

# Colorectal Cancer: Novel Insights

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## Colorectal Cancer: Novel Insights

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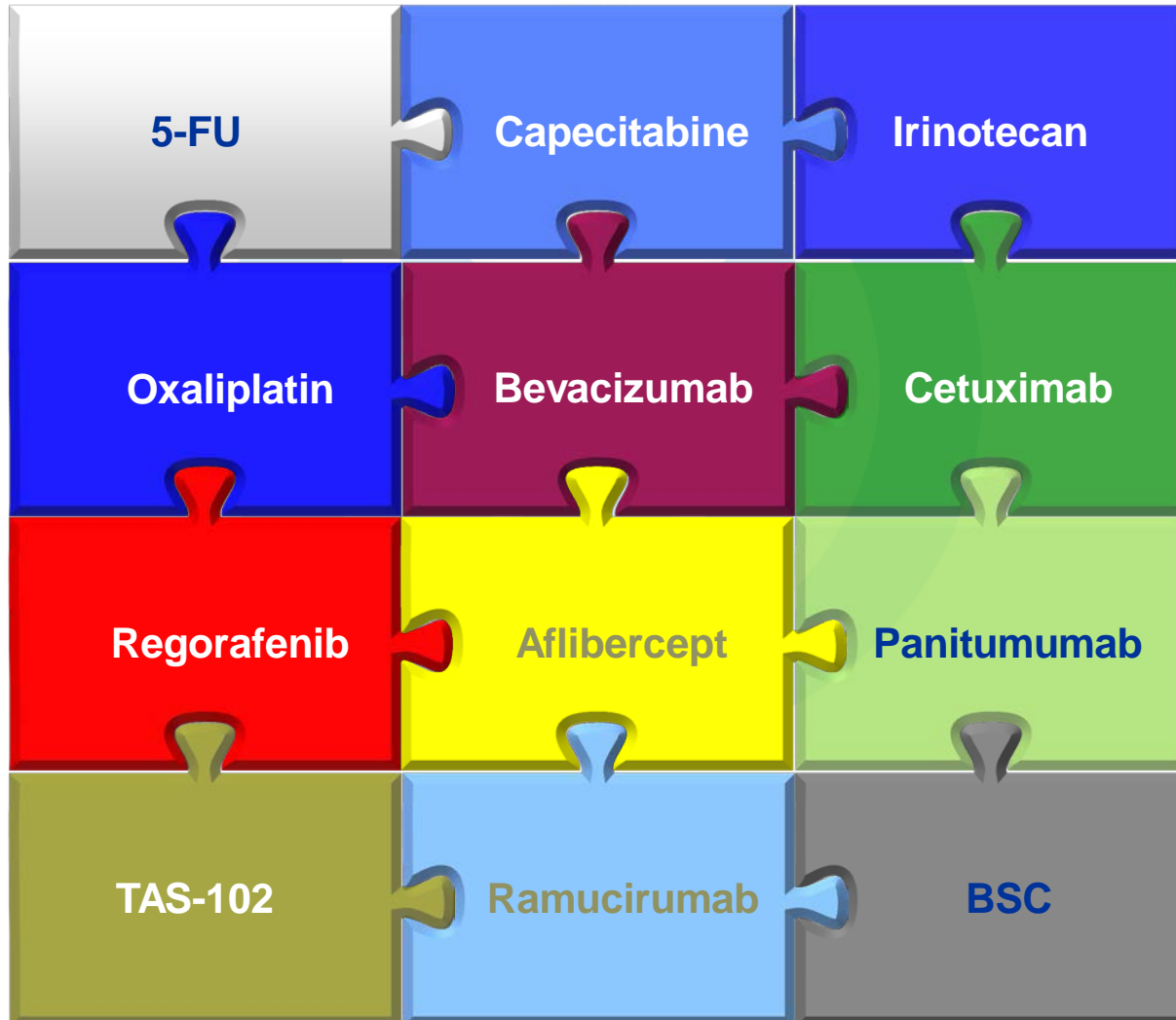
# Learning Objectives

- Know the current trend for treatment of advanced colorectal cancer
- Know the current status of personalized medicine in advanced CRC
- Know the differences between left vs right sided colon cancer

# Impact of CRC

- **CRC US statistics:**
  - 3<sup>rd</sup> highest incidence rate (~135,000/yr)
  - 2<sup>nd</sup> highest mortality rate (~49,000/yr)
- **CRC Global statistics:**
  - 3<sup>rd</sup> highest incidence rate ( ~ 1.2million/yr)
  - 4<sup>th</sup> highest mortality rate (~608,000/yr)
- ***The burden of disease is clearly evident...***

## A High Number of Agents Is Currently Available for the Treatment of mCRC



## Landscape in mCRC

- Bevacizumab and EGFR mAbs competing for first-line patients in KRAS wt CRC
- Bevacizumab, *ramucirumab* and Aflibercept competing for second-line patients with each other, and with EGFR mAbs in KRAS wt CRC
- Best sequence of therapies (VEGFi vs EGFRi) still to be established
- Regorafenib and TAS 102 as salvage therapy option
- Immunotherapy for pts with MMR deficient/MSI-H tumor

# Tools for Treatment Selection

Age  
PS  
Comorbidities  
Tumor burden  
Potential for cure?  
Symptoms?  
Tumor location

**Clinical Markers** **Molecular Markers**

Histologic grade  
CEA  
KRAS  
NRAS  
BRAF  
MSI/MMR

***Patient characteristics***

**+**

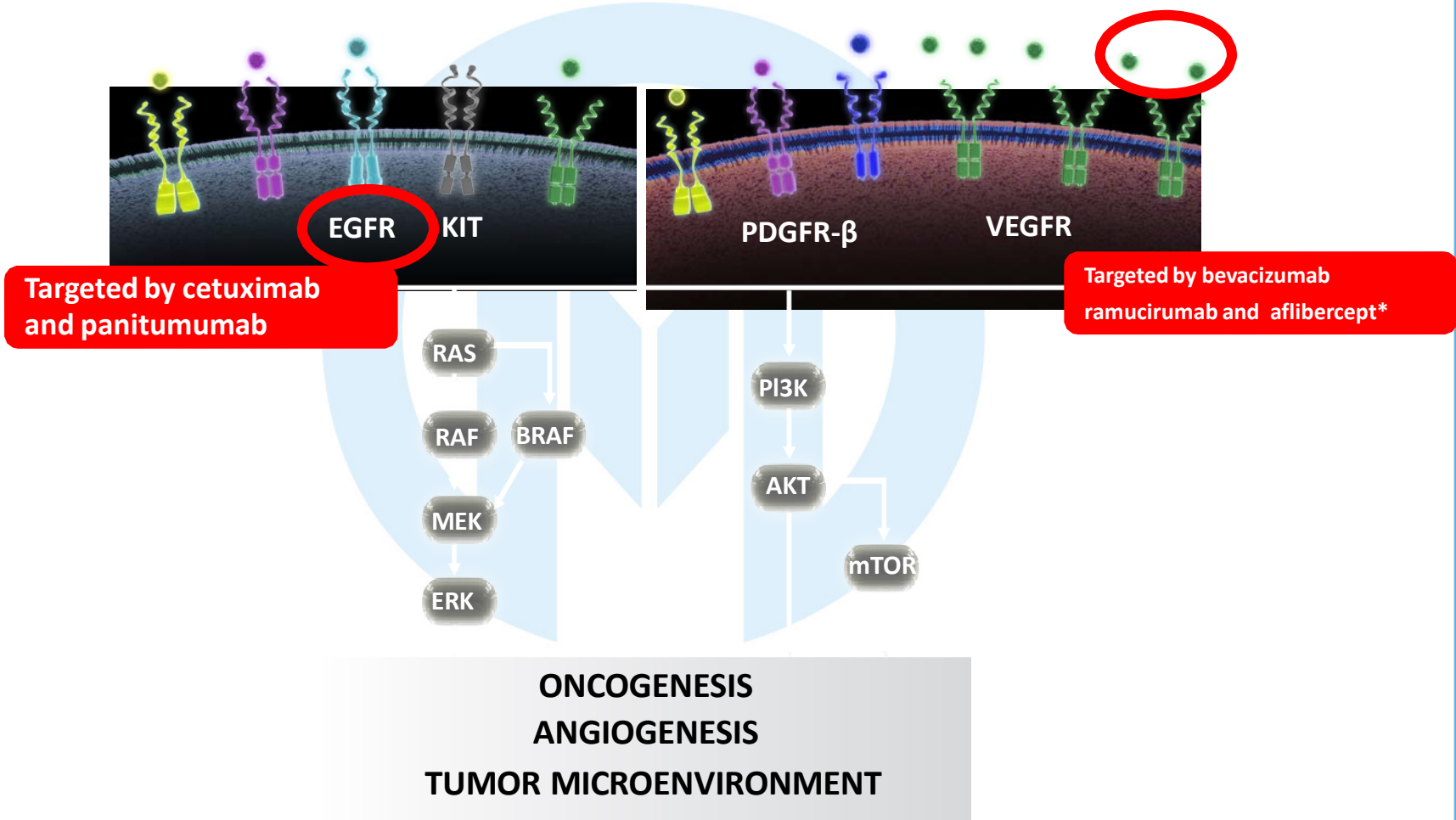
***Tumor characteristics***

# Questions

- How can biologics be used to their full potential?
  - Prognostic markers -Biomarker that correlates with clinical outcome ***regardless of therapy***
  - Predictive markers-Biomarker that is associated with the likelihood of ***response to therapy***
- Can a patient population be identified which would benefit most from one specific treatment strategy?



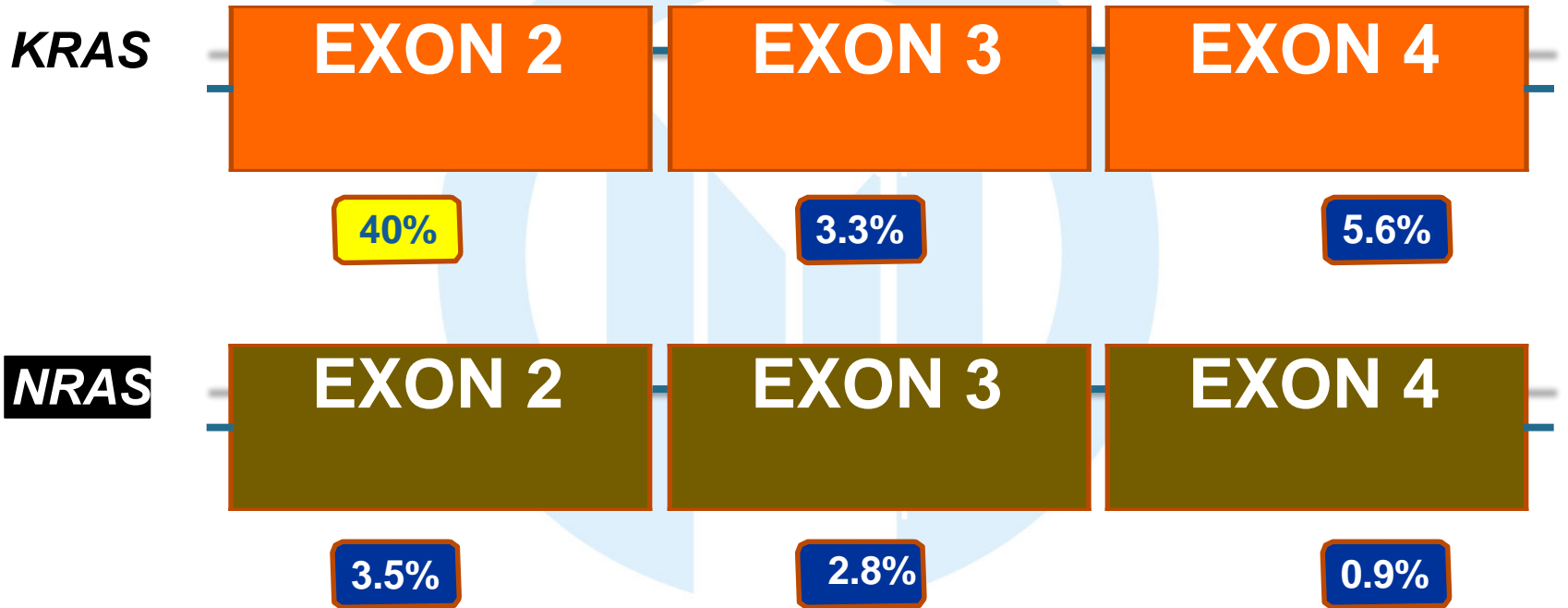
# Overview of EGFR and VEGFR Growth Signaling Pathways



# Biomarkers for anti- VEGF Drugs



# Spectrum of ras mutations



# Updated Analysis of PRIME study

## RAS and BRAF Status

	Panitumumab + FOLFOX4	FOLFOX4 Alone	Total
<b>KRAS exon 2 (codons 12/13)</b>			
Wild-type	325	331	656
Mutant	221	219	440
Wild-type KRAS exon 2 tumors tested for RAS and BRAF	N = 320	N = 321	N = 641
Wild-type KRAS exon 2/mutant other RAS – n (%)	51 (16)	57 (18)	108 (17)
<b>KRAS exon 3 (codon 61) – n (%)</b>			
Wild-type	308 (96)	306 (95)	614 (96)
Mutant	10 (3)	14 (4)	24 (4)
Failure	2 (1)	1 (0)	3 (0)
<b>KRAS exon 4 (codons 117/146) – n (%)</b>			
Wild-type	288 (90)	296 (92)	584 (91)
Mutant	21 (7)	15 (5)	36 (6)
Failure	11 (3)	10 (3)	21 (3)
<b>NRAS exon 2 (codons 12/13) – n (%)</b>			
Wild-type	308 (96)	307 (96)	615 (96)
Mutant	8 (3)	14 (4)	22 (3)
Failure	4 (1)	0 (0)	4 (1)
<b>NRAS exon 3 (codon 61) – n (%)</b>			
Wild-type	305 (95)	305 (95)	610 (95)
Mutant	12 (4)	14 (4)	26 (4)
Failure	3 (1)	2 (1)	5 (1)
<b>NRAS exon 4 (codons 117/146) – n (%)</b>			
Wild-type	316 (99)	313 (98)	629 (98)
Mutant	0 (0)	0 (0)	0 (0)
Failure	4 (1)	8 (2)	12 (2)
<b>BRAF exon 15 (codon 600) – n (%)</b>			
Wild-type	286 (89)	280 (87)	566 (88)
Mutant	24 (8)	29 (9)	53 (8)
Failure	10 (3)	12 (4)	22 (3)

**KRAS exon 2  
codon 12/13**

**40%**

**KRAS exon 3  
codon 61**

**4%**

**KRAS exon 4  
codon 117/146**

**6%**

**NRAS exon 2  
codon 12/13**

**3%**

**NRAS exon 3  
codon 61**

**4%**

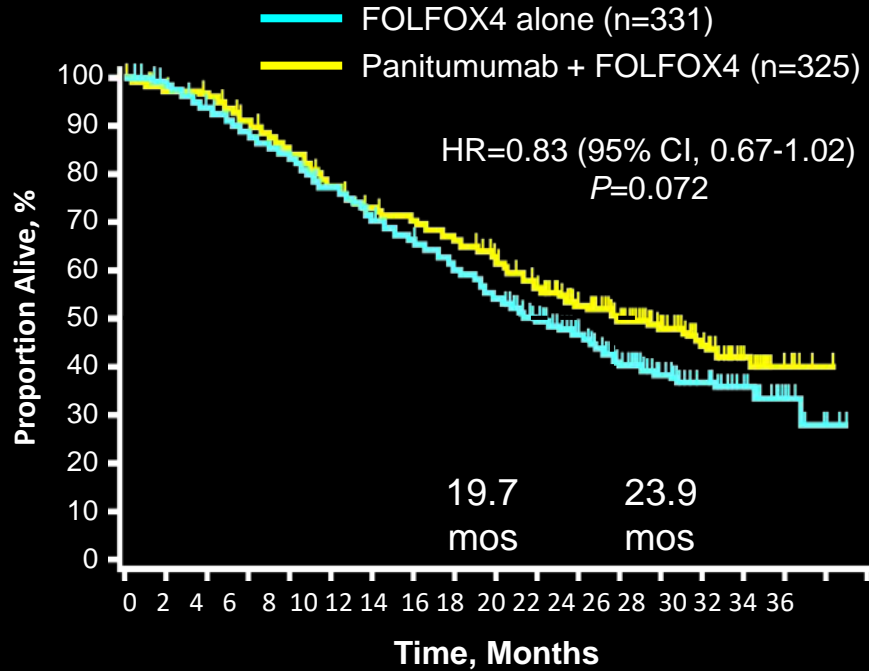
**BRAF exon 15  
codon 600**

**8%**

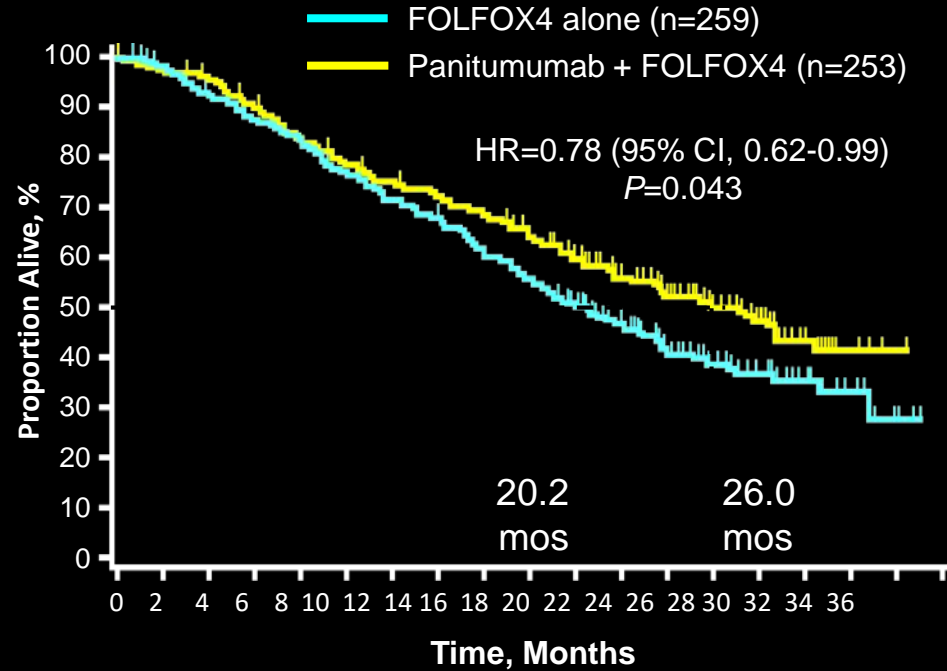
**17%**

# PRIME Biomarker Analysis: OS in Patients With WT *KRAS* Exon 2 and WT *RAS* mCRC

**WT *KRAS* Exon 2**

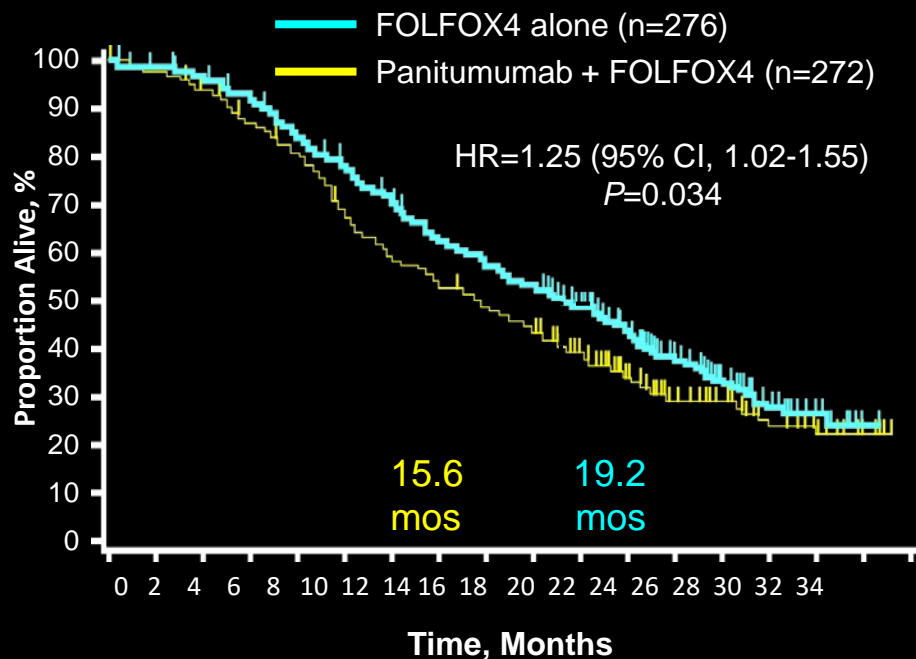


**WT *KRAS*/*NRAS***

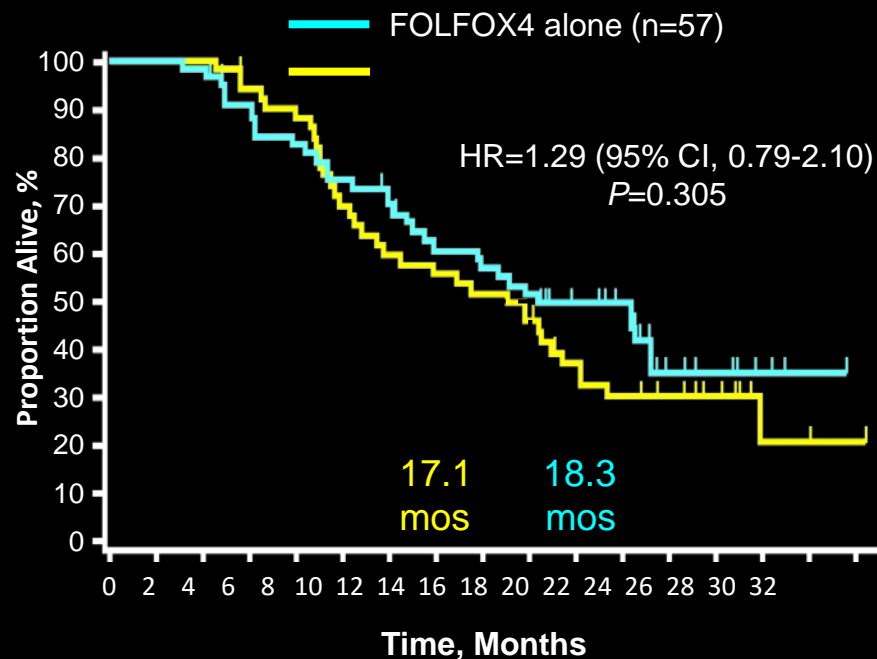


# PRIME Biomarker Analysis: OS in Patients With MT *RAS* or WT *KRAS* Exon 2/MT Other *RAS*

## MT *RAS*



## WT *KRAS* Exon 2/MT Other *RAS*

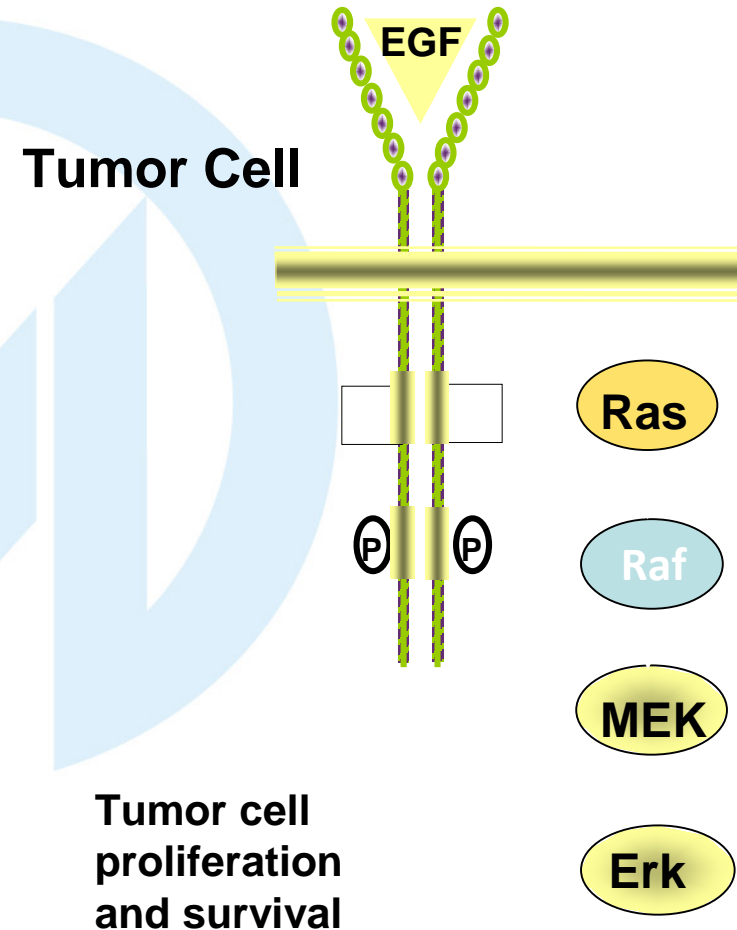


## Take home point

- testing for these additional codons could help screen 20% more patients with mCRC for treatment with EGFR inhibitors
- This method could help to more accurately select patients who will benefit from EGFR inhibitors.
- NCCN now recommends testing for all RAS mutations

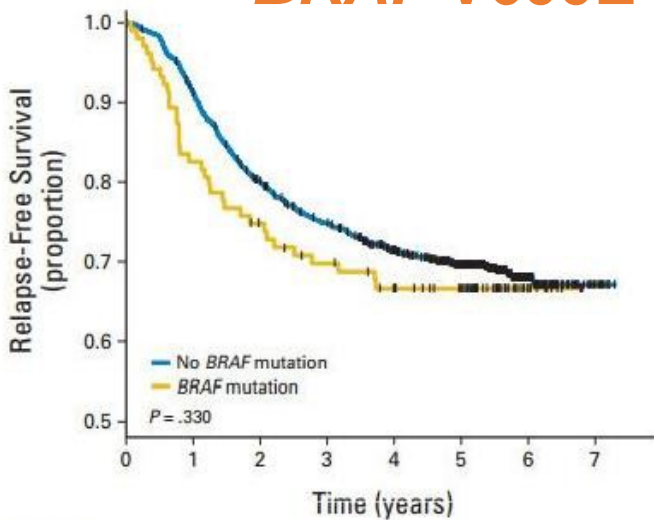
# BRAF Mutations in CRC

- BRAF is primary effector of KRAS signaling
- BRAF mutations:
  - Occur most frequently in exon 15 (V600E)
  - Found in 4%-14% of patients with CRC
  - Mutually exclusive with KRAS mutations

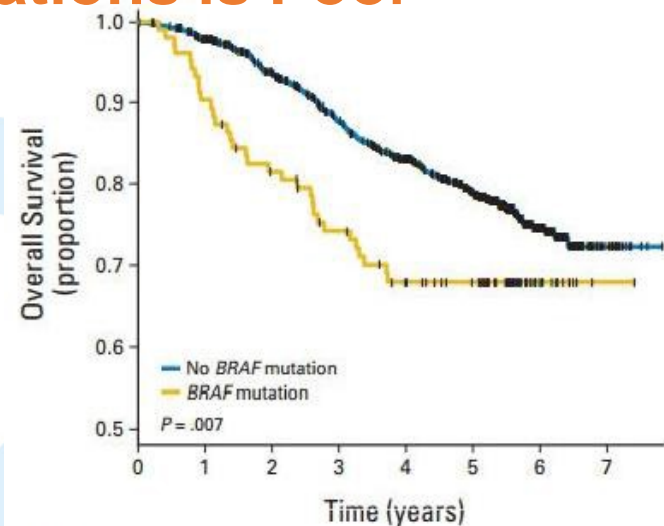




# The Prognosis of Patients With BRAF V600E Mutations is Poor



No. at risk	0	1	2	3	4	5	6	7
No BRAF	1,204	1,091	940	862	786	660	99	9
BRAF	103	85	75	68	61	53	11	0



No. at risk	0	1	2	3	4	5	6	7
No BRAF	1,204	1,152	1,048	937	842	697	160	18
BRAF	103	93	81	71	61	54	14	1

**Table 1. Association of the Mutation Status of the BRAF Oncogene with Progression-free Survival, Overall Survival, and Response Rate.\***

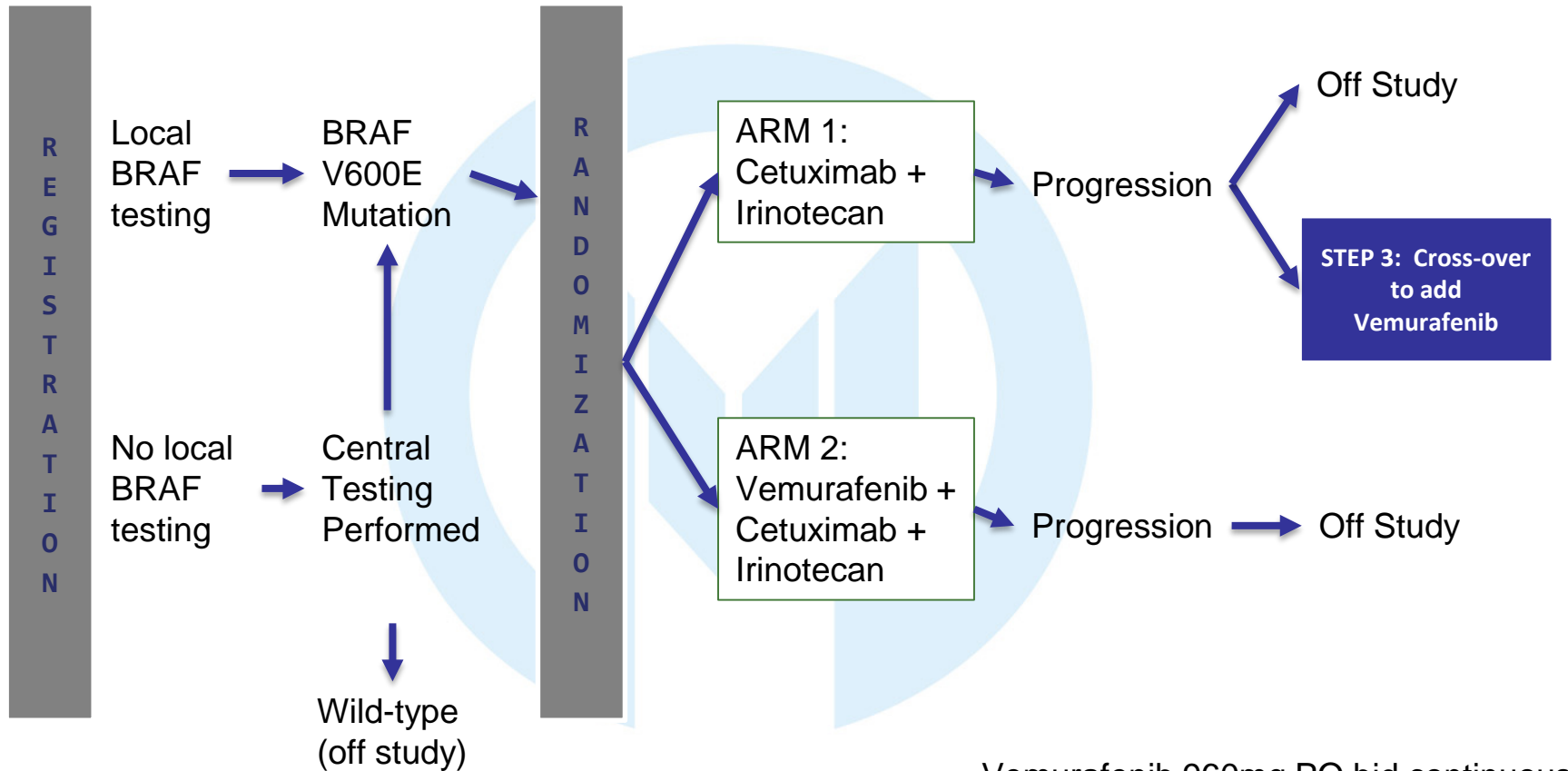
Variable	Wild-Type BRAF	Mutated BRAF	P Value
<b>No. of patients</b>			
CB group	243	17	
CBC group	231	28	
<b>Median progression-free survival (mo)</b>			
CB group	12.2	5.9	0.003
CBC group	10.4	6.6	0.010
<b>Median overall survival (mo)</b>			
CB group	24.6	15.0	0.002
CBC group	21.5	15.2	0.001
<b>Response rate (%)</b>			
CB group	50	35	0.32
CBC group	48	39	0.43

# Randomized trial of irinotecan and cetuximab with or without vemurafenib in *BRAF*-mutant metastatic colorectal cancer (SWOG S1406)

Scott Kopetz,<sup>1</sup> Shannon McDonough,<sup>2</sup> Heinz-Josef Lenz,<sup>3</sup> Anthony Magliocco,<sup>4</sup> Chloe Atreya,<sup>5</sup> Luis A. Diaz Jr.,<sup>6</sup> Carmen Allegra,<sup>7</sup> Kanwal Raghav,<sup>1</sup> Van Morris,<sup>1</sup> Stephen Wang,<sup>8</sup> Christopher Lieu,<sup>9</sup> Katherine A. Guthrie,<sup>2</sup> Howard S. Hochster<sup>10</sup>

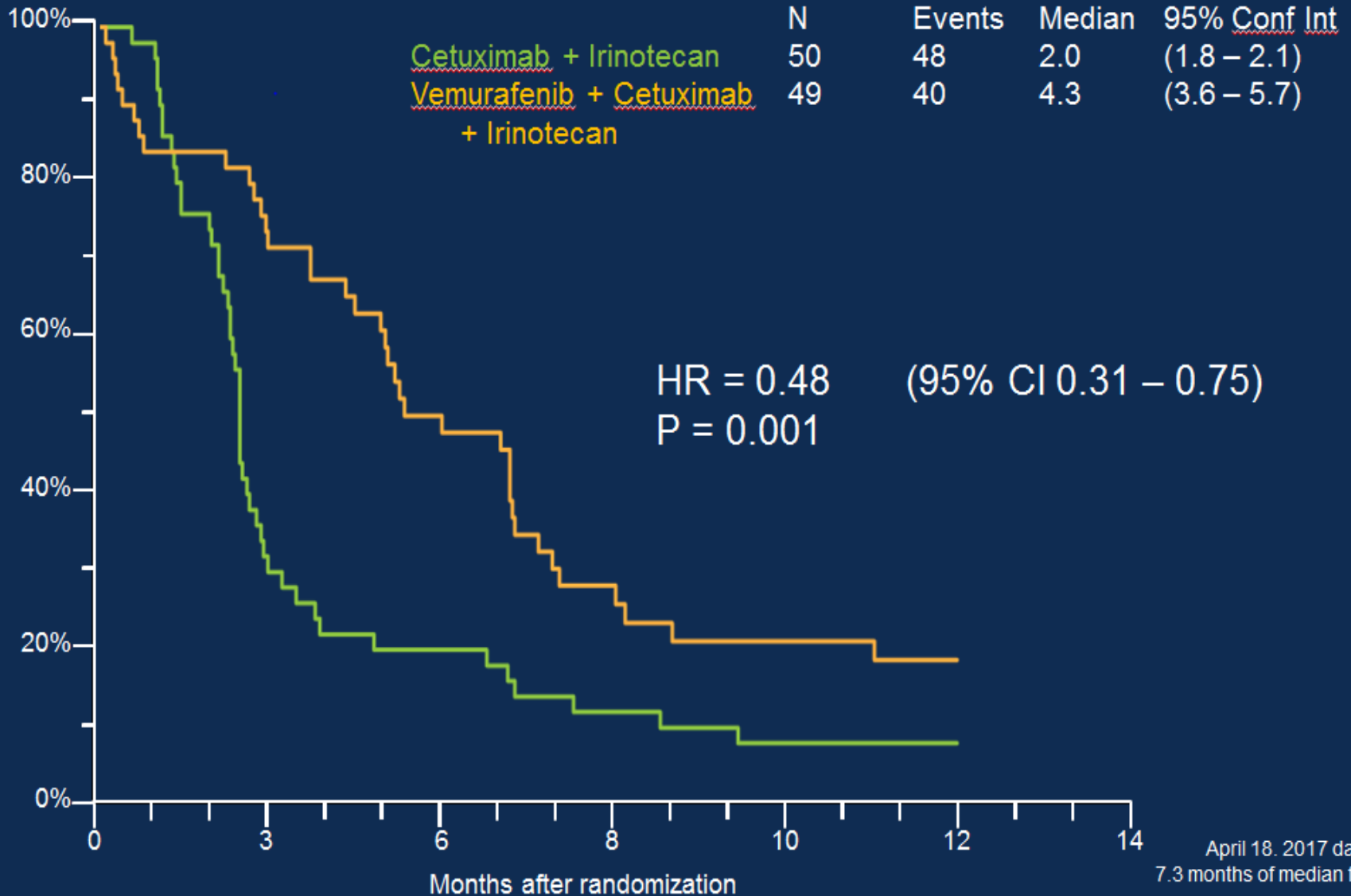
<sup>1</sup>The University of Texas MD Anderson Cancer Center, Houston, TX; <sup>2</sup>Fred Hutchinson Cancer Research Center, Seattle, WA; <sup>3</sup>USC Norris Comprehensive Cancer Center, Los Angeles, CA; <sup>4</sup>H. Lee Moffitt Cancer Center & Research Institute, Tampa, FL; <sup>5</sup>University of California, San Francisco, San Francisco, CA; <sup>6</sup>Memorial Sloan Kettering Cancer Center, New York, NY; <sup>7</sup>University of Florida, Gainesville, FL; <sup>8</sup>Kaiser Permanente, Sacramento, CA; <sup>9</sup>University of Colorado School of Medicine, Anschutz Medical Campus, Aurora, CO; <sup>10</sup>Yale Cancer Center, New Haven, CT

# Study Design



Vemurafenib 960mg PO bid continuous  
 Cetuximab 500mg/m<sup>2</sup> IV q2weeks  
 Irinotecan 180mg/m<sup>2</sup> IV q2weeks

# Primary Endpoint: Progression-free survival



# Response Rate

	<u>Cetuximab + Irinotecan</u> (n=47) <sup>a</sup>	<u>Vemurafenib + Cetuximab + Irinotecan</u> (n=44) <sup>a</sup>	<u>P-value</u> <sup>c</sup>
<u>Partial response</u> <sup>b</sup>	4%	16%	P=0.001
<u>Stable disease</u>	17%	50%	
<u>Progression</u> <sup>c</sup>	66%	18%	

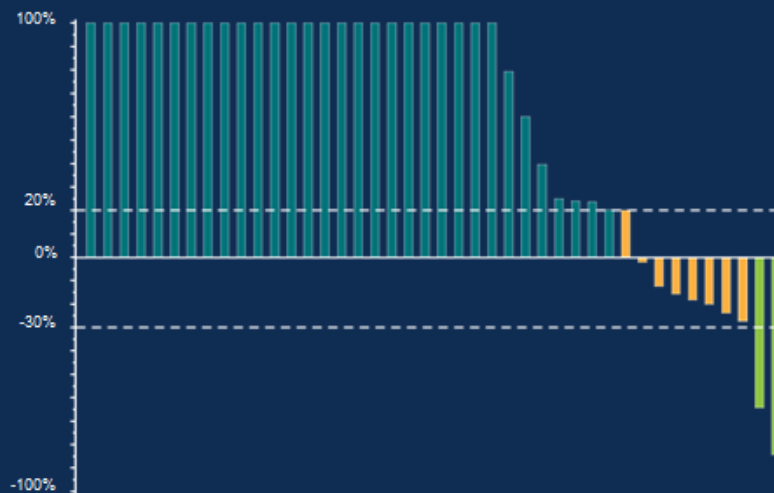
**Disease Control Rate**

22%

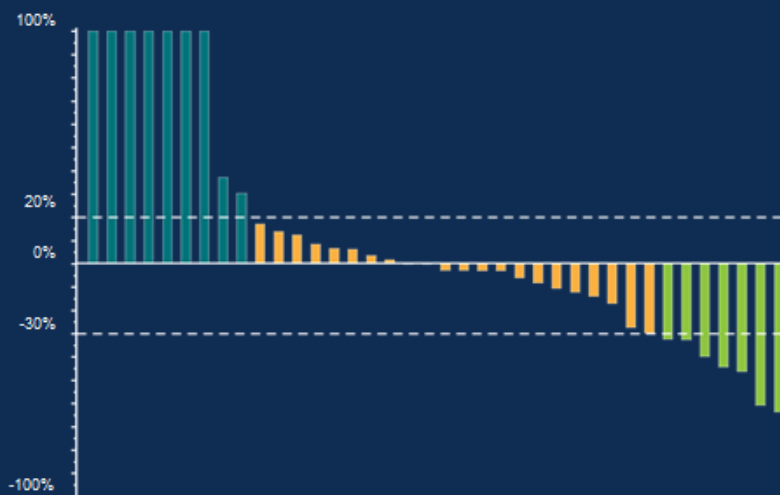
67%

<sup>a</sup>93 patients had measurable disease; <sup>b</sup>Confirmed and unconfirmed; PR for patients previously treated with irinotecan was 0% and 18%, respectively; <sup>c</sup>Including symptomatic deterioration; <sup>c</sup> Chi-squared

Cetuximab + Irinotecan

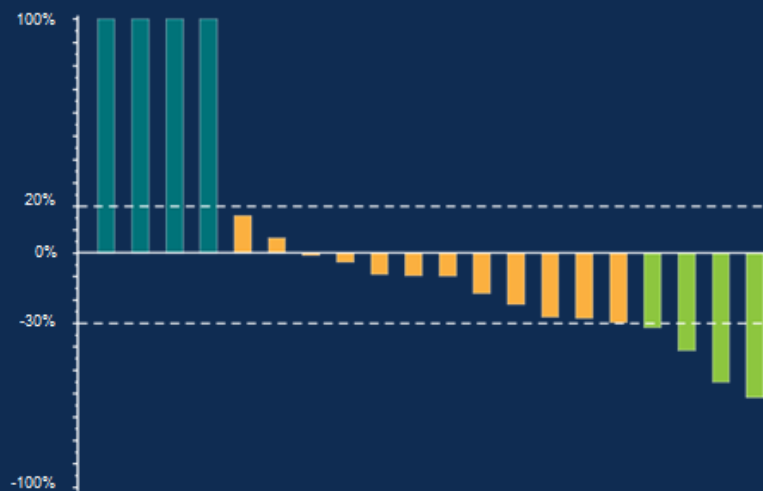
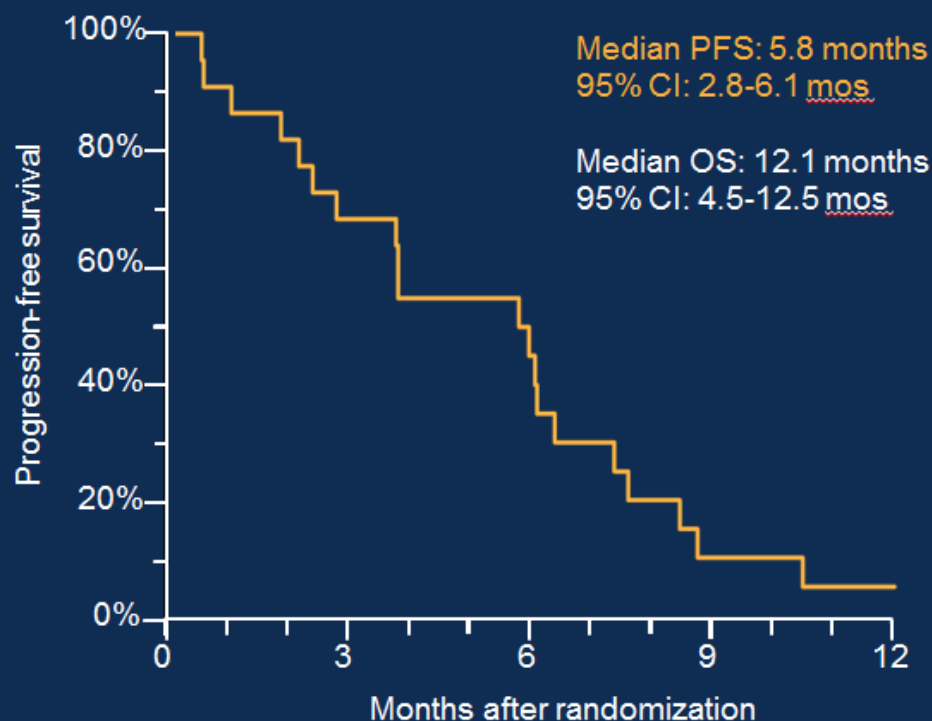


Vemurafenib + Cetuximab + Irinotecan



# Crossover to VIC after progression

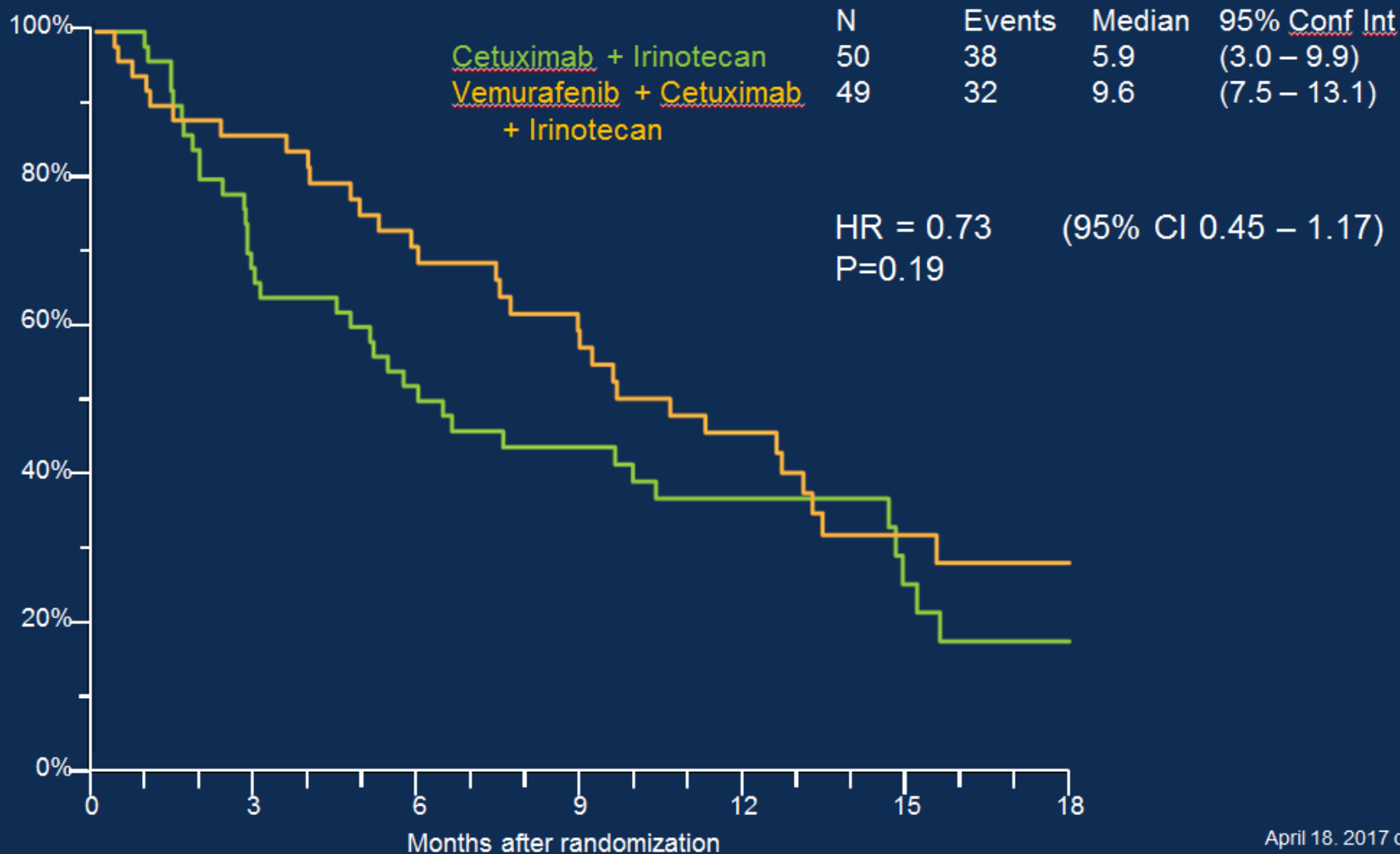
- Patients with radiographically documented progression on IC crossed over to receive VIC
- 48% of patients treated on IC arm crossed over



	<b>Crossover (n=24)<sup>a</sup></b>
Partial response	17%
Stable disease	55%
<b>Disease control rate</b>	<b>72%</b>

<sup>a</sup>2 patients did not progress prior to crossover; 4 did not have measurable disease; these patients are excluded from response rates

# Secondary Endpoint: Overall Survival



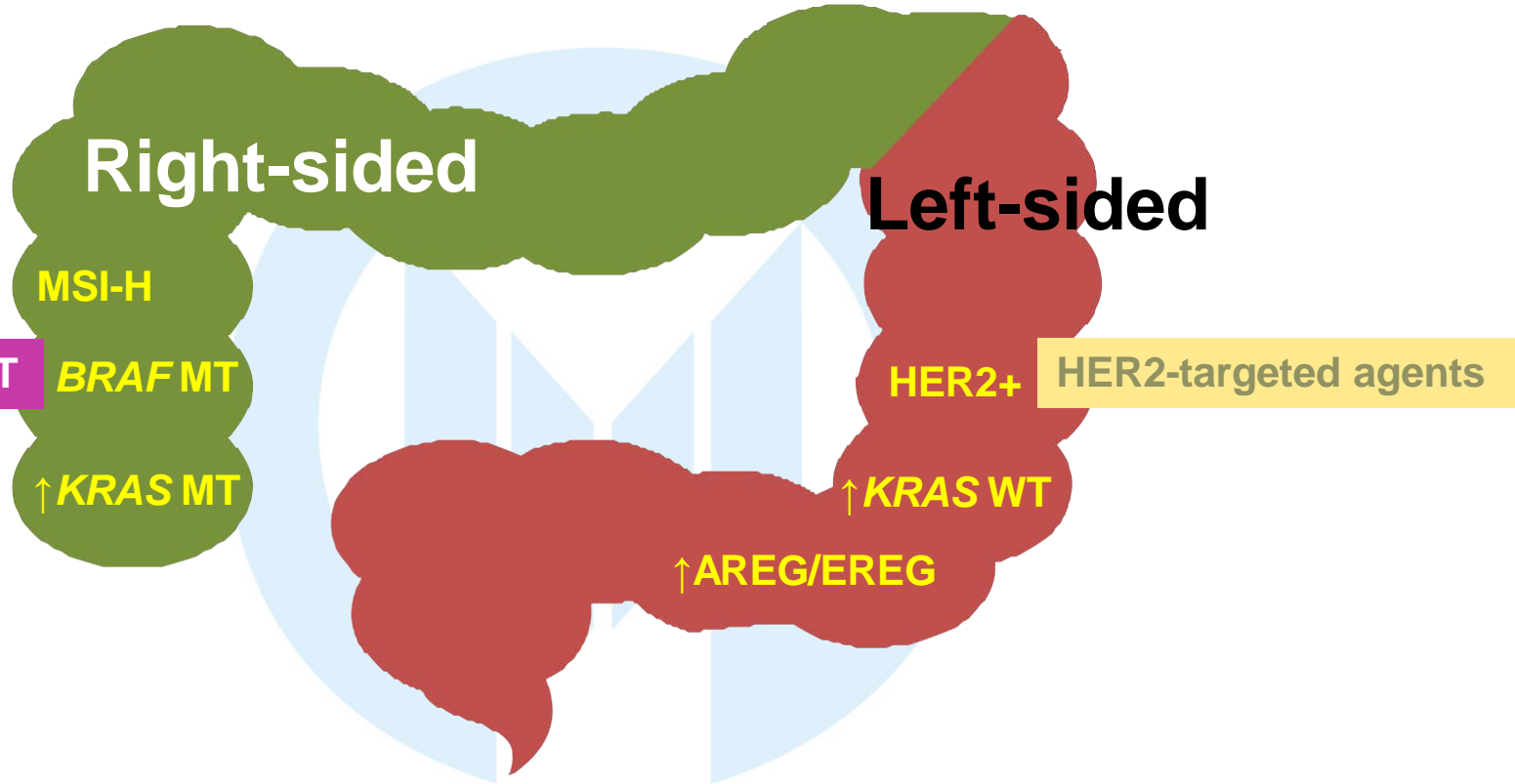
# Conclusions

- The combination of vemurafenib, cetuximab, and irinotecan (VIC) met its primary endpoint demonstrating improved progression-free survival in patients with BRAF<sup>V600E</sup> CRC
- Addition of Vemurafenib to IC showed activity even after progression on IC
- VIC represents a new treatment for metastatic BRAF<sup>V600E</sup> colorectal cancer



**What is the impact of site of primary tumour?**

# Primary Tumor Location and Potential Treatments



: *KRAS* wt

1<sup>ST</sup> LINE  
MET / ADVANCED  
COLORECTAL

*KRAS* wt  
Codons 12 & 13



**FOLFIRI**  
or  
**FOLFOX**  
  
MD choice

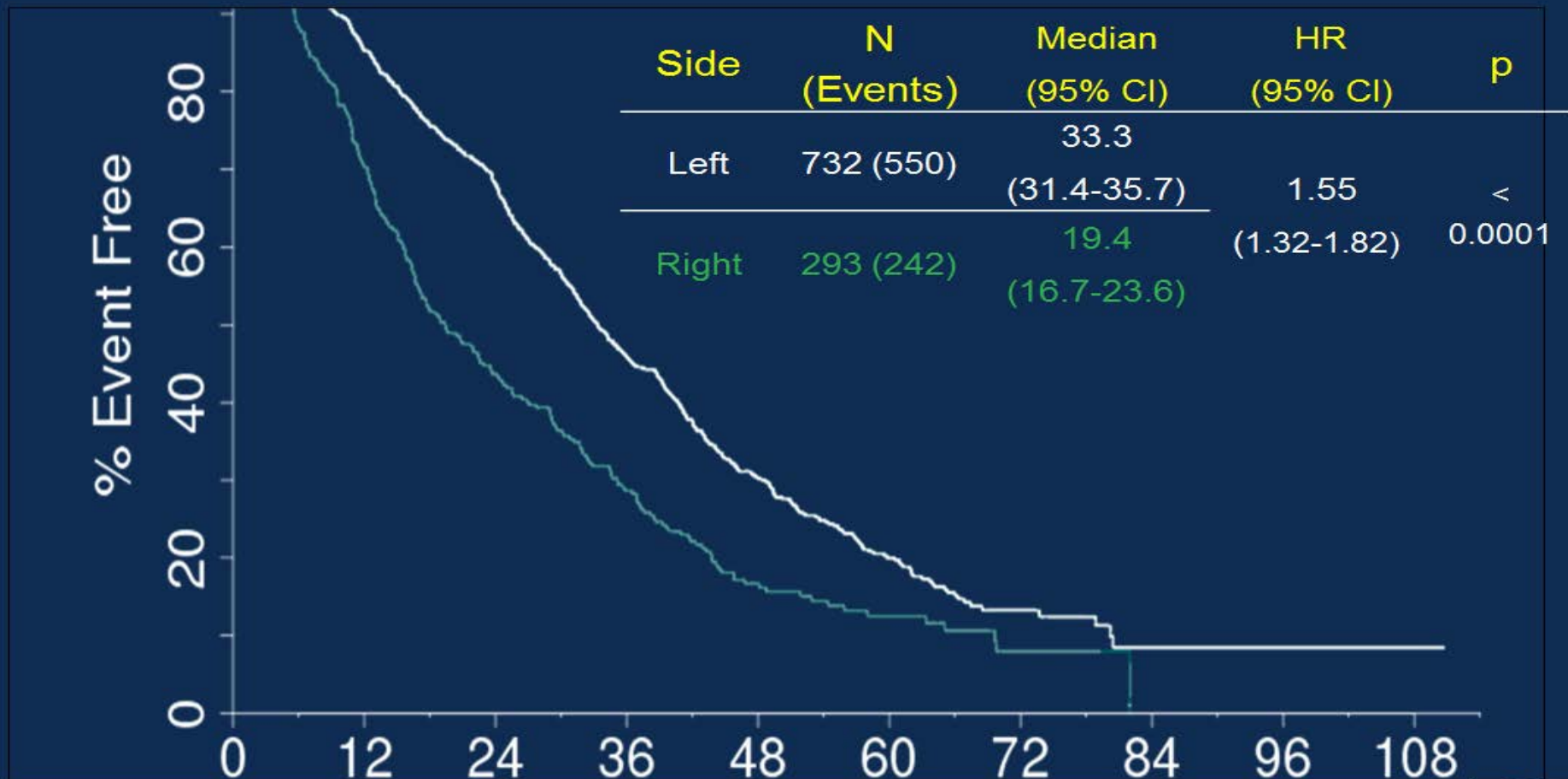
**N = 1137**

: NO DIFFERENCE

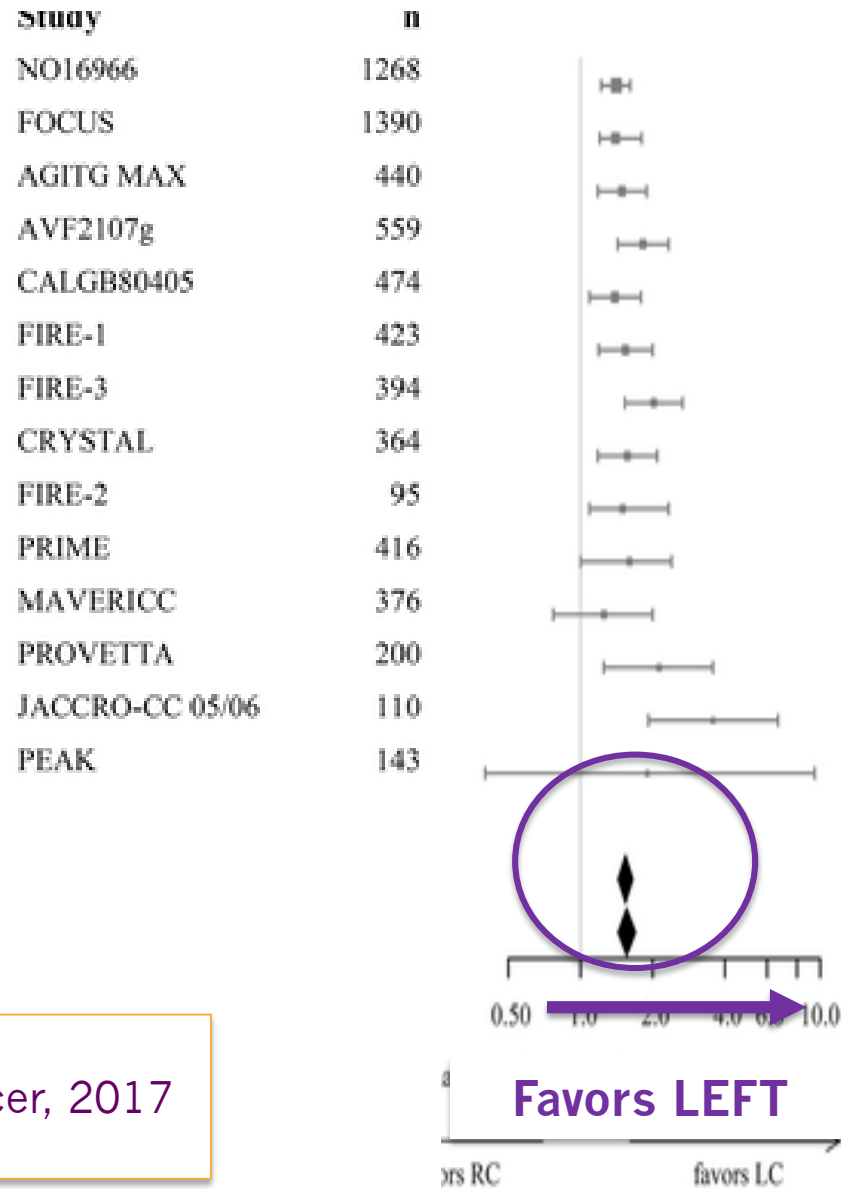
**Chemo + Cetuximab**  
OS = 29.9 mos  
PFS = 10.4 mos

**Chemo + Bevacizumab**  
OS = 29.0 mos  
PFS = 10.8 mos

## 80405: (KRAS WT) Overall Survival by Sidedness



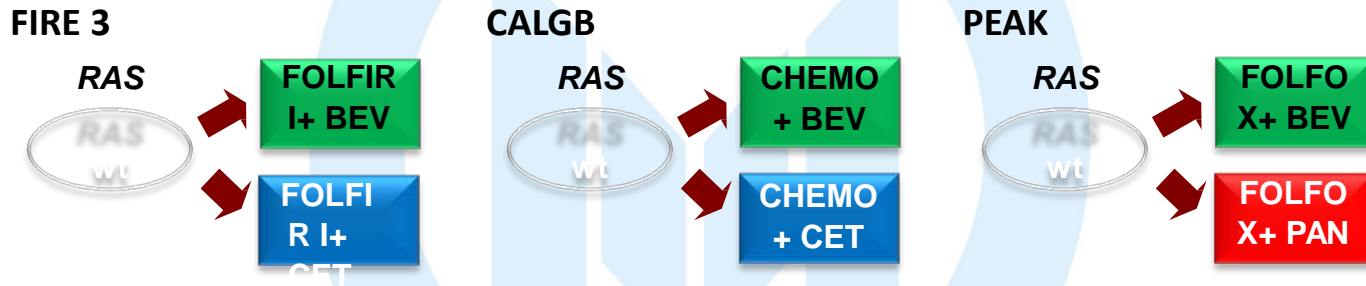
# Meta-analysis: Sidedness in MCRC



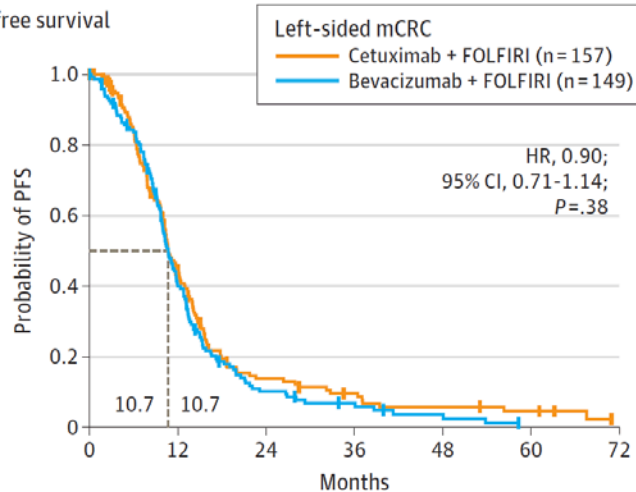
Holch et al, Eur J Cancer, 2017

# Predictive impact – bev versus anti-EGFRs

## Chemo+anti-EGFR vs Chemo+Bev



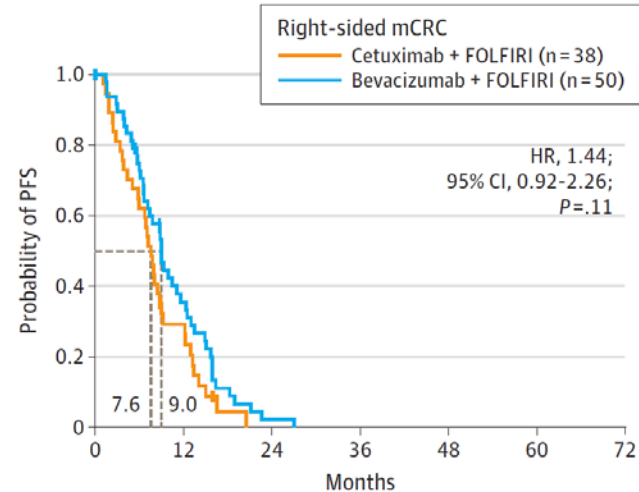
**A** Progression-free survival



No. at risk

Cetuximab + FOLFIRI	157	60	17	10	6	4	0
Bevacizumab + FOLFIRI	149	56	13	7	2	0	0

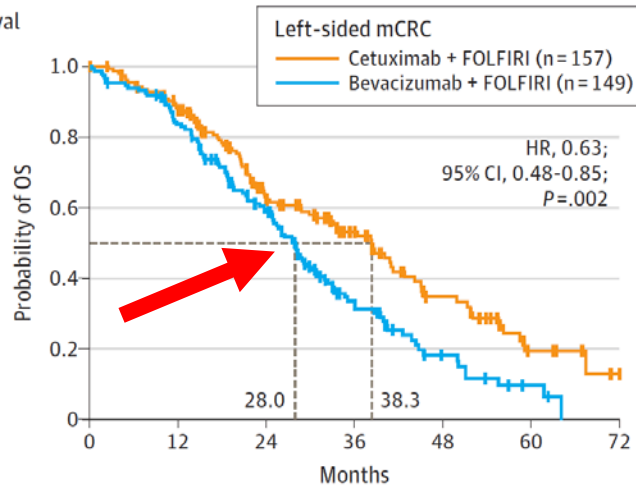
**Right-sided mCRC**



No. at risk

Cetuximab + FOLFIRI	38	10	0	0	0	0	0
Bevacizumab + FOLFIRI	50	16	1	0	0	0	0

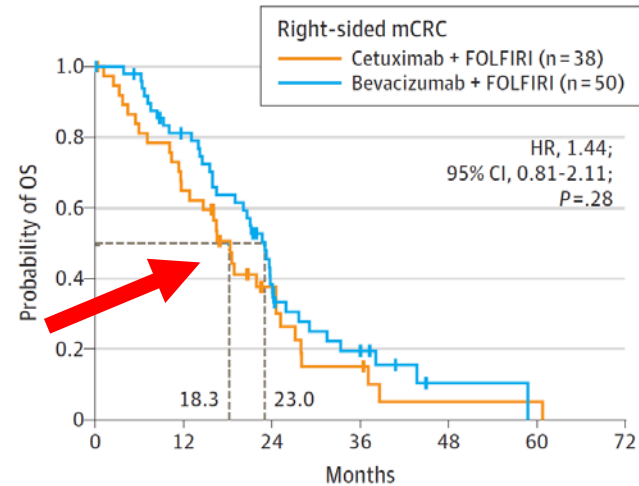
**B** Overall survival



No. at risk

Cetuximab + FOLFIRI	157	131	77	38	23	6	0
Bevacizumab + FOLFIRI	149	120	76	31	11	3	0

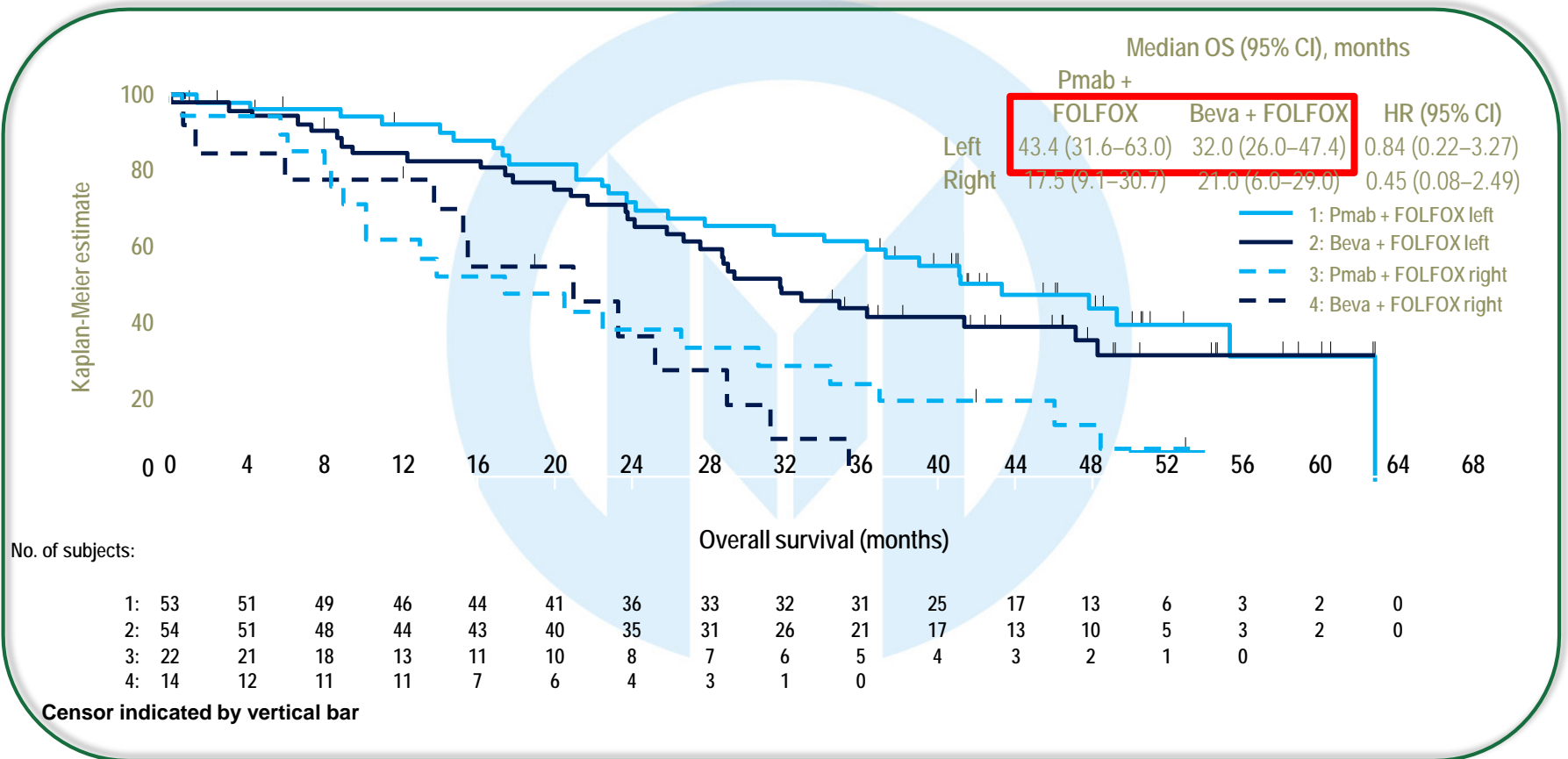
**Right-sided mCRC**



No. at risk

Cetuximab + FOLFIRI	38	24	10	4	1	1	0
Bevacizumab + FOLFIRI	50	37	16	7	1	0	0

# Right versus Left: PEAK – OS







# Right versus Left: CALGB80405 study - OS



31%



# Minimal difference between bev and cetux in left sided tumors

Population	Median OS (95% CI)		Log-rank	Adjusted HR
	L-sided	R-sided	p-value	(95%, CI)
80405 (N = 728)	32.9 (30.7, 35.3)	19.6 (7.0, 23.6)	< 0.0001	1.39 (1.03, 1.88)
<i>All RAS / BRAF wt (N = 225)</i>				
BV (N = 91)	38.7 (34.3, 42.3)	34.4 (23.6, 82.0)	0.918	0.62 (0.32, 1.23)
Cet (N = 96)	40.3 (34.0, 48.3)	18.4 (14.2, 30.1)	0.003	1.68 (0.85, 3.34)
<i>BRAF mut (N = 48)</i>				
BV (N = 23)	12.0 (4.8, 14.5)	23.7 (7.9, 36.9)	0.035	
Cet (N = 16)	9.6 (8.6, NE)	5.8 (1.9, 11.7)	0.508	

# Take Home Points

- Left sided primary colorectal cancers have better prognosis than right sided colon cancers
- Right sided colorectal cancers do not benefit from anti-EGFR therapy but do benefit from bevacizumab
- Left sided tumors benefit from both bevacizumab and anti-EGFR therapy

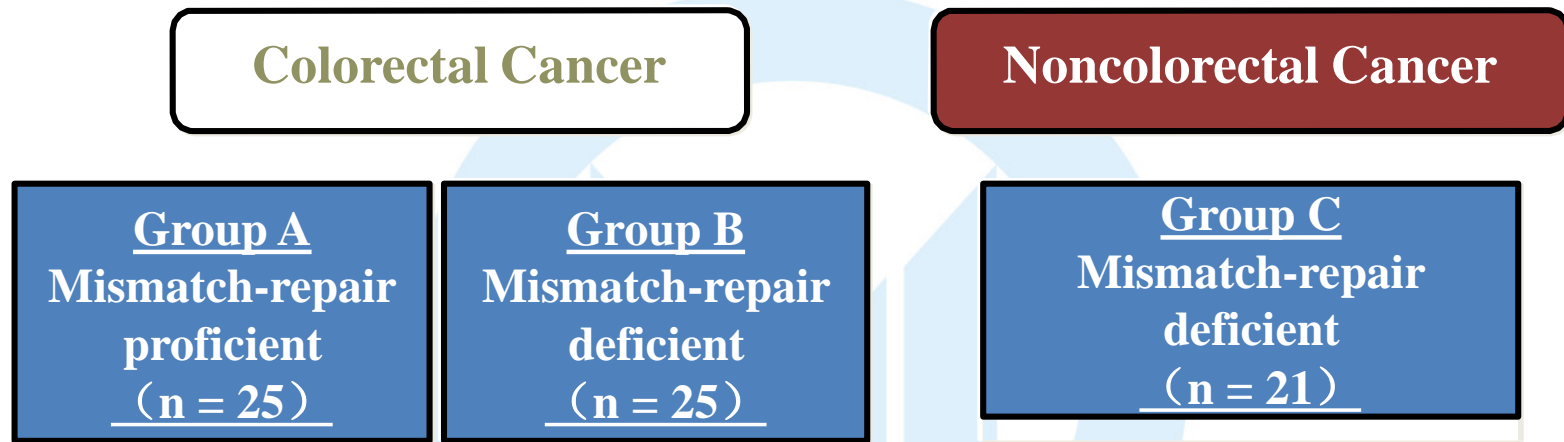


# **MSI-high CRC and Immune Checkpoint Blockade**

# Background

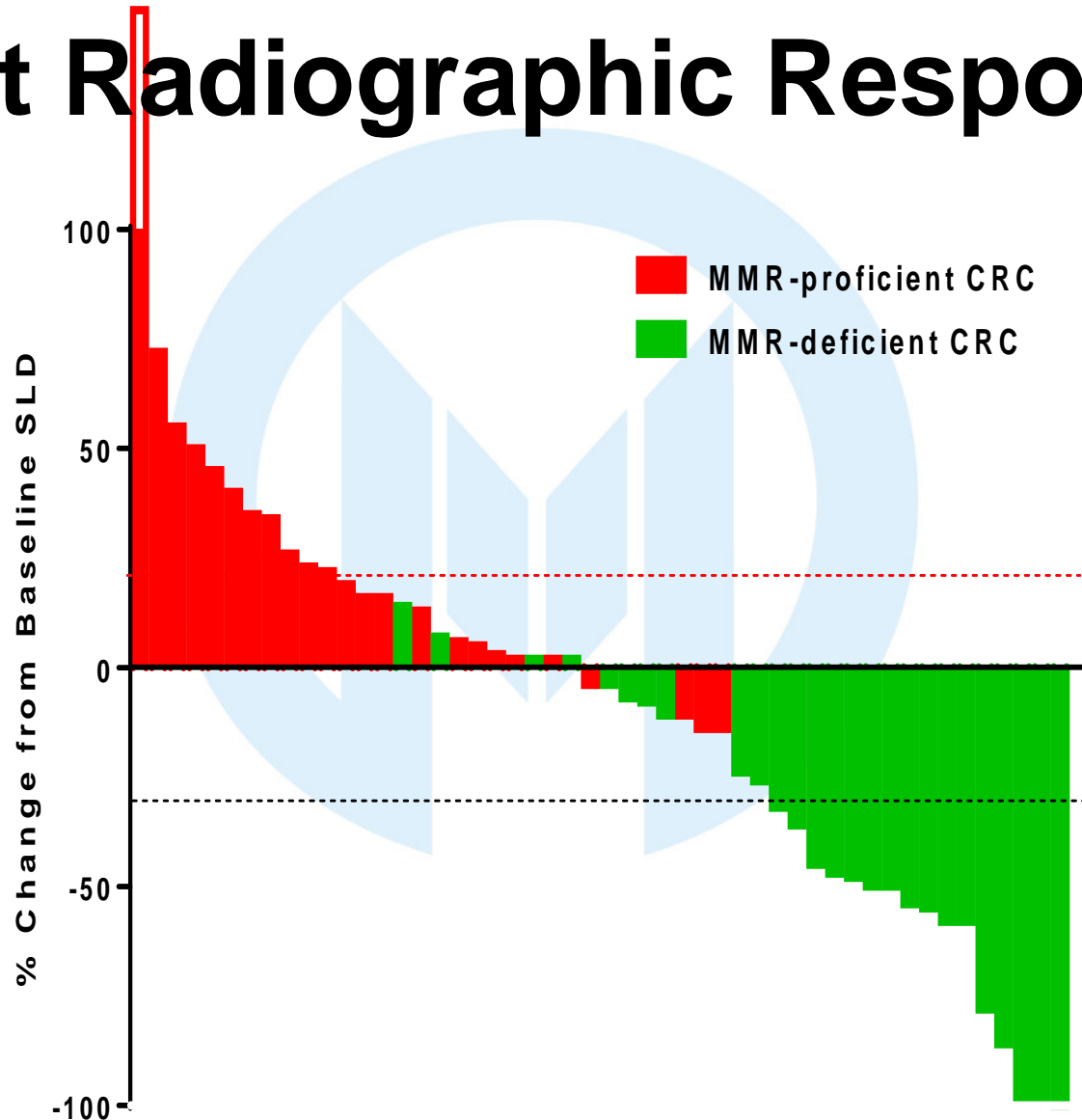
- In non selected colorectal cancer patients PD-1 blockade seems to be ineffective.
- Average tumor has dozens of somatic mutations.
- Mismatch repair deficient tumors harbor thousands of mutations
- Somatic mutations have the potential to generate neo-antigens which can be recognized by immune system.

# Clinical Trial Design: Basket Trial



- **Anti-PD-1 antibody (Pembrolizumab): 10 mg/kg q 2 week**
- **Primary endpoint: Immune-related ORR and the 20-week immune-related PFS rate**
- **Mismatch-repair status was assessed in tumors with the use of the MSI Analysis System**

# Best Radiographic Response



# Objective Responses According to RECIST Criteria

**Table 2.** Objective Responses According to RECIST Criteria.

Type of Response	Mismatch Repair–Deficient Colorectal Cancer (N = 10)	Mismatch Repair–Proficient Colorectal Cancer (N = 18)	Mismatch Repair–Deficient Noncolorectal Cancer (N = 7)
Complete response — no. (%)	0	0	1 (14)*
Partial response — no. (%)	4 (40)	0	4 (57)†
Stable disease at week 12 — no. (%)	5 (50)	2 (11)	0
Progressive disease — no. (%)	1 (10)	11 (61)	2 (29)
Could not be evaluated — no. (%)‡	0	5 (28)	0
Objective response rate (95% CI) — %	40 (12–74)	0 (0–19)	71 (29–96)
Disease control rate (95% CI) — %§	90 (55–100)	11 (1–35)	71 (29–96)
Median duration of response — wk	Not reached	NA¶	Not reached
Median time to response (range) — wk	28 (13–35)	NA¶	12 (10–13)

\* The patient had a partial response at 12 weeks, which then became a complete response at 20 weeks.

† One patient had a partial response at 12 weeks.

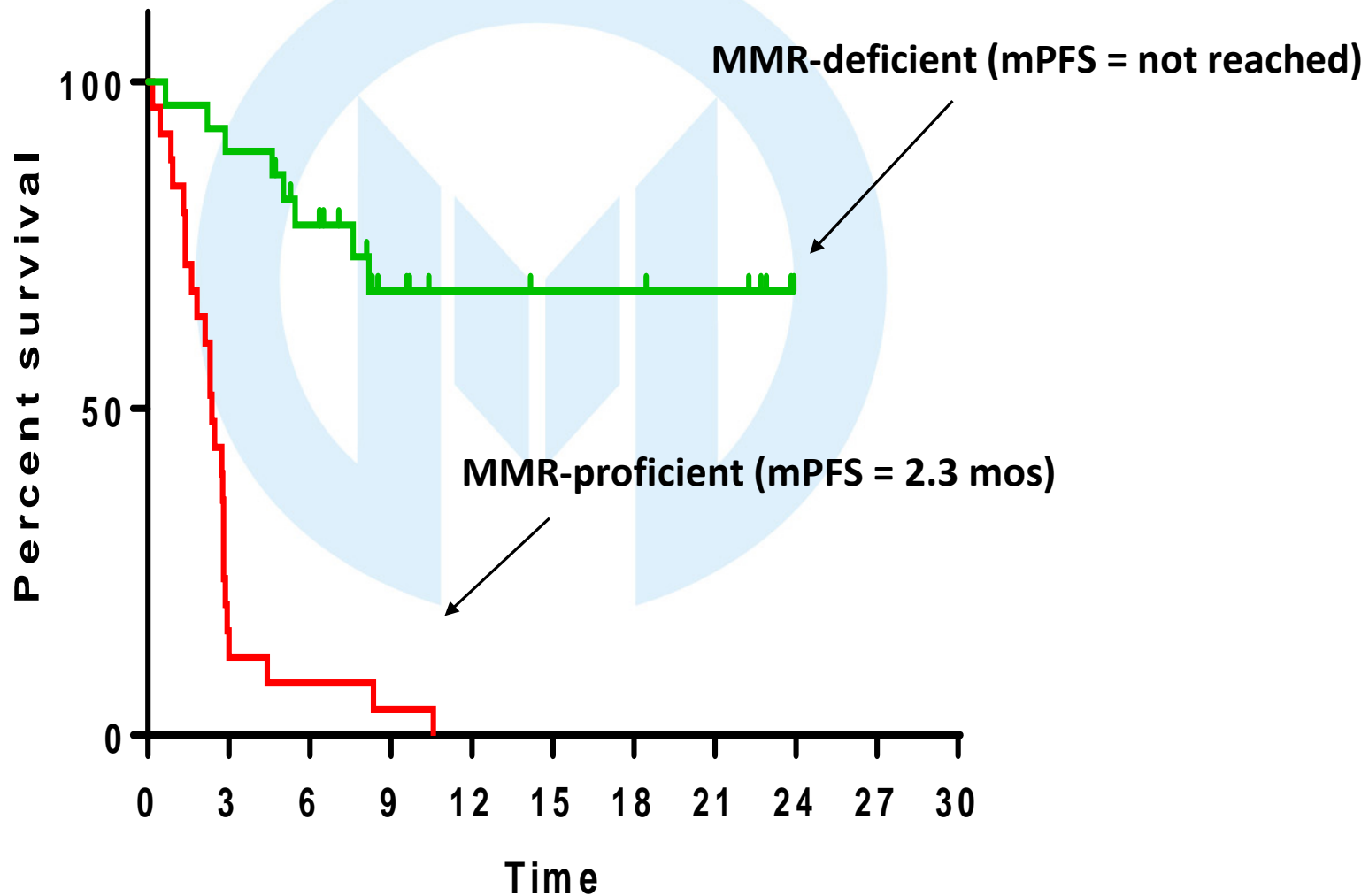
‡ Patients could not be evaluated if they did not undergo a scan at 12 weeks because of clinical progression.

§ The rate of disease control was defined as the percentage of patients who had a complete response, partial response, or stable disease for 12 weeks or more.

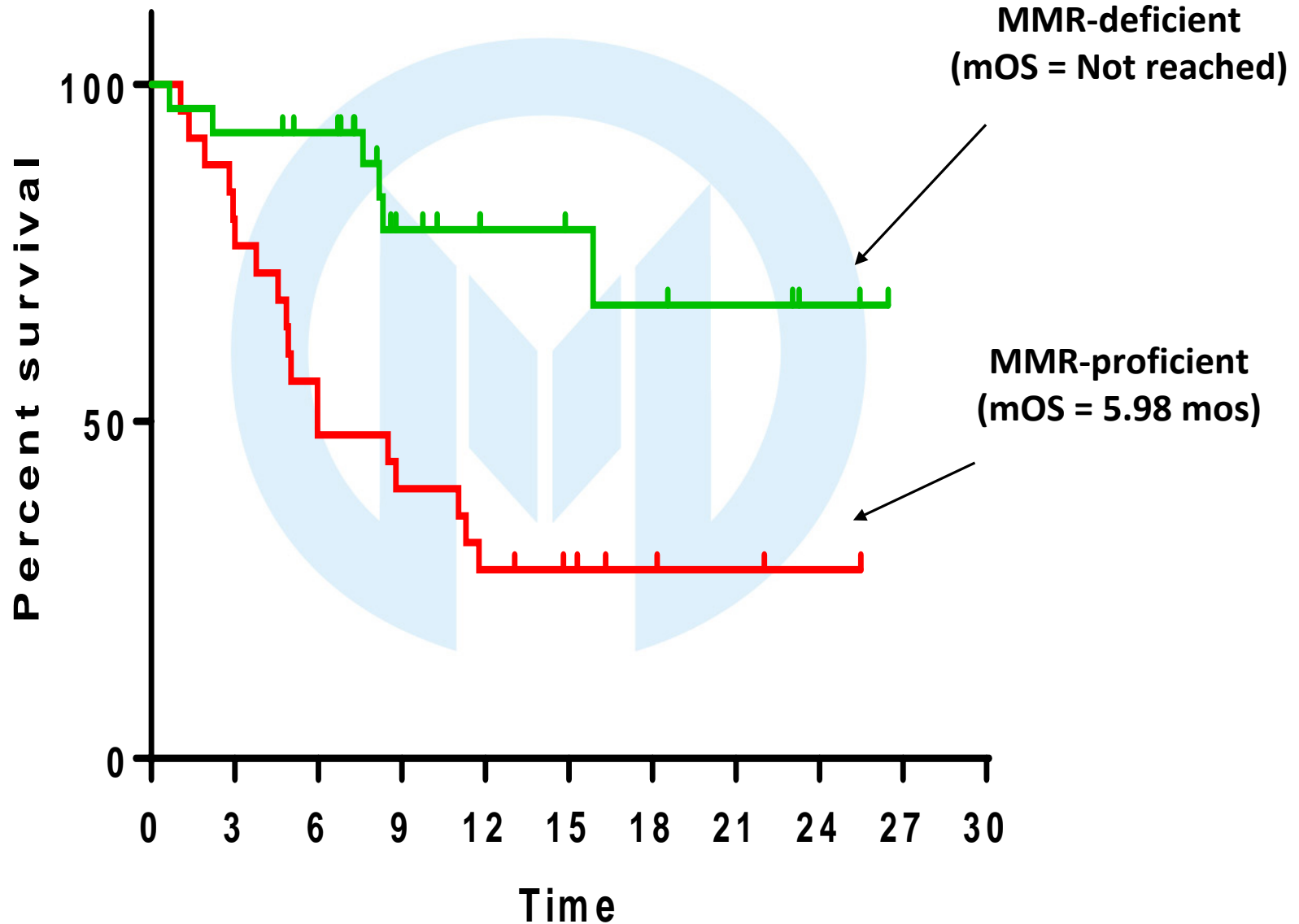
¶ The median time to response was not applicable (NA) because no responses were observed among patients with mismatch repair–proficient colorectal cancer.



# Progression-free Survival



# Overall Survival



# Adverse Events

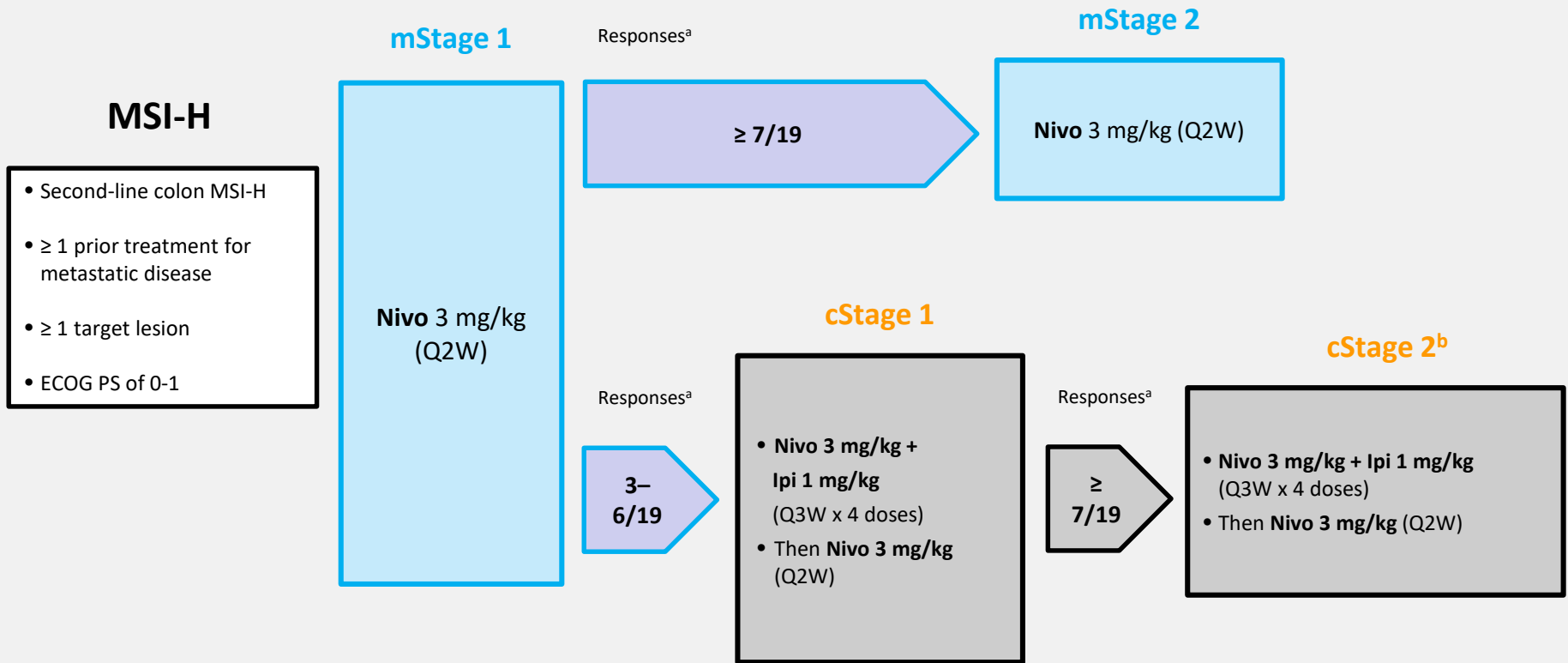
<i>Event-no. (%)</i>	All Grades N=53	Grade 3 or 4 N=53
<b>Generalized Symptoms</b>		
<i>Fatigue</i>	5 (9)	0 (0)
<i>Arthralgias</i>	8 (15)	0 (0)
<b>Gastrointestinal</b>		
<i>Nausea/vomiting</i>	4 (8)	0 (0)
<i>Diarrhea/colitis</i>	6 (11)	1 (2) ←
<b>Endocrine Disorders</b>		
<i>Thyroiditis/hypothyroidism</i>	6 (11)	0 (0)
<b>Hepatobiliary</b>		
<i>Pancreatitis</i>	4 (8)	2 (4) ←
<i>hyperbilirubinemia</i>	2 (4)	0 (0)
<b>Rash/pruritus</b>	13 (25)	1 (2) ←
<b>Respiratory</b>		
<i>Pneumonitis</i>	2 (4)	0 (0)
<b>Other</b>		
<i>Anemia</i>	2 (4)	1 (2) ←
<i>Flu-like symptoms</i>	2 (4)	0 (0)
<i>Leukopenia</i>	2 (4)	1 (2) ←
<i>Thrombocytopenia</i>	3 (6)	1 (2) ←

# Nivolumab ± Ipilimumab in Treatment of Patients With Metastatic Colorectal Cancer With and Without High Microsatellite Instability: CheckMate 142 Interim Results

Michael Overman,<sup>1</sup> Scott Kopetz,<sup>1</sup> Ray McDermott,<sup>2</sup> Joseph Leach,<sup>3</sup> Sara Lonardi,<sup>4</sup> Heinz-Josef Lenz,<sup>5</sup> Michael Morse,<sup>6</sup> Jayesh Desai,<sup>7</sup> Andrew Hill,<sup>8</sup> Michael Axelson,<sup>9</sup> Rebecca A. Moss,<sup>9</sup>  
Chen-Sheng Lin,<sup>9</sup> Monica Goldberg,<sup>9</sup> Thierry Andre<sup>10</sup>

<sup>1</sup>MD Anderson Cancer Center, Houston, TX, USA; <sup>2</sup>St Vincent's University Hospital, Dublin, Ireland; <sup>3</sup>Allina Health System, Minneapolis, MN, USA; <sup>4</sup>Istituto Oncologico Veneto IOV-IRCSS, Padova, Italy; <sup>5</sup>USC Norris Comprehensive Cancer Center, Los Angeles, CA, USA; <sup>6</sup>Duke University Office of Research Administration, Durham, NC, USA; <sup>7</sup>Royal Melbourne Hospital, Victoria, Australia; <sup>8</sup>Tasman Oncology Research Pty Ltd, Southport, Queensland, Australia; <sup>9</sup>Bristol-Myers Squibb, Princeton, NJ, USA; <sup>10</sup>Hopital Saint Antoine, Paris, France

# Phase 2 CheckMate 142 Study Design: MSI-H Cohort



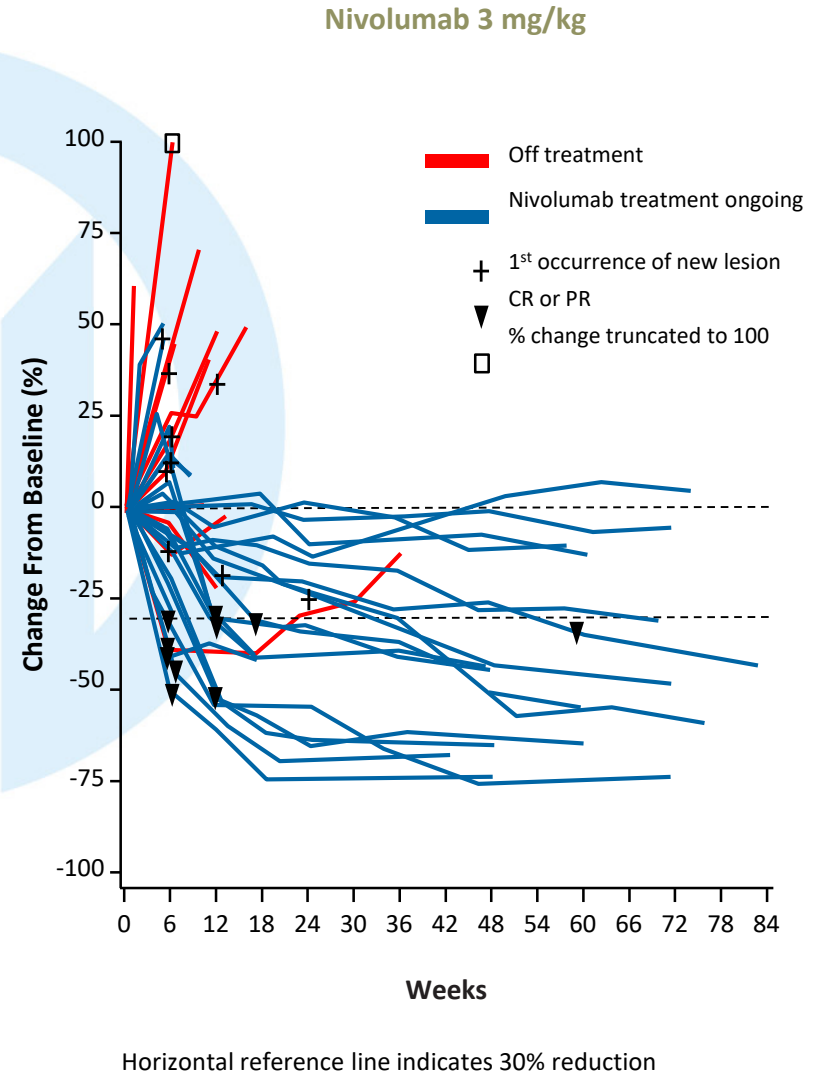
<sup>a</sup>In patients with centrally confirmed MSI-H status

<sup>b</sup>Currently enrolling

cStage 1 = combination therapy stage 1; cStage 2 = combination therapy stage 2; Ipi = ipilimumab; mStage 1 = monotherapy stage 1; mStage 2 = monotherapy stage 2; Nivo = nivolumab; Q2W = every 2 weeks; Q3W = every 3 weeks

## Investigator-Assessed Best Overall Response in Patients With MSI-H Receiving Nivolumab Monotherapy

	<b>Nivolumab 3 mg/kg (n = 47)<sup>a</sup></b>
ORR, n (%) (95% exact CI)	12 (25.5) (15.4, 38.1)
Complete response	0
Partial response	12 (25.5)
Stable disease	14 (29.8)
Progressive disease	17 (36.2)
Unable to determine	4 (8.5)
Median time to response, mo (range)	2.12 (1.3–13.6)
Median duration of response, mo (range)	NE (0.0 <sup>b</sup> –15.2 <sup>b</sup> )



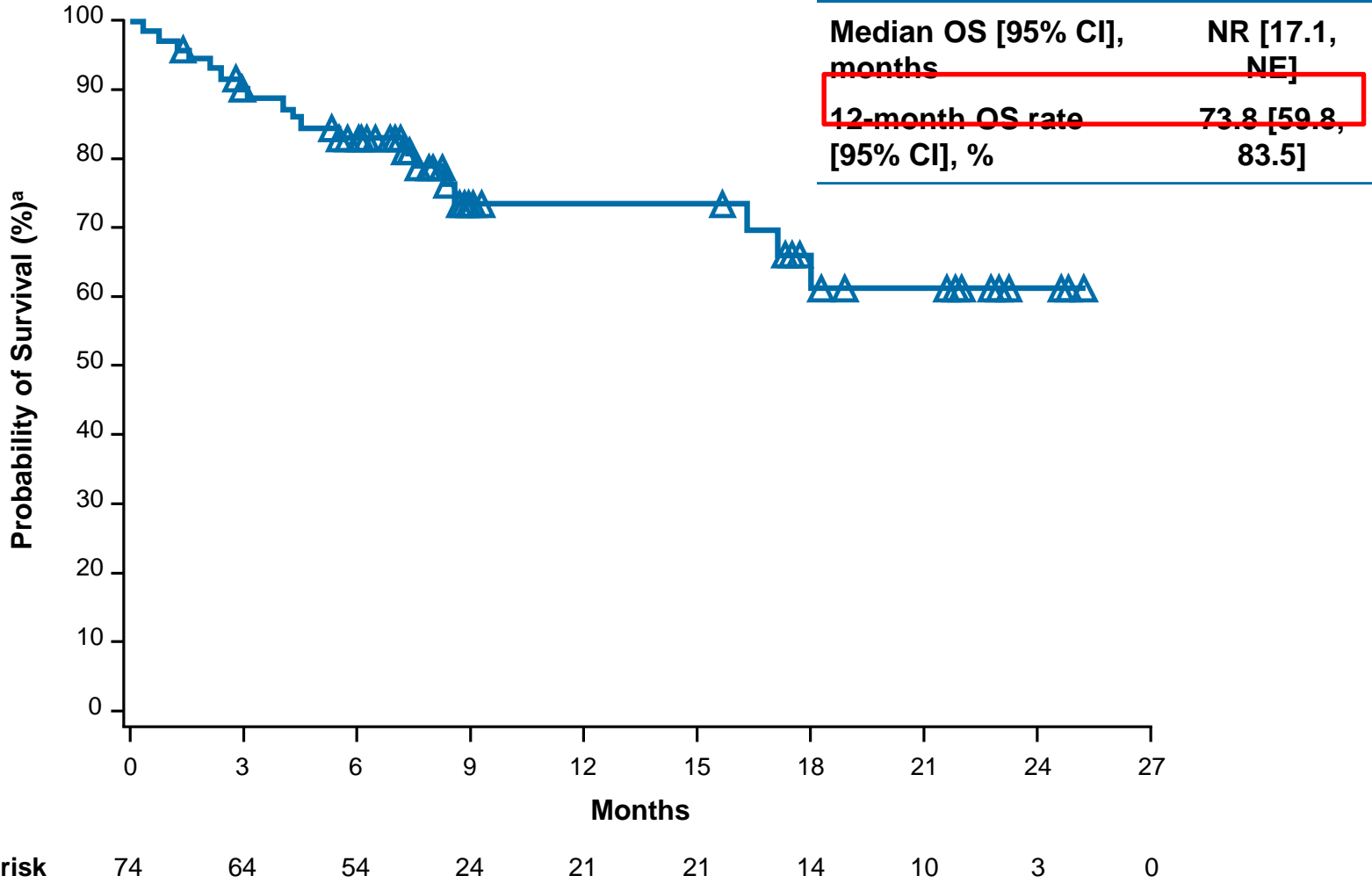
# Response and Disease Control

	dMMR/MSI-H per Local Laboratory (N = 74)		dMMR/MSI-H per Central Laboratory (n = 53)	
	Investigator	BICR	Investigator	BICR
ORR, n (%)	23 (31.1)	20 (27.0)	19 (35.8)	17 (32.1)
95% CI	20.8, 42.9	17.4, 38.6	23.1, 50.2	19.9, 46.3
Best overall response, n (%)				
CR	0	2 (2.7)	0	1 (1.9)
PR	23 (31.1)	18 (24.3)	19 (35.8)	16 (30.2)
SD	29 (39.2)	28 (37.8)	21 (39.6)	21 (39.6)
PD	18 (24.3)	20 (27.0)	10 (18.9)	12 (22.6)
Unable to determine	4 (5.4)	6 (11.1)	3 (5.7)	3 (5.7)
Disease control for $\geq 12$ weeks, n (%) <sup>a</sup>	51 (68.9)	46 (62.2)	39 (73.6)	37 (69.8)

BICR, blinded independent central review.

<sup>a</sup> Patients with CR, PR, or SD for  $\geq 12$  weeks.

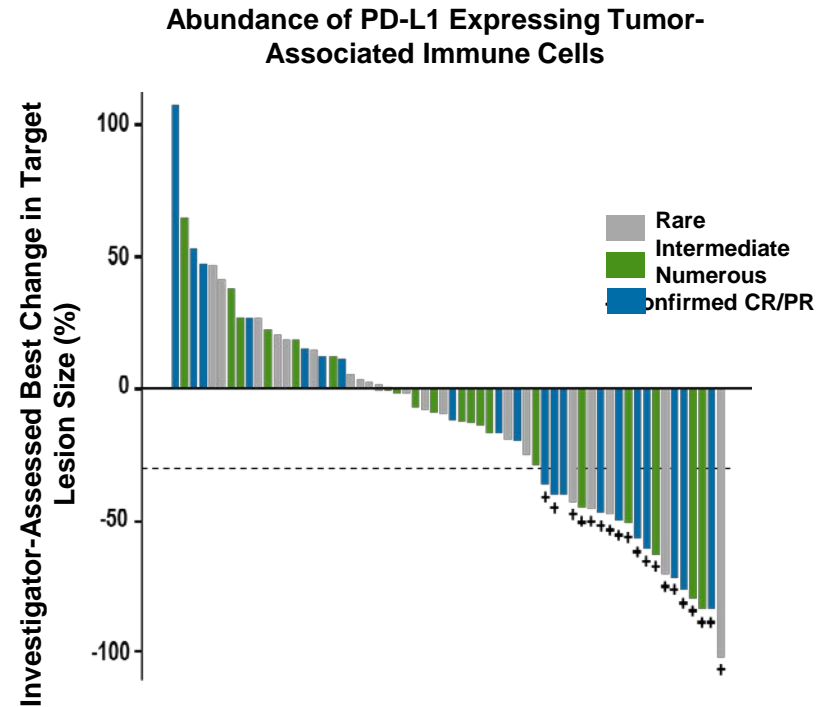
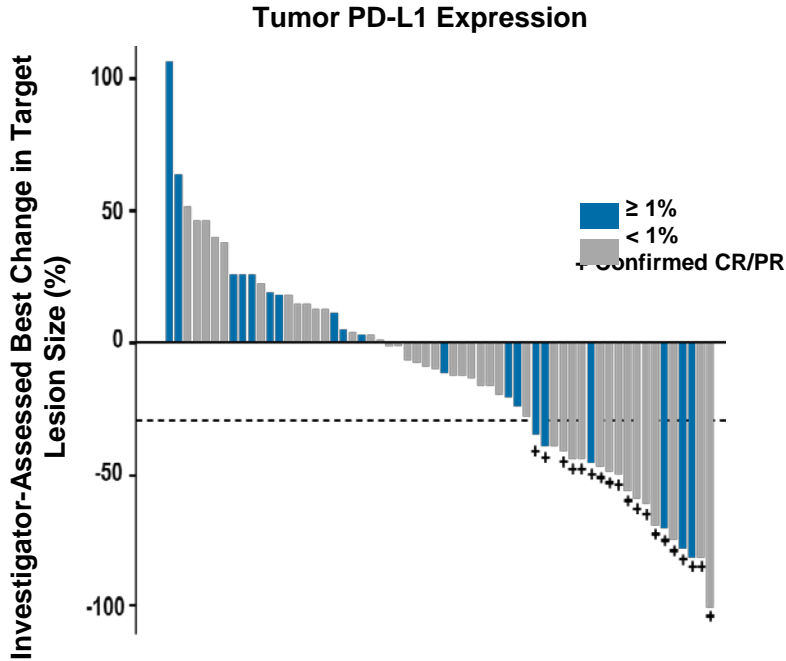
# Overall Survival



NR, not reached. <sup>a</sup> Patients evaluated as dMMR/MSI-H by local laboratory.



# Reduction in Target Lesion Size Regardless of PD-L1 Expression

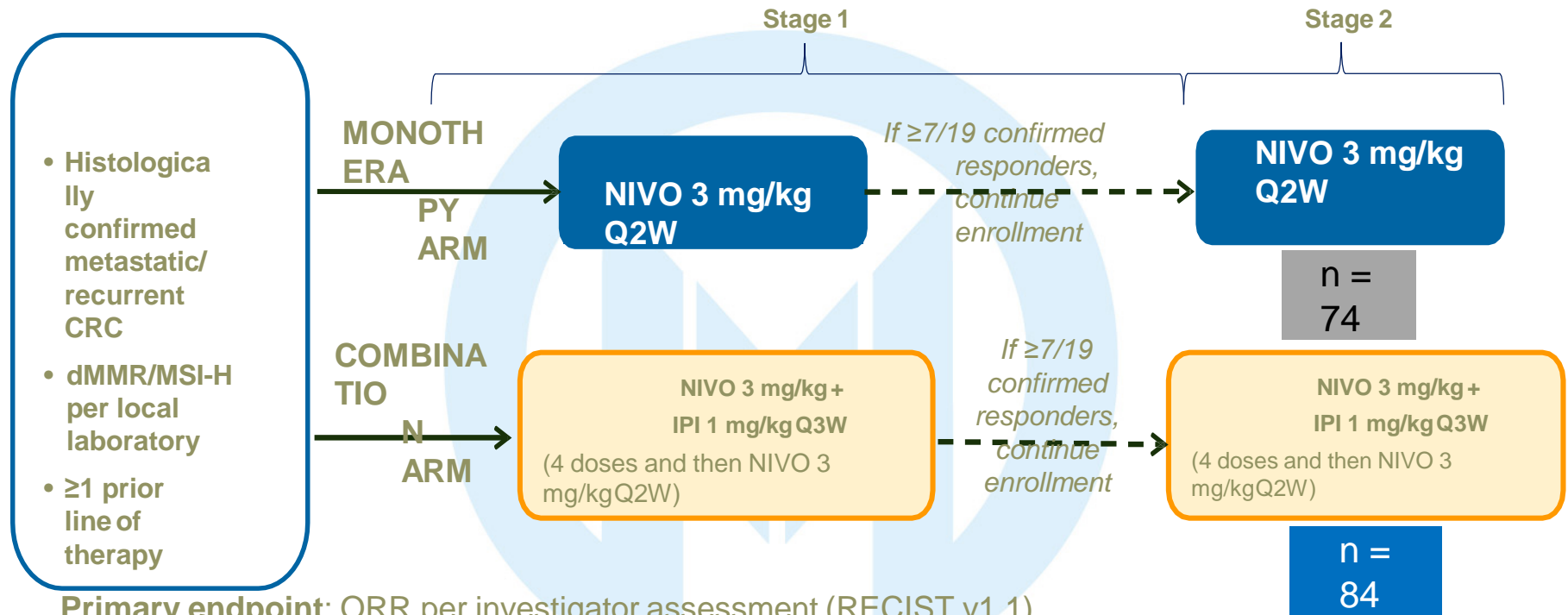


ORR, n/N (%)	Investigator	BICR
Tumor PD-L1 expression		
≥ 1%	6/21 (28.6)	7/20 (35.0)
< 1%	13/45 (28.9)	11/45 (24.4)

ORR, n/N (%)	Investigator	BICR
Abundance of PD-L1 expressing immune cells		
Rare	5/23 (21.7)	4/20 (20.0)
Intermediate	5/20 (25.0)	10/23
Numerous	9/23 (39.1)	(43.5)

# Study Design

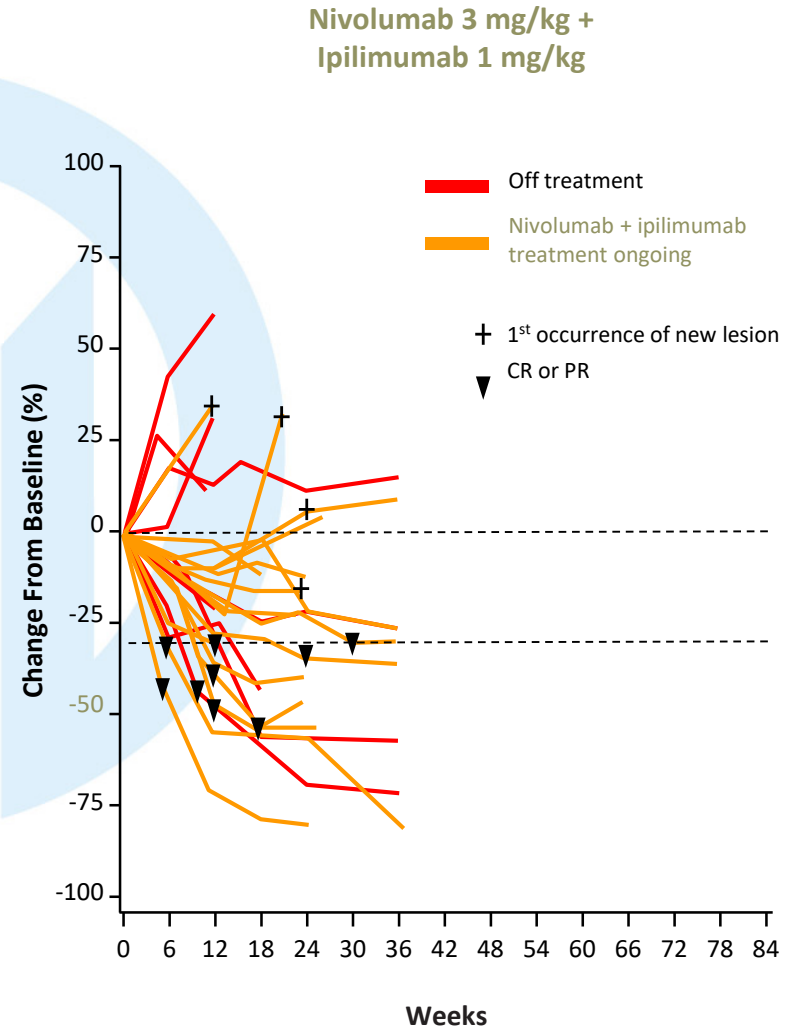
Patients



- Current analysis included all patients (n = 84) who received their first dose  $\geq 6$  months prior to the data cut-off
  - Median (range) time from first dose to data cut-off: 8.6 (6.3-19.4) months

# Investigator-Assessed Best Overall Response in Patients With MSI-H Receiving Nivolumab + Ipilimumab

	Nivolumab 3 mg/kg + Ipilimumab 1 mg/kg (n = 27) <sup>a</sup>
ORR, n (%) (95% exact CI)	11 (41.3) (18.6, 50.9)
Complete response	1
Partial response	10(37)
Stable disease	14 (51.9)
Progressive disease	2(7 )
Unable to determine	0
Median time to response, mo (range)	2.73 (1.2–6.9)
Median duration of response, mo (range)	NE (NE–NE)



<sup>a</sup>Patients with ≥ 12 weeks of follow-up

<sup>b</sup>Includes censored observations

CR = complete response; NE = not estimable; PR = partial response

Horizontal reference line indicates 30% reduction

## Overall Response and Disease Control

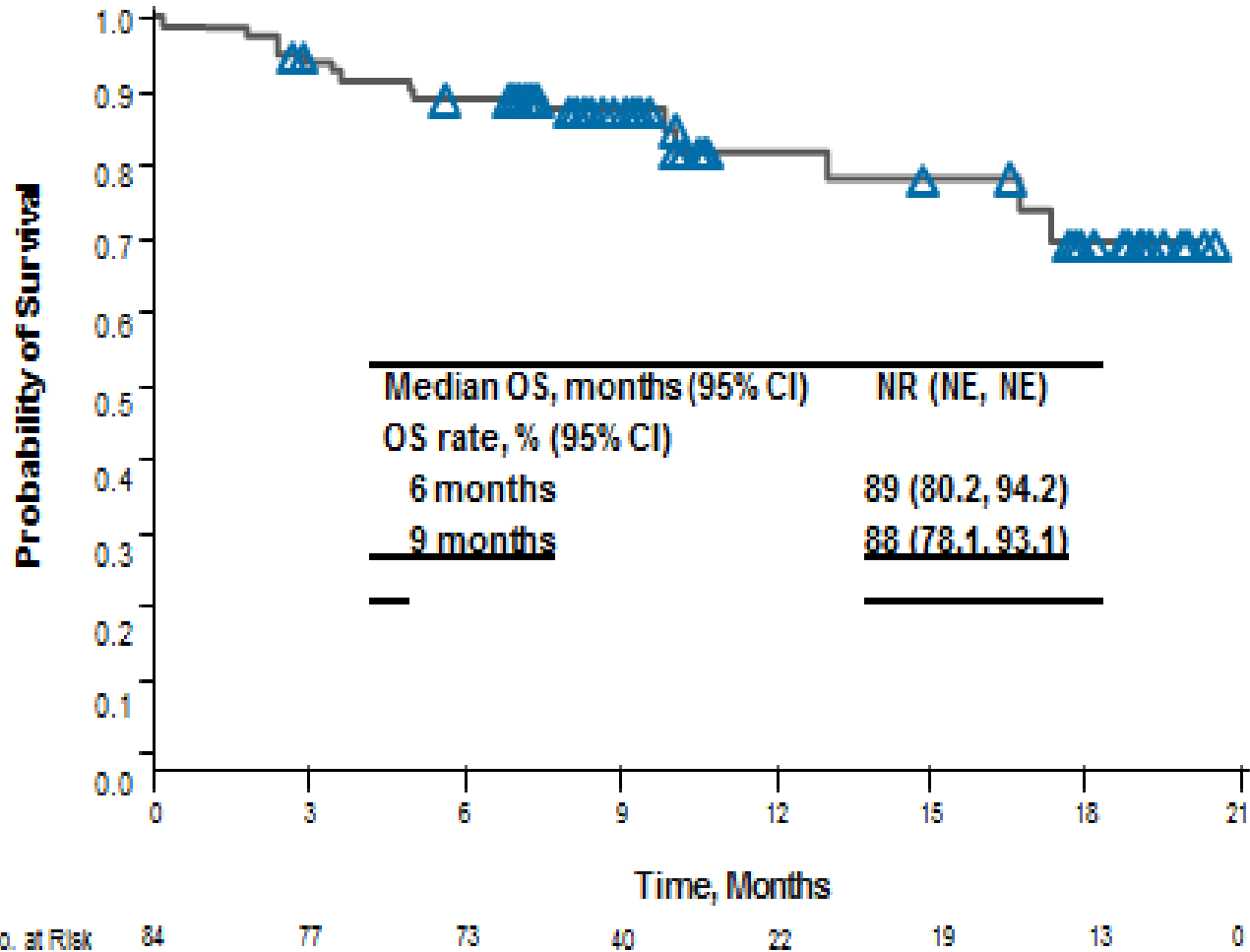
	NIVO + IPI (n = 84)	NIVO Monotherapy <sup>1</sup> (n = 74)
	Investigator-Assessed	Investigator-Assessed
ORR, n (%) [95% CI]	46 (55) [43.5, 65.7]	23 (31) [20.8, 42.9]
Best overall response, n (%)		
CR	2 (2)	0
PR	44 (52)	23 (31)
SD	26 (31)	29 (39)
PD	9 (11)	18 (24)
Not determined/reported	3 (4)	4 (5)
Disease control for ≥12 weeks, n (%) <sup>a</sup>	66 (79)	51 (69)

<sup>a</sup>Patients with CR, PR, or SD for ≥12 weeks

1. Overman M, et al. *J Clin Oncol* 2017;35(suppl 4 8): Abstract 618.

Andre T, et al. *J Clin Oncol*. 2017;35(suppl): Abstract 3531.

# Overall Survival



## Treatment-Related Adverse Events in $\geq 15\%$ of Patients With MSI-H

Event, n (%)	Nivolumab 3 mg/kg (n = 70)		Nivolumab 3 mg/kg + Ipilimumab 1 mg/kg (n = 30)	
	Any grade	Grade 3–4	Any grade	Grade 3–4
Any event	41 (58.6) <sup>a</sup>	10 (14.3)	25 (83.3)	8 (26.7)
Fatigue	13 (18.6)	1 (1.4)	6 (20.0)	0
Diarrhea	10 (14.3)	1 (1.4)	13 (43.3)	0
Pruritus	8 (11.4)	0	5 (16.7)	1 (3.3)
Nausea	5 (7.1)	0	6 (20.0)	0
Pyrexia	3 (4.3)	0	7 (23.3)	0
Any event leading to discontinuation	4 (5.7)	2 (2.9)	4 (13.3)	4 (13.3)

<sup>a</sup>One Grade 5 event of sudden death

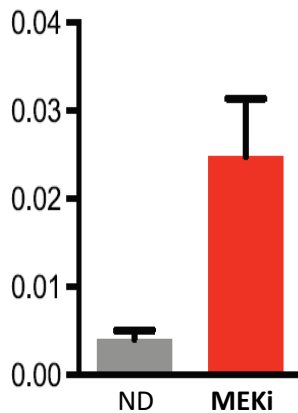
# Question??

- Can we convert non-immunogenic tumor into immunogenic tumor?

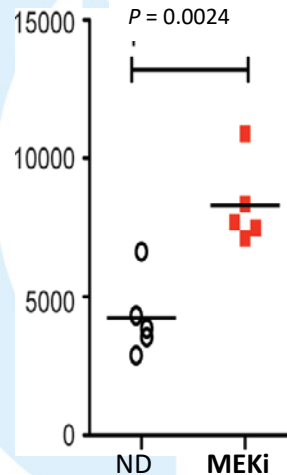
## PD-L1 and MEK Inhibition: A Rational Combination

- MEK inhibition alone can result in **intratumoral T-cell accumulation** and **MHC I upregulation**, and synergizes with an anti-PDL1 agent to promote **durable tumor regression**<sup>1</sup>

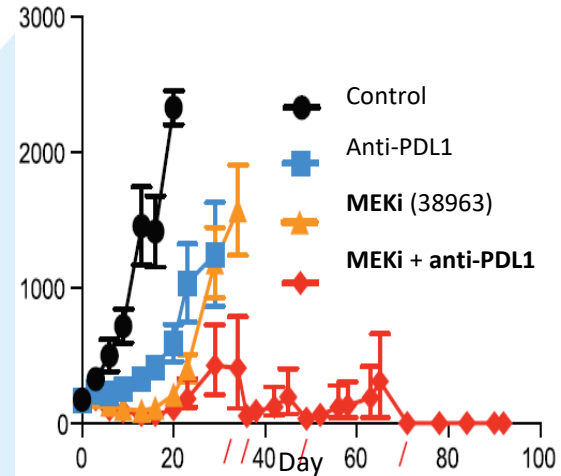
CD8<sup>+</sup> T cell  
per tumor cell



Class I MHC



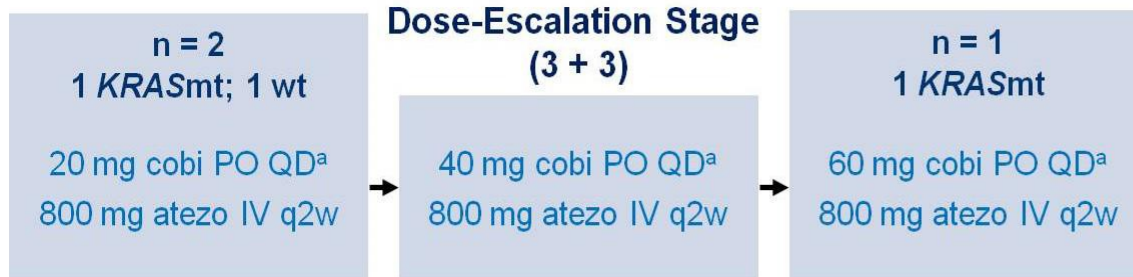
Tumor volume (mm<sup>3</sup>)



- To examine the possible benefits of MEK inhibition with an anti-PDL1 agent, we evaluated cobimetinib + atezolizumab in patients with advanced solid tumors



# Cobimetinib + Atezolizumab in CRC: Phase Ib Dose Escalation and Cohort Expansion Study

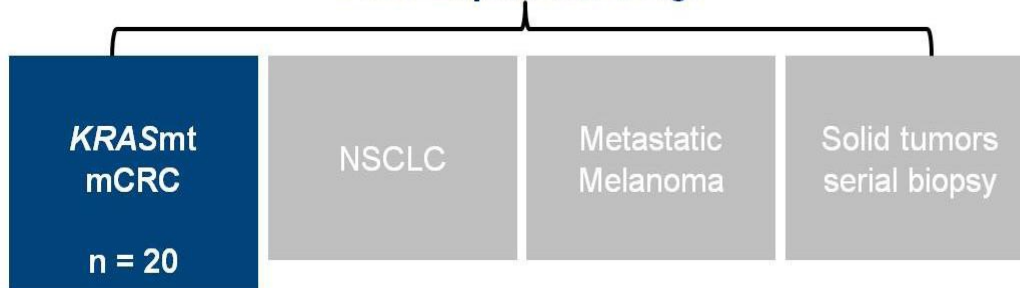


DLT window of 28 days until MTD for combination is defined

## Key eligibility Criteria

- ECOG PS of 0 or 1
- Measurable disease per RECIST v1.1

## Dose-Expansion Stage



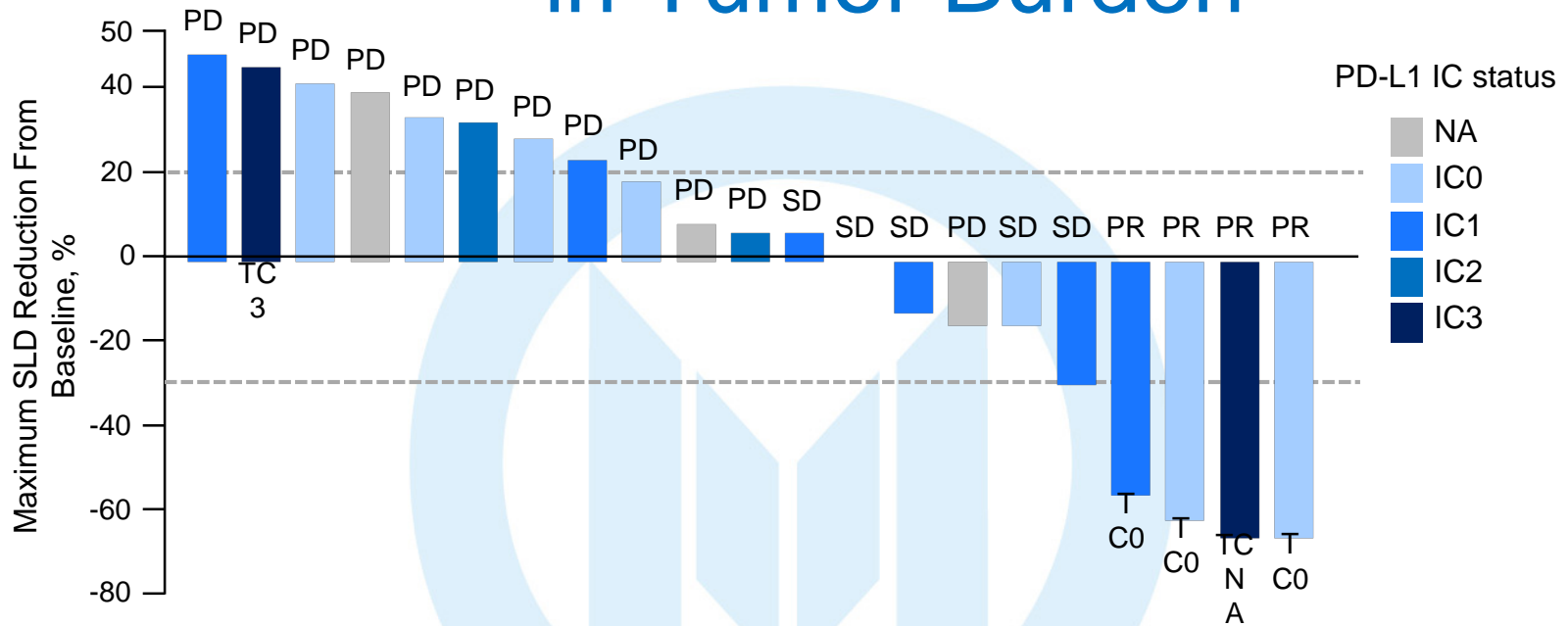
## Primary Objectives

- Safety and clinical activity of cobimetinib + atezolizumab

<sup>a</sup>Cobimetinib was administered on 21 days on/7 days off dosing schedule.

Atezo, atezolizumab; cobimetinib; DLT, drug limited toxicity; ECOG PS, Eastern Cooperative Oncology Group performance status; KRASmt, KRAS mutant; MTD, maximum tolerated dose; NSCLC, non-small cell lung cancer

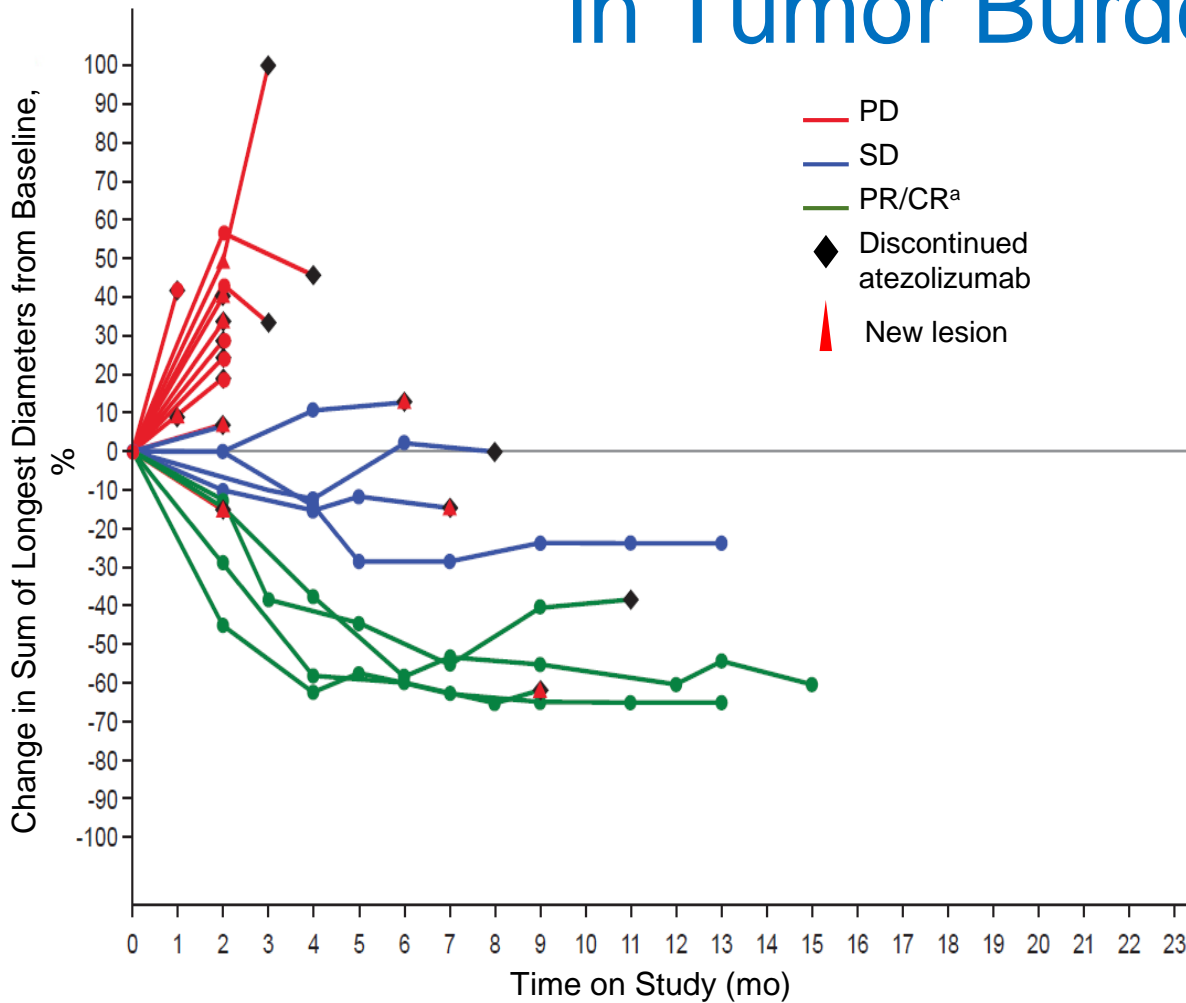
# Efficacy: Change in Tumor Burden



- 4 patients had partial responses (confirmed per RECIST v1.1)
- MSI status of CRC patients was examined by NGS-based scoring: 3 of 4 responders were mismatch-repair proficient (not MSI-H); 1 responder had unknown MSI status and was not evaluable
- Tumor volume reduction was not associated with PD-L1 status: TC3 (n = 1; PD), TC0 (n = 18), NA (n = 4)

PD-L1 IHC status on tumor cells (TC) and tumor-infiltrating immune cells (IC) defined as: TC3 = TC ≥ 50% PD-L1+ cells; IC3 = IC ≥ 10% PD-L1+ cells; TC2 = TC ≥ 5% and < 50% PD-L1+ cells; IC2 = IC ≥ 5% and < 10% PD-L1+ cells; TC1 = TC ≥ 1% and < 5% PD-L1+ cells; IC1 = IC ≥ 1% and < 5% PD-L1+ cells; TC0 = TC < 1% PD-L1+ cells; IC0 = IC < 1% PD-L1+ cells. NA, not available; NGS, next generation sequencing. Efficacy-evaluable patients. 2 patients missing or unevaluable are not included. Data cut-off February 12, 2016.

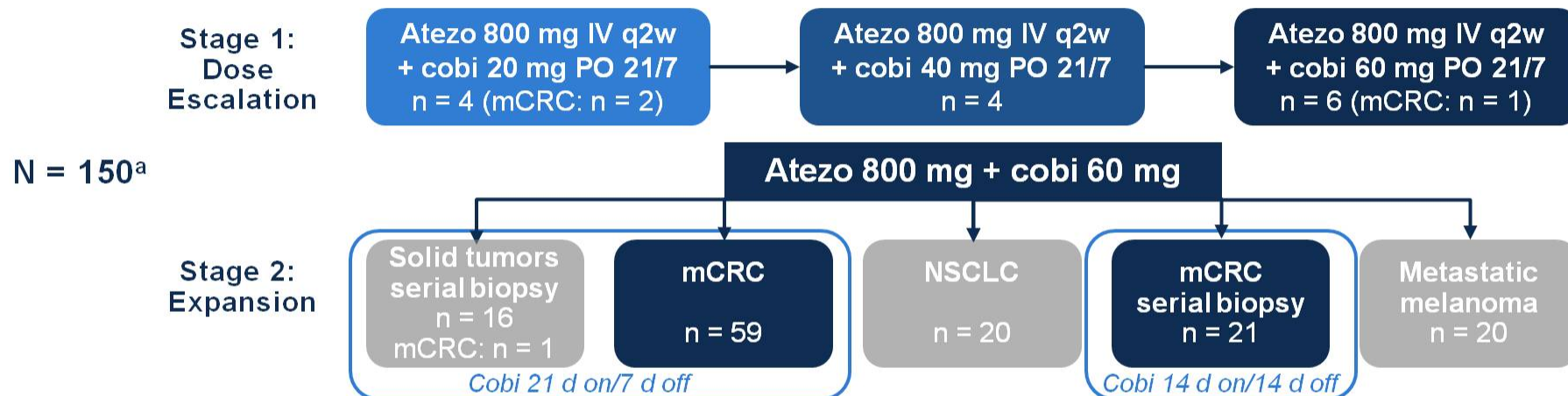
# Efficacy: Change in Tumor Burden Over Time



- Median duration of response was not reached (range: 5.4 to 11.1+ mo)
- Responses are ongoing in 2 of 4 responding patients

<sup>a</sup>Confirmed per RECIST v1.1. CR, complete response; PD, progressive disease; PR, partial response; SD, stable disease. Efficacy-evaluable patients. 2 patients missing or unevaluable are not included. Data cut-off February 12, 2016.

# Phase Ib Dose Escalation and Cohort Expansion Study (NCT01988896)



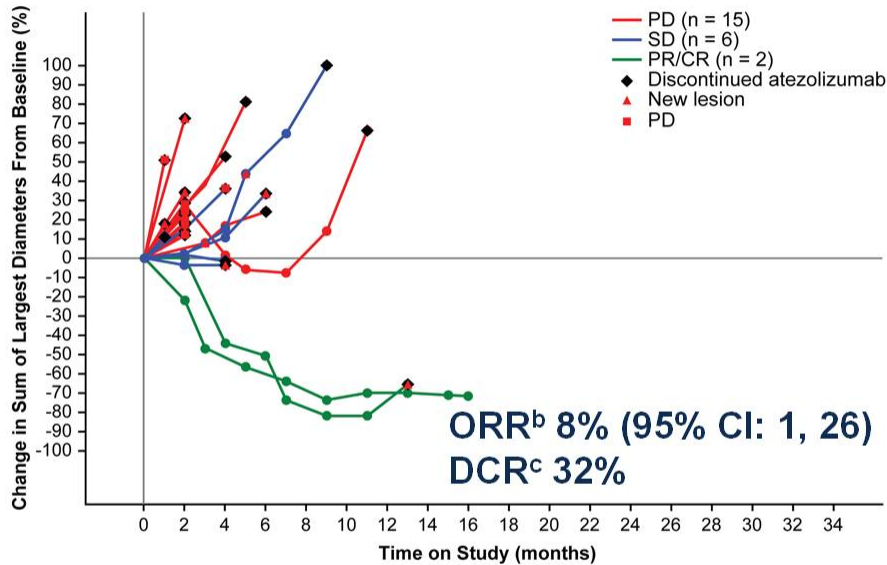
- **Endpoints:** Primary – Safety and tolerability  
Secondary – Investigator-assessed ORR and PFS by RECIST v1.1, and OS
- **Patients:** PD-L1 unselected  
MSI status was locally reported and centrally confirmed by NGS-based scoring

Atezo, atezolizumab; cobi, cobimetinib; IV, intravenously; NGS, next-generation sequencing; NSCLC, non-small cell lung cancer; ORR, overall response rate; OS, overall survival; PFS, progression-free survival; PO, orally; q2w, every 2 weeks.

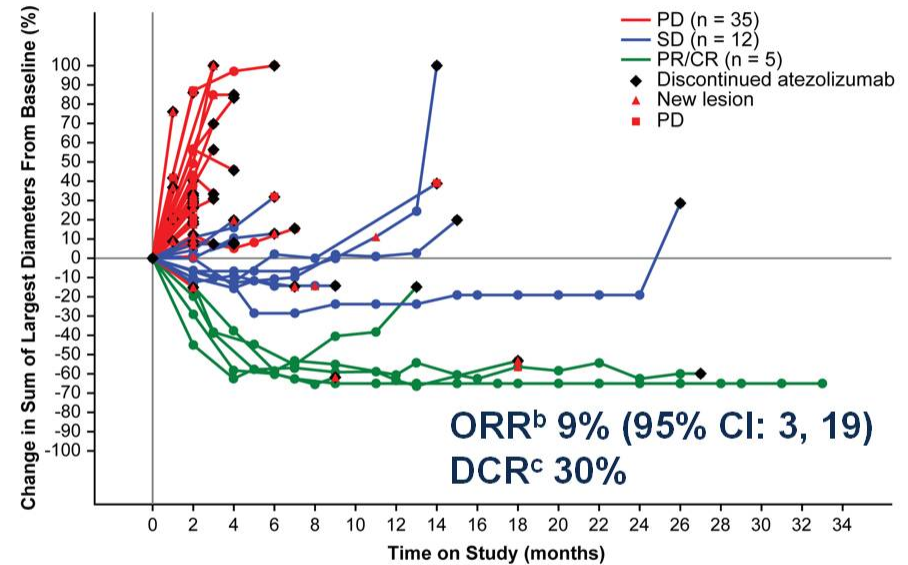
<sup>a</sup> Safety-evaluable population consisting of patients who received at least 1 dose of atezolizumab.

# Duration of Response

**KRAS Wild Type<sup>a,b</sup> (n = 23)**



**KRAS Mutant<sup>a,b</sup> (n = 52)**



- The median duration of response was 14.3 months (95% CI: 6.0, NE)

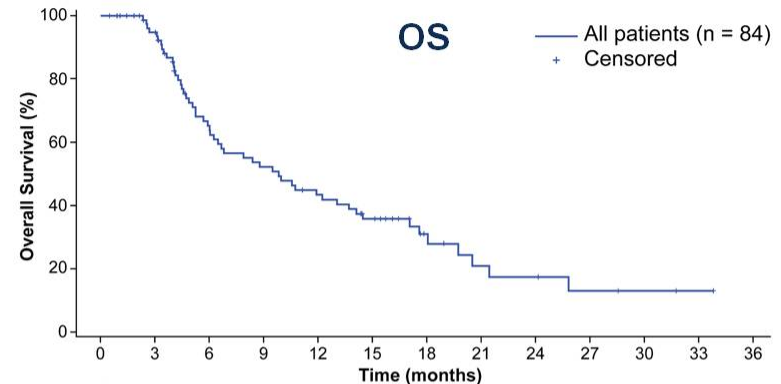
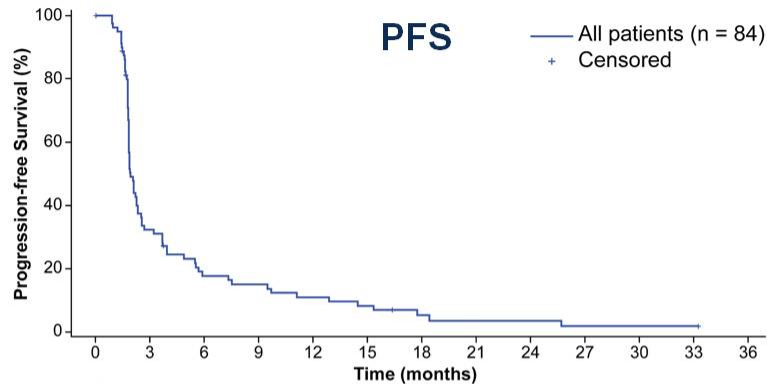
NE, not estimable.

Data cutoff: September 4, 2017.

<sup>a</sup> 2 patients had unknown *KRAS* mutation status and are not included in these graphs. <sup>b</sup> BOR was missing or unevaluable for 2 *KRAS* wild type and 5 *KRAS* mutant patients.

<sup>c</sup> DCR defined as PR or SD  $\geq$  6 weeks.

# Progression-Free Survival and Overall Survival



No. of Patients at Risk  
84 25 13 11 8 6 3 2 2 1 1 1

No. of Patients at Risk  
84 73 45 36 29 22 10 6 5 3 2 1

Patients	PFS		OS		
	Median (95% CI)	6-mo	Median (95% CI)	6-mo	12-mo
All (n = 84)	1.9 mo (1.8, 2.3)	18%	9.8 mo (6.2, 14.1)	65%	43%
MSS (n = 42) <sup>a</sup>	2.5 mo (1.8, 3.7)	27%	13.0 mo (6.0, 25.8)	71%	51%
<i>KRAS</i> mutant (n = 57) <sup>b</sup>	2.0 mo (1.8, 2.3)	22%	9.5 mo (6.0, 17.6)	67%	44%
<i>KRAS</i> wild type (n = 25) <sup>b</sup>	1.8 mo (1.8, 2.6)	9%	10.0 mo (4.9, 17.1)	65%	43%

Data cutoff: September 4, 2017.

<sup>a</sup> Of the remaining 42 non-MSS patients, 32 patients had unknown MSI status, 9 patients were MSI-low and 1 patient was MSI-high. <sup>b</sup> 2 patients had unknown *KRAS* mutation status.

### Eligibility Criteria:

- mCRC
  - $\geq 2$  prior therapies
  - Progression on or within 3 months of prior therapy
  - ECOG PS 0-1
- N = 360

2:1:1

R

**Atezolizumab + cobimetinib**  
840 mg IV days 1, 15 +  
60 mg days 1-21

**Atezolizumab**  
1200 mg IV on day 1  
28 day cycle

**Regorafenib**  
160 mg per d  
Days 1-21; 28 day cycle

Treatment until:  
Disease  
progression,  
unacceptable  
toxicity, or  
death

### Outcomes

- Primary endpoint: OS
- Secondary endpoint: PFS, ORR, DOR, QoL, safety

# Summary

- VEGF and EGFR mAbs competing for first-line patients in RAS wt CRC
- For anti-EGFR treatment, all *RAS* tests are required
- Primary tumor location is related to effect of cetuximab
- Bevacizumab, *Ramucirumab* and Aflibercept competing for second-line patients with each other, and with EGFR mAbs in KRAS wt CRC



# Summary

- Best sequence of therapies (VEGFi vs EGFRi) still to be established
- Regorafenib and TAS 102 as salvage therapy option
- Checkpoint inhibitors are highly active in select molecular subsets
- Rationale combination maybe able to covert “cold” tumor to “hot” tumor

Thank you !

GI oncology questions  
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