

Update zur Therapie des metastasierten kolorektalen Karzinoms

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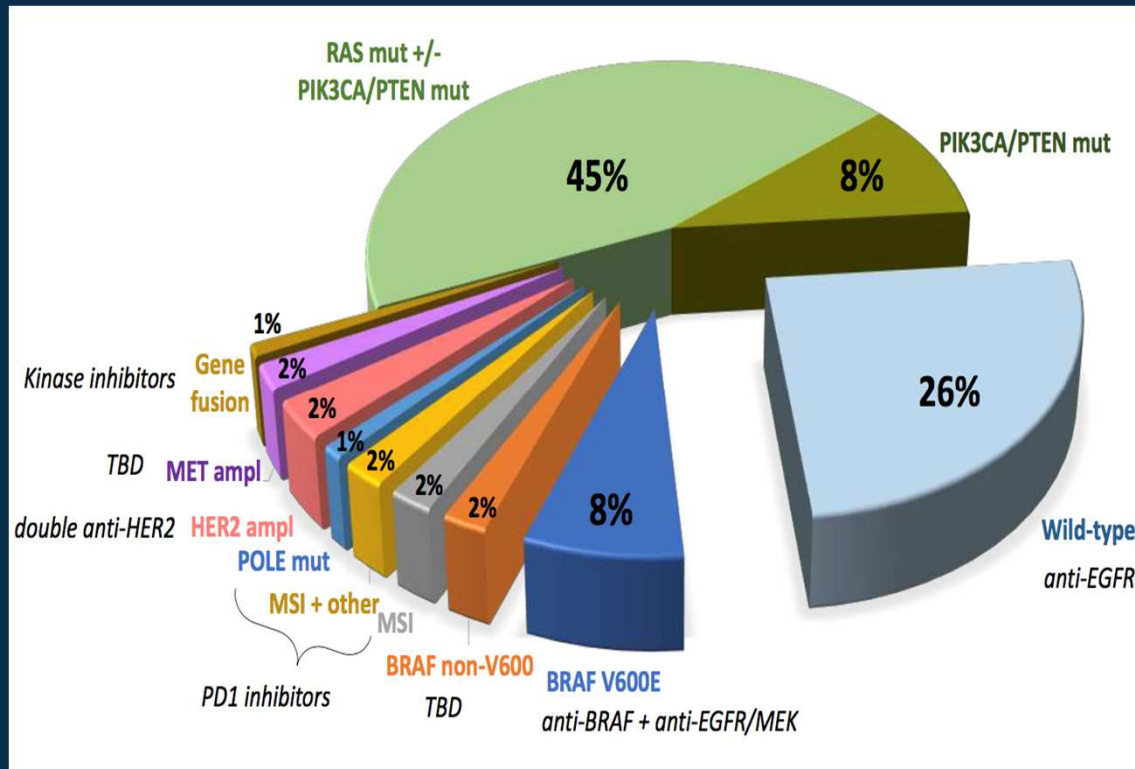
CCC MÜNCHEN
COMPREHENSIVE
CANCER CENTER

Molecular Subgroups of Colorectal Cancer

Genomic classification	• RAS mutation • BRAF V600 and non V600 mutation • HER2 amplification • Gene fusions • MSI and POLE mutations
Transcriptomic classification	• Consensus Molecular Subtypes (CMS)
Integrative classification (Sidedness of primary tumor)	• Right colon versus left colon

Extended molecular diagnostics: NGS-based panel sequencing ➔ molecular tumor board

Genomic Classification of mCRC: **New Avenues**



Specific treatments
for rare subtypes

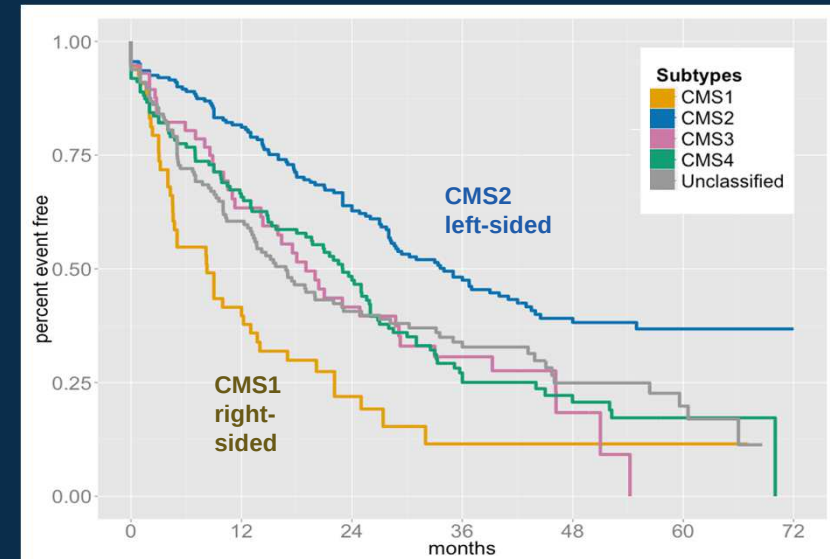
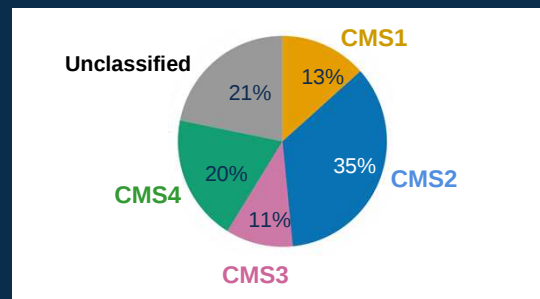
- MSI
- BRAF
- HER2 +
- TRK fusions

Rodrigo Dienstman, ESMO

Consensus Molecular Subtypes (CMS)

Transcriptomic Classification

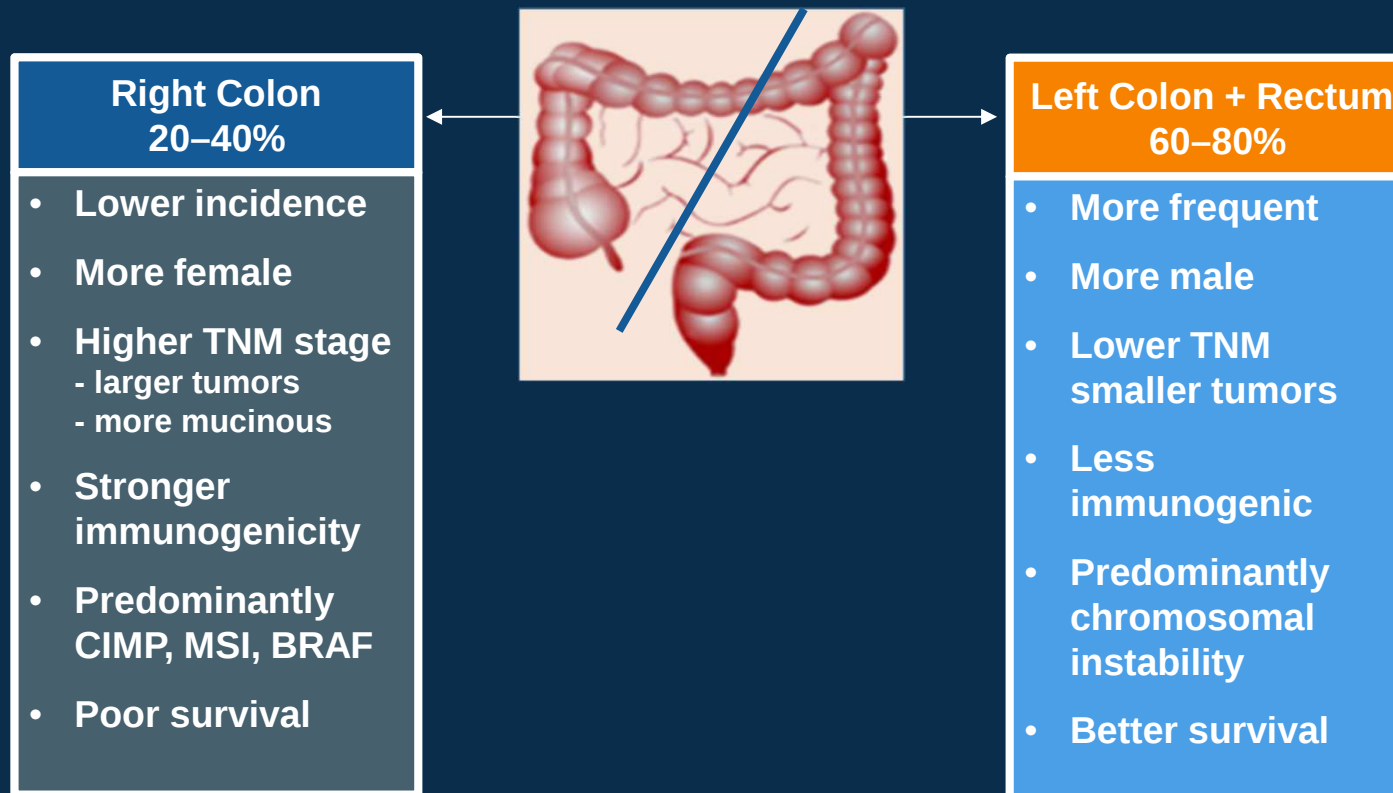
CMS1 MSI immune	CMS2 Canonical	CMS3 Metabolic	CMS4 Mesenchymal
MSI, CIMP high, hypermutation	SCNA high	Mixed MSI, SCNA low, CIMP low	SCNA high
BRAF mutations		KRAS mutations	
Immune infiltration and activation	WNT and MYC activation	Metabolic deregulation	Stroma infiltration, TNF β activation, angiogenesis
Worse survival			Worse RFS and OS



CMS has a strong prognostic relevance

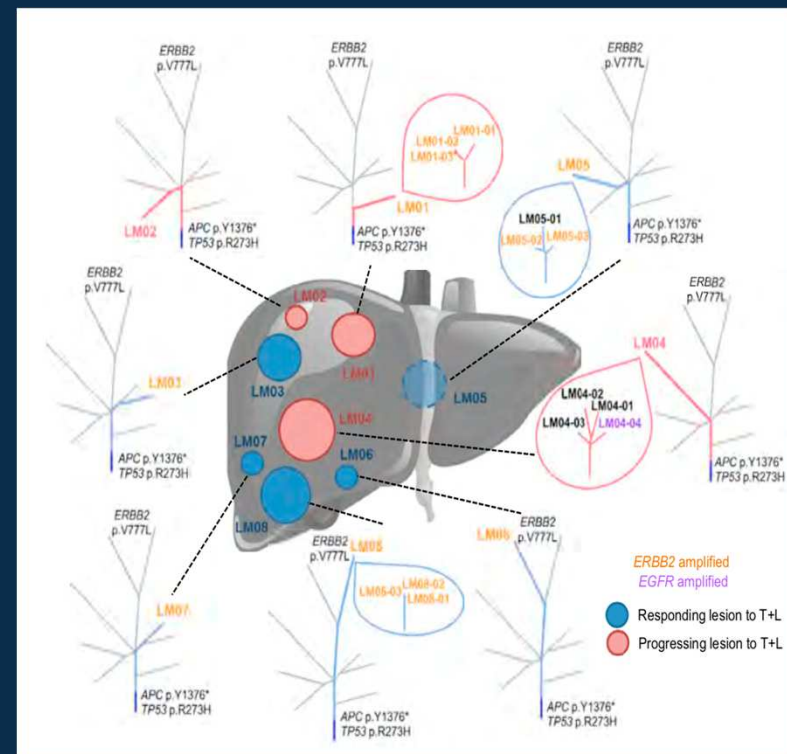
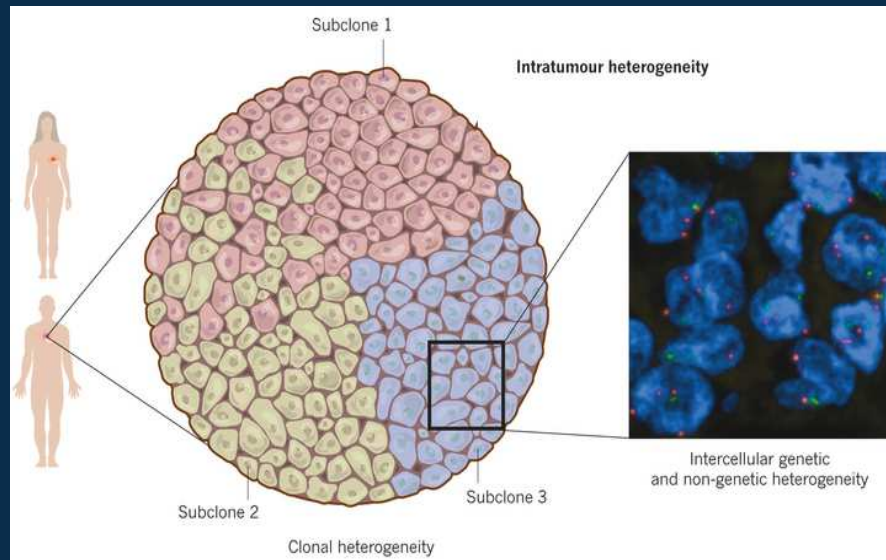
Guinney J et al. Nat Med 2015

Integrative Classification: Right- vs. Left-Sided Tumors



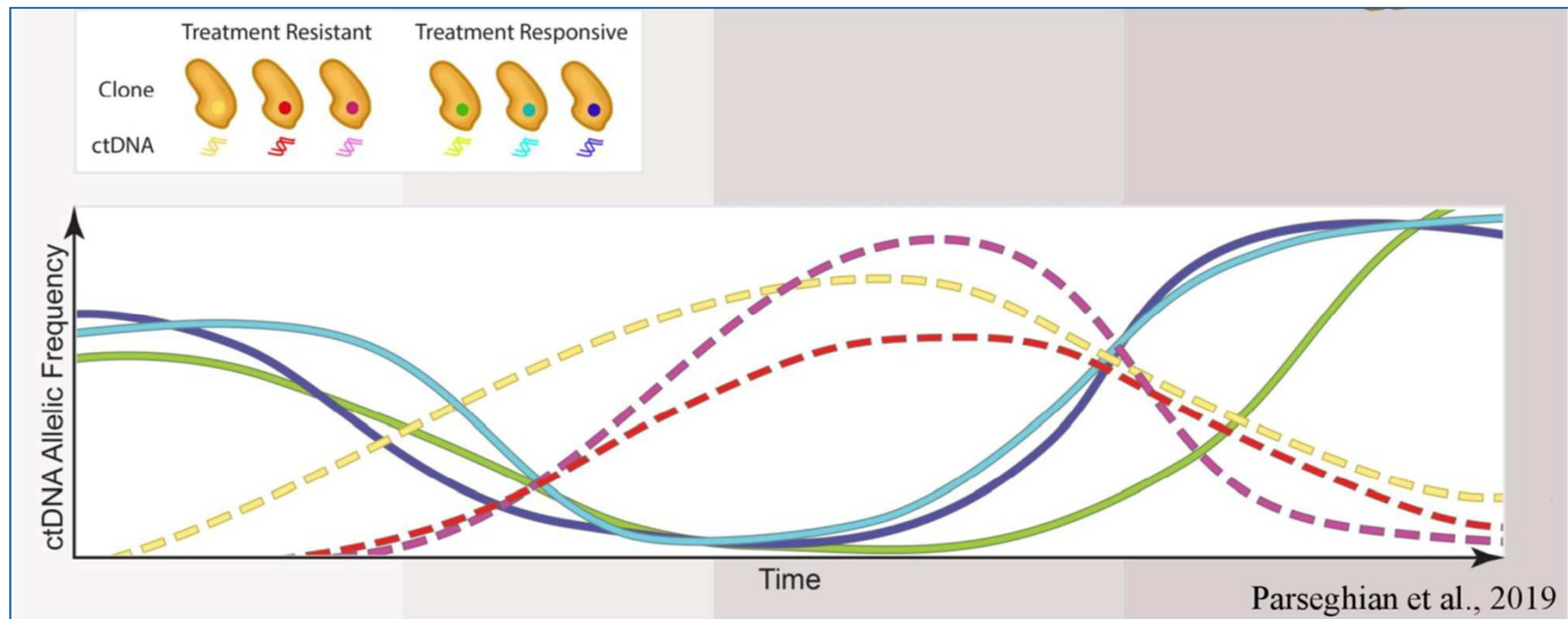
Heterogeneous clonal evolution

dependent on localisation of metastasis

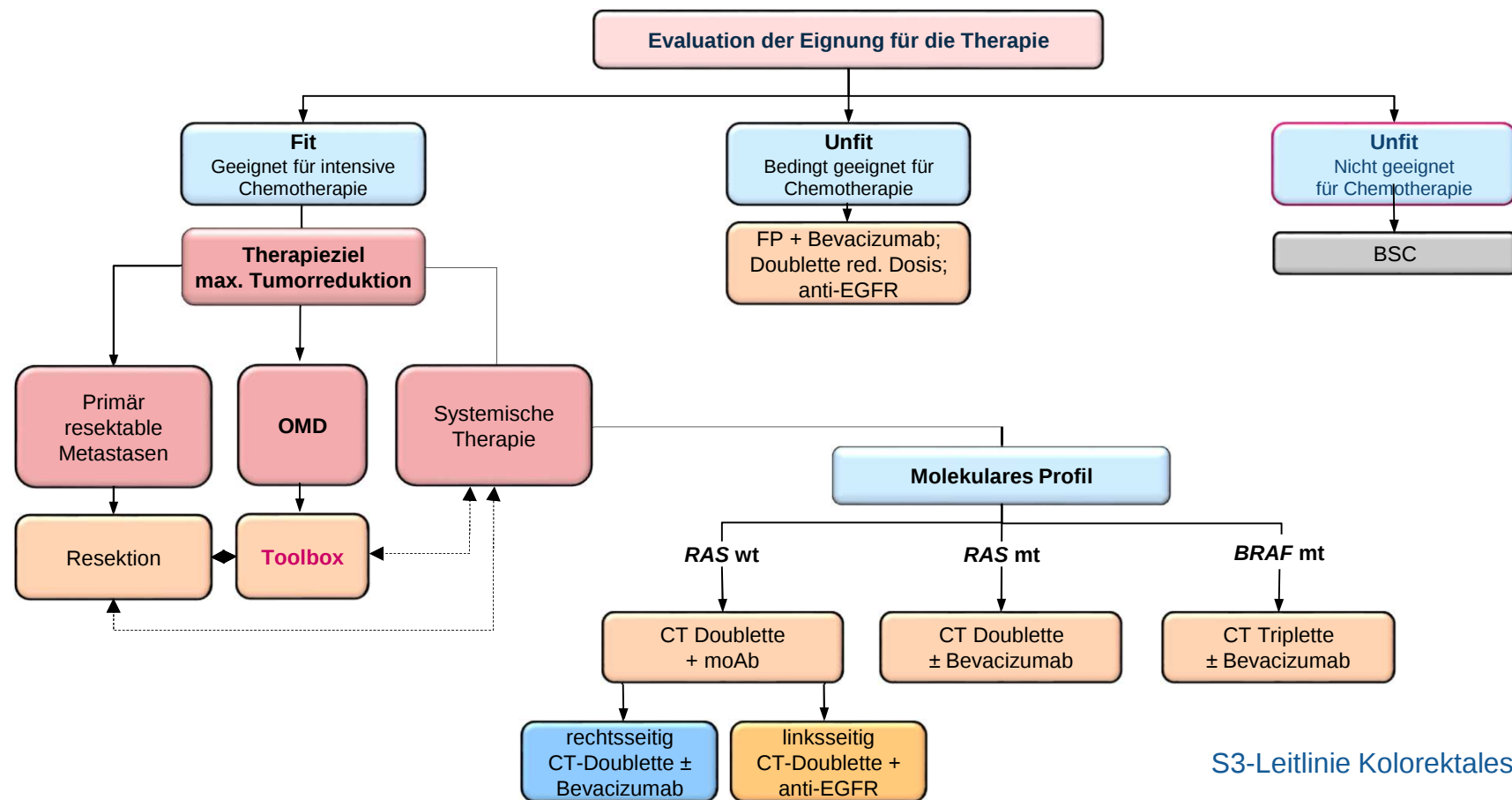


Siravegna G, et al. *Cancer Cell*. 2018;34(1):148-162.

Dynamische Evolution und Verschwinden von Mutationen in Abhängigkeit vom Selektionsdruck der Therapie



S3-LL: Therapiealgorithmus Erstlinientherapie des mKRK



1st-Line Behandlung bei RAS Wildtyp mCRC

rechtsseitig

schlechtere Prognose

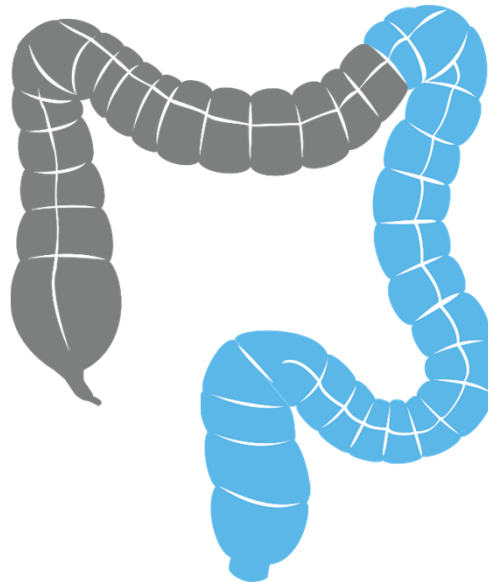
Ziel = OS

**Chemo ±
Bevacizumab**

FOLFOXIRI + Bev

Ziel = ORR / Conversion

Chemo + anti-EGFR



linksseitig

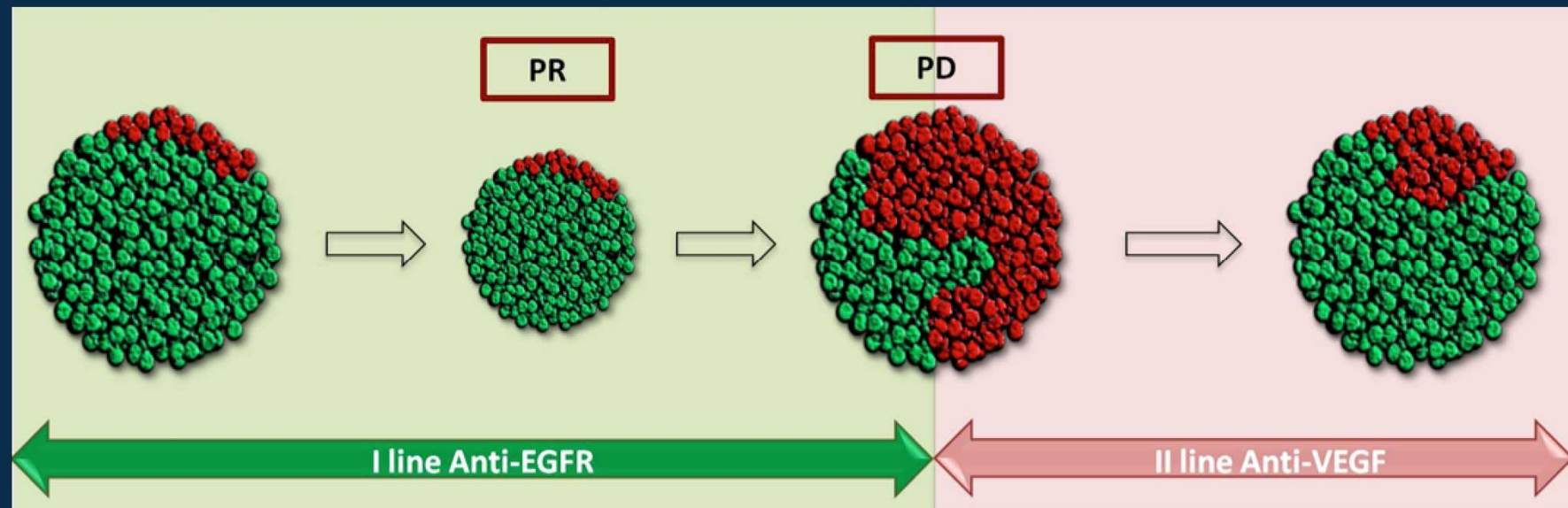
bessere Prognose

Ziel = OS / ORR

**Chemo +
anti-EGFR**

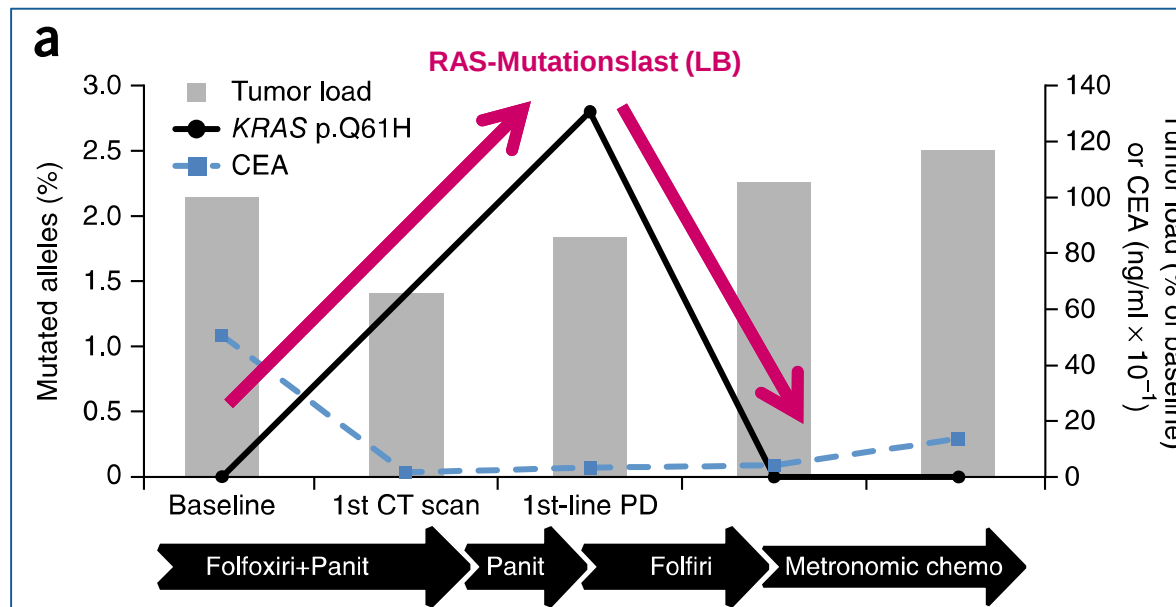
Temporary clonal evolution

due to selective pressure of therapy



Reversion of resistance after stop of anti-EGFR-therapy

Entwicklung von RAS Mutationen unter einer anti-EGFR Therapie



**Abfall der
RAS-Mutationslast
nach
Absetzen der
anti-EGFR Therapie**

Mögliche Relevanz der Liquid Biopsie in der
Therapieführung

Parseghia et al, Ann Oncol;30(2): 243-249, 2019
Siravegna et al, Nature Medicine 21, 795-801, 2015

Re-challenge

strategy after anti-EGFR pre-treatment

Anti-EGFR
induction

Doublet + anti-EGFR agent

Window
therapy

Doublet + bevacizumab

Re-challenge

Chemo + anti-EGFR agent

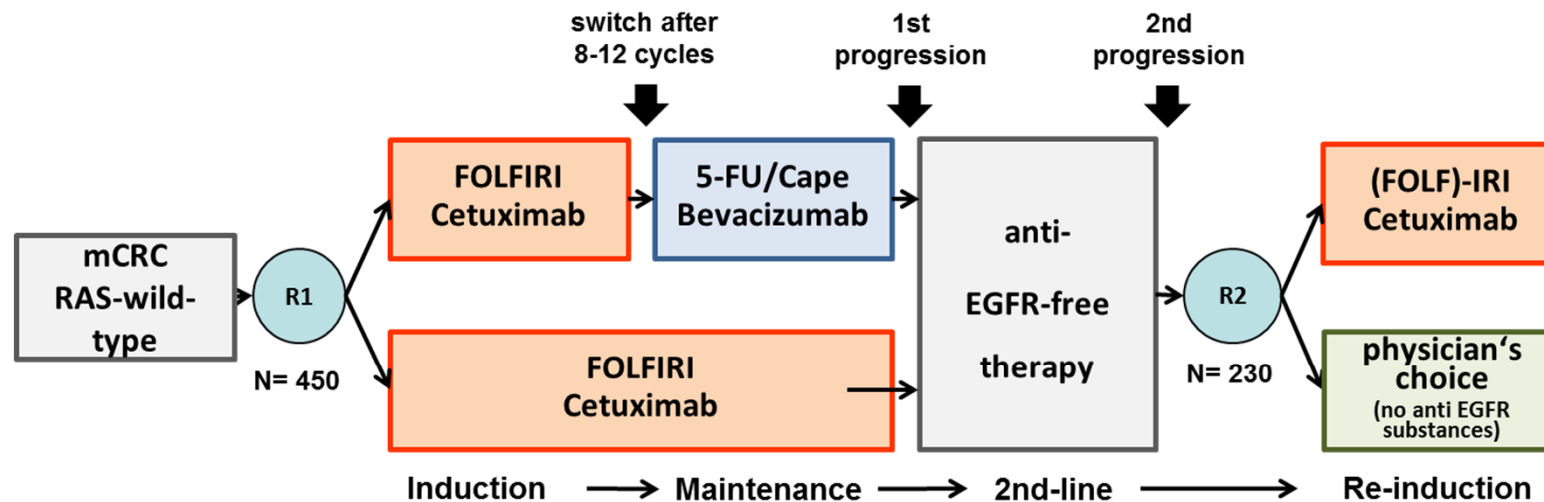
- if CR or PR in 1st line
- or SD $\geq$ 6 months in 1st line
- exclusion of RAS mutation by liquid or tumor biopsy

Questions:

- Optimal **duration** of window therapy
- Optimal **interval** from end of anti-EGFR

FIRE-4 Studie: anti-EGFR Re-challenge

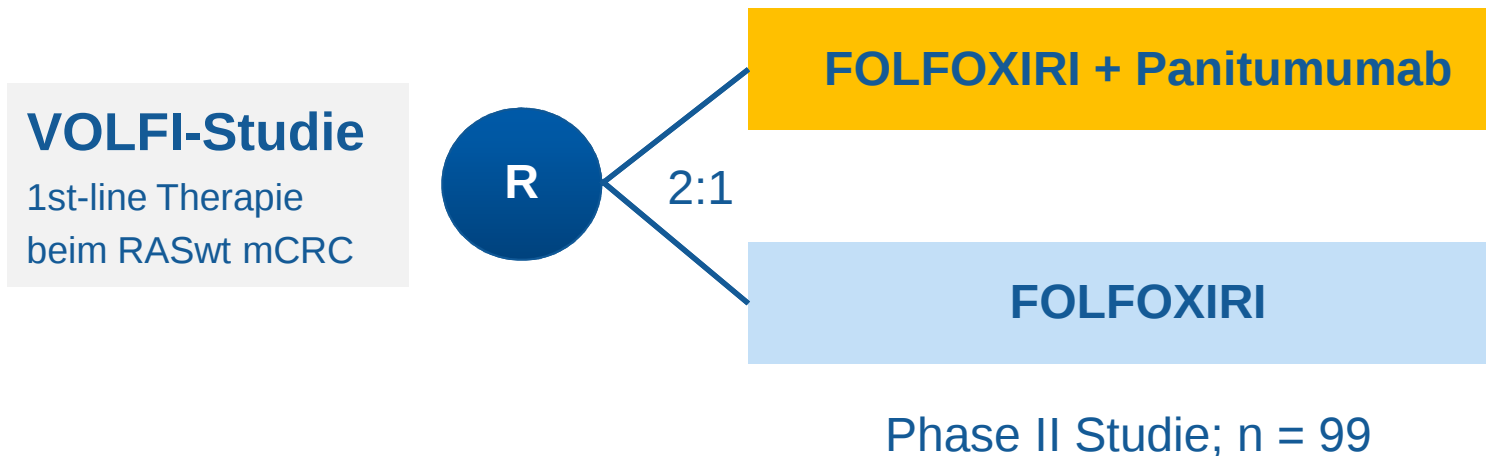
1st-line Therapie bei RAS wt mCRC



Primary Endpoint: OS3 after randomisation R2
Secondary Endpoint: PFS in 1st-line

Intensivierte Therapie

Triplette + anti-EGFR AK



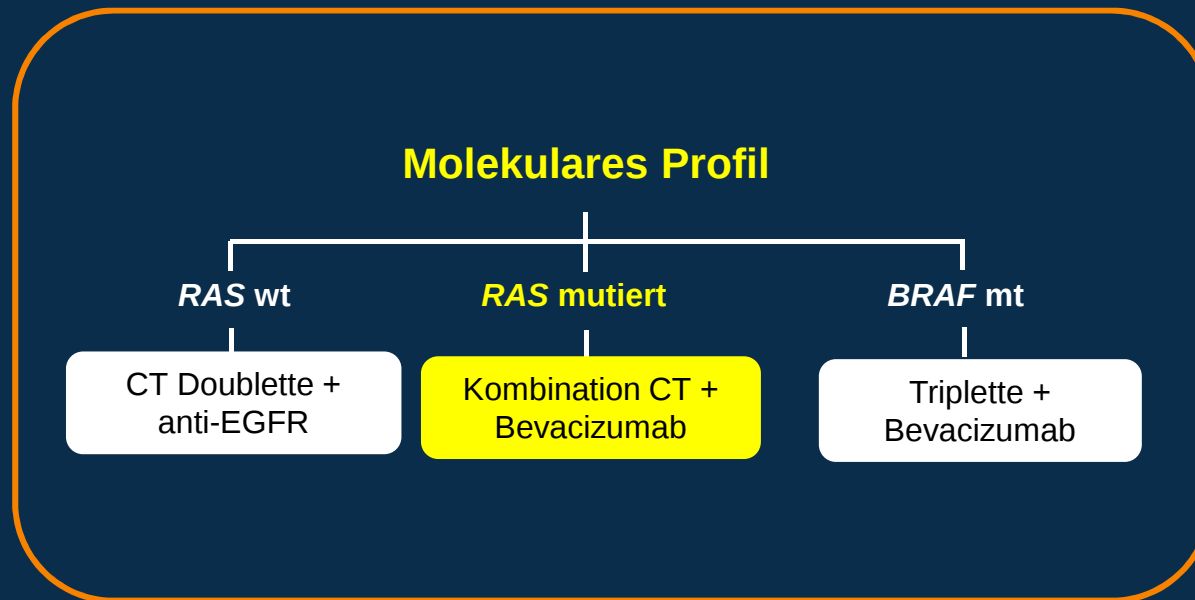
VOLFI: ORR in Abhängigkeit von Lokalisation / Biologie

	N	mFOLFOXIRI + panitumumab (%)	FOLFOXIRI (%)	Odds ratio	P
Full analysis set	96	87.3	60.6	4.47	0.004
Left	78	90.6	68.0	4.52	0.02
Right	18	70.0	37.5	3.89	0.34
RAS/BRAF wt	60	86.0	64.7	3.36	0.08
BRAF mut	16	85.7	22.2	21.00	0.04

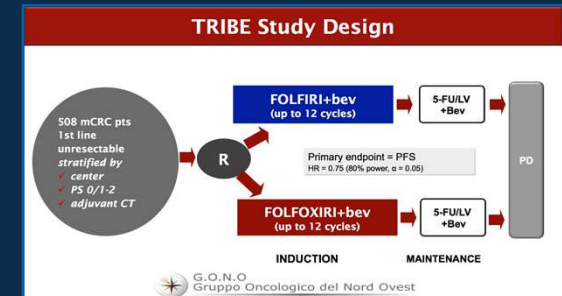
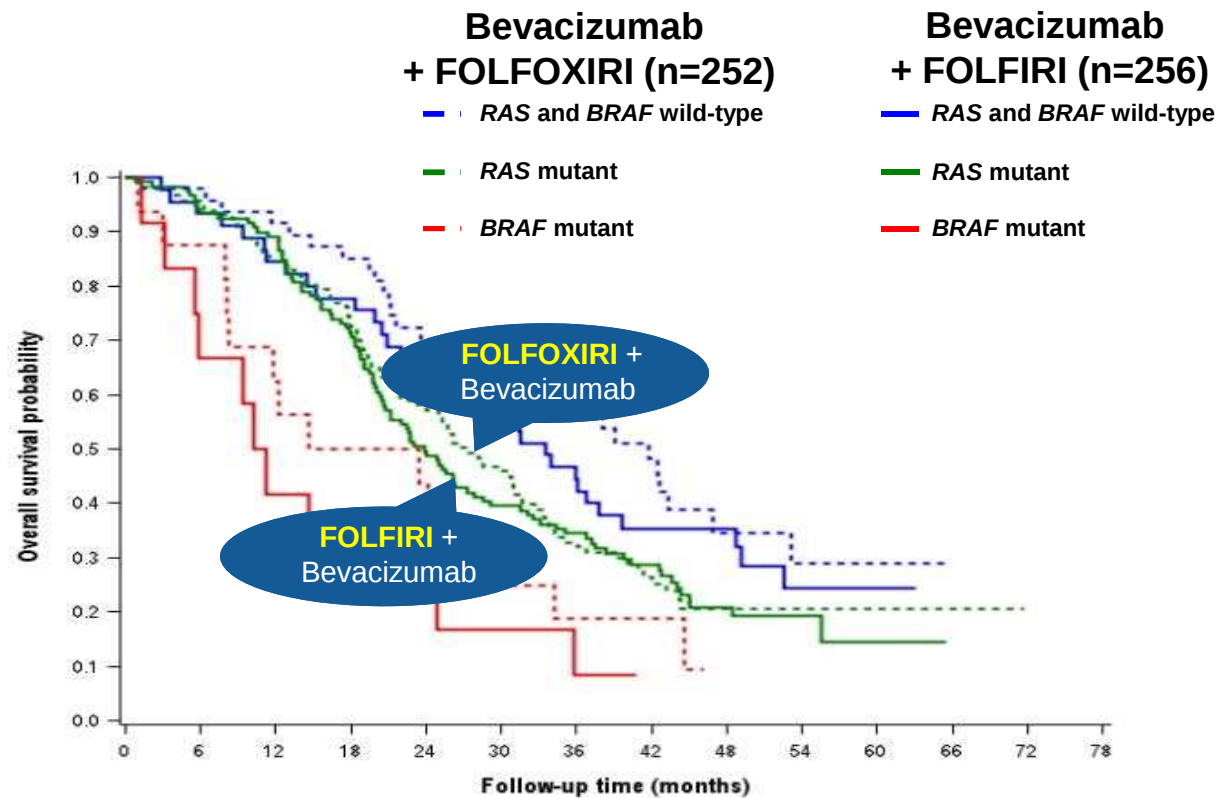
VOLFI: OS in Abhängigkeit von Lokalisation / Biologie

	N	mFOLFOXIRI + panitumumab (mo)	FOLFOXIRI (mo)
ITT	96	35.7	29.8
Left	78	39.9	35.3
Right	18	11.5	22.0
RAS/BRAF wt	60	43.5	35.3
BRAF mut	16	8.0	9.0

RAS mutiertes mCRC



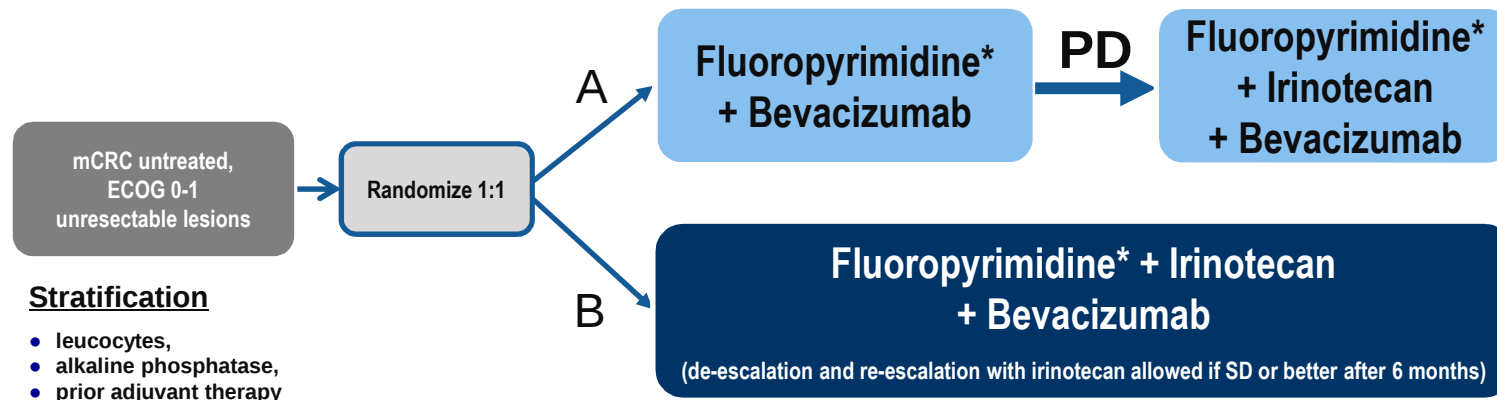
TRIBE Study: RAS mutant mCRC



No significant OS benefit from treatment intensification in RAS-mut mCRC

Cremolini C, et al. Lancet Oncol 2015;16:1306–1315

AIO: XELAVIRI



Arm A

Capecitabine plus bevacizumab q3w

capecitabine 2 x 1250 mg/m² day 1-14
bevacizumab 7.5 mg/kg day 1

5-FUFA plus bevacizumab q2w

folinic acid 400 mg/m² day 1
5-FU 400 mg/m² bolus day 1
5-FU 2400 mg/m² 46 h day 1-2
bevacizumab 5.0 mg/kg day 1

Arm B

CAPIRI plus bevacizumab q3w

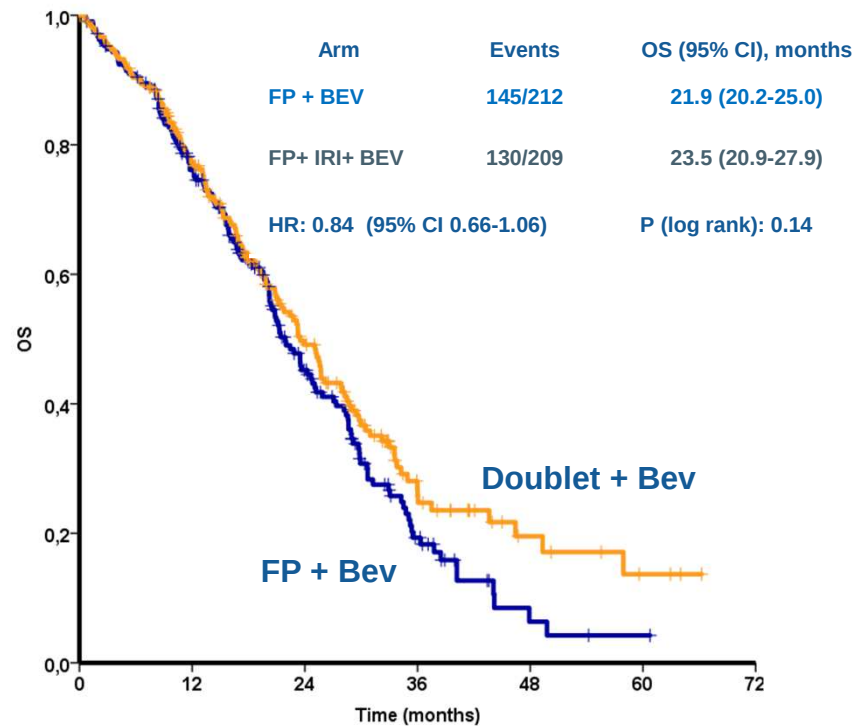
capecitabine 2 x 800 mg/m² day 1-14
irinotecan 200mg/m² day 1
bevacizumab 7.5 mg/kg day 1

FOLFIRI plus bevacizumab q2w

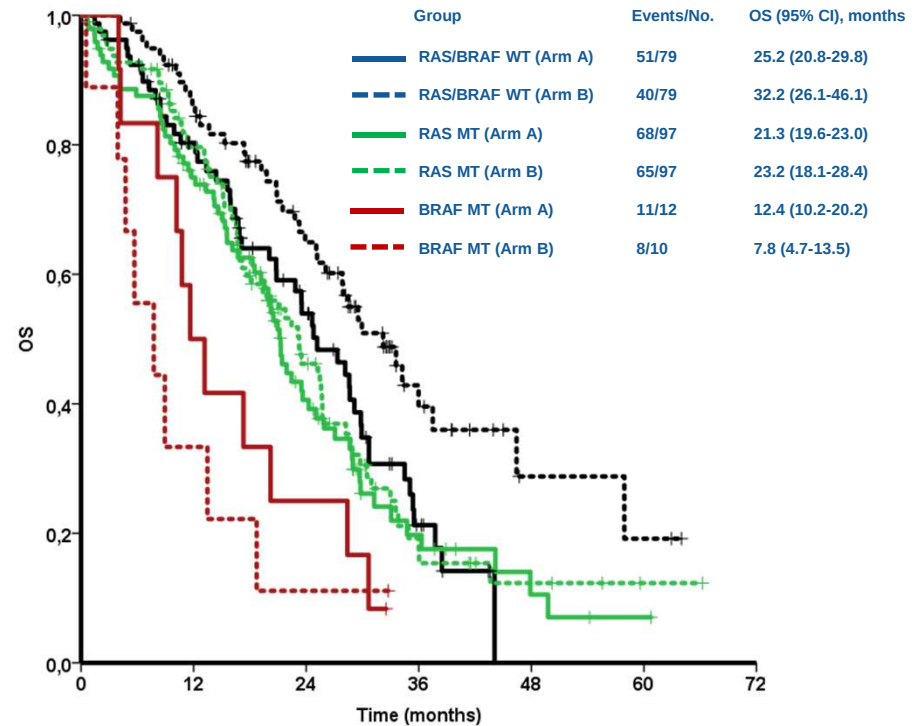
folinic acid 400 mg/m² day 1
5-FU 400 mg/m² bolus day 1
5-FU 2400 mg/m² 46 h day 1-2
irinotecan 180 mg/m² day1
bevacizumab: 5.0 mg/kg day 1

Modest D,Heinemann V, JCO accepted

XELAVIRI: Overall Survival

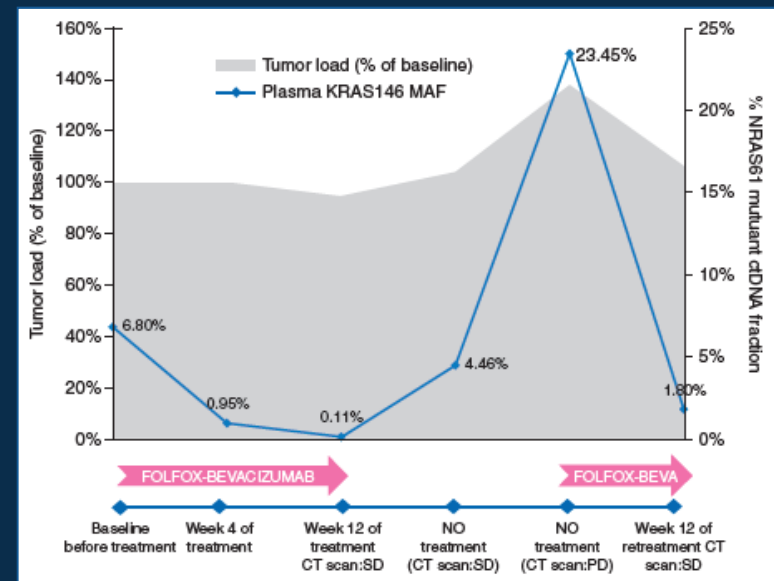
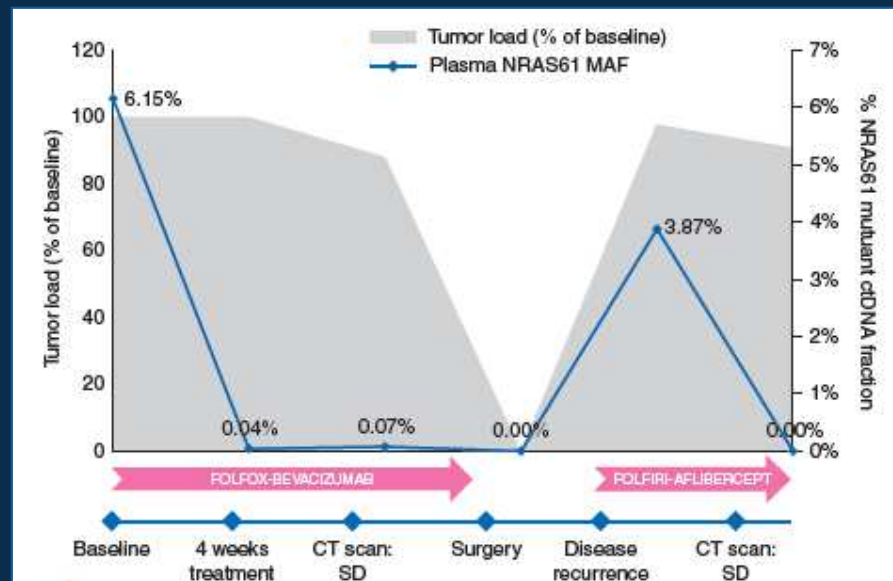


212	147	69	20	3	1
209	150	77	25	8	3



Doublet plus Bev not better than FP plus Bev in patients with RAS mutant tumors

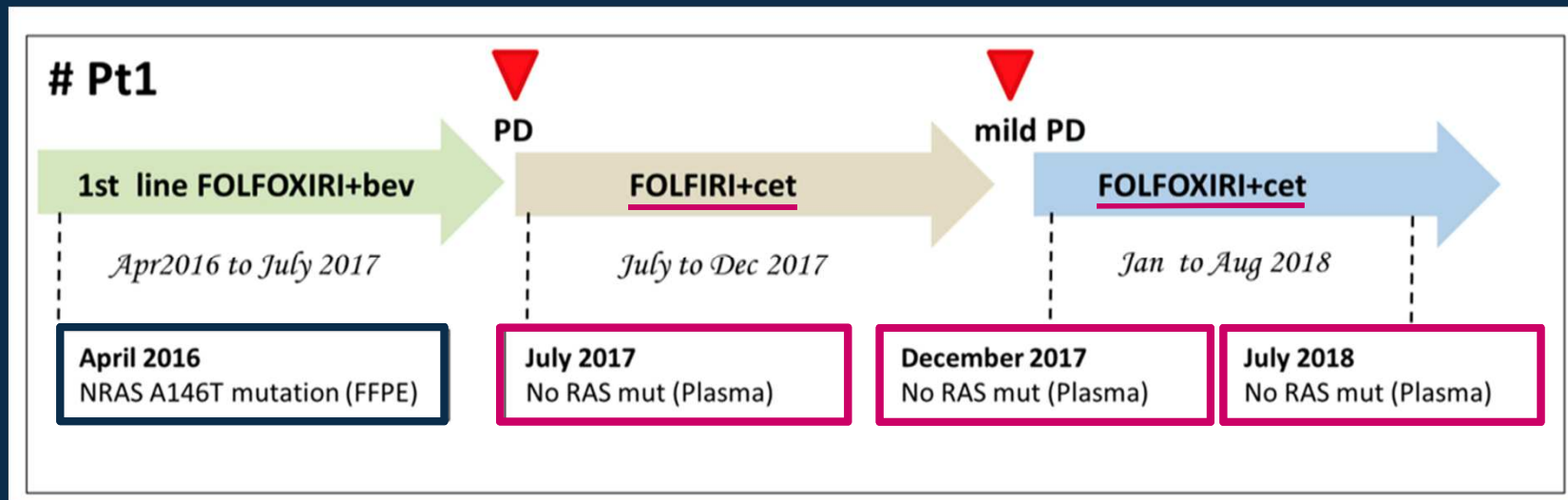
Elimination des RAS-mutierten Klons durch intensive 1st-line Chemotherapie



Longitudinale Analyse von Plasma RAS ctDNA

Vidal et al. Annals of Oncology

Elimination of RAS mutant clone by intensive therapy

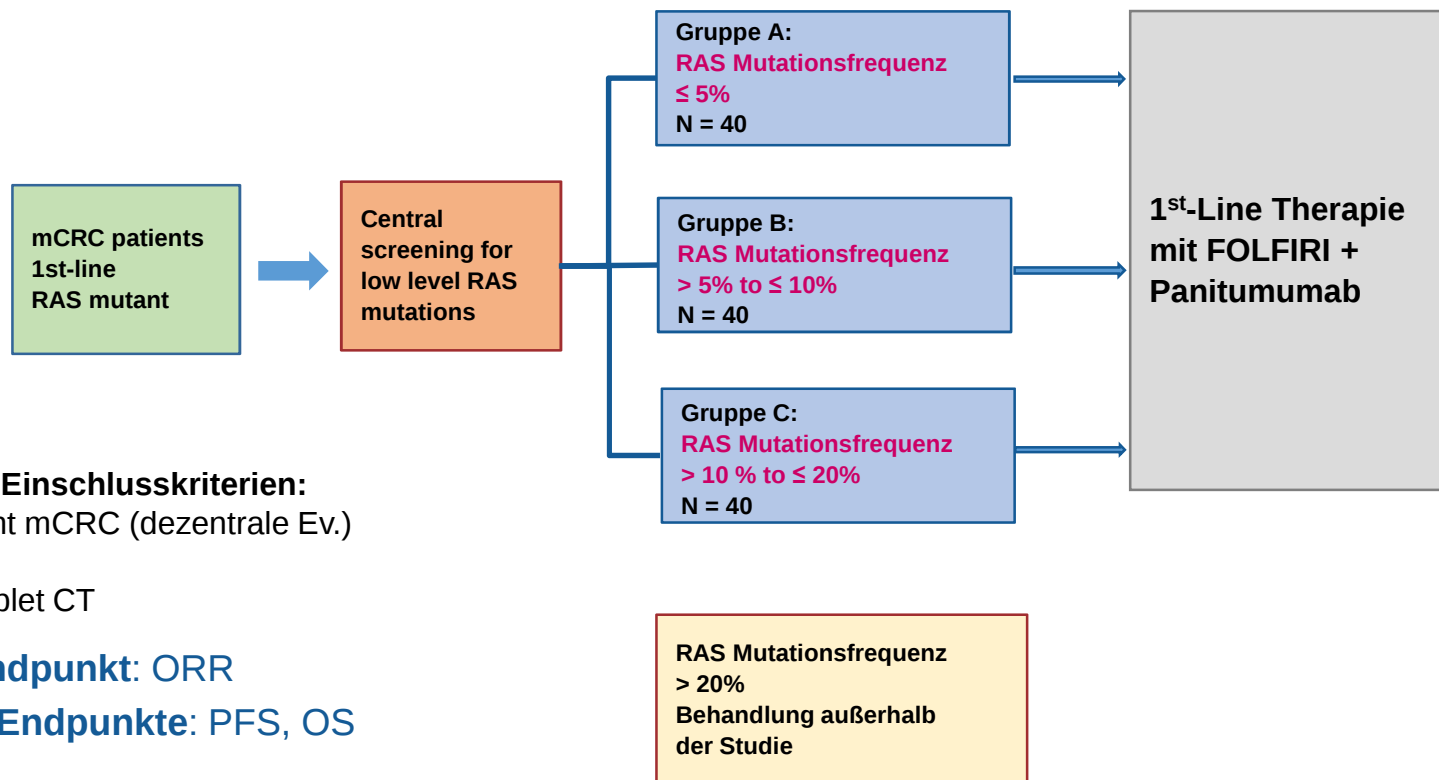


- Elimination of RAS-mutant clones by intensive chemotherapy in **5/11** patients (ctDNA)
- Benefit from anti-EGFR-based therapy

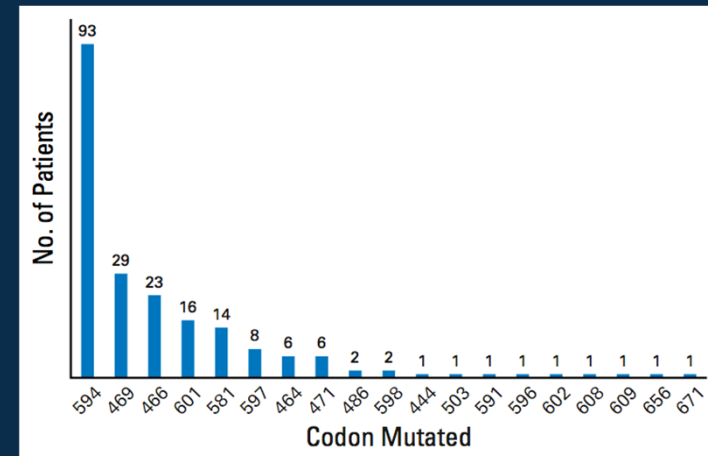
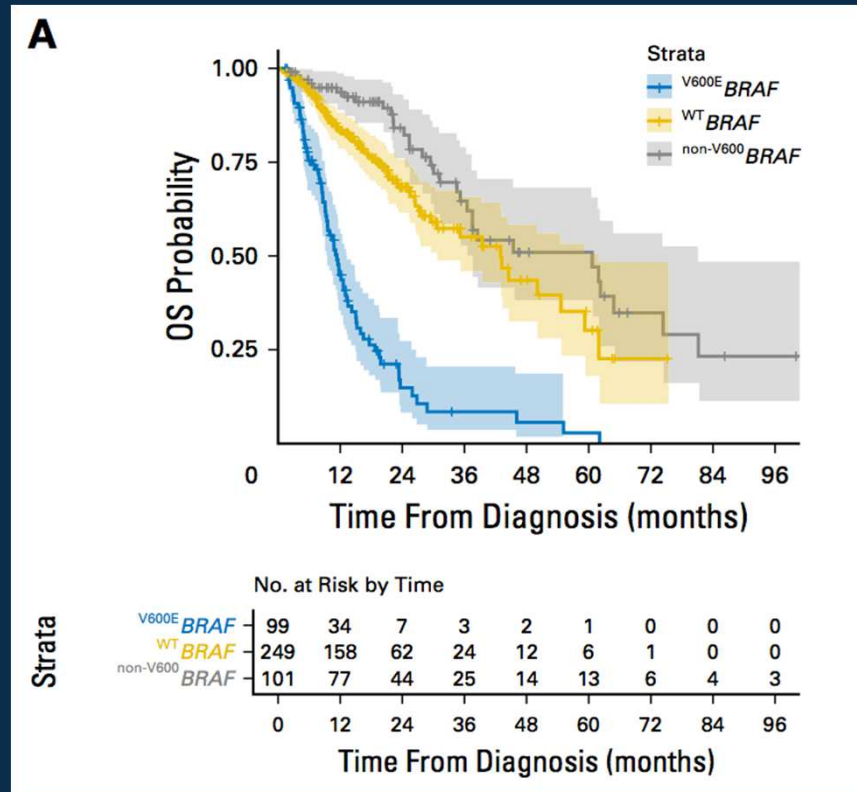
Raimondi C, et al. Cancers 2018

FIRE-5: Low-RAS Studie (AIO-TF-0118)

Erstlinientherapie des RAS-mutierten mKRC

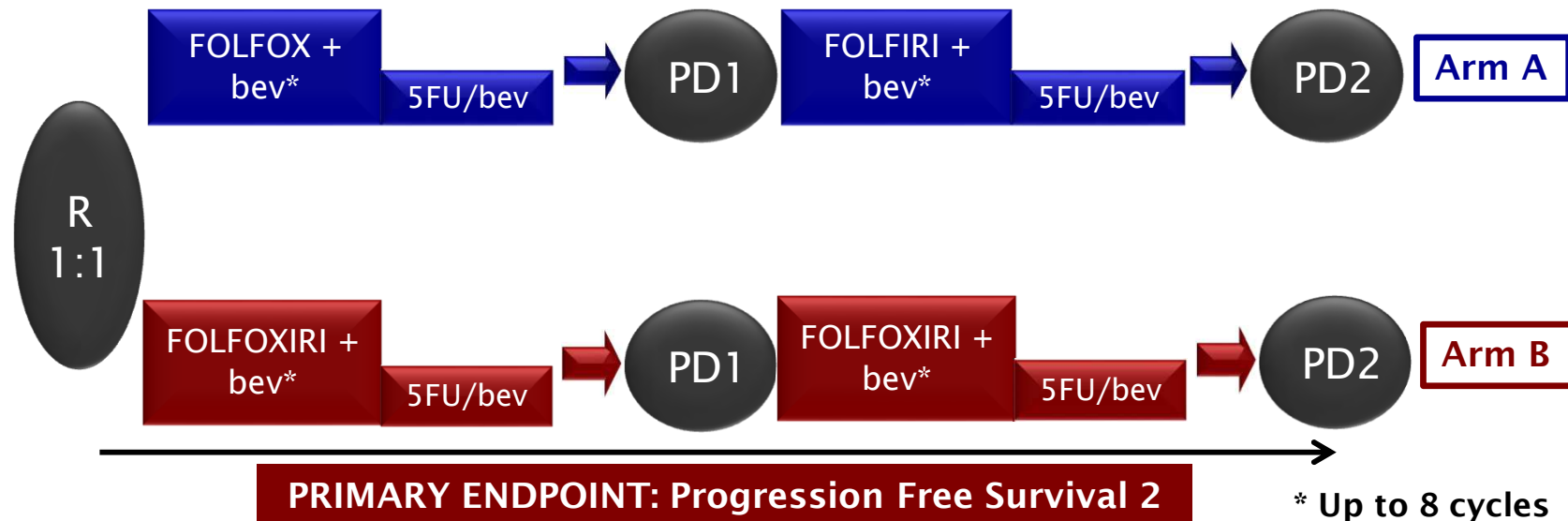


BRAF non-V600 in Metastatic Colorectal Cancer



- > male
- > younger patients
- > left-sided tumours
- < high grade tumours
- < peritoneal metastasis

TRIBE2: Study design and endpoints

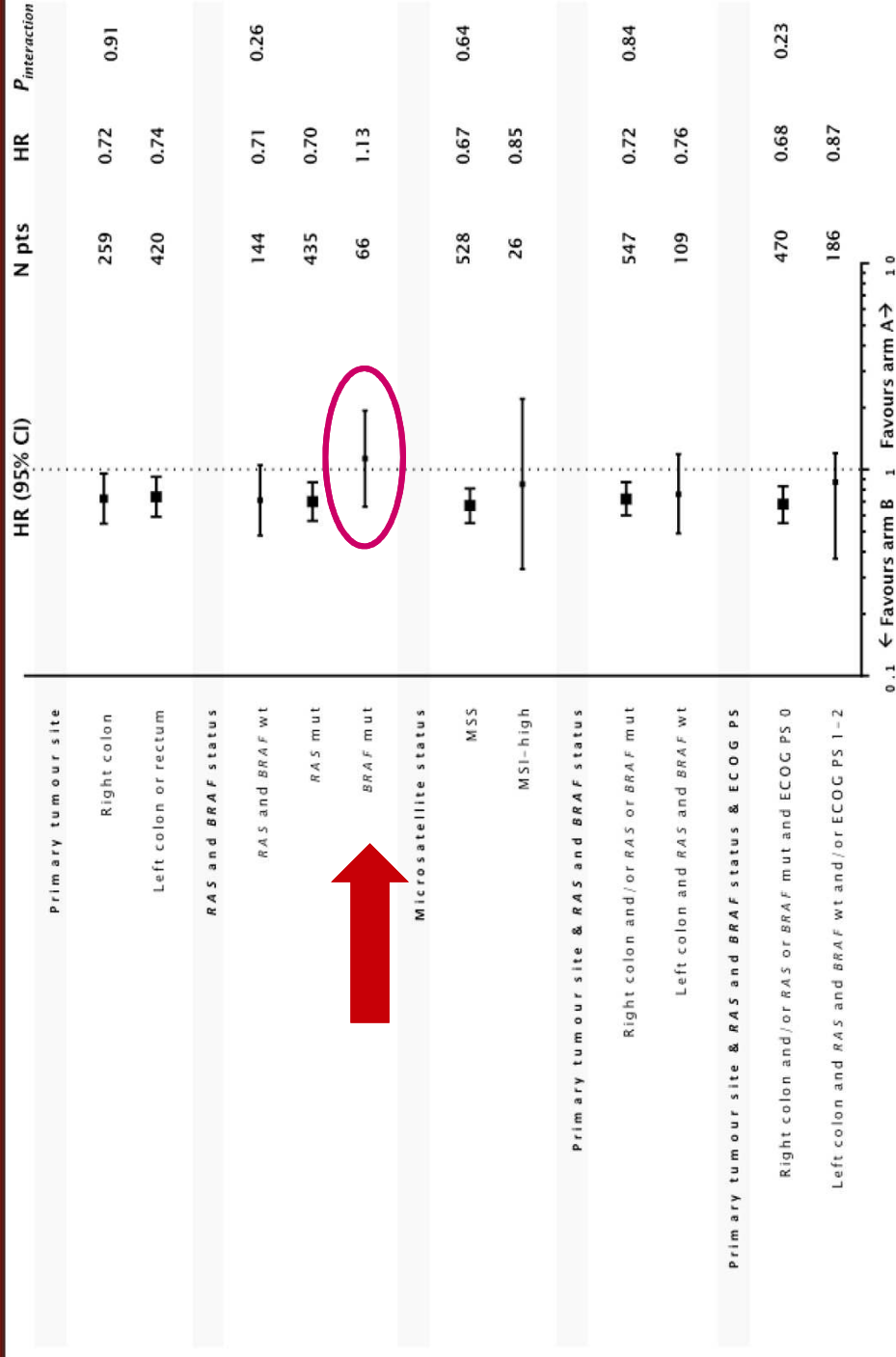


SECONDARY ENDPOINTS

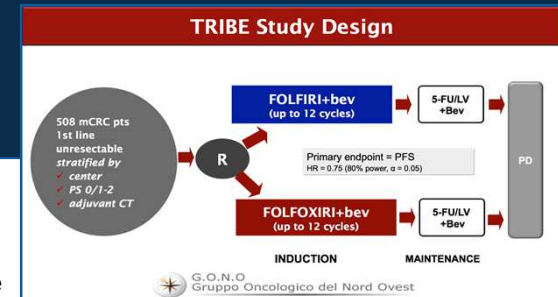
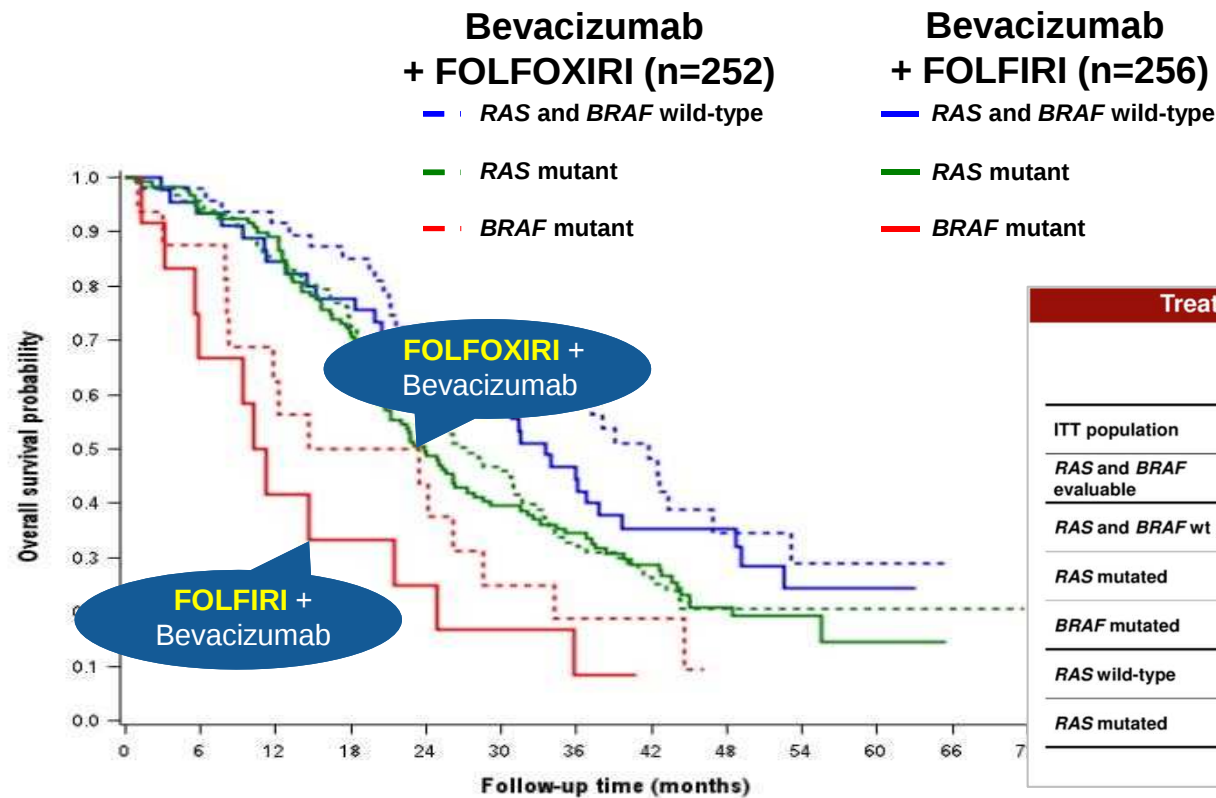
- 1st and 2nd Progression-Free Survival
- RECIST Response Rate in 1st and 2nd line
- Resection Rate
- Safety profile in 1st and 2nd line
- Overall Survival



Subgroup analyses – sidedness and molecular characteristics- PFS2



Behandlung bei BRAF-Mutation



Treatment effect in molecular subgroups					
	N	FOLFIRI + bev Median OS	FOLFOXIRI + bev Median OS	HR [95% CI]	p
ITT population	508	25.8	29.8	0.80 [0.65-0.98]	0.030
RAS and BRAF evaluable	357	24.9	28.6	0.84 [0.66-1.07]	0.159
RAS and BRAF wt	93	33.5	41.7	0.77 [0.46-1.27]	0.522*
RAS mutated	236	23.9	27.3	0.88 [0.65-1.18]	
BRAF mutated	28	10.7	19.0	0.54 [0.24-1.20]	
RAS wild-type	121	26.8	37.1	0.78 [0.51-1.20]	
RAS mutated	236	23.9	27.3	0.88 [0.65-1.18]	0.658*

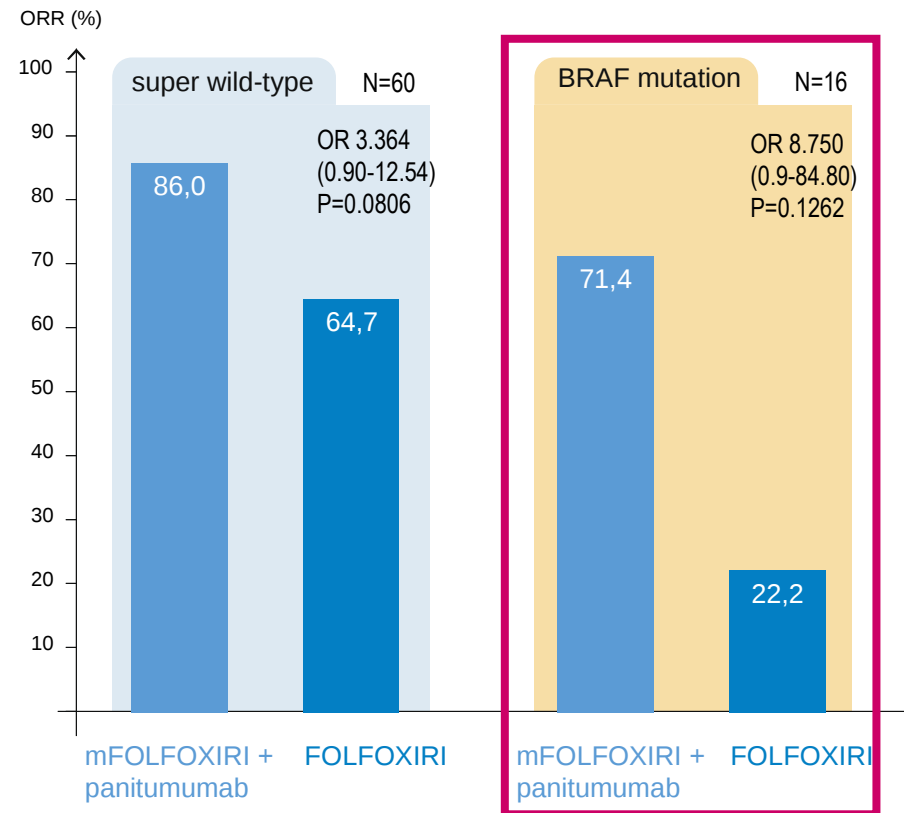
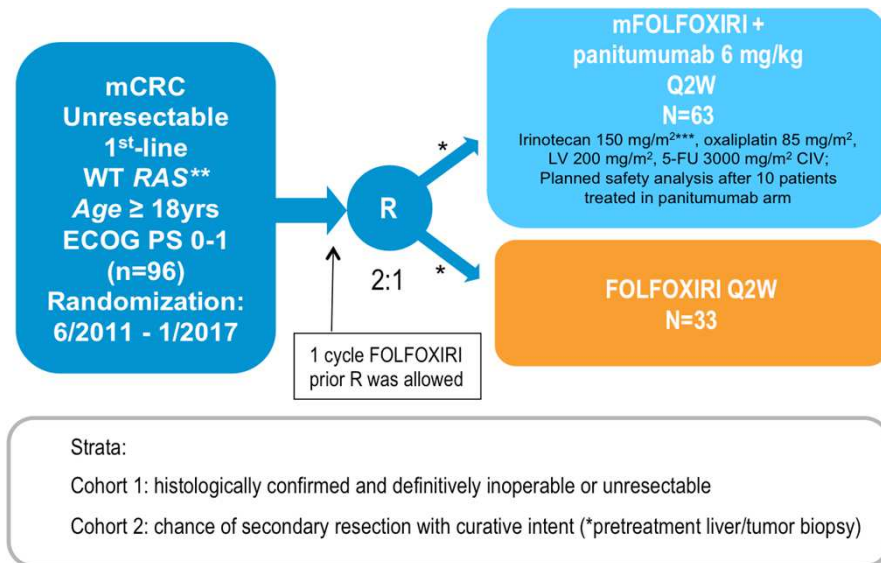
* p for interaction

Cremolini C, et al. Lancet Oncol 2015;16:1306–1315

EVALUATION OF RESPONSE SIDEDNESS + GENOTYPE

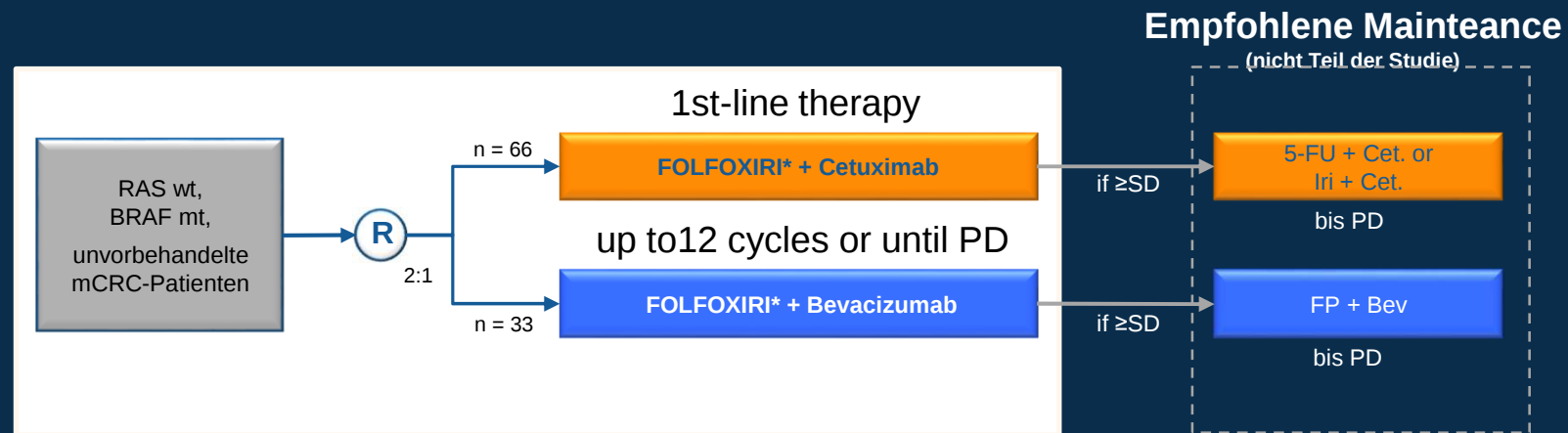
AIO

VOLFI Studie



FIRE 4.5:

Rekrutierende Studie beim BRAF mutierten mCRC



Phase II Studie

Primärer Endpunkt: ORR

Sekundäre Endpunkte: PFS, OS, ETS, DpR

Aktuell: **80/99** Patienten rekrutiert

FOLFOXIRI Dosierung:

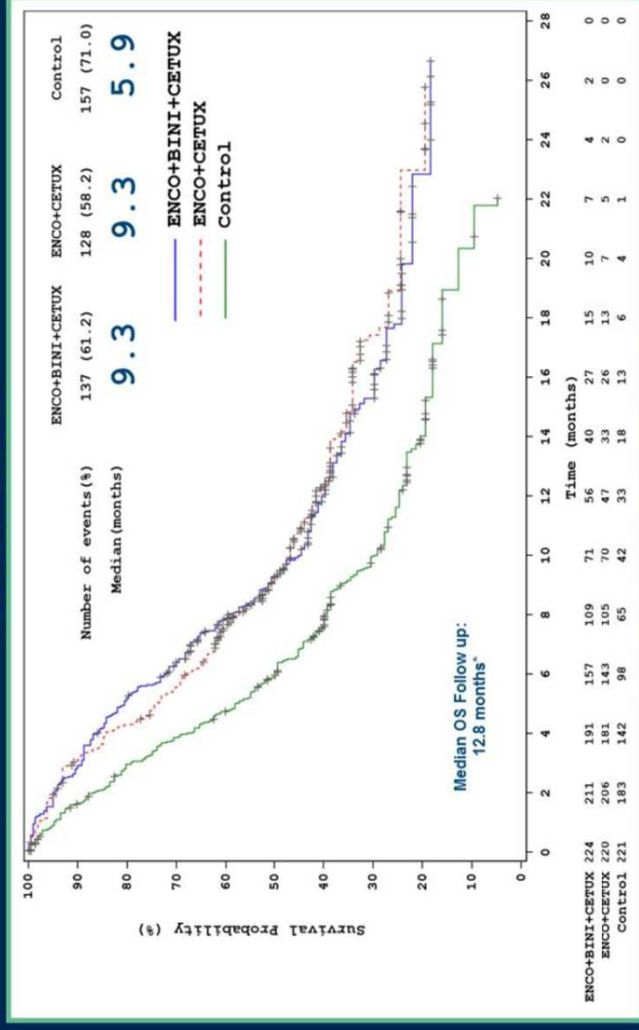
Irinotecan: 150mg/m²
FA (racemic): 400mg/m²
Oxaliplatin: 85 mg/m²
5-FU: 3000mg/m² iv für 48h

* de-escalation to FOLFIRI if toxicity is observed

BEACON CRC: Updated Analysis

- In this updated analysis of BEACON CRC (which includes ORR for all randomized patients and 6 additional 364 patients) and 6 months additional follow-up):
 - The triplet and doublet demonstrated improved OS and ORR in patients with BRAF V600E-mutant mCRC when compared with current standard of care chemotherapy

Overall Survival



Objective Response Rate

Confirmed Response by blinded central review	Triplet N=224	Doublet N=220	Control N=221
Objective Response Rate	27% (21%, 33%)	20% (15%, 25%)	2% (<1%, 5%)
95% (CI)			
p-value vs. Control	<0.0001	<0.0001	

The full updated BEACON results with subgroup analysis will be submitted to a future congress

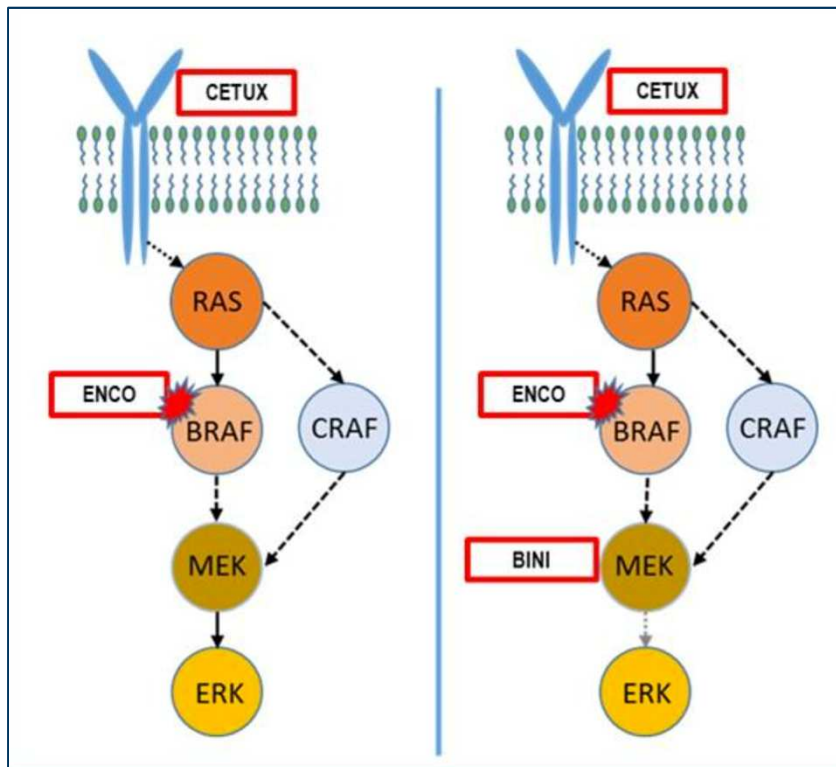
PRESENTED AT: Gastrointestinal Cancers Symposium

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PRESENTED BY: Scott Kopetz, MD

#G120

Therapie beim BRAFV600E - mutierten mCRC



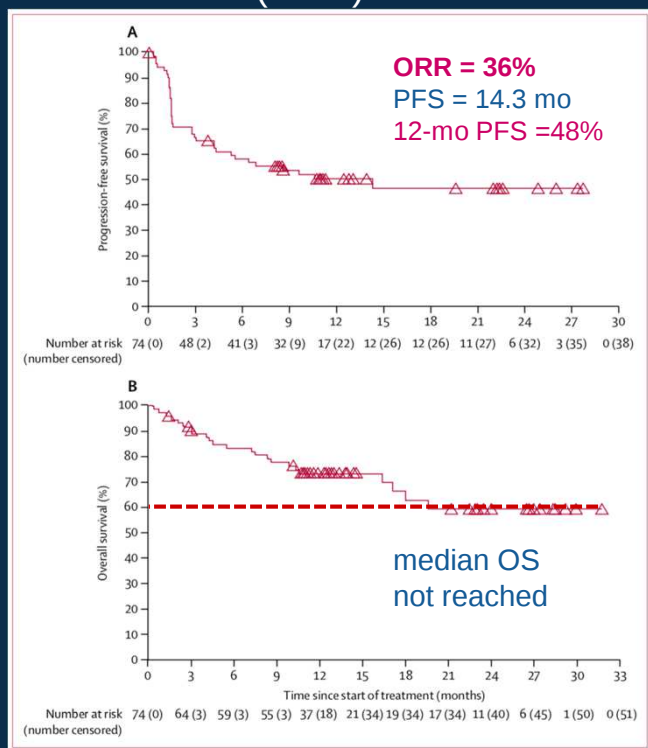
Schlussfolgerung

Encorafenib plus Cetuximab
mit oder ohne Binimetinib
induzierte einen längeren Erhalt
der QoL als die Standardtherapie
im Vergleichsarm

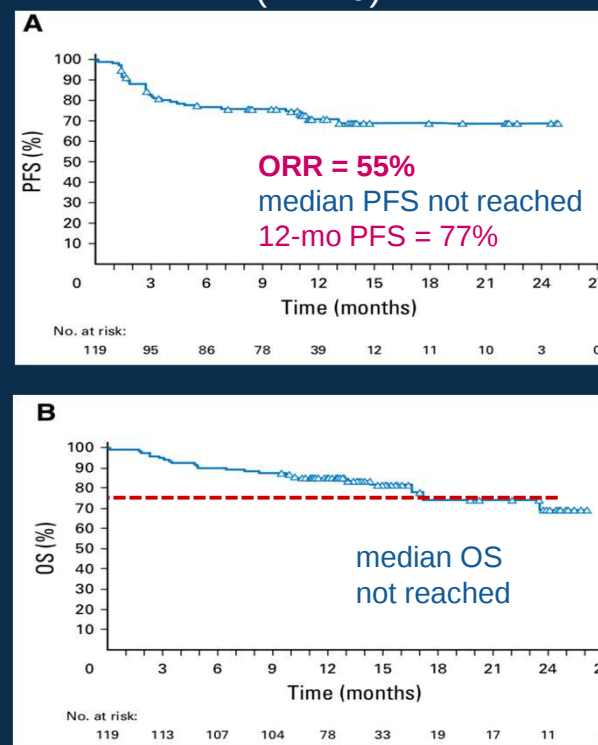
Scott Kopetz at 2020 Gastrointestinal Cancer Symposium

CheckMate 142: Nivolumab in MMRd mCRC

**Nivolumab 3mg/kg
(n=74)**



**Nivolumab 3mg/kg + Ipilimumab 1mg/kg
(n=119)**



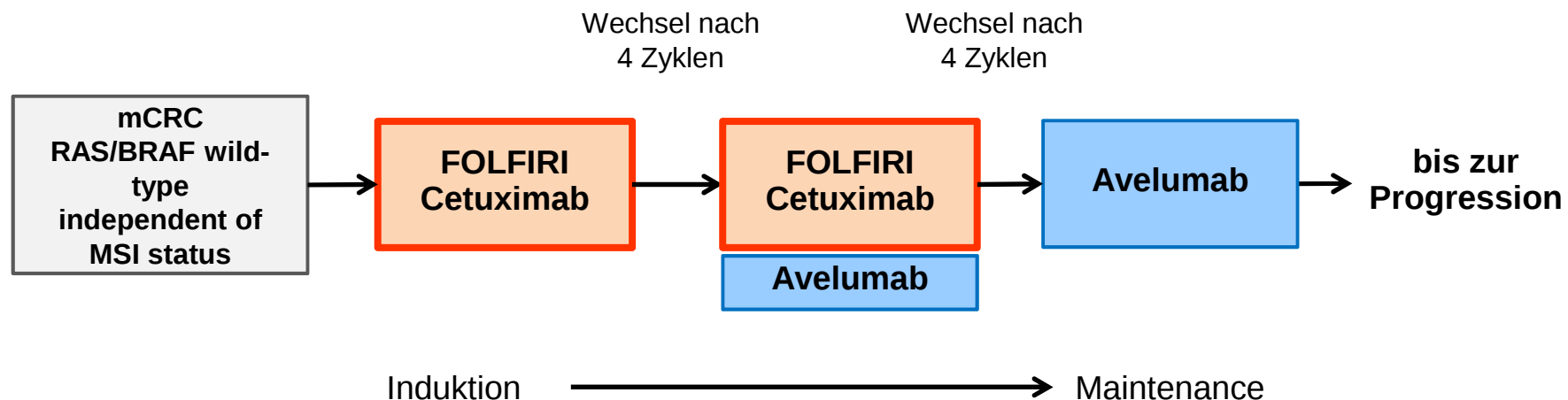
* Nivo 3mg/kg q2w until PD

** Nivo 3mg/kg plus Ipi 1 mg/kg q 3w for 4 doses, then Nivo 3mg/kg q2w

Overman MJ, Lancet Oncol 2017
Overman MJ, JCO 2017

FIRE-6 Avelumab in MMRp mCRC

Phase-IIa Design (n = 55)



Primärer Endpunkt:

PFS

Sekundäre Endpunkte:

Sicherheit, ORR, OS,
PFS Rate nach 12 Monaten
Translationale Forschung,

Herzlichen Dank für Ihre Aufmerksamkeit

