

*September 28<sup>th</sup> 2018*

# The Evolving Management of Metastatic Colorectal Cancer *Mt Tremblant*



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James Hamilton. c1840*



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**Good Afternoon**



# Disclosures

|                                  |   |
|----------------------------------|---|
| <b>Research Support/P.I.</b>     | Roche, Astra-Zeneca, Sanofi-Aventis   |
| <b>Employee</b>                  | N/A   |
| <b>Consultant</b>                | Lilly   |
| <b>Major Stockholder</b>         | N/A   |
| <b>Speakers Bureau</b>           | Lilly, Boehringer-Ingelheim, Merck, Astra-Zeneca  |
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| <b>Scientific Advisory Board</b> | N/A   |
| <b>Ownership</b>                 | Sarissa Inc   |
| <b>Board of Directors</b>        | Aptose BioSciences Inc  |

# Learning Objectives

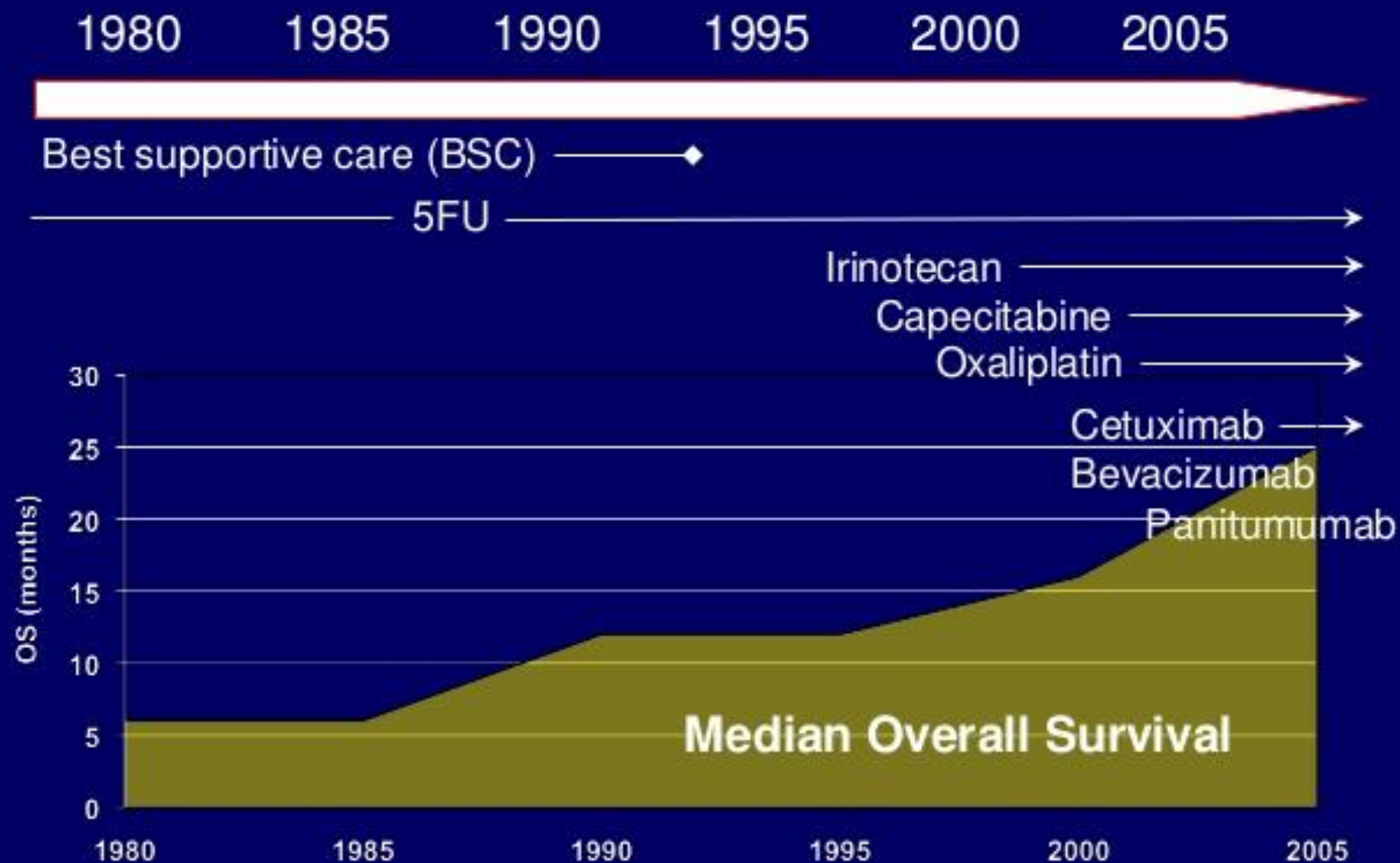
- Use of EGFR inhibitors in untreated pts with mCRC and CCO funding
- Prognostic and predictive value of primary tumor location
- Canadian Colorectal Cancer Sidedness Guidelines
- Biomarker role and impact in treatment selection for mCRC

# TOPIC ONE

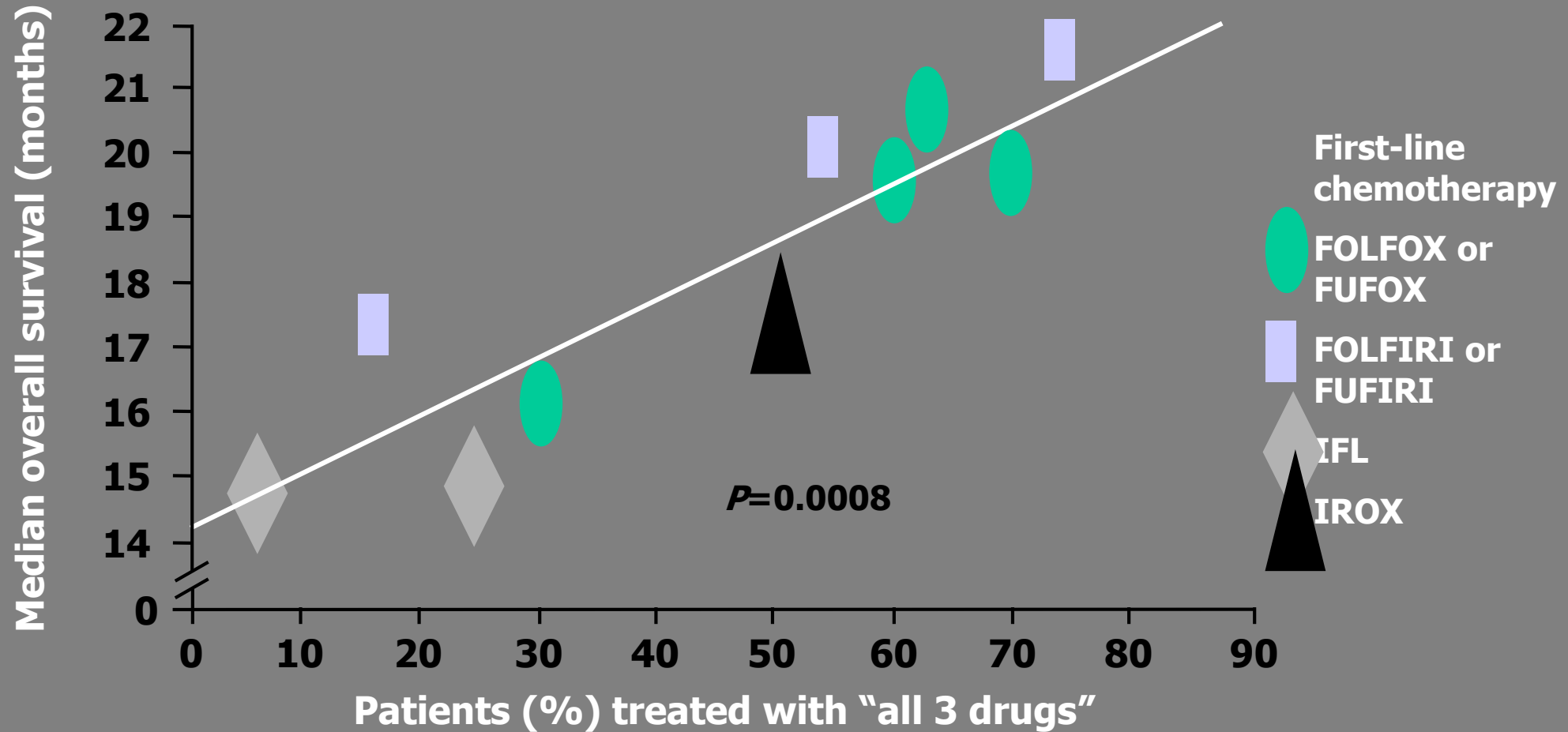
## The Evolution of Treatment in mCRC



# Advances in the Treatment of Stage IV CRC:



# THE VALUE OF CYTOTOXIC CHEMOTHERAPY: Correlation Between Exposure to “All-Three-Drugs” and Median Overall Survival



# Colorectal Cancer: 20 Years Later

meta-analysis 1992 80405 results

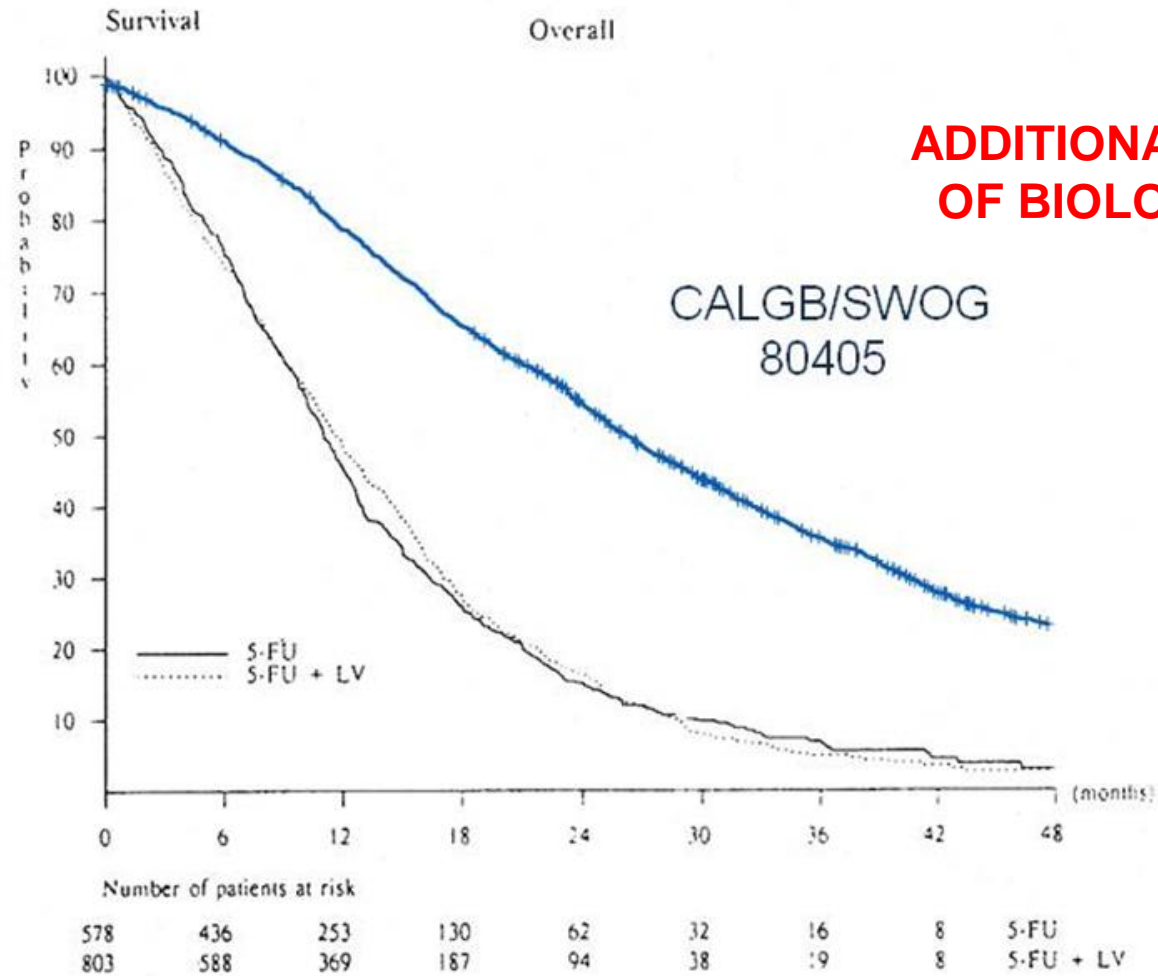


Fig 2. Overall survival. J Clin Oncol, 1992





## **TOPIC TWO**

Factors Influencing Treatment Selection

# PATIENT-FACTORS INFLUENCING Rx

- ECOG PS and Age
- Prior adjuvant FOLFOX
- Possible C/I to VEGF(R)-based Rx
- DYPD status (determines fluoropyrimidine tolerance)
- Goals and values
- Drug plan and perhaps home province
- Cognition and co-morbidities

# DISEASE-RELATED FACTORS INFLUENCING Rx

- Genetic status of the primary tumour: RAS, RAF, MSI
- Sidedness (Left vs Right)?
- Whether metastases are potentially resectable
- Primary in or out
- Primary in the rectum or colon

## OTHER FACTORS INFLUENCING Rx

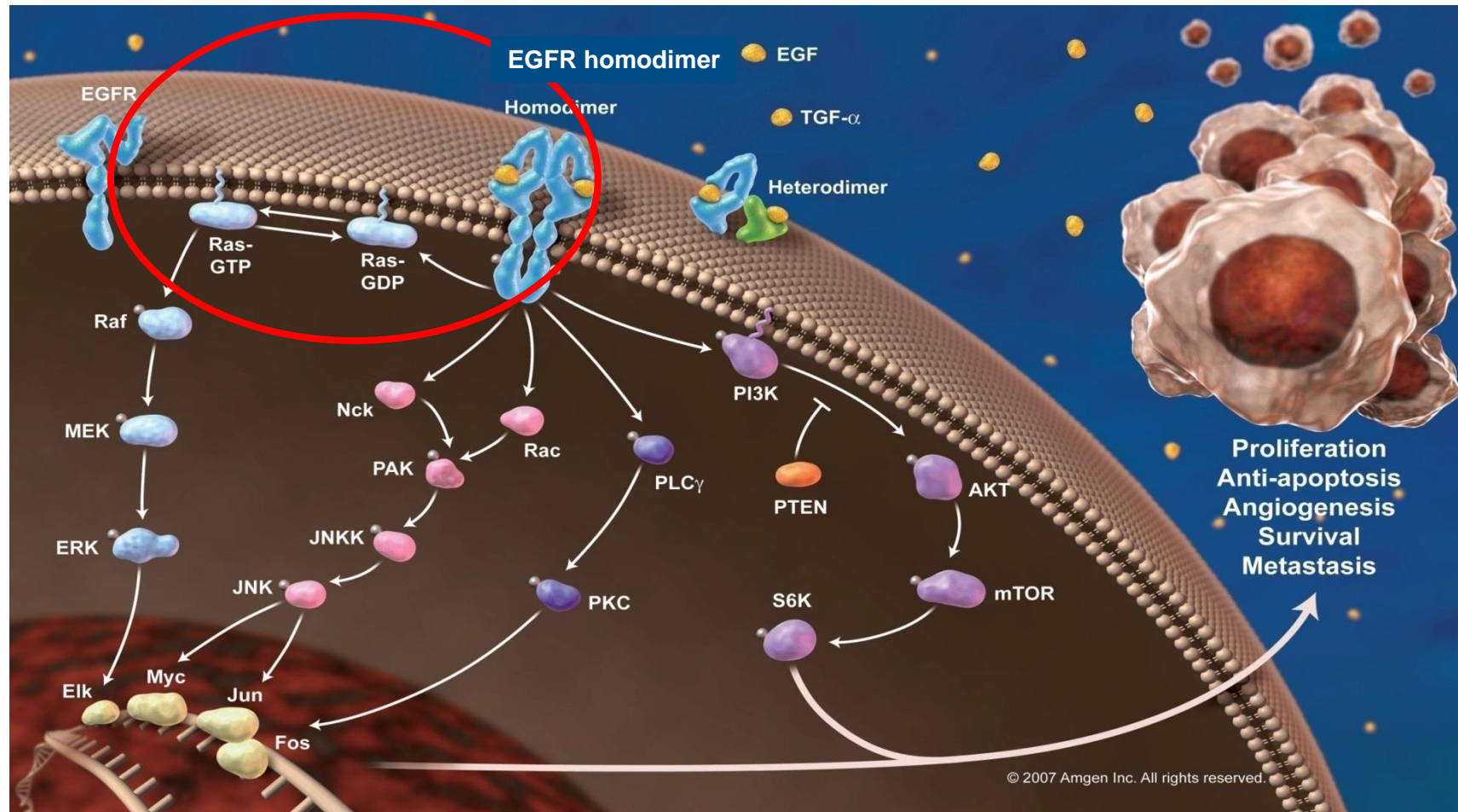
- Availability of a clinical trial
- Cost of drugs to the system (e.g. oxaliplatin now a 'we don't care' drug)

# TOPIC THREE

RAS and RAF



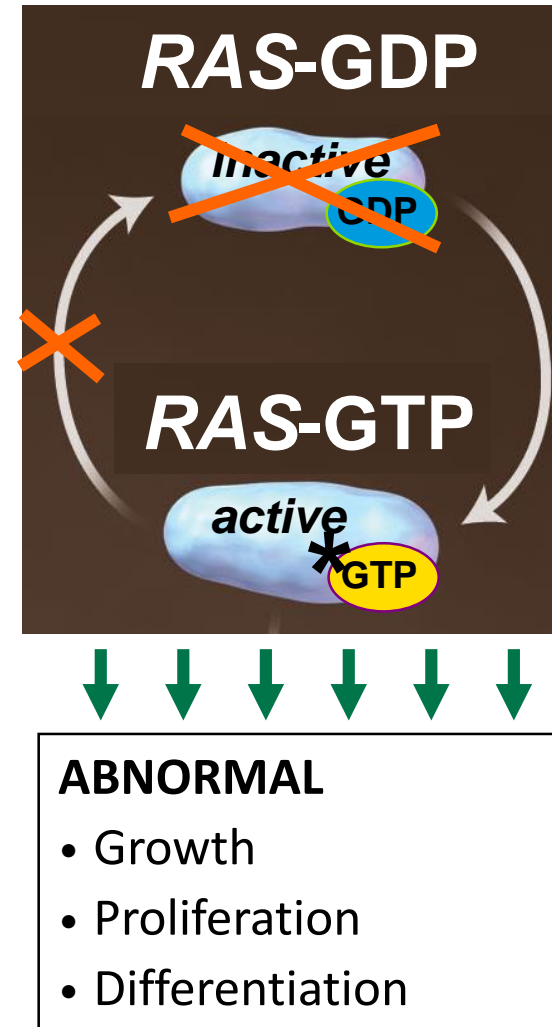
# EGFR Activation May Involve Downstream Signaling Pathways That Include *RAS*



Berg M, et al. *Discov Med* 2012;14:207-14; Di Fiore F, et al. *Br J Cancer* 2010;103:1765-72; Han W, et al. *Cancer Lett* 2012;318:124-34; Herbst RS, et al. *Cancer* 2002;94:1593-1611.

# Mutated *RAS* May Lead to Uncontrolled Cell Signaling and Cancer

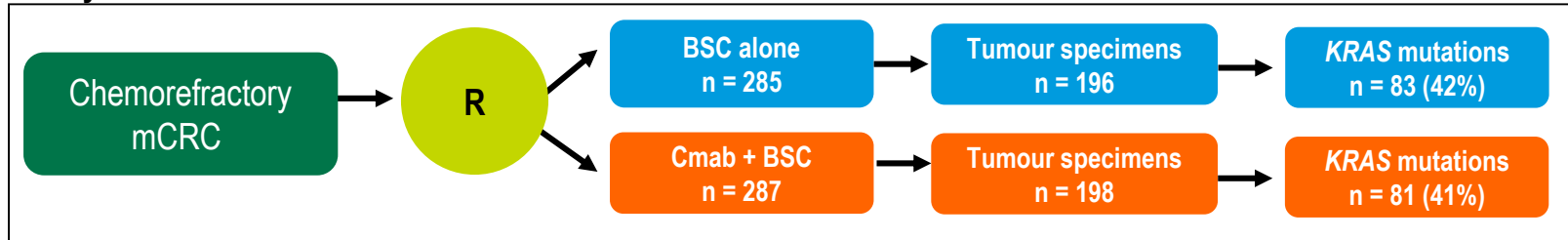
- Mutations in *KRAS* or *NRAS* are found in approximately 53% of CRCs<sup>1</sup>
- Specific mutations in *RAS* genes result in constitutively active proteins
- Patients with *KRAS* or *NRAS* mutations do not benefit from EGFR inhibition<sup>1-3</sup>



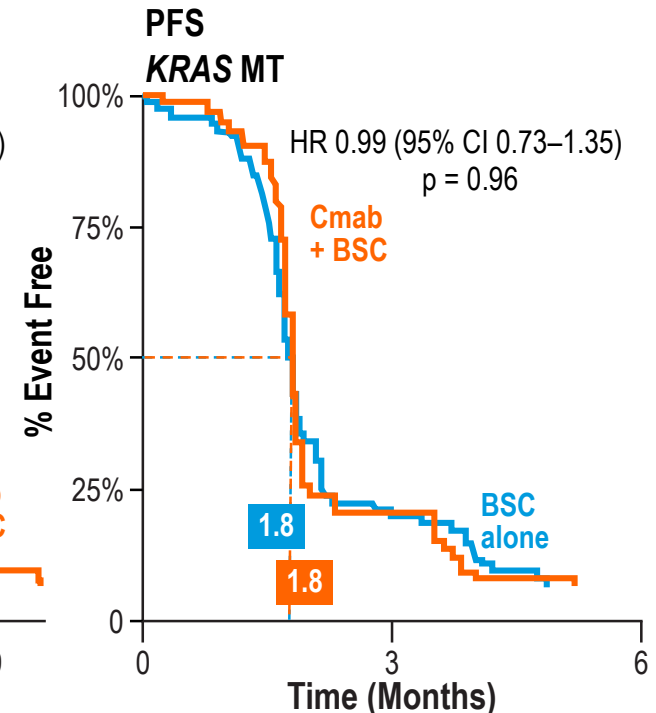
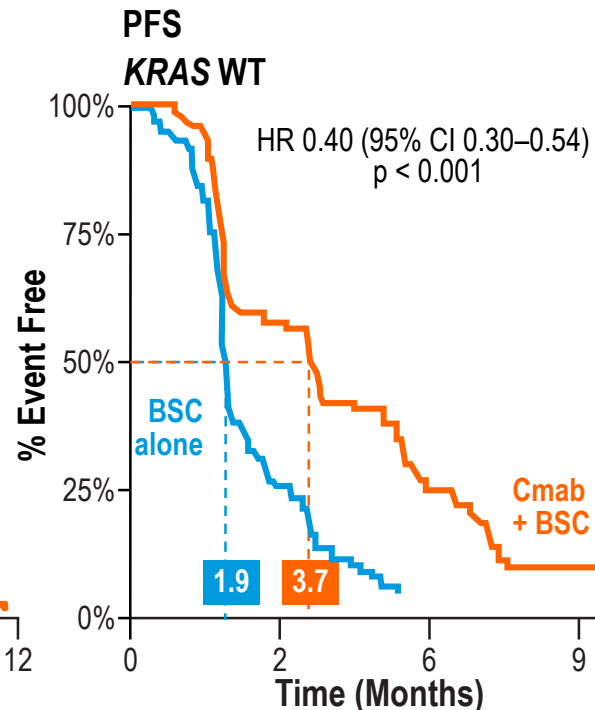
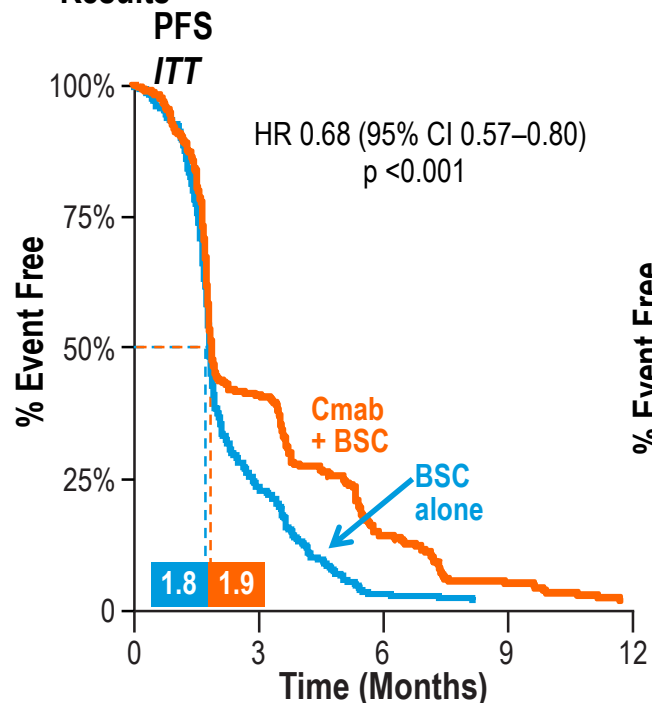
# NCIC CO.17 Trial

## Cetuximab and *KRAS* exon 2 Status

### Study Schema



### Results

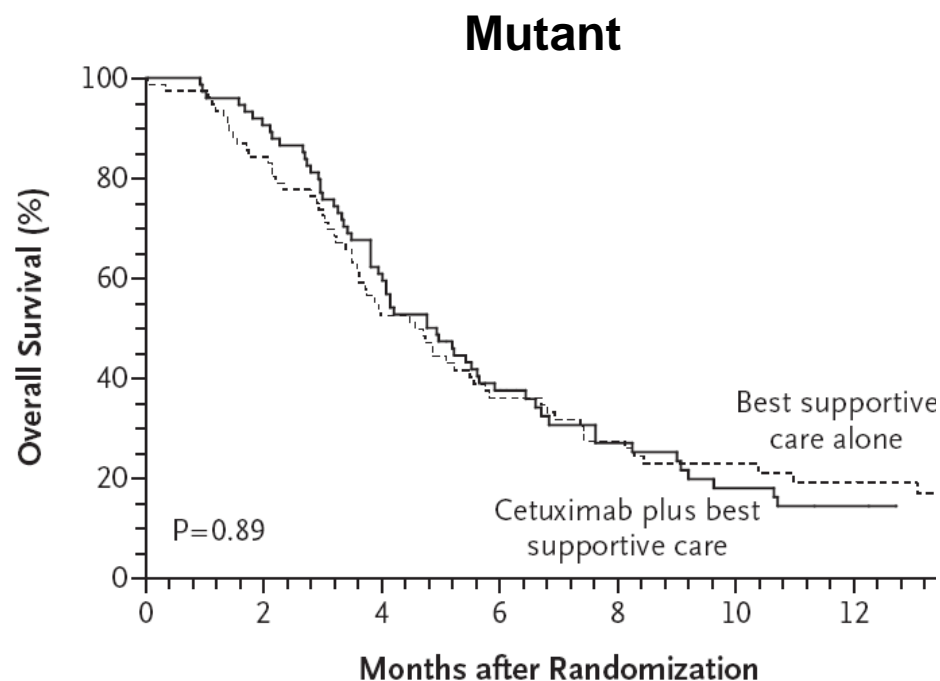


Adapted from Jonker et al. *N Engl J Med* 2008;357:2040-2048.  
Cmab = Cetuximab; NCIC = National Cancer Institute of Canada

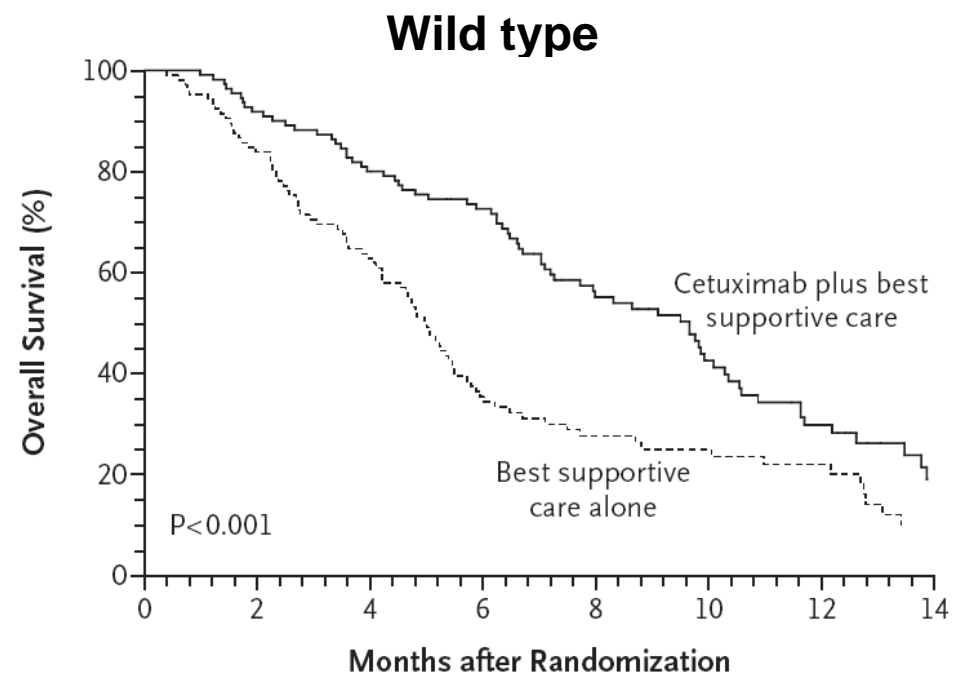
Adapted from Karapetis C, et al. *N Engl J Med* 2008;359:1757-1765.

# KRAS Mutational Status Associated with Outcome: Predictive Biomarker (but not Prognostic) – NCIC CO.17

## Overall Survival



| Study arm       | Median OS (months) | 95% CI    |
|-----------------|--------------------|-----------|
| Cetuximab + BSC | 4.5                | 3.8 – 5.6 |
| BSC alone       | 4.6                | 3.6 – 5.5 |

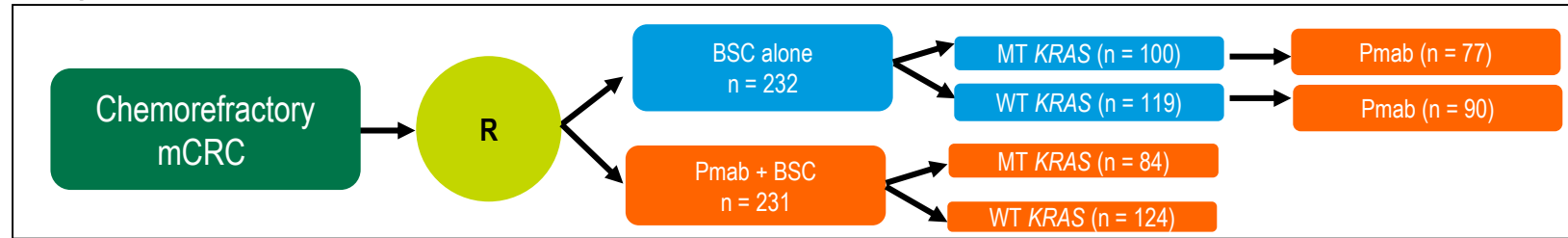


| Study arm       | Median OS (months) | 95% CI     |
|-----------------|--------------------|------------|
| Cetuximab + BSC | 9.5                | 7.7 – 10.3 |
| BSC alone       | 4.8                | 4.2 – 5.5  |

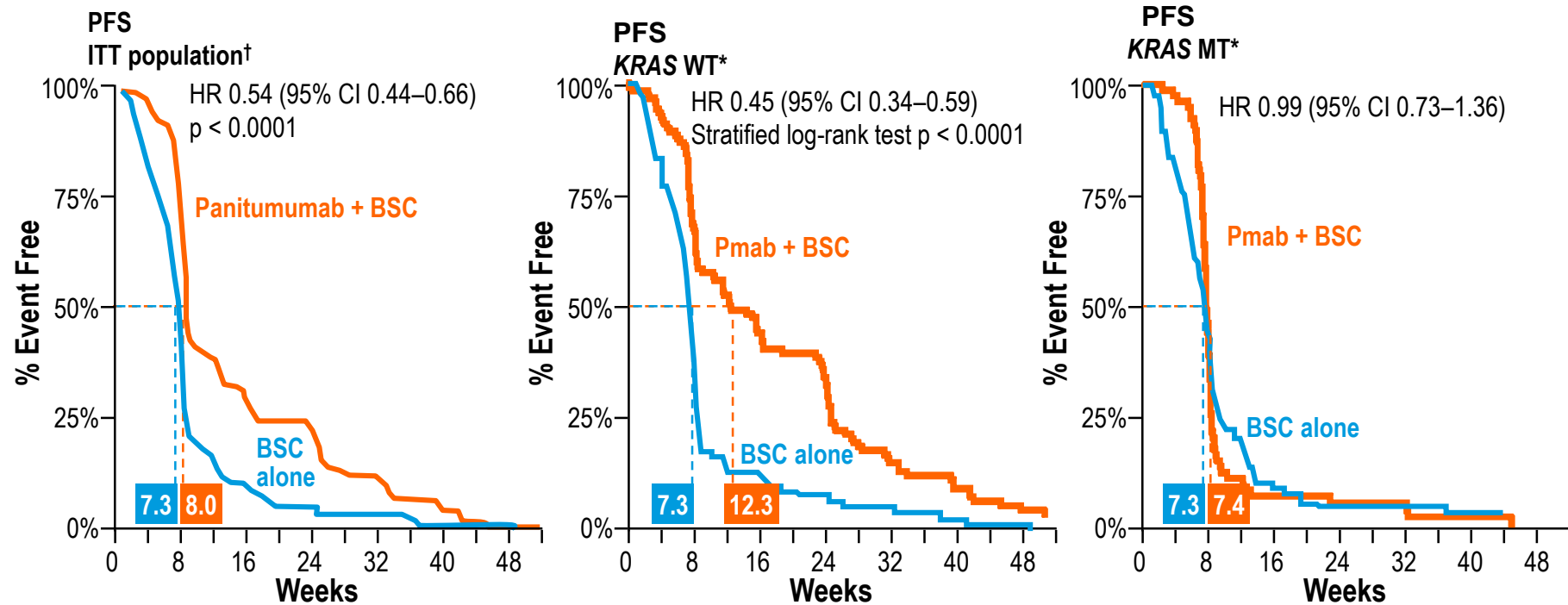
# 408 Study

## Panitumumab and *KRAS* Exon 2 Status

### Study Schema



### Results



BSC = Best supportive care; MT = Mutant; Pmab = Panitumumab; WT = Wild type

\*Adapted from Amado RG, et al. *J Clin Oncol* 2008;26:1626-1634.

†Adapted from Van Cutsem, et al. *J Clin Oncol* 2007;25:1658-1664.



# KRAS Exon 2 Status Predicts Response to EGFR Inhibitors in Most Studies

**NORDIC VII is the only study in which the results with regard to KRAS status were not clear in patients with wild-type tumours.**

| Study       |                           | n     | Key Results   | Improved Outcomes in KRAS WT patients |
|-------------|---------------------------|-------|---|---------------------------------------|
| First-line  | CRYSTAL <sup>1</sup>      | 1,198 | FOLFIRI + cetuximab > FOLFIRI   | ✓                                     |
|             | COIN <sup>2</sup>         | 1,630 | 5-FU/capecitabine + oxaliplatin + cetuximab > 5-FU/capecitabine + oxaliplatin | ✓                                     |
|             | PRIME <sup>3</sup>        | 1,183 | FOLFOX4 + panitumumab > FOLFOX4   | ✓                                     |
|             | NORDIC VII <sup>4</sup>   | 566   | FLOX = FLOX + cetuximab   | ✗                                     |
| Second-line | EPIC <sup>5,6</sup>       | 1,298 | Irinotecan + cetuximab > Irinotecan   | ✓                                     |
|             | 181 <sup>7</sup>          | 1,186 | FOLFIRI + panitumumab > FOLFIRI   | ✓                                     |
| Third-line  | NCIC CO.17 <sup>8,9</sup> | 572   | Cetuximab + BSC > BSC   | ✓                                     |
|             | 408 <sup>10, 11</sup>     | 463   | Panitumumab + BSC > BSC   | ✓                                     |
|             | FIRE-3 <sup>12</sup>      | 592   | Cetuximab + FOLFIRI > Bevacizumab + FOLFIRI                                   | ✓                                     |
|             | CALGB/SWOG 80405          | 526   | Cetuximab + Chemo = Bevacizumab + Chemo                                       | ✓                                     |

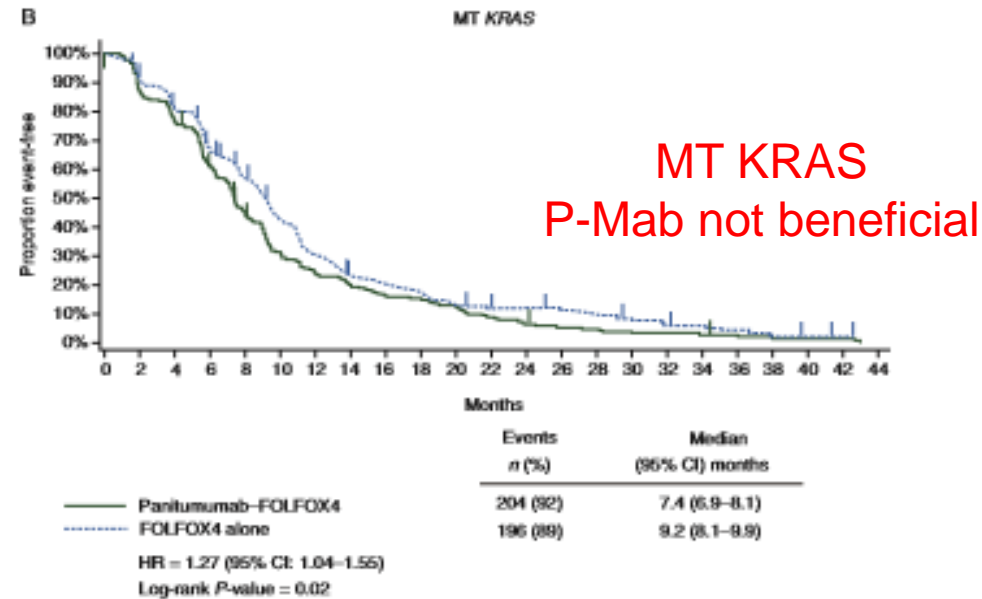
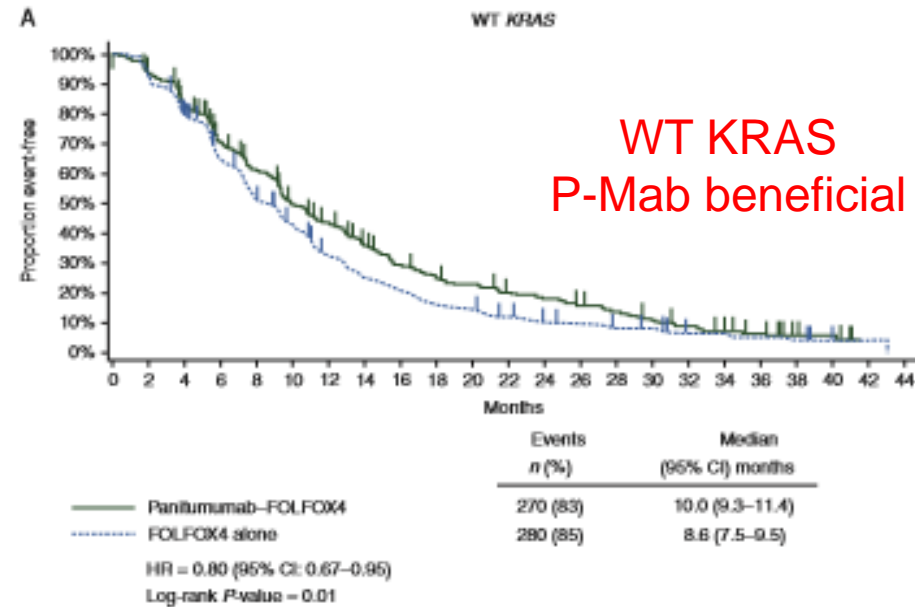
BSC = Best supportive care; FLOX = Fluorouracil (5-FU), leucovorin (LV), and oxaliplatin; FOLFIRI = 5-FU/LV + irinotecan; FOLFOX = 5-FU + LV + oxaliplatin

1. Van Cutsem E, et al. *J Clin Oncol* 2011;29:2011; 2. Maughan TS, et al. *Lancet* 2011;377:2103-2114; 3. Douillard JY, et al. *J Clin Oncol* 2010;28:4697-4705; 4. Tveit K, et al. *J Clin Oncol* 2012;30:1755-62; 5. Sobrero AF, et al. *J Clin Oncol* 2008;26:2311-2319; 6. Langer C, et al. *Ann Oncol* 2008;19(suppl 8) (Abstr 385P); 7. Peeters M, et al. *J Clin Oncol* 2010;28:4706-4713; 8. Karapetis C, et al. *N Engl J Med* 2008;359:1757-1765; 9. Jonker DJ, et al. *N Engl J Med* 2007;357:2040-2048; 10. Van Cutsem E, et al. *J Clin Oncol* 2007;25:1658-1664; 11. Amado RG, et al. *J Clin Oncol* 2008;26:1626-1634. 12. Heinemann V, et al. *Lancet Oncol* 2014;15:1065-1075. 13. Venook A. ASCO 2014. *J Clin Oncol* 2014;32:5s(suppl). Abstr LBA3; 14. Lenz H. *Ann Oncol* 2014;25 (suppl 4). Abstr 501O.

PFS the primary  
endpoint: met

# PRIME: ANALYSIS A/C to KRAS STATUS

## PRIME: FOLFOX ± P-MAb



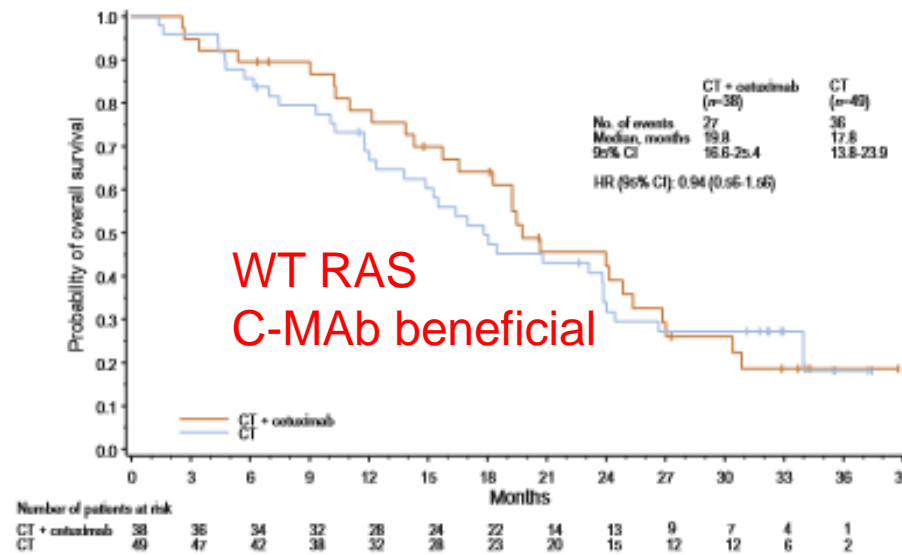
**Figure 1.** (A) Kaplan-Meier plot of PFS (WT KRAS, Panitumumab-FOLFOX4 versus FOLFOX4 alone). (B) Kaplan-Meier plot of PFS (MT KRAS, Panitumumab-FOLFOX4 versus FOLFOX4 alone). (C) PFS Forest plot—WT KRAS efficacy analysis set.

Overall Survival

THE SAME PATTERN  
SEEN WITH CETUXIMAB

OPUS:  
FOLFOX ±  
C-MAb

A. RAS wild-type (all loci)



B. RAS mutation (any locus)

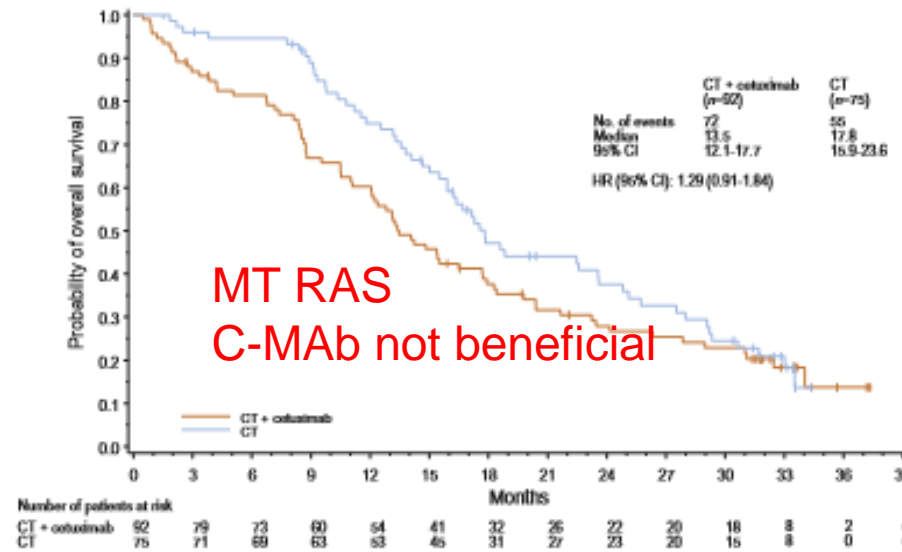
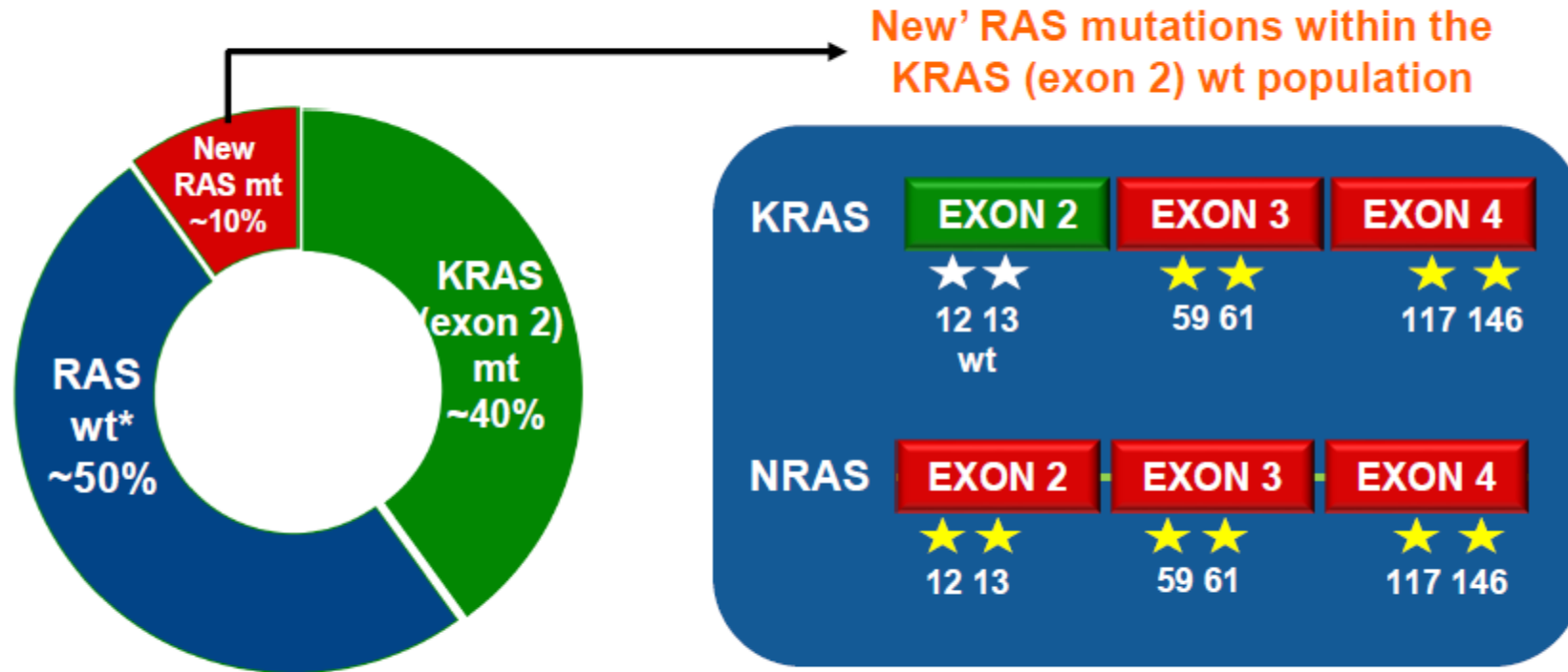


Fig. 3. Overall survival for patients with RAS wild-type (A) and any RAS mutant (B) tumours. CT, chemotherapy; HR, hazard ratio; NR, not reached.

# EXPANDED RAS ANALYSIS

## Beyond KRAS (exon 2): Additional RAS mutations in mCRC

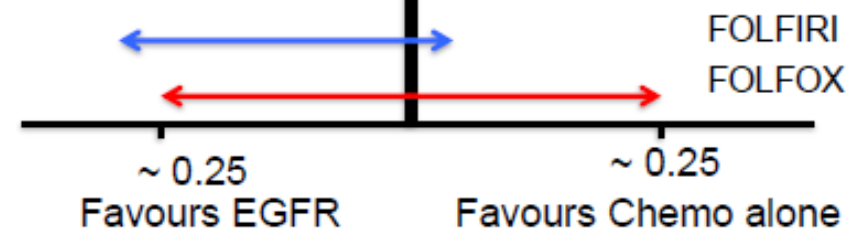


\*Wt at KRAS exons 2, 3, 4 and NRAS exons 2, 3, 4

# MEASURING EXPANDED RAS PROVIDES MORE BENEFIT FROM EGFR MABS THAN KRAS

## Influence of KRAS and RAS mutational status on survival Randomised trials of EGFR antibodies – 1<sup>st</sup> line

| Trial                     | Therapy                           | OS (mo)<br>KRAS <sup>wt</sup> |       | OS (mo)<br>RAS <sup>wt</sup> |       | OS (mo)<br>RAS <sup>mut</sup> |       |
|---------------------------|-----------------------------------|-------------------------------|-------|------------------------------|-------|-------------------------------|-------|
|                           |                                   | CTx                           | +EGFR | CTx                          | +EGFR | CTx                           | +EGFR |
| <b>CRYSTAL</b><br>(n=666) | FOLFIRI<br>+/- Cetux <sup>*</sup> | HR 0.80<br>0.67-0.95          |       | HR 0.69<br>0.54-0.88         |       | HR 1.05<br>0.85-1.25          |       |
| <b>PRIME</b><br>(n=656)   | FOLFOX<br>+/- Panit <sup>*</sup>  | HR 0.83<br>0.67-1.02          |       | HR 0.78<br>0.62-0.99         |       | HR 1.25<br>1.02-1.55          |       |
| <b>OPUS</b><br>(n=197)    | FOLFOX<br>+/- Cetux <sup>*</sup>  | HR 0.86<br>0.60-1.22          |       | HR 0.83<br>0.49-1.42         |       | HR 1.35<br>0.95-1.92          |       |



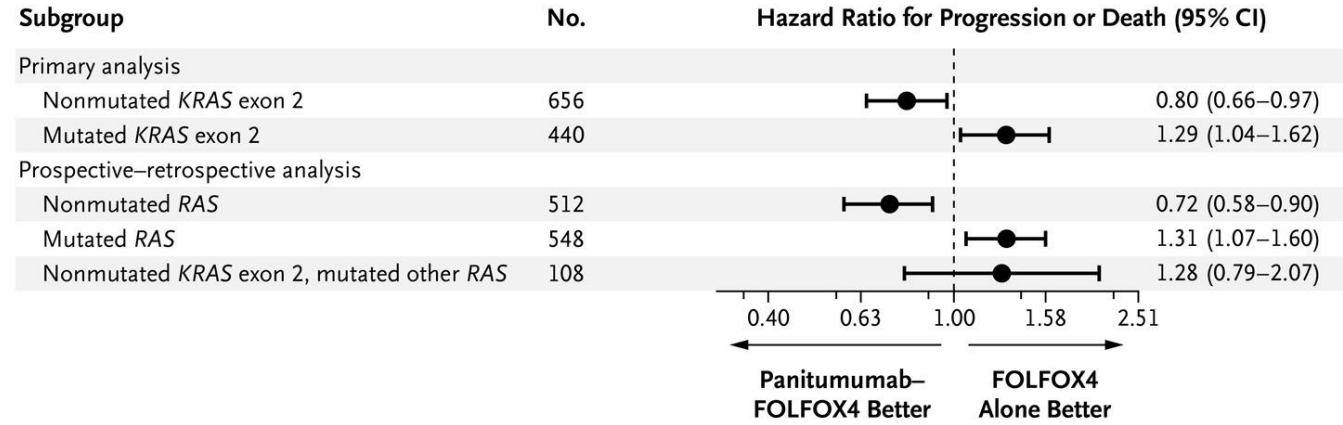
Van Cutsem, Ciardiello, Köhne et al. ASCO 2014  
 Douillard et al. NEJM 2014  
 Bokemeyer, Köhne et al. 2014 ASCO 2014



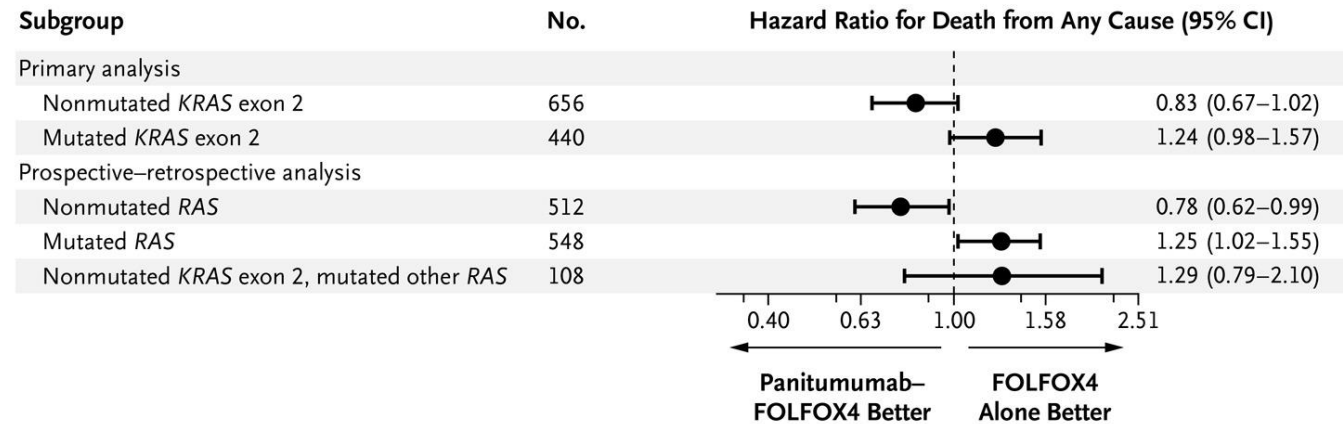
# PRIME

## Hazard Ratio for Disease Progression or Death and Hazard Ratio for Death from Any Cause, According to *KRAS* and *RAS* Mutation Status.

### A Progression-free Survival



### B Overall Survival



Douillard J-Y et al. N Engl J Med 2013;369:1023-1034

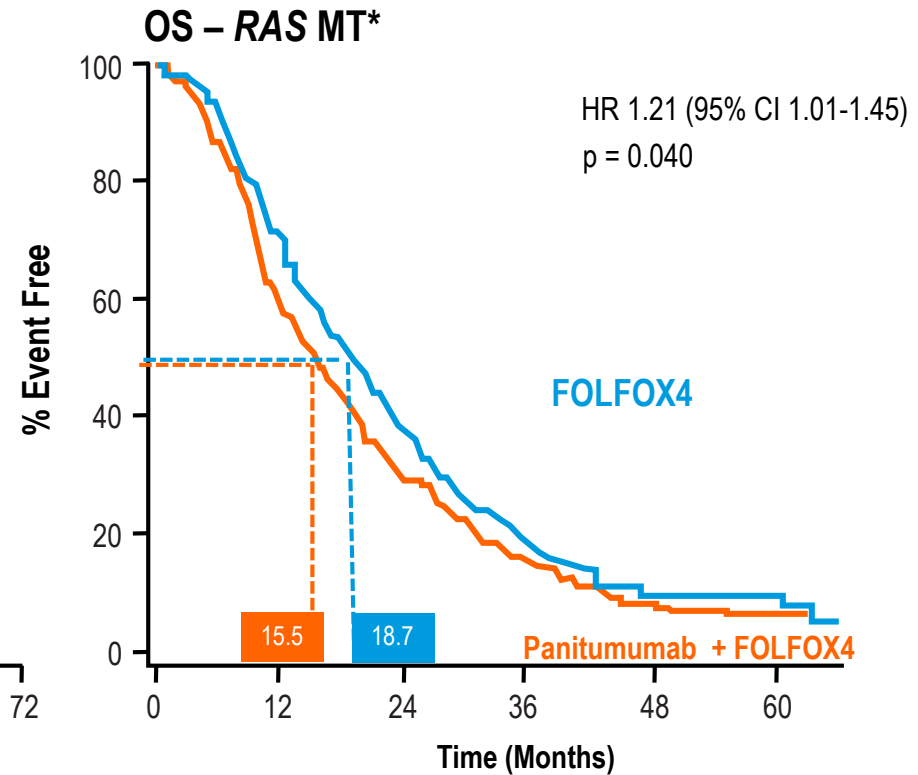
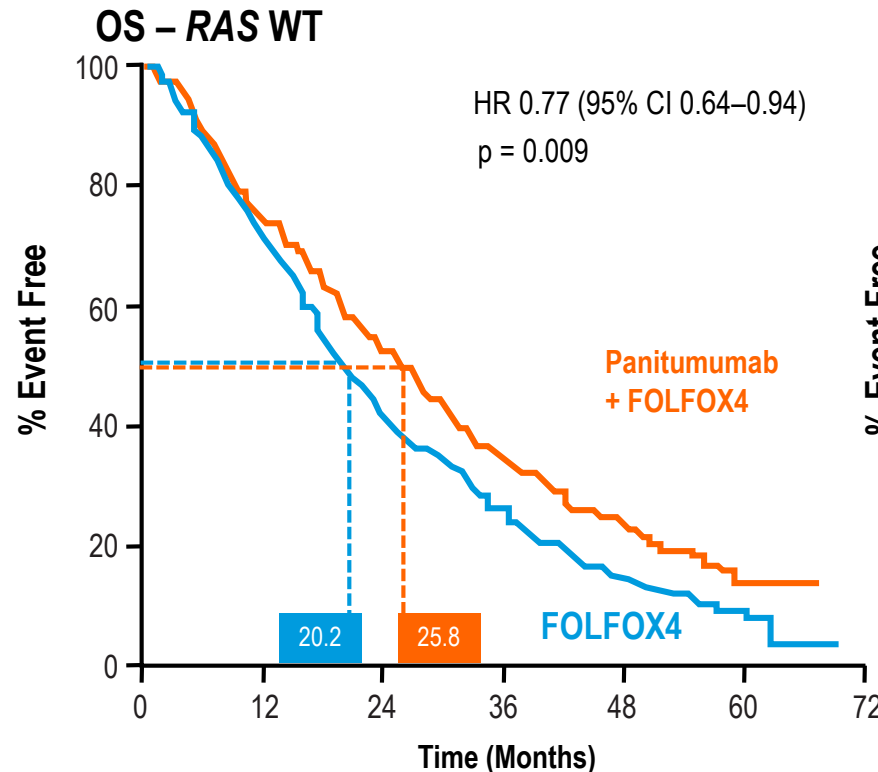
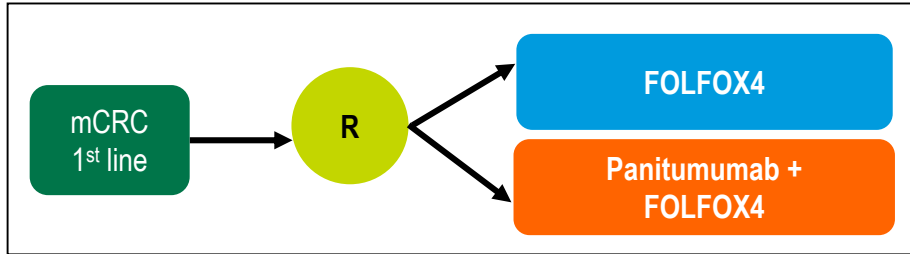


The NEW ENGLAND  
JOURNAL of MEDICINE

# PRIME Phase III Study

RAS Status<sup>†</sup>: Panitumumab + FOLFOX4 vs. FOLFOX Alone (Updated Analysis<sup>‡</sup>)

## Study Schema



\*KRAS exon 3, 4 ; NRAS exon 2, 3, 4; or BRAF exon 15;

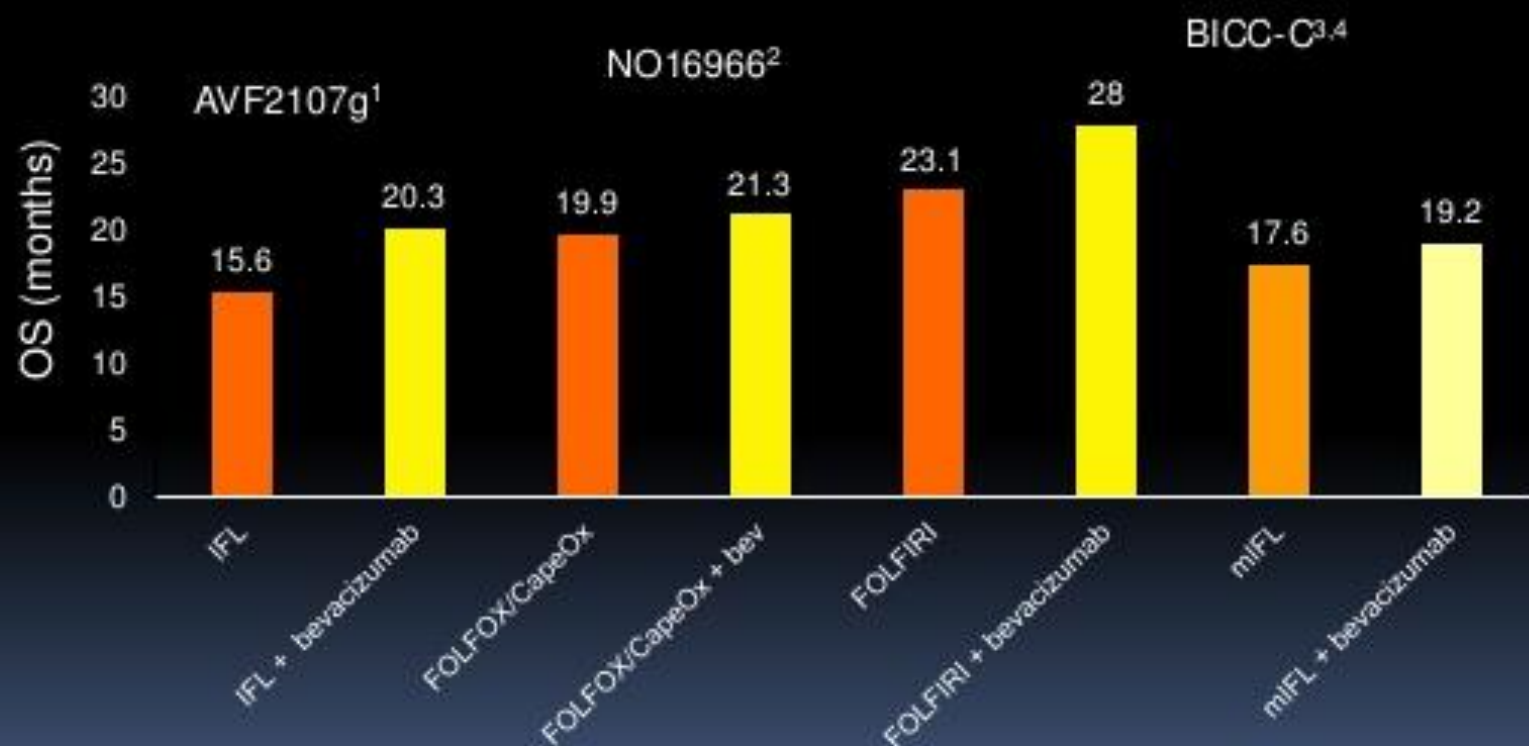
<sup>†</sup>Prospective-retrospective analysis; <sup>‡</sup>Exploratory analysis

Adapted from Douillard JY, et al. *N Engl J Med* 2013;369:1023-1034.

## **TOPIC FOUR**

**WHICH BIOLOGICAL: BEV OR anti-EGFR?**

# First-Line Bevacizumab in mCRC: Overall Survival



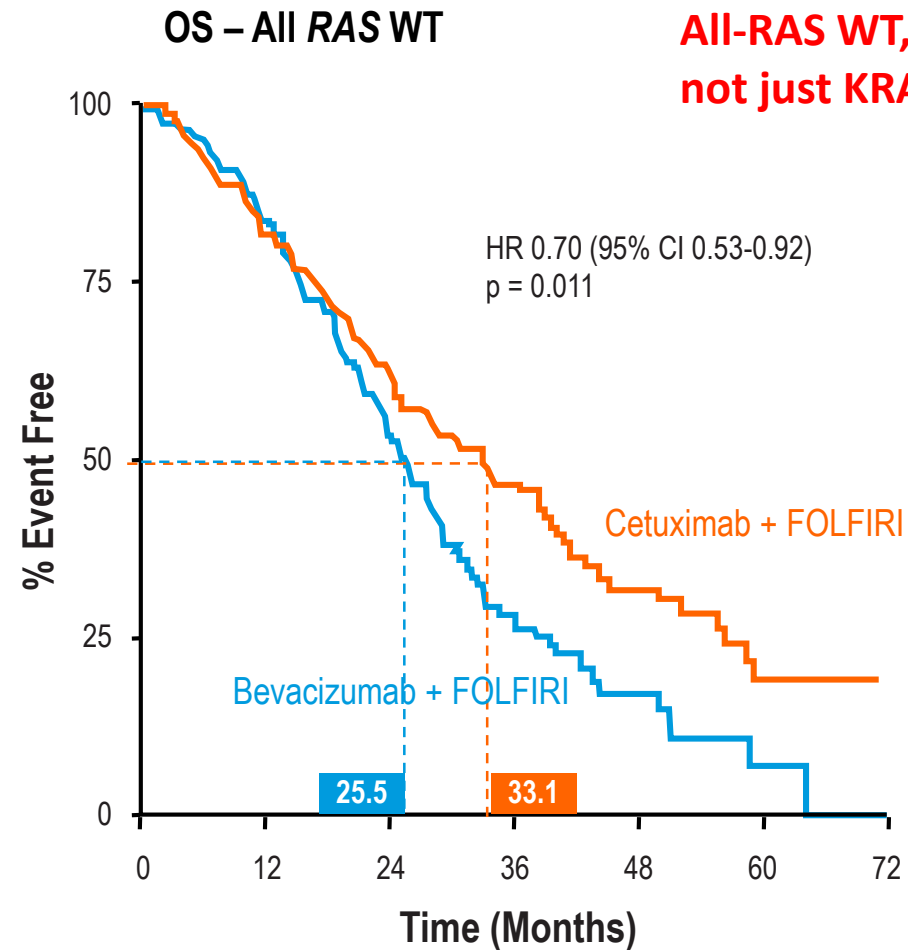
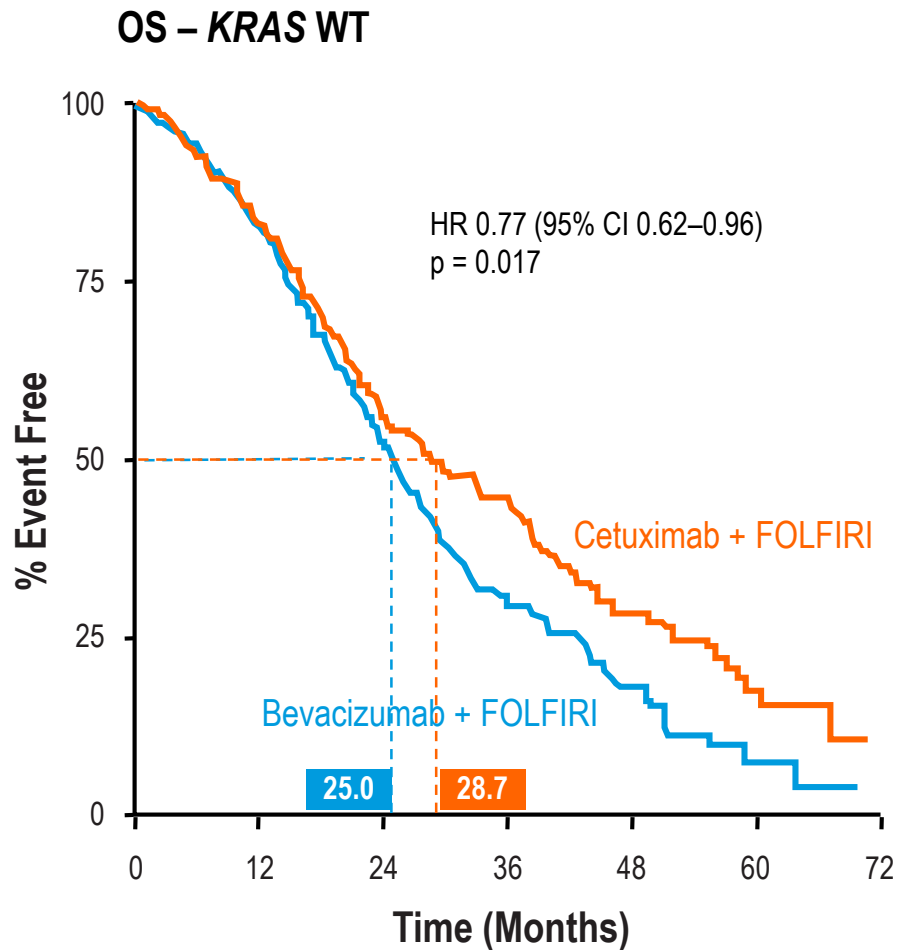
\* $P < 0.001$ ; † $P = 0.0769$ .

1. Hurwitz H et al. *N Engl J Med*. 2004;350:2335-2342; 2. Saltz LB et al. *J Clin Oncol*. 2008;26:2013-2019;

3. Fuchs C et al. *J Clin Oncol*. 2007;25:4779-4786; 4. Fuchs C et al. *J Clin Oncol*. 2008;26:689-690;

# FIRE-3 Phase III Trial: Cetuximab vs. Bevacizumab with FOLFIRI as First-line Treatment in *KRAS* WT Patients

**EGFR MAb better than  
VEGF MAb, especially in  
All-RAS WT,  
not just *KRAS* WT**

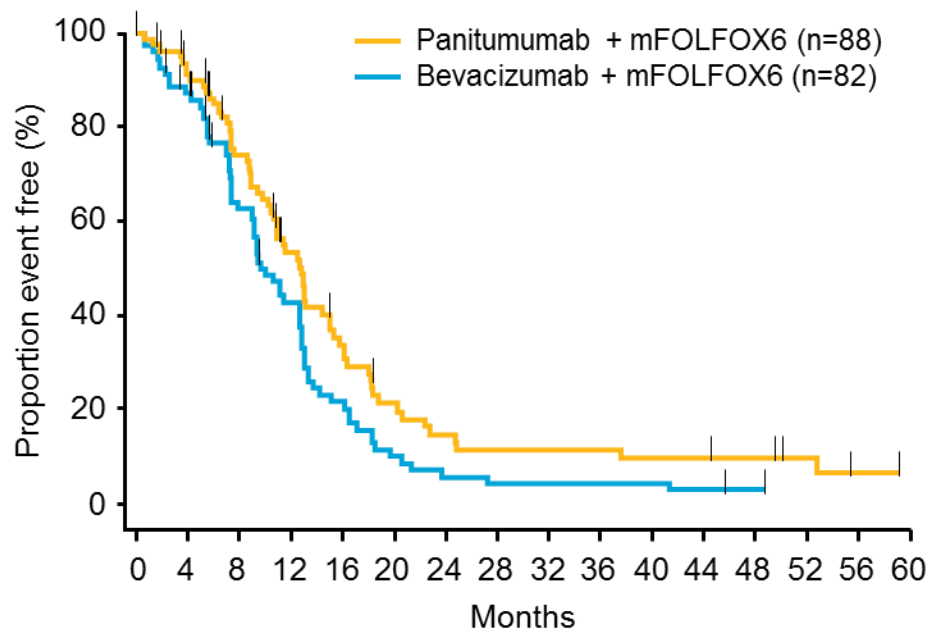


# PFS – RAS population

**PEAK:  
FOLFOX/PMAb  
VS  
FOLOFOX/BEV**

## RAS WT

|                              | Panitumumab +<br>mFOLFOX6 (n=88) | Bevacizumab +<br>mFOLFOX6 (n=82) |
|------------------------------|----------------------------------|----------------------------------|
| Median PFS, (95% CI), months | 12.8 (10.7, 15.1)                | 10.1 (9.0, 12.7)                 |
| HR (95% CI)                  | 0.68 (0.48, 0.96)                |                                  |
| p-value                      | 0.029                            |                                  |

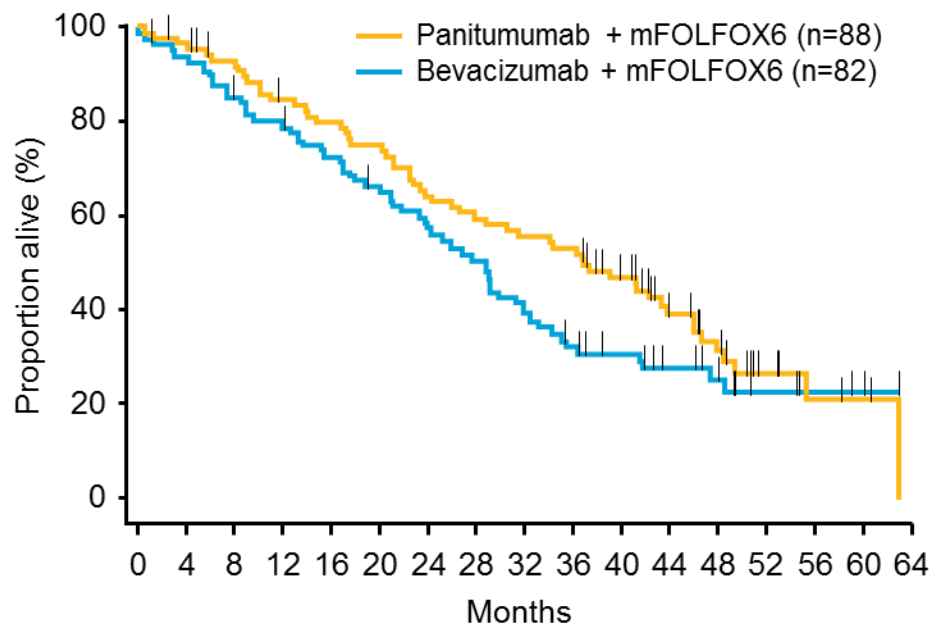


|       |    |    |    |    |    |    |   |   |   |   |   |   |   |   |   |   |
|-------|----|----|----|----|----|----|---|---|---|---|---|---|---|---|---|---|
| Pmab* | 88 | 72 | 55 | 36 | 22 | 13 | 9 | 7 | 7 | 7 | 6 | 6 | 5 | 3 | 1 | 0 |
| Bmab* | 82 | 68 | 45 | 30 | 14 | 7  | 4 | 3 | 3 | 3 | 3 | 2 | 1 | 0 | 0 | 0 |

# OS – RAS WT population

## RAS WT

|                             | Panitumumab +<br>mFOLFOX6 (n=88) | Bevacizumab +<br>mFOLFOX6 (n=82) |
|-----------------------------|----------------------------------|----------------------------------|
| Median OS, (95% CI), months | 36.9 (27.9, 46.1)                | 28.9 (23.3, 32.0)                |
| HR (95% CI)                 | 0.76 (0.53, 1.11)                |                                  |
| p-value                     | 0.15                             |                                  |



|       |    |    |    |    |    |    |    |    |    |    |    |    |    |   |   |   |   |
|-------|----|----|----|----|----|----|----|----|----|----|----|----|----|---|---|---|---|
| Pmab* | 88 | 84 | 78 | 70 | 66 | 62 | 53 | 49 | 46 | 44 | 34 | 22 | 15 | 7 | 3 | 2 | 0 |
| Bmab* | 82 | 76 | 69 | 63 | 57 | 51 | 44 | 39 | 31 | 23 | 19 | 14 | 10 | 5 | 3 | 2 | 0 |

**PEAK:  
FOLFOX/PMAb  
Vs  
FOLOFOX/BEV**

# Tumour response-related efficacy – RAS WT population

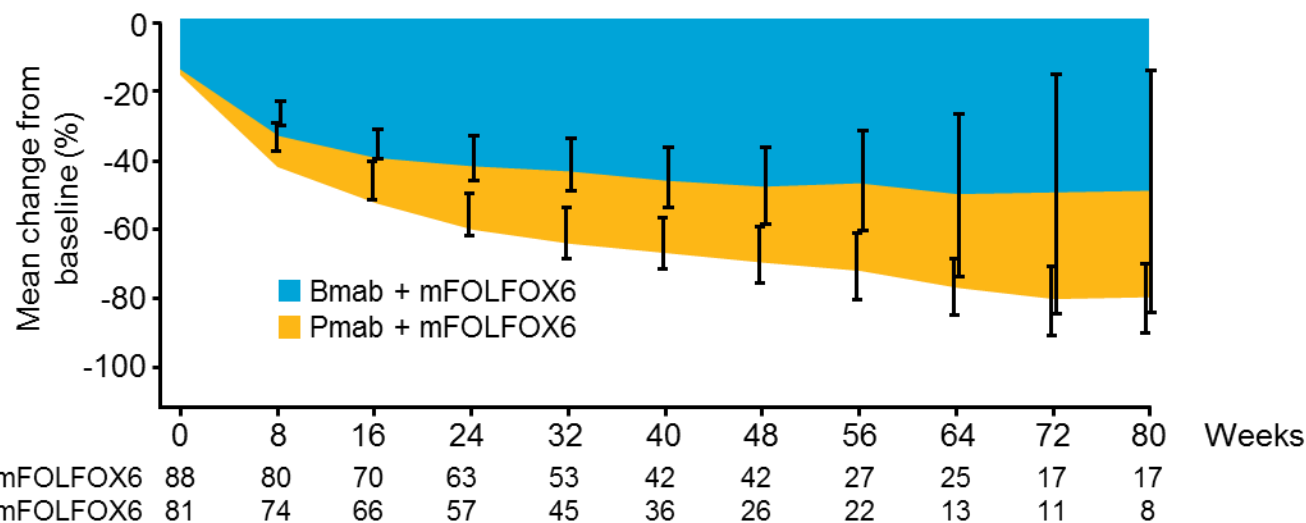
|                             | Pmab + mFOLFOX6 (n=88)   | Bmab + mFOLFOX6 (n=82) |
|-----------------------------|--------------------------|------------------------|
| Median DoR, months (95% CI) | 11.4 (10.0, 16.3)        | 9.0 (7.6, 9.5)         |
| HR (95% CI); p-value        | 0.59 (0.39, 0.88); 0.011 |                        |
| Median TTR, months (95% CI) | 2.3 (1.9, 3.7)           | 3.8 (2.1, 5.7)         |
| HR (95% CI); p-value        | 1.19 (0.81, 1.74); 0.37  |                        |
| Median DpR, months (Q1, Q3) | 65.0 (45.7, 89.5)        | 46.3 (29.5, 63.3)      |
| p-value                     | 0.0018                   |                        |

**Longer**

**Quicker**

**Deeper**

Mean (95% CI) percentage change from baseline in tumour load<sup>a</sup> over time



<sup>a</sup>Sum of all target lesions - includes those patients evaluable for objective response who had baseline tumour shrinkage data

DoR = duration of response;  
DpR = depth of response;  
ETS = early tumour shrinkage;  
TTR = time to response



## THE BENEFITS OF ACHIEVING A PARTIAL RESPONSE

### Between-treatment comparisons of PFS and OS outcomes by tumour shrinkage at week 8 – RAS WT population

|   | Tumour shrinkage at week 8 |                        |                            |                        |  |
|---|----------------------------|------------------------|----------------------------|------------------------|--|
|   | <30                        |                        | ≥30                        |                        |  |
|   | Panitumumab + mFOLFOX6     | Bevacizumab + mFOLFOX6 | Panitumumab + mFOLFOX6     | Bevacizumab + mFOLFOX6 |  |
| n (%)   | 29 (36)                    | 41 (55)                | 51 (64)                    | 33 (45)                | More pts achieve 30%+ shrinkage on pmab      |
| Odds ratio <sup>a</sup> , (95% CI)<br>p-value | —                          |                        | 1.99 (0.99, 4.10)<br>0.052 |                        |  |
| Median PFS, months (95% CI)                   | 11.6 (7.5, 15.4)           | 9.7 (7.5, 12.9)        | 13.0 (10.9, 18.1)          | 11.1 (9.0, 16.6)       | Pts achieving 30%+ shrinkage have better PFS |
| HR (95% CI)<br>p-value                        | 0.79 (0.45, 1.38)<br>0.40  |                        | 0.74 (0.45, 1.23)<br>0.24  |                        |  |
| Median OS, months (95% CI)                    | 34.2 (17.5, 42.3)          | 23.9 (20.1, 29.0)      | 43.8 (36.4, 63.0)          | 35.1 (29.9, NE)        | Pts achieving 30%+ shrinkage have better OS  |
| HR (95% CI)<br>p-value                        | 0.75 (0.43, 1.31)<br>0.31  |                        | 0.77 (0.42, 1.42)<br>0.41  |                        |  |

***Both PFS and OS are longer if the tumor shrinkage exceeds 30% by week 8***

<sup>a</sup>Odds ratio is defined as the odds of having ≥30% tumour shrinkage in the panitumumab + mFOLFOX6 arm relative to the odds in the bevacizumab + mFOLFOX6 arm

# REMOVING BRAF PTS MEANS EVEN MORE BENEFIT FOR P-MAb

## Summary of PFS, OS and ORR results – RAS WT and RAS WT/BRAF WT populations

|                               | RAS WT                       |                              | RAS WT / BRAF WT             |                              |
|-------------------------------|------------------------------|------------------------------|------------------------------|------------------------------|
|                               | Pmab +<br>mFOLFOX6<br>(n=88) | Bmab +<br>mFOLFOX6<br>(n=82) | Pmab +<br>mFOLFOX6<br>(n=77) | Bmab +<br>mFOLFOX6<br>(n=79) |
| <b>PFS</b>                    |                              |                              |                              |                              |
| Patients with event, n (%)    | 64 (73)                      | 70 (85)                      | 55 (71)                      | 67 (85)                      |
| Median, months<br>(95% CI)    | 12.8<br>(10.7, 15.1)         | 10.1<br>(9.0, 12.7)          | 13.1<br>(11.6, 16.2)         | 10.1<br>(9.0, 12.7)          |
| HR (95% CI); p-value          | 0.68 (0.48, 0.96); 0.029     |                              | 0.61 (0.42, 0.88); 0.0075    |                              |
| <b>OS</b>                     |                              |                              |                              |                              |
| Patients with event, n (%)    | 57 (65)                      | 58 (71)                      | 48 (62)                      | 55 (70)                      |
| Median, months<br>(95% CI)    | 36.9<br>(27.9, 46.1)         | 28.9<br>(23.3, 32.0)         | 41.3<br>(31.6, 46.7)         | 28.9<br>(23.9, 33.1)         |
| HR (95% CI); p-value          | 0.76 (0.53, 1.11); 0.15      |                              | 0.70 (0.48, 1.04); 0.08      |                              |
| <b>ORR</b>                    |                              |                              |                              |                              |
| Responders, n (%)<br>[95% CI] | 57 (65)<br>[54, 75]          | 49 (60)<br>[49, 71]          | 49 (64)<br>[52, 74]          | 46 (59)<br>[47, 70]          |
| Odds ratio (95% CI); p-value  | 1.12 (0.56, 2.22); 0.86      |                              | 1.17 (0.58, 2.38); 0.76      |                              |

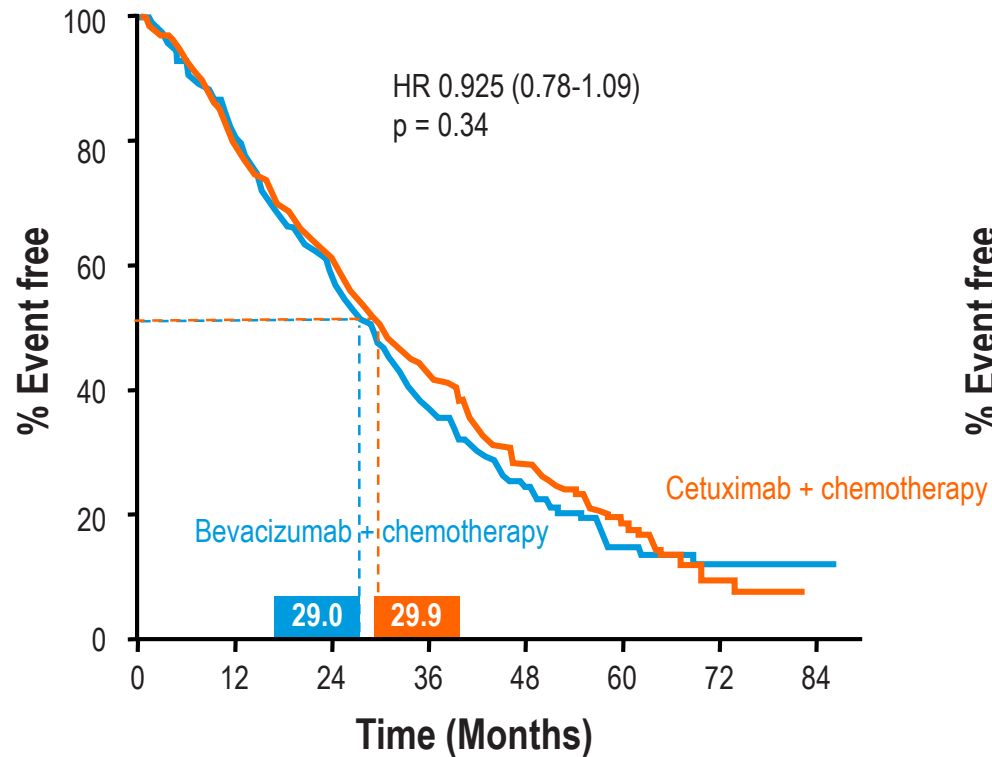
# Conclusions

- This final analysis of the PEAK trial found a significant improvement in PFS for patients with *RAS* WT mCRC receiving first-line treatment with panitumumab + mFOLFOX6 vs bevacizumab + mFOLFOX6 (HR: 0.68; p=0.029)
- Median OS was numerically longer in the panitumumab vs bevacizumab arm
  - 8.0 months longer in the *RAS* WT population (36.9 vs 28.9 months)
  - 12.4 months longer in the *RAS* WT/*BRAF*WT population (41.3 vs 28.9 months)
- ORR was similar between treatments, but tumour responses with panitumumab occurred earlier, lasted longer and were deeper vs bevacizumab
- Based on PEAK, panitumumab + mFOLFOX6 is an effective first-line treatment for patients with *RAS* WT mCRC

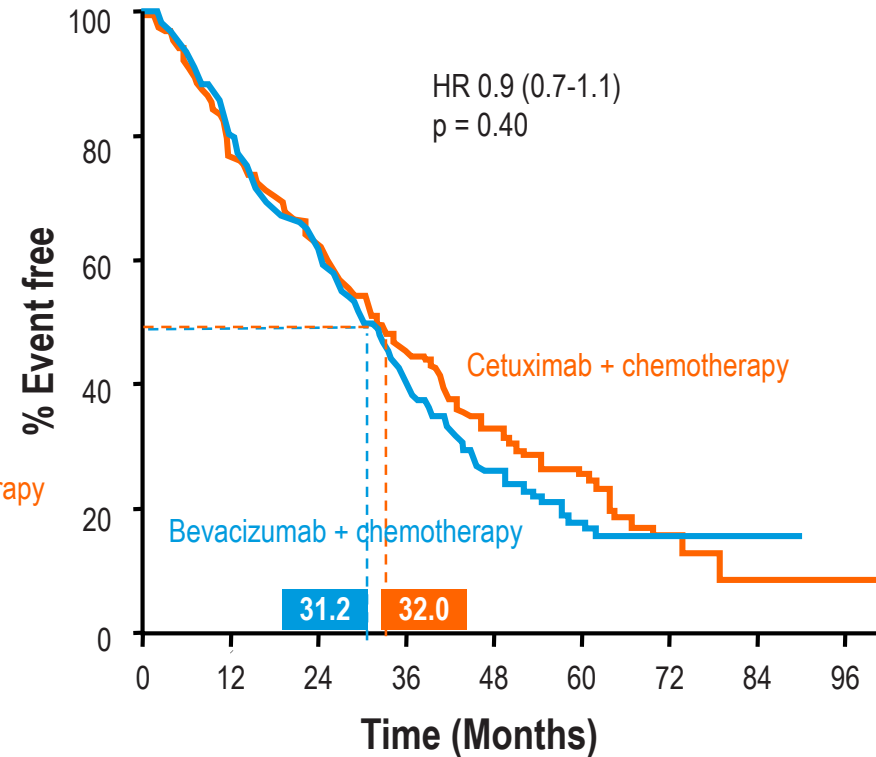
# CALGB/SWOG 80405:

Phase III trial of FOLFIRI or FOLFOX with Bevacizumab or Cetuximab for patients with *KRAS* exon 2 WT untreated mCRC

OS – *KRAS* WT

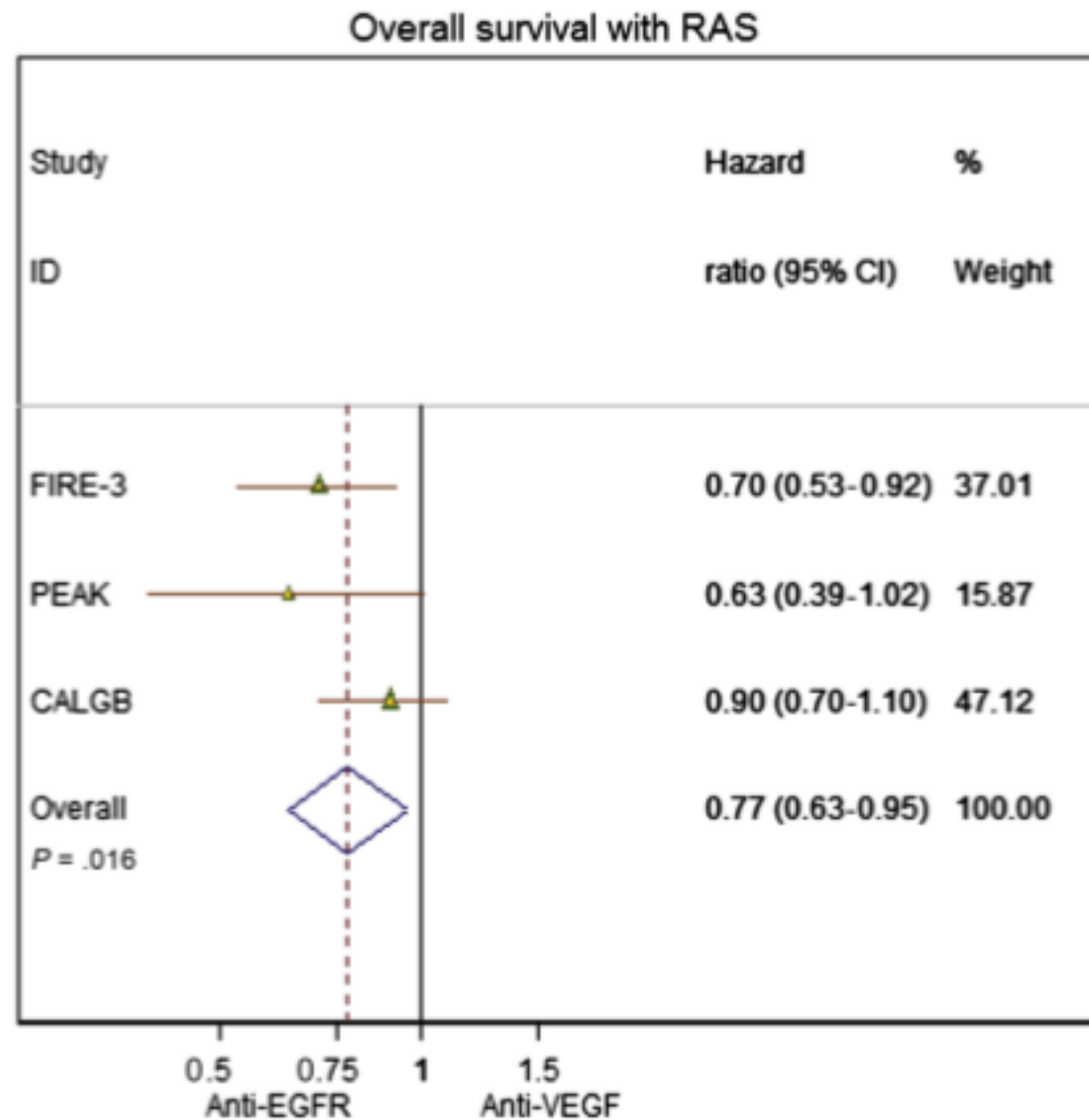


OS – All *RAS* WT



# Three major trials of CT + anti-EGFR Mab vs Bev a/c to ras

Figure 7 Forest Plot for Overall Survival (OS) for All RAS Wild Type (RAS-WT) Patients

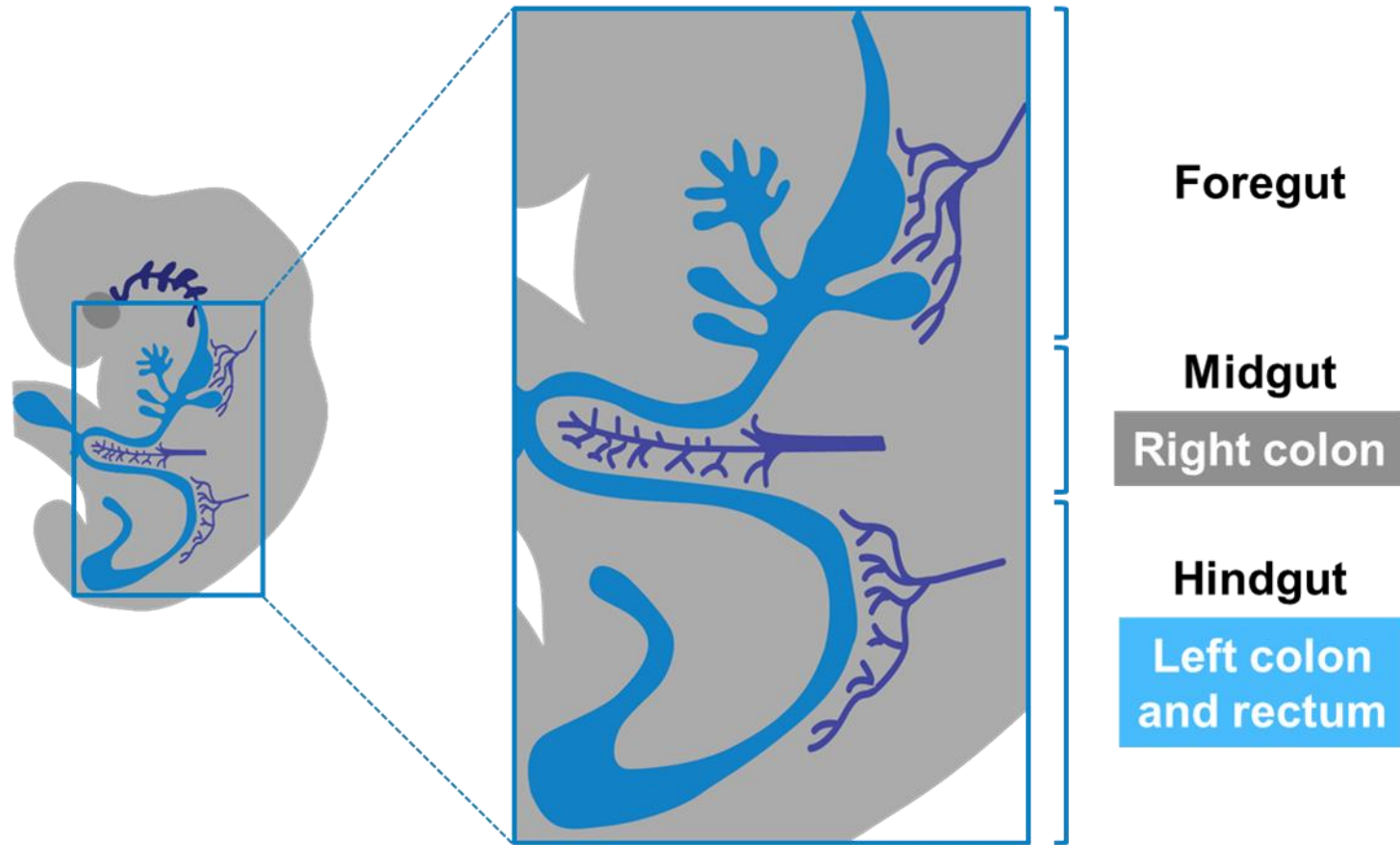


**TOPIC FIVE**

**SIDEDNESS**

# Anatomy, embryology and physiology

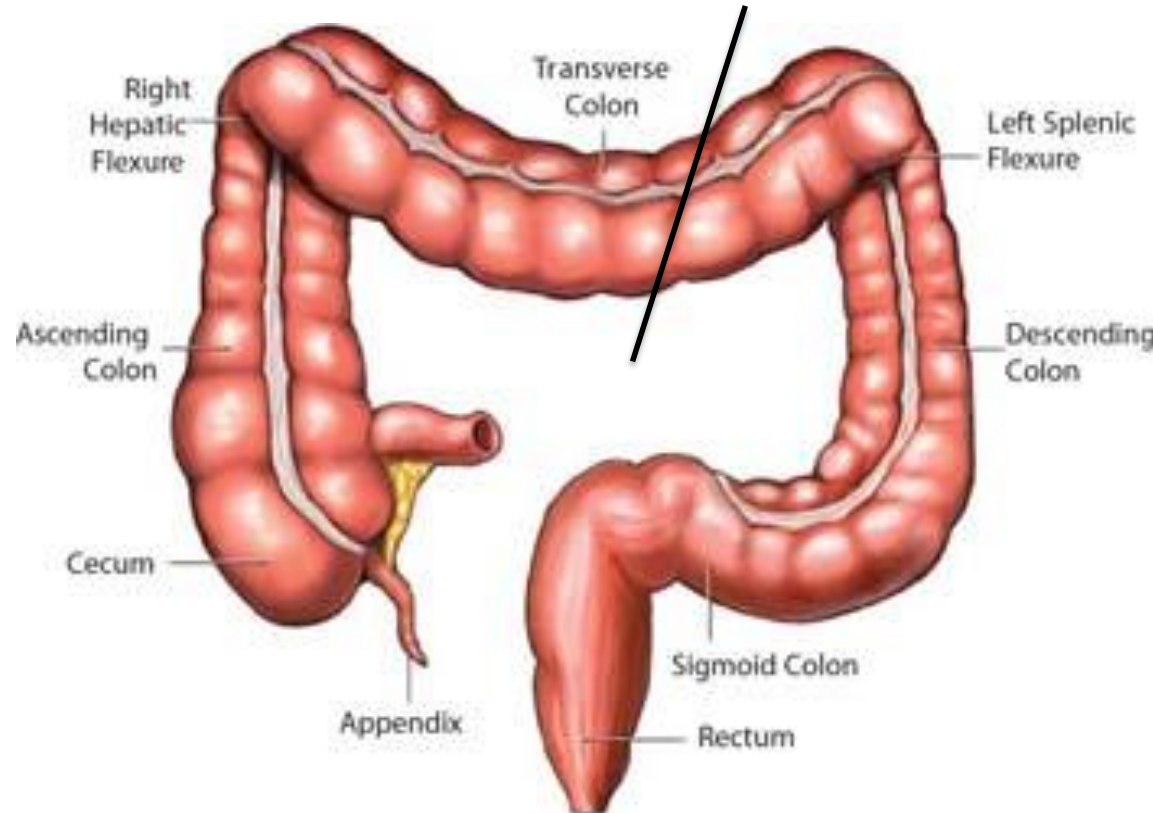
- ☰ The right colon and left colorectum have different embryonic origins and vascular supplies



# Right and Left Colon: Different Cancers, Different Biology, Different Outcomes.

## Right Cancers:

- Female
- Older
- More BRAF mutations
- MSI high
- More co-morbidities
- Less acute presentation
- WORSE prognosis



## Left Cancers:

- Male
- Younger
- Earlier Diagnosis
- More acute presentation
- Better Prognosis

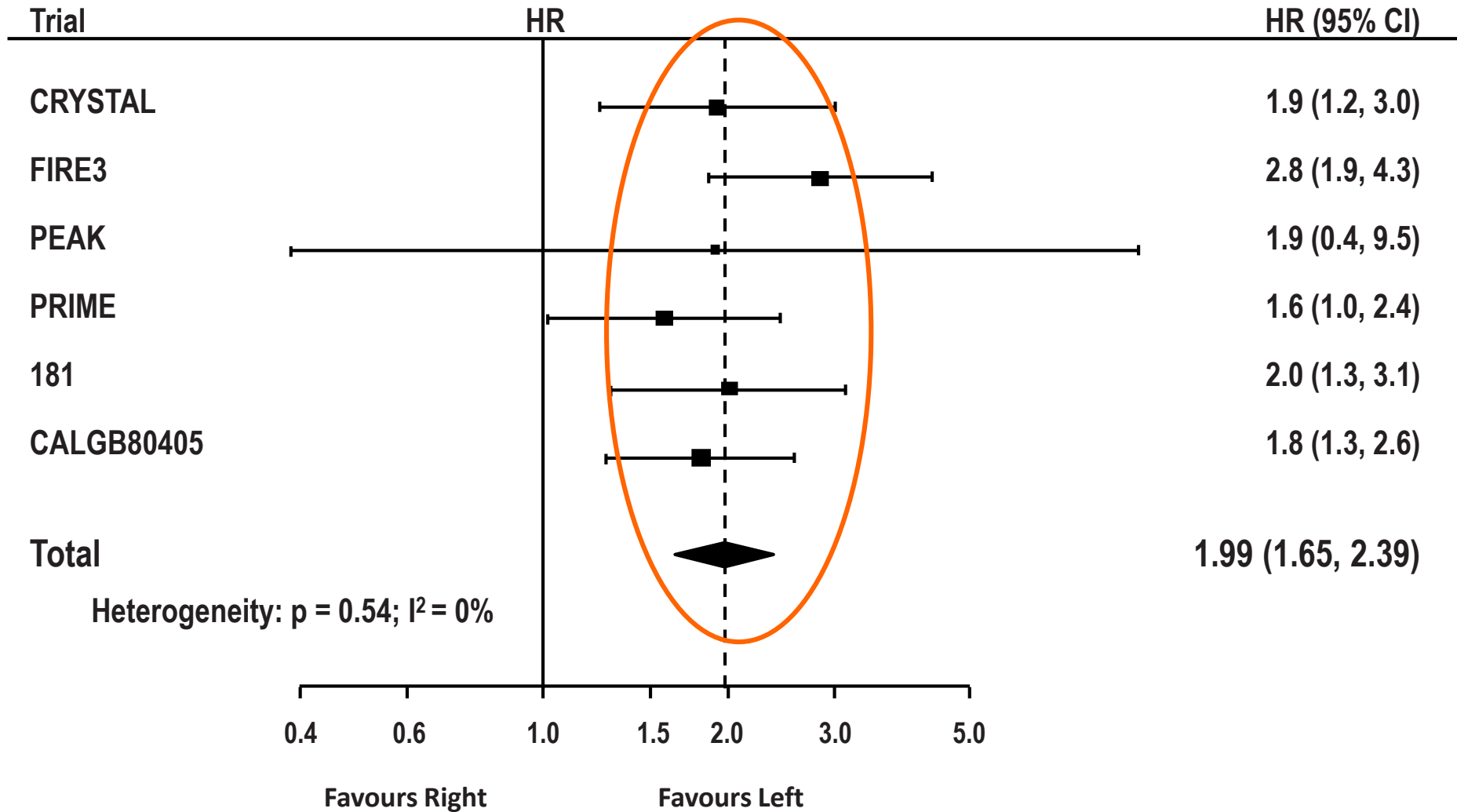
Anatomy of Large Intestine

Is right and left too simplistic? Is this instead a continuum?



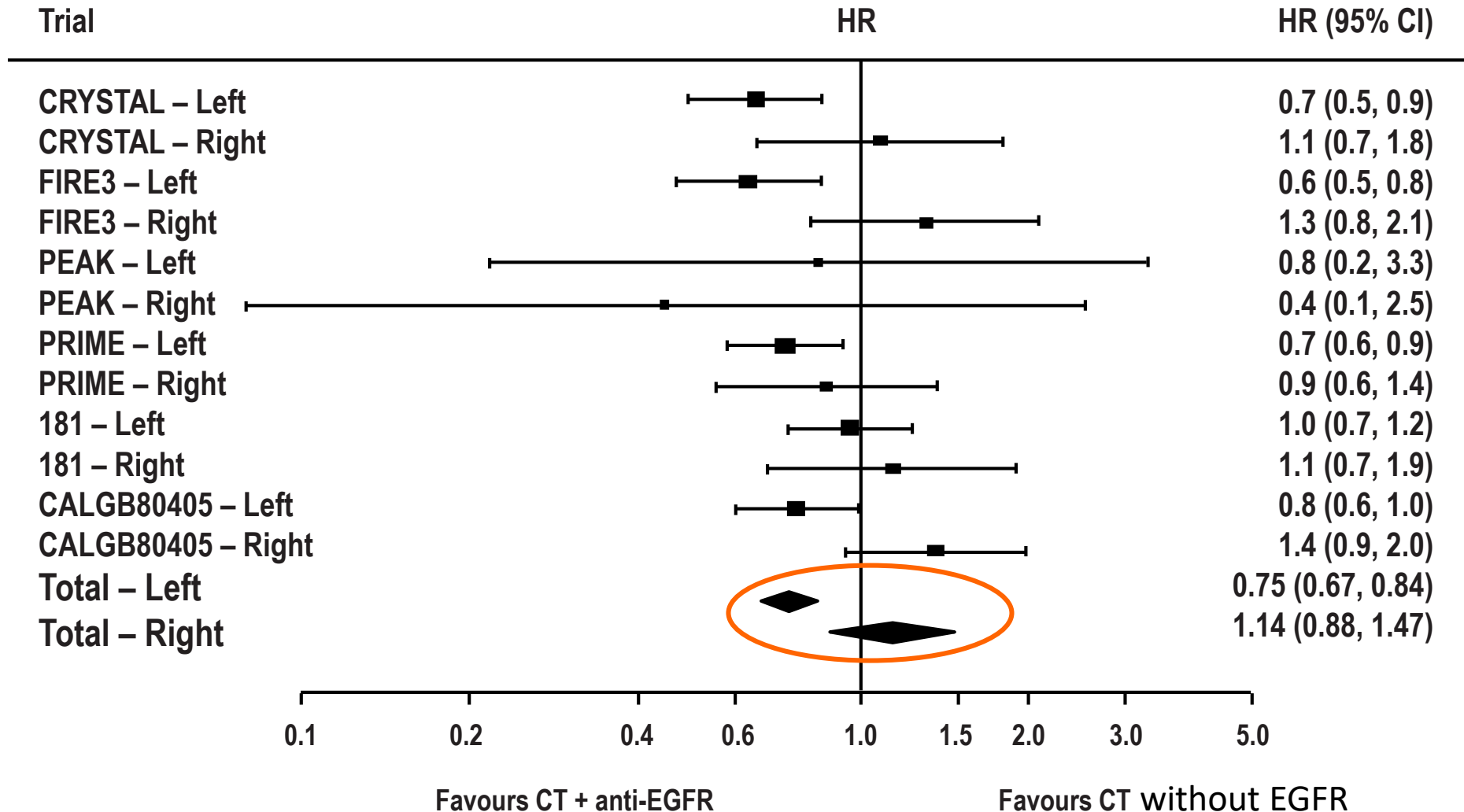
# Prognostic Analysis: OS

In all trials, OS favours left- over right-sided cancer



# Predictive Analysis: OS

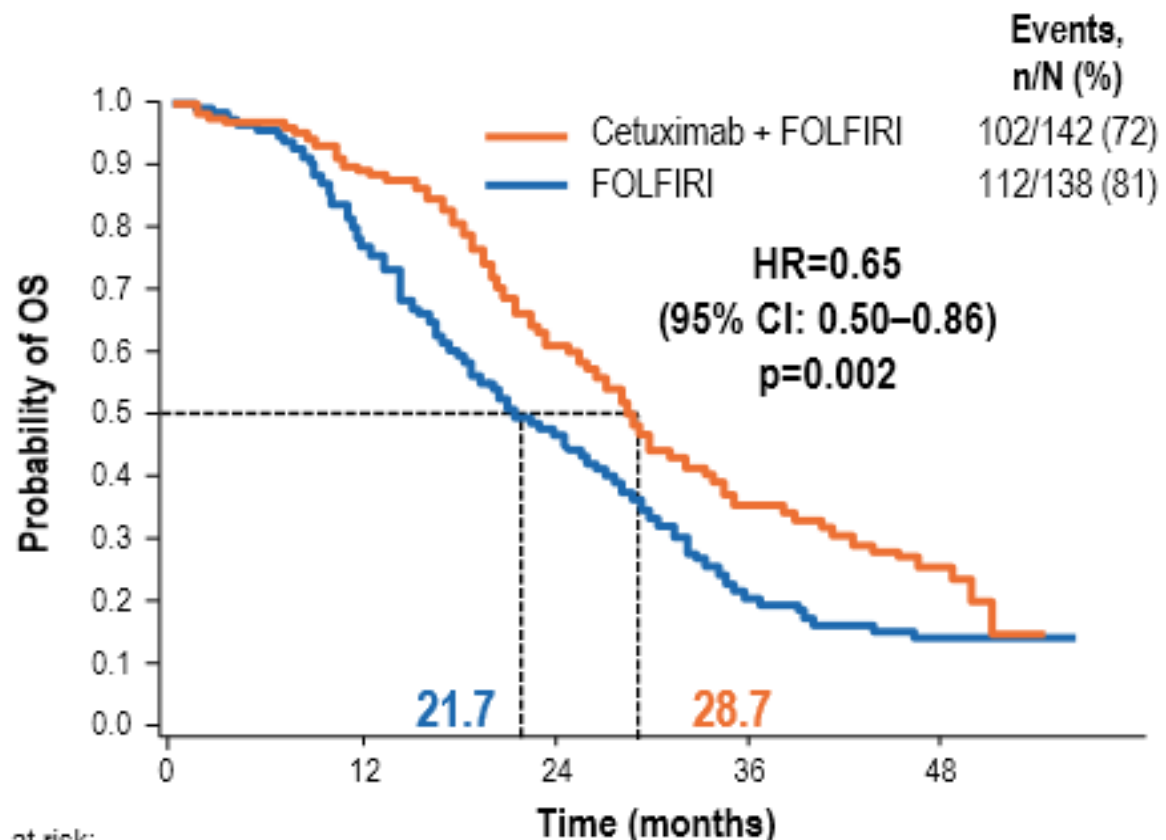
LEFT-SIDED TUMORS(RAS WT) DO BETTER ON EGFR Rx



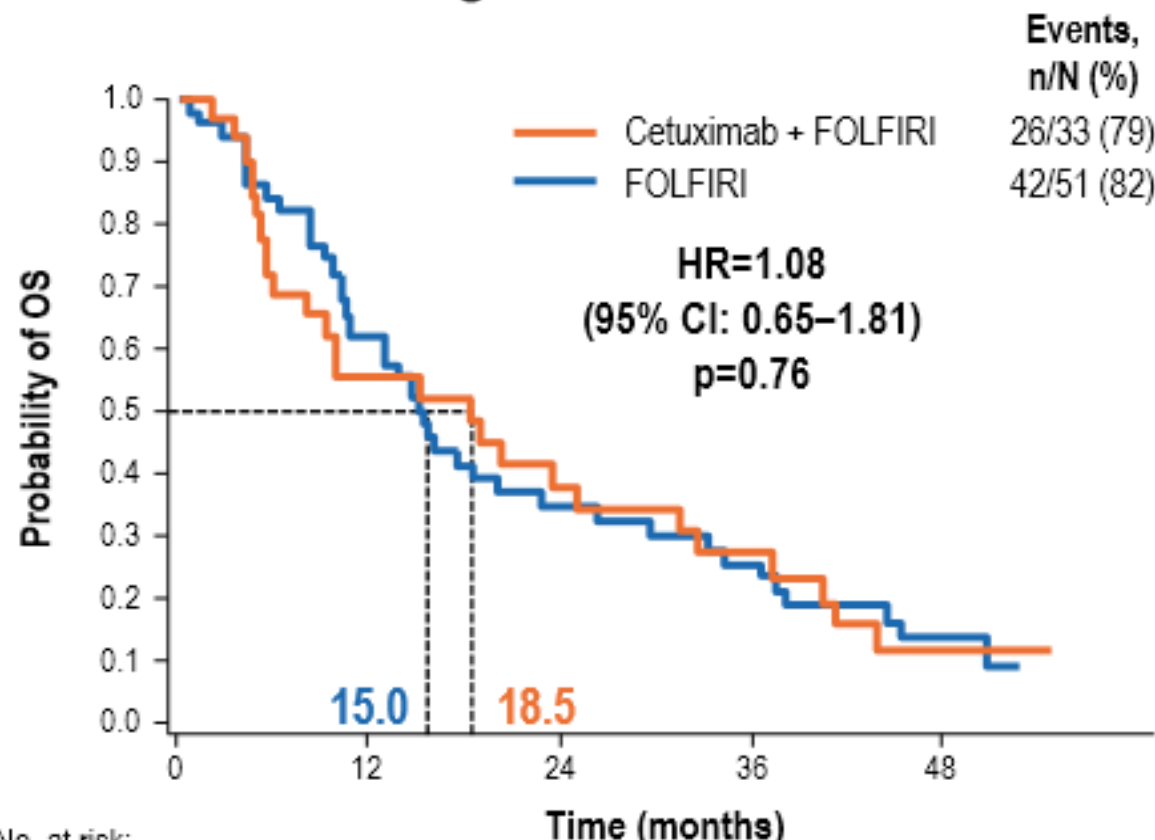
# CRYSTAL

## Relevant predictive value of primary tumor location: OS (*RAS*-wt)

### Left-sided tumors



### Right-sided tumors



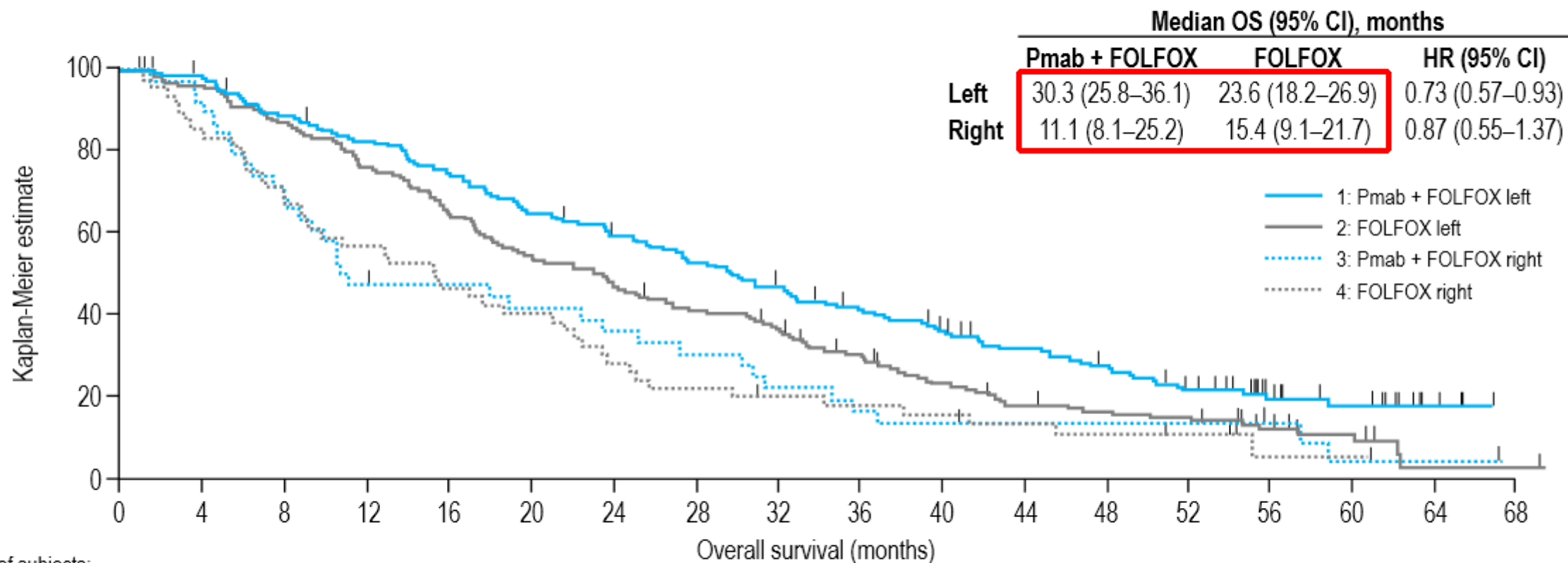
No. at risk:

|               | 0   | 12  | 24 | 36 | 48 |
|---------------|-----|-----|----|----|----|
| Cet + FOLFIRI | 142 | 123 | 83 | 47 | 14 |
| FOLFIRI       | 138 | 104 | 63 | 27 | 7  |

No. at risk:

|               | 0  | 12 | 24 | 36 | 48 |
|---------------|----|----|----|----|----|
| Cet + FOLFIRI | 33 | 16 | 11 | 7  | 1  |
| FOLFIRI       | 51 | 31 | 16 | 11 | 3  |

# PRIME – OS – 1st line

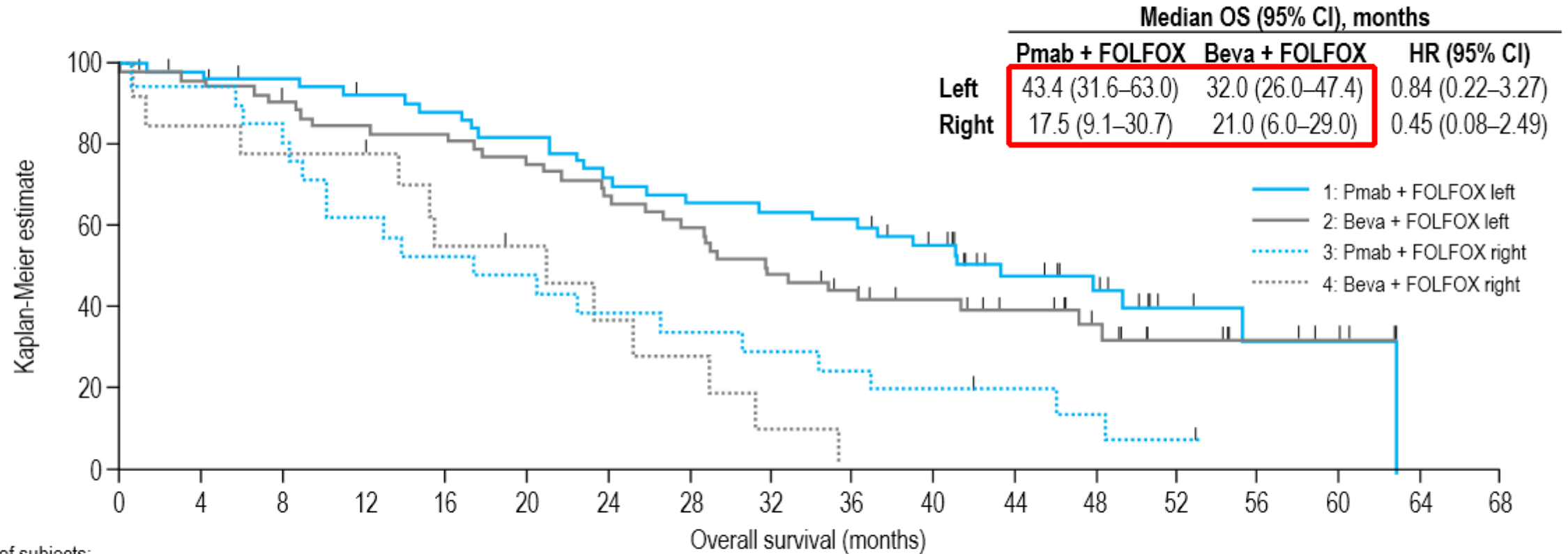


No. of subjects:

|    |     |     |     |     |     |     |    |    |    |    |    |    |    |    |    |    |   |   |
|----|-----|-----|-----|-----|-----|-----|----|----|----|----|----|----|----|----|----|----|---|---|
| 1: | 169 | 164 | 147 | 136 | 124 | 107 | 97 | 86 | 77 | 66 | 56 | 46 | 39 | 30 | 16 | 11 | 3 | 0 |
| 2: | 159 | 151 | 137 | 120 | 103 | 86  | 76 | 64 | 56 | 44 | 32 | 24 | 21 | 19 | 11 | 6  | 1 | 1 |
| 3: | 39  | 36  | 26  | 18  | 17  | 15  | 13 | 11 | 8  | 6  | 5  | 4  | 4  | 4  | 3  | 1  | 1 | 0 |
| 4: | 49  | 42  | 34  | 28  | 23  | 20  | 14 | 11 | 9  | 8  | 7  | 6  | 5  | 4  | 1  | 1  | 0 |   |

Censor indicated by vertical bar

# PEAK – OS – 1st line



No. of subjects:

|    |    |    |    |    |    |    |    |    |    |    |    |    |    |   |   |   |   |
|----|----|----|----|----|----|----|----|----|----|----|----|----|----|---|---|---|---|
| 1: | 53 | 51 | 49 | 46 | 44 | 41 | 36 | 33 | 32 | 31 | 25 | 17 | 13 | 6 | 3 | 2 | 0 |
| 2: | 54 | 51 | 48 | 44 | 43 | 40 | 35 | 31 | 26 | 21 | 17 | 13 | 10 | 5 | 3 | 2 | 0 |
| 3: | 22 | 21 | 18 | 13 | 11 | 10 | 8  | 7  | 6  | 5  | 4  | 3  | 2  | 1 | 0 |   |   |
| 4: | 14 | 12 | 11 | 11 | 7  | 6  | 4  | 3  | 1  | 0  |    |    |    |   |   |   |   |

Censor indicated by vertical bar

# Panitumumab efficacy by tumour location analysis

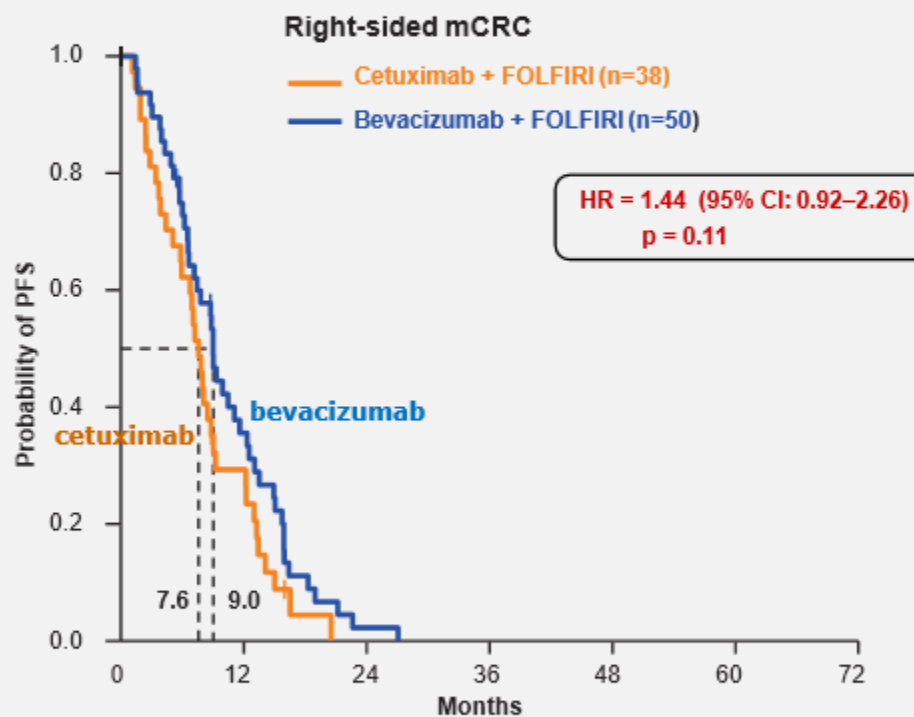
## Prognostic effect of primary tumour location

| WT RAS  | PRIME                 |       | PEAK                   |       |
|---|-----------------------|-------|------------------------|-------|
|   | Panitumumab + FOLFOX4 |       | Panitumumab + mFOLFOX6 |       |
| Panitumumab arm                                     | Left                  | Right | Left                   | Right |
| Median OS, months                                   | 30.3                  | 11.1  | 43.4                   | 17.5  |
| Adjusted HR, <sup>†</sup> right vs left<br>(95% CI) | 1.58<br>(1.02–2.45)   |       | 2.68<br>(1.31–5.46)    |       |
| Comparator arm                                      | FOLFOX4               |       | Bevacizumab + mFOLFOX6 |       |
|   | Left                  | Right | Left                   | Right |
| Median OS, months                                   | 23.6                  | 15.4  | 32.0                   | 21.0  |
| Adjusted HR, <sup>†</sup> right vs left<br>(95% CI) | 1.27<br>(0.88–1.84)   |       | 2.86<br>(1.40–5.84)    |       |

- Right-sided primary tumours were associated with worse prognosis compared with left-sided tumours, regardless of treatment received

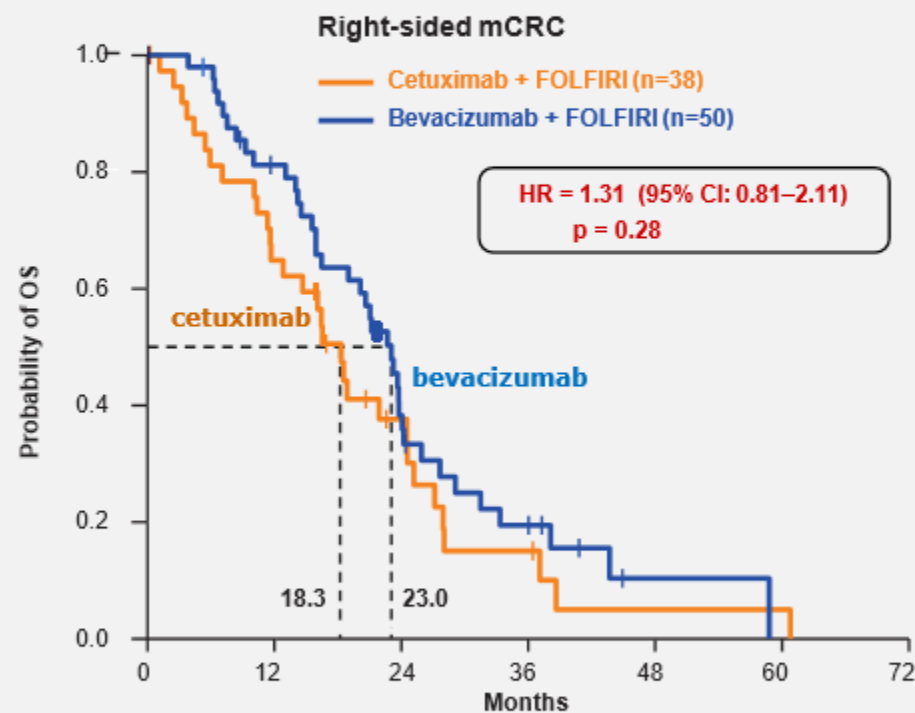
# FIRE-3: Right-sided tumors

## Progression-free survival



|                       |    | Numbers atRisk |    |    |    |    |    |    |
|-----------------------|----|----------------|----|----|----|----|----|----|
|                       |    | 0              | 12 | 24 | 36 | 48 | 60 | 72 |
| Cetuximab + FOLFIRI   | 38 | 10             | 0  | 0  | 0  | 0  | 0  | 0  |
| Bevacizumab + FOLFIRI | 50 | 16             | 1  | 0  | 0  | 0  | 0  | 0  |

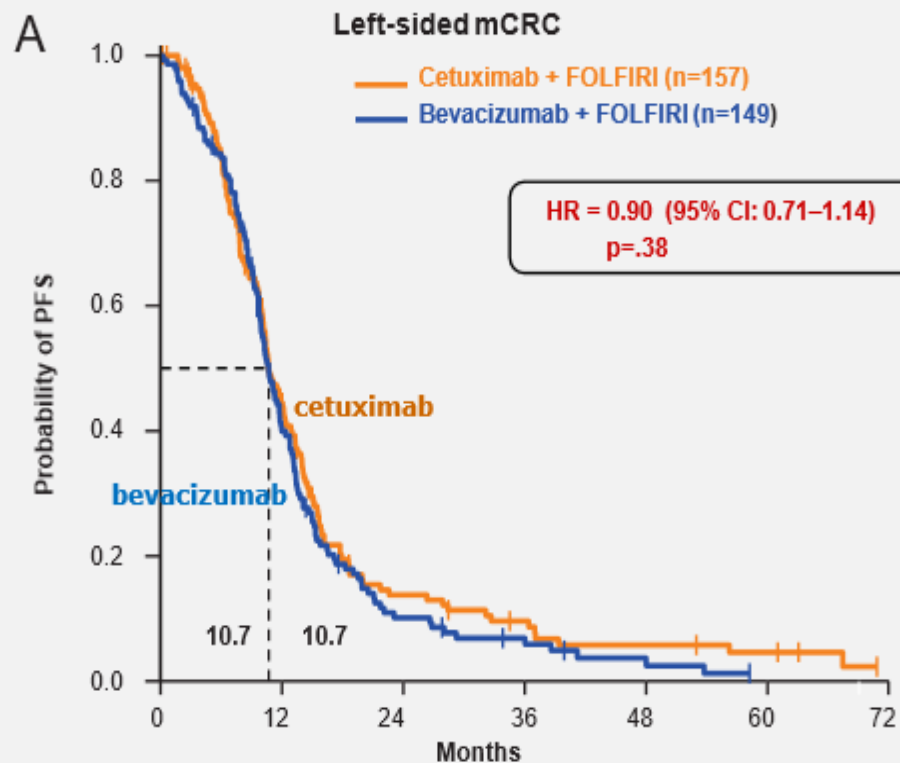
## Overall survival



|                       |    | Numbers atRisk |    |    |    |    |    |    |
|-----------------------|----|----------------|----|----|----|----|----|----|
|                       |    | 0              | 12 | 24 | 36 | 48 | 60 | 72 |
| Cetuximab + FOLFIRI   | 38 | 24             | 10 | 4  | 1  | 1  | 0  | 0  |
| Bevacizumab + FOLFIRI | 50 | 37             | 16 | 7  | 1  | 0  | 0  | 0  |

# FIRE-3: Left-sided tumors

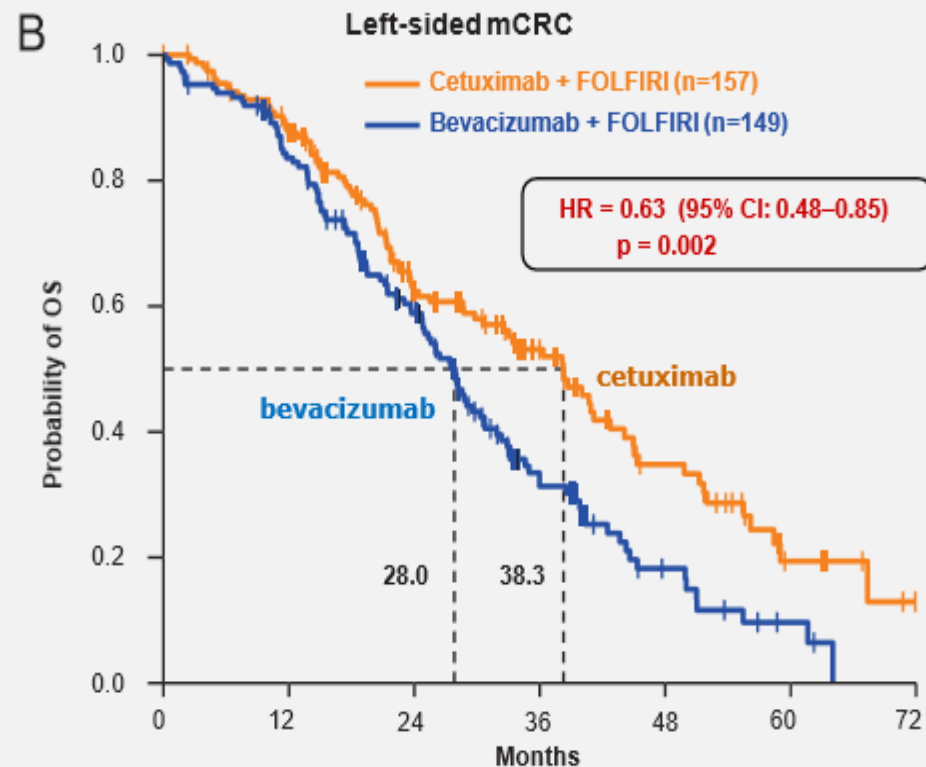
## Progression-free survival



Numbers at Risk

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

## Overall survival

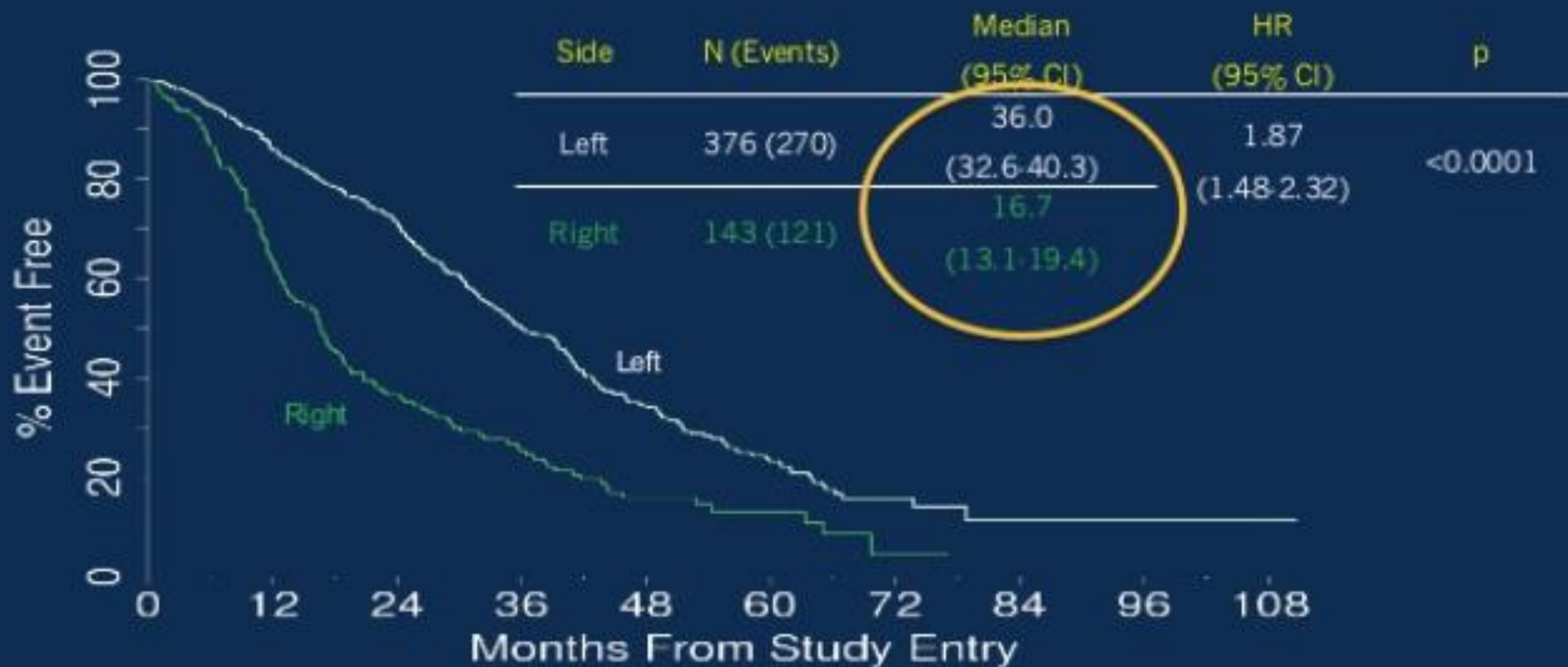


Numbers at Risk

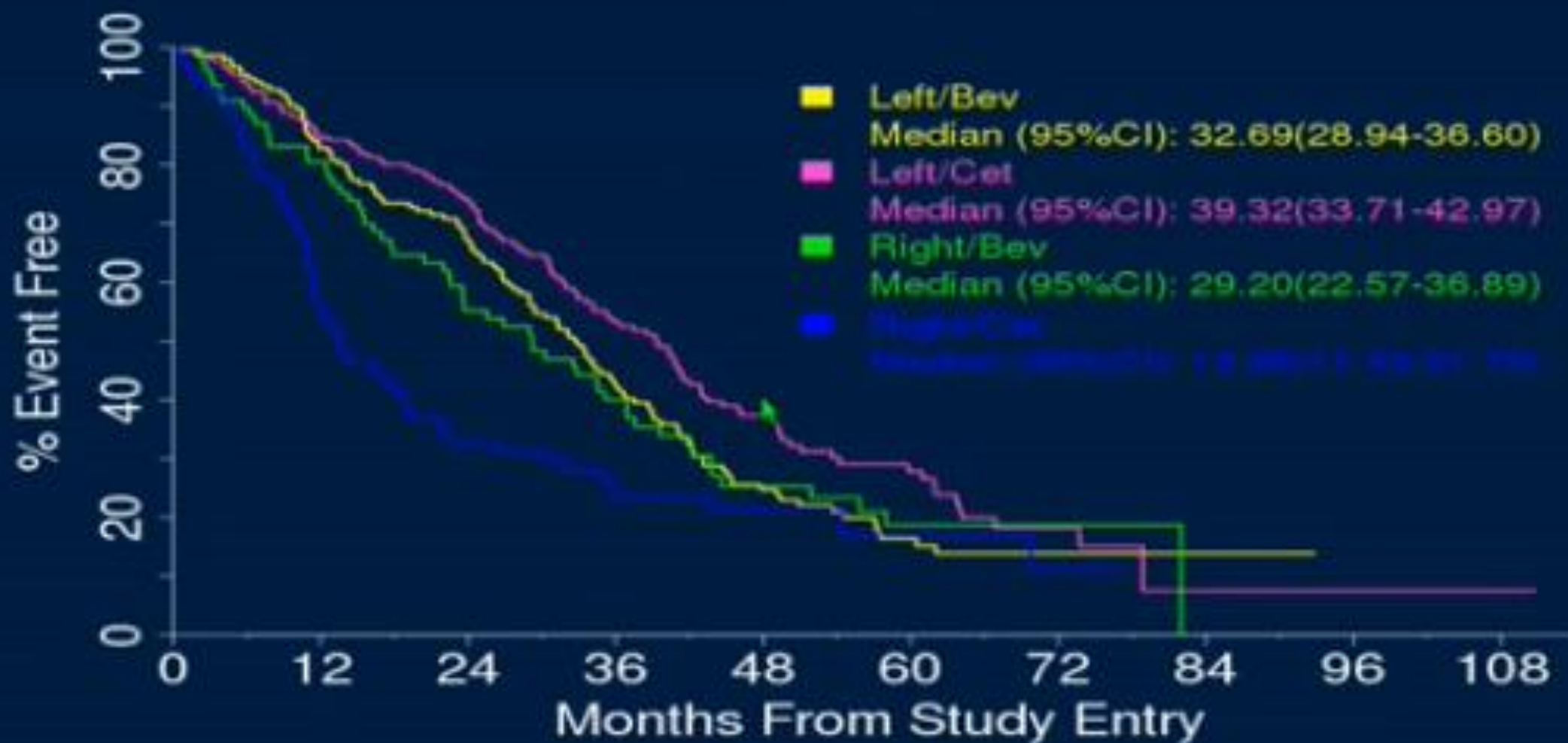
|                       |     |     |    |    |    |   |   |
|-----------------------|-----|-----|----|----|----|---|---|
| Cetuximab + FOLFIRI   | 157 | 131 | 77 | 38 | 23 | 6 | 0 |
| Bevacizumab + FOLFIRI | 149 | 120 | 76 | 31 | 11 | 3 | 0 |



# 80405: OS by Sidedness (Cetuximab)



# 80405: Overall Survival by Sidedness and Biologic



# Potential explanatory factors

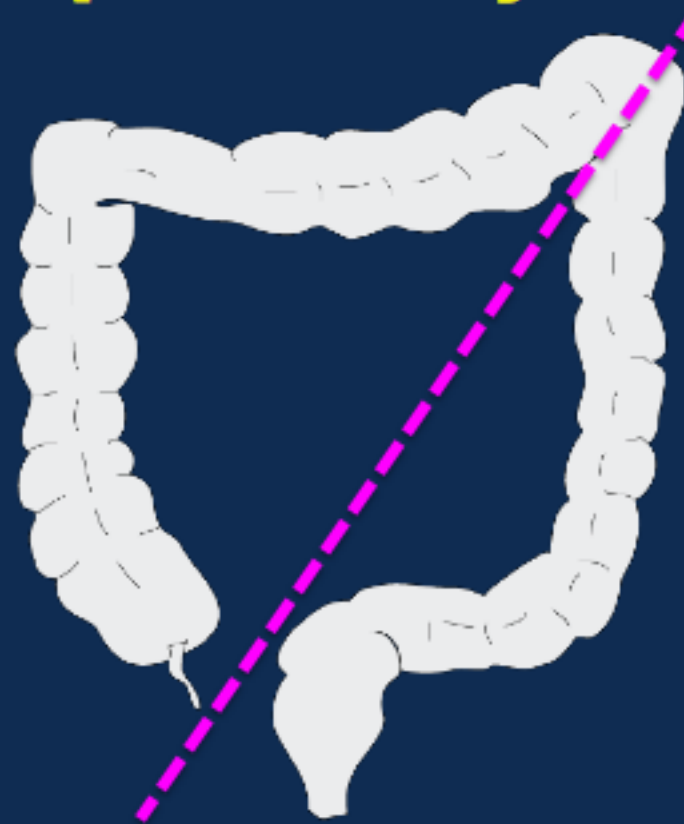
Clinical

Embryologic /  
Physiologic

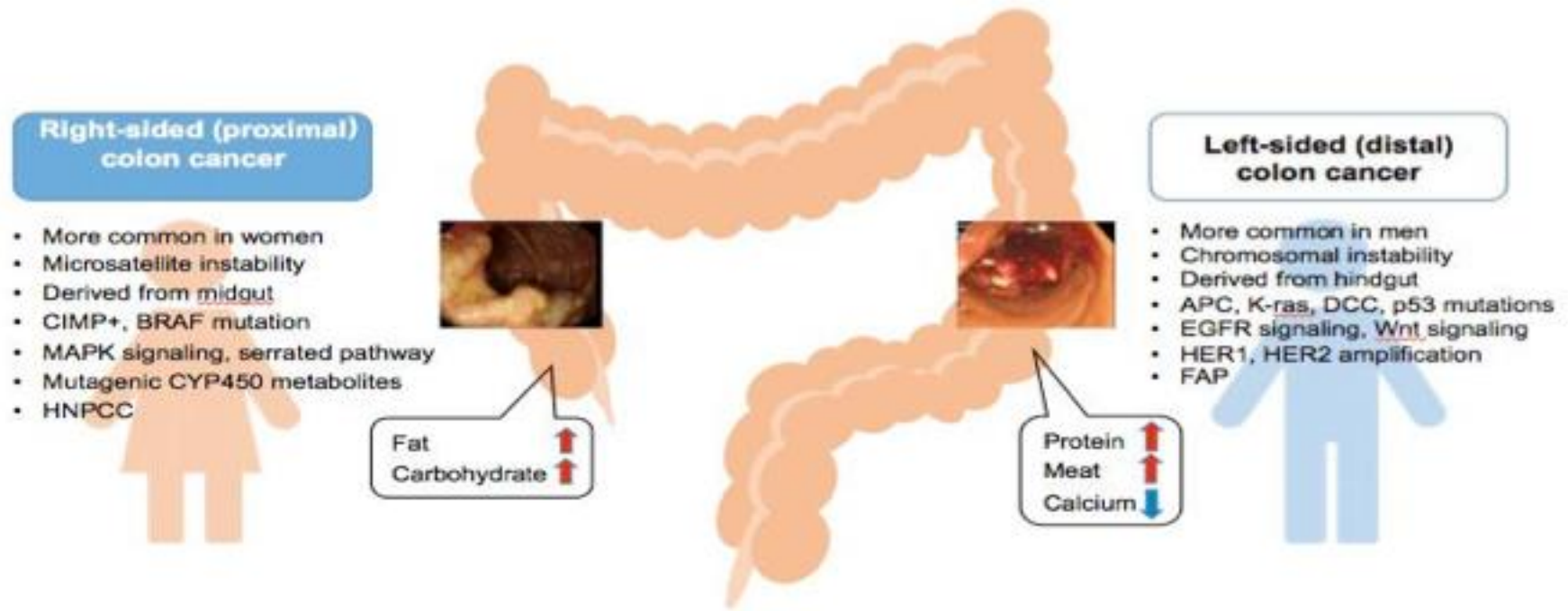
Epidemiologic

Environmental

Molecular /  
Genetic



# Midgut vs Hindgut



**Summary of common clinical and molecular characteristics of right- and left-sided colon tumors and associations with dietary factors.** CIMP = CpG island methylator phenotype; HNPCC = hereditary non-polyposis colorectal cancer; APC = adenomatous polyposis coli; K-ras = Kirsten-ras; DCC = deleted in colorectal cancer; FAP = familial adenomatous polyposis.

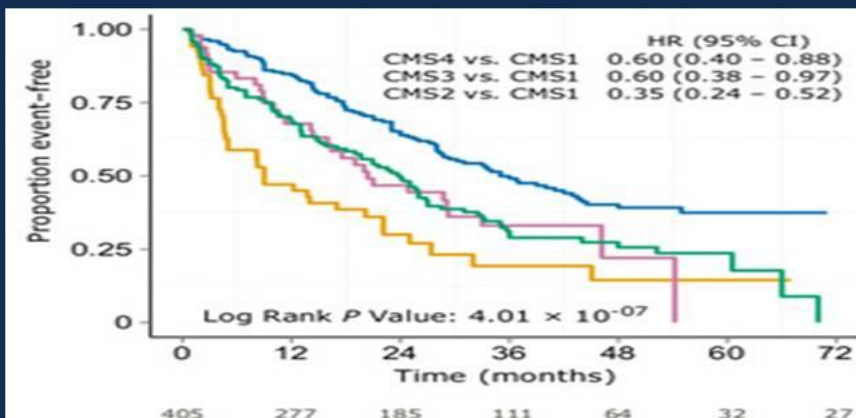
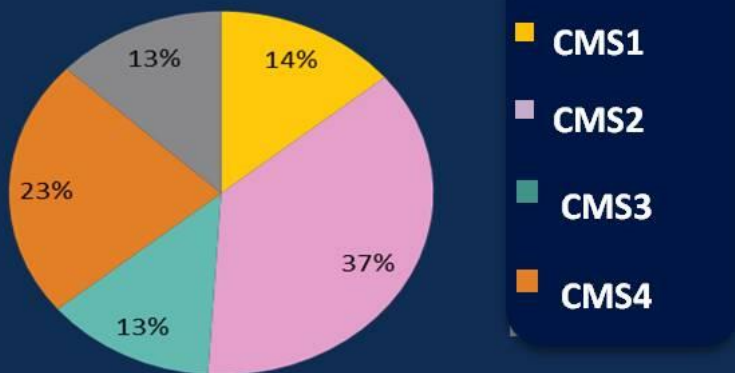


# Discussion: Relating Phenotype to Genotype

## Consensus Molecular Subtypes in Colorectal Cancer

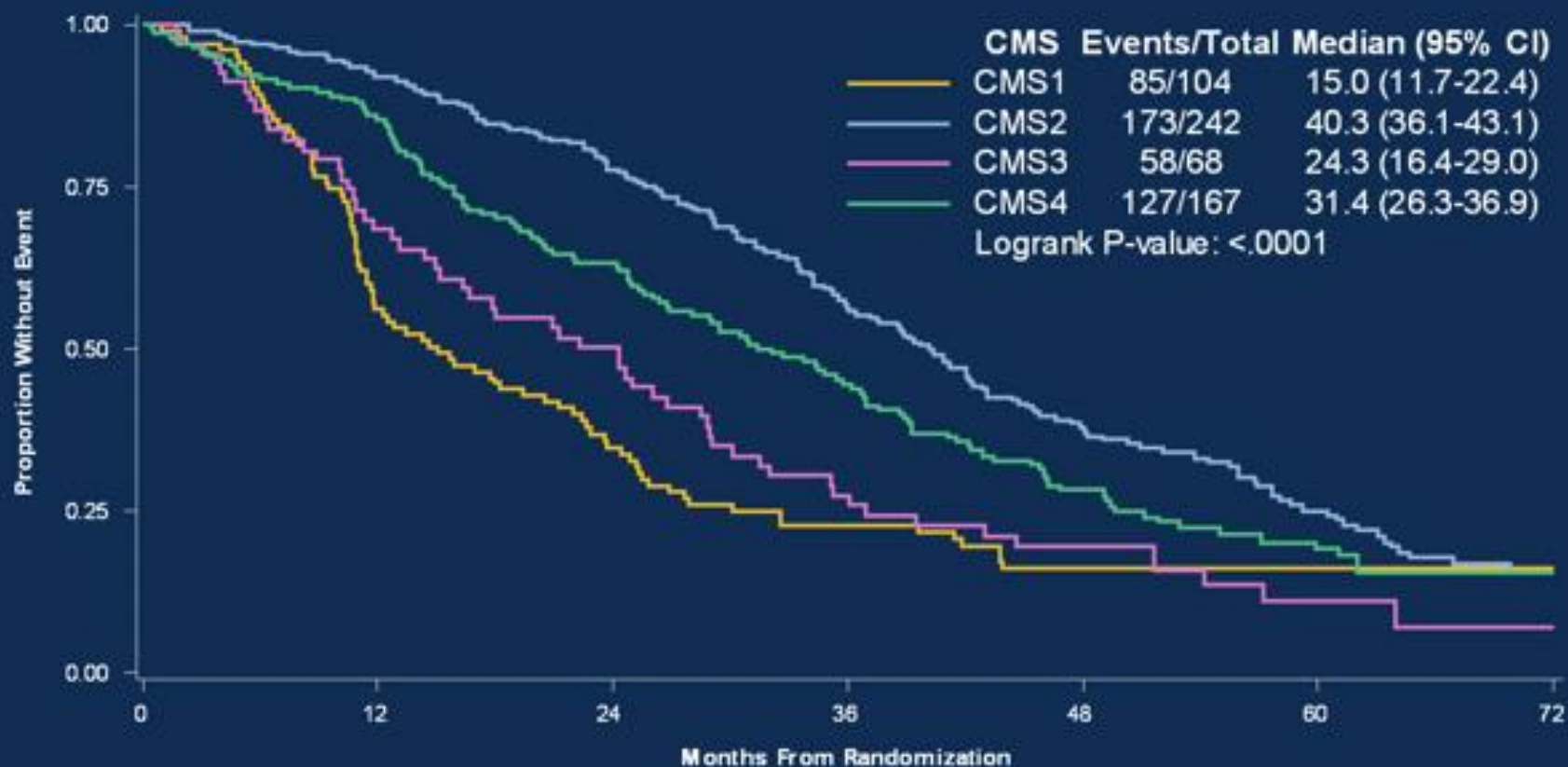
Guinney et al. *Nature Medicine* Nov 2015

### 4 Consensus Molecular Subtypes



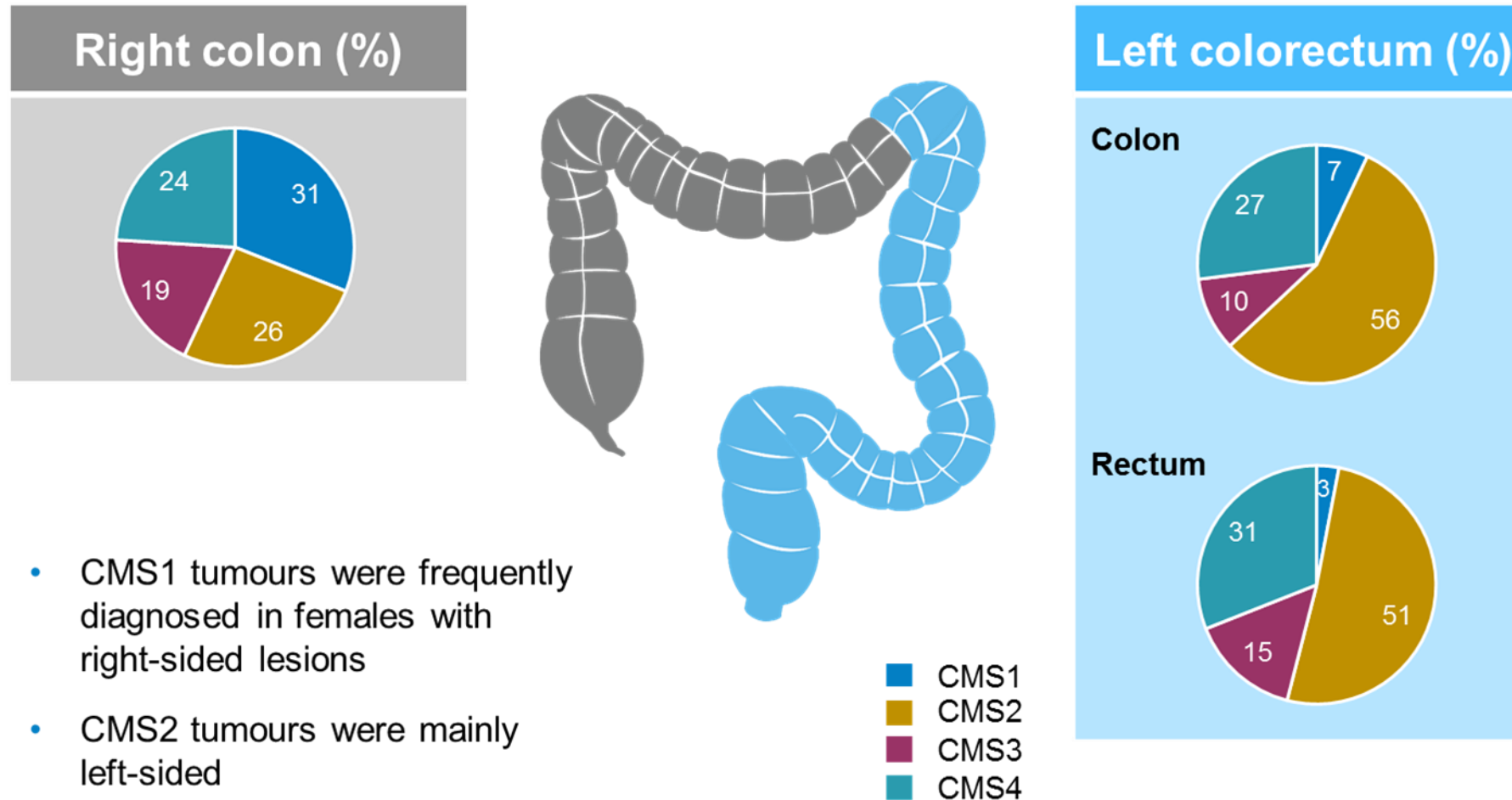
| CMS1<br>MSI Immune                 | CMS2<br>Canonical      | CMS3<br>Metabolic                    | CMS4<br>Mesenchymal                                 |
|------------------------------------|------------------------|--------------------------------------|---|
| 14%                                | 37%                    | 13%                                  | 23%   |
| MSI, CIMP high, hypermutation      | SCNA high              | Mixed MSI status, SCNA low, CIMP low | SCNA high   |
| <i>BRAF</i> mutations              |                        | <i>KRAS</i> mutations                |   |
| Immune infiltration and activation | WNT and MYC activation | Metabolic deregulation               | Stromal infiltration, TGFβ activation, angiogenesis |
| Worse survival after relapse       |                        |                                      | Worse relapse-free and overall survival             |

## OS – All Patients by CMS Subtype



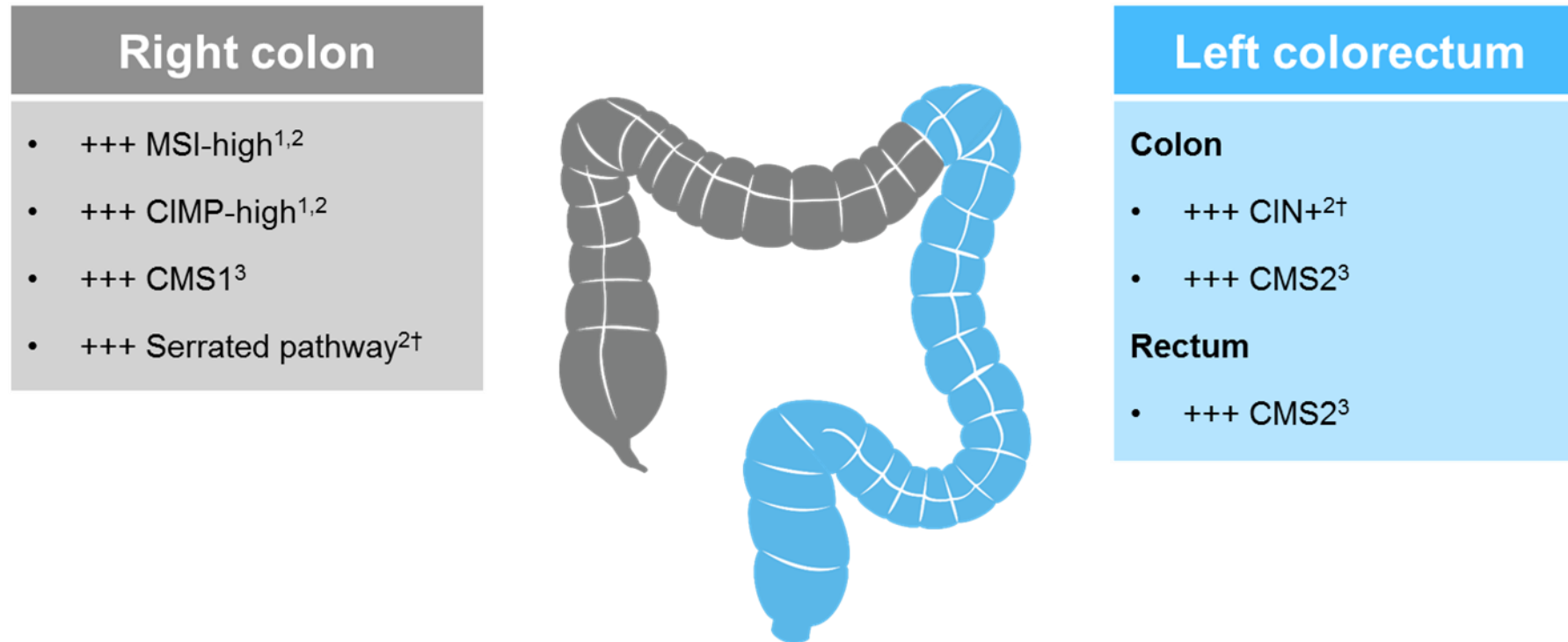
# Molecular pathways/tumour classification models

## ☰ Distribution of CMS groups varies by colorectal tumour location



# Molecular pathways/tumour classification models

## ☰ CIMP-high, MSI-high and CMS distribution varies between colorectal locations



- Molecular heterogeneity may be captured by the anatomical location of the tumour

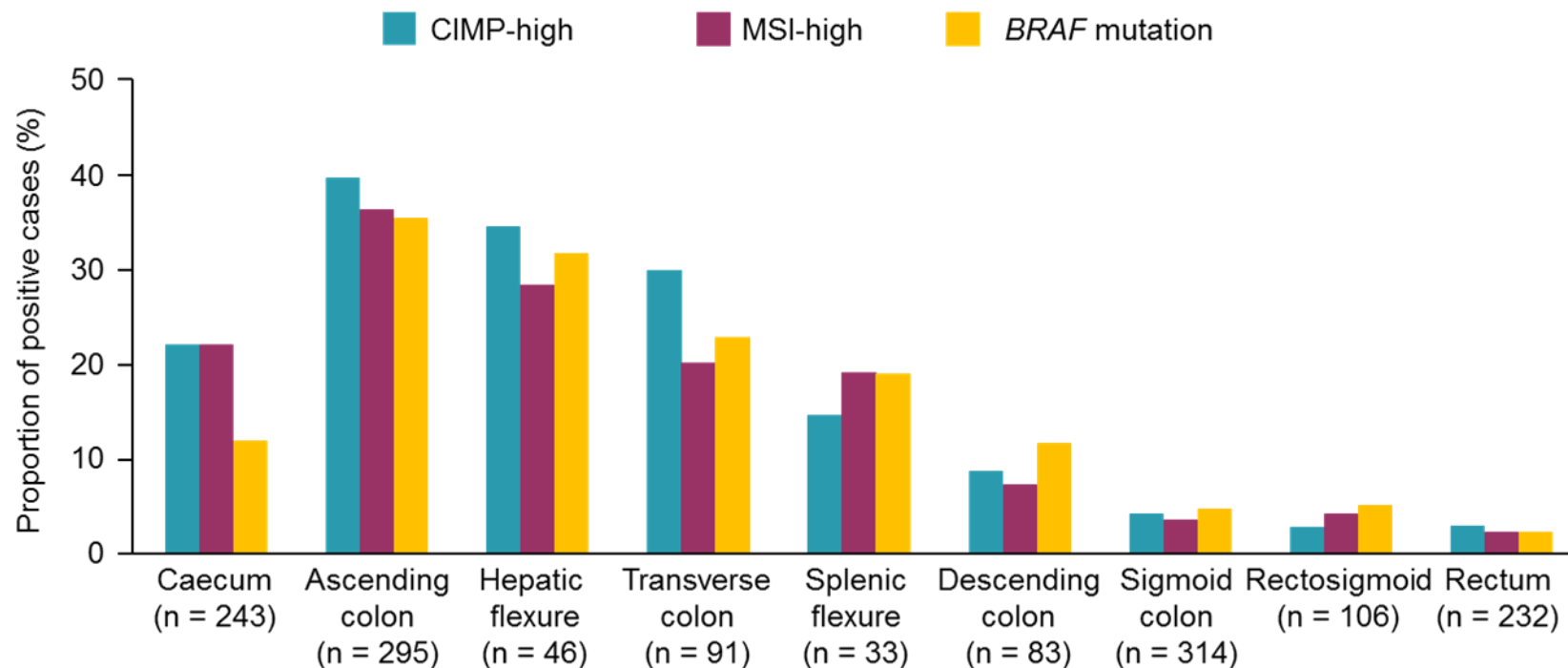
1. Yamauchi M, et al. Gut 2012;61:847–54;  
2. Missiaglia E, et al. Ann Oncol 2014;25:1995–2001;  
3. Guinney J, et al. Nat Med 2015;21:1350–6 (and Supplementary Figure 10).

+++ denotes that a molecular or morphological subtype is more prevalent compared with other colorectal segments.  
†CIN expression and serrated pathway data are not available for rectal tumours.  
CIMP, CpG island methylator phenotype; CIN, chromosomal instability;  
CMS, Consensus Molecular Subtypes; MSI, microsatellite instability.



# Is classification by side an over-simplification?

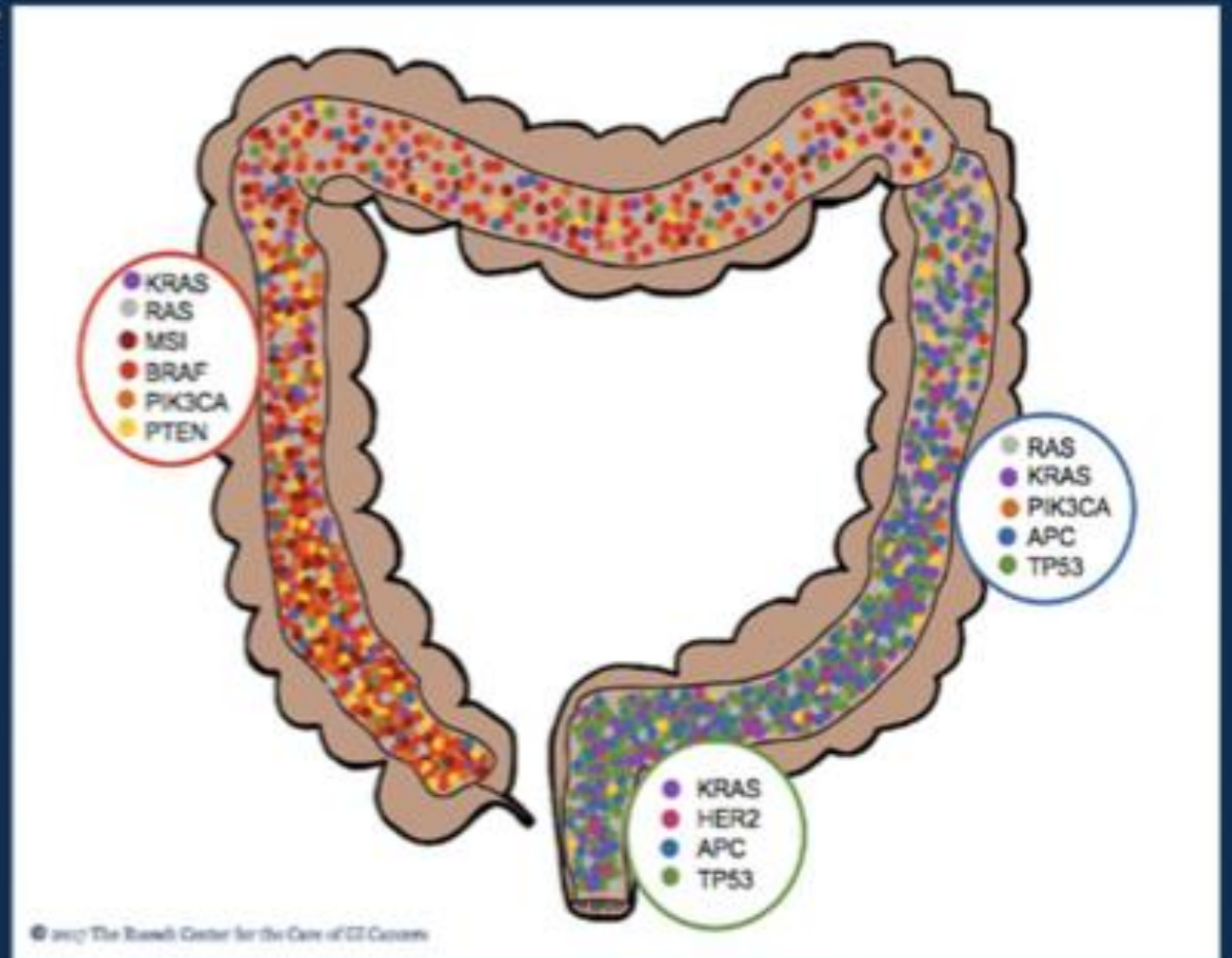
## ☰ Distribution of molecular characteristics varies across colorectal cancer subsites



Variation in molecular characteristics across bowel subsites challenges the concept of discrete right colon and left colorectal tumour characteristics

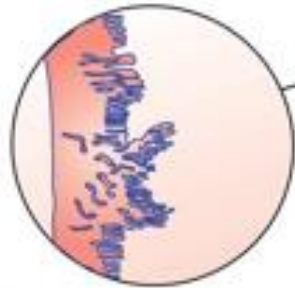
# Conclusions

- CRCs carry a continuum of molecular alterations from right to left, rather than having a sharp, clear-cut distinction



**RIGHT-SIDED CRC**  
Inferior outcomes  
with cetuximab  
Poorer prognosis

Sessile serrated polyps  
CMS1 and CMS3  
CIMP-high  
Midgut  
*BRAF* mutant  
MSI-high  
Bile acid exposure  
Invasive bacteria biofilms



**LEFT-SIDED CRC**  
Superior outcomes  
with cetuximab  
Better prognosis

Tubular adenoma  
CMS2 and CMS4  
Higher *EREG/AREG*  
expression  
Hindgut



Figure 2. Summary of key biologic differences between right- and left-sided CRCs.

Abbreviations: AREG, amphiregulin; CIMP, CpG island methylator phenotype; CMS, consensus molecular subtype; CRCs, colorectal cancers; EREG, epiregulin; MSI, microsatellite instability.

# TOPIC SIX

MSI



# Two different pathways of carcinogenesis

85%



## Chromosomal instability (CIN):

Proficient DNA Mismatch Repair (pMMR)  
and Microsatellite stable (MSS)

15%



## Microsatellite instability (MSI):

Deficient Mismatch Repair (dMMR)

2/3



*BRAF*<sup>V600E</sup>  
mutation  
≈50%

Sporadic cases  
Epigenetic *MLH1* inactivation  
(*MLH1* promoter methylation)

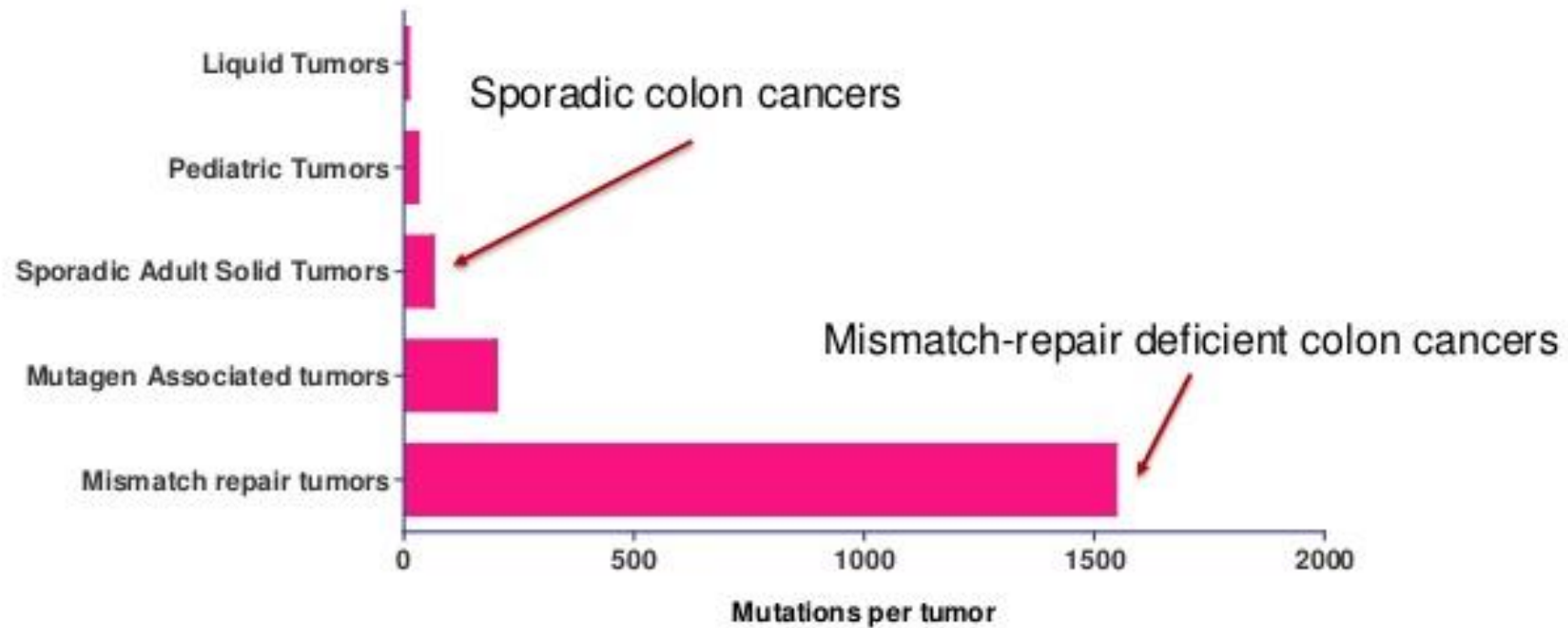
1/3



No  
*BRAF*<sup>V600E</sup>  
mutation

Familial cases  
Lynch syndrome  
germline mutation:  
(*MLH1*, *MSH2*, *MSH6*, *PMS2*)

## Mutations per tumor



The James

 THE OHIO STATE UNIVERSITY  
WEXNER MEDICAL CENTER

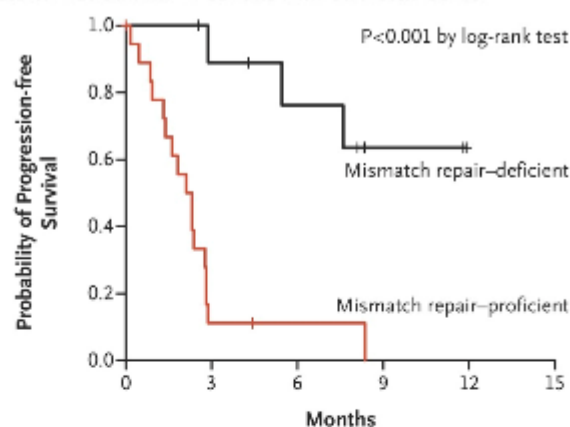
# Mismatch repair deficiency

- Proficient mismatch repair (pMMR) = microsatellite stable (MSS)
- Deficient mismatch repair (dMMR) = microsatellite instable (MSI)
- Detected by:
  - IHC- missing mismatch repair proteins
  - PCR- microsatellite instable
- Present clinical use:
  - Lynch syndrome
  - Determine whether to offer adjuvant chemotherapy for Stage II colon cancer patients



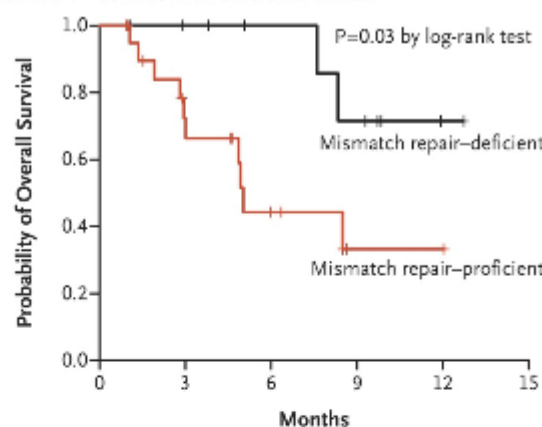


**A Progression-free Survival in Cohorts with Colorectal Cancer**



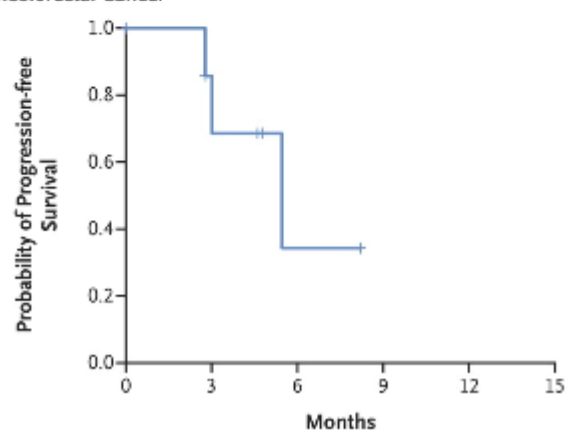
| No. at Risk                | 0  | 3 | 6 | 9 | 12 | 15 |
|----------------------------|----|---|---|---|----|----|
| Mismatch repair-deficient  | 11 | 8 | 6 | 2 | 0  | 0  |
| Mismatch repair-proficient | 21 | 2 | 1 | 0 | 0  | 0  |

**B Overall Survival in Cohorts with Colorectal Cancer**



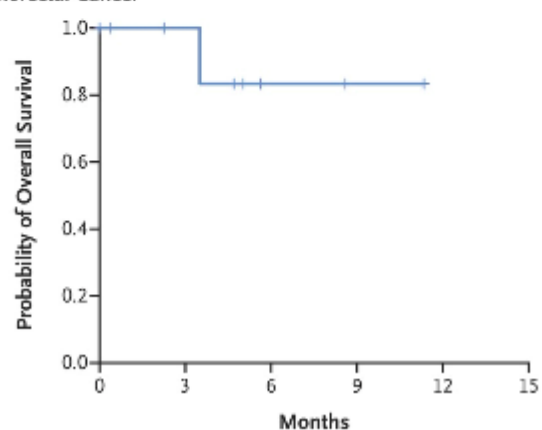
| No. at Risk                | 0  | 3  | 6 | 9 | 12 | 15 |
|----------------------------|----|----|---|---|----|----|
| Mismatch repair-deficient  | 11 | 9  | 7 | 5 | 1  | 0  |
| Mismatch repair-proficient | 21 | 12 | 5 | 1 | 1  | 0  |

**C Progression-free Survival in Cohort with Mismatch Repair-Deficient Noncolorectal Cancer**



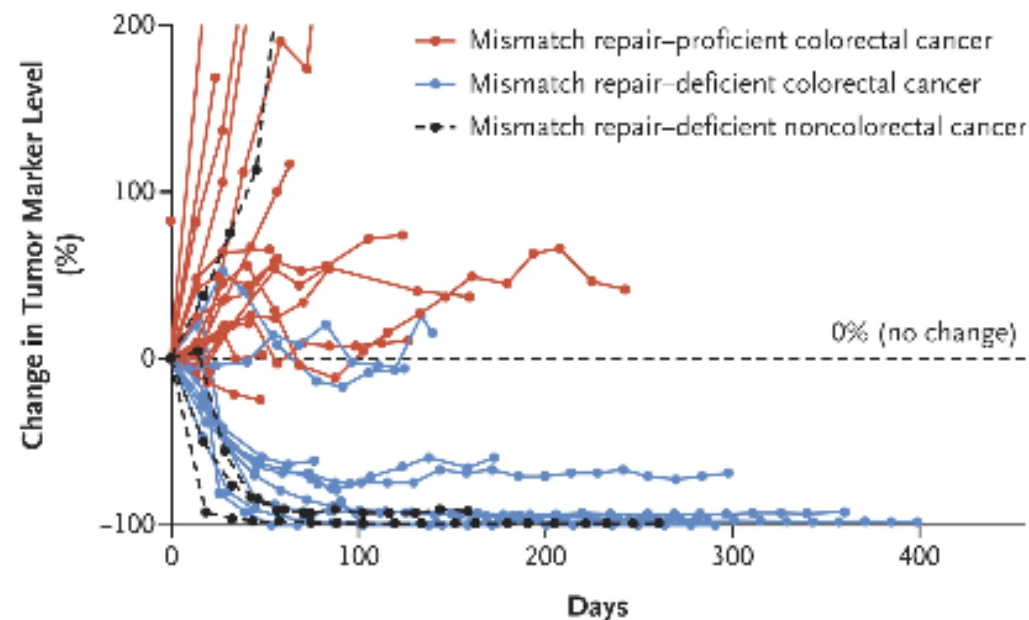
| No. at Risk | 0 | 3 | 6 | 9 | 12 | 15 |
|-------------|---|---|---|---|----|----|
|             | 9 | 5 | 1 | 0 | 0  | 0  |

**D Overall Survival in Cohort with Mismatch Repair-Deficient Noncolorectal Cancer**

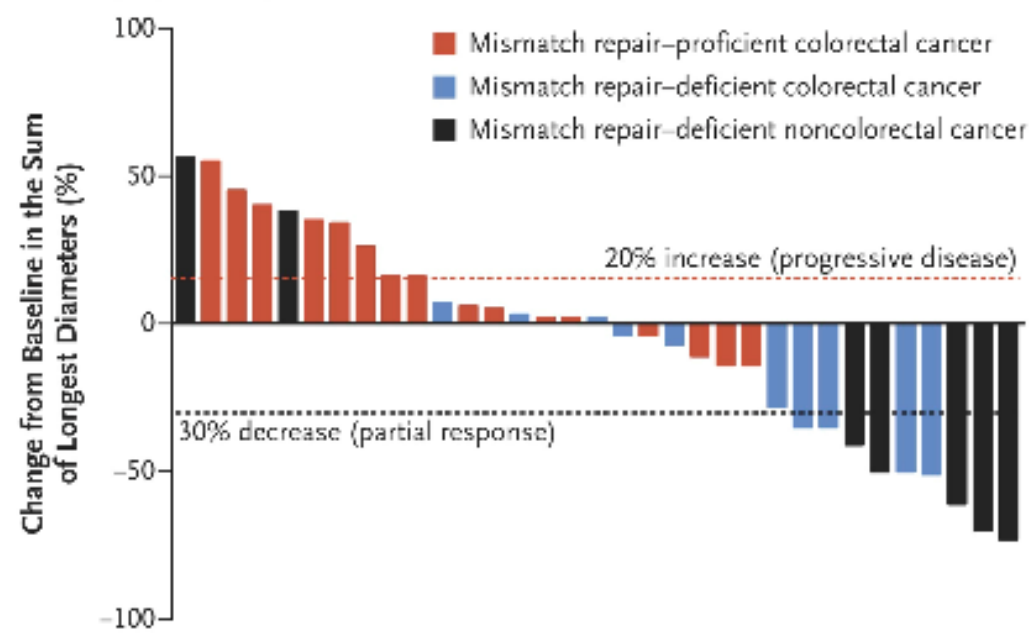


| No. at Risk | 0 | 3 | 6 | 9 | 12 | 15 |
|-------------|---|---|---|---|----|----|
|             | 9 | 6 | 2 | 1 | 0  | 0  |

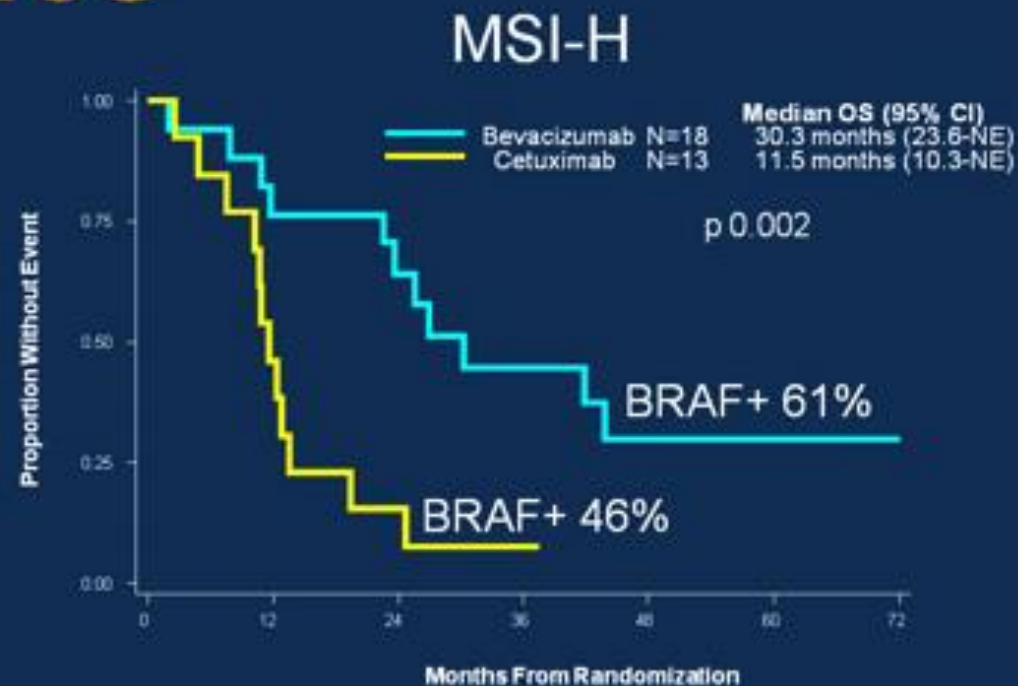
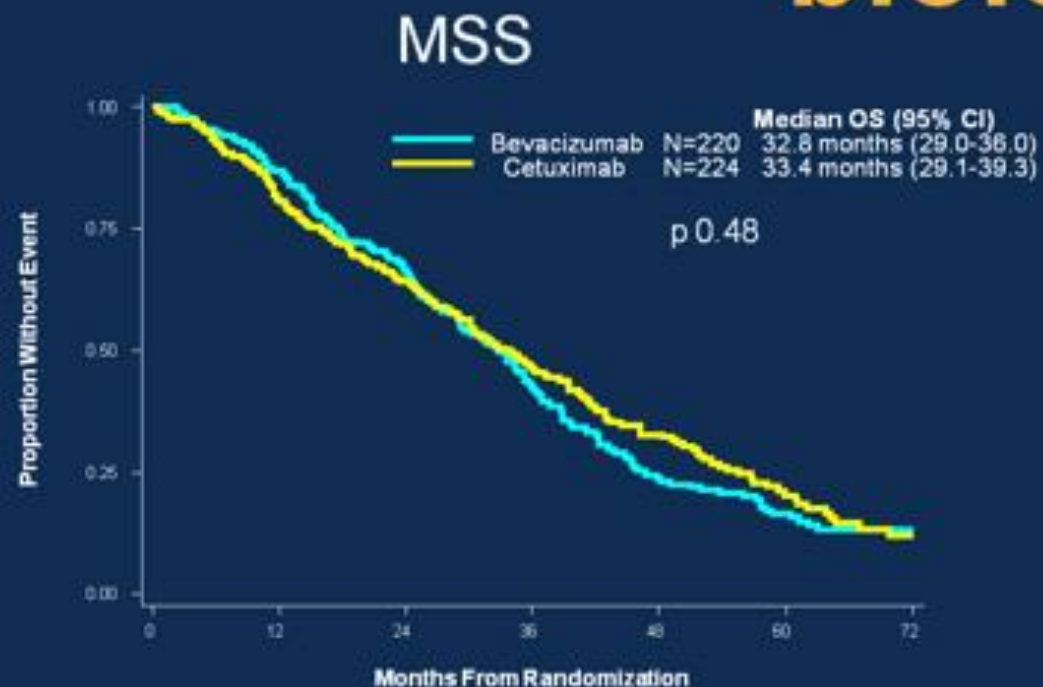
**A Biochemical Response**



**B Radiographic Response**



# Interaction of MSI status with biologics

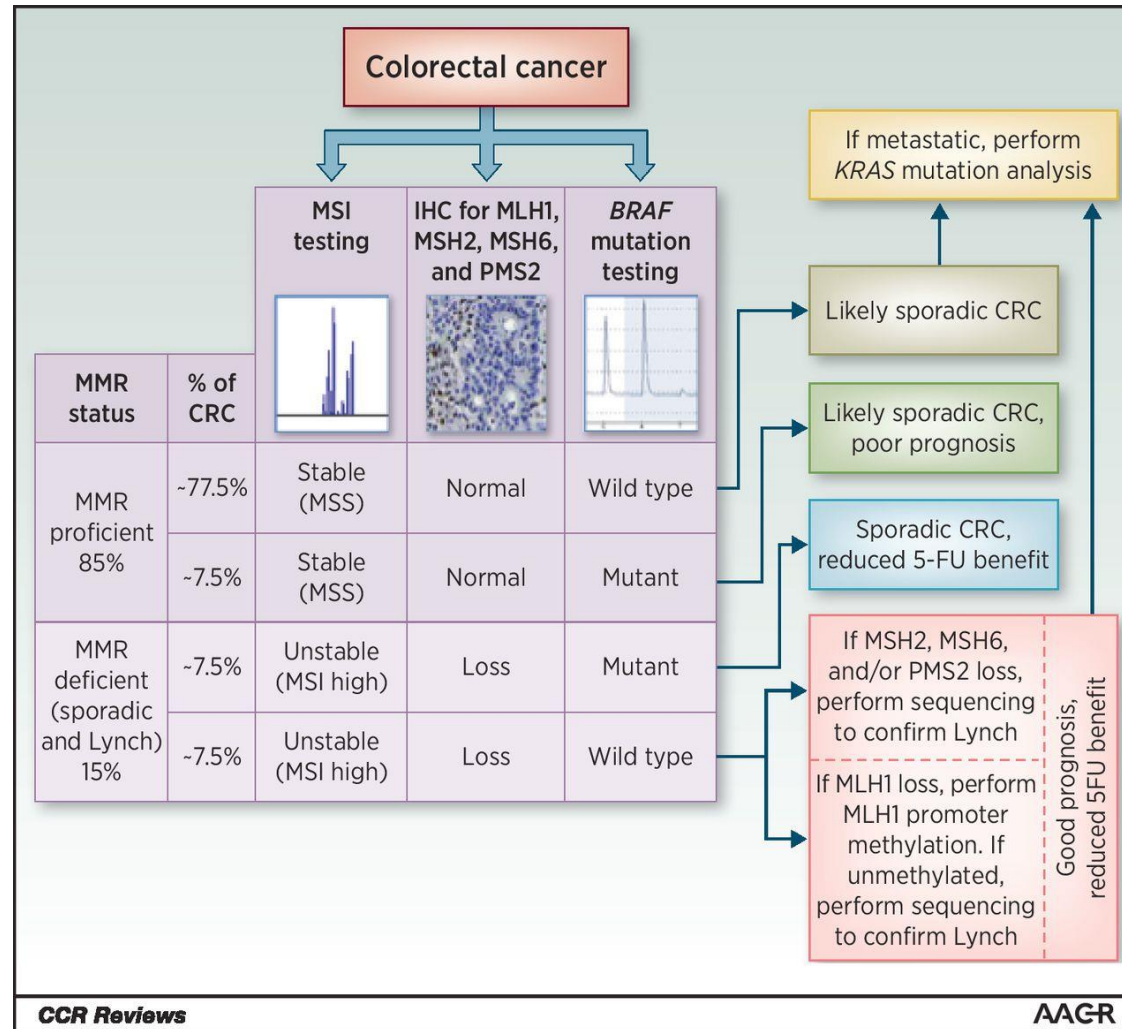


HR<sub>adj</sub> 1.03 (95% CI 0.82-1.30)

HR<sub>adj</sub> 0.17 (95% CI 0.07-0.41)

Interaction p 0.0002

# MSI testing algorithm proposed in 2012 by the Association for Molecular Pathology.



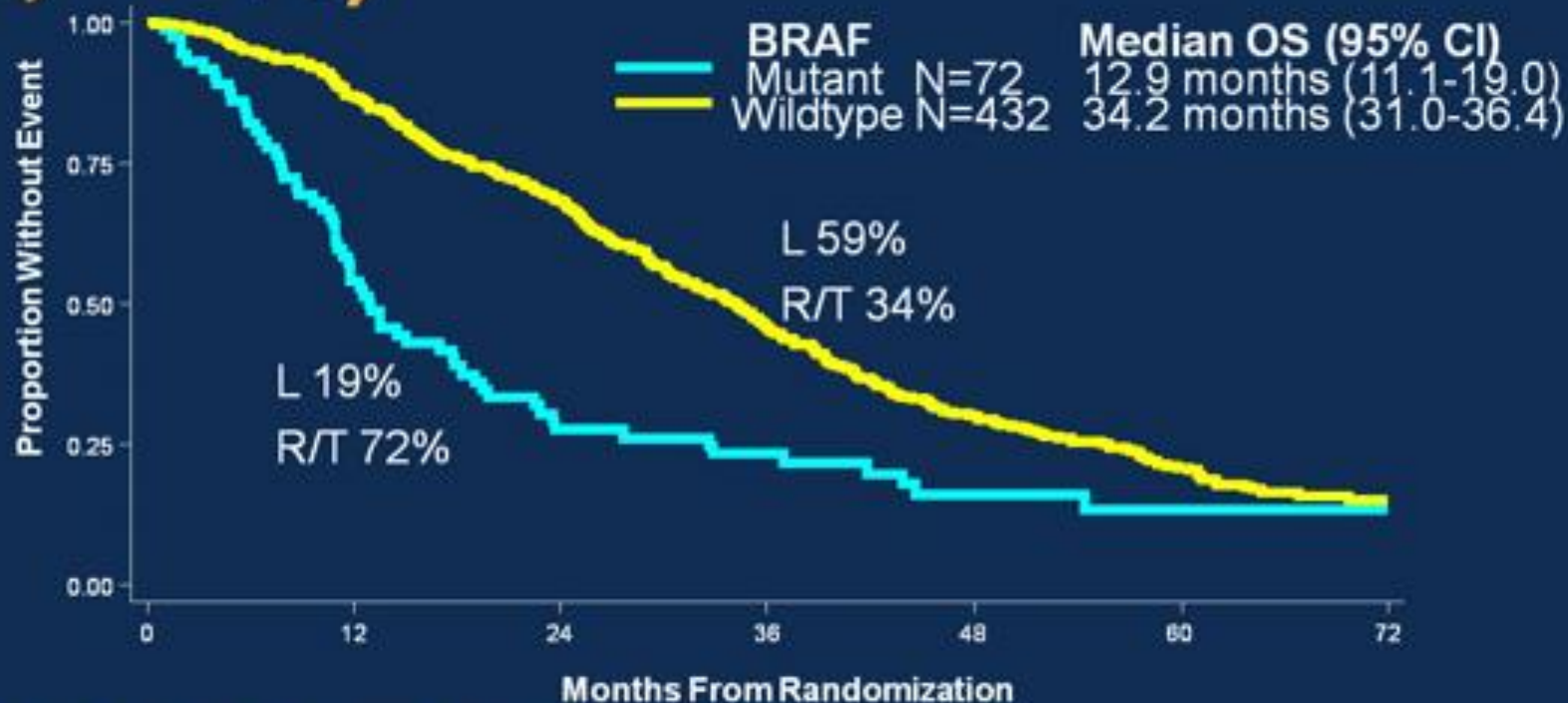
Jonathan C. Dudley et al. Clin Cancer Res 2016;22:813-820

# TOPIC SEVEN

BRAF M+



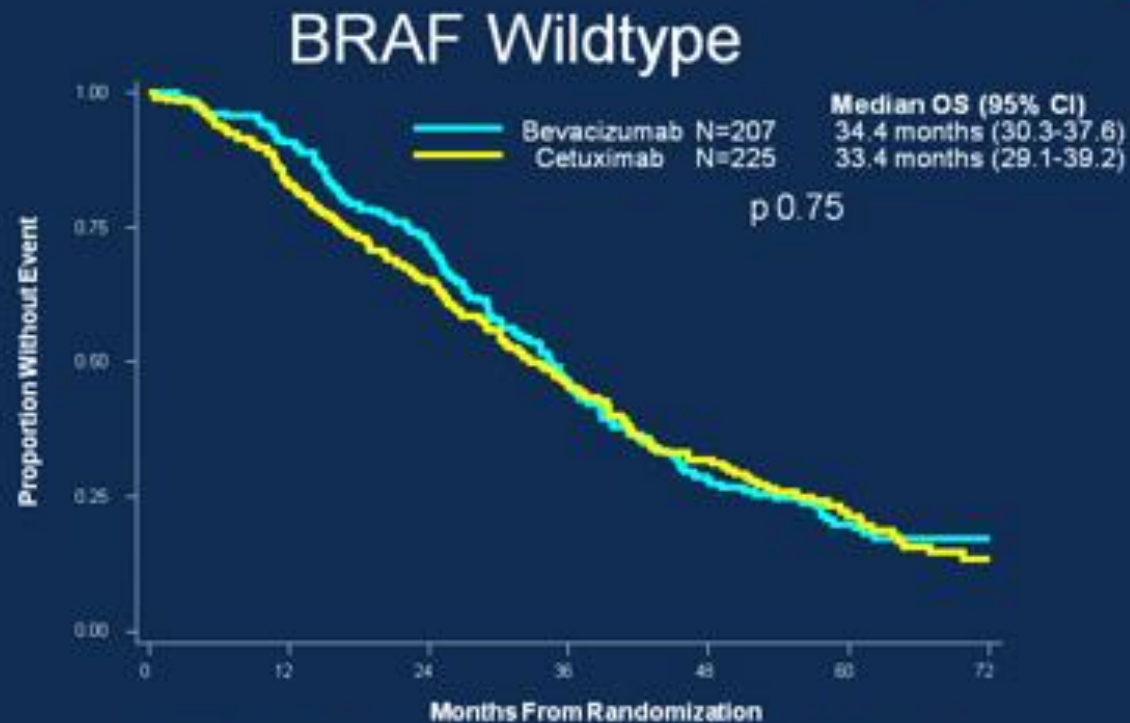
# OS is affected by BRAF mutations (72/504, 14%)



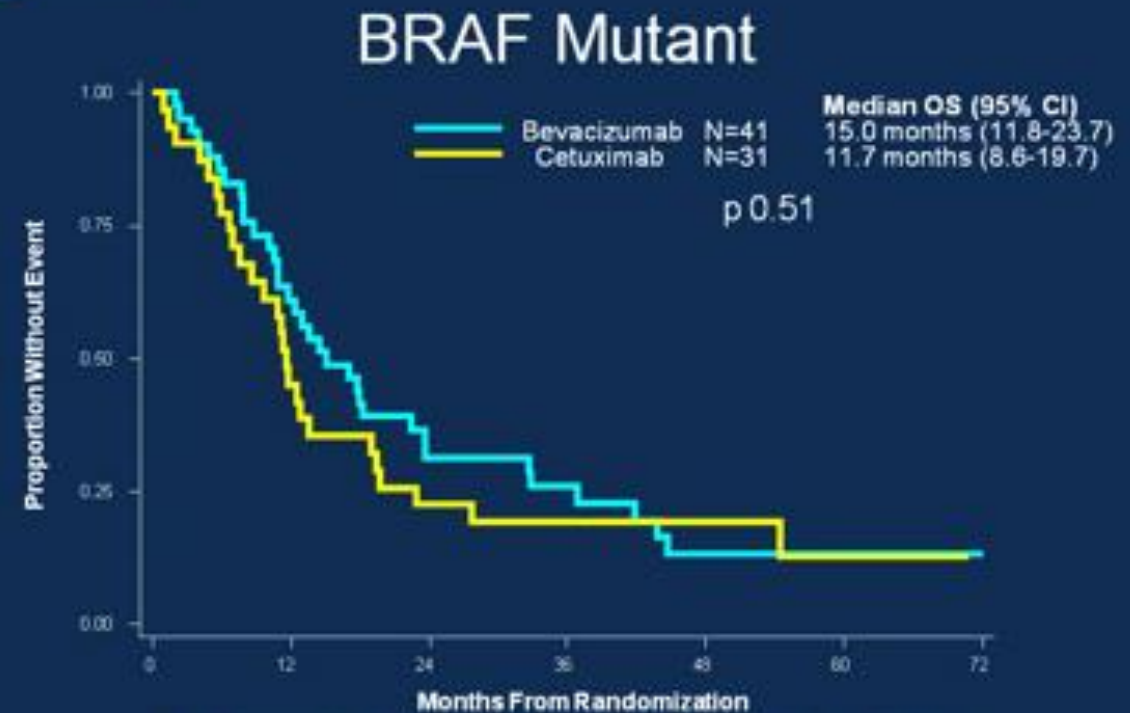
HR<sub>adj</sub> 1.67 (95% CI 1.20-2.33) p 0.0035

Without sidedness: HR<sub>adj</sub> 1.82 (95% CI 1.37-2.44) p 0.0001

# Interaction of BRAF status with biologics



$HR_{adj}$  0.97 (95% CI 0.77-1.23)



$HR_{adj}$  0.61 (95% CI 0.35-1.06)

Interaction p 0.13

# Efficacy Results According to RAS and BRAF Mutation Status in the Primary-Analysis Population.

**PRIME**

**Table 3.** Efficacy Results According to RAS and BRAF Mutation Status in the Primary-Analysis Population.\*

| Variable  | Panitumumab–<br>FOLFOX4 | FOLFOX4<br>Alone | Hazard Ratio<br>(95% CI) | P<br>Value |
|---|-------------------------|------------------|--------------------------|------------|
| <b>No RAS or BRAF mutations</b>                                   |                         |                  |                          |            |
| No. of patients   | 228                     | 218              |                          |            |
| Months of progression-free survival<br>— median (95% CI)          | 10.8 (9.4–12.4)         | 9.2 (7.4–9.6)    | 0.68 (0.54–0.87)         | 0.002      |
| Months of overall survival<br>— median (95% CI)                   | 28.3 (23.7–NE)          | 20.9 (18.4–23.8) | 0.74 (0.57–0.96)         | 0.02       |
| <b>No RAS mutation, BRAF mutation</b>                             |                         |                  |                          |            |
| No. of patients   | 24                      | 29               |                          |            |
| Months of progression-free survival<br>— median (95% CI)          | 6.1 (3.7–10.7)          | 5.4 (3.3–6.2)    | 0.58 (0.29–1.15)         | 0.12       |
| Months of overall survival<br>— median (95% CI)                   | 10.5 (6.4–18.9)         | 9.2 (8.0–15.7)   | 0.90 (0.46–1.76)         | 0.76       |
| <b>RAS or BRAF mutation</b>                                       |                         |                  |                          |            |
| No. of patients   | 296                     | 305              |                          |            |
| Months of progression-free survival<br>— median (95% CI)          | 7.3 (6.3–7.7)           | 8.0 (7.5–9.0)    | 1.24 (1.02–1.49)         | 0.03       |
| Months of overall survival<br>— median (95% CI)                   | 15.3 (12.7–17.6)        | 18.0 (15.9–20.8) | 1.21 (0.99–1.47)         | 0.06       |
| <b>No KRAS mutation in exon 2, other RAS<br/>or BRAF mutation</b> |                         |                  |                          |            |
| No. of patients   | 75                      | 86               |                          |            |
| Months of progression-free survival<br>— median (95% CI)          | 6.7 (5.3–8.2)           | 7.3 (5.7–8.0)    | 1.05 (0.73–1.52)         | 0.80       |
| Months of overall survival<br>— median (95% CI)                   | 14.5 (10.4–18.5)        | 15.8 (11.9–18.8) | 1.14 (0.78–1.66)         | 0.51       |

\* NE denotes not evaluated.

← RAS WT/RAF WT do better, and do benefit from P-mab

← RAF M+ pts do poorly but still marginally benefit from P-mab



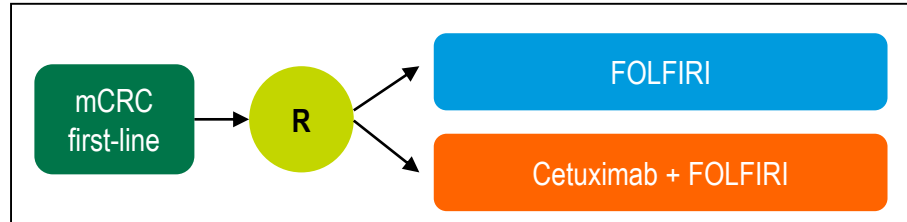


# CRYSTAL Phase III Study

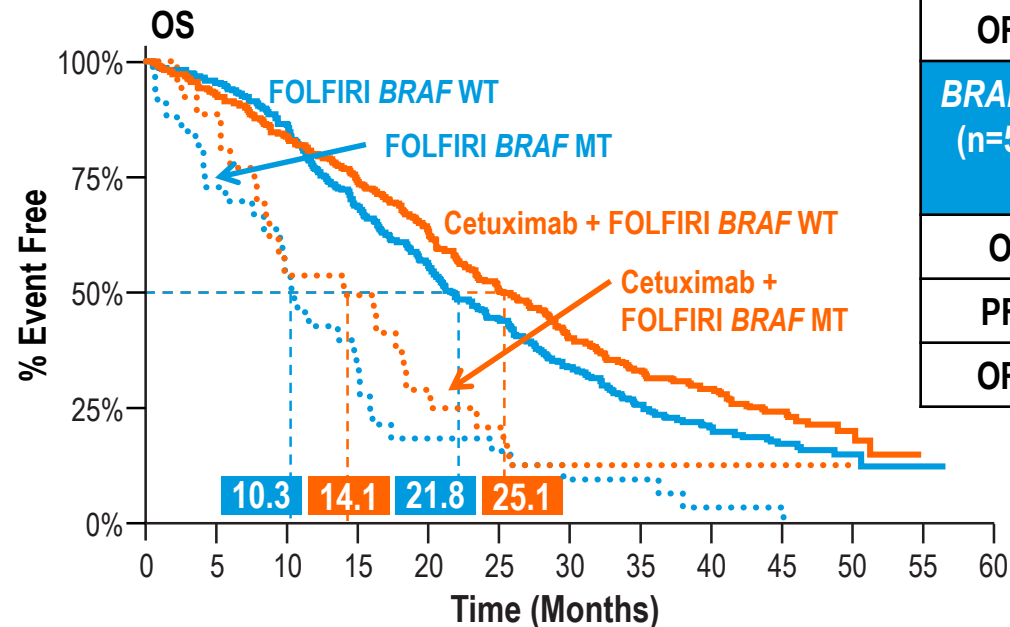
## BRAF Status and Response with Cetuximab

**RAF M+ pts have an overall poor outcome, but still can benefit from EGFR inhibitors (in this case, C-mab)**

### Study Schema



### Results



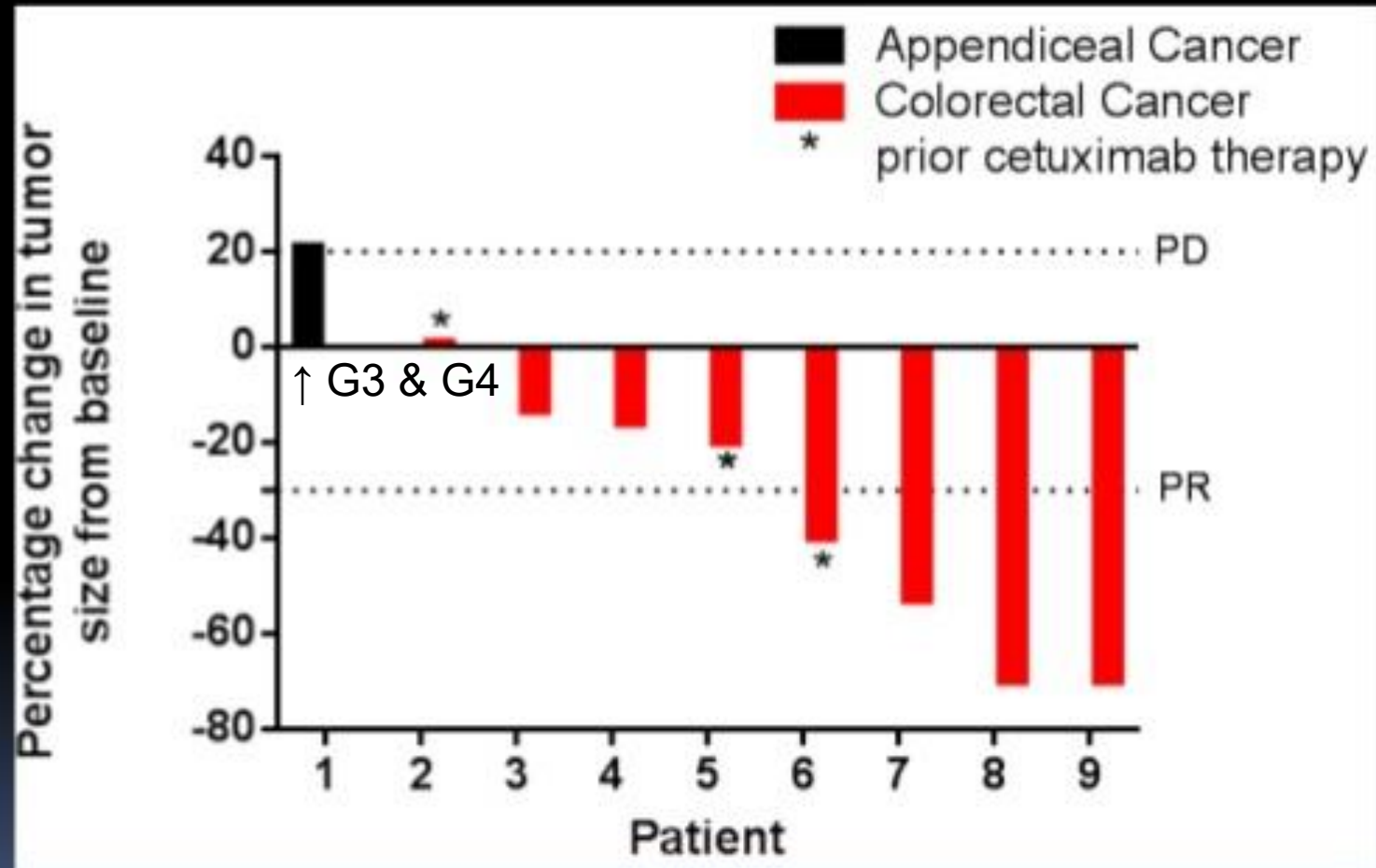
| BRAF MT (n = 59) | Cmab + FOLFIRI (n = 26)  | FOLFIRI (n = 33)  | HR    | p       |
|------------------|--------------------------|-------------------|-------|---------|
| OS               | 14.1                     | 10.3              | 0.908 | 0.74    |
| PFS              | 8.0                      | 5.6               | 0.934 | 0.87    |
| ORR              | 19.2                     | 15.2              | NR    | 0.91    |
| BRAF WT (n=566)  | Cmab + FOLFIRI (n = 277) | FOLFIRI (n = 289) | HR    | p       |
| OS               | 25.1                     | 21.6              | 0.830 | 0.0547  |
| PFS              | 10.9                     | 8.8               | 0.673 | 0.0013  |
| ORR              | 61.0                     | 42.6              | NR    | < 0.001 |

CRYSTAL = Cetuximab Combined With Irinotecan in First-Line Therapy for Metastatic Colorectal Cancer;  
 NR = not reported; ORR = Overall response rate

Adapted from Van Cutsem E, et al. *J Clin Oncol* 2011;29:2011-2019.

# Responses by RECIST 1.1 in all restaged patients

VEMURAFENIB,  
IRINOTECAN  
AND C-Mab in  
BRAF M+ pts



## TRIBE Phase III Study Design

### Patients

- Unresectable mCRC
- No prior mCRC treatment
- Adjuvant oxali-containing chemotherapy allowed if >12 mo between tx and relapse

FOLFIRI + Bev (up to 12 cycles)  
5-FU/LV + Bev (Maintenance)

1:1 Randomization

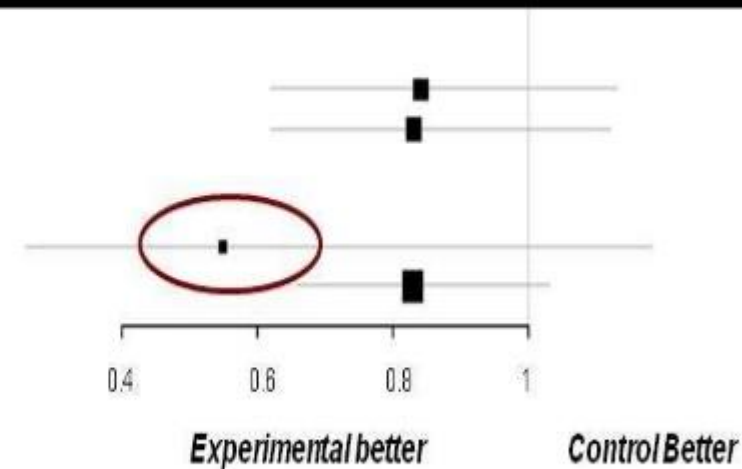
FOLFOXIRI + Bev (up to 12 cycles)  
5-FU/LV + Bev (Maintenance)

Treat to progression

## TRIBE: PFS Subgroup Analyses

### MOLECULAR CHARACTERISTICS

|             |     |      |       |
|-------------|-----|------|-------|
| KRAS status |     |      |       |
| mut         | 200 | 0.84 | 0.973 |
| wt          | 193 | 0.83 |       |
| BRAF status |     |      |       |
| mut         | 28  | 0.55 | 0.323 |
| wt          | 365 | 0.83 |       |



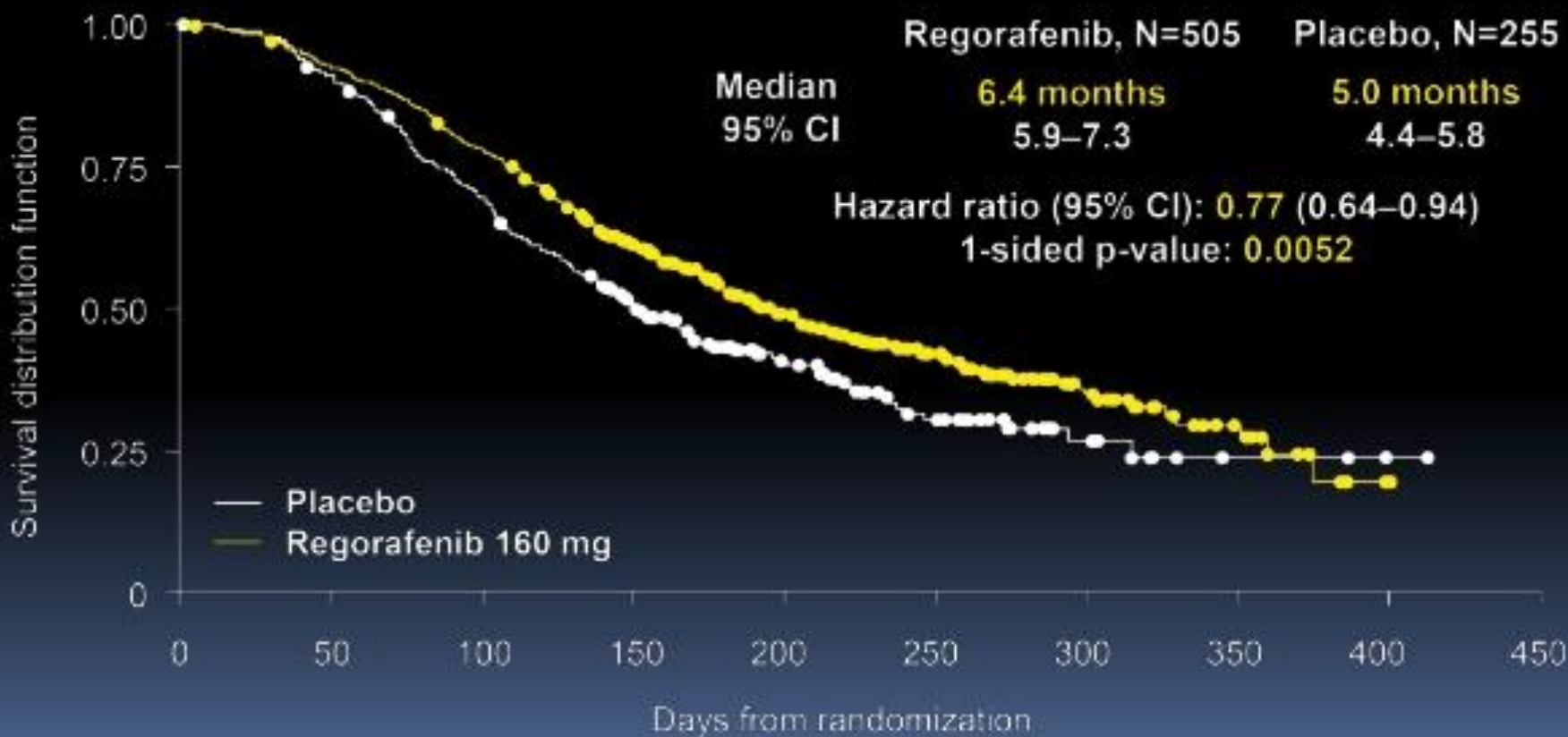
# **TOPIC EIGHT**

## **OTHER NEW DRUGS**

## REGORAFENIB IN REFRACTORY CRCa

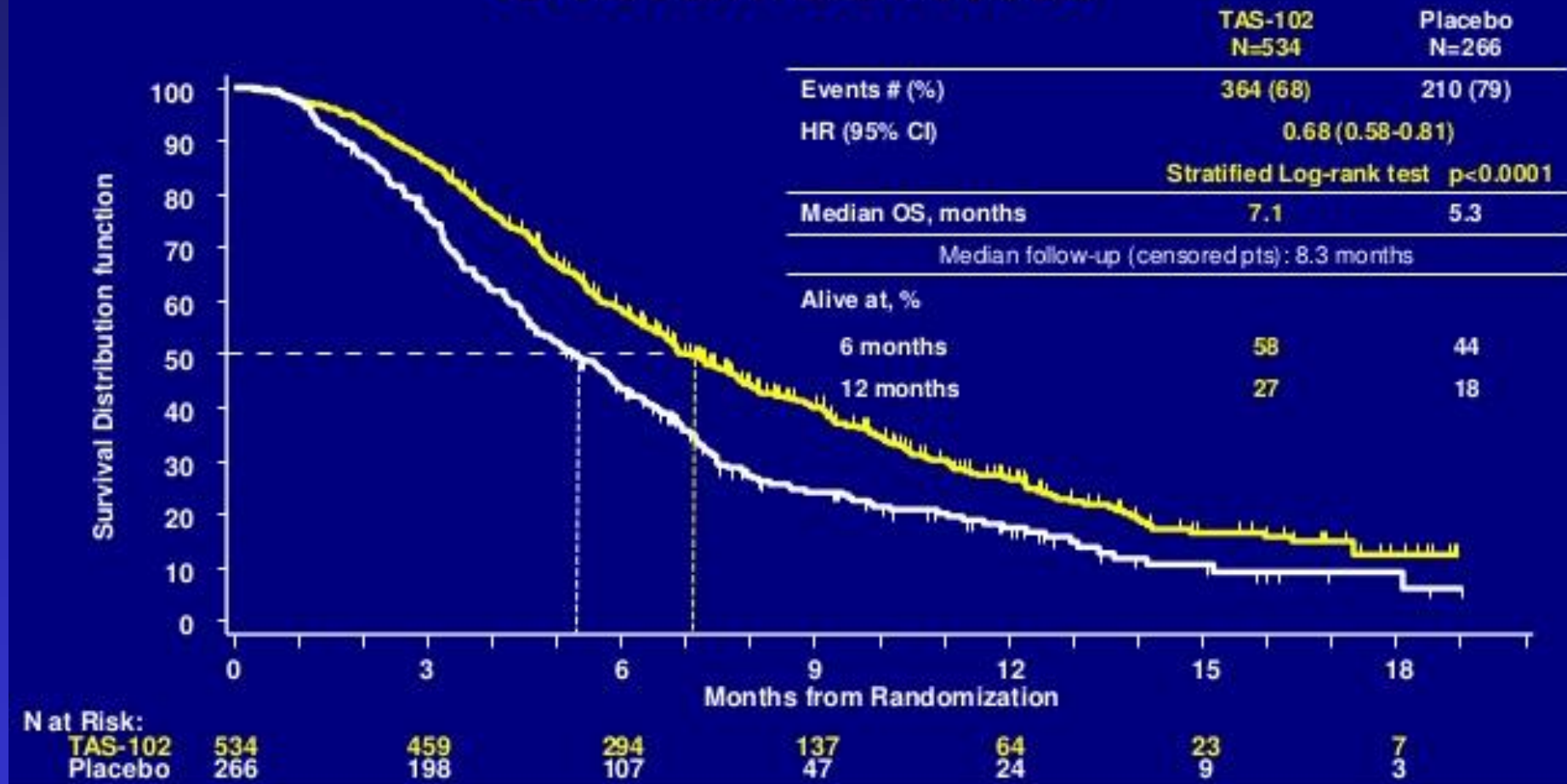
# CORRECT: OS (primary endpoint)

Primary endpoint met prespecified stopping criteria at interim analysis  
(1-sided  $p < 0.009279$  at approximately 74% of events required for final analysis)



# RECOURSE: TAS-102 IN REFRACTORY CRCa

## Overall Survival





# Canadian Colorectal Cancer Consensus Statement

**CURRENT**  
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**PRACTICE GUIDELINE**

## **The predictive effect of primary tumour location in the treatment of metastatic colorectal cancer: a Canadian consensus statement**

A.B.K. Abrahao MD,<sup>#a</sup> S. Karim MD,<sup>†a</sup> B. Colwell MD,<sup>§</sup> S. Berry MD MHSc,<sup>‡</sup> and J. Biagi MD\*



TABLE VI Analysis of primary tumour location in first-line trials comparing EGFR monoclonal antibodies with bevacizumab

| Variable     | CALGB/SWOG 80405 <sup>12</sup><br>(FOLFOX or FOLFIRI with cetuximab<br>vs. FOLFOX or FOLFIRI with bevacizumab) |                |                |               | FIRE-3 <sup>33</sup><br>(FOLFIRI with cetuximab<br>vs. FOLFIRI with bevacizumab) |                |                |               | PEAK <sup>34</sup><br>(FOLFOX with panitumumab<br>vs. FOLFOX with bevacizumab) |               |                |               |
|--------------|--|----------------|----------------|---------------|--|----------------|----------------|---------------|--|---------------|----------------|---------------|
|              | Left-sided   |                | Right-sided    |               | Left-sided   |                | Right-sided    |               | Left-sided   |               | Right-sided    |               |
|              | Cmab<br>(n=173)  | Bev<br>(n=152) | Cmab<br>(n=71) | Bev<br>(n=78) | Cmab<br>(n=157)  | Bev<br>(n=149) | Cmab<br>(n=38) | Bev<br>(n=50) | Pmab<br>(n=53)   | Bev<br>(n=54) | Pmab<br>(n=22) | Bev<br>(n=14) |
| ORR (%)      | 69.4   | 57.9           | 42.3           | 39.7          | 69.0   | 62.0           | 52.6           | 50.0          | 64.2   | 57.4          | 63.6           | 50.0          |
| OR           | 1.65   |                | 1.11           |               | 1.37   |                | 1.11           |               | 1.33   |               | 1.75           |               |
| 95% CI       | 1.16 to 2.34   |                | 0.61 to 2.01   |               | 0.85 to 2.19   |                | 0.48 to 2.59   |               | 0.57 to 3.11   |               | 0.36 to 8.39   |               |
| p Value      | 0.005  |                | 0.73           |               | 0.23   |                | 0.83           |               |  |               |                |               |
| PFS (months) | 12.7   | 11.2           | 7.5            | 10.5          | 10.7   | 10.7           | 7.6            | 9.0           | 14.6   | 11.5          | 8.7            | 12.6          |
| HR           | 0.84   |                | 1.64           |               | 0.90   |                | 1.44           |               | 0.68   |               | 1.04           |               |
| 95% CI       | 0.66 to 1.06   |                | 1.15 to 2.36   |               | 0.71 to 1.14   |                | 0.92 to 2.26   |               | 0.45 to 1.04   |               | 0.50 to 2.18   |               |
| p Value      | 0.15   |                | 0.006          |               | 0.38   |                | 0.11           |               | 0.07   |               | 0.90           |               |
| OS (months)  | 39.3   | 32.6           | 13.9           | 29.2          | 38.3   | 28.0           | 18.3           | 23.0          | 43.4   | 32.0          | 17.5           | 21.0          |
| HR           | 0.77   |                | 1.36           |               | 0.63   |                | 1.31           |               | 0.77   |               | 0.67           |               |
| 95% CI       | 0.59 to 0.99   |                | 0.93 to 1.99   |               | 0.48 to 0.75   |                | 0.81 to 2.11   |               | 0.46 to 1.28   |               | 0.30 to 1.50   |               |
| p Value      | 0.04   |                | 0.10           |               | 0.002  |                | 0.28           |               | 0.31   |               | 0.32           |               |

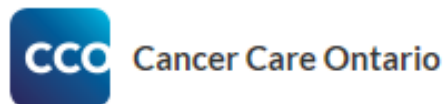
FOLFOX = folinic acid–5-fluorouracil–oxaliplatin; FOLFIRI = folinic acid–5-fluorouracil–irinotecan; Cmab = cetuximab; Bev = bevacizumab; Pmab = panitumumab; ORR = overall response rate; OR = odds ratio; CI = confidence interval; PFS = progression-free survival; HR = hazard ratio; OS = overall survival.

**TABLE VIII** Recommendations in *RAS* wild-type metastatic colorectal cancer (mCRC)

| Scenario        | Recommendation  |
|-----------------|---|
| First line      | <ul style="list-style-type: none"> <li>■ For patients with left-sided disease, standard chemotherapy (FOLFOX or FOLFIRI) in combination with an EGFR mAb (cetuximab or panitumumab) is recommended in the first-line setting.</li> <li>■ For patients with right-sided disease, first-line EGFR monoclonal antibodies are not recommended. The combination of bevacizumab with standard chemotherapy remains the standard of care for these patients.</li> <li>■ Extended <i>RAS</i> testing should be available in a timely manner to allow for the appropriate selection of a biologic for first-line treatment decisions.</li> </ul> |
| Second line     | <ul style="list-style-type: none"> <li>■ There is no evidence to recommend the selective use of EGFR monoclonal antibodies in the second-line setting based on the location of the primary tumour.</li> <li>■ Patients who have not been treated with bevacizumab in the first line should be offered bevacizumab in combination with standard chemotherapy.</li> </ul>   |
| Third line      | <ul style="list-style-type: none"> <li>■ All <i>RAS</i> wild-type patients who have not previously been treated with an EGFR monoclonal antibody should be offered one.</li> </ul>  |
| Tumour response | <ul style="list-style-type: none"> <li>■ At this time, there is insufficient evidence for the selective use of EGFR monoclonal antibodies based on primary tumour location if tumour response is the primary goal of therapy.</li> </ul>  |
| Future research | <ul style="list-style-type: none"> <li>■ Primary tumour location should be factored into the design of future clinical trials in the treatment of <i>RAS</i> wild-type mCRC.</li> <li>■ Given that primary tumour location is a surrogate for more complex biologic mechanisms, ongoing research should try to understand the patient- and tumour-related factors that underlie the differential benefit of biologics that have been observed based on primary tumour location.</li> </ul>  |

FOLFOX = folinic acid–5-fluorouracil–oxaliplatin; FOLFIRI = folinic acid–5-fluorouracil–irinotecan.

# Canadian Consensus Statement endorsed by GI Disease Site Committee of Cancer Care Ontario



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The Role of Primary Tumour Location in the Selection of Biologics for the Treatment of Unresectable Metastatic Colorectal Cancer: An Endorsement of a Canadian Consensus Statement

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### The Role of Primary Tumour Location in the Selection of Biologics for the Treatment of Unresectable Metastatic Colorectal Cancer: An Endorsement of a Canadian Consensus Statement

ID: GL END 2-31 Jul 2018

Type of Content: Guidelines & Advice, Clinical  
Document Status: Current

**Authors:** R. Goodwin, C. Agbassi, E. Kennedy, J. Biagi, R. Wong, S. Welch, S. Berry, and the Gastrointestinal Disease Site Group

## RECOMMENDATIONS

The Gastrointestinal Disease Site Group of Cancer Care Ontario endorses the following recommendations from *The Predictive Effect of Primary Tumour Location in the Treatment of Metastatic Colorectal Cancer: a Canadian Consensus Statement* [1]. Rectal cancer was included in the analysis of left-sided colon cancer. See Abrahao et al [1] for more details on the development of the recommendations and the evidence that supports them.

### 1. First-Line

- a. In patients with *RAS* wild-type left-sided colon cancer, standard chemotherapy (FOLFOX or FOLFIRI) in combination with an EGFR mAb (cetuximab or panitumumab) is recommended in the first-line setting.

#### Qualifying statement

Standard chemotherapy in combination with an EGFR mAb is the preferred option over the current standard of standard chemotherapy and bevacizumab.

- b. In patients with *RAS* wild-type right-sided colon cancer, the use of an EGFR mAb first-line is not recommended. The combination of bevacizumab plus standard chemotherapy remains the standard of care for these patients.

## Guideline Endorsement 2-31

- c. Extended *RAS* testing should be available in a timely manner to allow for the appropriate selection of biologic for first-line treatment decisions.

### 2. Second-Line

- a. At this time, there is no evidence to recommend the selective use of EGFR mAbs in the second-line setting based on PTL.
- b. In the second-line setting, patients who were treated with an EGFR mAb instead of bevacizumab in the first line of therapy can be considered to receive bevacizumab in combination with standard chemotherapy.

### 3. Third-Line

All patients with *RAS* wild-type disease who have not previously been treated with an EGFR mAb should be offered one.

### 4. Tumour Response

At this time, in cases where tumour response is the primary goal of therapy, the evidence is insufficient for the selective use of EGFR mAbs based on PTL.

Does not mention sidedness (yet)

Disallows use of bevacizumab in 2<sup>nd</sup> line after 1<sup>st</sup> line EGFR mabs



Cancer Care Ontario

### 1. New Drug Funding Program and Ministry Funding Announcement

The following drug funding policy will be added under Cancer Care Ontario's (CCO's) *New Drug Funding Program* effective September 1, 2017, according to specific criteria:

| Disease Site     | Drug        | Indication   | New/<br>Revised |
|------------------|-------------|--|-----------------|
| Gastrointestinal | Panitumumab | First-line metastatic colorectal, small bowel, or appendiceal cancer for patients who have contraindications or intolerance to bevacizumab | New             |

- Examples of contraindications or intolerance to bevacizumab include:
  - High risk of bleeding or wound healing issues due to temporal proximity to surgery – recently received or planned for resectable/potentially resectable liver metastases.
  - A history of cardiovascular disease, or established class-specific side effects to bevacizumab such as hypertension, thromboembolic events, atrial fibrillation, as well as, proteinuria, risk of or presence of fistulae, risk of or current GI perforation, primary tumour in place, active bleeding, non-healing wound, ulcer, recent trauma, etc.
- Treatments administered prior to RAS testing will not be reimbursed.
- Patients who use panitumumab in the first line setting will not be eligible for bevacizumab, cetuximab, or panitumumab in later lines of therapy.
- Switches between bevacizumab and panitumumab will only be considered within the first 3 months of starting therapy with either agent, provided there is no disease progression on treatment. Patients will only be approved for one switch (i.e., from bevacizumab to panitumumab or vice versa). A clinic note indicating the reason(s) for switching and contraindication(s) to bevacizumab is required.
- Panitumumab must be used in addition to combination chemotherapy. Single agent treatments will not be reimbursed.

# BASE CASE

## UNRESECTABLE, NO RECENT ADJUVANT, 5FU TOLERANT, FIT, YOUNG, NO C/I TO BEV

| RAS M+                          | RAF M+                                       | RAS WT (R)                      | RAS WT (L)                          | MSI-H                           |                   |
|---------------------------------|--|---------------------------------|-------------------------------------|---------------------------------|-------------------|
| FOLFOX/Bev<br>or<br>FOLFIRI/Bev | FOLFOXIRI/Bev                                | FOLFOX/Bev<br>or<br>FOLFIRI/Bev | FOLFOX/P-mab<br>or<br>FOLFIRI/P-mab | Pembrolizumab                   | 1 <sup>st</sup> L |
| FOLFIRI<br>or<br>FOLFOX         | Pembrolizumab if<br>associated with<br>MSI-H | FOLFIRI<br>or<br>FOLFOX         | FOLFIRI/Bev<br>or<br>FOLFOX/Bev     | FOLFOX/Bev<br>or<br>FOLFIRI/Bev | 2 <sup>nd</sup> L |
| TAS-102<br>or<br>Regorafenib    | Regorafenib<br>or<br>P-mab                   | P-mab<br>or<br>C-mab/irinotecan | TAS -102<br>or<br>Regorafenib       | FOLFIRI<br>or<br>FOLFOX         | 3 <sup>rd</sup> L |
| SoL                             | SoL  | TAS -102<br>or<br>Regorafenib   | SoL                                 | Regorafenib<br>or<br>TAS-102    | 4 <sup>th</sup> L |



# Monitoring and Management of EGFR-related Hypomagnesemia

- Magnesium levels should be monitored closely (*e.g.*, every two weeks)
- Infusions of magnesium sulfate, as well as oral supplementation, may be required
  - Administer 2 g/hr in a solution
  - Can administer up to 10 g over 5 hours
  - Supplements may not always be effective

**AFTER ONC CYCLE OF CHEMO...**

**CXR 30/I/2018**

**CXR 7/III/2018**



1 June 2017



6 Feb 2018





**THE END**

