

XXIII JORNADA DE REVISIÓN DEL
**congreso
americano[®]
de
oncología**

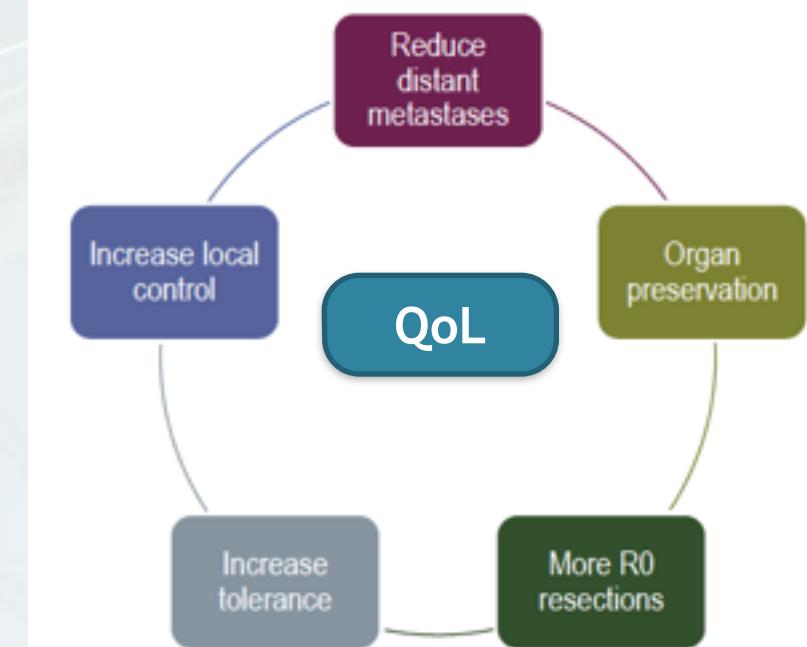
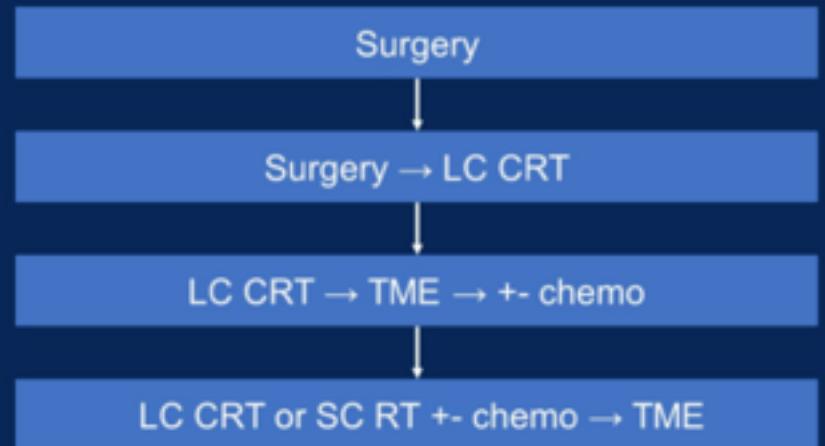
DIGESTIVO COLORRECTAL

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Hospital Universitario de Burgos
23 de junio de 2023



ENFERMEDAD LOCALIZADA

Evolution of management of LARC



“TNT”

- Induction
- Consolidation

QT

- Double?
- Triplet?

RT

- Short?
- Long?
- Omitted?
- Boost?

Rectal Preservation

- Watch and wait
- Local excision

Preoperative Chemotherapy with Selective Chemoradiation versus Chemoradiation for Locally Advanced Rectal Cancer: The PROSPECT Trial (Alliance N1048)

PROSPECT Study Motivation

- Long term toxicity from pelvic chemoradiation:
 - Impaired bowel, bladder, and sexual function¹
 - Increased risk of pelvic fracture and second malignancy²
 - Impaired marrow reserve³
 - Infertility and premature menopause⁴
 - Increasing diagnosis of rectal cancer before age 50⁵

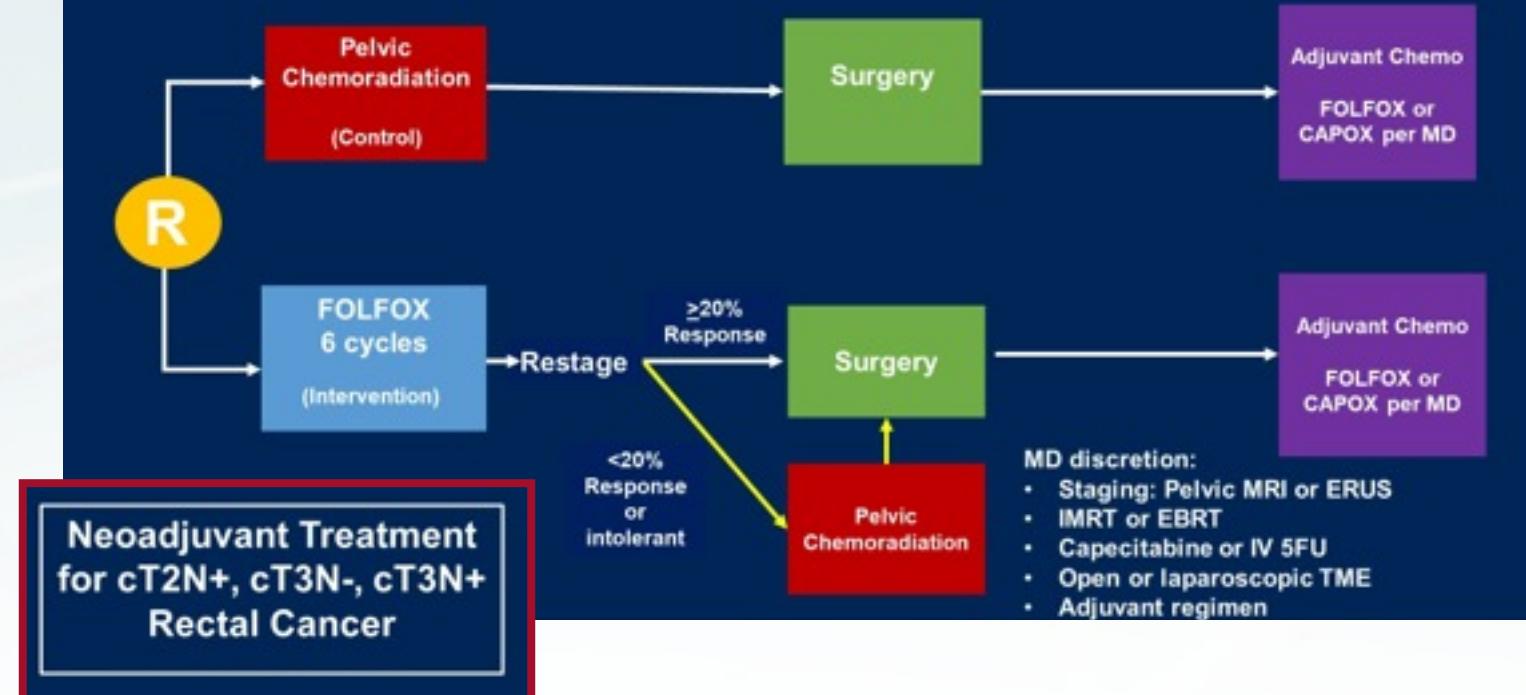
PROSPECT Trial Hypothesis

- Neoadjuvant chemotherapy with FOLFOX and only selective use of pelvic chemoradiation will be noninferior to routine use of pelvic chemoradiation for locally advanced rectal cancer

ORIGINAL ARTICLE

Preoperative Treatment of Locally Advanced Rectal Cancer

PROSPECT Study Full Schema



- Primary Endpoint:
- Disease Free Survival

Inclusion:

- Clinical Stage T2N+, T3N-, T3N+
- Chemoradiation is indicated
- Candidate for sphincter-sparing surgery
- Tumor requiring an APR
- cT4 tumor
- ≥ 4 pelvic lymph nodes $\geq 1\text{cm}$ in short axis

Exclusion:

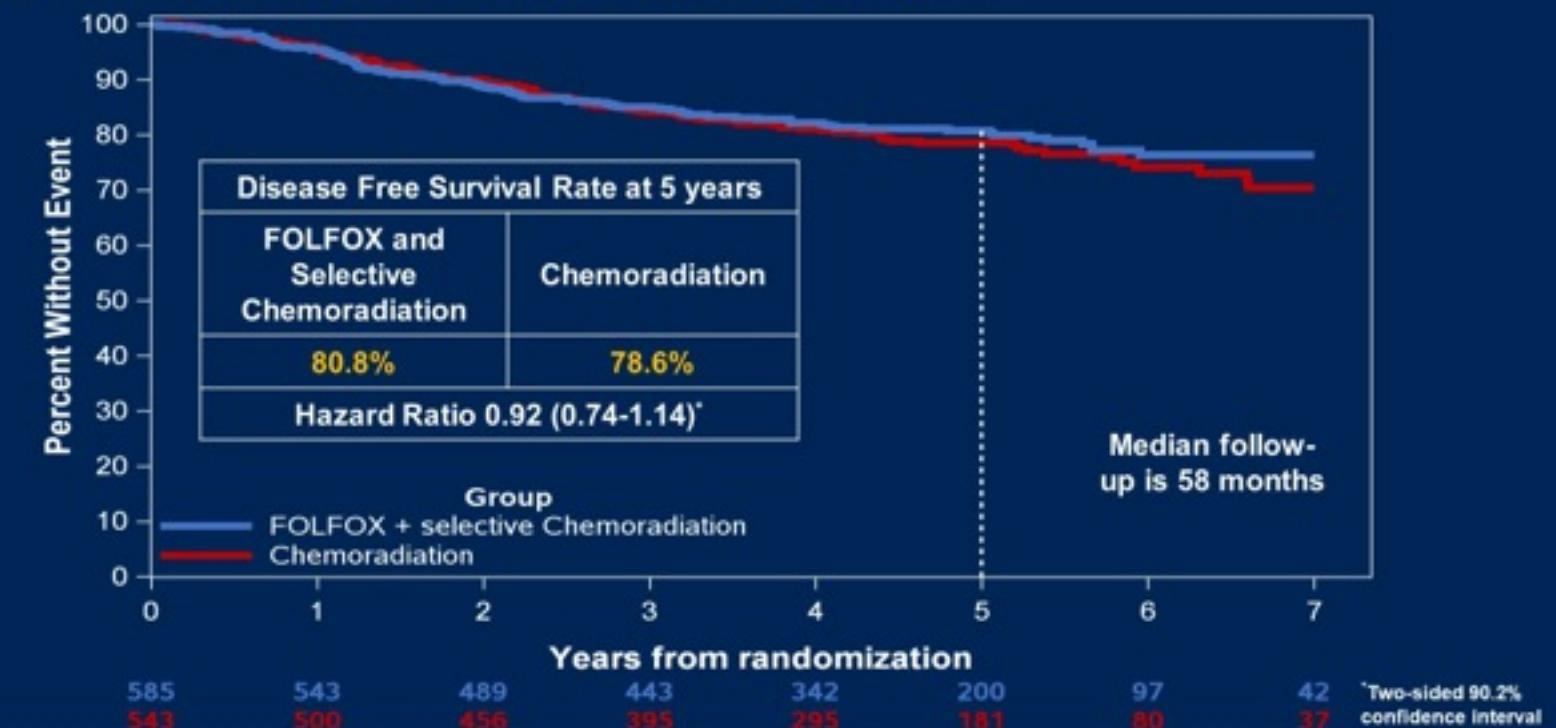
Non-inferiority Hypothesis for Disease Free Survival

Non-inferiority could be claimed if the upper limit of the two-sided 90.2% confidence interval of the hazard ratio (HR) did not exceed 1.29.

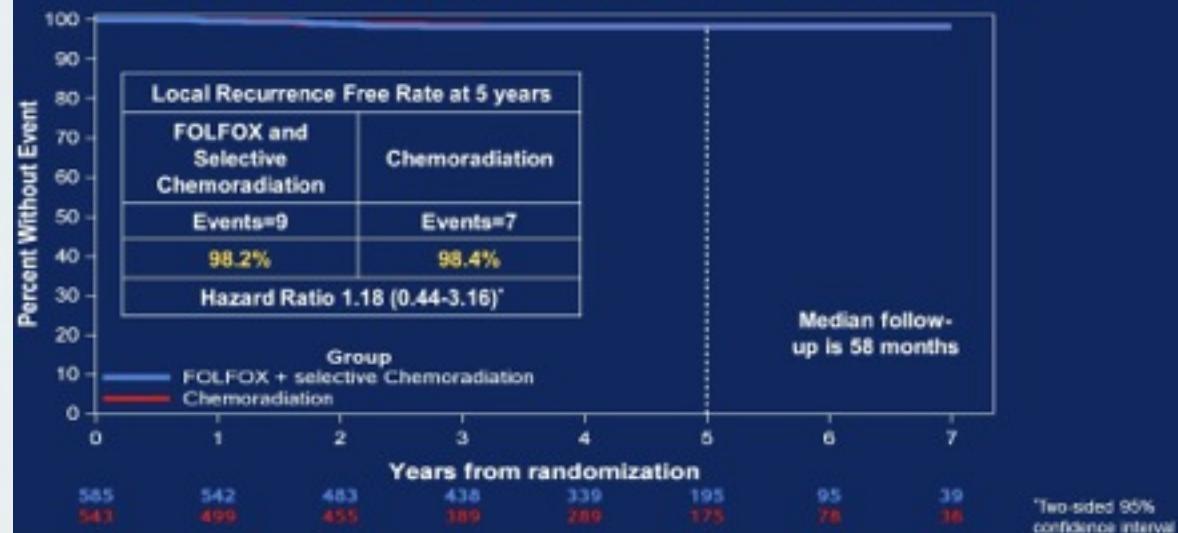
This corresponds to an absolute difference in 5-year DFS of <5%.



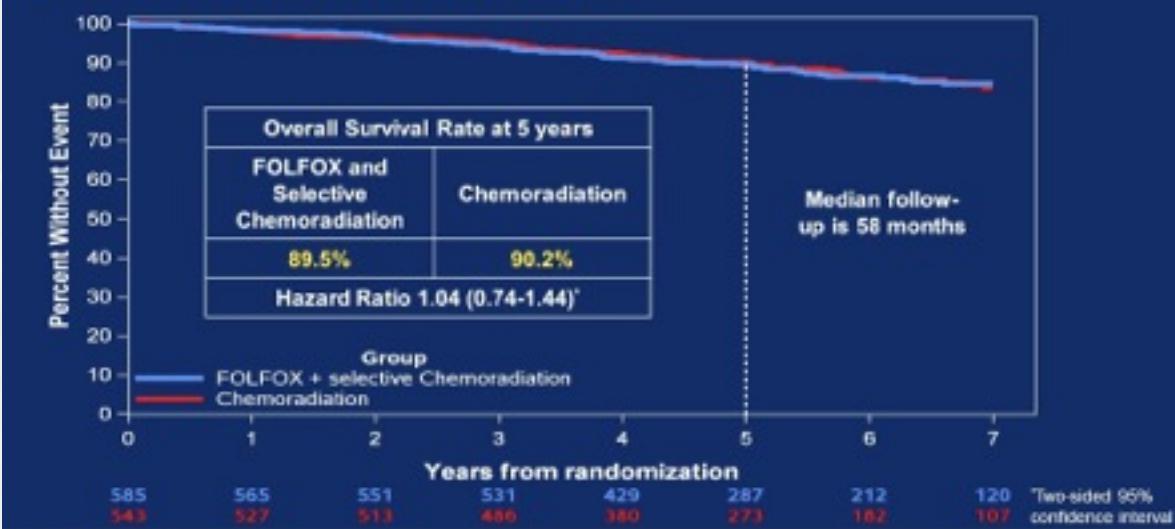
PROSPECT: Disease Free Survival



PROSPECT: Freedom from Local Recurrence



PROSPECT: Overall Survival



Surgical and Pathologic Endpoints

Secondary endpoints in participants who completed Surgery	FOLFOX and Selective Pelvic Chemoradiation N=535	Pelvic Chemoradiation N=510
Complete (R0) Rectal Resection	99%	97%
Low Anterior Resection Rate	98%	98%
Pathologic Complete Response	22%	24%
Positive Radial margin	1.2%	1.5%

Adjuvant Treatment and Therapy Duration

Adjuvant treatment in patients who had surgery	FOLFOX and Selective Pelvic Chemoradiation N=535	Pelvic Chemoradiation N=510
Received any adjuvant chemotherapy	82%	83%
Median duration (IQR) from randomization to last dose of postoperative therapy (weeks)	35 (33, 39)	37 (34, 40)

Use of Pelvic Chemoradiation in patients randomized to FOLFOX

9% (53/585) of participants randomized to FOLFOX received neoadjuvant chemoradiation either because:

Restaging demonstrated clinical response <20% or

They did not tolerate at least 5 cycles of FOLFOX

PROSPECT: Clinician-Reported Toxicity

Most severe toxicity during observation period based on CTCAE v. 4.0	FOLFOX and Selective Chemoradiation 12 weeks*	Chemoradiation 6 weeks
Neoadjuvant grade ≥3 adverse events	41%	23%
Adjuvant grade ≥3 adverse events	25%	39%

*22 weeks if also treated with chemotherapy

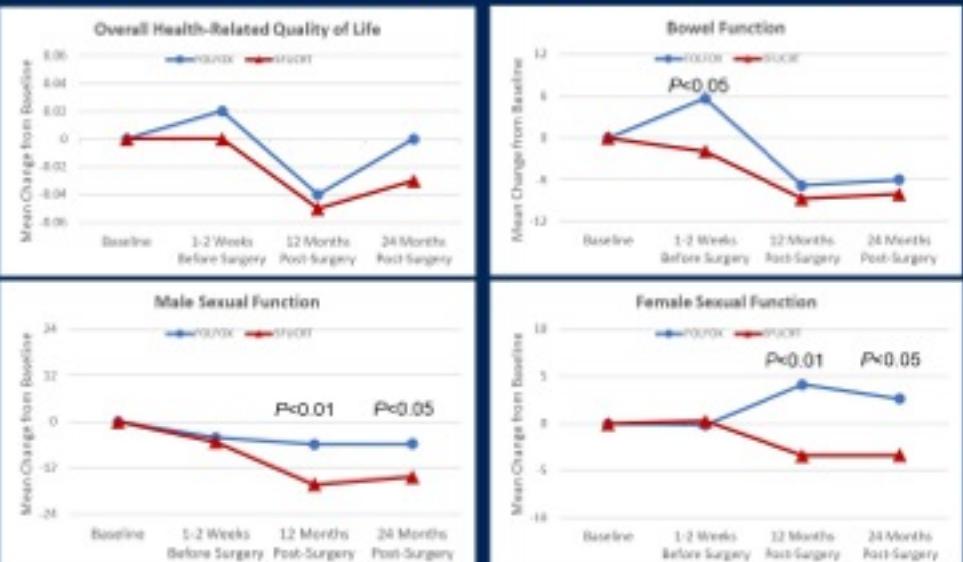
During Neoadjuvant treatment:

- More diarrhea in the RT group
- More neuropathy in the FOLFOX group

During Adjuvant treatment:

- More diarrhea in the RT group
- More neuropathy in the RT group

PROSPECT: Quality of Life Evaluation



Quality of Life:
Trend, but no
significant
difference
between groups

Bowel function
and sexual
function favor
FOLFOX group

N=373
Positive values represent
improvement compared to
baseline

Neoadjuvant FOLFOX, with only selective use of pelvic chemoradiation, is a safe and effective treatment option for patients with cT2N+, cT3N-, or cT3N+ rectal cancer

Long-term Outcome of Neoadjuvant mFOLFOX6 with or without Radiation versus Fluorouracil plus Radiation for Locally Advanced Rectal Cancer(FOWARC): A Multicenter, Randomized Phase III Trial

Study design



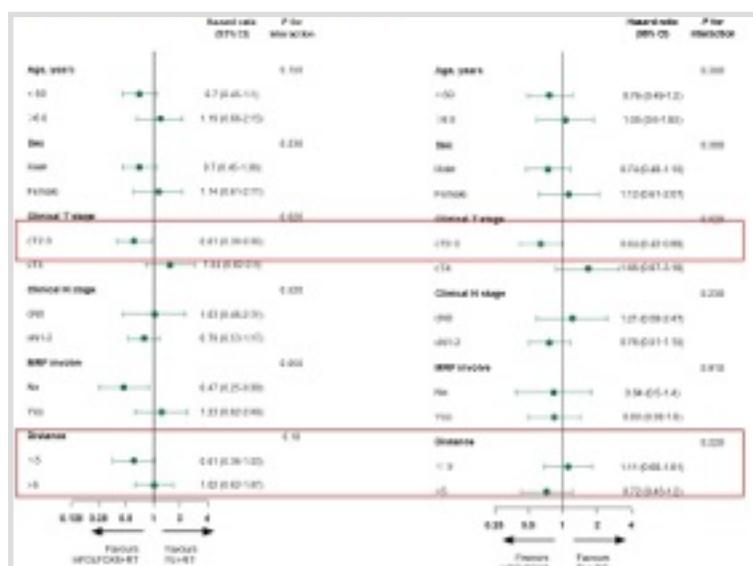
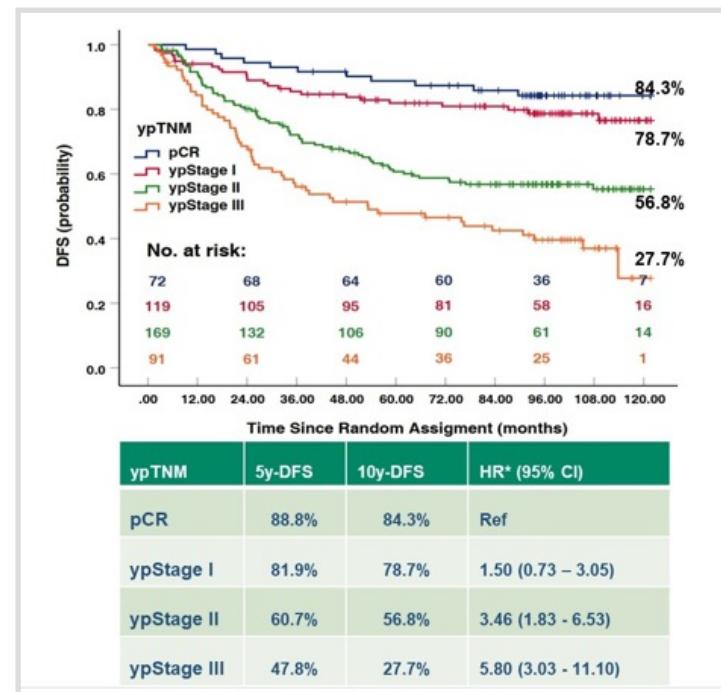
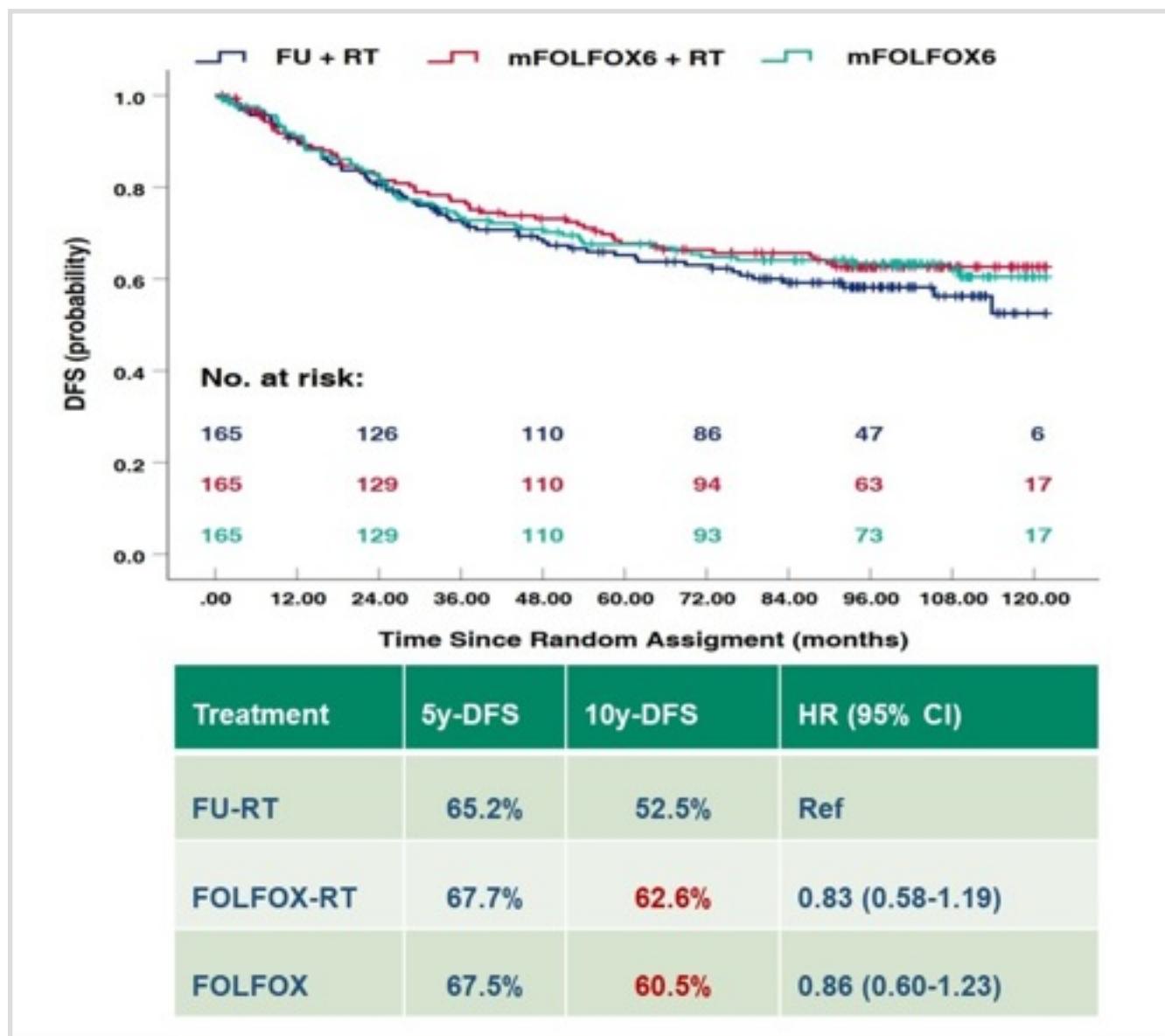
Study endpoints

- Primary endpoint:
3-year disease free survival (3y-DFS)
- Secondary endpoints:
pCR, OS, morbility, QOL ect

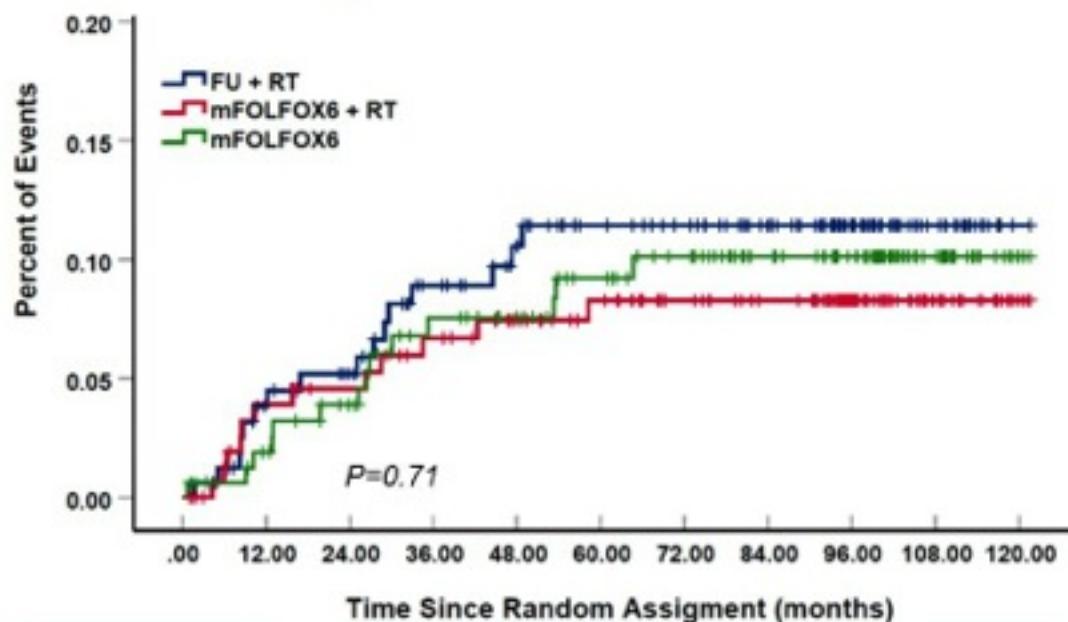
Clinical characteristics (ITT)

Items	FU-RT N=165	FOLFOX-RT N=165	FOLFOX N=165
Age median,yrs (mean±SD)	53.99±11.91	52.16±11.81	54.10±12.13
Gender, Male (%)	103 (62.4)	114 (69.1)	108 (65.5)
cT4 (%)	57 (34.5)	56 (33.9)	50 (30.3)
cN1 (%)	84 (50.9)	88 (53.3)	76 (46.1)
cN2 (%)	44 (26.7)	47 (28.5)	43 (26.1)
MRF positive (%)	32 (31.5)	38 (37.0)	33 (25.9)
Distance from anal verge ≤5cm	90 (54.5)	83 (50.3)	70 (42.4)

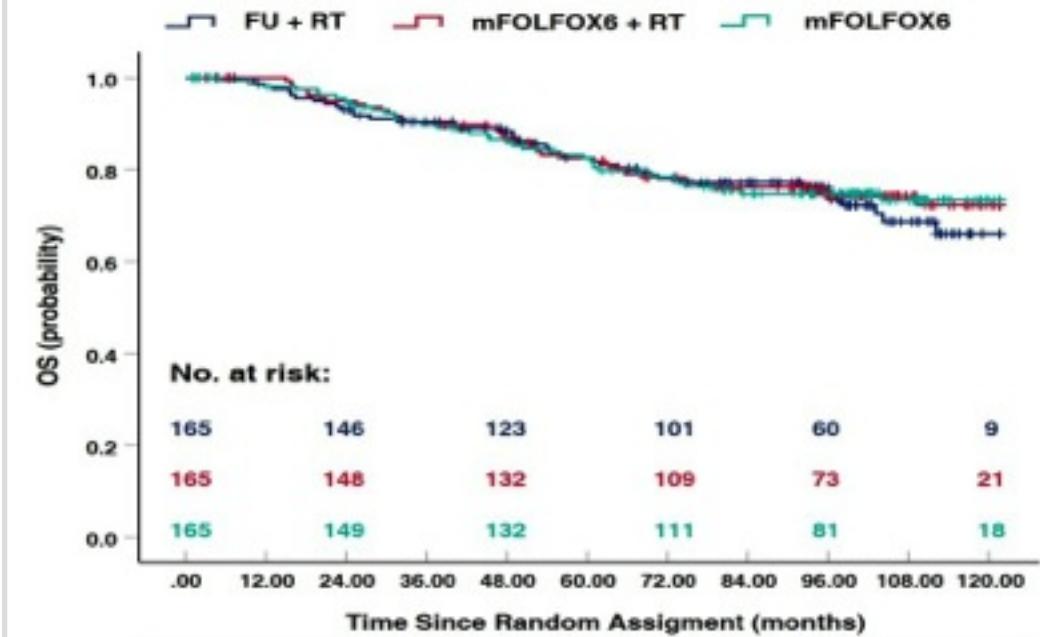
Long term Disease-Free Survival (DFS)



Locoregional recurrence (LR)



Overall survival

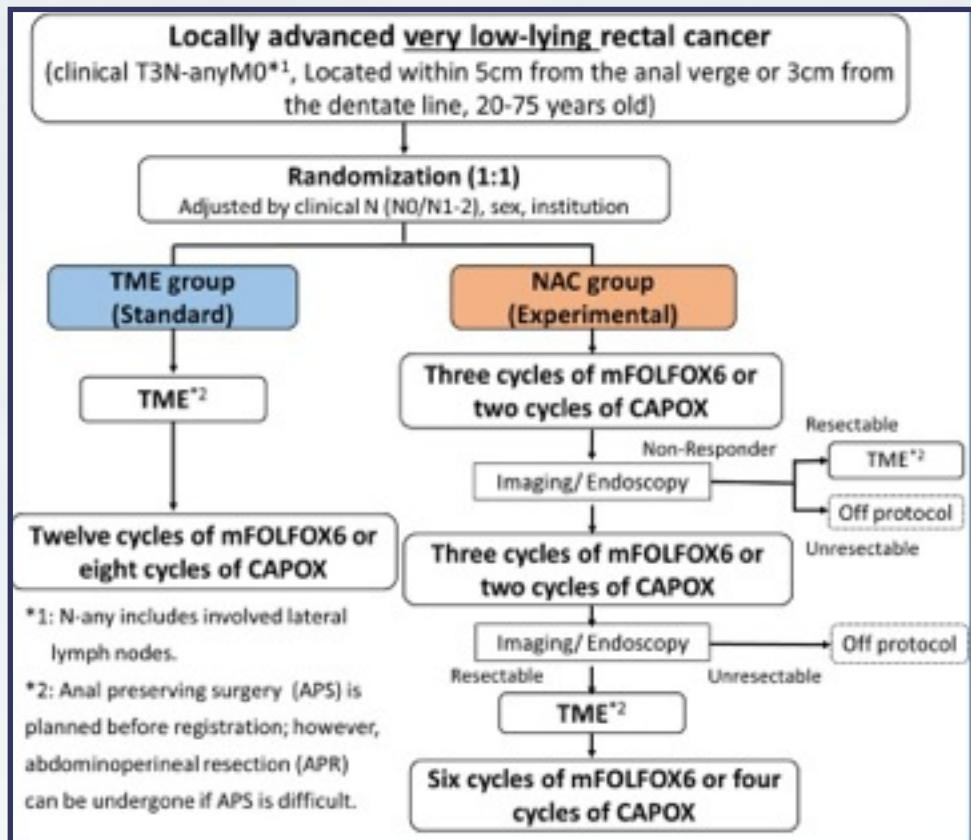


Treatment	5y-OS	10y-OS	HR (95% CI)
FU-RT	82.5%	65.9%	Ref
FOLFOX-RT	81.8%	72.3%	0.91 (0.58-1.41)
FOLFOX	81.8%	73.4%	0.91 (0.58-1.41)

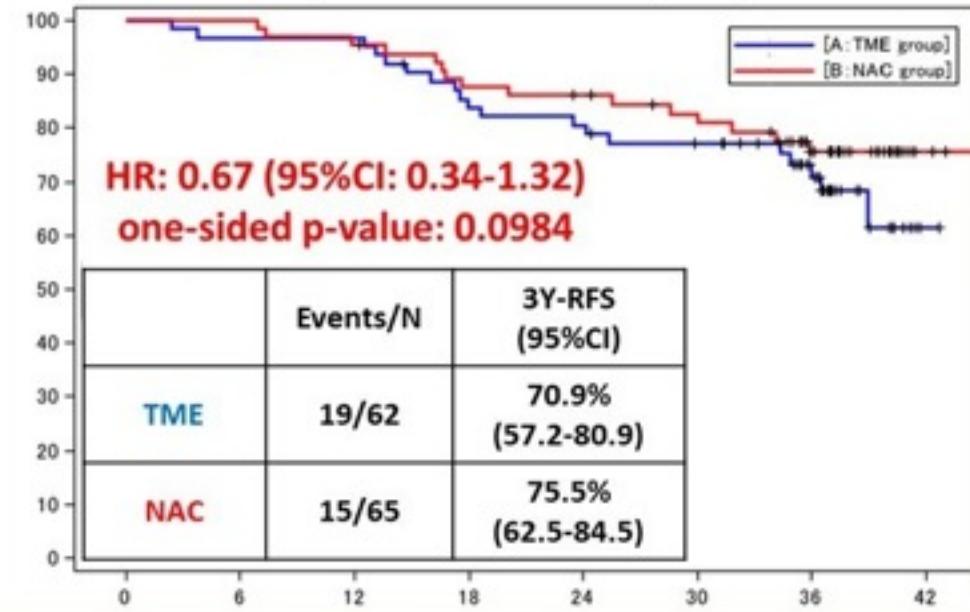
Neoadjuvant chemotherapy (NAC) followed by total mesorectal excision and adjuvant chemotherapy versus TME followed by adjuvant chemotherapy in very low-lying cT3 rectal cancer (NAIR): a multicenter, randomized, open-label, phase 2/3 trial

Primary Endpoint:

- Recurrence-free survival (3yrs)



Relapse-free Survival (RFS)



➤ Median Wexner incontinence score

Pre: Preoperatively, M: month(s)

	Pre	3M	6M	12M	24M	36M
TME group	1.0	14.0	12.0	10.0	9.0	10.0
NAC group	1.0	15.0	13.5	12.0	11.0	11.0

At 3 years after randomization, 61 pts in the NAC group (93.8%) and 52 pts in the TME group (83.9%) could defecate via their anuses (P=0.092).

➤ Grade 3 or higher post-operative complications

	TME (N=62)	NAC (N=65)	p value
Any grade 3 or higher complications	13 (21.0%)	13 (20.0%)	1.000
Anastomotic leakage	2 (3.2%)	2 (3.1%)	
Pelvic abscess	5 (8.1%)	6 (9.2%)	
Bowel obstruction	1 (1.6%)	0	

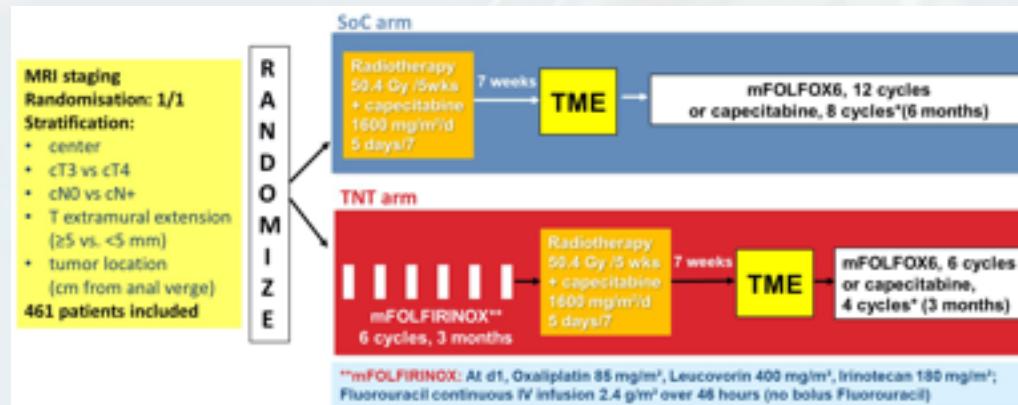
No treatment-related deaths occurred in either of the groups.

TNT trial results

Outcomes	RAPIDO		STELLAR		PRODIGE-23	
	TNT-SCRT	LCRT	TNT-SCRT	LCRT	TNT-LCRT	LCRT
pCR	28%	14%	21.8% (incl cCR)	12.3%	28%	12%
R0	90%	90%	91.5%	87.8%	95%	94%
3-yr DrTF	23.7%	30.4%	-	-	-	-
3-yr DFS	-	-	64.5%	62.3%	76%	69%
3-yr LR	8.3%	6.0%	8.4%	11.0%	LR: 4.3%	5.7%
3-yr DM	20.0%	26.8%	23.9%	24.7%	21%	28%
3-yr OS	89%	89%	86.5%	75.1%	91%	88%

Total neoadjuvant therapy with mFOLFIRINOX versus preoperative chemoradiation in patients with locally advanced rectal cancer: 7-year results of PRODIGE 23 phase III trial, a UNICANCER GI trial.

Primary: Disease-Free Survival (DFS)

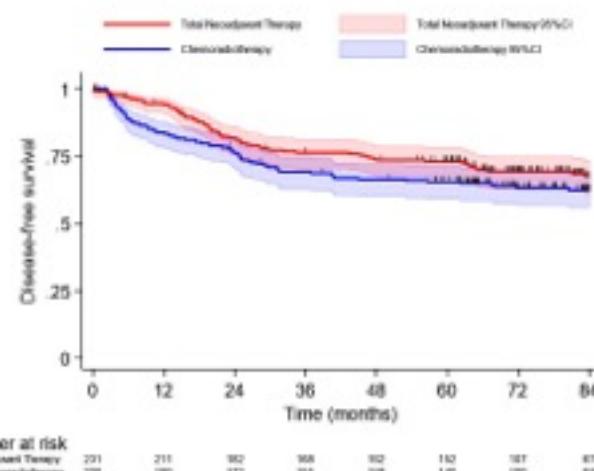


Cumulative incidence of rectal cancer recurrences

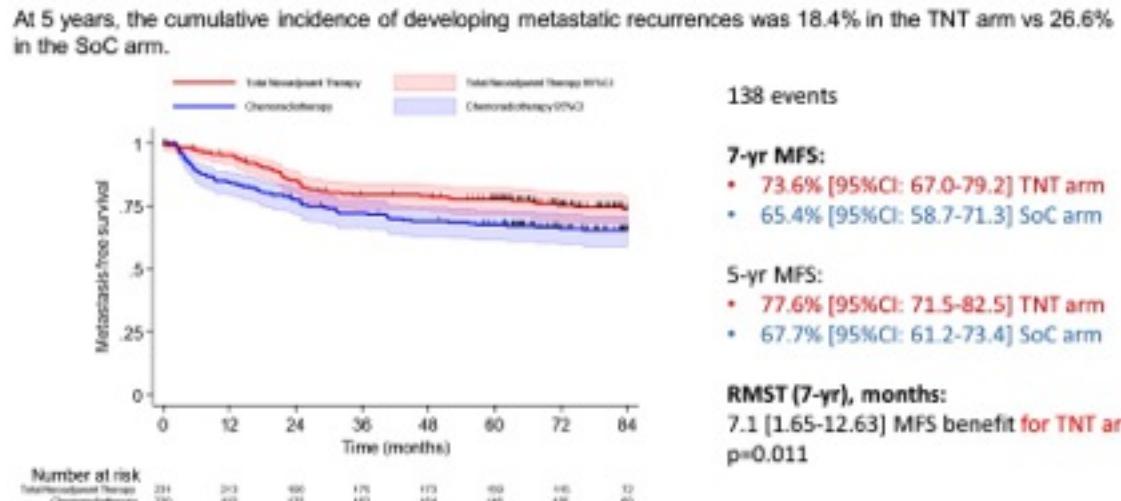
Results	TNT	SoC
Local		
At 5 years	4.7% [95%CI: 2.5-8.5]	6.4% [95%CI: 3.8-10.8]
At 7 years	5.3% [95%CI: 2.9-9.3]	8.1% [95%CI: 4.9-13.3]
Metastatic*		
At 5 years	18.4% [95%CI: 13.8-24.2]	26.6% [95%CI: 21.2-33.0]
At 7 years	20.7% [95%CI: 15.6-27.0]	27.7% [95%CI: 22.2-34.2]
Alive with metastases	19/44 (43%)	21/60 (35%)

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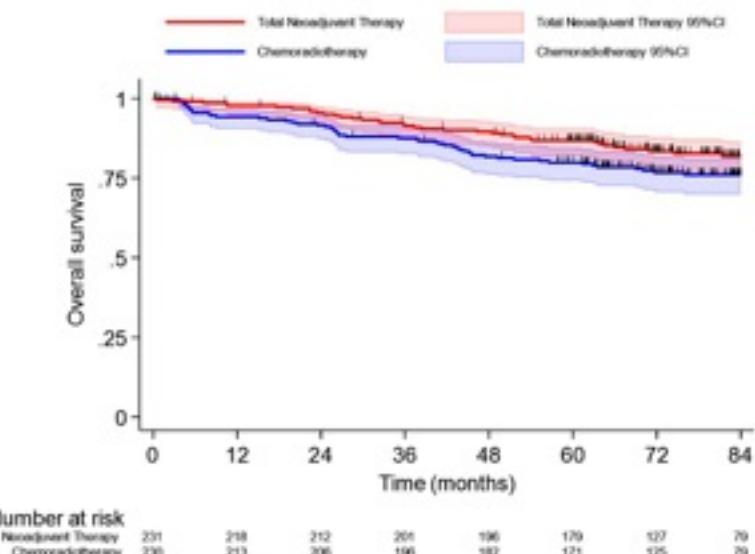
Disease-Free Survival



Metastasis-free Survival

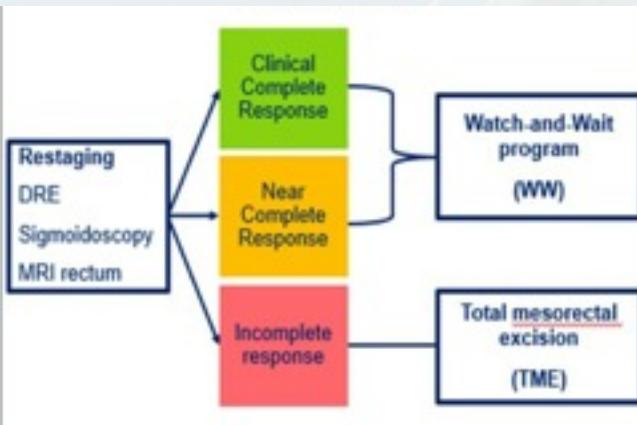
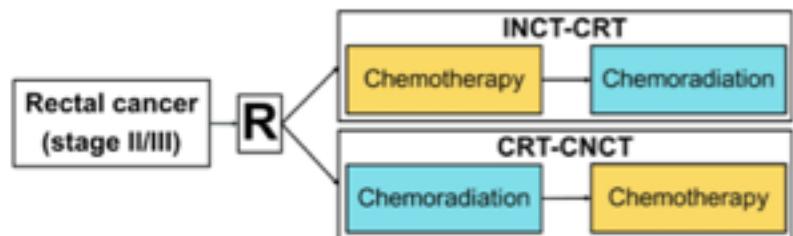


Overall Survival



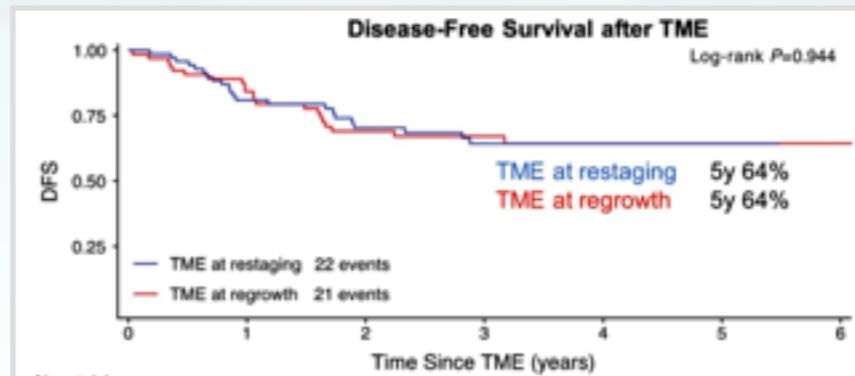
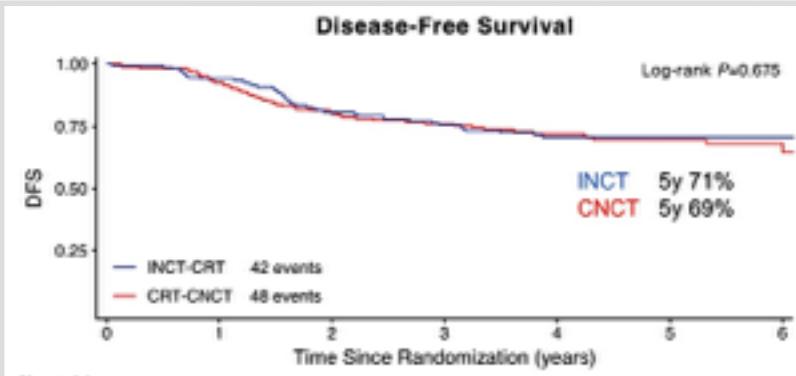
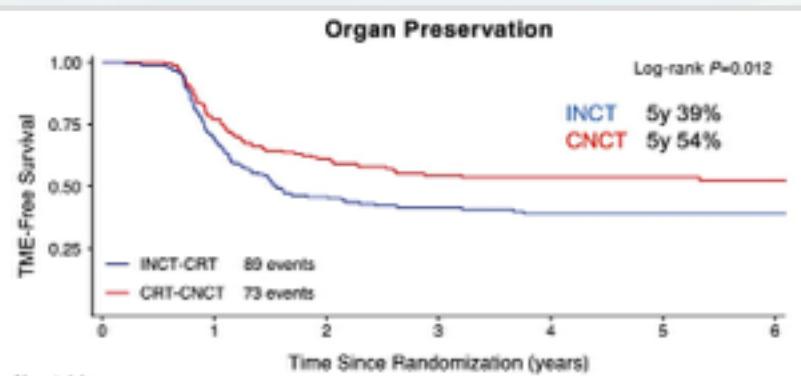
Sustained organ preservation in rectal cancer patients treated with total neoadjuvant therapy: Long-term results of the OPRA trial

- Randomized phase II trial.



3- and 5-year rates with 95% CI.

	Induction		Consolidation		P
	3-year, %	5-year, %	3-year, %	5-year, %	
DFS	77 (70-84)	72 (65-80)	76 (70-83)	71 (63-78)	0.60
TME-free survival	41 (34-50)	39 (32-48)	55 (48-63)	54 (47-63)	0.01
LRFS	95 (91-99)	94 (90-98)	94 (90-98)	90 (85-95)	0.27
DMFS	83 (78-90)	82 (75-89)	83 (77-89)	79 (72-86)	0.66
OS	95 (92-99)	88 (82-94)	94 (91-98)	88 (82-94)	0.73

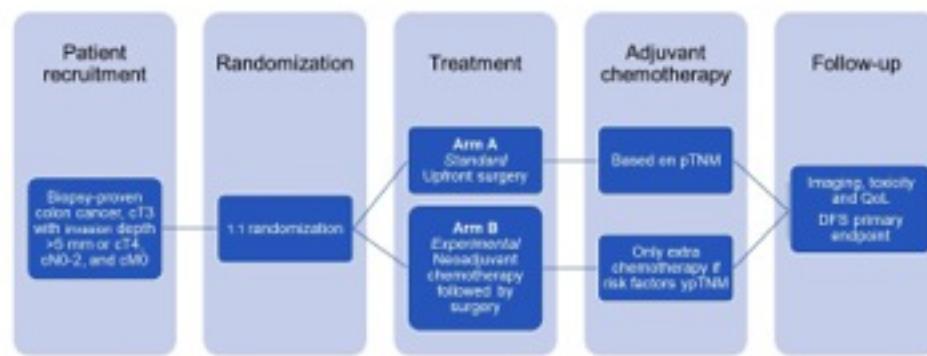


36% patients who started WW developed a regrowth
94% occurred within 2 years and 99% occurred within 3 years

Phase III randomized clinical trial comparing the efficacy of neoadjuvant chemotherapy and standard treatment in patients with locally advanced colon cancer

The Scandinavian NeoCol trial

Study design



- Primary endpoint was Disease free survival (DFS)
- Secondary endpoints
 - Rate of patients fulfilling the criteria for adjuvant chemotherapy
 - Overall survival (OS)
 - Toxicity
 - Quality of life

Patients

	Upfront surgery (standard)	Neoadj. treatment (experimental)	All
N	122	126	248
Age (median, min;max)	69.4 (30.3;81.8)	63.8 (23.8;84.3)	66.3 (23.8;84.3)
Sex, N (%female)	45 (37%)	66 (52%)	111 (45%)
Localization of tumor, N (%)			
Right	52 (43%)	59 (47%)	111 (45%)
Left	70 (57%)	66 (52%)	136 (54%)
Right or left not specified	-	1 (1%)	1 (1%)
T-category at baseline CT scan, N (%)			
T3	89 (74%)	91 (72%)	180 (73%)
Median extramural invasion (mm)	7 mm	8 mm	7 mm
T4	31 (25%)	33 (26%)	65 (26%)
T3 or T4 not specified	1 (1%)	2 (2%)	3 (1%)
Performance status, N (%)			
0	107 (88%)	115 (91%)	222 (90%)
1	13 (10%)	8 (6%)	21 (8%)
2	-	1 (1%)	1 (1%)
0, 1 or 2 not specified	2 (2%)	2 (2%)	4 (2%)

Pathology

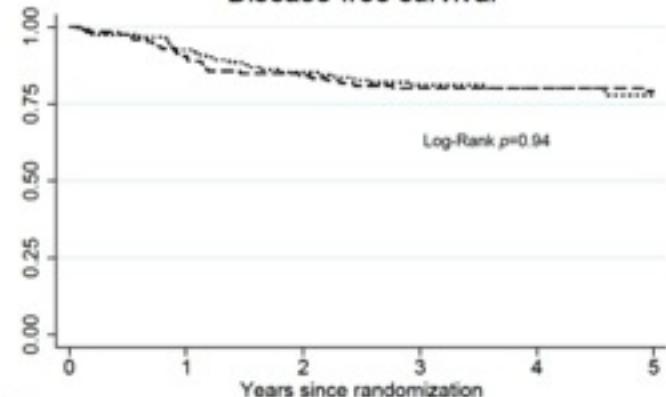
	Upfront surgery (standard)	Neoadj. treatment (experimental)	All
Involvement of resection margin, N (%) R1	10 (9%)	9 (7%)	19 (8%)
R0	109 (90%)	114 (93%)	223 (91%)
Unknown	2 (1%)	-	1 (1%)
pypT-category at surgery*, N (%)			
0	-	4 (3%)	4 (2%)
1	2 (2%)	1 (1%)	3 (1%)
2	3 (2%)	7 (6%)	10 (4%)
3	75 (62%)	76 (62%)	151 (62%)
4	39 (32%)	35 (28%)	73 (30%)
pypN-category at surgery, N (%)			
0	57 (48%)	72 (59%)	129 (54%)
1	43 (35%)	31 (25%)	73 (30%)
2	20 (17%)	19 (16%)	39 (16%)
Perineural invasion, N (%yes)			
	21 (18%)	19 (15%)	40 (17%)
Vascular invasion, N (%yes)			
	48 (39%)	30 (25%)	77 (32%)
Perforation, N (%yes)			
	2 (2%)	3 (2%)	5 (2%)

Adjuvant chemotherapy indication

	Upfront surgery (standard)	Neoadj. Treatment (experimental)	p
Criteria for adjuvant chemotherapy fulfilled*, N (%yes)	88 (73%)	72 (59%)	0.03

*according to national guidelines applied to p ypTNM, investigator

Disease-free survival

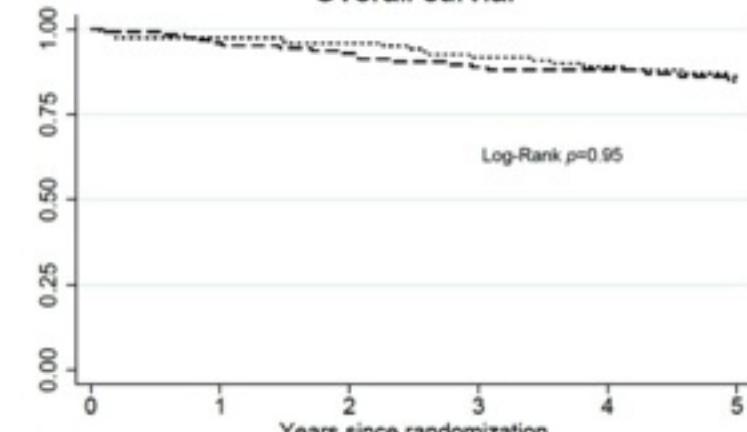


Number at risk

Adjuv.	122	113	104	95	79	64
Neoadj.	126	114	106	99	86	75

..... Adjuv. - - - Neoadj.

Overall survival

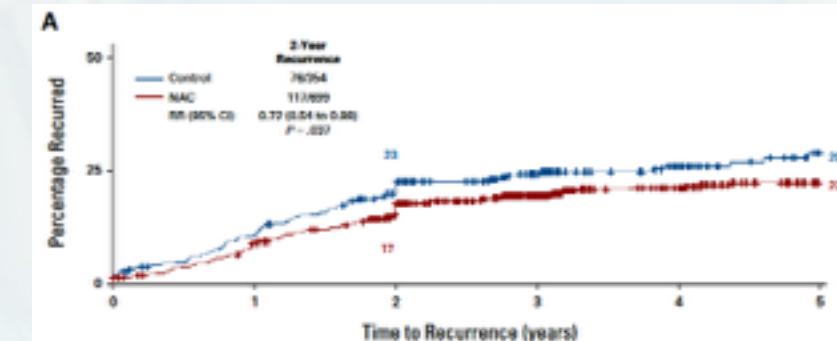
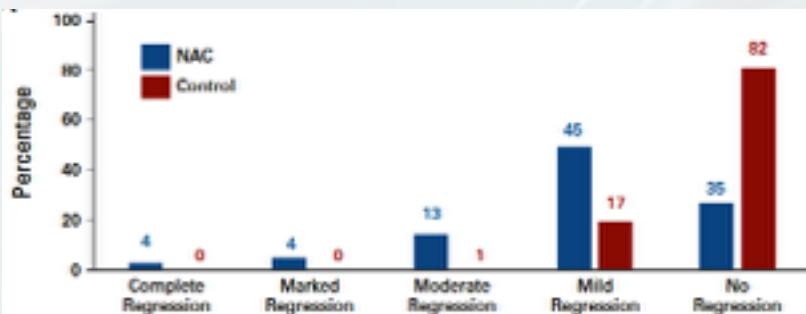
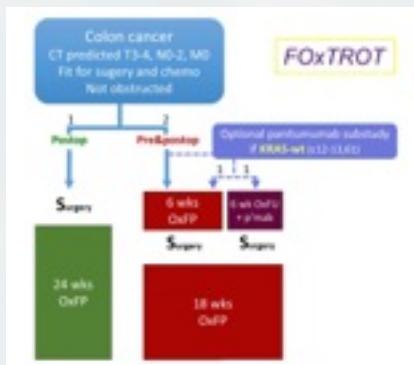


Number at risk

Adjuv.	122	119	117	108	92	75
Neoadj.	126	121	117	110	94	82

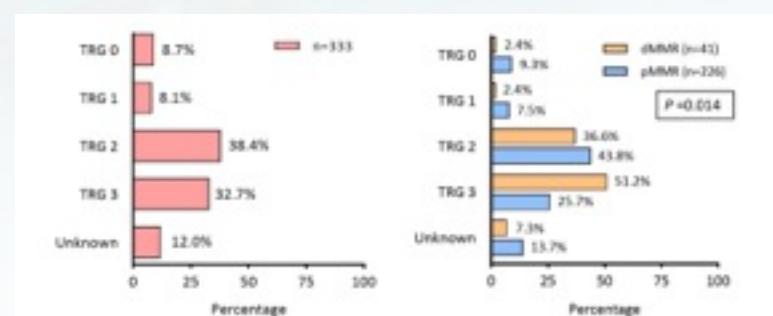
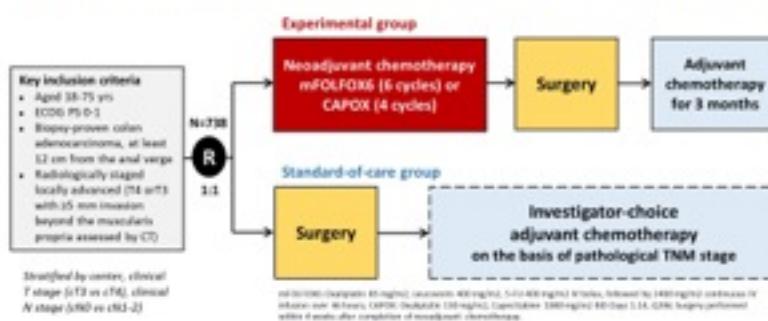
..... Adjuv. - - - Neoadj.

Preoperative Chemotherapy for Operable Colon Cancer: Mature Results of an International Randomized Controlled Trial

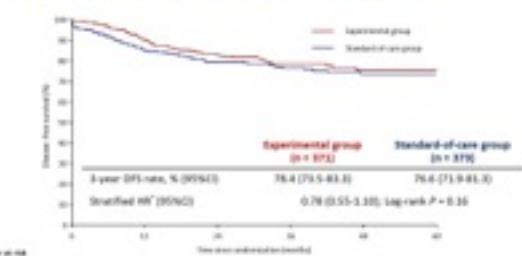


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Perioperative Chemotherapy With mFOLFOX6 or CAPOX for Patients With Locally Advanced Colon Cancer (OPTICAL): A Multicenter, Randomized, Phase III Trial

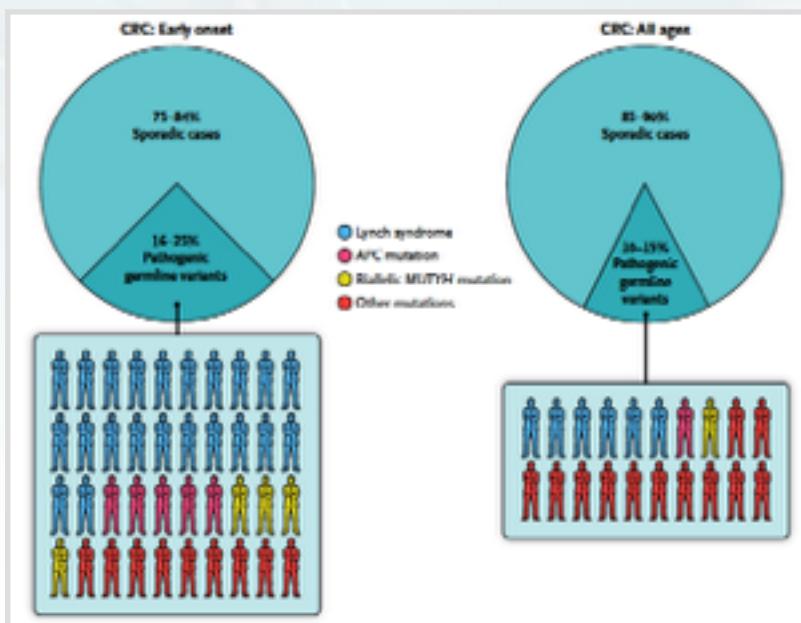
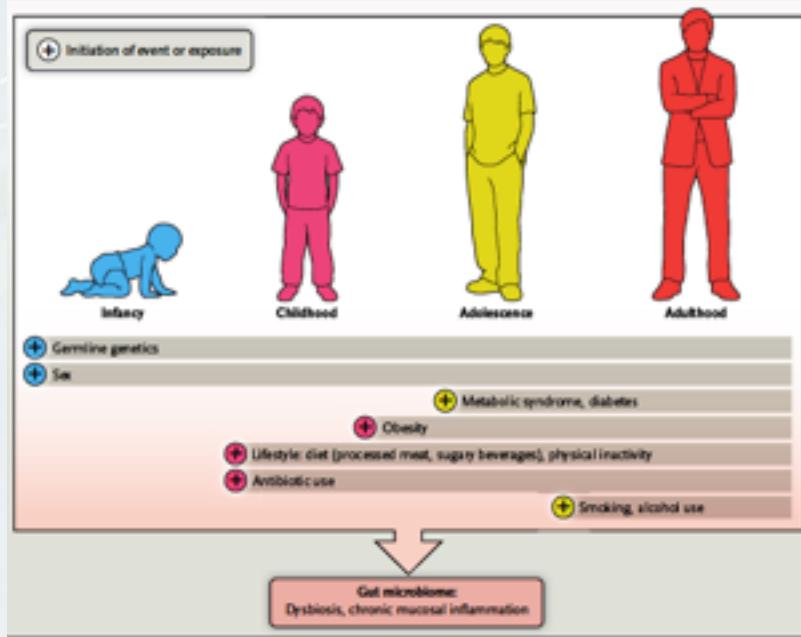
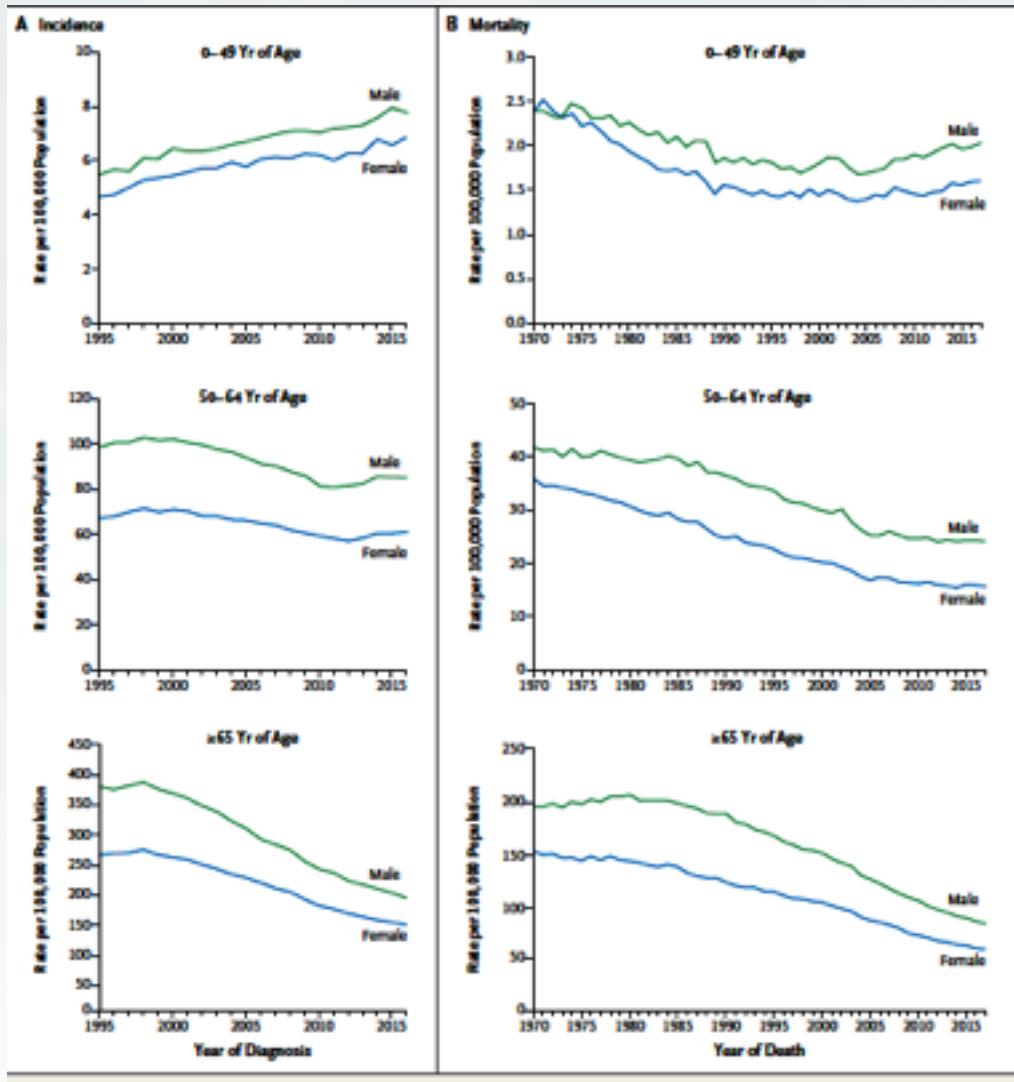


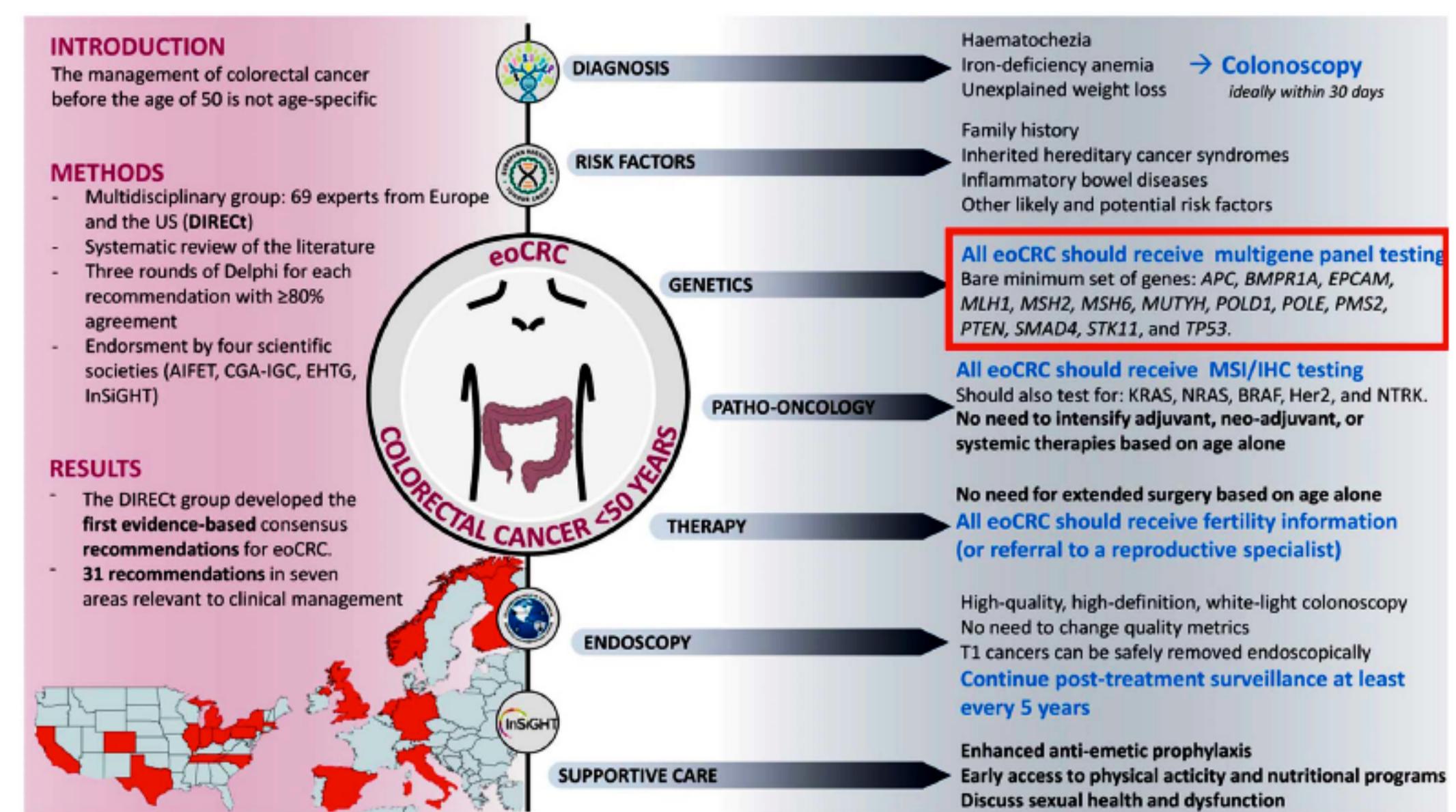
Disease-Free Survival (DFS) in mITT Population



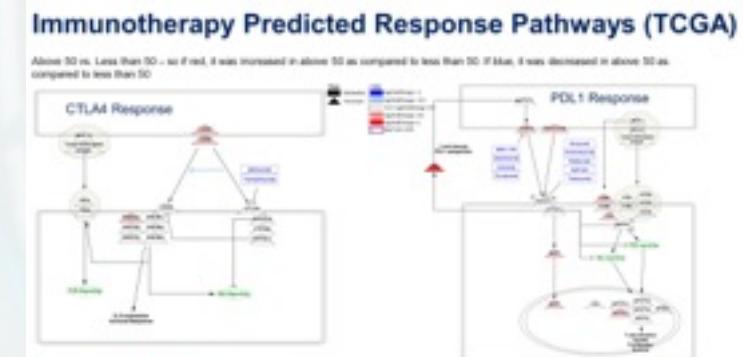
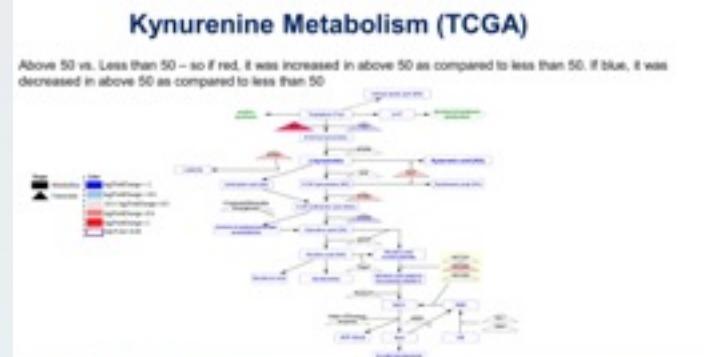
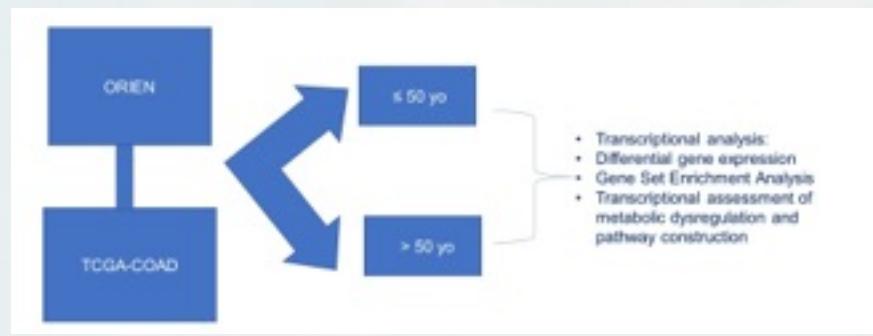
Huabin Hu.. ASCO 2022

Increasing Incidence of Early-Onset Colorectal Cancer

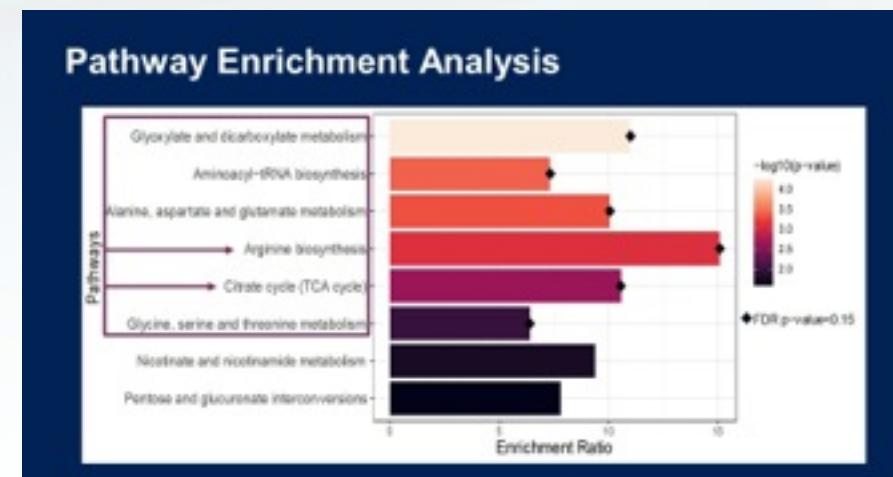
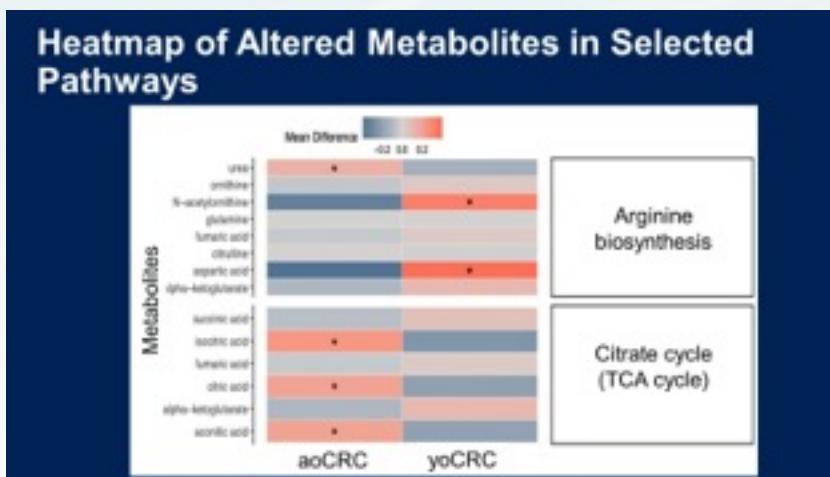




Abstract 3509: Transcriptional metabolic profiling in young onset colorectal cancer (YOCRC) patients

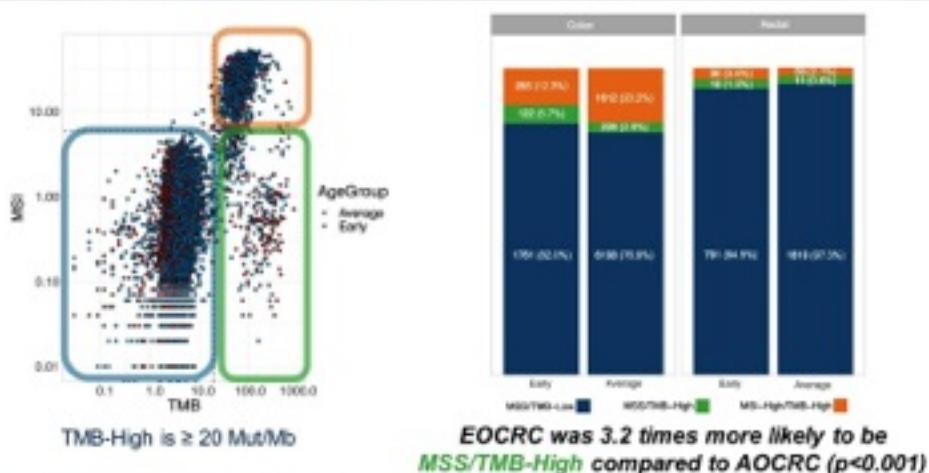
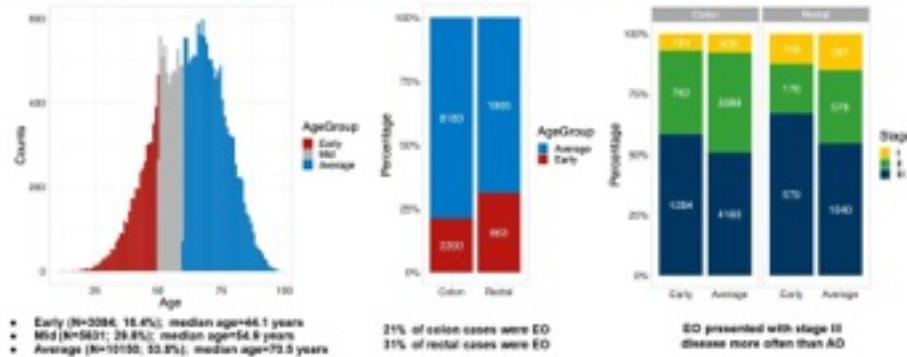


Metabolomic Differences in Young-onset versus Average-onset Colorectal Adenocarcinoma



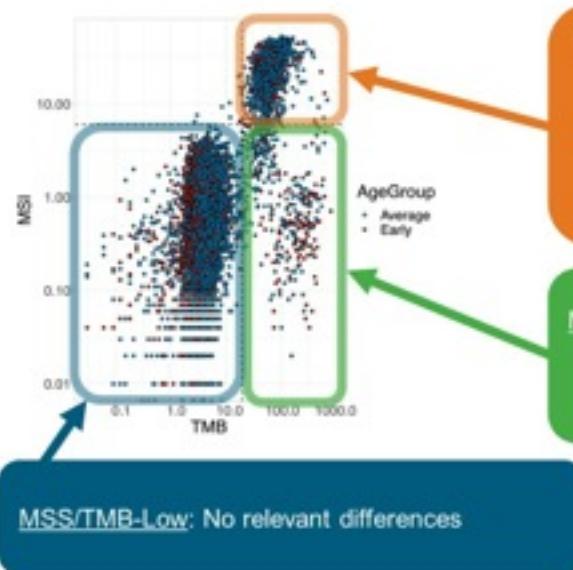
Evaluation of genomic alterations in early-onset versus average-onset colorectal cancer

EOCRC was more frequently rectal primary and stage III compared to AOCRC.



Key genomic differences between EO and AO

(significance: adjusted p-value < 0.05)



MSI-H/TMB-High: EO compared to AO has:

- More KRAS mutations (46% vs 15%)
- More PIK3CA mutations (47% vs 28%)
- More ERBB2/3 mutations (16% and 13% vs 7% and 5%)
- Fewer BRAF mutations (8% vs 65%)
- Fewer RNF43 mutations (25% vs 59%)

MSS/TMB-High: EO compared to AO has:

- More POLE mutations (65% vs 35%)
- Fewer ACVR2A mutations (32% vs 51%)

MSS/TMB-Low: No relevant differences



ENFERMEDAD AVANZADA

Efficacy of panitumumab in left-sided patients with MSS/MSI-L and RAS/BRAF WT: a biomarker study of the phase III PARADIGM trial

Chemotherapy-naïve patients with RAS WT mCRC (N=823)

Key eligibility criteria:

- Unresectable disease
- Age: 20–79 years
- ECOG PS: 0–1
- ≥1 evaluable lesion

Enrolled across 197 sites (May 2015 to June 2017)



Primary endpoint

- OS: Left-sided^a population; if significant, analyzed in overall population

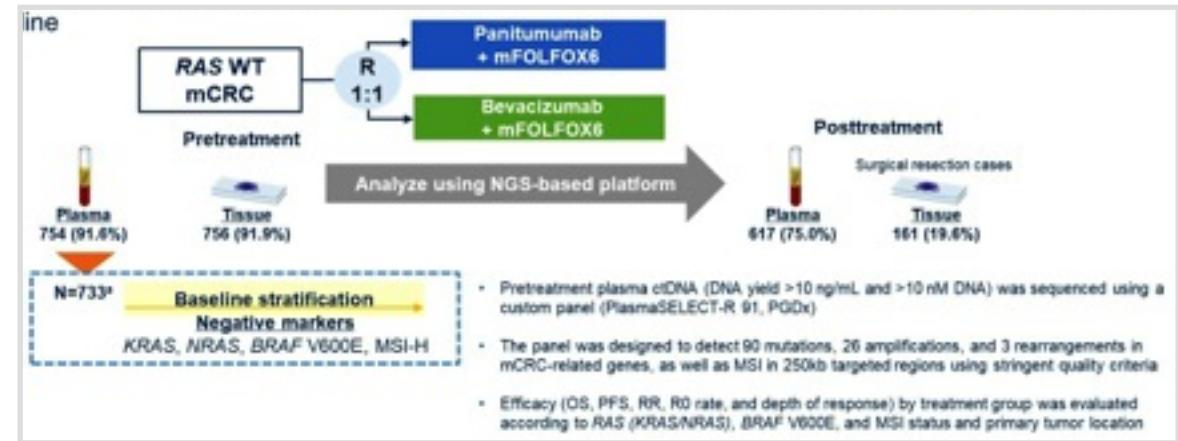
Secondary endpoints

- PFS, RR, depth of response, R0 rate: Left-sided^a and overall populations
- Safety: All treated patients

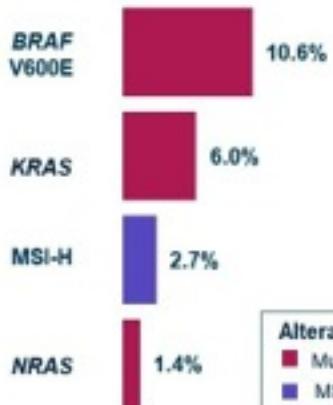
Exploratory endpoints

- ETS, depth of response, DCR: Left-sided^a and overall populations

Data cutoff date: January 14, 2022



Overall (N=733)



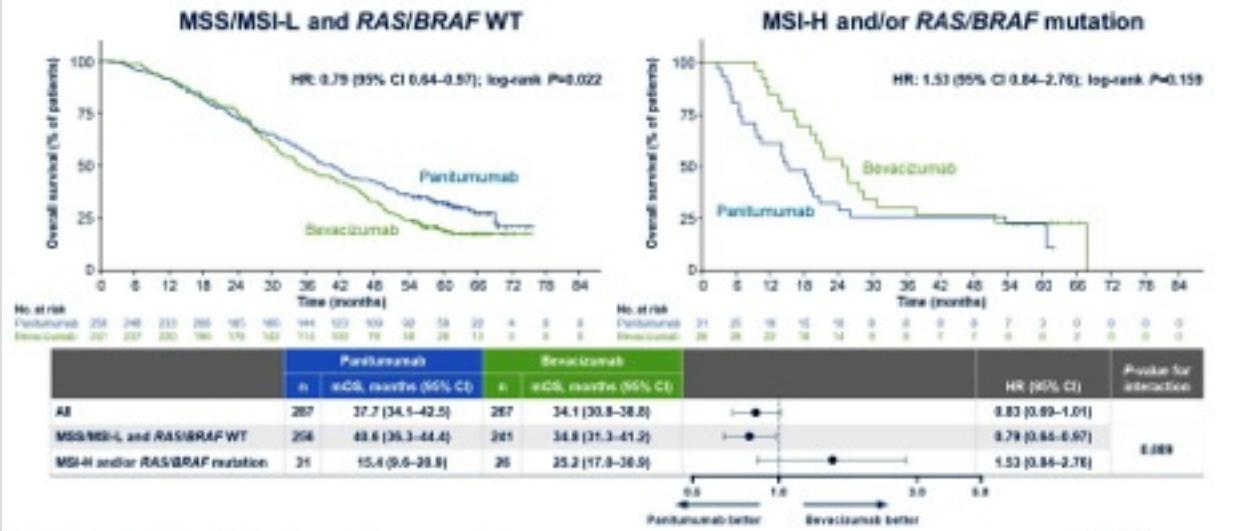
Co-occurring gene alterations by tumor sidedness

Gene alteration status, n (%)	Left-sided mCRC (n=554)	Right-sided mCRC (n=169)	Other ^a (n=10)	Overall (n=733)
MSI-H or RAS/BRAF mutation	57 (10.3)	73 (43.2)	5 (50%)	135 (18.4)
MSS/MSI-L and RAS/BRAF WT	497 (89.7)	96 (56.8)	5 (50%)	598 (81.6)

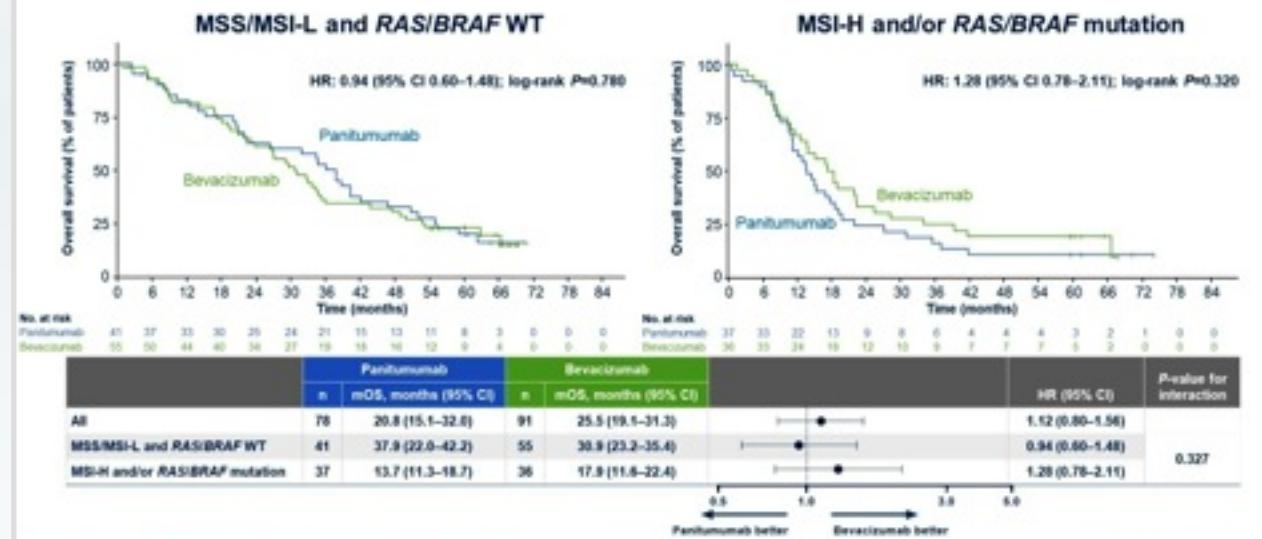
Incidence of gene alterations in baseline ctDNA

Gene alteration, n (%)	Left-sided mCRC (n=554)		Right-sided mCRC (n=169)		Overall population (N=733)	
	Panitumumab + mFOLFOX6 (n=287)	Bevacizumab + mFOLFOX6 (n=267)	Panitumumab + mFOLFOX6 (n=96)	Bevacizumab + mFOLFOX6 (n=91)	Panitumumab + mFOLFOX6 (n=368)	Bevacizumab + mFOLFOX6 (n=368)
BRAF V600E	16 (5.6)	8 (3.0)	26 (33.3)	27 (29.7)	42 (11.4)	36 (9.9)
KRAS	11 (3.8)	16 (6.0)	9 (11.6)	8 (8.8)	21 (5.7)	23 (6.2)
NRAS	6 (2.1)	2 (0.7)	1 (1.2)	0	7 (1.9)	3 (0.8)
MSI-H	1 (0.3)	2 (0.7)	8 (10.3)	8 (8.8)	9 (2.4)	11 (3.0)

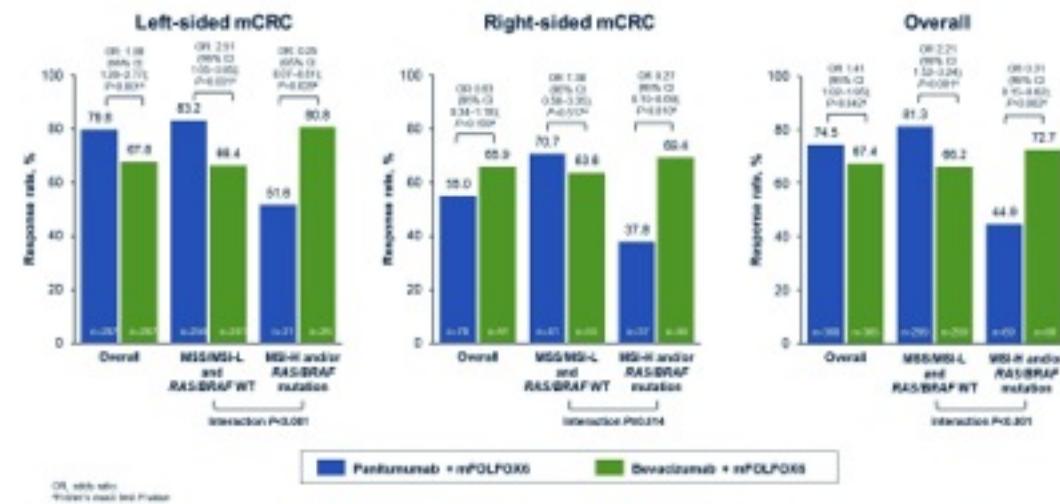
Overall survival by MSS/MSI and RAS/BRAF status in left-sided mCRC



Overall survival by MSS/MSI and RAS/BRAF status in right-sided mCRC

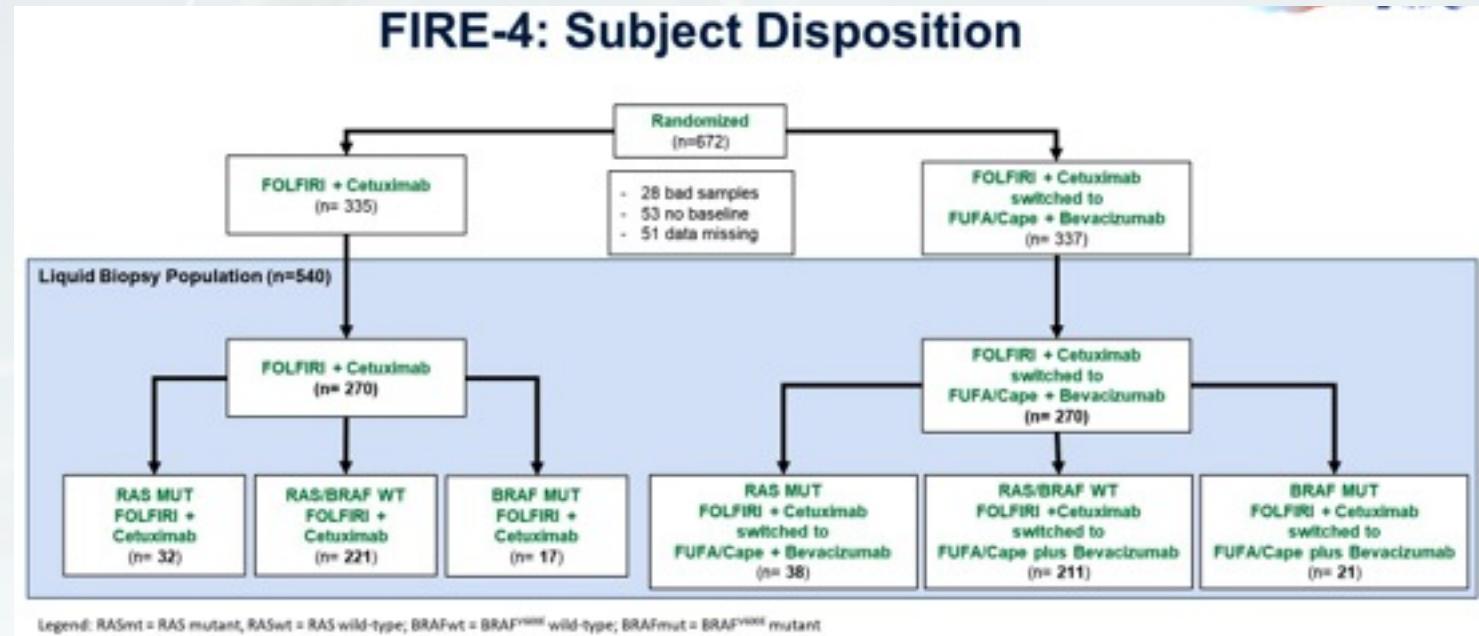
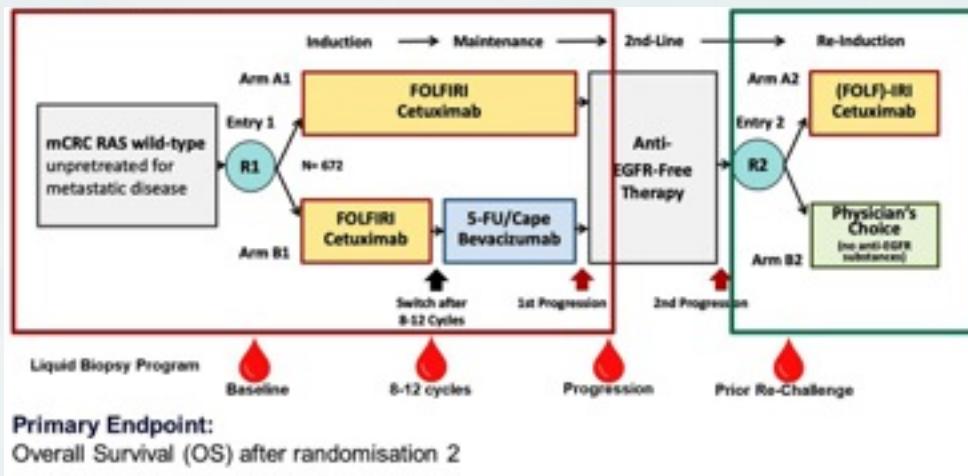


Response rate by MSS/MSI and RAS/BRAF status

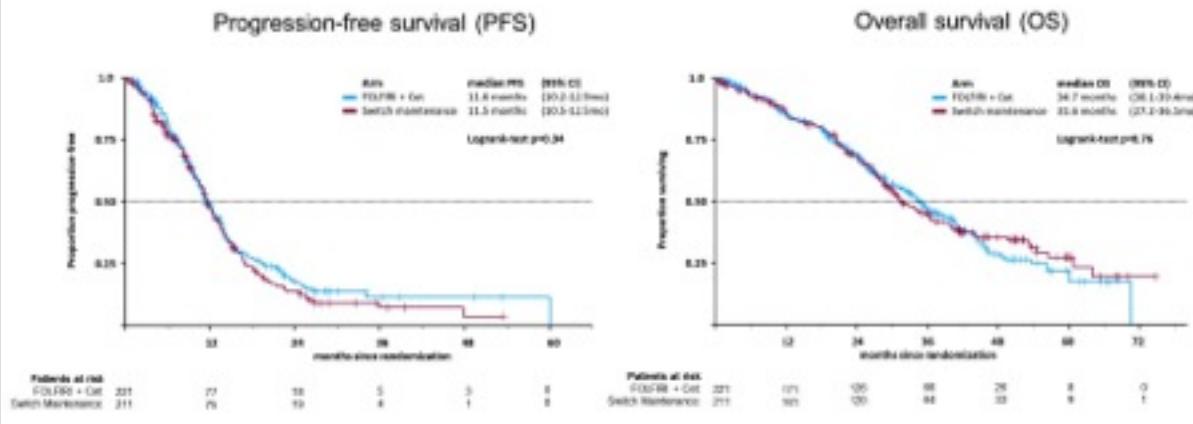


Influence of baseline liquid biopsy results on first-line treatment efficacy of FOLFIRI plus Cetuximab in patients with tissue RAS-WT mCRC

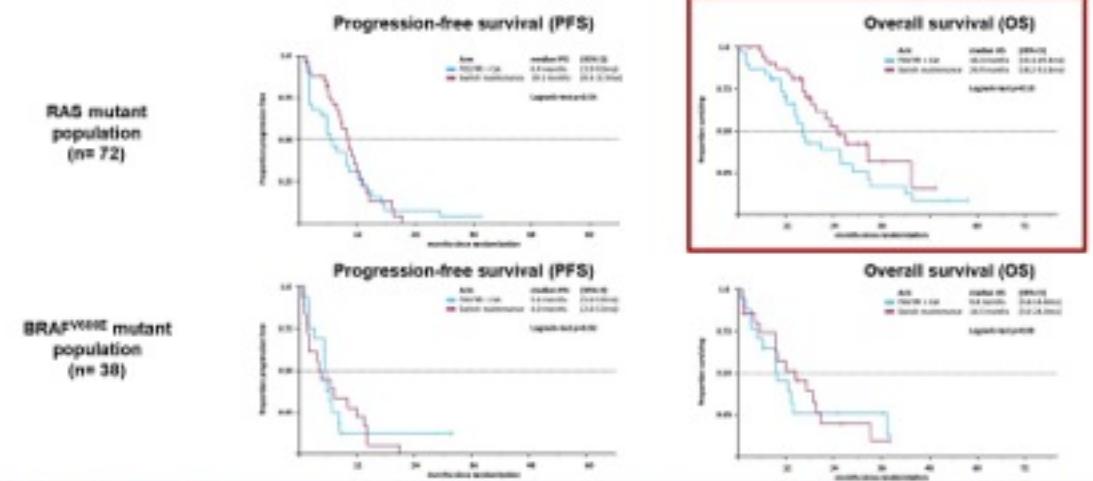
Data of the phase III FIRE-4 study (AIO KRK-0114):



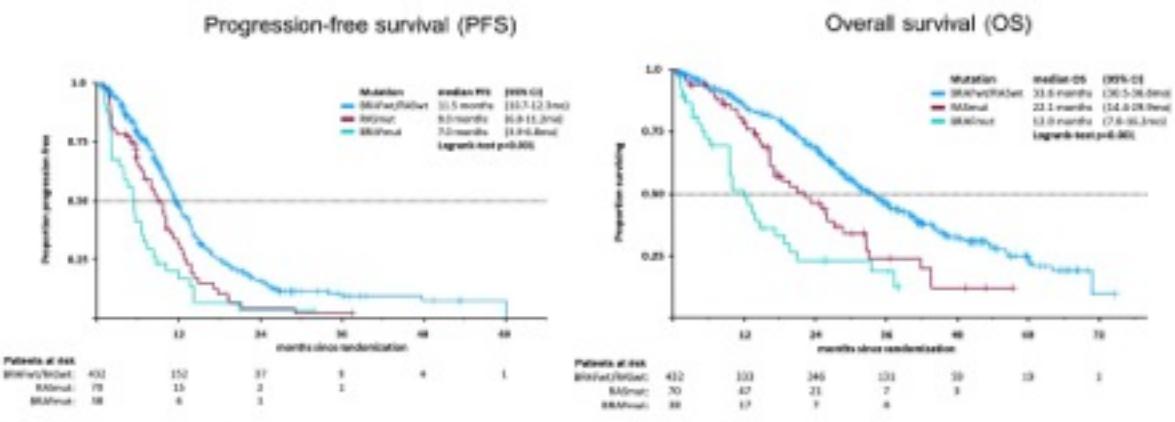
FIRE-4: Effect of treatment arm on survival RAS/BRAF wild-type population (n=432)



FIRE-4: Effect of treatment arm on survival



FIRE-4: Effect of baseline liquid biopsy result on survival



FIRE-4: Impact of emerging RAS mutations on survival

	Liquid Biopsy Baseline → Follow Up	PFS months (95%CI)	OS months (95%CI)
Arm A	RASwt → RASwt (n=112) RASwt → RAS mutant (n= 27)	10.4 (8.3 - 11.6) 15.1 (11.9-19.1) $P=0.04$	36.0 (28.7-41.0) 35.4 (32.5-44.1) $P=0.73$
Arm B	RASwt → RASwt (n= 142) RASwt → RAS mutant (n= 19)	12.5 (11.2-13.6) 13.8 (8.5 - 16.8) $P=0.47$	31.1 (28.5-36.4) 28.0 (20.6-39.8) $P=0.06$
All	RASwt → RASwt (n= 254) RASwt → RAS mutant (n= 46)	11.4 (10.5-12.5) 13.9 (11.9-16.6) $P=0.31$	32.7 (29.1-36.4) 33.6 (27.2-40.0) $P=0.22$

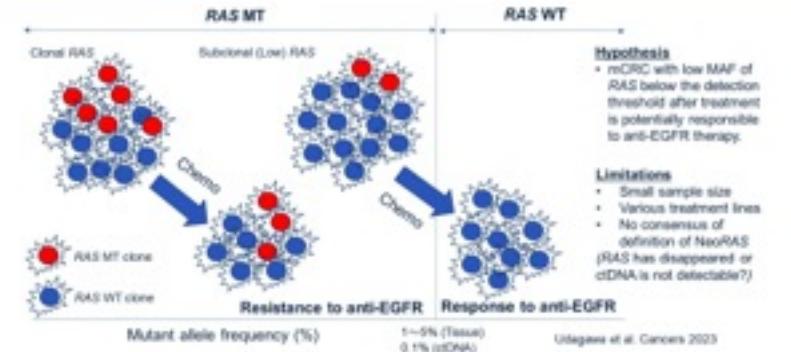
Arm A Continuous Cetuximab administration: 19.4%
Arm B Switch 5FU Bevacizumab maintenance: 11.8% $p=0.049$

Emerging RAS mutations:

- were associated with a significantly longer median Cetuximab exposure (27.6 weeks vs. 48.3 weeks (Logrank test $P=0.001$))
- were associated with a longer median PFS in the Cetuximab continuation arm (10.4 months vs. 15.1 months (Logrank test $P=0.04$))
- did not impact median OS

NeoRAS wild-type metastatic colorectal cancer in the SCRUM-Japan GOZILA study

The concept of NeoRAS WT mCRC



Patients and methods

- mCRC patients with tissue RAS MT (KRAS or NRAS exon 2, 3, 4 MT) prior to initiation of 1st line chemotherapy are eligible, regardless of treatment lines (n=647).



Taniguchi et al. Neoplasia 2016
Nakamura et al. Nat Med 2020

The incidence of NeoRAS WT mCRC depend on the RAS variants

Group A		Site of original KRAS MT in tissue (codon)						Site of original NRAS MT in tissue (codon)					
		Exon 2	Exon 3	Exon 4		Exon 2	Exon 3	Exon 4		Exon 2	Exon 3	Exon 4	
Plasma RAS MT	All	12	13	59	61	117	146		12	13	59	61	117
	Not detected	91	64	12	0	5	0	6	3	0	0	1	0
	Detected	387	278	66	0	11	4	11	5	3	0	9	0
NeoRAS (%)		19.0	18.7	15.4	0	31.3	0	35.3	37.5	0	0	10	0

- KRAS codon 12/13 (18.1%) vs other RAS MTs (25.1%). P = 0.21

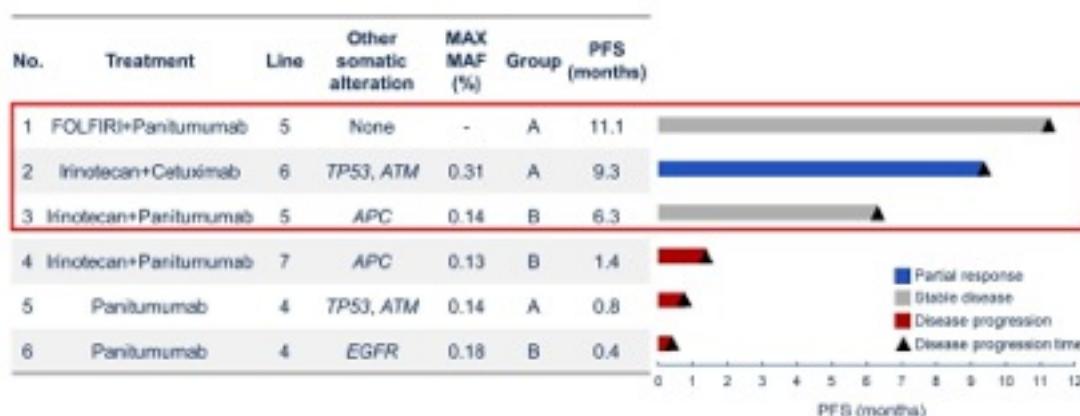
Group B		Site of original KRAS MT in tissue (codon)						Site of original NRAS MT in tissue (codon)					
		Exon 2	Exon 3	Exon 4		Exon 2	Exon 3	Exon 4		Exon 2	Exon 3	Exon 4	
Plasma RAS MT	All	12	13	59	61	117	146		12	13	59	61	117
	Not detected	42	27	7	0	2	0	4	1	0	0	1	0
	Detected	387	278	66	0	11	4	11	5	3	0	9	0
NeoRAS (%)		9.8	8.9	9.6	0	15.4	0	26.7	16.7	0	0	10	0

- KRAS codon 12/13 (9.0%) vs other RAS MTs (15.4%). P = 0.14

Logistic regression analysis for NeoRAS WT : Group A/B

*Ref	GroupA	Odds Ratio	95% CI Lower limit	95% CI Upper limit	P. value
	Liver metastasis (Negative* or Positive)	0.17	0.09	0.33	<0.00001
	Lung metastasis (Negative* or Positive)	0.75	0.39	1.44	0.39
	Lymph node metastasis (Negative* or Positive)	0.45	0.22	0.92	0.028
	Peritoneal metastasis (Negative* or Positive)	1.45	0.75	2.83	0.27
	Bone metastasis (Negative* or Positive)	0.28	0.06	1.29	0.10
	Multiple organ metastases (Negative* or Positive)	0.94	0.49	1.80	0.86
	Treatment lines (5th or Later* or 2nd, 3rd and 4th)	1.59	0.77	3.28	0.21
	Treatment (Without* or Regorafenib)	0.79	0.19	3.31	0.74
	Treatment (Without* or VEGF inhibitors)	1.72	0.85	3.48	0.13
	Tissue RAS MT (KRAS codon 12, 13* or Others)	1.82	0.90	3.70	0.098
	GroupB	Odds Ratio	95% CI Lower limit	95% CI Upper limit	P. value
	Liver metastasis (Negative* or Positive)	0.37	0.18	0.79	0.0097
	Lymph node metastasis (Negative* or Positive)	0.66	0.27	1.59	0.35
	Peritoneal metastasis (Negative* or Positive)	1.35	0.63	2.87	0.44
	Multiple organ metastases (Negative* or Positive)	0.66	0.30	1.45	0.30
	Treatment (Without* or VEGF inhibitors)	2.27	0.96	5.35	0.061
	Tissue RAS MT (KRAS codon 12, 13* or Others)	2.34	1.00	5.49	0.048

Clinical outcomes of anti-EGFR therapy for NeoRAS WT mCRC (n=6)



#3567: Reversion of RAS mutations in metastatic colorectal cancer in the CCTG CO.26 clinical trial

Authors: Florence T.H. Wu¹, James T. Topham¹, Chris J. O'Callaghan², Harriet Feilotter², Hagen F. Kennecke³, Kimberly C. Banks⁴, Daniel J. Renouf⁵, Derek J. Jonker⁶, Dongsheng Tu², Eric X. Chen⁹, Jonathan M. Loree¹

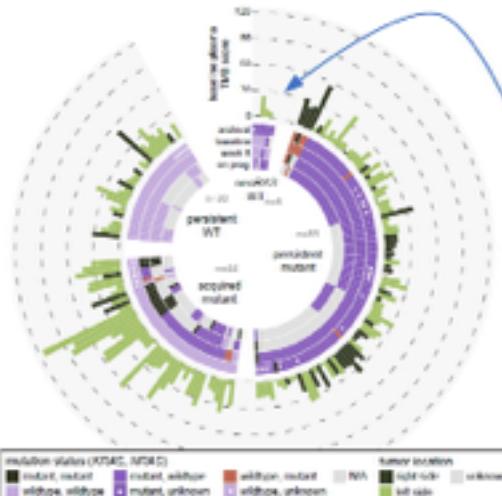
¹BC Cancer, University of British Columbia, Vancouver, BC, Canada. ²Canadian Clinical Trials Group, Kingston, ON, Canada. ³Virginia Mason Cancer Center, Seattle, WA, USA. ⁴Guardant Health, Redwood City, CA, USA. ⁵The Ottawa Hospital, University of Ottawa, Ottawa, ON, Canada. ⁶Princess Margaret Cancer Centre, Toronto, ON, Canada.

Background:

- KRAS/NRAS mutations in metastatic colorectal cancer (mCRC) drive resistance to anti-EGFR antibodies.
- It is unclear if RAS mutations are ever clonally lost, potentially uncovering a new therapeutic option for patients over time.

Methods:

- CCTG CO.26 was a phase II clinical trial that assessed durvalumab + tremelimumab in patients with heavily pretreated mCRC.
- For 169 patients enrolled on CO.26, initial tissue based KRAS and NRAS mutation status (wildtype vs. mutated) was determined by whole exome sequencing (WES) of archival tumor tissue or standard of care clinical assays.
- Patients were then divided into four categories (Fig. 1) based on the temporal evolution of their mCRC RAS mutation status, as determined by Guardant OMNI panel next-generation sequencing of circulating tumor DNA (ctDNA) from plasma collected at trial baseline, week 8, and on progression.



Results:

- Of the 95 cases that were RAS^{mut} at the time of cancer diagnosis, 6 cases (6.3%) became RAS^{wt} by ctDNA analysis at 'baseline' or 'week 8' of the CO.26 trial. This is comparable with the 1.6% to 8.8% rates of 'neo-RAS-wildtype' reversions previously reported [1].

Conclusions

- Serial ctDNA sampling is useful for detecting and delineating the duration of RAS mutation reversions in mCRC.
- In CO.26, mCRC RAS mutation reversions were rare (6.3%) and mostly transient (4.2%), likely reflecting clonal dynamics in action.

Future Directions:

- Is there a RAS-mutant MAF threshold below which the bulk of tumor burden can still reliably yield response to anti-EGFR therapy?
- Future studies should incorporate contemporaneous re-biopsy of tumor tissue to confirm ctDNA-detected RAS reversions.
- There are at least four Phase II trials underway (MoLiMoR, KAIROS, ConVertix, and CETIDYL) testing anti-EGFR therapy after detection of 'neo-RAS-wildtype' reversions [2].

References: [1] Henry et al., Kopetz, ASCO GI, abstract #180, Journal of Clinical Oncology 38, no. 4_suppl (February 01, 2020) 180-180. [2] Osumi et al. Eur J Cancer. 2021;153:86-95.

Canadian Cancer Trials Group



CCTG CO.26
NCT02870920



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Results:

- Reversions were often transient. Fig. 2A,B,E,F re-acquired tumour RAS-mutant status by 55 to 293 (median 162.5) days after baseline.
- Four cases showed continued strong presence of other clonal somatic mutations when 'neo-RAS-wildtype' reversion occurred (Fig. 2A,B,C,E), making these less likely to be false negatives from low-volume or non-secreting tumor burden.

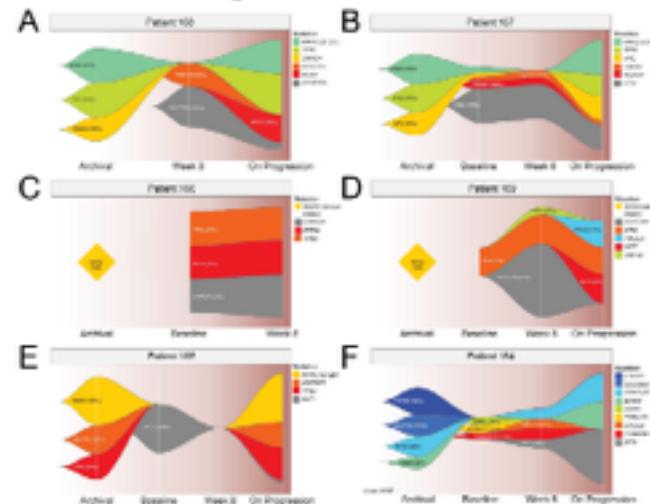


Figure 2. Clonal variant dynamics during neo-RAS-wildtype reversions.

Relative variant allele frequency (rVAF) = VAF of a given variant divided by the maximum VAF of any somatic variant in the same patient sample. N next to gene names indicate the rVAF of that variant at the first timepoint it was observed for that sample. Synchronous mutations are not shown in this figure but were used to calculate rVAF.

- 'Neo-RAS-wildtype' group had lower rates of synchronous metastases, but overall survival (OS) was not significantly different from overall population once they reach stage IV.

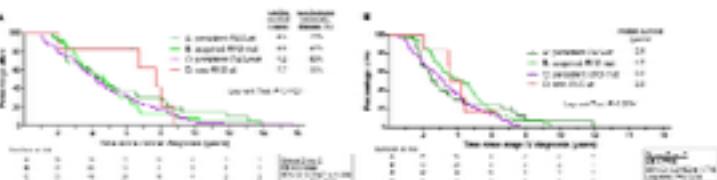


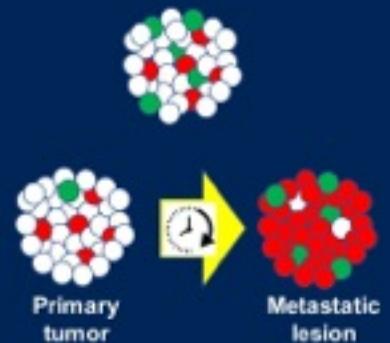
Figure 3. Kaplan-Meier analyses of OS.

Optimizing anti-EGFR therapy for metastatic colorectal cancer

- 1) Can ctDNA predict outcomes with 1st line chemotherapy + anti-EGFR?
- 2) Can a patient with a tissue RAS mutation lose that RAS mutation and benefit from anti-EGFR therapy?
- 3) ctDNA vs tissue NGS: What is the optimal approach?

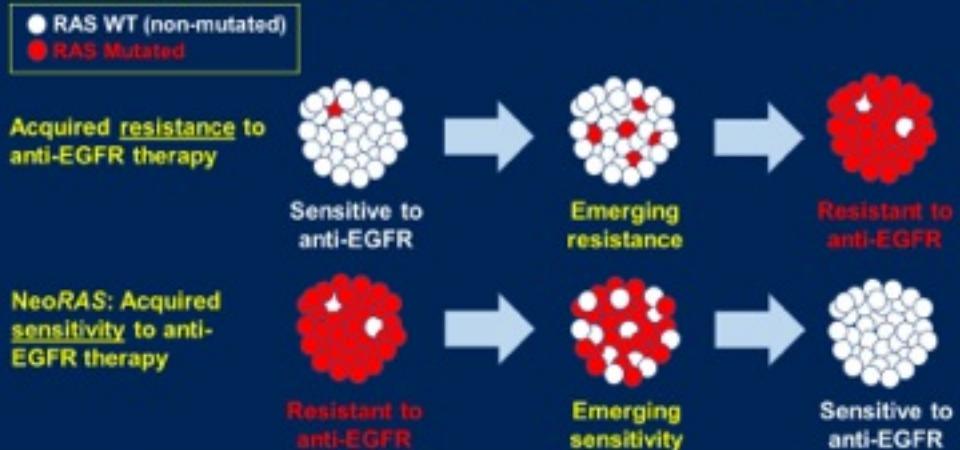
Potential explanations for plasma-tissue assay discordance

- Spatial heterogeneity of RAS/BRAF mutations in tumor tissue
- Temporal heterogeneity: RAS/BRAF mutations emerged in metastatic lesions after surgical resection of the primary
- Assay sensitivity



- What is the optimal absolute/ relative allele frequency for a subclonal RAS/BRAF alteration to predict resistance to anti-EGFR therapy?

"NeoRAS" = Loss of driver RAS mutations



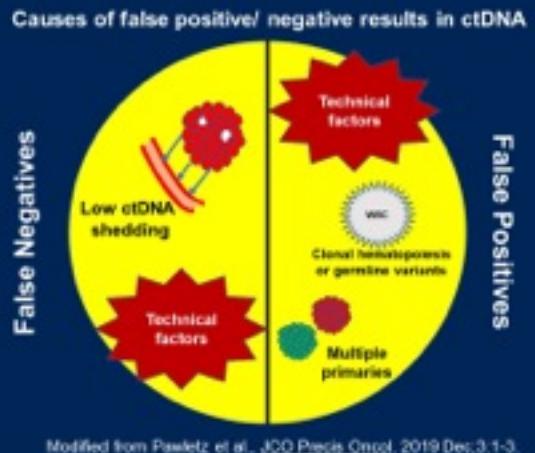
- How to distinguish between NeoRAS WT vs "RAS mutation not detected" in plasma?
- Anti-EGFR therapy vs SOC?
 - Trial participation is encouraged
 - Outside of trial, consider a tissue biopsy to confirm loss of the RAS mutation before treating with anti-EGFR

Clinical trials evaluating anti-EGFR therapy for NeoRAS WT metastatic colorectal cancer

Study	Location	ctDNA Assay	Treatment line	Study design (N)	Description
MolMiR (NCT04554836)	Germany	ddPCR	1L	Randomized phase II (N=144)	Active, not recruiting
CONVERTIX (NCT04100240-001)	Spain	SEANing	2L	Single-arm phase II (N=40)	Closed: No patients met eligibility
CETIDYL (NCT04189055)	France	Real-time PCR	≥ 3L	Single-arm phase II (N=72)	Recruiting
KAIROS (NCT04100240-001)	Italy	Real-time PCR	2L	Single-arm phase II (N=112)	Recruiting
C-PROWESS (NCT03121058)	Japan	ddPCR	≥ 3L	Single-arm phase II (N=112)	Recruiting

Potential explanations for NeoRAS WT phenomenon

- **False negatives**
 - Low ctDNA shedding
 - Other technical factors
- **Spatial heterogeneity** of RAS mutations in tumor tissue
- **Temporal heterogeneity** RAS wild-type disease has a fitness advantage(?)



Modified from Pawletz et al., JCO Precis Oncol. 2019 Dec;3:1-3.

Liquid vs tissue biopsy:

- Tissue NGS and ctDNA are complementary
- Both approaches have strengths/ limitations
- Adding ctDNA to tissue testing may capture clinically relevant heterogeneity
- Choice of assay should be tailored to clinical circumstances

Does serial circulating tumor DNA (ctDNA) monitoring identify additional acquired actionable alterations in metastatic colorectal cancer (mCRC)

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Introduction

Traditionally, biomarker ascertainment occurred once for patients with mCRC. Assessment of biomarkers by liquid biopsy enables serial monitoring of disease progression and acquired resistance for patients with mCRC without repeated tissue biopsy.

Here we explore the changes in genomic profiles as detected by serial liquid biopsies to identify actionable alterations and their clonal architecture.

Methods

Detected variants from serial Guardant360 ctDNA assays for 3350 patients with mCRC with ≥1 detected somatic alterations were used to assess variant gain or loss across timepoints.

Variant changes were normalized by patient-specific limits of detection (LoD), established by the 0.1% MAF threshold of the serial assay with the lowest max somatic mutant allele frequency (MAF) per patient to control for changes in total ctDNA concentrations at each time point¹.

Relative MAF (rMAF) was defined as the variant MAF divided by the max detected MAF in that sample. Variants were defined as clonal if detected at somatic rMAF ≥ 10%.

MSI scores and status ("MSI High", "MSI Low") was evaluable for 3,030 patients with mCRC across multiple assays. Blood TMB (bTMB) score was evaluable for 1,387 patients across assays. Patients were selected for "MSI-low" and "bTMB-Low" status at their initial Guardant360 assay and "MSI-High" and "bTMB-High" at subsequent assays.

We also assessed whether the presence of DNA damage repair pathway mutations at an initial timepoint predisposed patients to having greater changes in MSI score and bTMB score compared to mutation negative patients. Variants evaluated as DNA damage repair pathway mutations included: BRCA1/2, ATM, CHEK2, MLH1, RAD51D.

Serial monitoring reveals emergence of anti-EGFR resistance and secondary therapeutic mutations for initially MAPK mutation negative mCRC patients

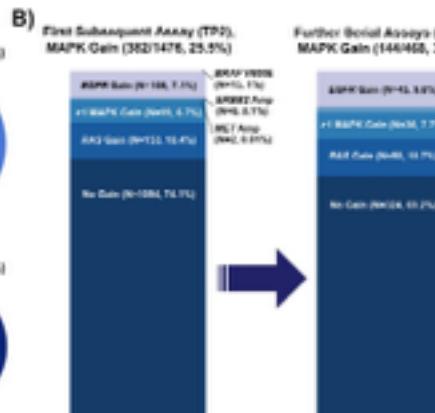
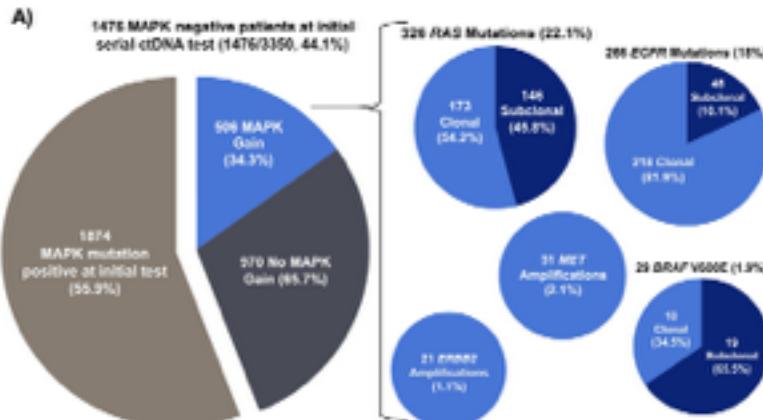


Figure 1. Patient level mutation gain across serial liquid biopsy testing time points. **A)** Of 3350 screened patients, 1476 were negative for MAPK clonal and subclonal mutations at initial assay. Mutations developed in 506 (25.6%) patients, with 382 gains (76.5%) occurring at the first subsequent test. MAPK gains across genes were more often clonal (including amplifications) than subclonal (453:213; 68%:32%). **B)** Subsequent assays demonstrate similar rate of MAPK mutation development beyond the first serial test.

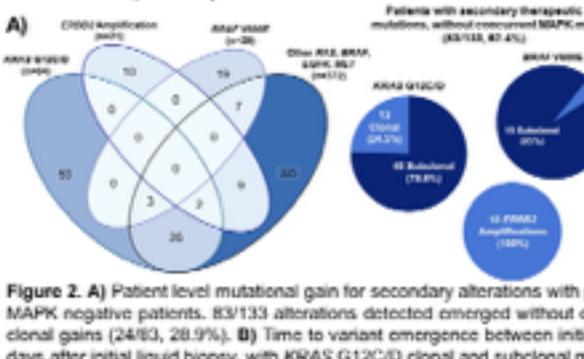
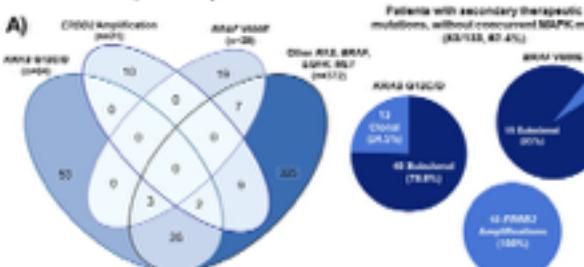


Figure 2. **A)** Patient level mutational gain for secondary alterations with potential therapeutic relevance (BRAF V600E, KRAS G12C/D, ERBB2 amp) in initially MAPK negative patients. 83/133 alterations detected emerged without concurrent MAPK mutation. Subclonal gains (59/83, 71.1%) were more common than clonal gains (24/83, 28.9%). **B)** Time to variant emergence between initial assay and subsequent serial test. Therapeutic mutations emerged a median of 372 days after initial liquid biopsy, with KRAS G12C/D clonal and subclonal mutations detected in assays significantly longer than assays with no gains.

Results

Impact of DNA repair pathway alterations (BRCA1/2, ATM, CHEK2, MLH1, RAD51D) on acquisition of MSI and blood TMB (bTMB) ≥20M/mb at a future time point for patients without these alterations at baseline.

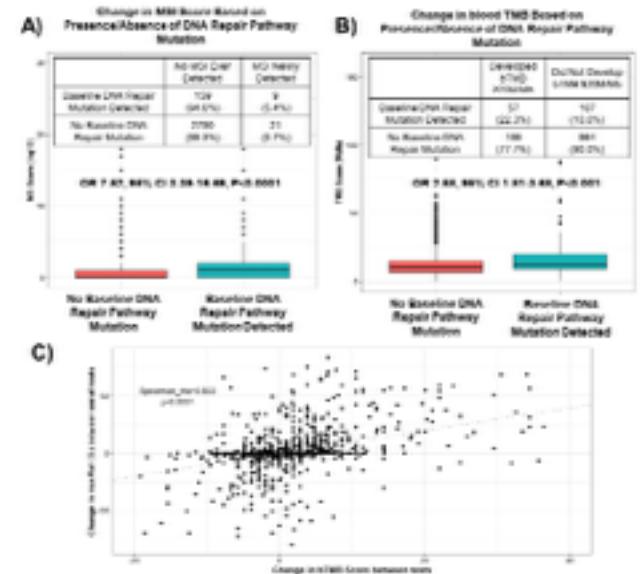


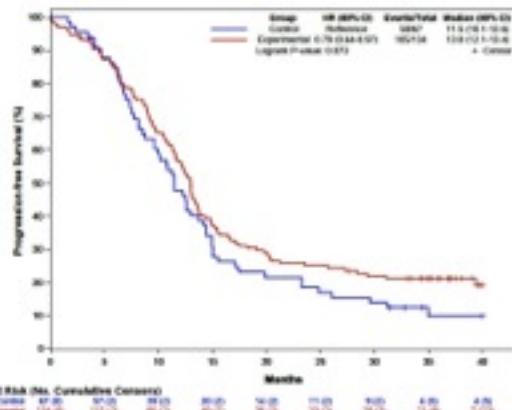
Figure 3. Changes in **A)** MSI scores and **B)** bTMB were more common for patients with DNA repair pathway mutations on their initial assay. We also noted that **C)** increase in bTMB is significantly correlated with increasing max somatic MAF ($p=0.504$, $p<0.0001$).

Conclusions

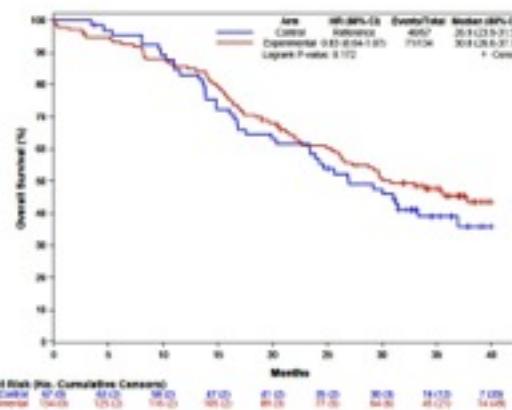
- In this large cohort, serial ctDNA appears feasible to identify acquired alterations. Many of these alterations have therapeutic implications and further work is needed to understand if and how to target acquired alterations.

FOLFOXIRI plus bevacizumab and atezolizumab as upfront treatment of unresectable mCRC patients: updated and overall survival results of the phase II randomized AtezoTRIBE study

Updated PFS – pMMR cohort

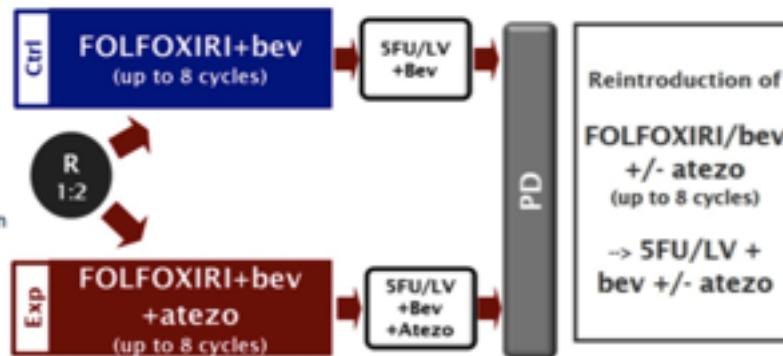


Overall Survival – pMMR cohort

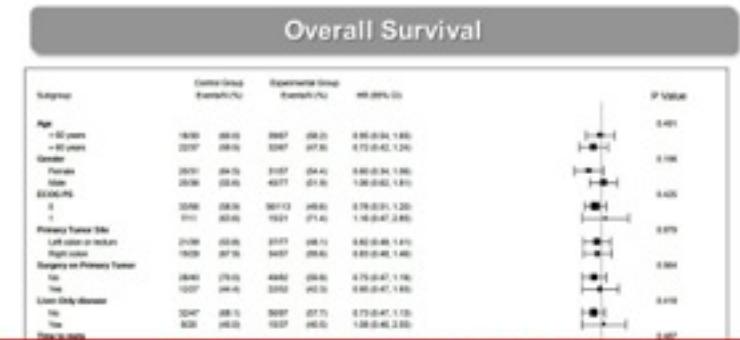
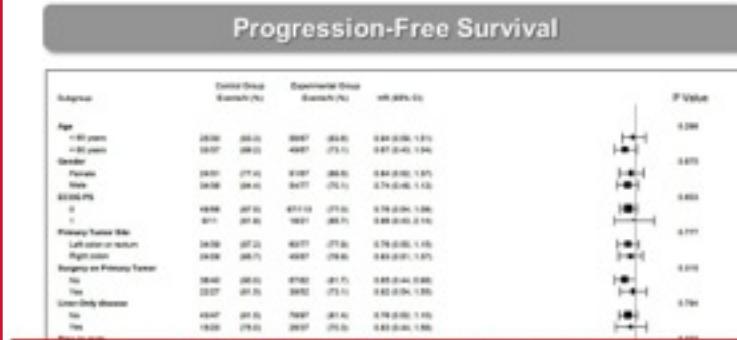


Key eligibility criteria

- Previously untreated, unresectable and RECIST v1.1-measurable mCRC
- Age 18-75 years
- ECOG PS ≤ 2 (ECOG PS= 0 if age= 71-75 years)
- Adjuvant oxaliplatin-containing chemotherapy not allowed
- Adjuvant fluoropyrimidine monotherapy allowed if more than 6 months elapsed between the end of adjuvant and first relapse
- Adequate bone marrow, liver and renal functions
- No contraindications to ICI



Subgroup analyses for PFS and OS – pMMR cohort



Multivariate Cox proportional hazard regression model

- TMB: P value= 0.054
- Immunoscore IC: P value= 0.060



Multivariate Cox proportional hazard regression model

- TMB: P value= 0.033
- Immunoscore IC: P value= 0.022



Federica Marmorino, Gianmarco Piccinno, Daniele Rossini, Filippo Ghelardi, Sabina Murgioni, Lisa Salvatore, Vincenzo Nasca, Carlotta Antoniotti, Francesca Daniel, Francesco Schietroma, Veronica Conca, Carolina Alves Costa Silva, Emiliano Tamburini, Stefano Tamberi, Alessandro Passardi, Lorenzo Antonuzzo, Raffaella D'Onofrio, Laurence Zitvogel, Chiara Cremolini and Lisa Derosa

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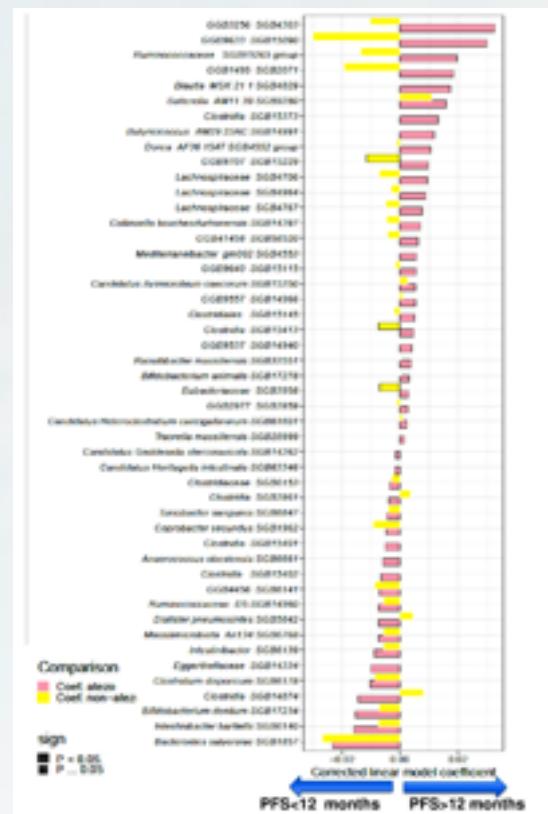


Figure 3. KM for PFS according to treatment arm and *Ruminococcaceae unclass* prevalence

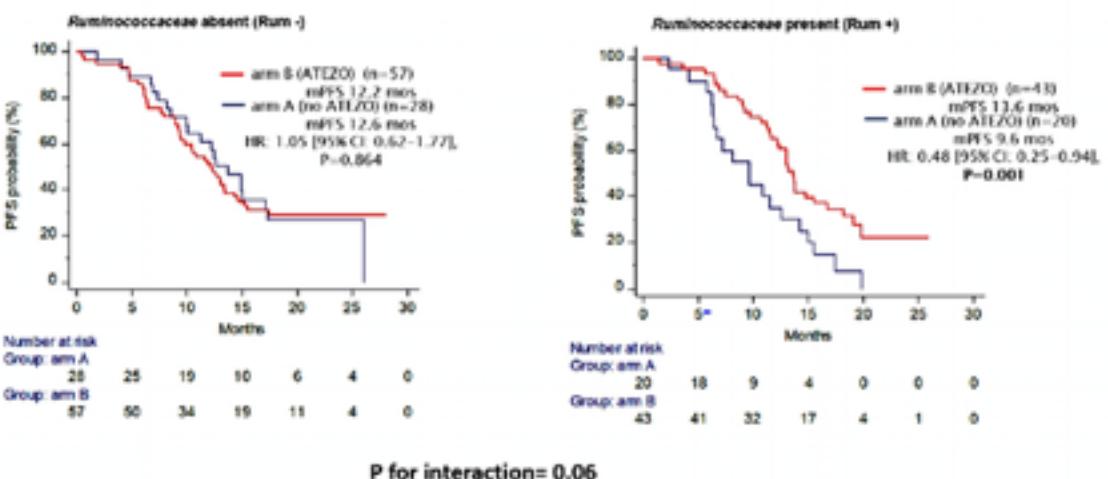
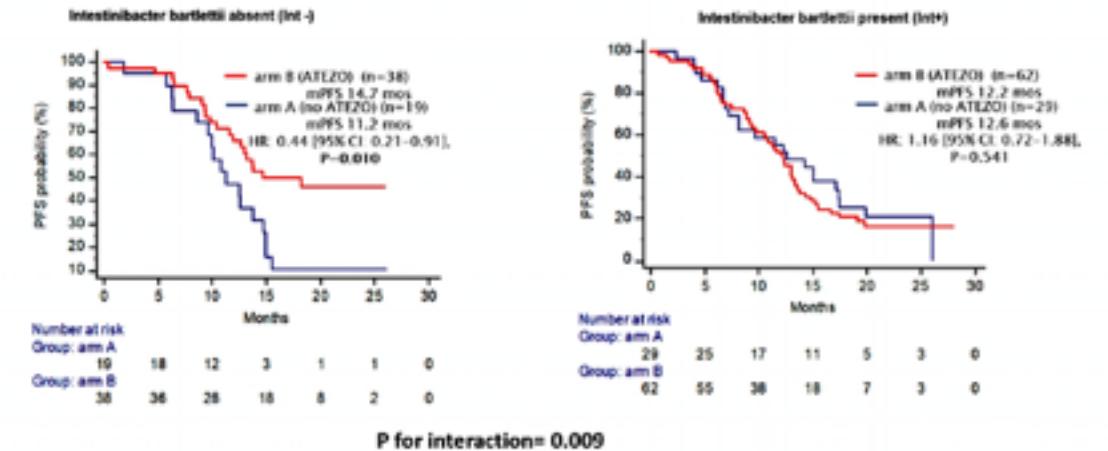


Figure 4. KM for PFS according to treatment arm and *Intestinibacter bartlettii* prevalence



First-line nivolumab + ipilimumab in patients with microsatellite instability-high/mismatch repair-deficient metastatic colorectal cancer: 64-month follow-up from CheckMate 142

Umar Iqbal, 1 Michael J. Quezada, 2 Eric Van Cutsem, 3 Maria-José Llorente, 4 Koen Van Oosterom, 5 Michael Herberman, 6 Michael S. Giltnane, 7 Diane Gosselin, 8 Georges A. Rizk, 9 Hassan 10 Fabio Colombo, 11

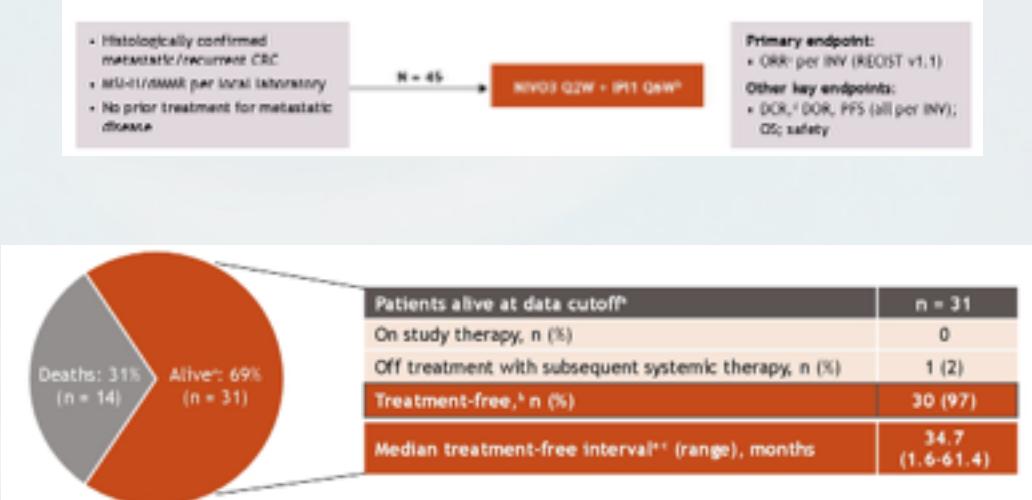
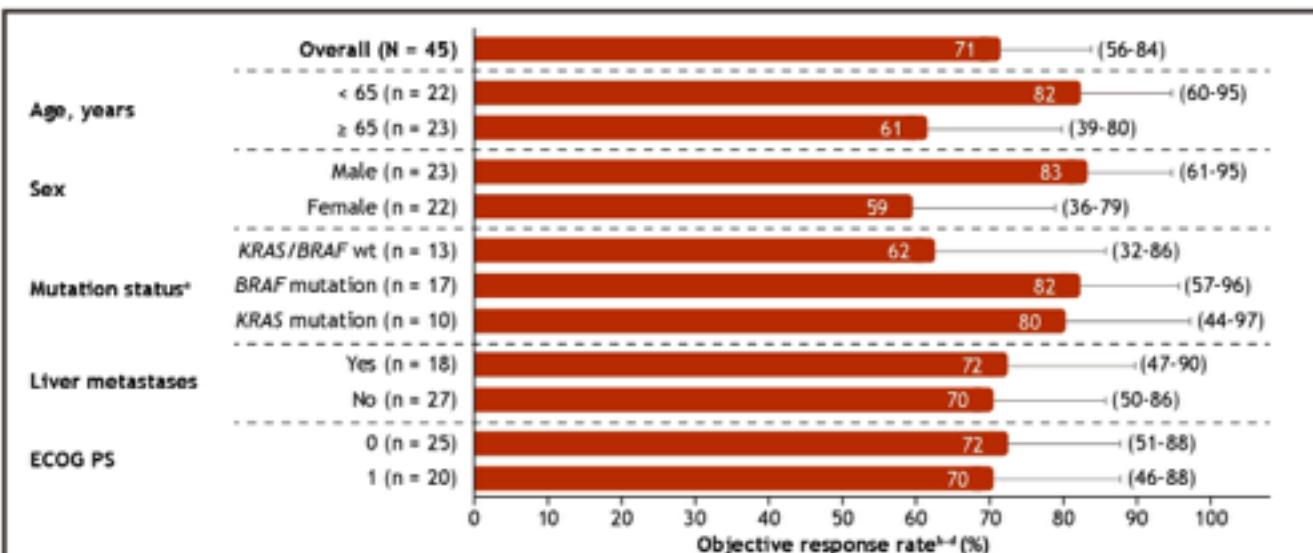


Figure 2. Objective response rate by subgroup



^aExcludes 5 patients with unknown mutation status; ^bPer Investigator; ^cPatients with CR + PR divided by the number of treated patients; ^dError bars and numbers in parentheses indicate 95% CIs; evaluated subgroups had overlapping 95% CIs for ORR; wt, wild-type.

- At 60 months, PFS and OS rates were 55% and 67%, respectively

Exploratory biomarker analysis

- In TMB-evaluable patients (n = 20; Figure 7):
 - ORR was 50% in patients with TMB-low tumors and 70% in those with TMB-high tumors
 - Median PFS was 17.4 months in patients with TMB-low tumors and not reached in those with TMB-high tumors
 - Median OS was 28.8 months in patients with TMB-low tumors and not reached in those with TMB-high tumors

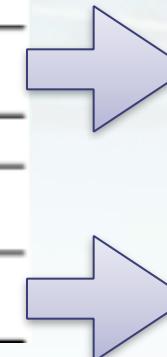
Molecular Targets in Colorectal Cancer

Moving Beyond EGFR, BRAF, and dMMR

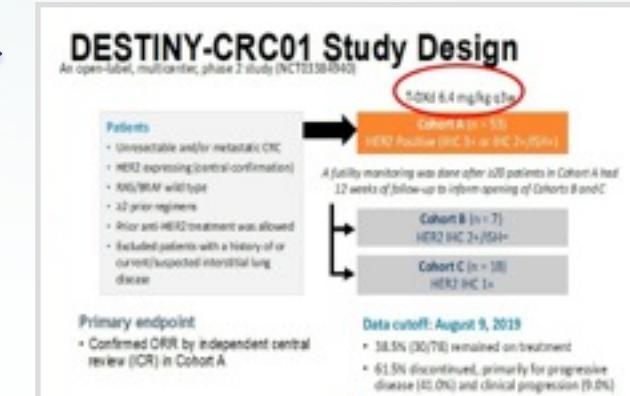
- Targets:
 - HER2 amplification
 - KRAS G12C mutation
 - NTRK fusion
 - RET fusion

Anti-HER2 strategies in mCRC

	Treatment	n	Prior Lines of Treatment	Mutational Status	mPFS (m)	ORR (%)
MyPathway	Trastuzumab + pertuzumab	57	≥1	RAS WT	2.9	32
TAPUR	Trastuzumab + pertuzumab	28	≥0	NS	NR	14
TRIUMPH	Trastuzumab + pertuzumab	27 (Tissue) 25 (ctDNA)	≥1	RAS WT	4.0 3.1	30 25
Monoclonal antibody + TKI						
HERACLES-A	Trastuzumab + lapatinib	35	≥2	KRAS WT	4.7	28
Yuan et al.	Trastuzumab + pyrotinib	11	≥2	RAS WT and mutated	NR	27
MOUNTAINEER	Trastuzumab + tucatinib	23	≥2	RAS WT	8.1	52
ADCs						
HERACLES-B	Pertuzumab + T-DM1	31	≥2	RAS/BRAF WT	4.1	10
DESTINY-CRC01	TD	53 (Cohort A)	≥2	RAS/BRAF WT	6.9	45



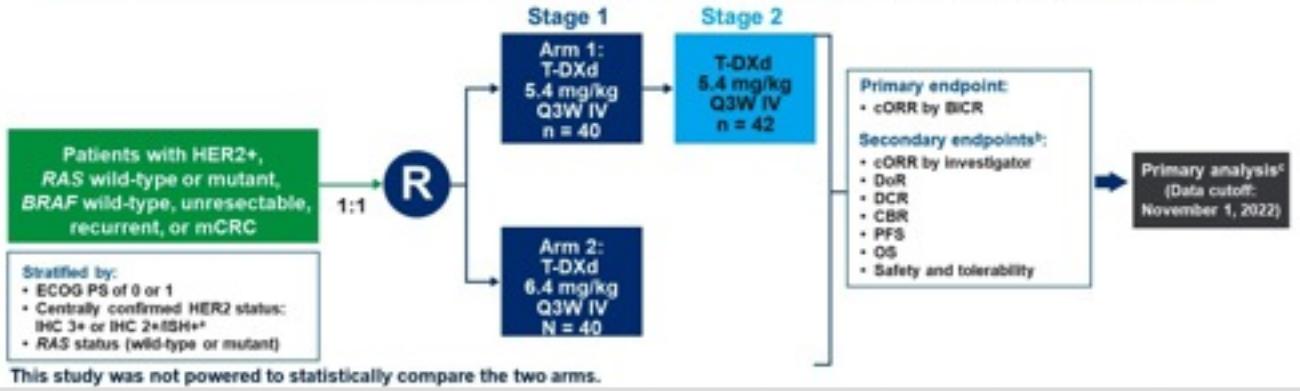
FDA Approval Jan 2023



T-DXd in Patients With HER2-Overexpressing/Amplified (HER2+) Metastatic Colorectal Cancer (mCRC): Primary Results From the Multicenter, Randomized, Phase 2 DESTINY-CRC02 Study

A randomized, blinded, 2-stage, 2-arm, multicenter, global, phase 2 study (NCT04744831)

- Stage 1 (randomized) was followed by Stage 2 (nonrandomized), which enrolled an additional 42 patients



K. Raghav. ASCO 2023 # 3501

Baseline Characteristics

	T-DXd 5.4 mg/kg Q3W		T-DXd 6.4 mg/kg Q3W	
	Stage 1 n = 40	Stage 2 n = 42	Total N = 82	Stage 1 N = 40
Median age, years (range)	58.2 (26-78)	60.6 (30-84)	59.1 (26-84)	62.3 (35-81)
Sex, n (%)				
Male	21 (52.5)	24 (57.1)	45 (54.9)	19 (47.5)
Region, n (%)				
Asia-Pacific	25 (62.5)	22 (52.4)	47 (57.3)	24 (60.0)
US	5 (12.5)	1 (2.4)	6 (7.3)	2 (5.0)
Europe	10 (25.0)	19 (45.2)	29 (35.4)	14 (35.0)
HER2 status, n (%)				
IHC 3+	32 (80.0)	32 (76.2)	64 (78.0)	34 (85.0)
IHC 2+ESH+	8 (20.0)	10 (23.8)	18 (22.0)	6 (15.0)
ECOG PS, n (%)				
0	22 (55.0)	24 (57.1)	46 (56.1)	22 (55.0)
1	18 (45.0)	18 (42.9)	36 (43.9)	18 (45.0)
RAS status, n (%)				
Wild-type	34 (85.0)	34 (81.0)	68 (82.9)	34 (85.0)
Mutant	6 (15.0)	8 (19.0)	14 (17.1)	6 (15.0)
HER2/RAS status, n (%)				
IHC 2+ESH+/wild-type	7 (17.5)	5 (11.9)	12 (14.6)	6 (15.0)
IHC 2+ESH+/mutant	1 (2.5)	5 (11.9)	6 (7.3)	0
IHC 3+/wild-type	27 (67.5)	29 (69.0)	56 (68.3)	28 (70.0)
IHC 3+/mutant	5 (12.5)	3 (7.1)	8 (9.8)	6 (15.0)
Liver metastases at baseline, n (%)	29 (72.5)	30 (71.4)	59 (72.0)	26 (65.0)
CNS metastases at baseline, n (%)	3 (7.5)	0	3 (3.7)	1 (2.5)
Primary tumor site, n (%)				
Left colon ^a	32 (80.0)	29 (69.0)	61 (74.4)	34 (85.0)
Rectum	15 (37.5)	12 (28.6)	27 (32.9)	19 (47.5)
Right colon ^b	8 (20.0)	13 (31.0)	21 (25.6)	6 (15.0)

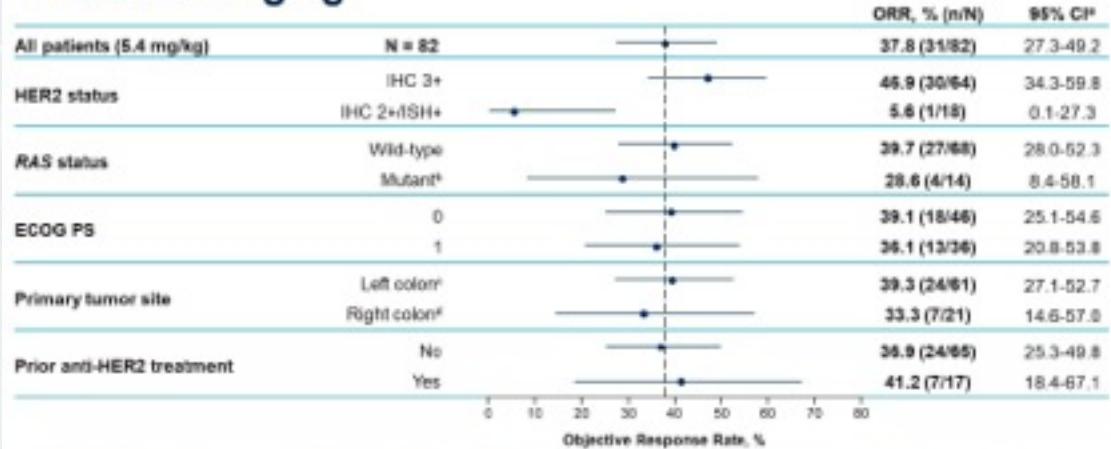
Prior Treatment

	T-DXd 5.4 mg/kg Q3W		T-DXd 6.4 mg/kg Q3W	
	Stage 1 n = 40	Stage 2 n = 42	Total N = 82	Stage 1 N = 40
Median prior lines of systemic therapy, n (range)	4 (1-12)	3 (1-7)	3 (1-12)	4 (1-8)
Systemic chemotherapy, n (%)				
Irinotecan	39 (97.5)	40 (95.2)	79 (96.3)	40 (100)
Fluoropyrimidines ^c	40 (100)	42 (100)	82 (100)	40 (100)
Oxaliplatin	40 (100)	41 (97.6)	81 (98.8)	40 (100)
Anti-EGFR, n (%)	29 (72.5)	28 (66.7)	57 (69.5)	31 (77.5)
Anti-HER2, n (%)				
HER2 TKP	11 (27.5)	6 (14.3)	17 (20.7)	10 (25.0)
Anti-HER2 antibodies	6 (15.0)	4 (9.5)	10 (12.2)	7 (17.5)
Anti-VEGF, n (%)	36 (90.0)	38 (90.5)	74 (90.2)	38 (95.0)
Regorafenib and tipiracitibrium, n (%)	20 (50.0)	14 (33.3)	34 (41.5)	13 (32.5)
Other systemic therapy, n (%)	5 (12.5)	6 (14.3)	11 (13.4)	10 (25.0)

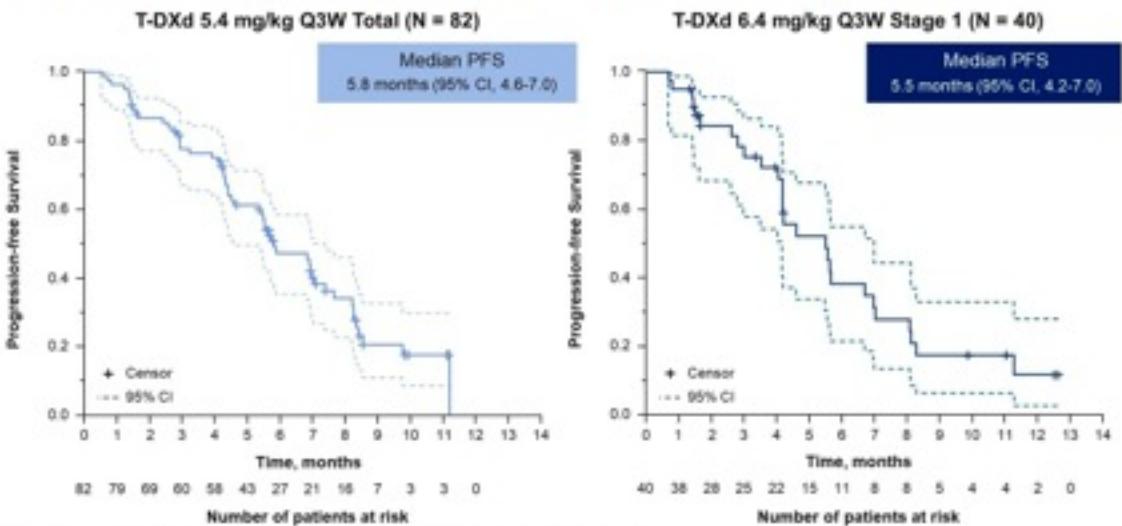
Efficacy Results

	T-DXd 5.4 mg/kg Q3W		T-DXd 6.4 mg/kg Q3W	
	Stage 1 n = 40	Stage 2 n = 42	Total N = 82	Stage 1 N = 40
cORR, n (%) [95% CI]	18 (45.0) [29.3-61.5]	13 (31.0) [17.6-47.1]	31 (37.8) [27.3-49.2]	11 (27.5) [14.6-43.9]
CR	0	0	0	0
PR	18 (45.0)	13 (31.0)	31 (37.8)	11 (27.5)
SD	20 (50.0)	20 (47.6)	40 (48.8)	23 (57.5)
PD	2 (5.0)	6 (14.3)	8 (9.8)	4 (10.0)
NE	0	3 (7.1)	3 (3.7)	2 (5.0)
Confirmed DCR, n (%) [95% CI]	38 (95.0) [83.1-99.4]	33 (78.6) [63.2-89.7]	71 (86.6) [77.3-93.1]	34 (85.0) [70.2-94.3]
Median DoR, mo (95% CI)	8.1 (4.2-NE)	4.6 (4.1-7.0)	5.5 (4.2-8.1)	5.5 (3.7-NE)

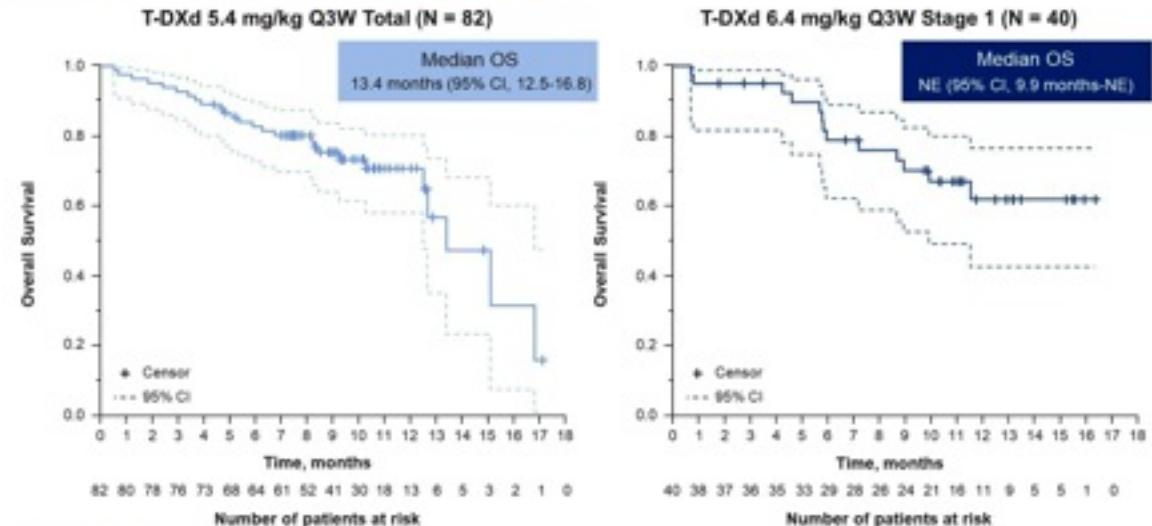
Best Overall Response by BICR by Subgroup With T-DXd 5.4 mg/kg



Median Progression-Free Survival by BICR



Median Overall Survival



Overall Safety Summary

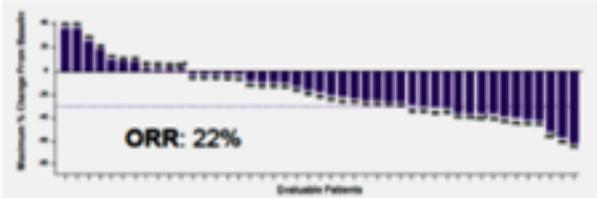
n (%)	T-DXd 5.4 mg/kg Q3W			T-DXd 6.4 mg/kg Q3W
	Stage 1 n = 41 ^a	Stage 2 n = 42	Total N = 83	Stage 1 N = 39
TEAEs	40 (97.6)	42 (100)	82 (98.8)	39 (100)
Drug-related	38 (92.7)	38 (90.5)	76 (91.6)	37 (94.9)
TEAEs grade ≥3	20 (48.8)	21 (50.0)	41 (49.4)	23 (59.0)
Drug-related	16 (39.0)	18 (42.9)	34 (41.0)	19 (48.7)
Serious TEAEs	8 (19.5)	12 (28.6)	20 (24.1)	12 (30.8)
Drug-related	4 (9.8)	7 (16.7)	11 (13.3)	6 (15.4)

Adjudicated Drug-Related ILD/Pneumonitis by Independent Adjudication Committee

Adjudicated as drug-related ILD/pneumonitis, n (%)	T-DXd 5.4 mg/kg Q3W			T-DXd 6.4 mg/kg Q3W
	Stage 1 n = 41 ^a	Stage 2 n = 42	Total N = 83	Stage 1 N = 39
Any grade	4 (9.8)	3 (7.1)	7 (8.4)	5 (12.8)

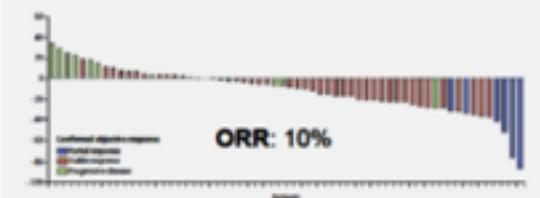
KRAS G12C inhibitors in mCRC

Adagrasib



Weiss et al, ESMO 2021

Sotorasib



Fakih et al, Lancet Oncol 2022

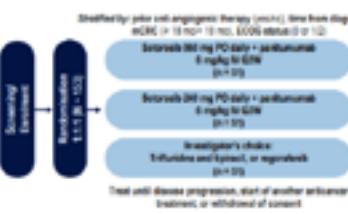
CodeBreak 200: Phase 2 Multicentre, Randomized, Active-controlled Study of Sotorasib + Panitumumab in KRAS G12C-mutant⁴

Key inclusion criteria:

- ≥ 18 years of age
- KRAS G12C-mutant mCRC, identified through next-generation sequencing of tumor biopsy
- ≤ 1 prior line of therapy for mCRC, progressed or after first-line cytotoxic, radiosens, and immunotherapy
- ECOG 0-1
- Measurable disease per RECIST 1.1

Key exclusion criteria:

- Radiotherapy within 2 weeks, palliative therapy within 4 weeks of study
- Prior treatment with EGFR monoclonal antibody or anti-EGFR monoclonal antibody and targeted agents, unless the investigator deems these agents

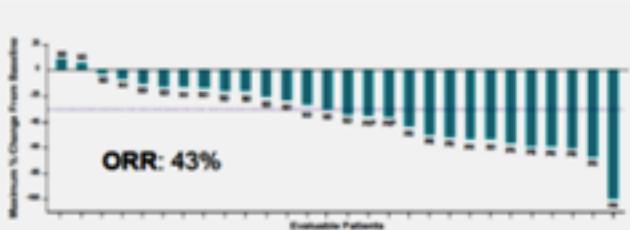


- Primary endpoint:**
- PFS per BICR measured by CTCAE and assessed by RECIST v1.1

- Secondary endpoints:**
- OS, ORR, DCR, TTR
- Safety/tolerability
- PROs, QoL
- PK

KRAS G12C inhibitors + anti-EGFR in mCRC

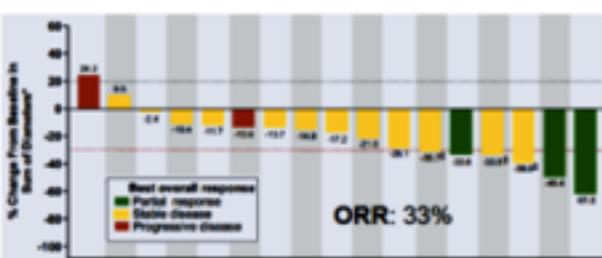
Adagrasib + Cetuximab



Updated ORR (n=28): 46%

Weiss et al, ESMO 2021
Klempner et al, ESMO 2022

Sotorasib + Panitumumab



Updated ORR (n=40): 30%

Fakih et al, ESMO 2021
Kuboki et al, ESMO 2022

KRYSTAL-10: Phase 3 Study of Adagrasib + Cetuximab vs Chemotherapy in KRAS G12C-mutant CRCm (201)²

Patients with previously treated advanced CRC harboring KRAS^{G12C} mutations (n=420)

Adagrasib^a 600 mg BID + cetuximab^b 500 mg/m² Q2W

Chemotherapy (FOLFIRI or mFOLFOX6)^c

KRYSTAL-1

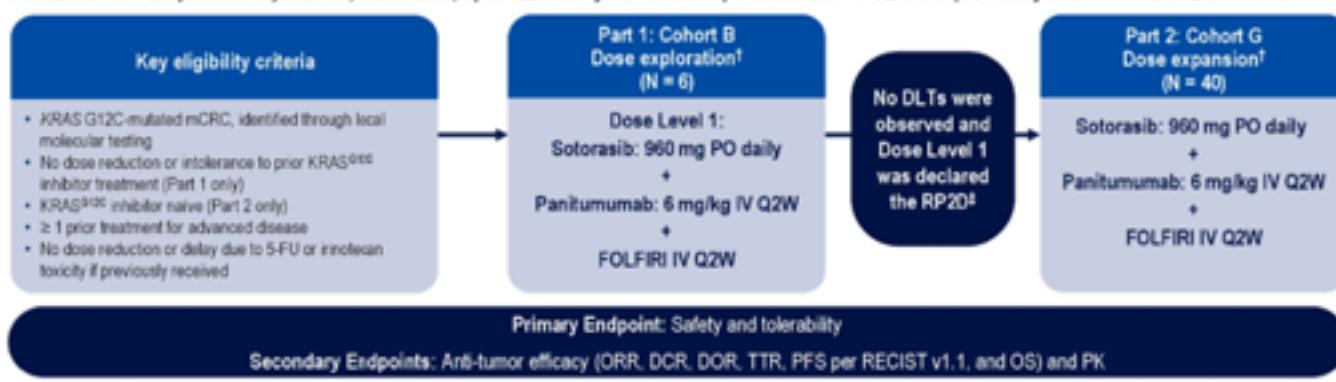
CodeBreak 101

Sotorasib (Soto) Plus Panitumumab (Pmab) and FOLFIRI for Previously Treated KRAS G12C-mutated Metastatic Colorectal Cancer (mCRC): CodeBreak 101 Phase 1b Safety and Efficacy

David S. Hong,¹ Yasutoshi Kuboki,² John H. Strickler,³ Marwan Fakih,⁴ Hélène Houssiau,⁵ Timothy J. Price,⁶ Elena Élez,⁷ Salvatore Siena,⁸ Emily Chan,⁹ Jane Hippenmeyer,⁹ Panli Cardona,⁹ Qui Tran,⁹ Toshiki Masuishi¹⁰

STUDY SCHEMA

CodeBreak 101 Subprotocol H phase 1b, multicenter, open-label study^a: sotorasib + panitumumab + FOLFIRI in previously treated KRAS G12C-mutated mCRC



- All patients received prior fluoropyrimidine, and 70% of patients received prior irinotecan
- Patients received a median of 2 prior lines of therapy with approximately one-third each receiving the study regimen as second-line, third-line or ≥ fourth-line therapy

- No clinically meaningful PK interaction was observed between sotorasib and irinotecan
- No DLTs were observed in dose exploration and sotorasib 960 mg daily, panitumumab 6 mg/kg IV Q2W and FOLFIRI IV Q2W was determined as the RP2D
- Safety findings were consistent with known profiles of sotorasib, panitumumab, and FOLFIRI

- Confirmed ORR (all partial responses) was 55% (95% CI: 38.7, 70.2) and DCR was 93% (95%CI: 80.5, 98.5), with 2 additional patients with unconfirmed responses awaiting confirmatory scan

Response by investigator assessment ^a	Part 1 Sotorasib + Panitumumab + FOLFIRI (n = 6)	Part 2 Sotorasib + Panitumumab + FOLFIRI (n = 36)	Total (N = 42 ^a)
ORR confirmed (95% CI)	3 (50) (11.8, 88.2)	20 (56) (38.1, 72.1)	23 (55) (38.7, 70.2)
CR	0	0	0
PR	3 (50)	20 (56) [†]	23 (55) [†]
SD	3 (50)	13 (36)	16 (38)
PD	0	2 (6)	2 (5)
Unavailable	0	1 (3)	1 (2)
DCR (95% CI)	6 (100) (54.1, 100.0)	33 (92) (77.5, 98.3)	39 (93) (80.5, 98.5)

Efficacy and safety of IBI351 (GFH925) monotherapy in metastatic colorectal cancer harboring KRASG12C mutation: preliminary results from a pooled analysis of two phase I studies

Ying Yuan¹, Yanhong Deng², Yongdong Jin³, Yuejin Pan⁴, Cunji Wang⁵, Zhiwu Wang⁶, Zhiye Zhang⁷, Xianglao Meng⁸, Yi Hu⁹, Mingfang Zhao¹⁰, Huijuan Wang¹¹, Nong Yang¹², Dongde Wu¹³, Xiaorong Dong¹⁴, Dingthi Huang¹⁵, Meili Sun¹⁶, Lian Liu¹⁷, Dong Hu¹⁸, Zengqin Guo¹⁹, Kefeng Ding¹

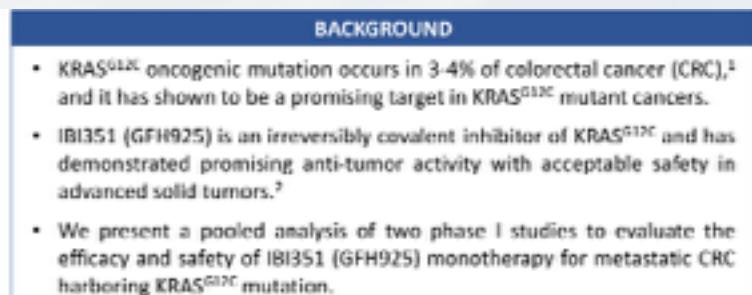
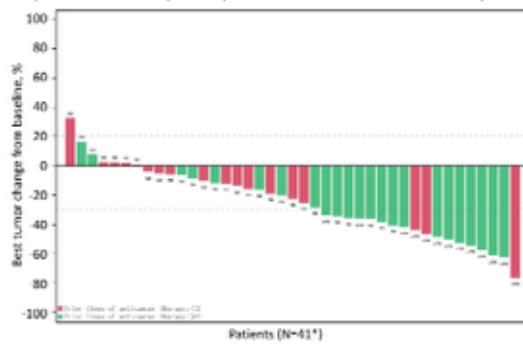


Figure 1. Tumor change in target lesion size from baseline at 600 mg BID



Patients (N=41*)

*One patient without post-baseline measurement is not shown on the graph.

Abstract #3563

Safety and Efficacy of D-1553 in KRAS G12C-mutated Colorectal Cancer: Results from a Phase I/II Study



Dan-yun Ruan¹, Myung Ah Lee², Yanhong Deng³, Keun-Wook Lee⁴, Michael Millward⁵, Jaspreet Singh Grewal⁶, Shirish M. Gadgeel⁷, Rachel E. Sanborn⁸, Xinfang Hou⁹, Shaozhong Wei¹⁰, Seok Jae Hu¹¹, Yu-rong Liu¹, Xiaoxi Xie¹², Ziyong Xiang¹², Zhe Shi¹², Yaolin Wang¹², Ling Zhang (ling.zhang@inventisbio.com)¹², Gary Edward Richardson¹³, Rui-hua Xu¹

METHODS

Figure 1. Study Design

Key Eligibility

- Refractory to or intolerant of standard therapy.
- KRAS G12C identified by molecular testing.
- Measurable disease.

Phase Ia (Dose Escalation)*

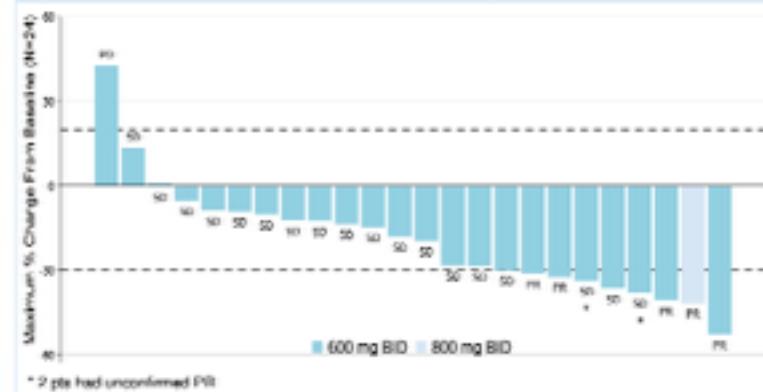
D-1553 Single Agent
(150 mg/day – 1600 mg/day)

RP2D

Phase II

Arm B: Solid tumor
D-1553 Single Agent
(600 mg BID)

Figure 3. Best Tumor Response and Tumor Burden Change



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Muchas gracias