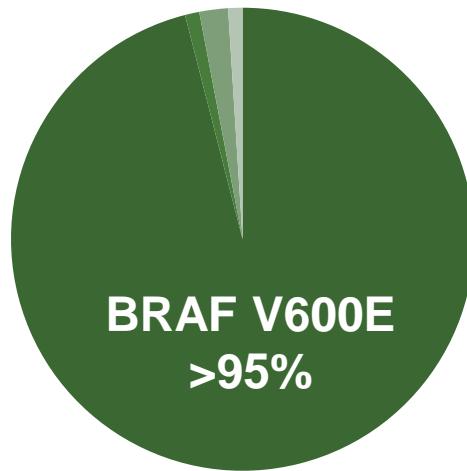
- Advisory board, research funding: Amgen, Roche/Genentech
- Research funding: Biocartis, Sysmex
- Advisory board: Taiho, BMS, Merrimack, Bayer, Array, Sanofi, GSK

Classic Mechanisms of Carcinogenesis in Colorectal Cancer



Serrated Adenoma / BRAF^{mut} Subgroup

- BRAF mutations reflect a unique subset of CRC
 - Vast majority are BRAF V600E
 - Unique precursor lesion: Sessile serrated adenoma
 - Prognostic importance in early and late stage

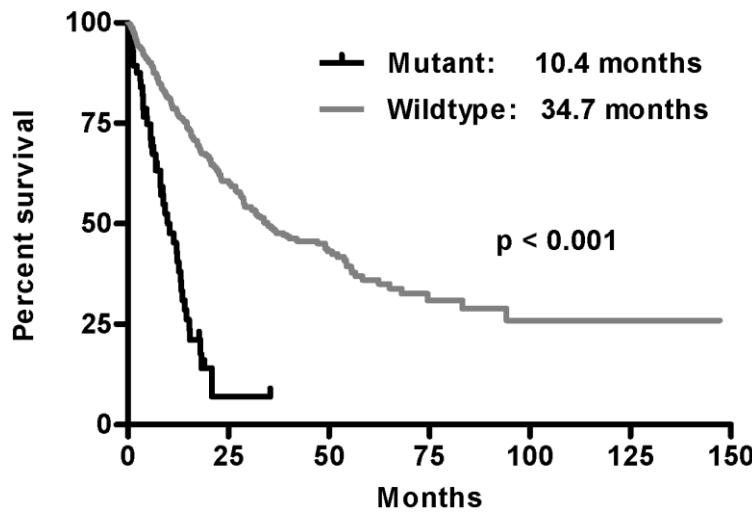


Clinical Utility of BRAF^{mut} in Standard of Care for CRC

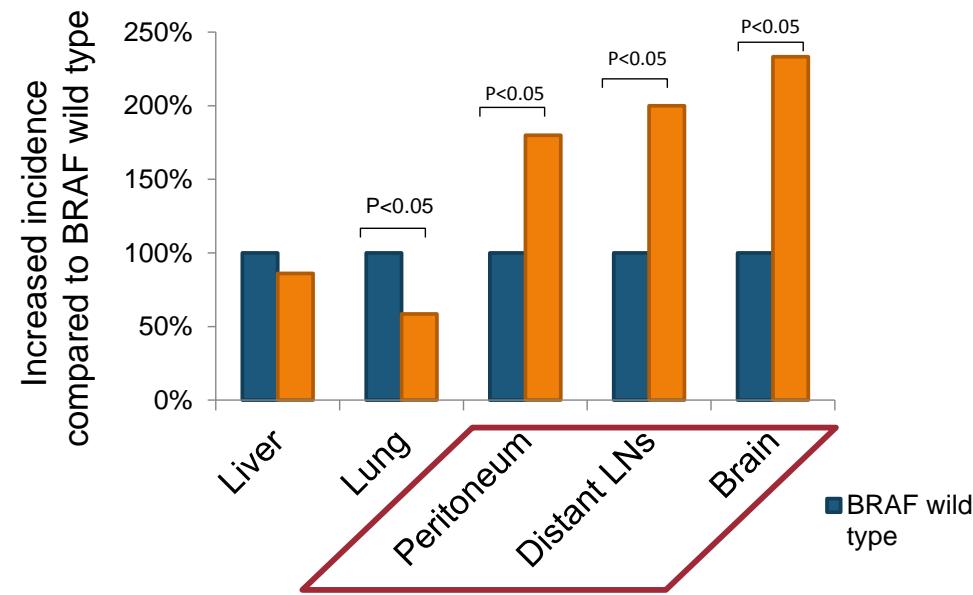
- Determining Sporadic vs Familial MSI-H
 - Part of the recommended algorithm.
- Prognosis for Stage II / III
 - Yes, likely. Not yet in guidelines.
- Prognosis for Stage IV
 - Yes, in guidelines
- Predictive for EGFR Inhibitor sensitivity
 - Mixed data

Unique BRAF^{mut} Clinical Behavior: Metastatic Colorectal Cancer

Very short overall survival

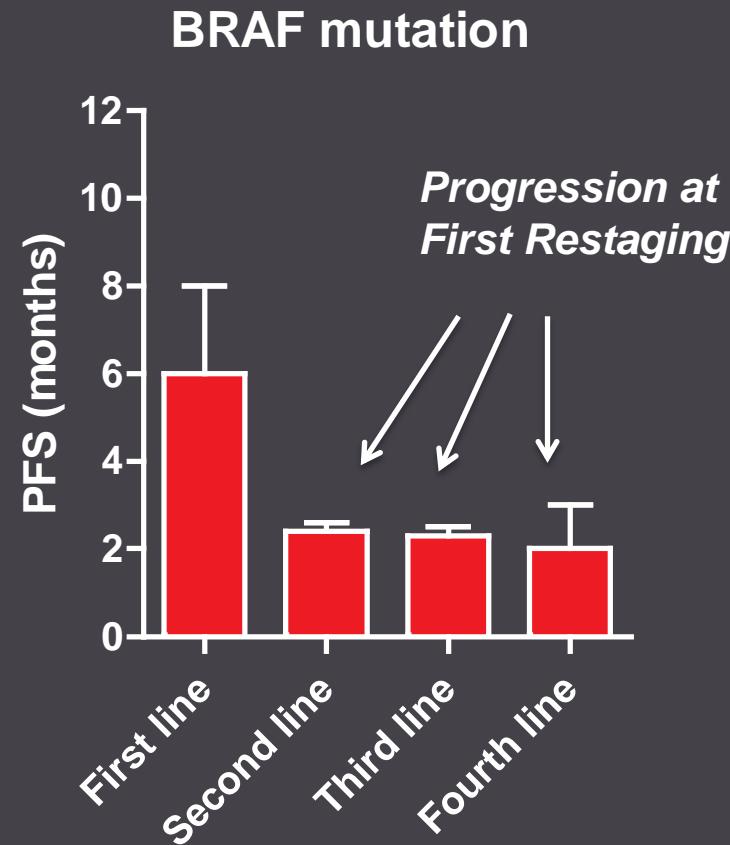
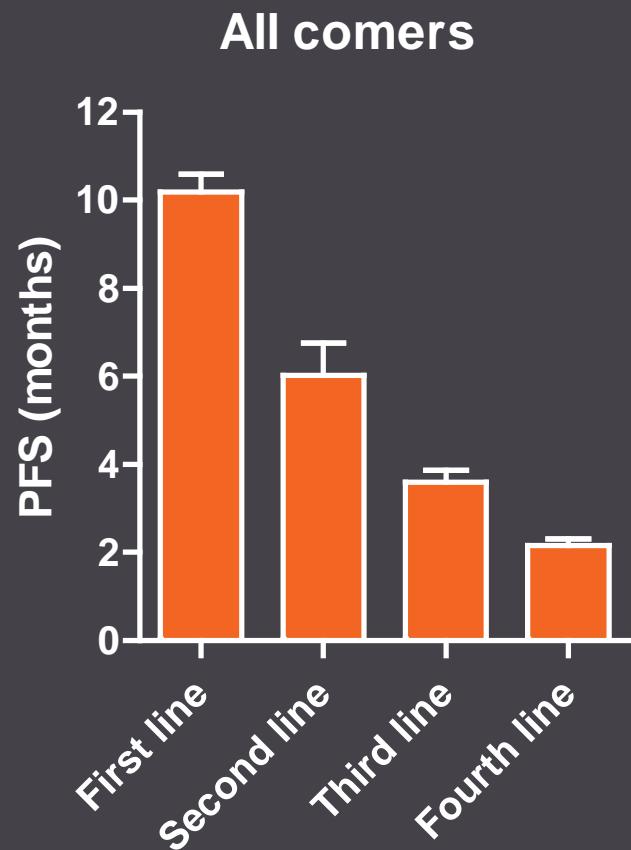


Atypical patterns of metastases



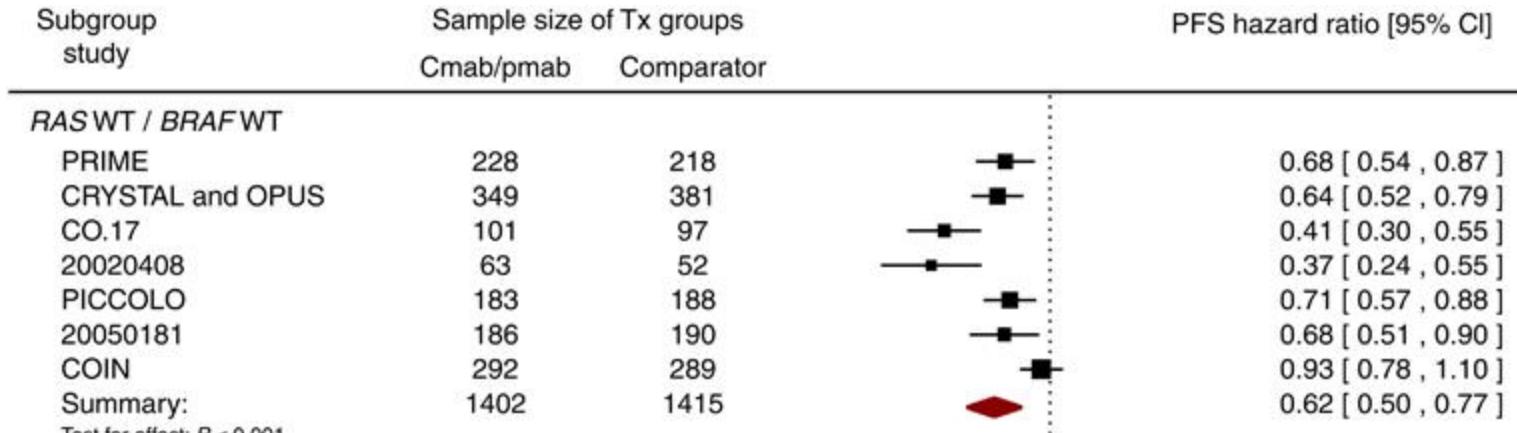
Hazard Ratio of 10.6 for OS
Less than 1 year OS

Limited Benefit of Standard of Care for BRAF^{mut} CRC Patients



BRAF mutation and EGFR inhibition

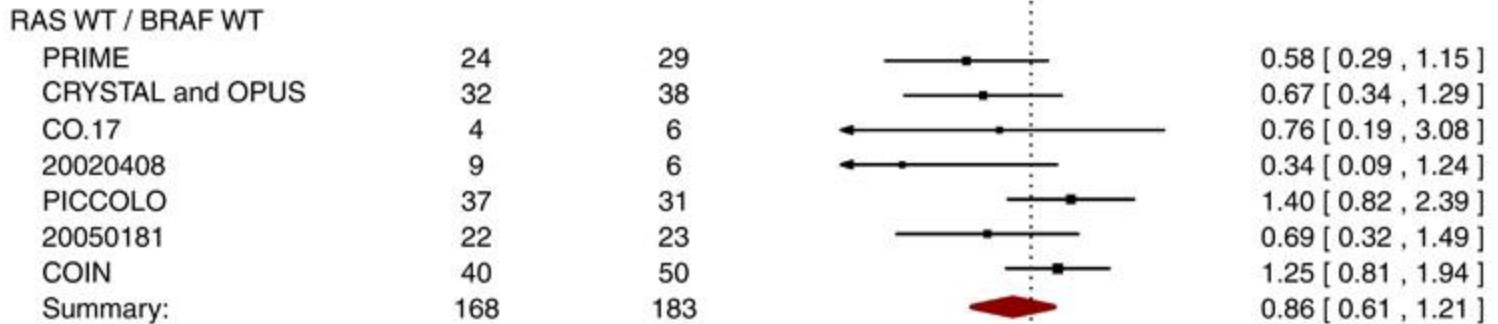
BRAF WT



Test for effect: $P < 0.001$

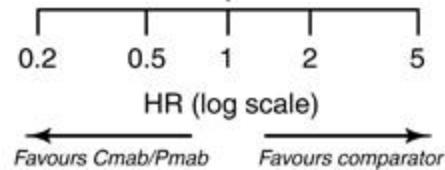
Heterogeneity: $I^2 = 82\%$, $P < 0.001$

BRAF mut



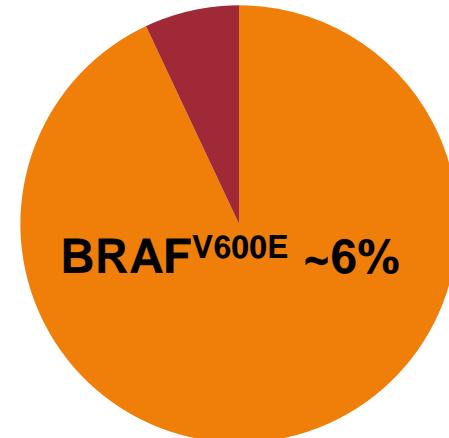
Test for effect: $P = 0.38$

Heterogeneity: $I^2 = 39\%$, $P = 0.13$

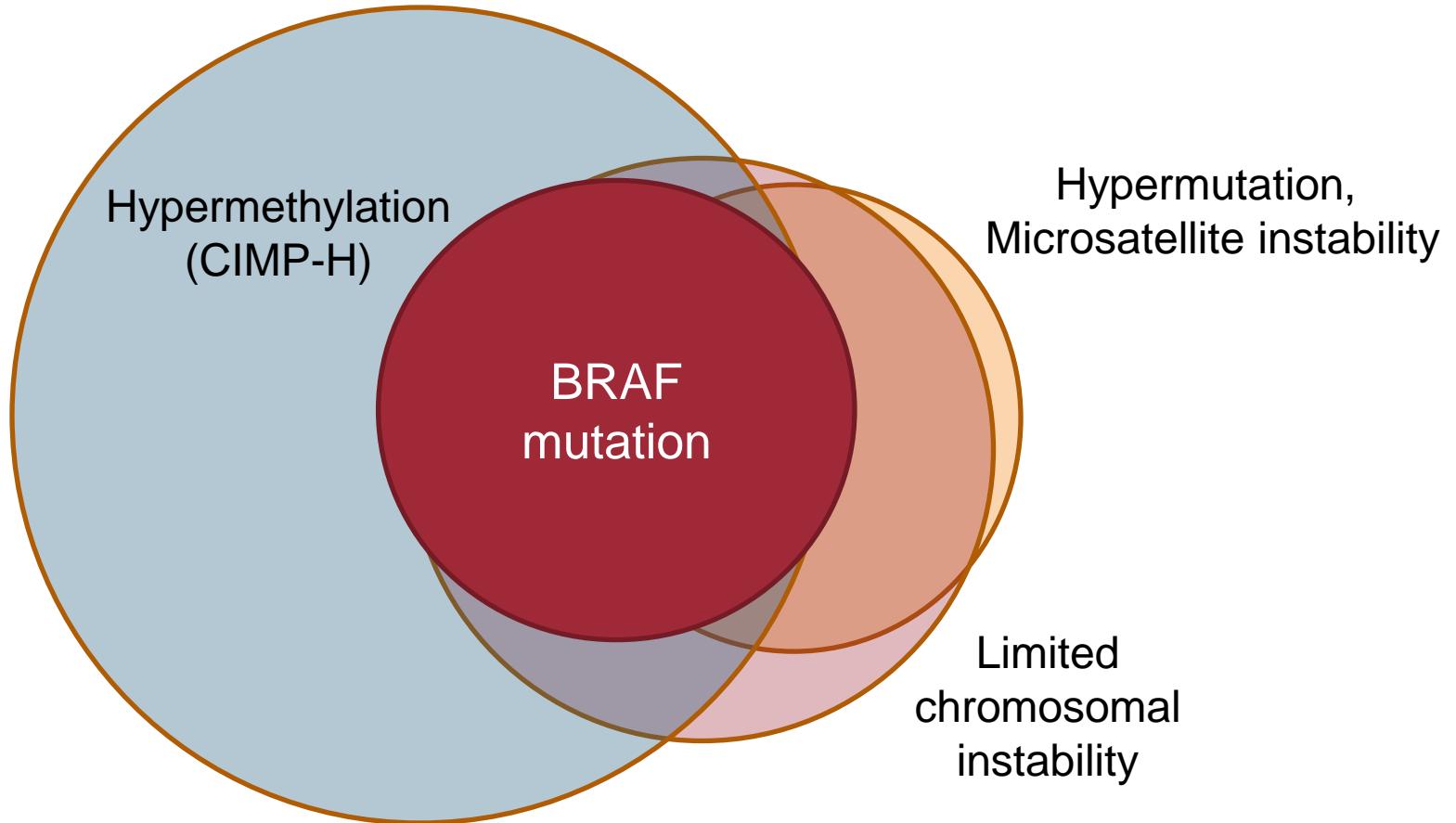


Why BRAF in CRC provides an example for discussion

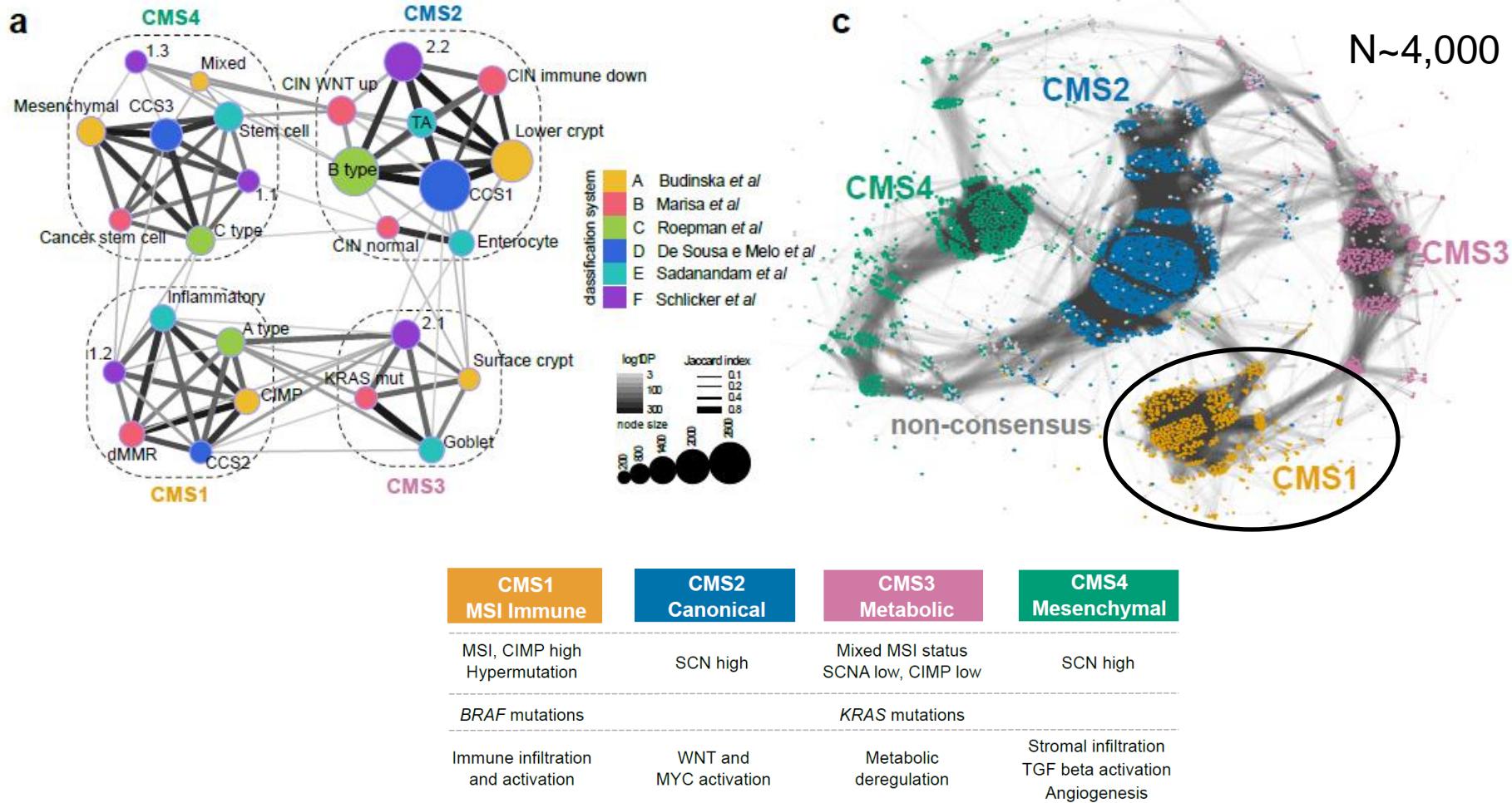
- Defines a unique molecular AND clinical subset
 - Substantial clinical need
- Compensatory activity uncovers at least two key drivers: BRAF and EGFR
- Proof of concept is established
 - With strong preclinical rationale established
 - In multiple single arm studies
 - In ongoing randomized CRC trials
- Lessons learned for iterative drug development



Landscape of BRAF^{mut} Colorectal Cancer



Consensus Molecular Subtypes: CMS1



How to treat these patients?

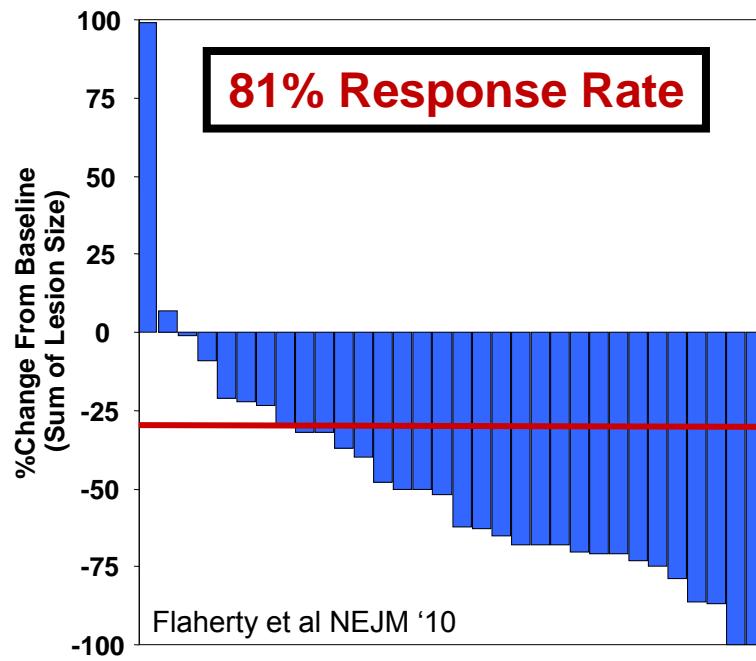
BRAF inhibitor



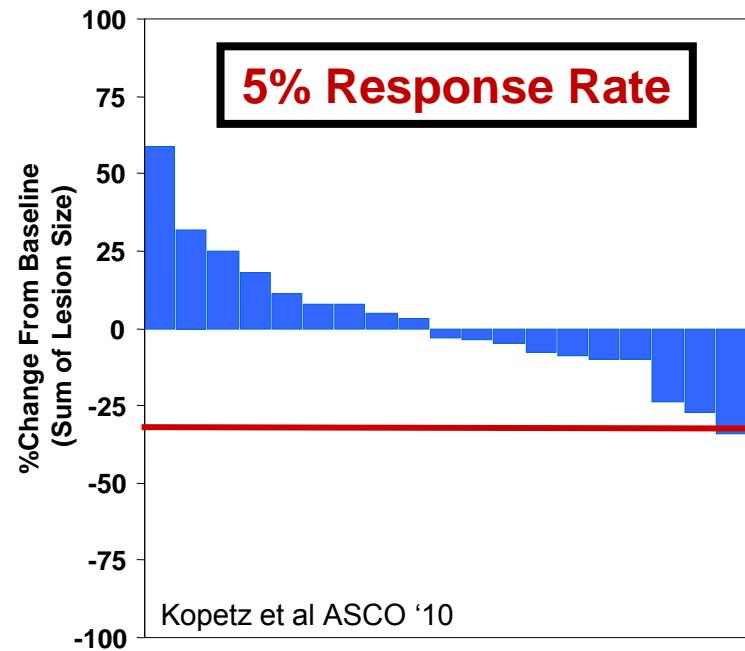
BRAF mutation

Vemurafenib (PLX4032)

Refractory Melanoma



Refractory Colorectal

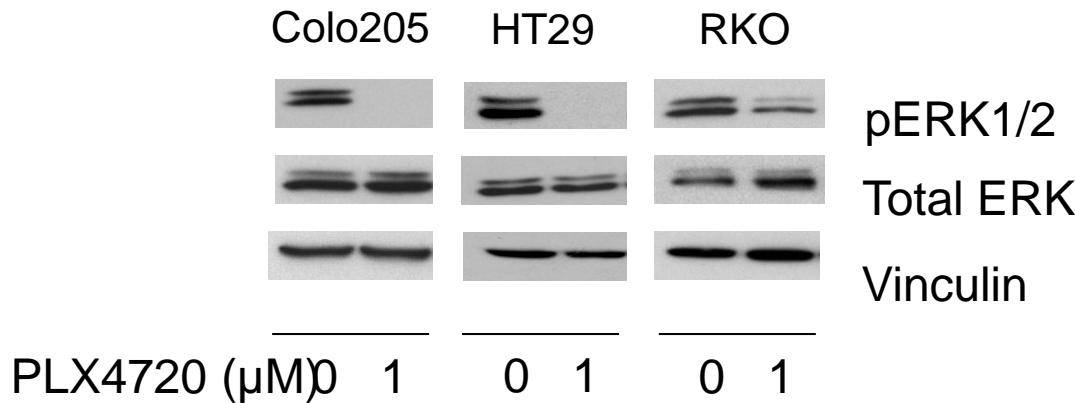


Preclinical data fails to demonstrate sufficient MAPK inhibition in CRC, unlike melanoma

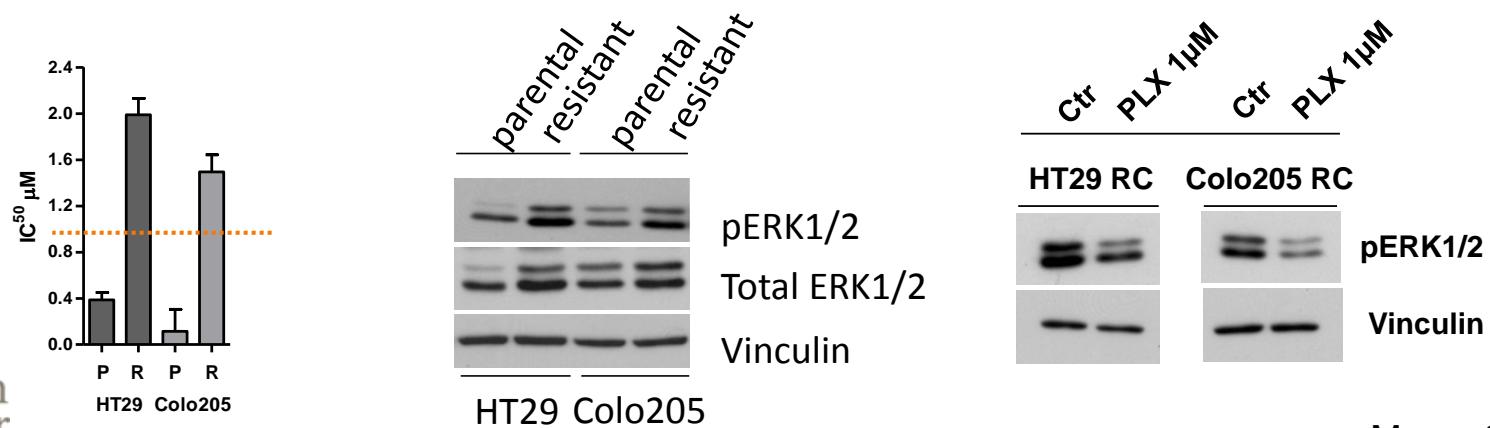
Mao, et al CCR '14

Persistent MEK activation: Expansion from melanoma concepts

ERK inhibition is incomplete in some CRC cell lines



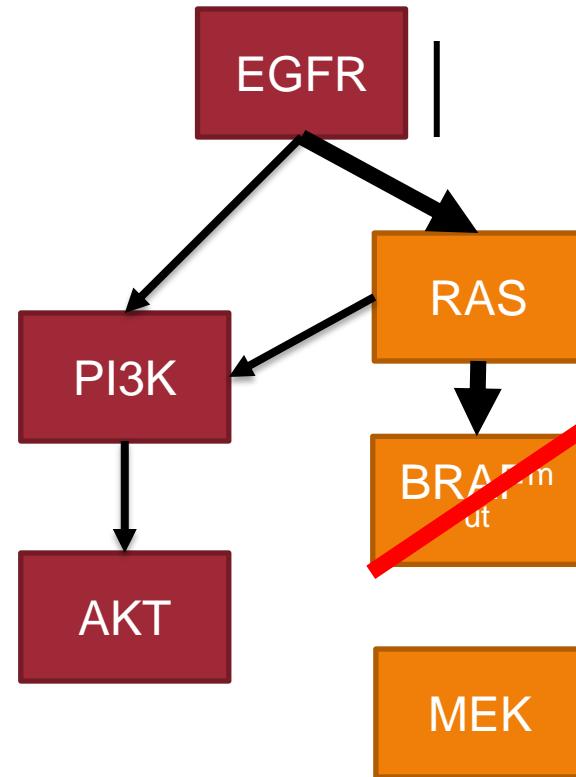
Acquired resistance is associated with increased pERK and incomplete inhibition



BRAF resistance through MEK reactivation

IF the problem is only incomplete MEK inhibition....

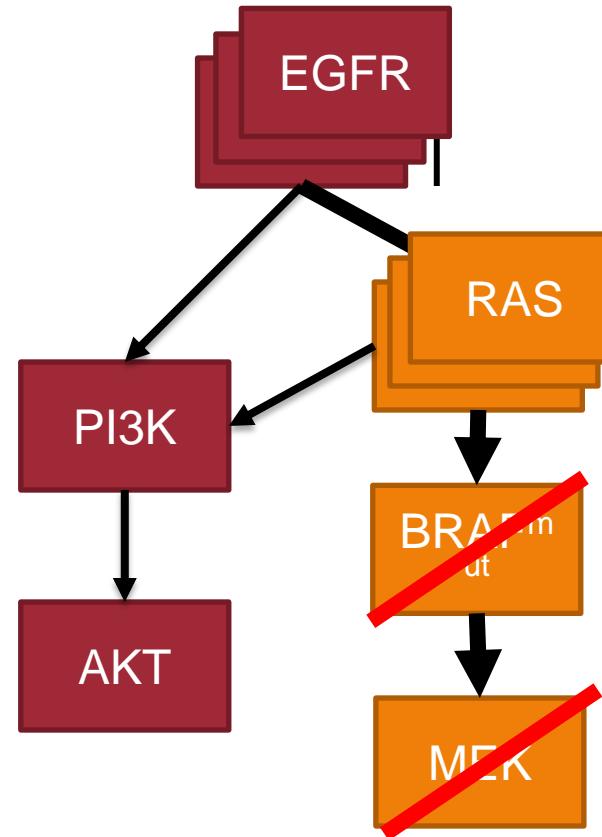
Then treat with dual inhibition of BRAF + MEK



BRAF resistance through MEK reactivation

IF the problem is only incomplete MEK inhibition....

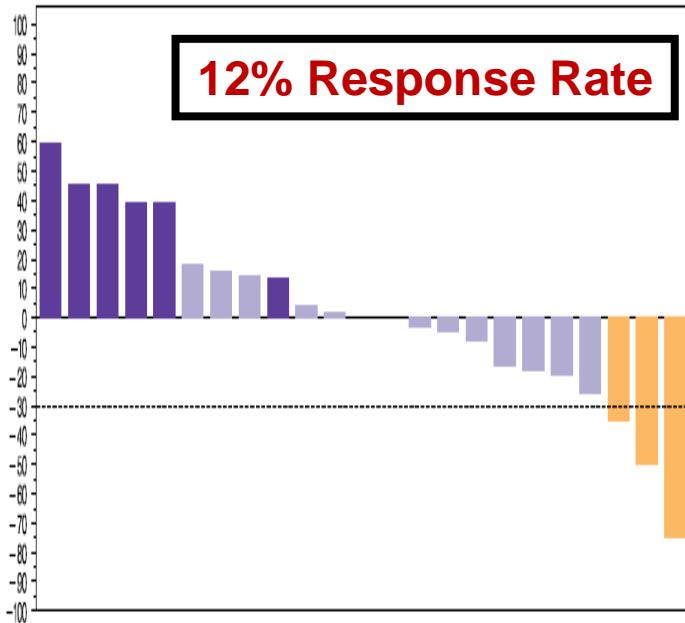
Then treat with dual inhibition of BRAF + MEK



Minimal Improved Efficacy with Dual BRAF + MEK Inhibition

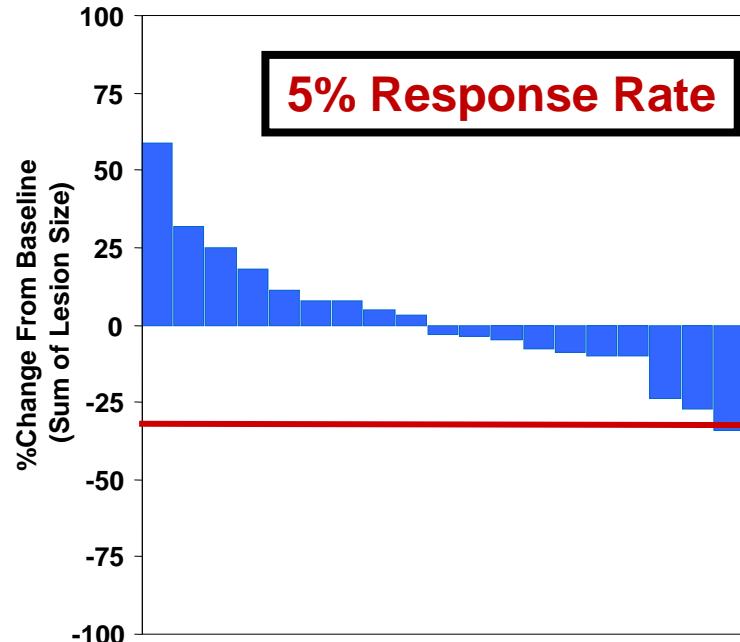


BRAF + MEK inhibition



GSK212 + GSK436

BRAF inhibition

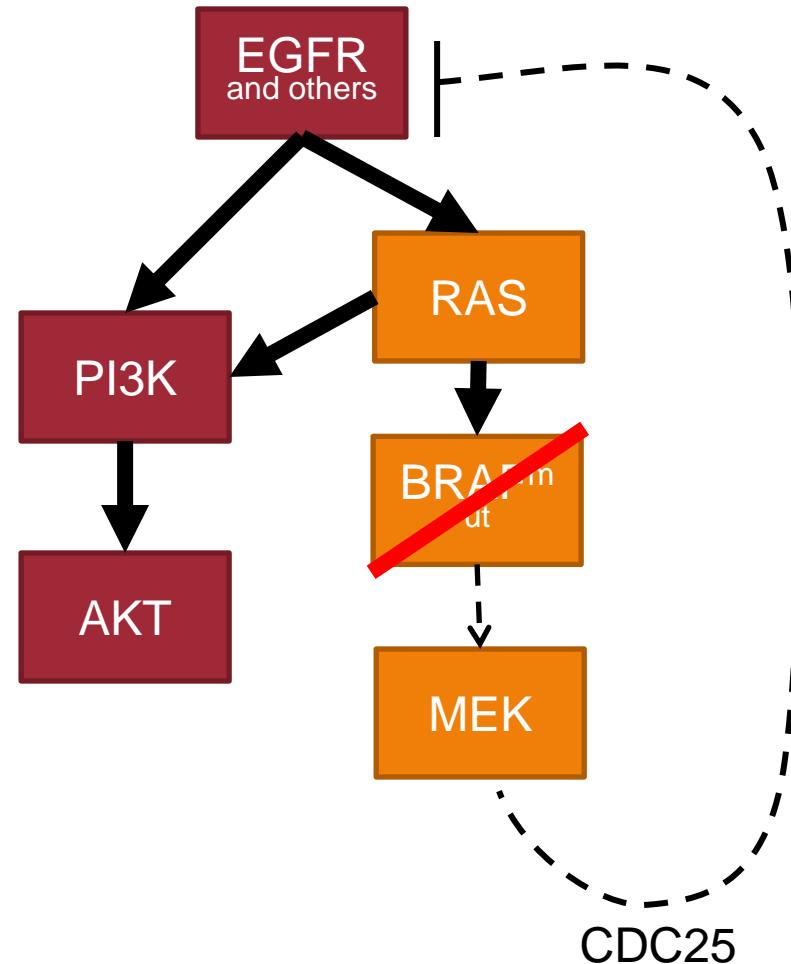


Vemurafenib

Key Finding: Feedback EGFR Signaling

Perhaps the problem
isn't ONLY incomplete
MEK inhibition....

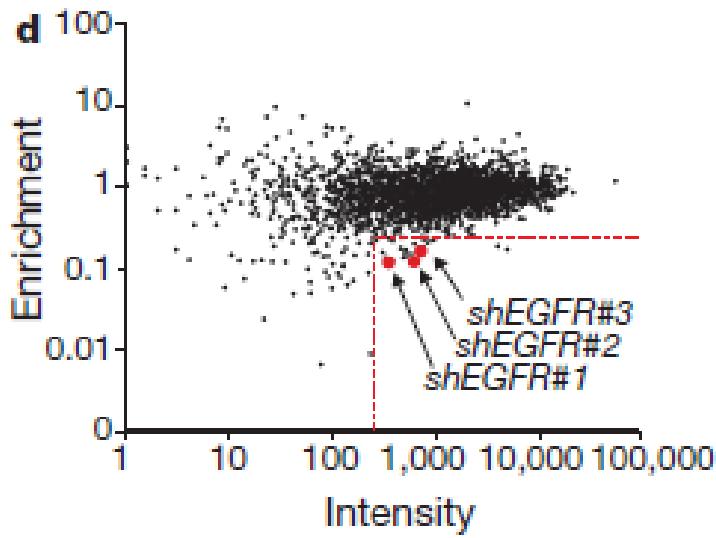
Treat with EGFR +
BRAF inhibitors



Prahallas et al Nature '12, Corcoran et al Can Disc '12

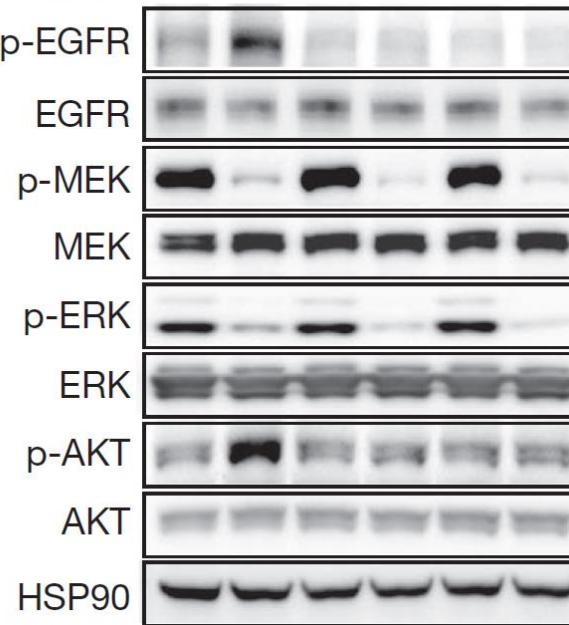
Unbiased Synthetic Lethality Screen: EGFR Identified as a Synergistic Partner

HT29 cell line (Sensitive)



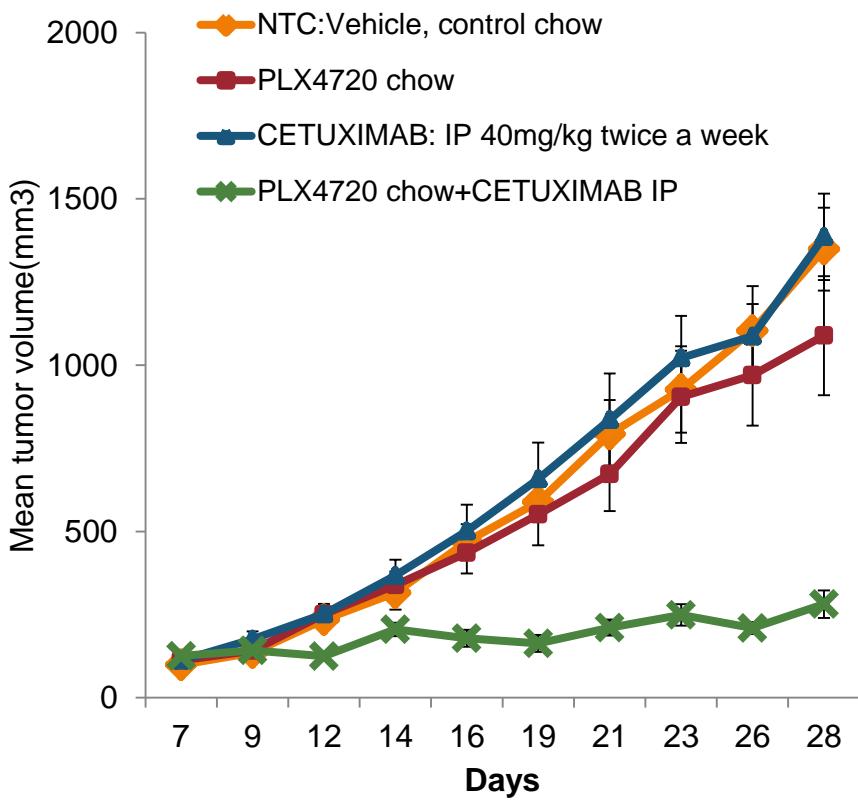
EGFR identified

Gefitinib:	-	-	-	-	+	+
Cetuximab:	-	-	+	+	-	-
PLX4032:	-	+	-	+	-	+

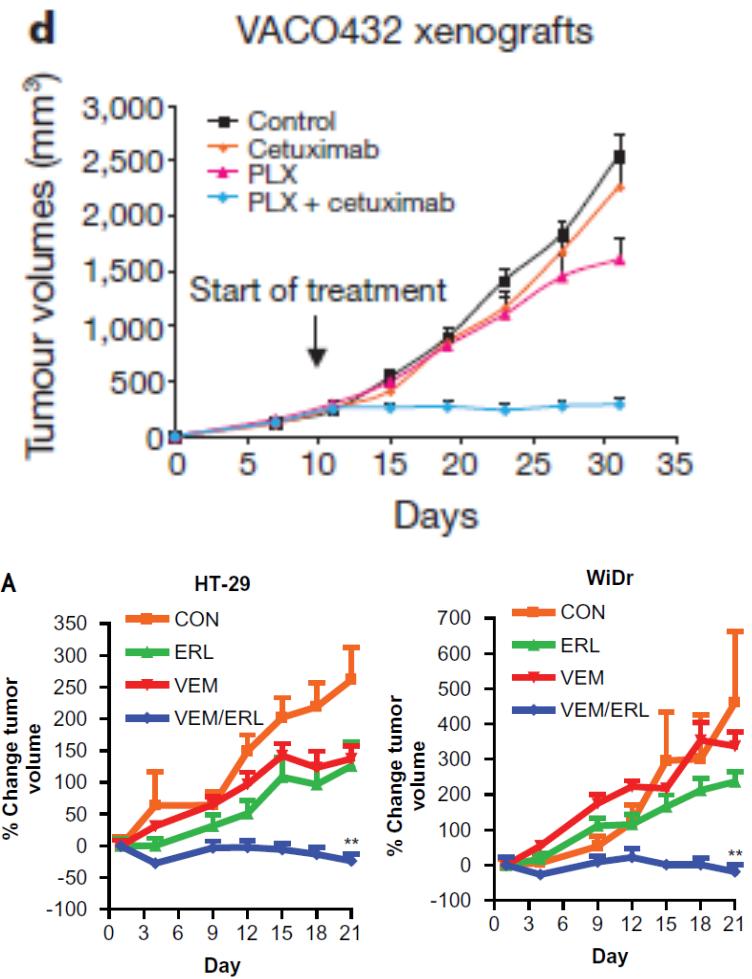


Synergy in Murine Models: BRAF + EGFR

Cell line and Patient-derived xenograft models

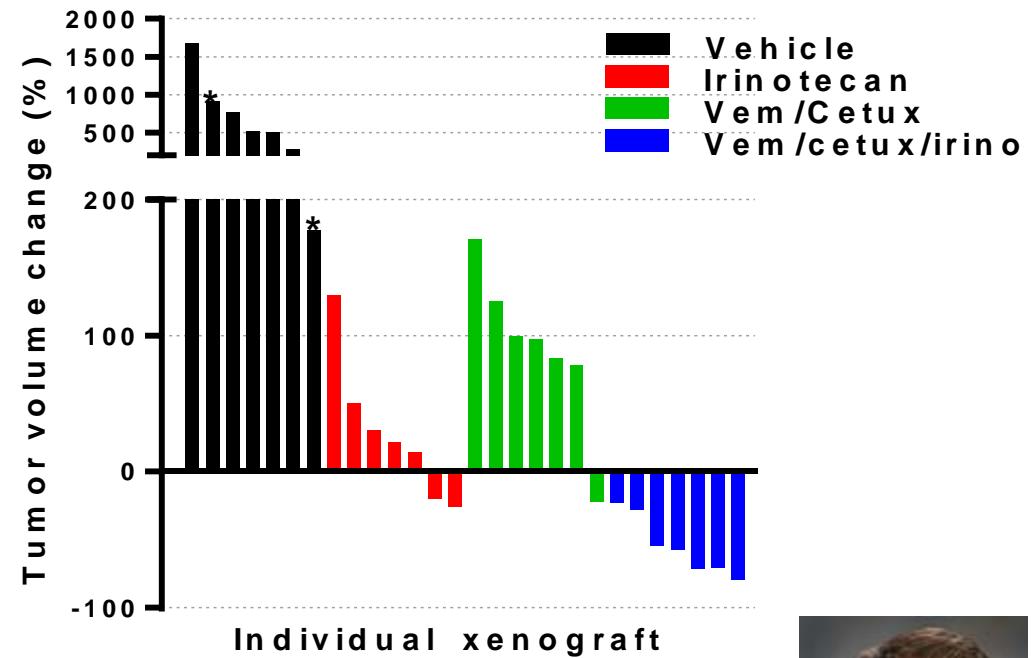
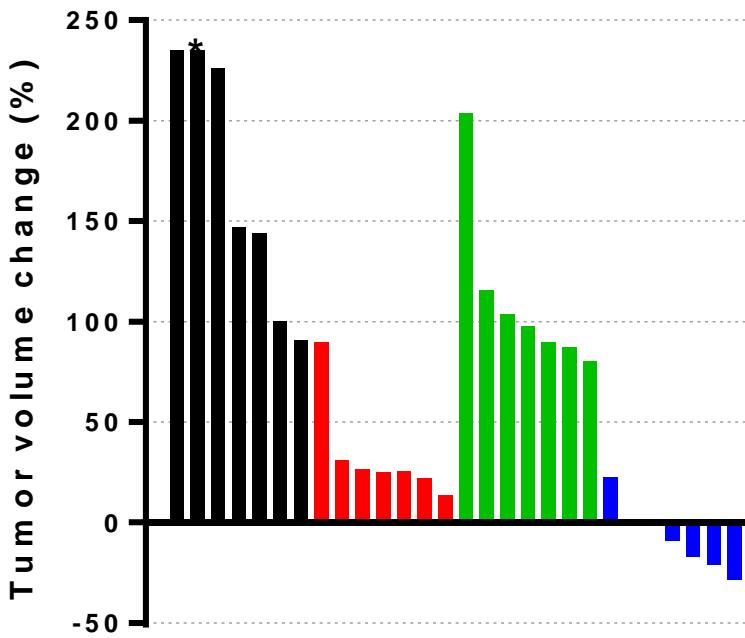


Kopetz, unpublished; Yang et al Can Res '11

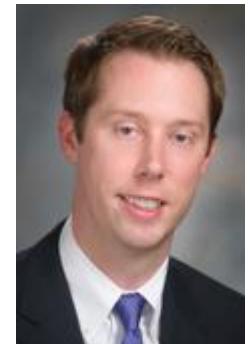


Prahallad et al Nature '12; Corcoran et al Can Disc '12

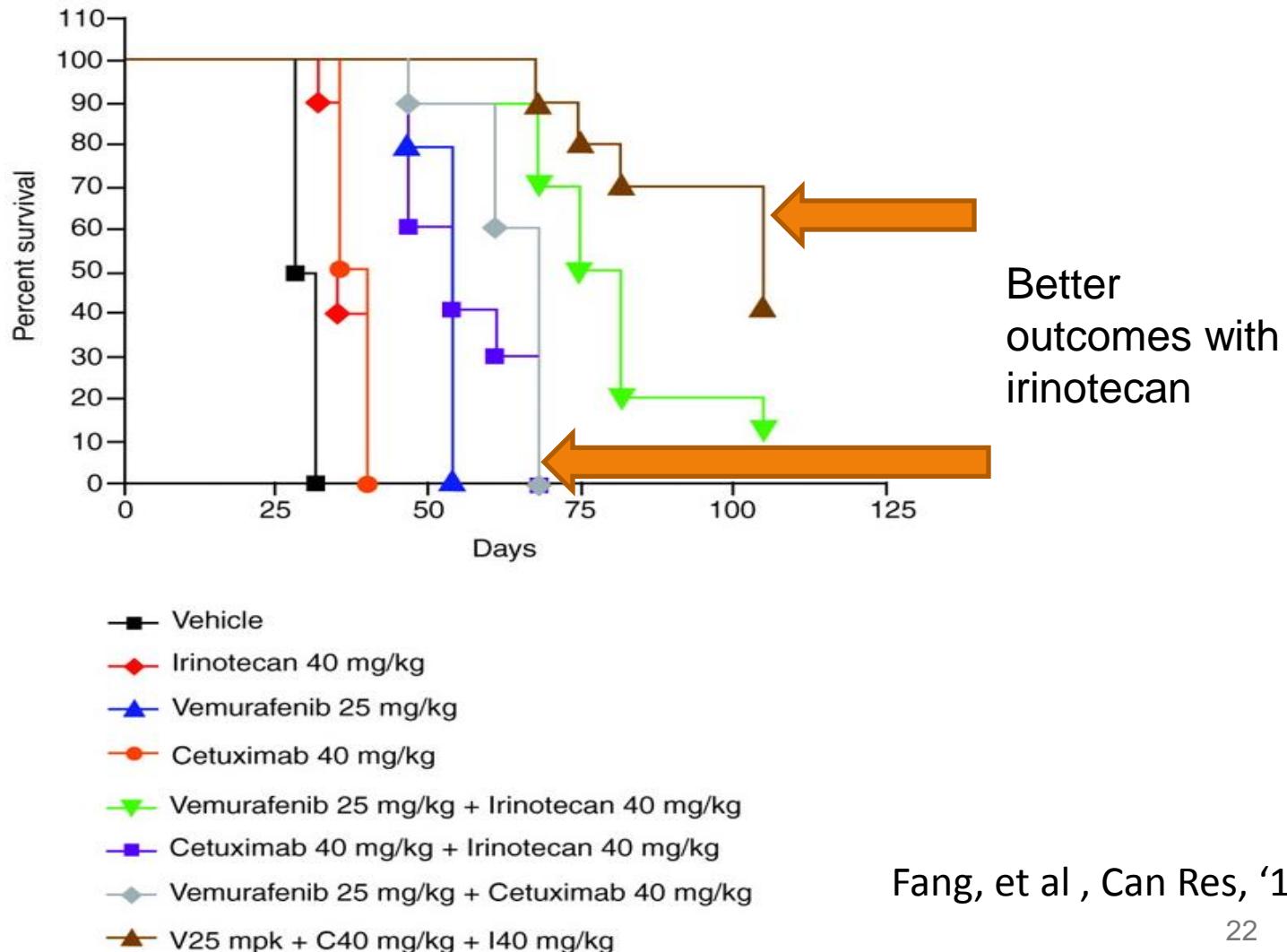
Irinotecan combined with EGFR/BRAF inhibition induces regressions



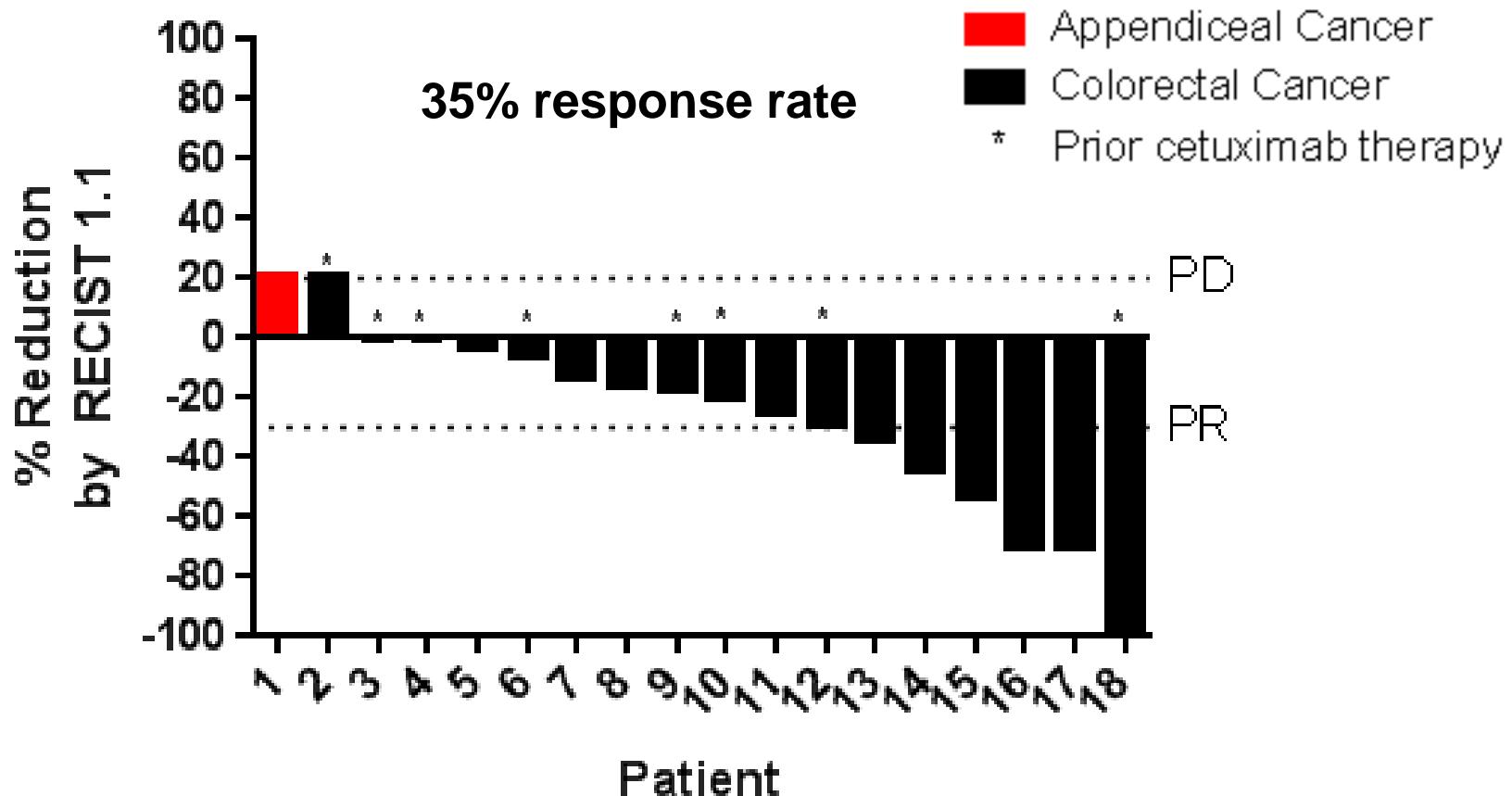
Van Morris, Alex Sorokin



Improved *Survival* with Cetuximab and Vemurafenib combined with Irinotecan



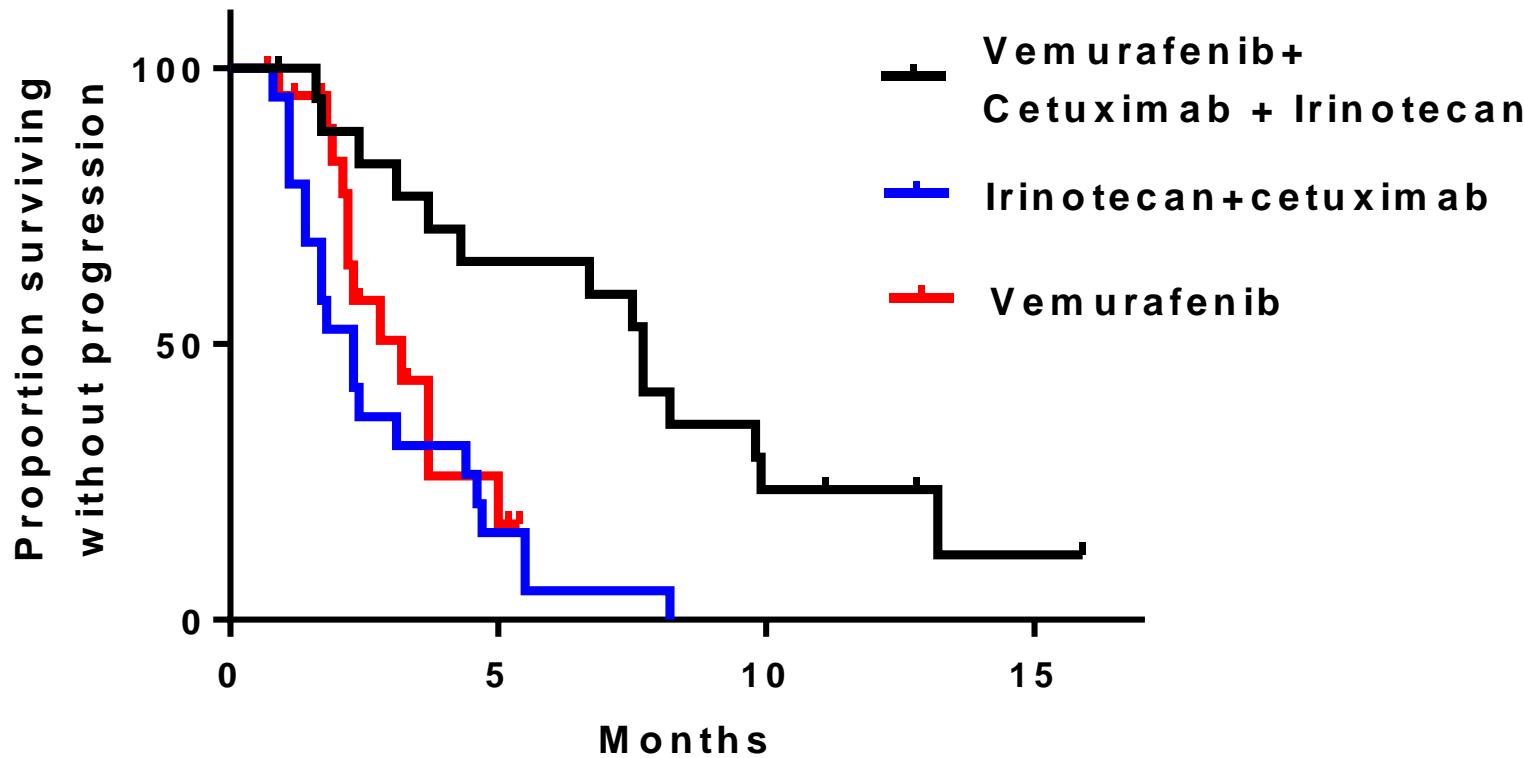
Phase 1B of Cetuximab, Vemurafenib, Irinotecan: High Response Rate



Historical response rate is <10% for cetuximab and irinotecan with PFS of 2 months.

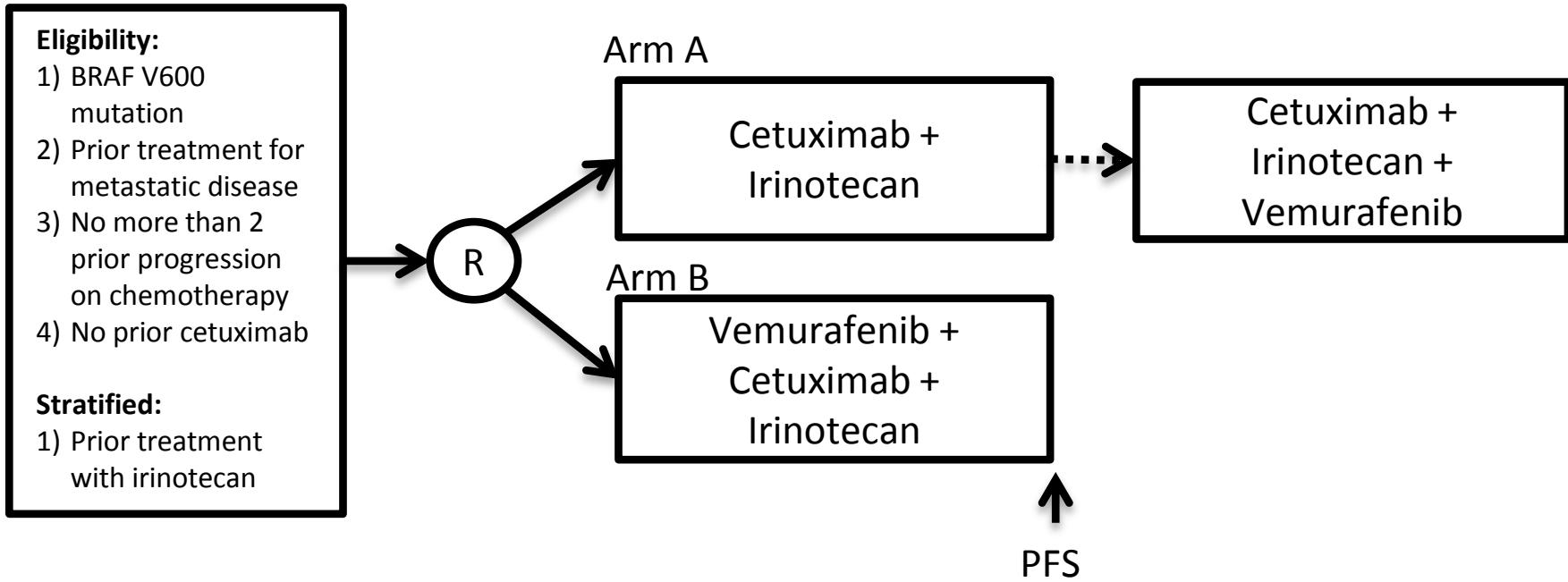
David Hong et al. ASCO, '15; Morris, CCC, '14

Cross-Trial Comparison: Phase 1B vs Historic Control for mBRAF CRC



SWOG 1406: BRAF + EGFR

with availability through other cooperative groups

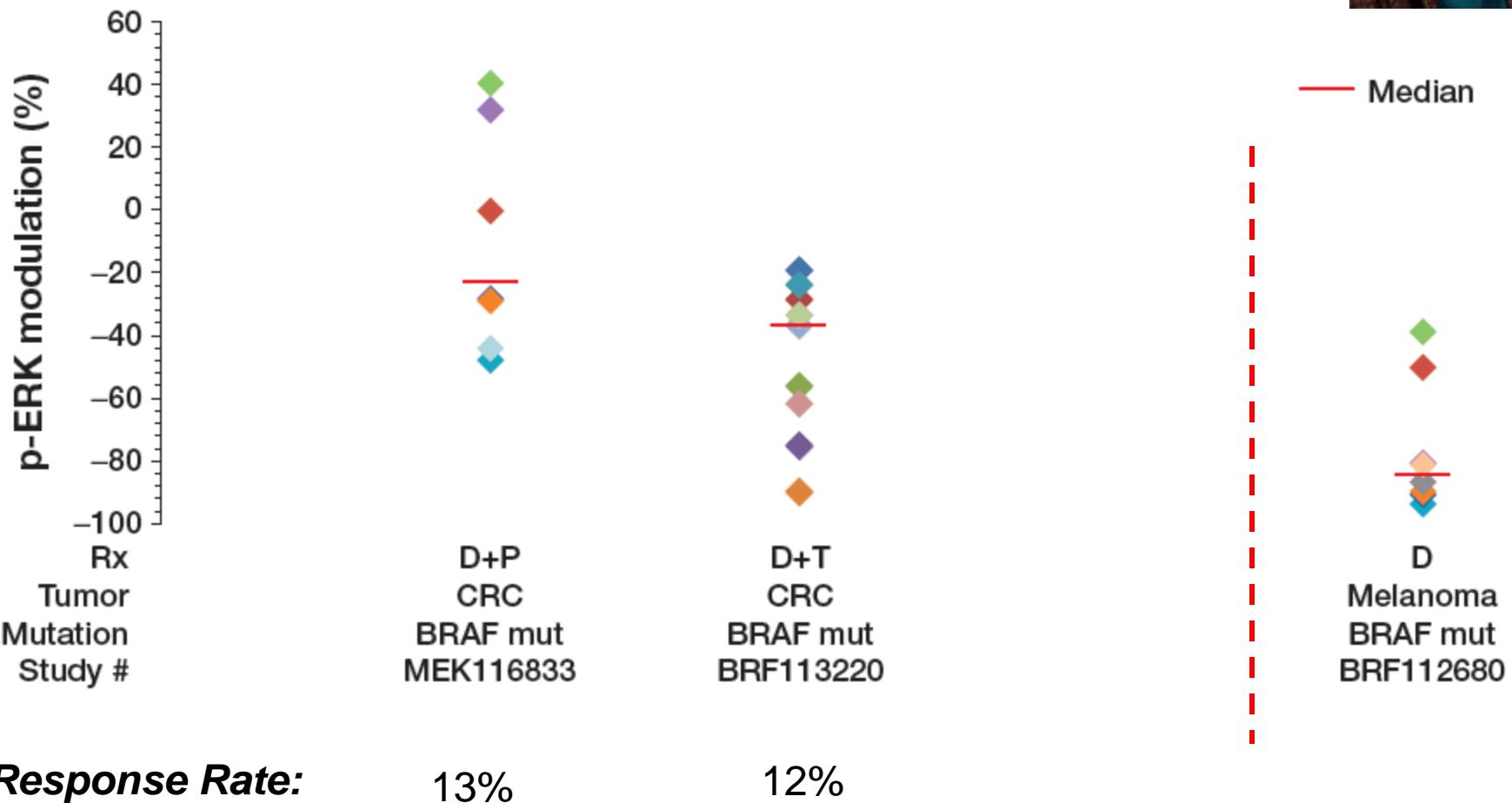


Historical response rate is <10% for cetuximab and irinotecan, with PFS of 2.4 months for BRAF^{mut}

Target HR 0.5 for PFS, with 2-sided alpha 5%, power 90%
N= 105 patients

Results Fall '16, Kopetz, PI

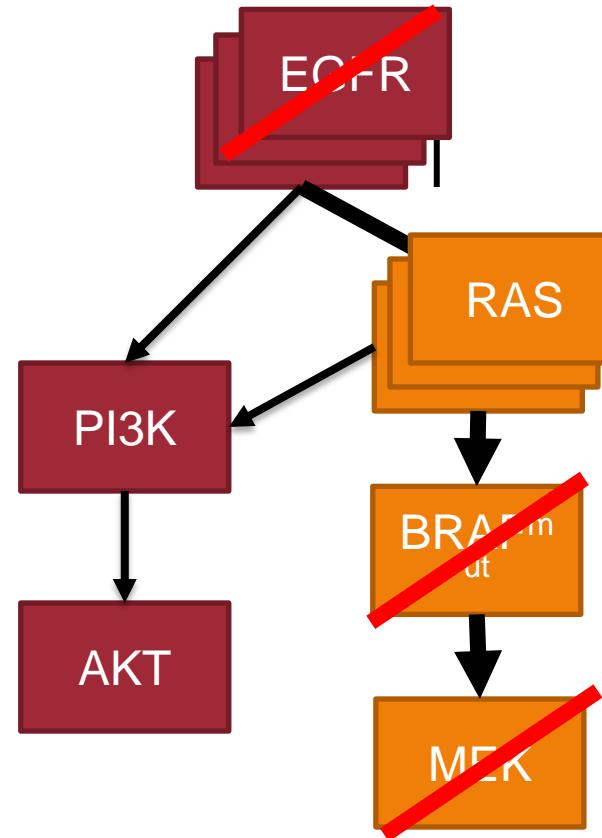
Cross-Study Comparison of Phospho-ERK Modulation



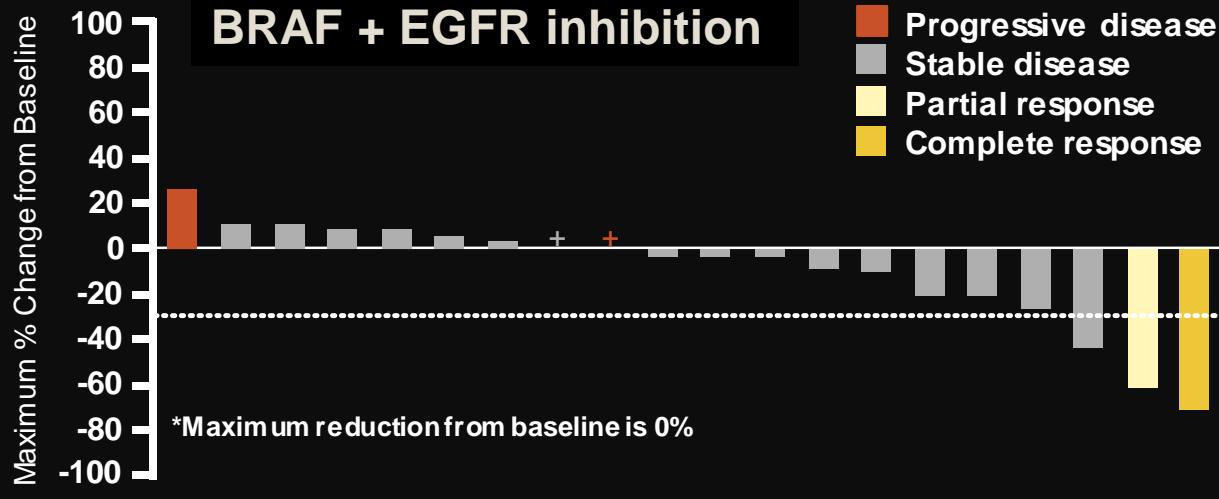
BRAF resistance through MEK reactivation

IF the problem is only incomplete MEK inhibition....

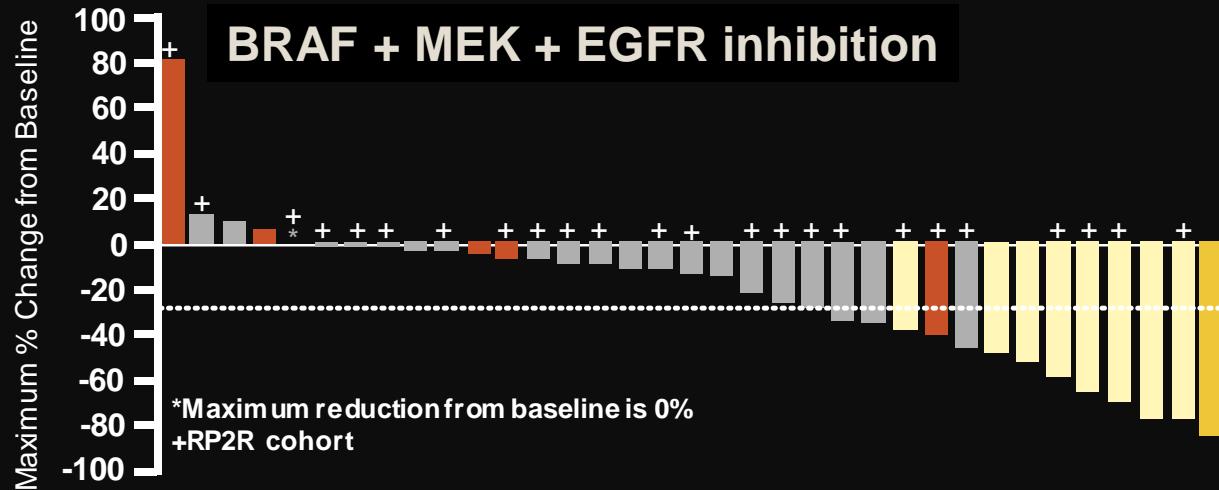
Then treat with triplet inhibition of
BRAF + MEK + EGFR



Dabrafenib + Panitumumab +/- Trametinib

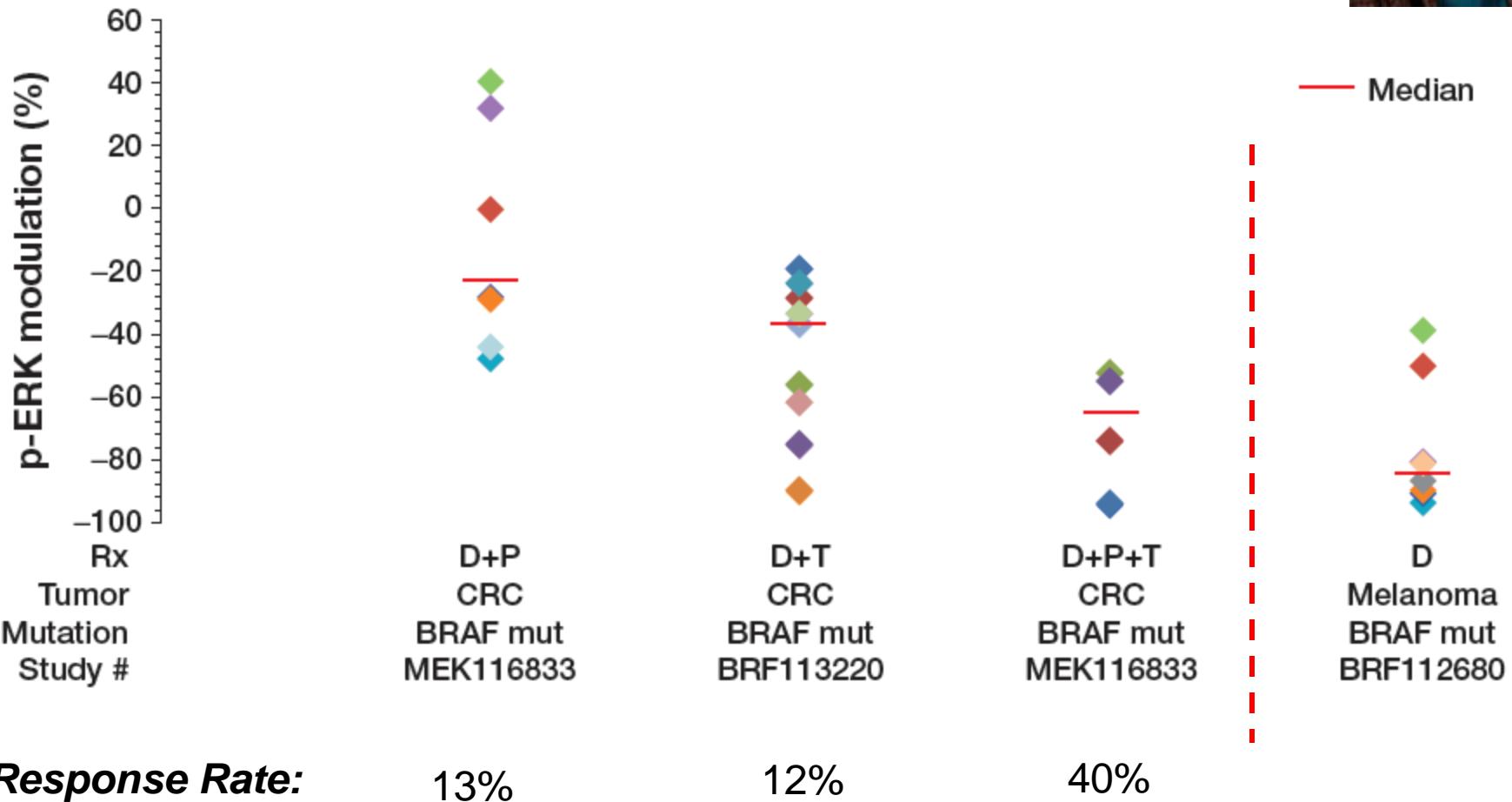


D+P (N=20)
CR+PR: 2 (10%)

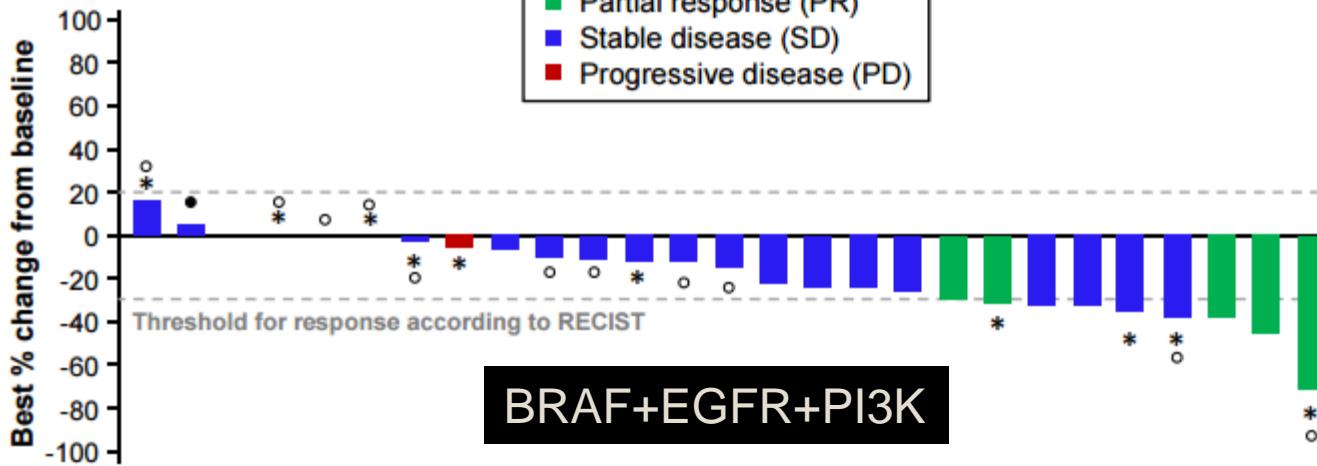
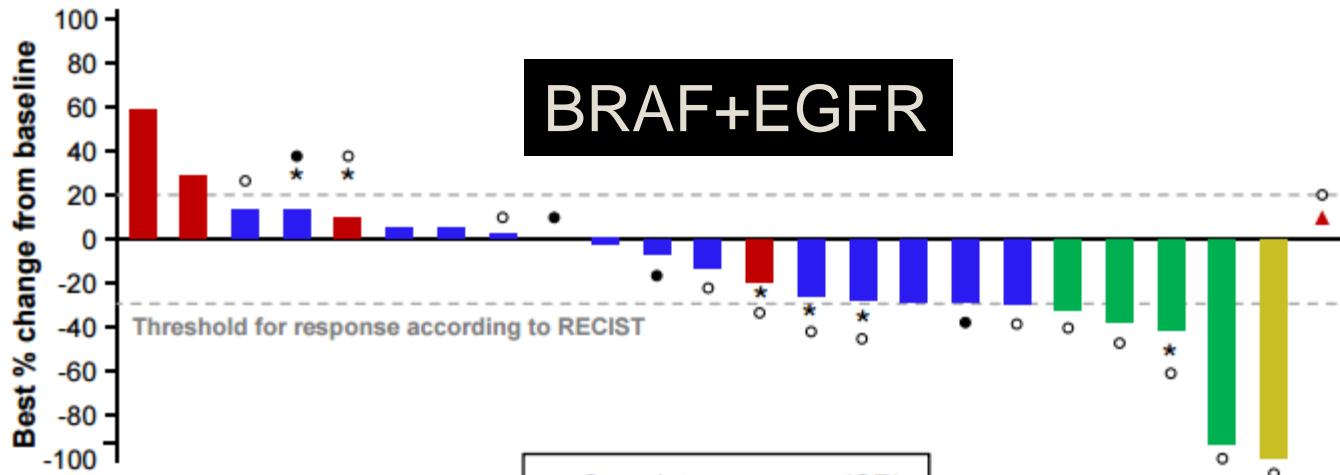


D+P+T (N = 35)
CR+PR: 9 (26%)

Cross-Study Comparison of Phospho-ERK Modulation



Encorafenib + Cetuximab +/- BLY719



Dual Combination

PD: 4 (15%)
SD: 14 (54%)
PR: 5 (19%) ^a
CR: 1 (4%)

Triple Combination

PD: 1 (4%)
SD: 17 (61%)
PR: 9 (32%) ^b

See survival data update tomorrow (J. Tabernero)

Geel et al ASCO '14

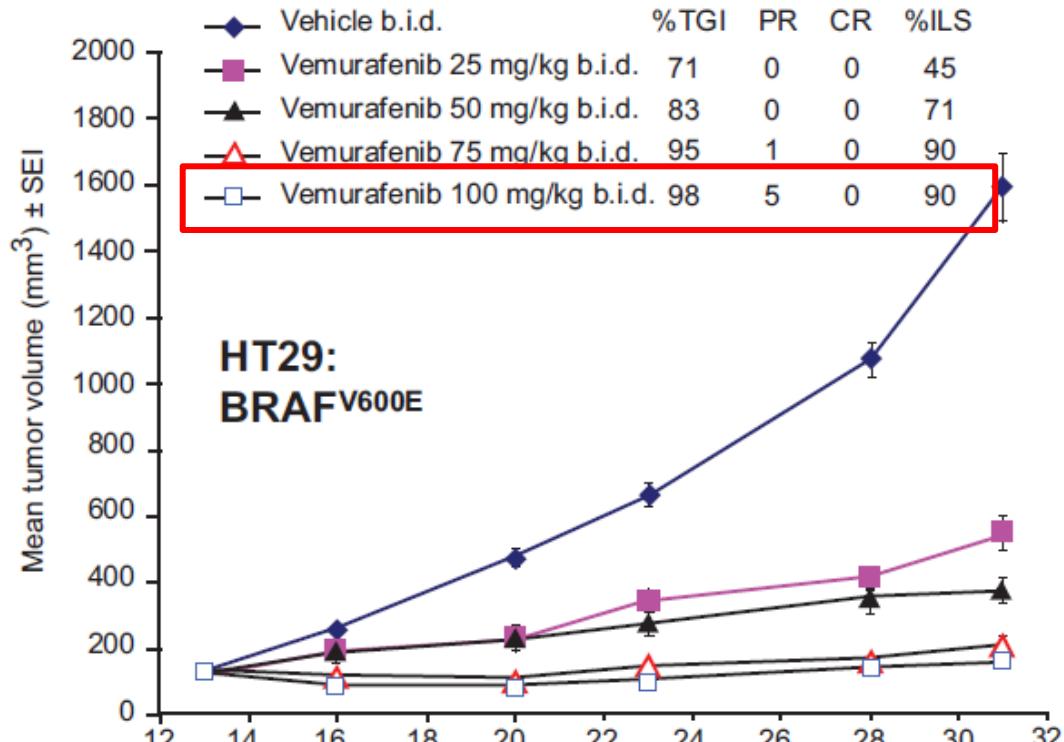
Comparison of RR and PFS for BRAF^{mut} CRC

Regimen	Response	PFS	Citation
Single/Doublet RAF/MEK			
Vemurafenib	5%	2.1 months	Kopetz, JCO'15
Dabrafenib	11%	NR	Falchook, Lancet '08
Encorafenib	16%	NR	Gomez-Roca, ESMO '14
Dabr +Tramet	12%	3.5 months	Corcoran, JCO '15
Doublet with EGFR			
Vem + Panit	13%	3.2 months	Yeager et al CCR '14
Vem + Cetux	20%	3.2 months	Tabernero et al ASCO '14
Encoraf + Cetux	23%	3.7 months	Schellen et al AACR '15
Dabr + Panit	10%	3.4 months	Atreya, ASCO '15
Triplet with EGFR			
Vem + Cetux + Irino	35%	7.7 months	Hong, ASCO '15
Dabr + Tramet + Panit	26%	4.1 months	Atreya, ASCO '15
Encoraf+Cetux+Alpelisib	32%	4.3 months	Schellen et al AACR '15

Learning from BRAF Drug Development Story

- Better preclinical models
- Improving interrogation of response
- Beyond paired biopsies... improved options for interrogating biology of patients under therapy

Vemurafenib *in vivo* in colorectal cancer: Too optimistic?



Representative Cell Line Model

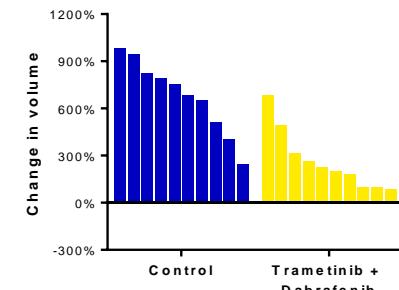
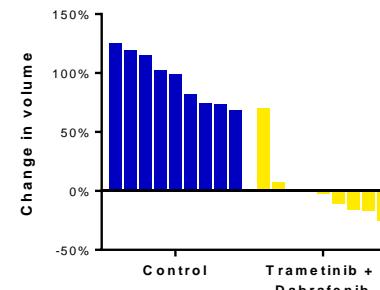
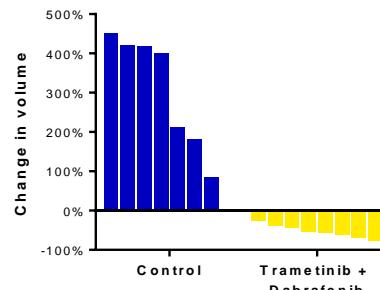
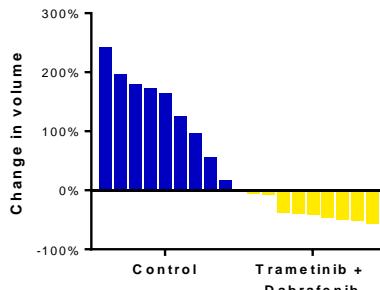
Fang, et al , Can Res, '12

Co-clinical trials: BRAF+MEKi Study



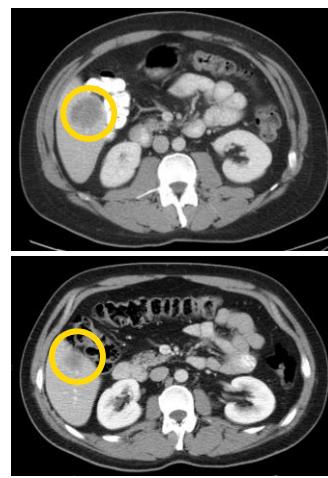
PDX response

A



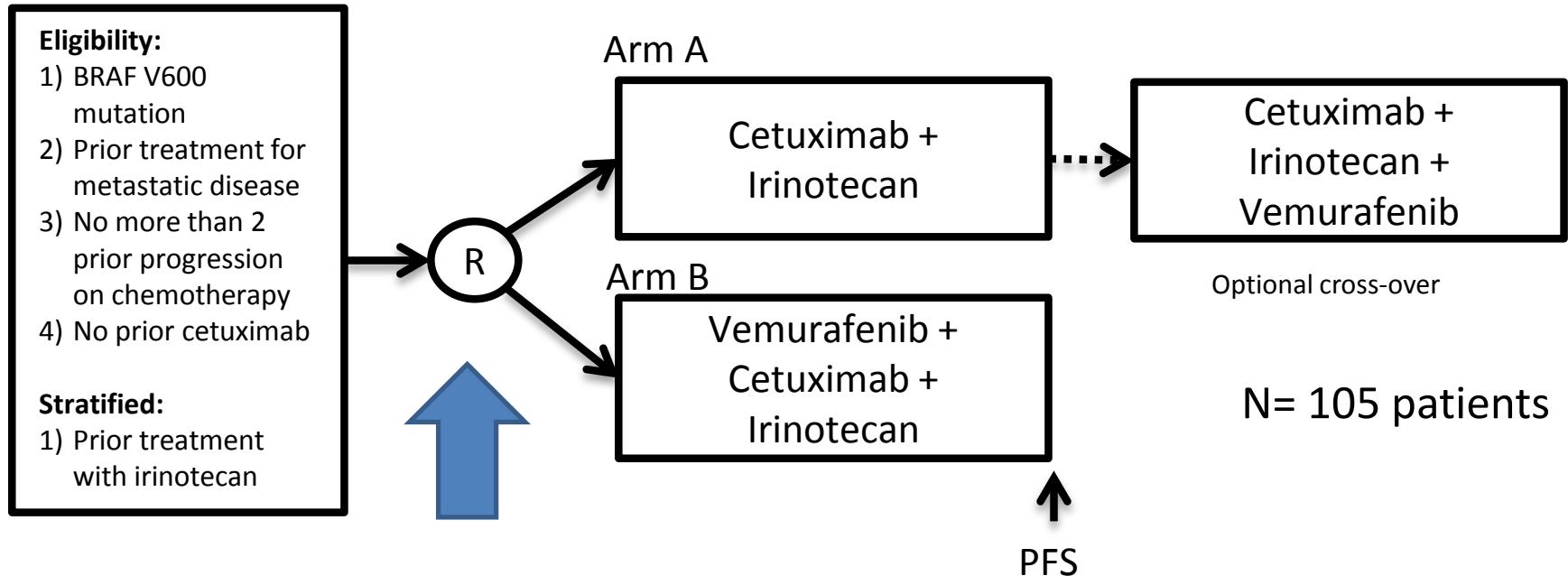
Patient response

B



Corcoran, Atreya et al JCO '15

Prospective PDX Clinical Trial Integration: S1406: BRAF + EGFR Inhibition in mCRC

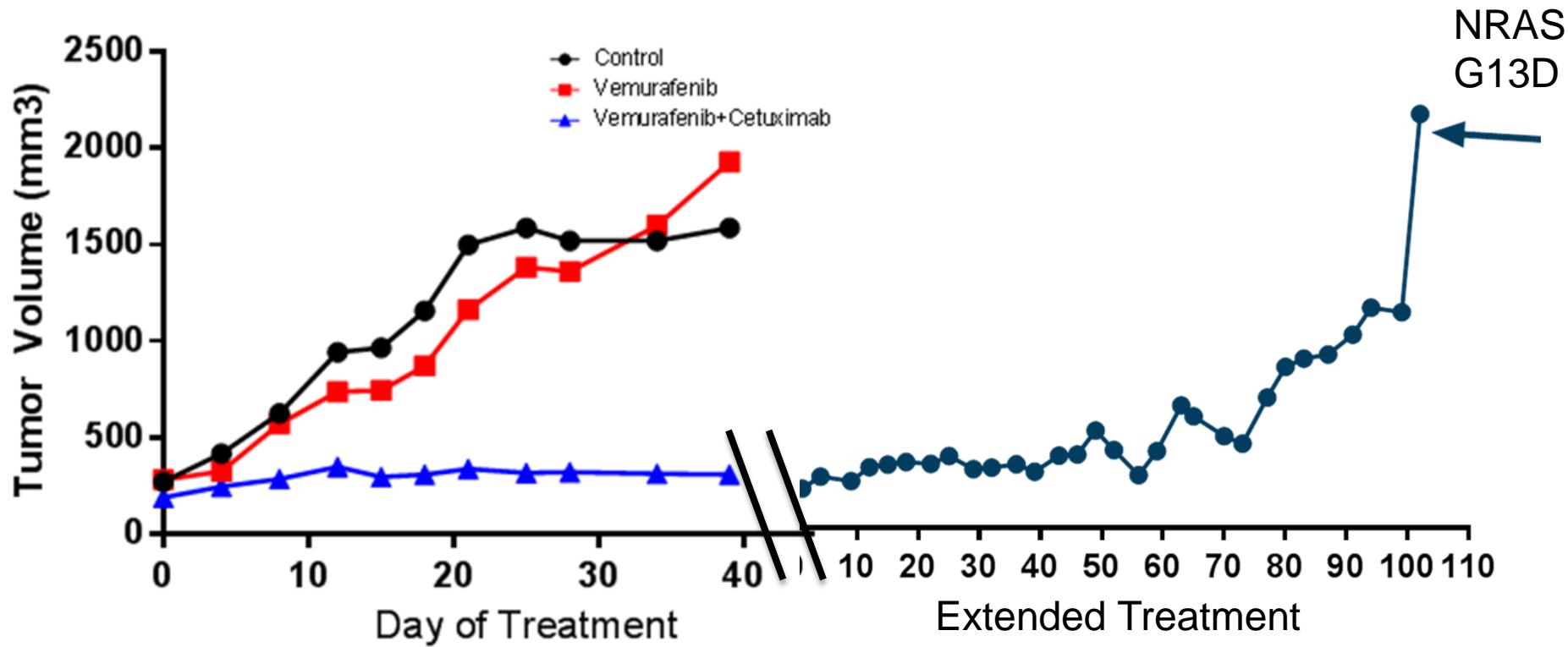


Embedded Patient-Derived Xenograft Co-Clinical Trial

- **On study biopsies to derive PDXs**, serial cfDNA
- **Co-clinical trial** to interrogate pharmacodynamics and mechanisms of resistance, and correlate outcomes

SWOG Hope, NIH R01 Funding, JAX Collaboration
MDACC, UCSF, USC, UCSD, U Colorado, Yale

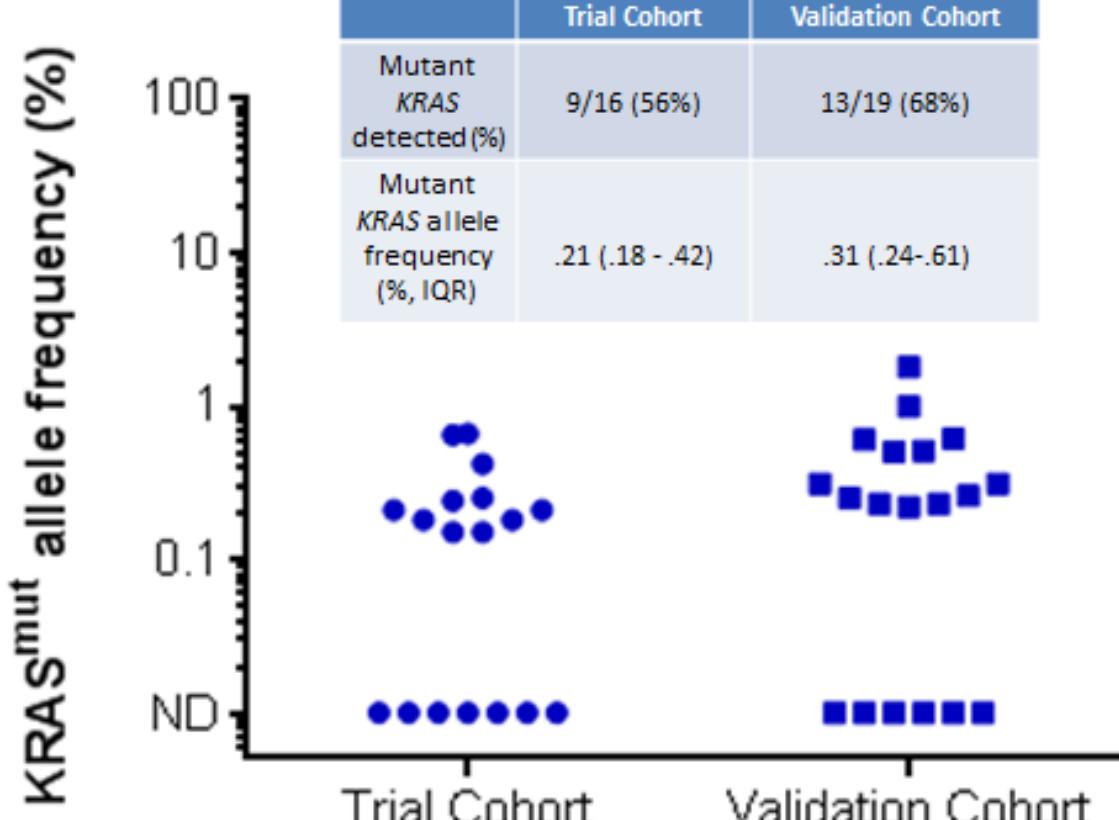
Resistance to vemurafenib + cetuximab is associated with MAPK pathway mutation



Classically, KRAS and BRAF mutations are mutually exclusive
So where are these rare clones coming from?

Kopetz, et al JCO '15

Rare KRAS^{mut} Cells Exist in BRAF^{mut} Colorectal Tumors



Jayesh Desai
Kopetz, et al JCO '15

Acquired RAS after BRAF+EGFR

- Baseline mutation analysis
 - Hotspots in AKT1, AKT2, BRAF, CDK, EGFR, ERBB2, FGFR1, FGFR3, FLT3, HRAS, JAK2, KIT, KRAS, MET, NRAS, PDGFRA, PIK3CA, RET
 - **BRAF V600E**
- Start study treatment:
 - cetuximab+ encorafenib
 - confirmed PR after 10 weeks
 - After 4 months of therapy: New progressive lesion
- Mutation analysis new lesion (same gene panel):
 - **BRAF V600E** and **KRAS G12R**

Post-progression cfDNA demonstrates clones that reactivate MAPK

Identified in cfDNA

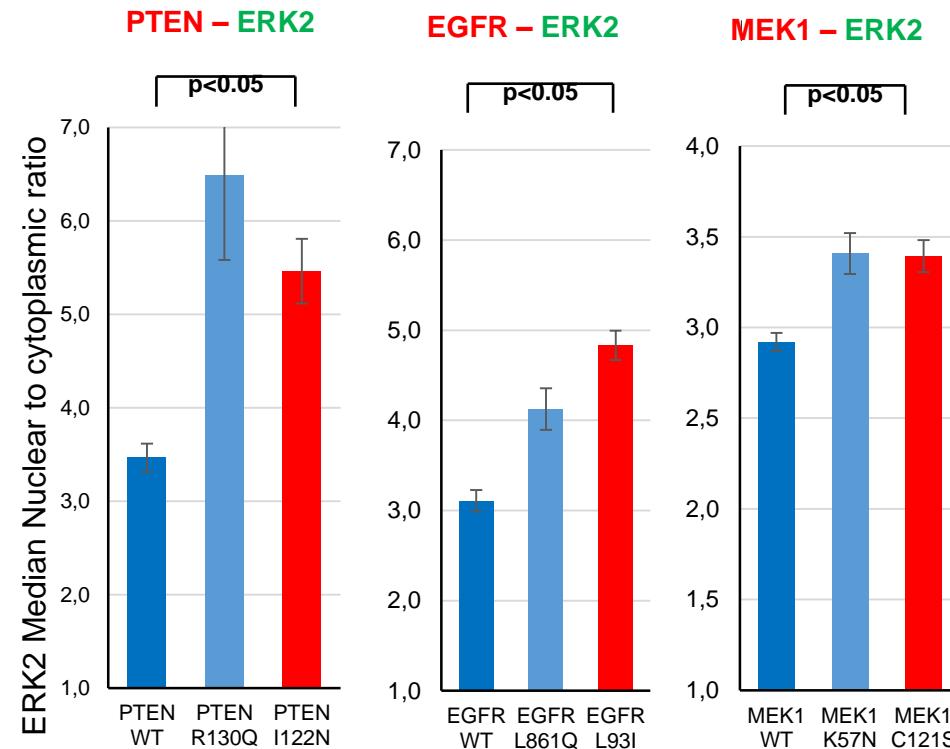
PTEN I122N

MEK1 C121S

GNAS R201C

EGFR L93I

ARAF S490T

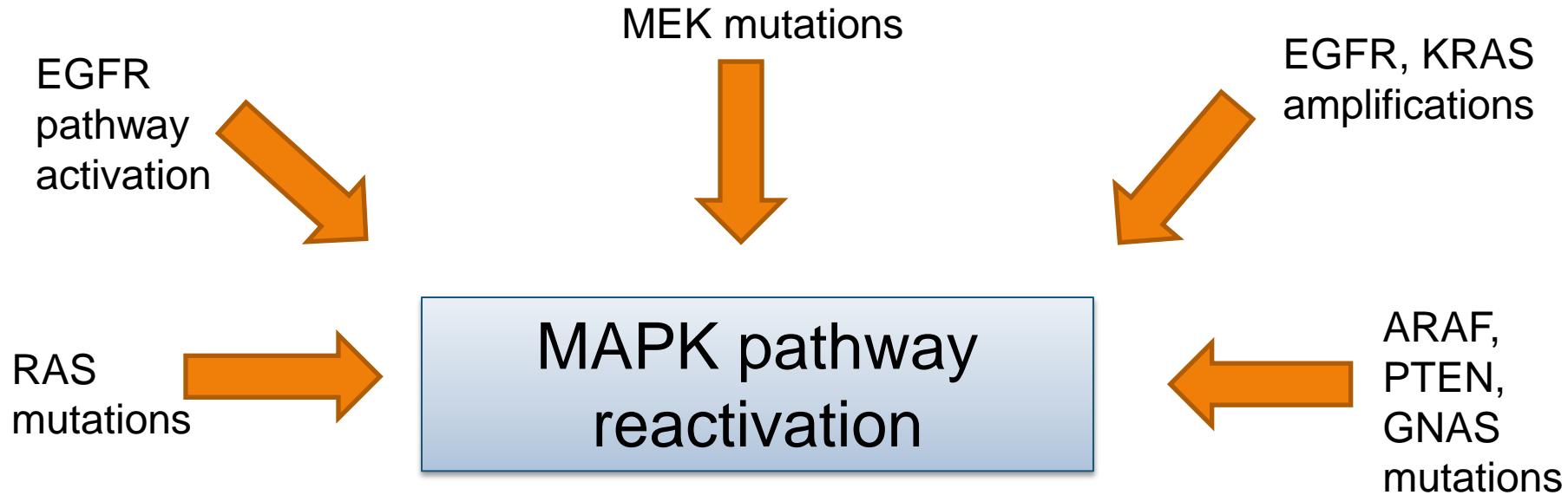


Gabi Tarnic, Novellus

If this is the case, then MEK inhibition may reverse resistance to EGFR/BRAF

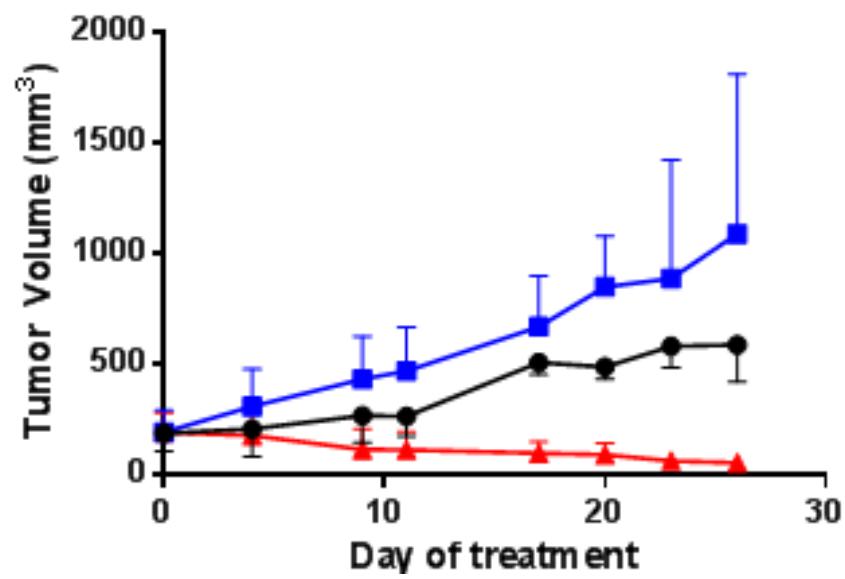
But may not substantially improve initial activity

Resistance to BRAF +/- EGFR

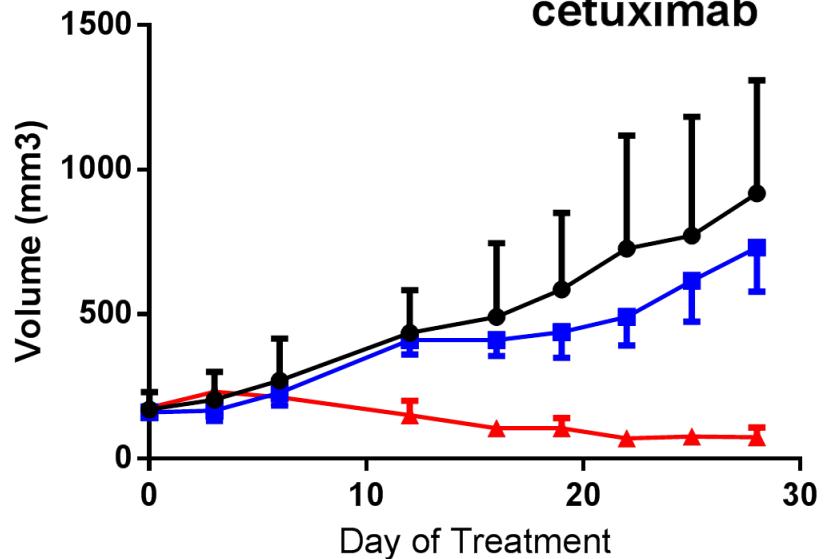


Another example of **convergent evolution** in colorectal cancer resistance to targeted therapy

PDX model: $BRAF^{V600E}/KRAS^{\text{mut}}$: $BRAF+EGFR$ vs $MEK+EGFR$ inhibition



$BRAF^{V600E}/KRAS^{G12R}$



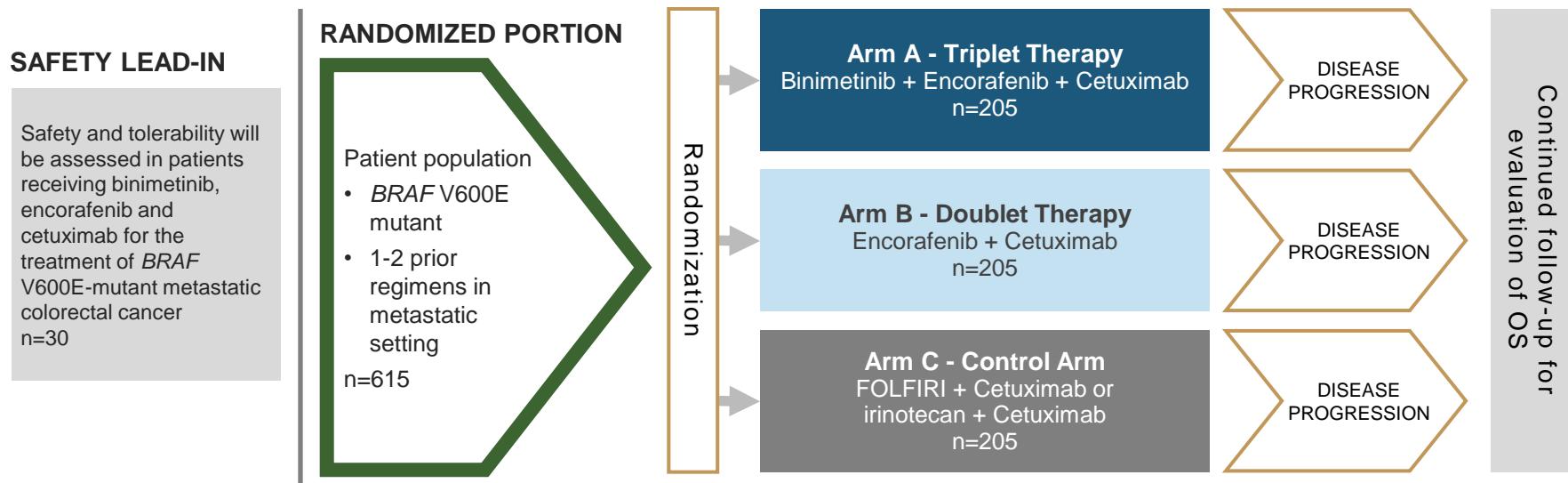
$BRAF^{V600E}/KRAS^{G12D}$

MEK inhibition reverses resistance to $BRAF/EGFR$

Van Morris

Encorafenib + Cetuximab ± Binimetinib

Phase 3 *BRAF^{mut}* Colorectal Cancer Study Design

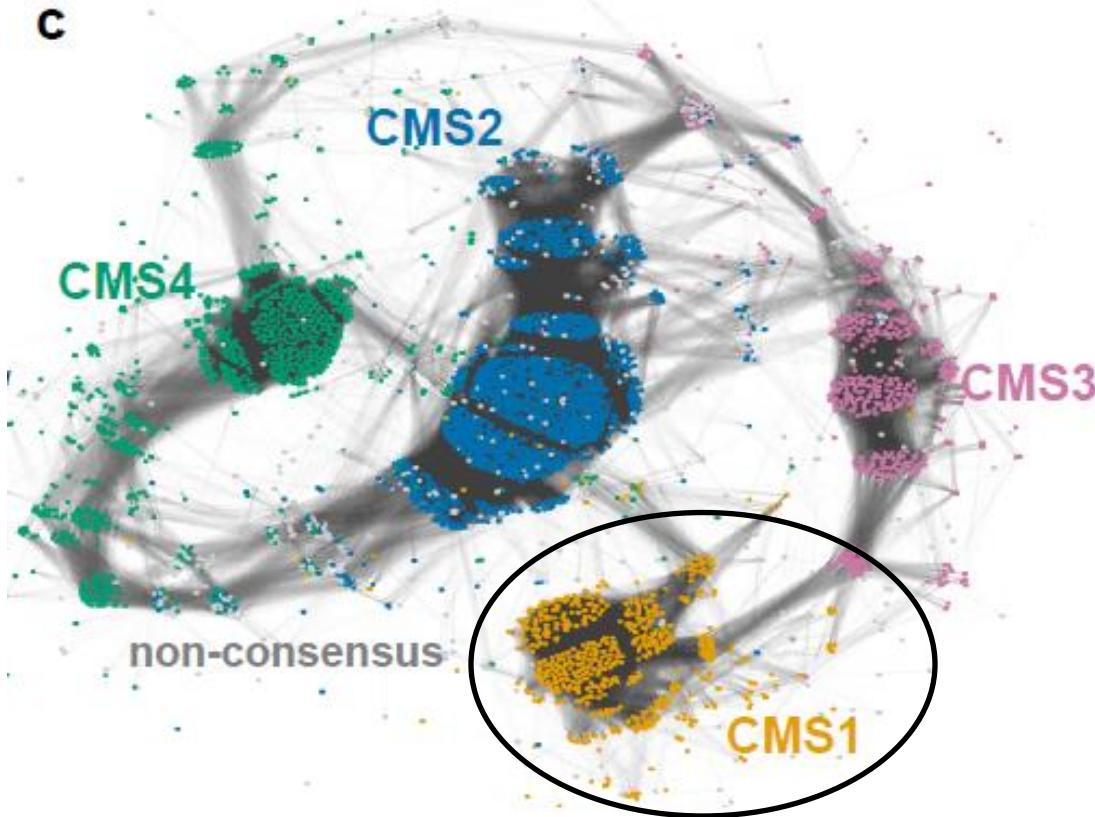


Primary Endpoint: Overall survival (OS) of the triplet therapy compared to the control arm.

Secondary Endpoints: Address efficacy of the doublet therapy compared to the control arm, and the triplet therapy compared to the doublet therapy.

Patient enrollment is expected to be completed in 2018.

Consensus Molecular Subtypes: CMS1



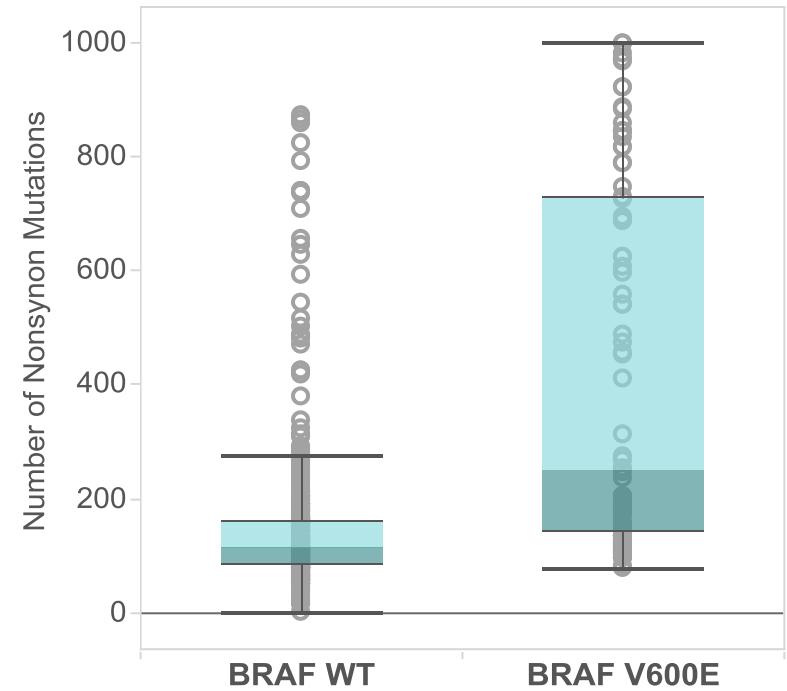
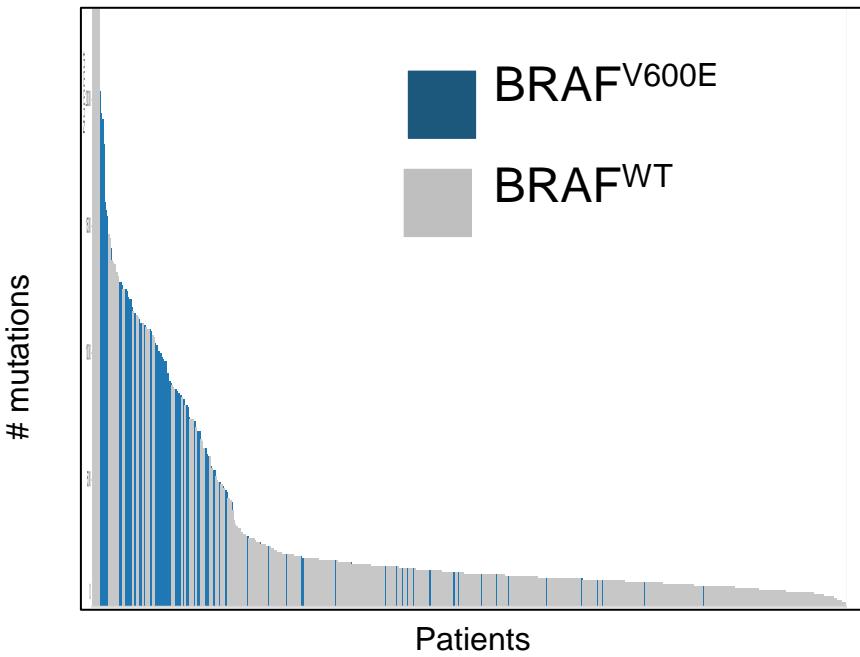
CMS1 MSI Immune

MSI, CIMP high
Hypermutation

BRAF mutations

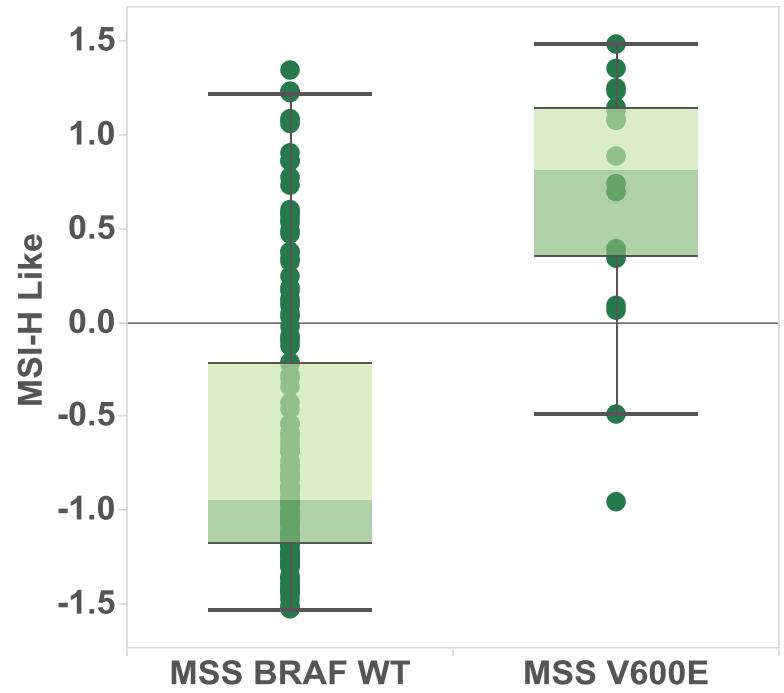
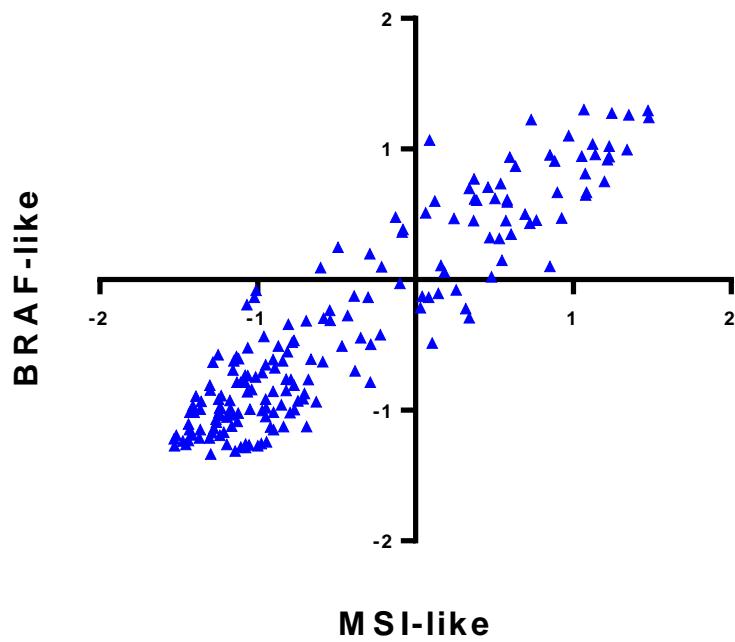
Immune infiltration
and activation

“BRAF^{mut} biology” is more than a mutated kinase



Sequencing data from NHS/HP Study (Giannakos, et al Cell Report '16)

BRAF and MSI share similar gene expression signatures



Role for immunotherapy in CMS1/BRAF mutated context?

Conclusions

- Testing for BRAF mutation is standard of care
 - Strong prognostic information that is useful for clinical management
- Patients with BRAF^{mut} CRC have distinct biology and limited benefit with standard therapy
- Patients should be enrolled in clinical trials when available:
 - BRAF+EGFR+irinotecan or BRAF+EGFR+MEK appear most promising
 - Immunotherapy studies may be particularly relevant for patients with BRAF^{mut} CRC (with or without MSI-H)

Acknowledgements

Kopetz Lab

- Van Morris, MD
- Feng Tian, PhD
- Camilla Jiang, MD, PhD
- Mike Lee, MD
- Jian Song, DVM
- Riham Katkhuda, MD
- Chris Lieu, MD
- Ali Kazmi, MD
- Pia Morelli, MD, PhD
- Michael Overman, MD

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Acknowledgements

Kopetz Lab

- Van Morris, MD
- Feng Tian, PhD



- Michael Overman, MD

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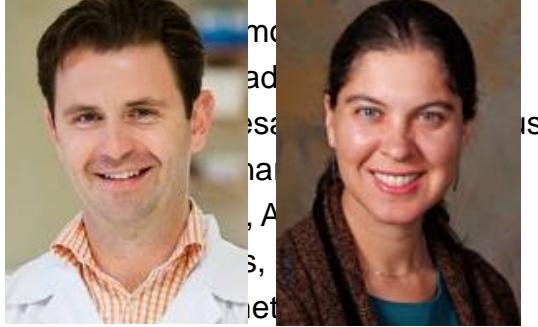
- Yvonne Lassere
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