

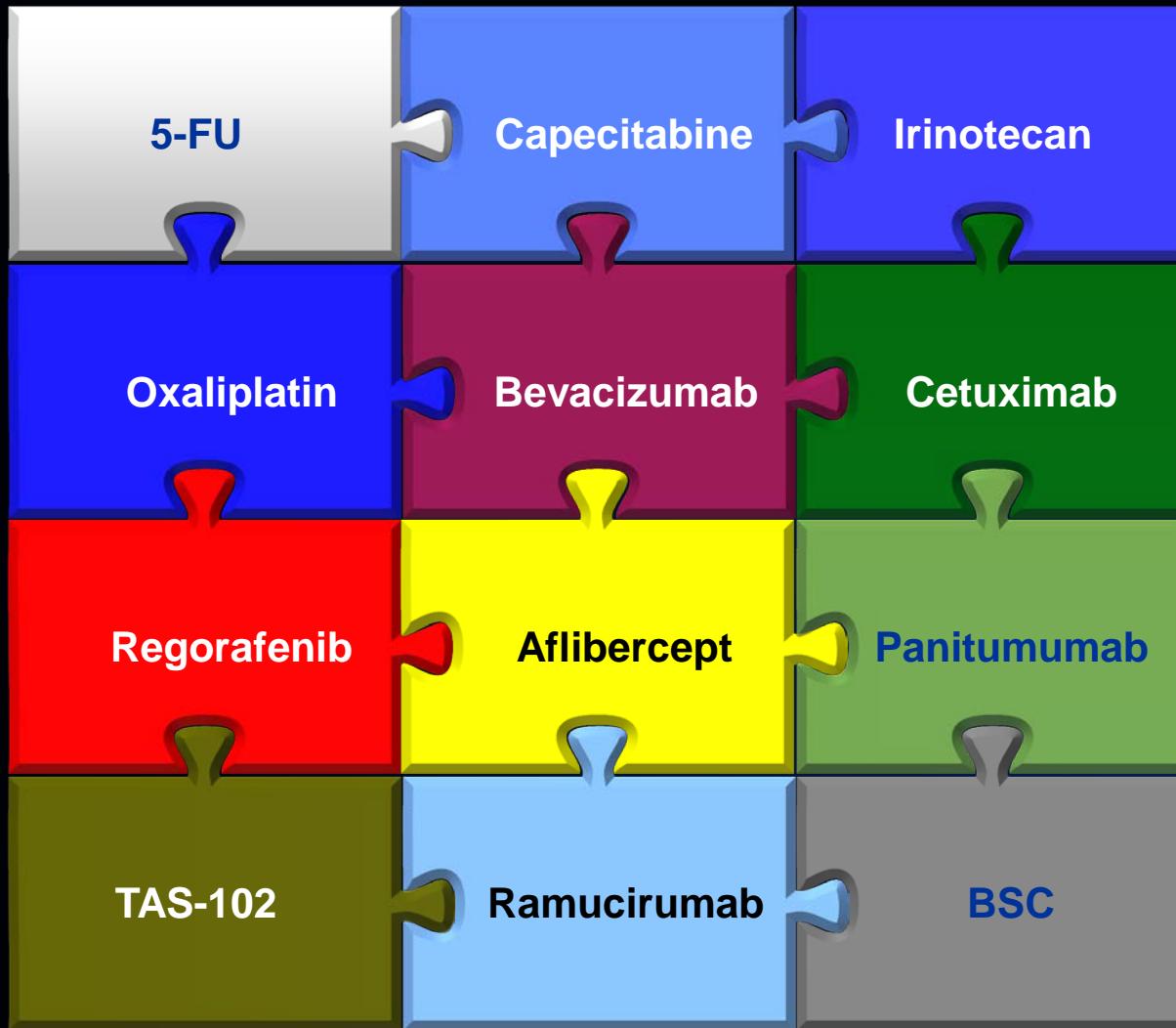
Optimizing treatment outcomes in WT RAS mCRC

A/Prof Niall Tebbutt

Director Medical Oncology

Olivia Newton-John Cancer Research and Wellness Centre

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



The Goals of Systemic Therapy

- Extending OS
- Maintaining quality of life as long as possible
- Tumour response; especially if symptomatic or potentially resectable
 - Consider intensity of therapy/toxicities
 - Consider patient wishes
 - Which situation needs more aggressive and which a more gentle therapy?

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

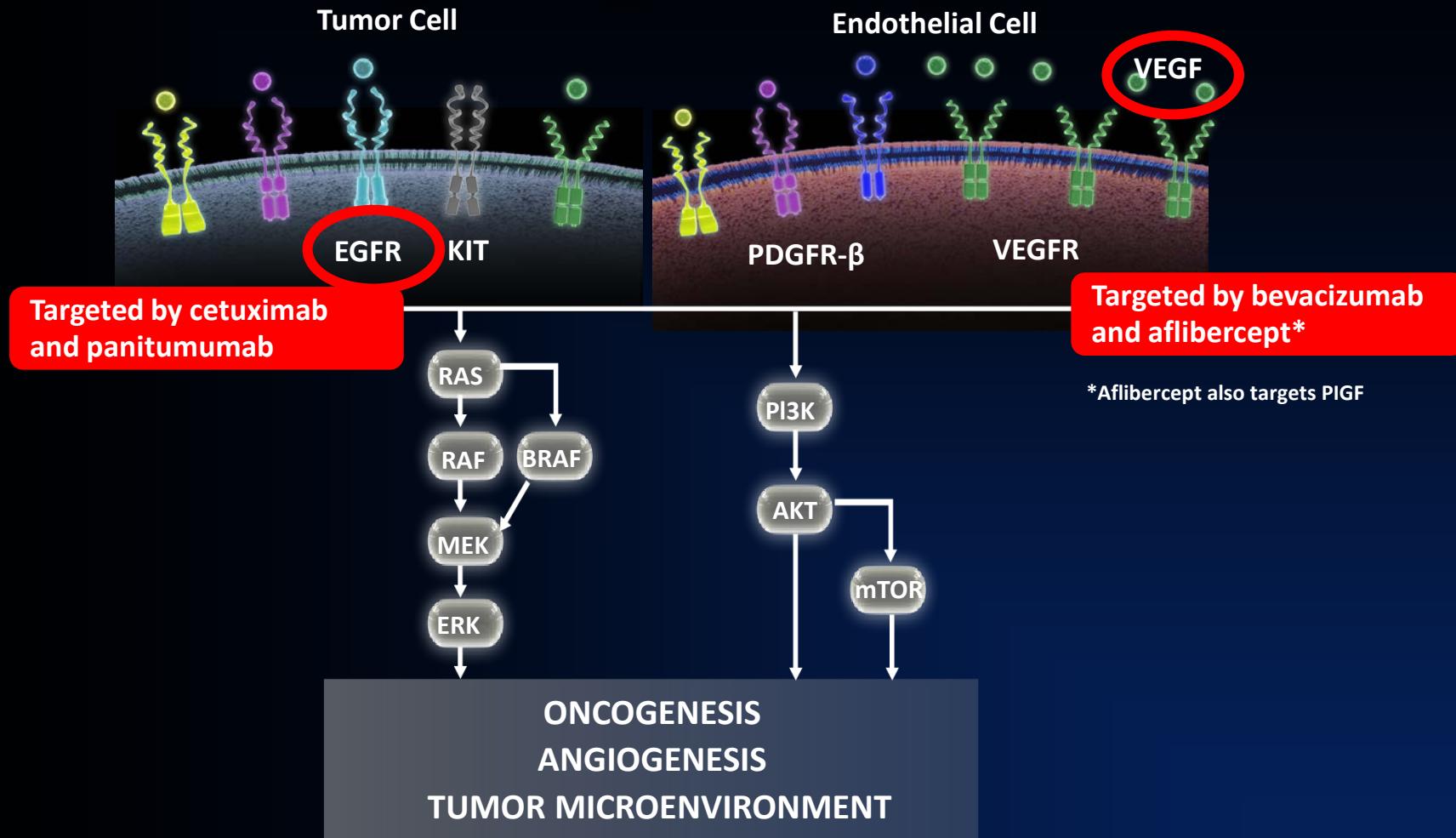
Why personalized medicine?

“

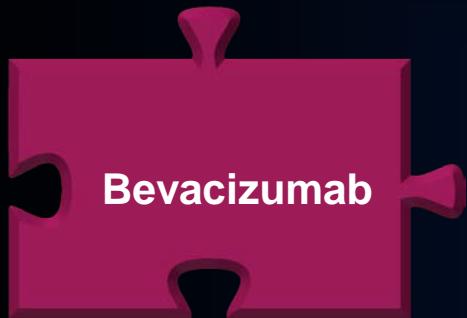
Personalized management is considered the future of cancer care: medicine aiming at giving patients the best treatment according to their personal medical history, their physiological status, and the molecular characteristics of their tumour.”

European Society for Medical Oncology (ESMO)

Overview of EGFR and VEGFR Growth Signaling Pathways



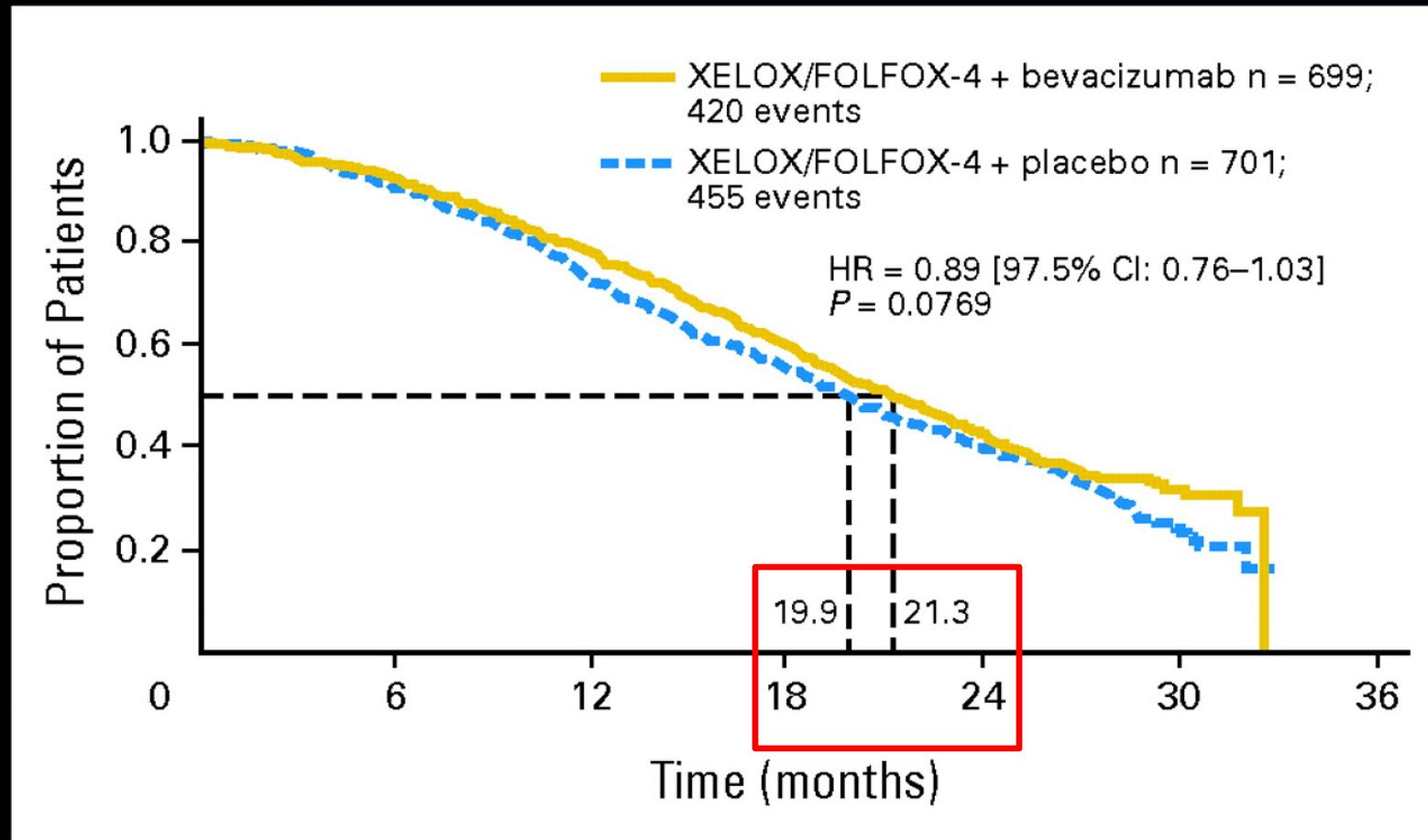
Initial choice of biologic agent for 1st line therapy of mCRC



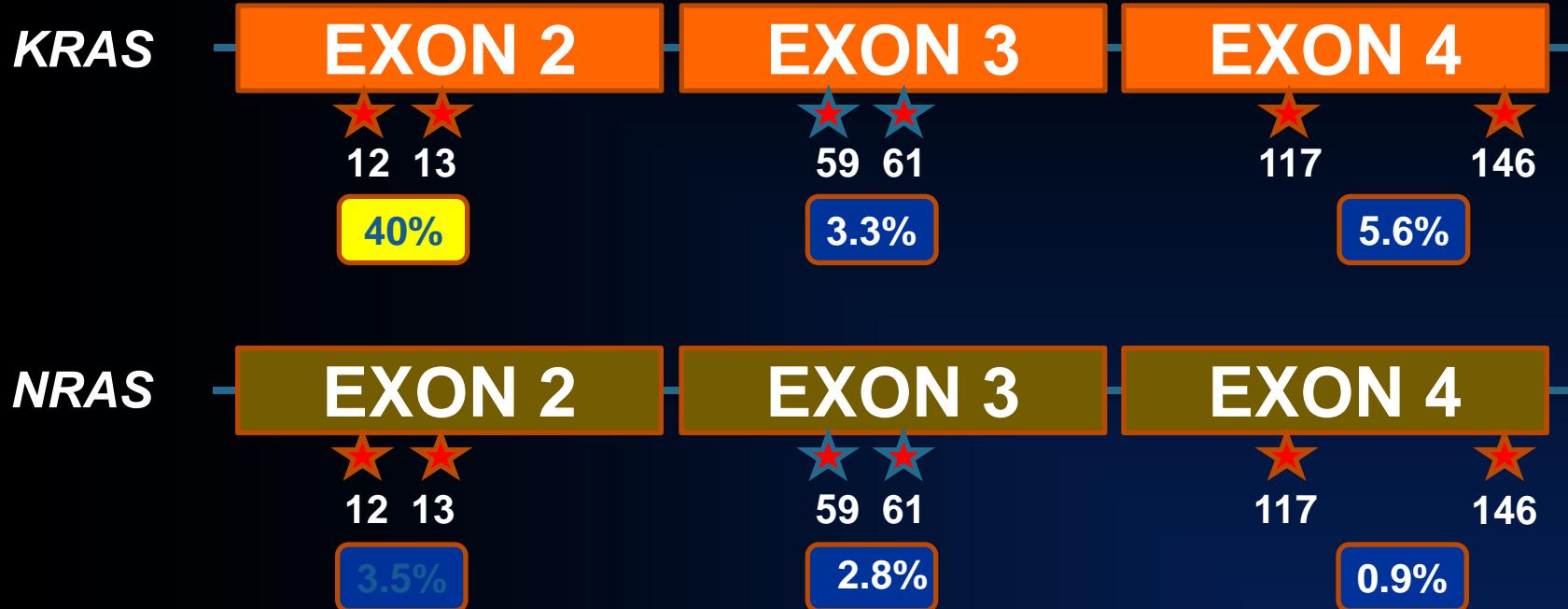
VS



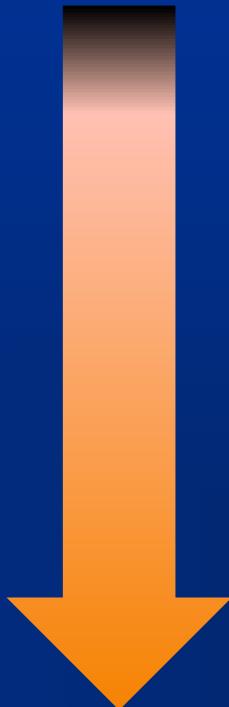
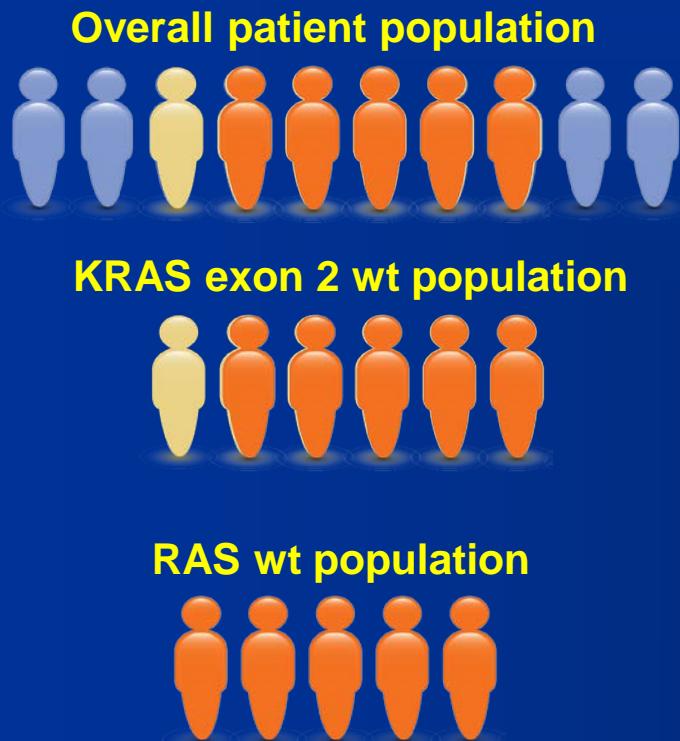
FOLFOX/XELOX and bevacizumab: PFS gain but modest effect on OS



Spectrum of ras mutations



Progress: Improving patient selection extends the benefit with anti-EGFR therapies

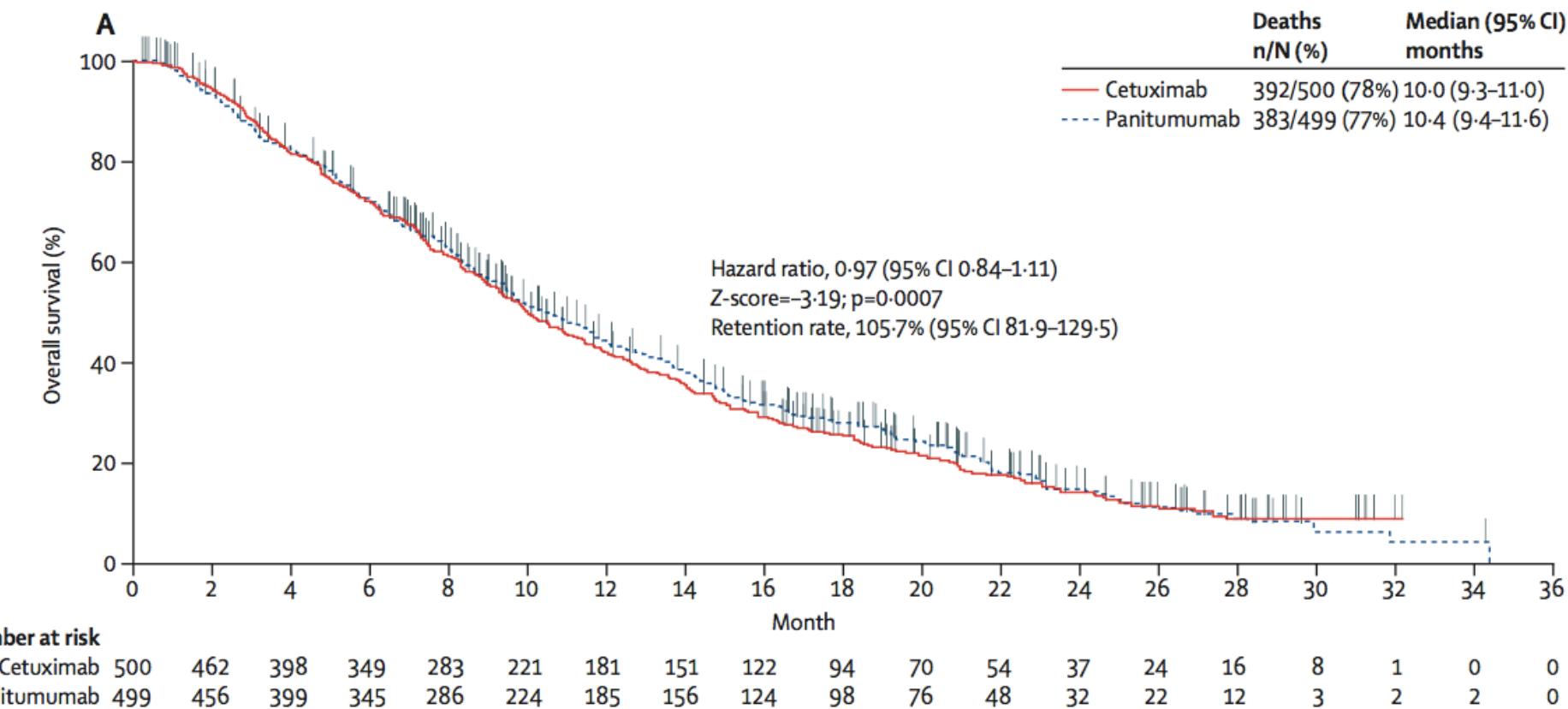


OS benefit*
CRYSTAL^{1,2}
PRIME³
FIRE-3^{4,5}

PFS benefit*
OPUS^{6,7}
PEAK⁸

*Decreased HR in the RAS wt population compared with the KRAS exon 2 wt population or ITT population

ASPECCT panitumumab v cetuximab: OS (non-inferiority)



Main differences between panitumumab and cetuximab relate to safety

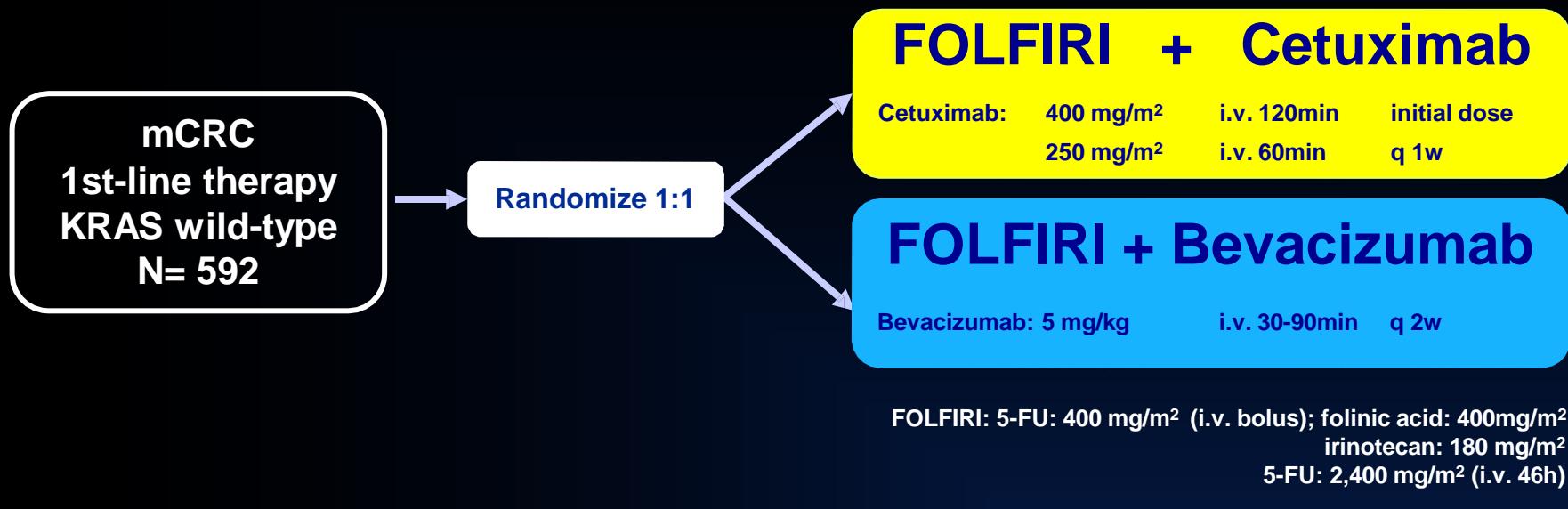
Adverse Event – n (%)	Panitumumab N=496	Cetuximab N =503
Fatal AE's		
Colon cancer	29 (5.8)	50 (9.9)
Others	20 (4.0) 9 (1.8)	34 (6.8) 16 (3.2)
Treatment-related fatal AE's	0 (0)	1 (0.2)
Skin and Subcutaneous tissue AE's		
Any grade	430 (86.7)	440 (87.5)
Grade 3	60 (12.1)	48 (9.5)
Grade 4	2 (0.4)	0 (0)
Serious	1 (0.2)	0 (0)
Hypomagnesemia		
Any grade	143 (28.8)	95 (18.9)
Grade 3	27 (5.4)	10 (2.0)
Grade 4	9 (1.8)	3 (0.6)
Infusion reactions		
Any grade	14 (2.8)	63 (12.5)
Grade 3	1 (0.2)	5 (1.0)
Grade 4	0 (0)	4 (0.8)
Diarrhea		
Any grade	91 (18.3)	89 (17.7)
Grade 3	7 (1.4)	9 (1.8)
Grade 4	3 (0.6)	0 (0.0)

First-line Chemotherapy + EGFR Inhibitor Regimens in mCRC: Efficacy Summary From Phase 3 Trials

Trial	Comparative Regimens	Median PFS, Mos	Median OS, Mos
CRYSTAL ¹	FOLFIRI/Cetux vs FOLFIRI	9.9 vs 8.4	23.5 vs 20.0
OPUS ²	FOLFOX4/Cetux vs FOLFOX4	8.3 vs 7.2	22.8 vs 18.5
PRIME ³⁻⁵	FOLFOX4/Pmab vs FOLFOX4*	9.6 vs 8.0	23.8 vs 19.4
	FOLFOX4/Pmab vs FOLFOX4 (KRAS/NRAS wild-type)*	10.1 vs 7.9	26.0 vs 20.2
COIN ⁶	FOLFOX/XELOX/Cetux vs FOLFOX/XELOX	8.6 vs 8.6	17.0 vs 17.9

Head to head comparisons of chemotherapy + EGFR inhibitor vs chemotherapy + bevacizumab

FIRE-3 study design



- Primary endpoint: **Overall response rate (ORR) (inv assessed)**
- Designed to detect a difference of 12% in ORR induced by FOLFIRI + cetuximab (62%) as compared to FOLFIRI + bevacizumab (50%)
- 284 evaluable patients per arm needed to achieve 80% power for an one-sided Fisher's exact test at an alpha level of 2.5%

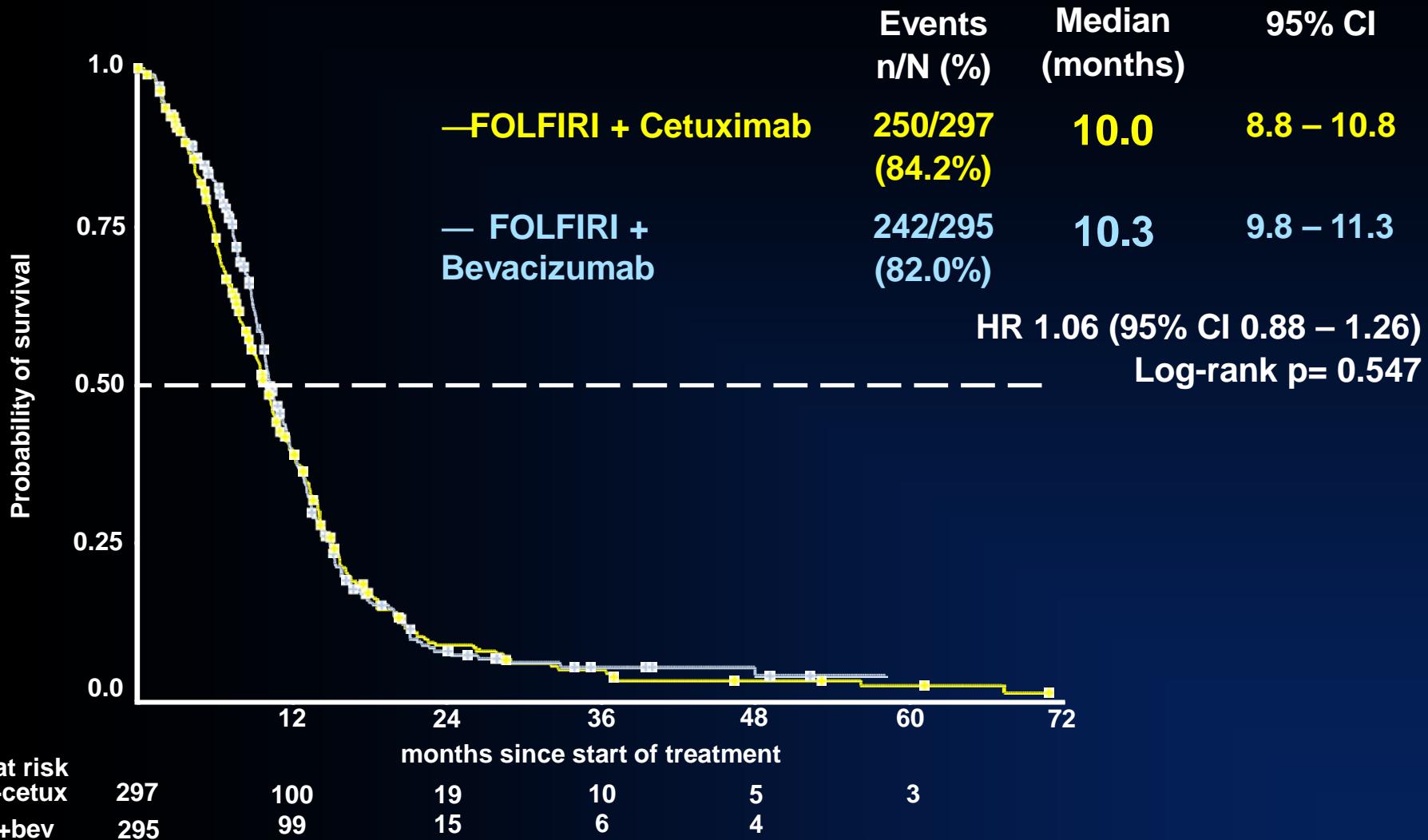
FIRE-3 ORR Primary Endpoint

	FOLFIRI + Cetuximab	FOLFIRI + Bevacizumab	Odds ratio	p	
ORR	%	95%-CI	%	95%-CI	
ITT population (N= 592)	62.0	56.2 – 67.5	58.0	52.1 – 63.7 1.18 0.85-1.64	0.183

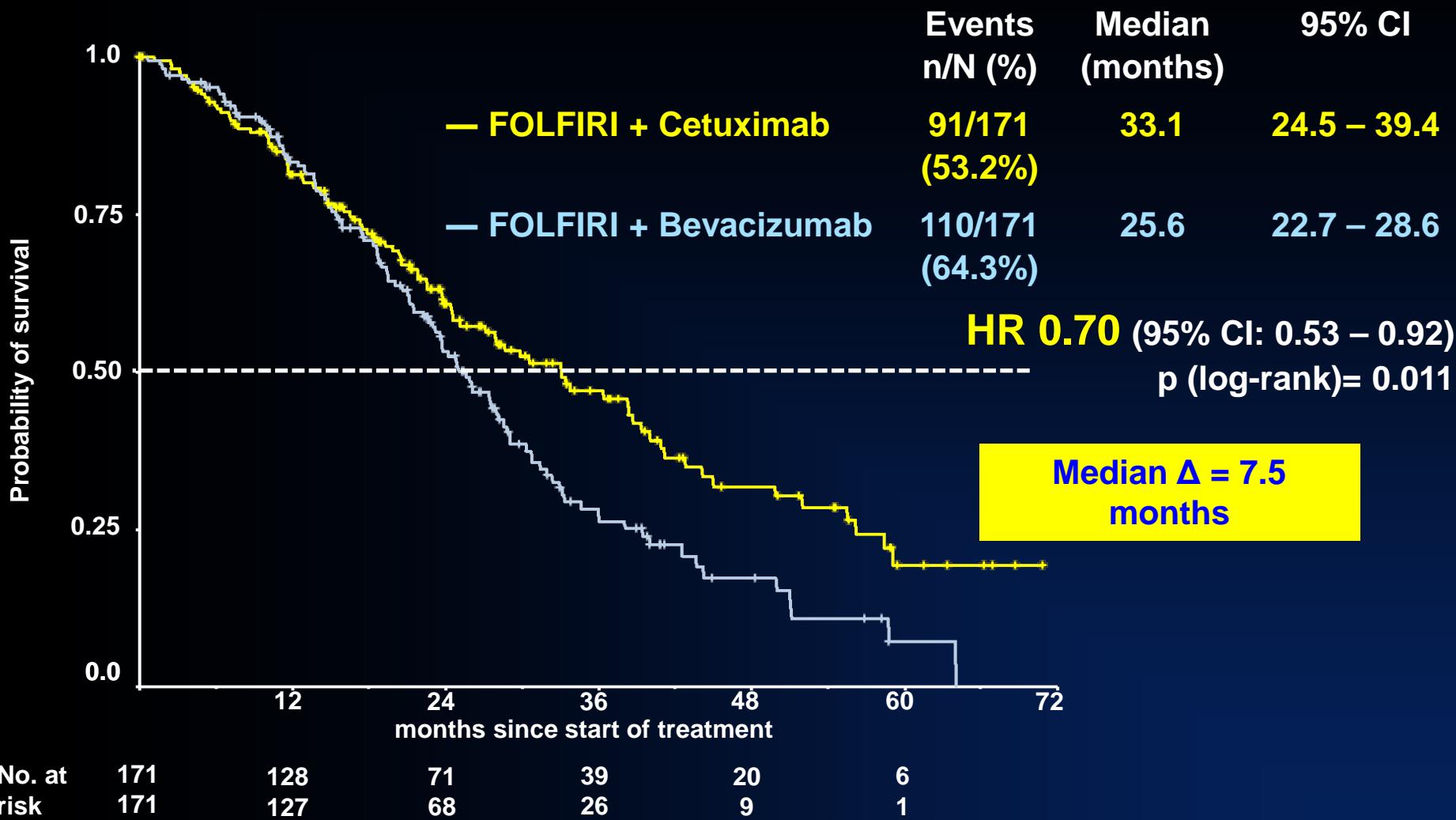
p = Fisher's exact test (one-sided)

Heinemann et al., Lancet Oncol 2014

FIRE-3 PFS



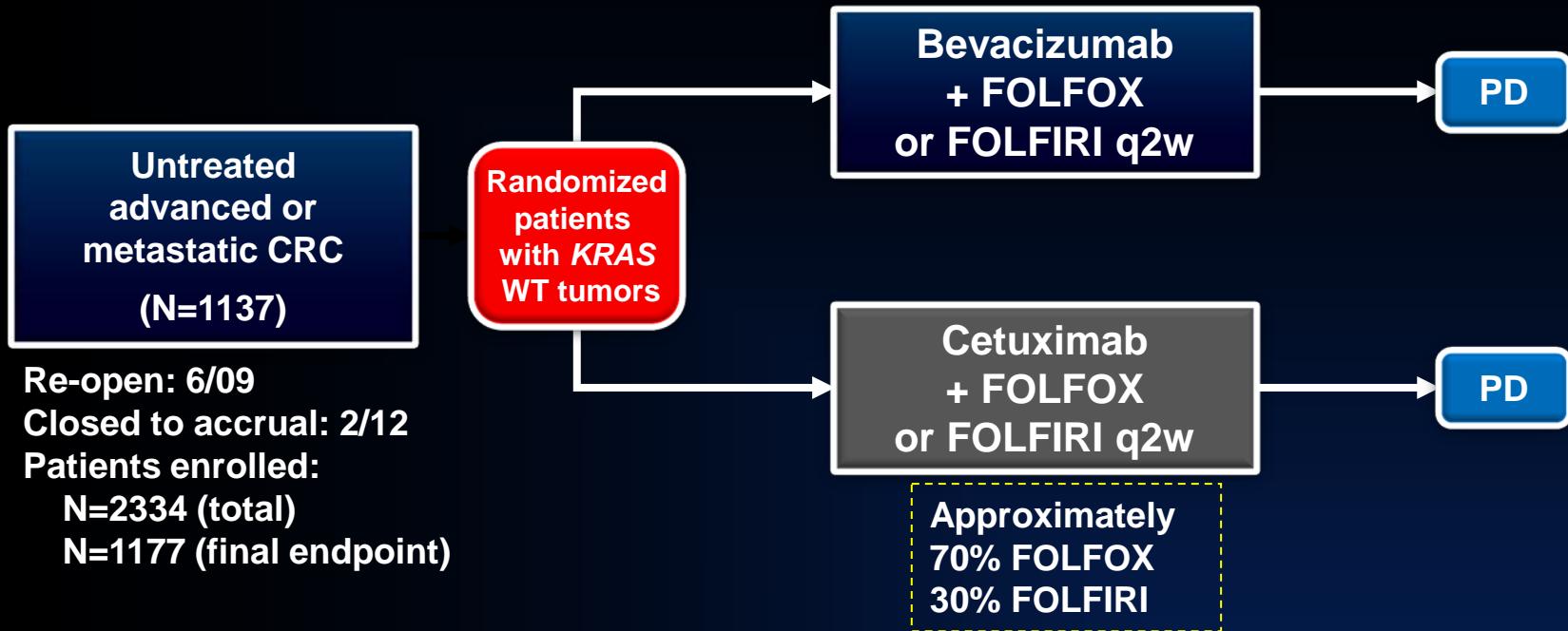
FIRE-3 ESMO/ECCO Update Overall survival All-RAS* wild-type



RAS* wild-type: KRAS 61/146; NRAS Exon2, NRAS Exon3

Heinemann et al., Lancet Oncol 2014

CALGB/SWOG 80405: H2H Bevacizumab vs Cetuximab in First-line *KRAS* WT mCRC

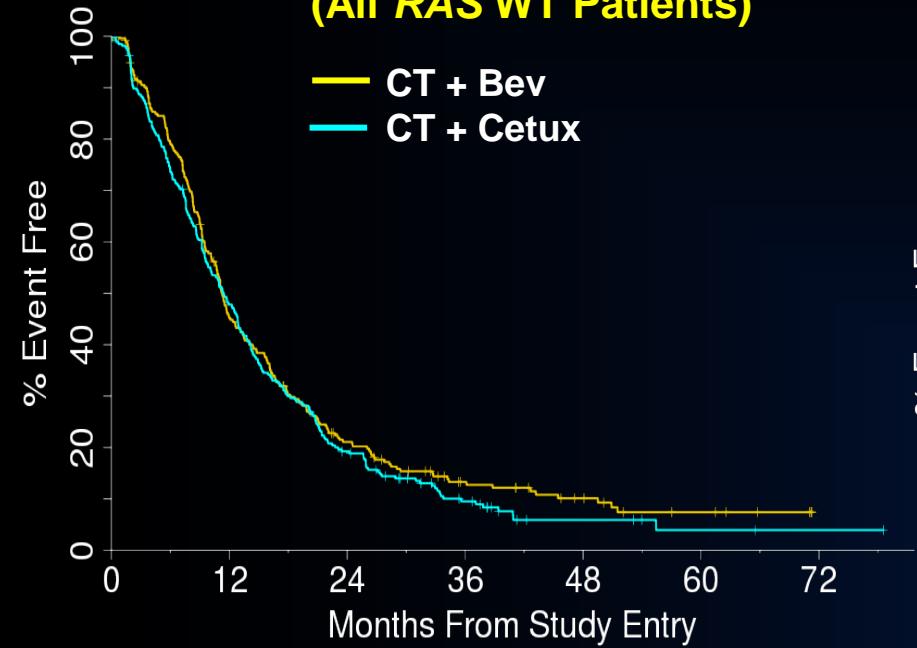


- **Primary endpoint: OS**
 - Superiority trial with 90% power to detect an OS HR of 1.25 (2-sided $\alpha=0.05$)
- **Secondary endpoints: ORR, PFS, TTF, DOR, and safety**

CALGB/SWOG 80405: PFS and OS in All RAS WT Patients

Progression-Free Survival
(All RAS WT Patients)

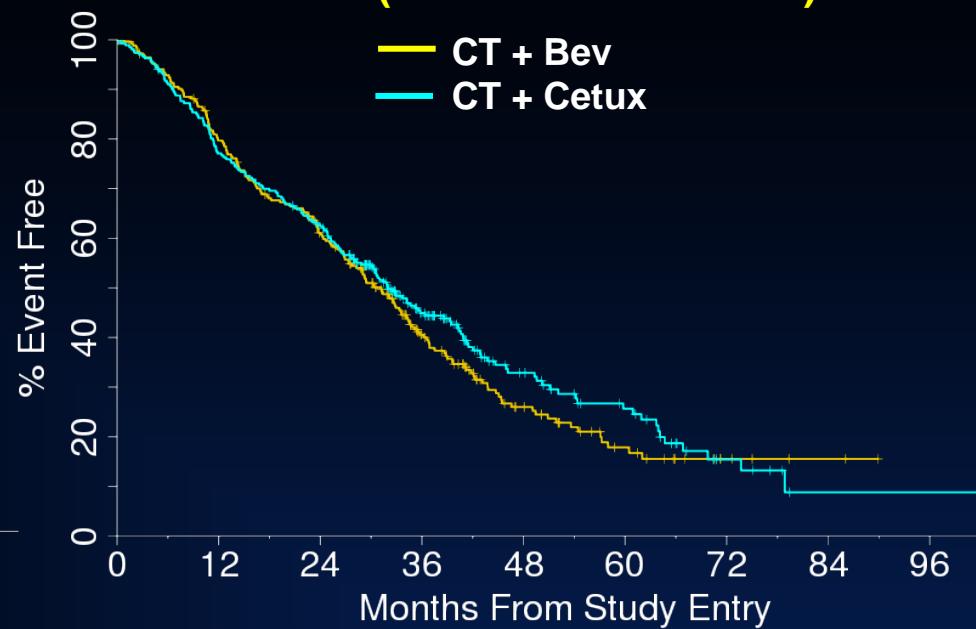
— CT + Bev
— CT + Cetux



# At Risk	256	112	49	23	13	6	2	1
# At Risk	270	126	49	18	5	2		

Overall Survival
(All RAS WT Patients)

— CT + Bev
— CT + Cetux



# At Risk	256	199	147	77	35	16	5	2	1
# At Risk	270	205	164	88	41	24	7	1	1

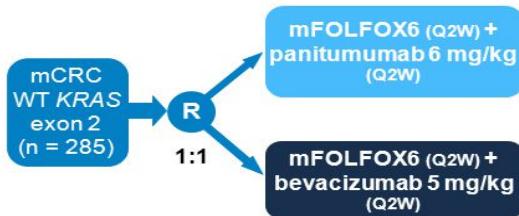
Arm	N (Events)	mPFS, Mos	HR (95% CI) P Value
CT + Bev	256 (221)	11.3 (10.3-12.6)	1.1 (0.9-1.3)
CT + Cetux	270 (241)	11.4 (9.6-12.9)	P=0.31

Arm	N (Events)	mOS, Mos	HR (95% CI) P Value
CT+ Bev	256 (178)	31.2 (26.9-34.3)	0.9 (0.7-1.1)
CT + Cetux	270 (177)	32.0 (27.6-38.5)	P=0.40

PEAK Study

Phase 2 PEAK study

mFOLFOX6 + panitumumab or bevacizumab in 1st-line treatment of WT KRAS exon 2 mCRC



The diagram illustrates the study design. A box labeled "mCRC WT KRAS exon 2 (n = 285)" has an arrow pointing to a central circle labeled "R". From this central circle, two arrows branch out to two separate boxes: "mFOLFOX6 (Q2W) + panitumumab 6 mg/kg (Q2W)" and "mFOLFOX6 (Q2W) + bevacizumab 5 mg/kg (Q2W)". The ratio "1:1" is indicated between the two treatment boxes.

WT RAS ¹	Panitumumab + mFOLFOX6 (n = 88)	Bevacizumab + mFOLFOX6 (n = 82)
Median PFS, mo*	13.0	9.5
HR (95% CI) P-value	0.65 (0.44–0.96) P = 0.029	
Median OS, mo†	41.3	28.9
HR (95% CI) P-value	0.63 (0.39–1.02) P = 0.058	
ORR, %* (95% CI)	63.6 (52.7–73.6)	60.5 (49.0–71.2)

• Primary endpoint: PFS

- No planned formal hypothesis testing

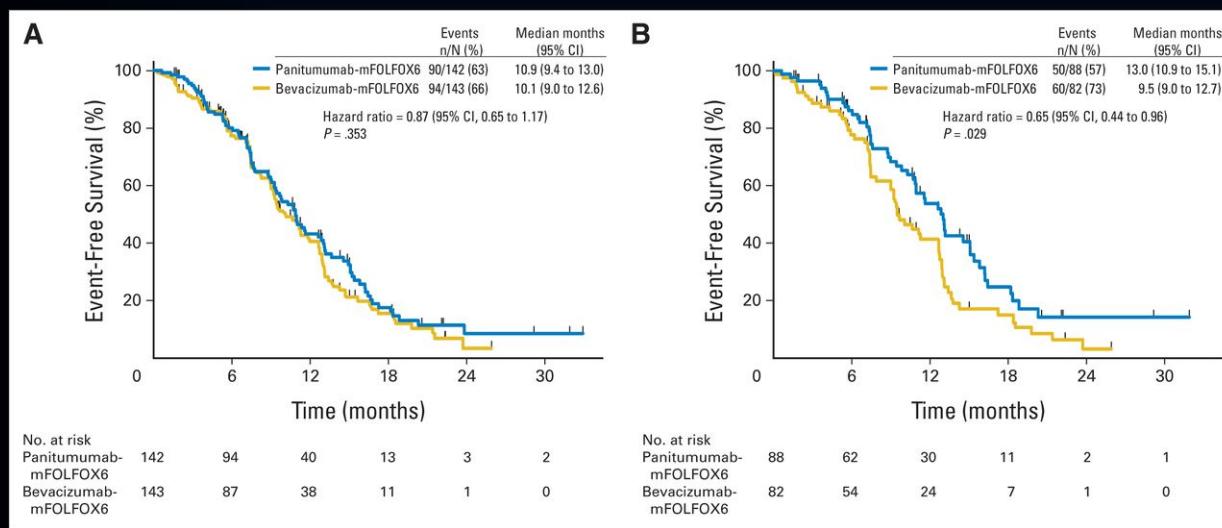
• Prespecified extended RAS analysis

1. Schwartzberg LS, et al. J Clin Oncol 2014;32:2240–7;
Protocol ID: 20070509; ClinicalTrials.gov identifier: NCT00819780.

*Primary analysis; †longer follow-up analysis.
RAS ascertainment rate: 82%.
WT RAS = WT KRAS/NRAS exons, 2, 3, 4.

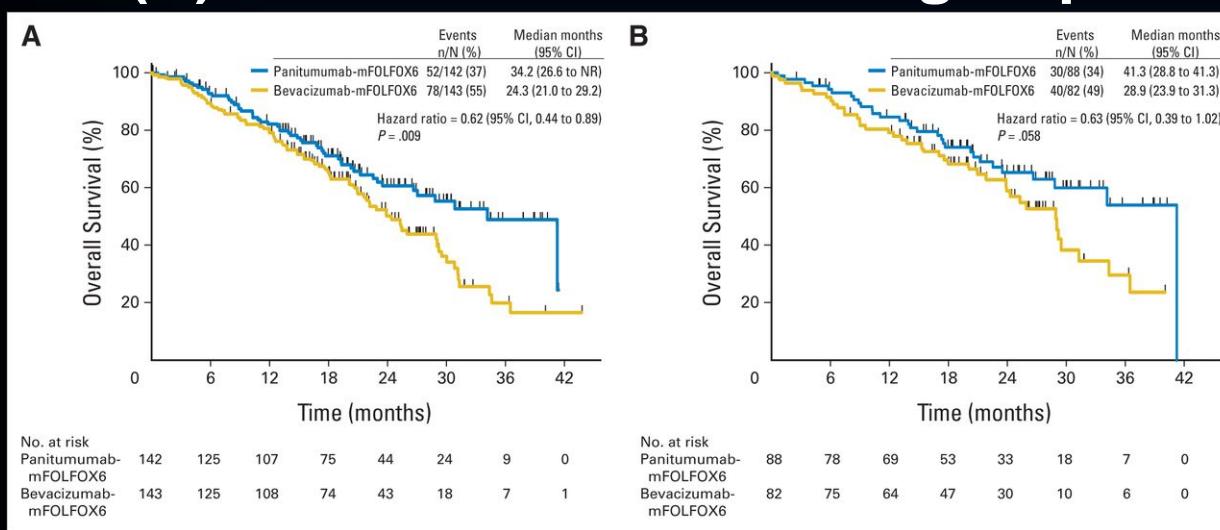
Presented by:

PEAK: Progression-free survival in (A) wild-type (WT) KRAS exon 2 intent-to-treat group and (B) extended WT RAS subgroup.



Lee S. Schwartzberg et al. JCO 2014;32:2240-2247

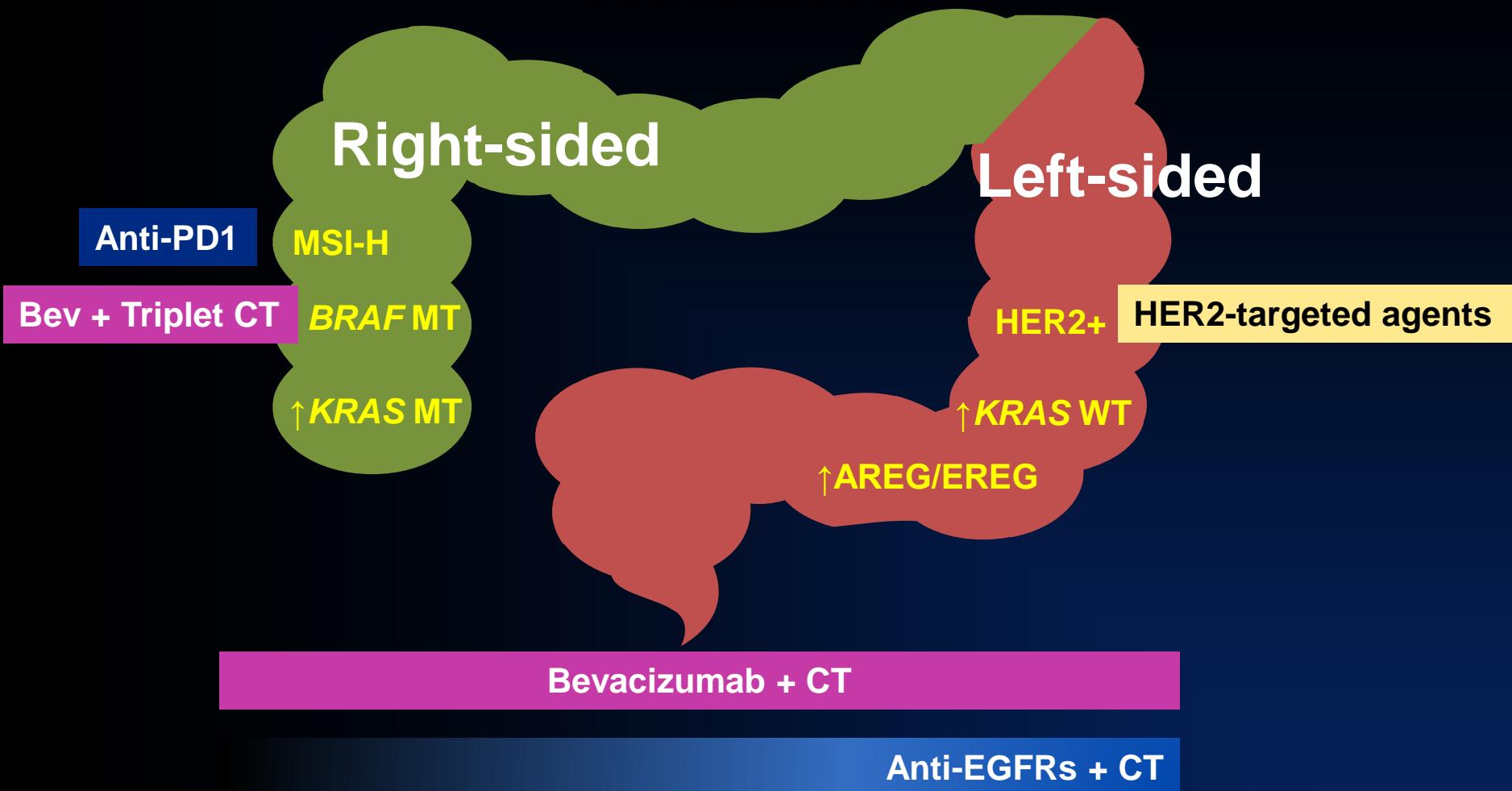
PEAK: Overall survival in (A) wild-type (WT) KRAS exon 2 intent-to-treat group and (B) extended WT RAS subgroup.



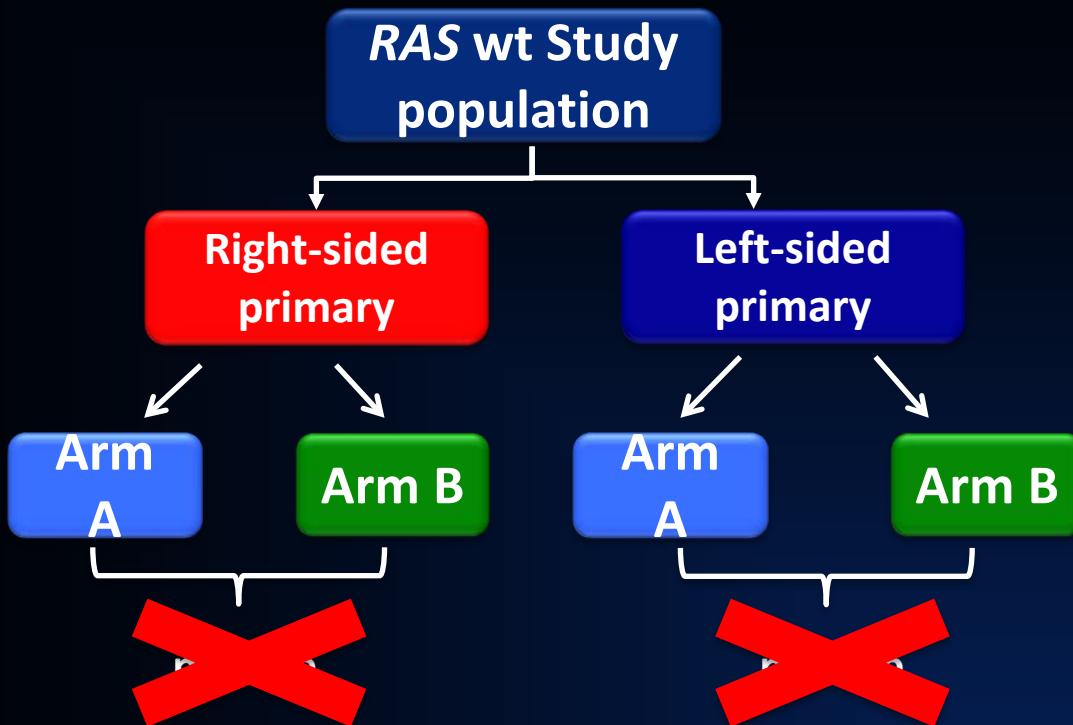
Lee S. Schwartzberg et al. JCO 2014;32:2240-2247

What is the impact of site of primary tumour?

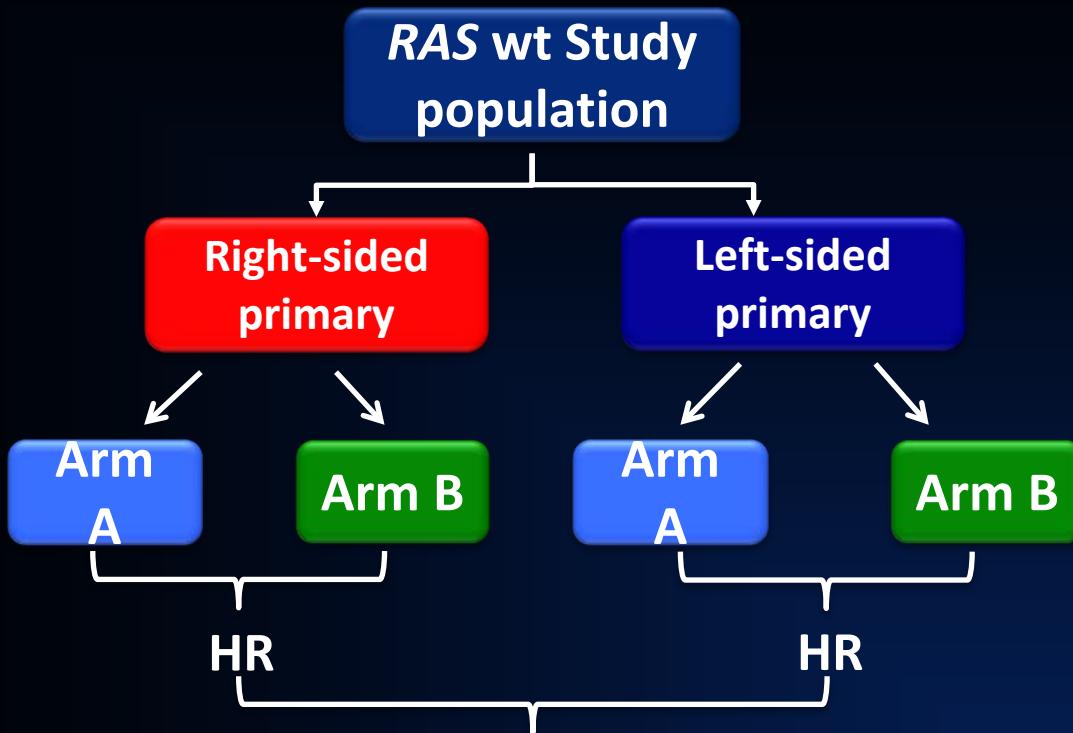
Primary Tumor Location and Potential Treatments



How to interpret subgroup analyses



How to interpret subgroup analyses



p for interaction

If significant ($p < 0.10$) → hypothesis-generating

Right versus Left: summary

1) Prognostic impact

Right-sided tumours have worse prognosis

Prognostic impact

JAMA Oncology | Original Investigation

Prognostic Survival Associated With Left-Sided vs Right-Sided Colon Cancer A Systematic Review and Meta-analysis

Fausto Petrelli, MD; Gianluca Tomasello, MD; Karen Borgonovo, MD; Michele Ghidini, MD; Luca Turati, MD; Pierpaolo Dallera, MD; Rodolfo Passalacqua, MD; Giovanni Sgroi, MD; Sandro Barni, MD

1.437.846 patients in 66 studies (stage I→IV)

HR (left vs right): 0.82 [95%CI: 0.79-0.84], p<0.001

“... Left sided primary tumor location was associated with a significantly reduced risk of death (HR, 0.82; 95% CI, 0.79-0.84; $P < .001$) and this was independent of stage, race, adjuvant chemotherapy, year of study, number of participants, and quality of included studies ... ”

Right versus Left: summary

- 1) Prognostic impact

Right-sided patients have worse prognosis

STRATIFY CLINICAL TRIALS

- 2) Predictive impact

Predictive impact – bevacizumab

Chemo +/- bevacizumab



*p for interaction OS=0.38;
PFS=0.59*

*p for interaction OS=0.29;
PFS=0.62*

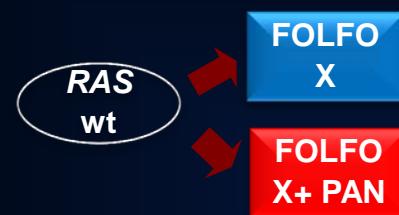
Predictive impact – anti-EGFRs

Chemo +/- anti-EGFR

CRYSTAL



PRIME



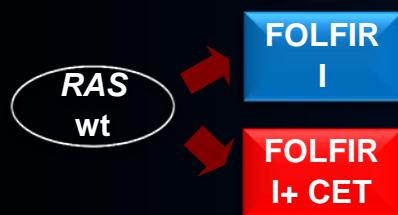
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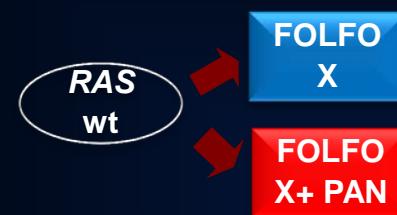
Predictive impact – anti-EGFRs

Chemo +/- anti-EGFR

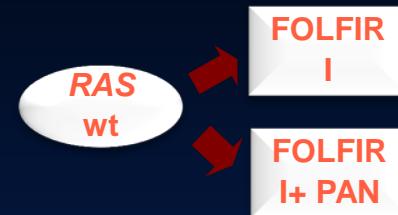
CRYSTAL



PRIME

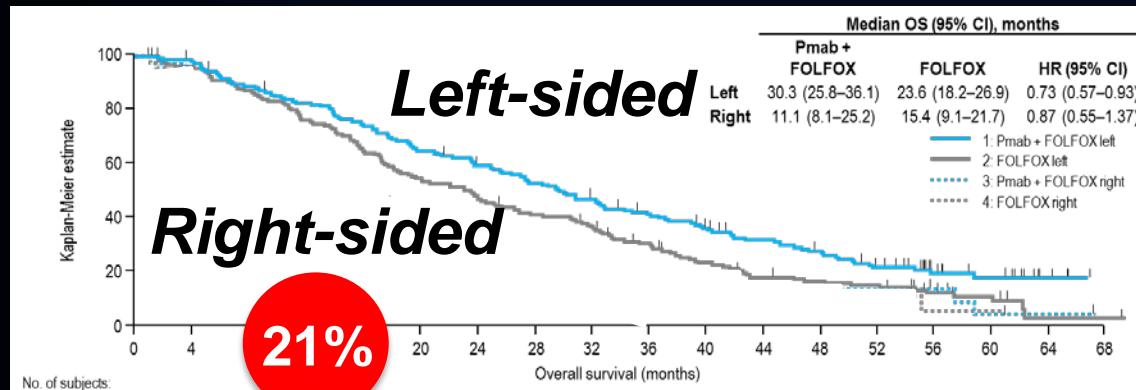


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Right versus Left: PRIME study - OS

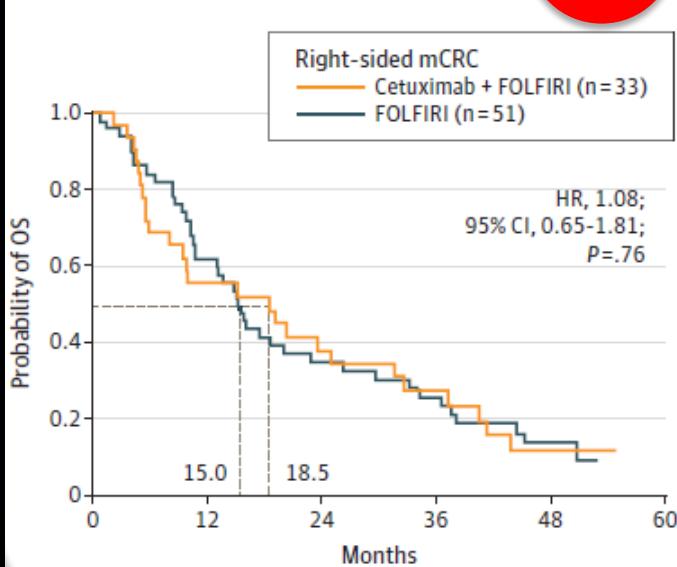
PRIME Trial: FOLFOX+PAN vs FOLFOX



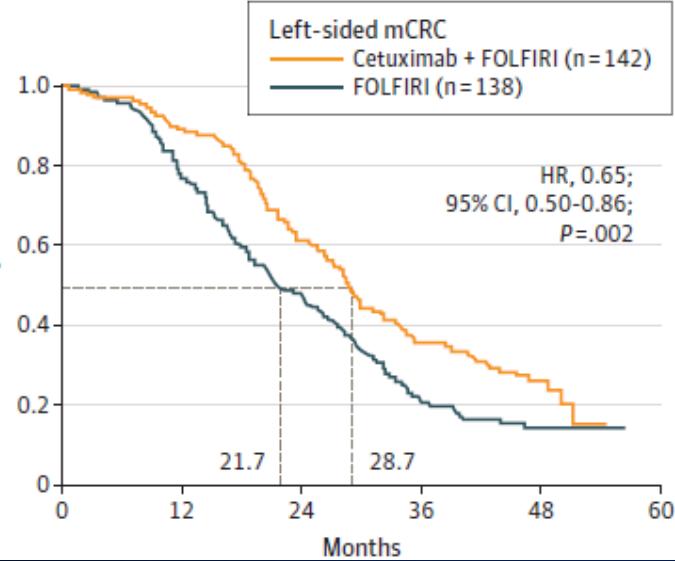
Right versus Left: CRYSTAL study - OS

CRYSTAL Trial: FOLFIRI + cetuximab vs FOLFIRI

Right-sided 23%



Left-sided



p for interaction 0.17

Right versus Left: summary

1) Prognostic impact

Right-sided tumours have worse prognosis

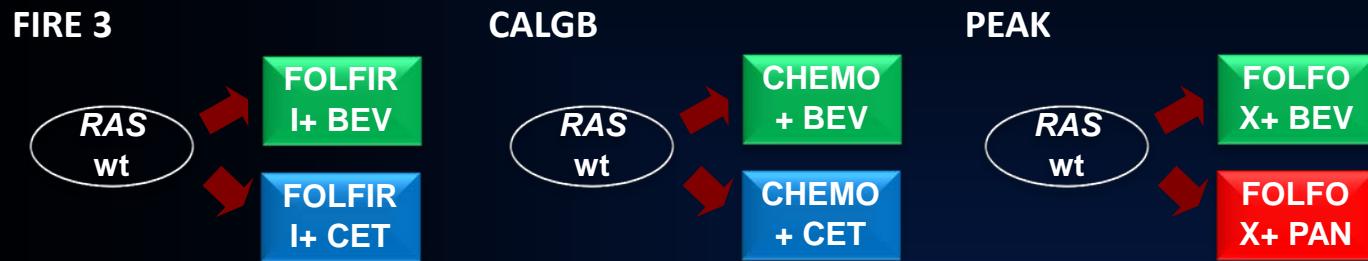
2) Predictive impact

No different benefit from bev for right versus left

*More benefit from anti-EGFRs in left-sided than right-sided mCRC
tumours*

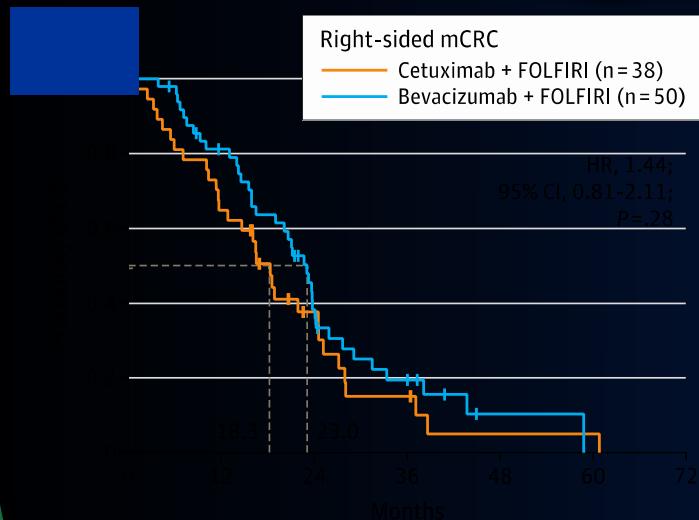
Predictive impact – bev versus anti-EGFRs

Chemo+anti-EGFR vs Chemo+Bev

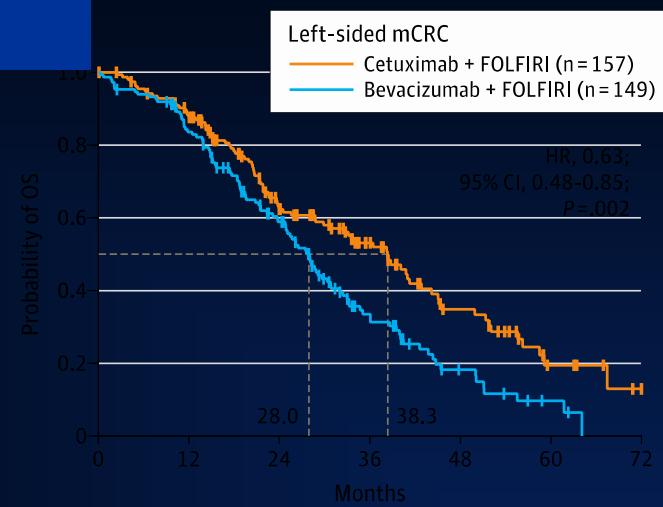


Right versus Left: FIRE-3 study - OS

Right-sided 22%



Left-sided

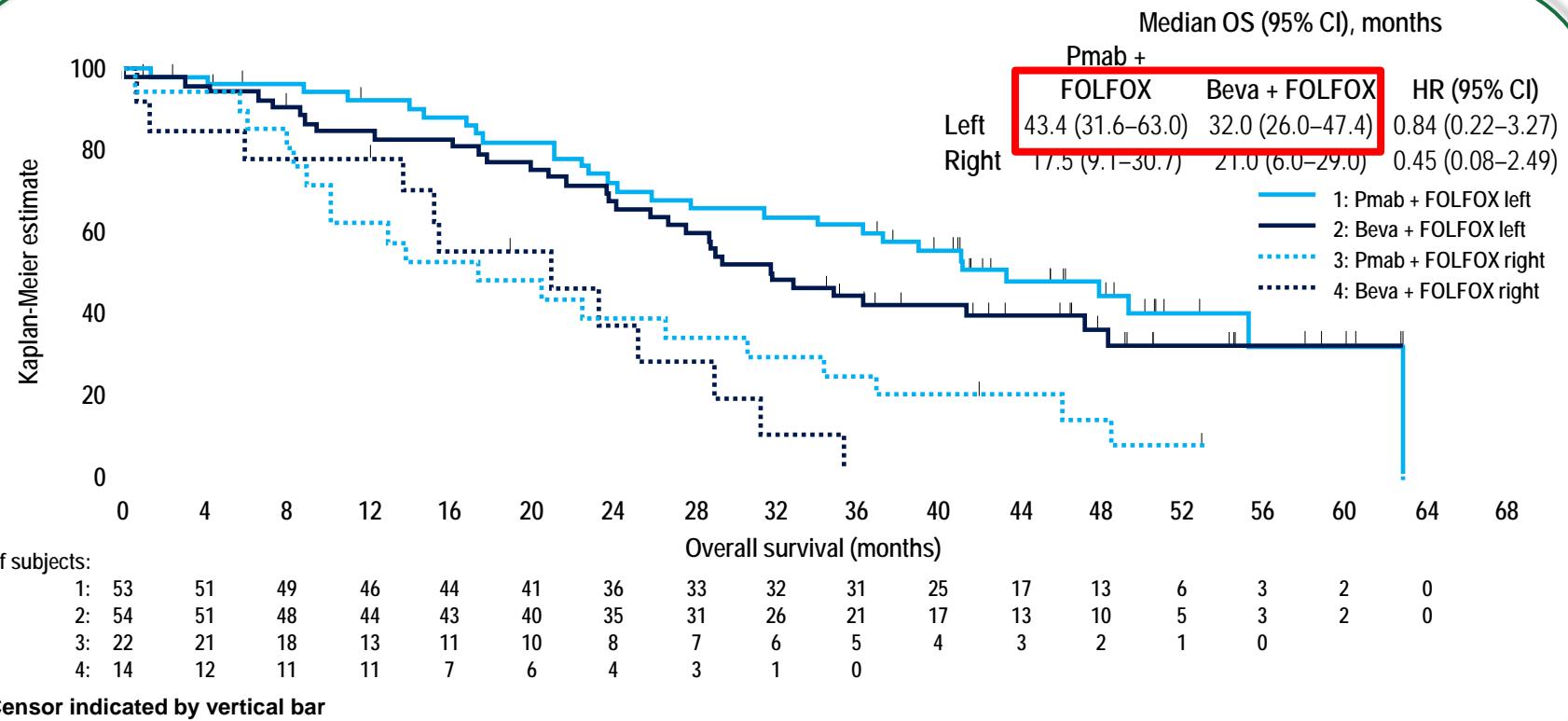


p for interaction 0.009

Right versus Left: CALGB80405 study - OS



Right versus Left: PEAK – OS



Beva, bevacizumab; HR, hazard ratio; OS, overall survival; Pmab, panitumumab

Right versus Left: meta-analysis of H2H trials - OS



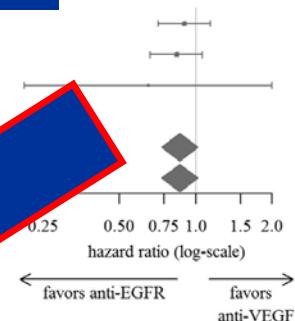
Right versus Left: meta-analysis of H2H trials - PFS

C

Left-sided

study	n	PFS HR	95% CI	P-value
FIRE-3	306	48.9	0.9 (0.71 , 1.14)	
CALGB/SWOG 80405	325	48.9	0.84 (0.66 , 1.06)	
PEAK	107	2.2	0.65 (0.21 , 2)	
Summary (FE)		0.86	(0.73 , 1.02)	0.084
Summary (RE)		0.86	(0.73 , 1.02)	

Heterogeneity: $I^2 = 0\%$, 95% CI = (0% , 97.2%)
 P-value = 0.813 (χ^2 test)

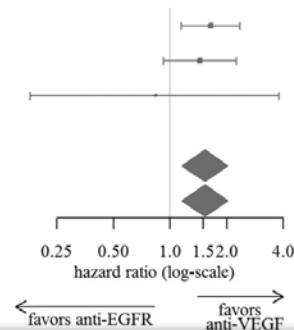


D

Right-sided

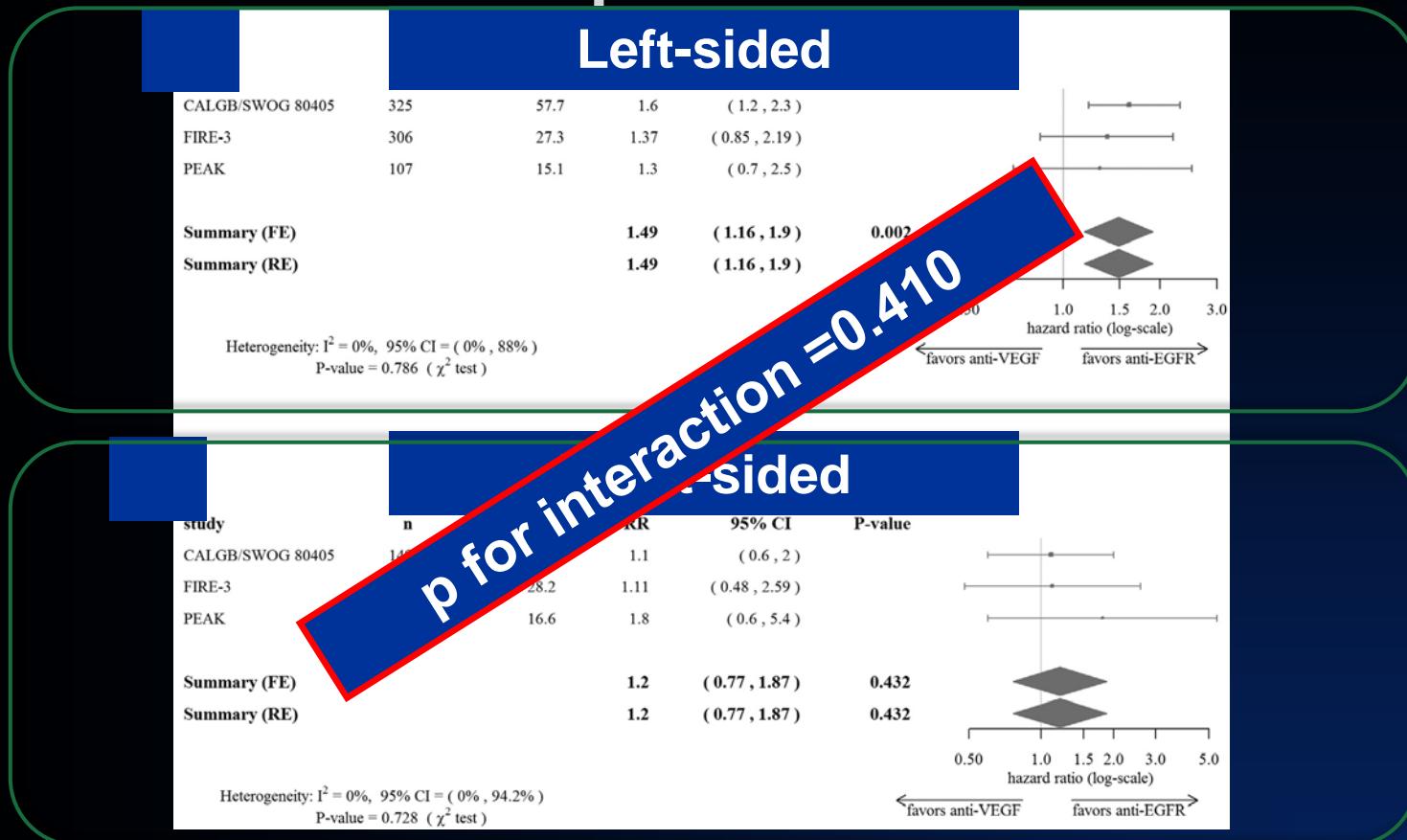
study	n	PFS HR	95% CI	P-value
CALGB/SWOG 80405	325	1.64	(1.15 , 2.36)	
FIRE-3	37.7	1.44	(0.92 , 2.26)	
PEAK	3.3	0.84	(0.18 , 3.79)	
Summary (FE)		1.53	(1.16 , 2.01)	0.003
Summary (RE)		1.53	(1.16 , 2.01)	0.003

Heterogeneity: $I^2 = 0\%$, 95% CI = (0% , 98.3%)
 P-value = 0.668 (χ^2 test)

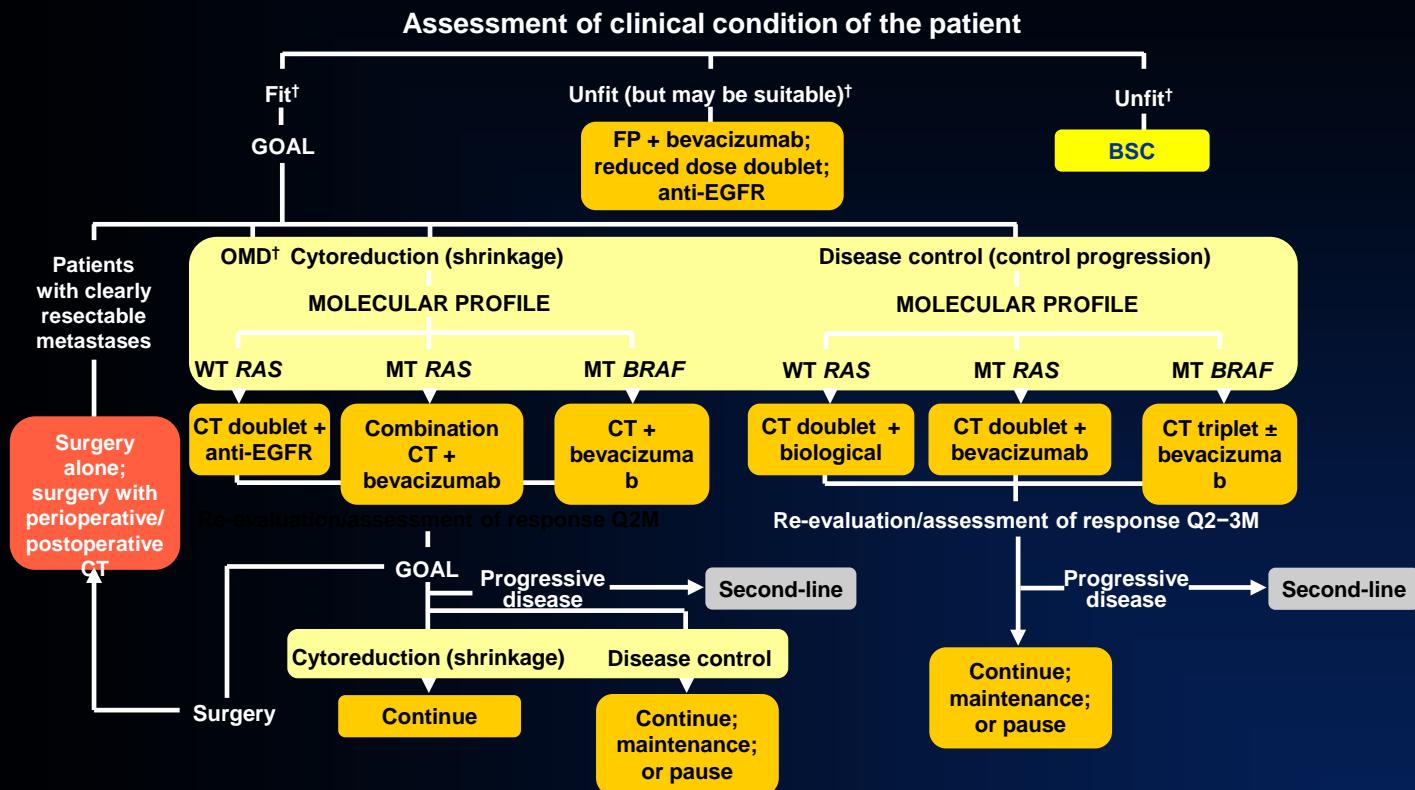


p for interaction <0.001

Right versus Left: meta-analysis of H2H trials - response rate



ESMO recommendations: 2016



[†]Patients assessed as 'fit' or 'unfit' according to medical condition, not due to malignant disease; [‡]Separate algorithm. OMD, oligometastatic disease.

ESMO Recommendations: 2016

Goal / Condition	Molecular	Preferred 1st line regimen
Disease stabilization	all WT	Doublet plus EGFR or Doublet plus beva
	RAS mut	Doublet plus beva
	BRAF mut	FOLFOXIRI +/- beva
Cytoreduction	all WT	Doublet plus EGFR (FOLFOXIRI plus beva)
	RAS mut	Doublet or triplet + beva
	BRAF mut	FOLFOXIRI + beva
“Frail”, or chosen sequential treatment	no BRAF !	Cape or FU + beva

After Right vs Left

Goal / Condition	Molecular	Preferred 1st line regimen
Disease stabilization	all WT	Left: Doublet/EGFR Right: Doublet (FOLFOXIRI)/beva
	RAS mut	(Doublet)/FOLFOXIRI/beva
	BRAF mut	FOLFOXIRI/(Doublet)/beva
Cytoreduction	all WT	Left: Doublet/EGFR Right: FOLFOXIRI/beva (Doublet/EGFR)
	RAS mut	FOLFOXIRI/beva
	BRAF mut	FOLFOXIRI/beva
“Frail”, or chosen sequential treatment	no BRAF !	Capecitabine/beva

Should we consider primary tumour location among other drivers of the first-line therapy?



Candidate predictors of resistance to anti-EGFRs

Different molecular alterations, supported by a strong and sound biologic rationale, have been suggested as predictors of primary resistance to anti-EGFRs

Candidate alterations:

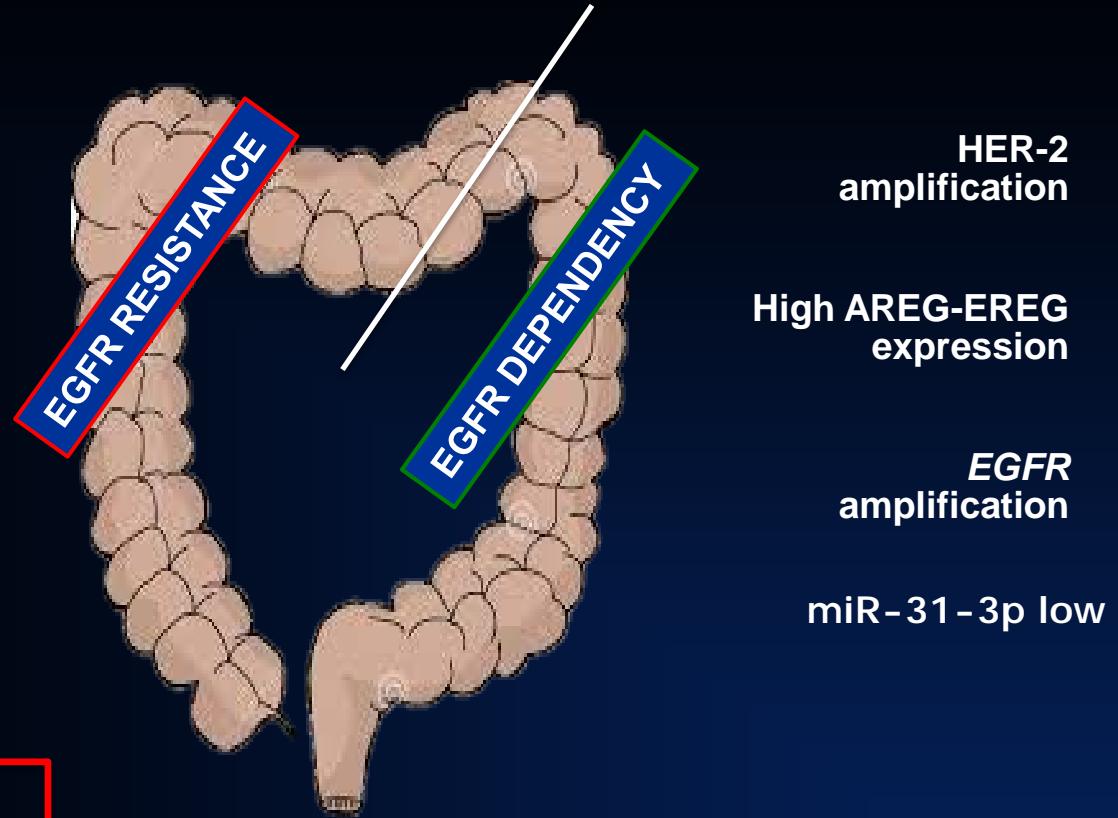
- *HER2* amplification or mutations
- *MET* amplification
- *ALK/ROS1/NTRKs* and *RET* fusions
- *PIK3CA/PTEN/Akt* and *MAPKs* pathways' activating mutations

BUT...

Up to today their potential impact has been demonstrated only in pre-clinical experiences and retrospective series

Right versus Left: Molecular make-up

BRAF V600E
mutation
BRAF-like signature
RAS mutations
PIK3CA mutations
dMMR
CIMP-high
Low AREG-EREG expression
CMS1(Immune)
miR-31-3p high
EGFR promoter methylation
ALK/ROS1/NTRK rearrangements



Conclusions

1. Few new classes of therapeutics; refinement of patient selection
2. Data relating to tumour site; improves patient selection
3. Reinforce the use of EGFR antibody therapy in patients with mCRC and left-sided RAS *wt* tumours
4. Patients with right-sided RAS *wt* tumours might be better treated with triplet chemotherapy alone or triplet chemotherapy plus bevacizumab except maybe doublet chemotherapy plus anti-EGFR if the goal is tumour size reduction as the ORRs were higher (but not PFS and OS)
5. Knowledge of site of primary tumor does not replace molecular characterisation