



# Young Onset CRC: What's New about Biology and Treatment in Advanced Disease?

*Virtual Patient Conference – May 20, 2023*

*Benny Johnson, DO  
Assistant Professor  
Department of Gastrointestinal Medical Oncology  
The University of Texas - MD Anderson Cancer Center*

THE UNIVERSITY OF TEXAS  
**MD Anderson  
Cancer Center**

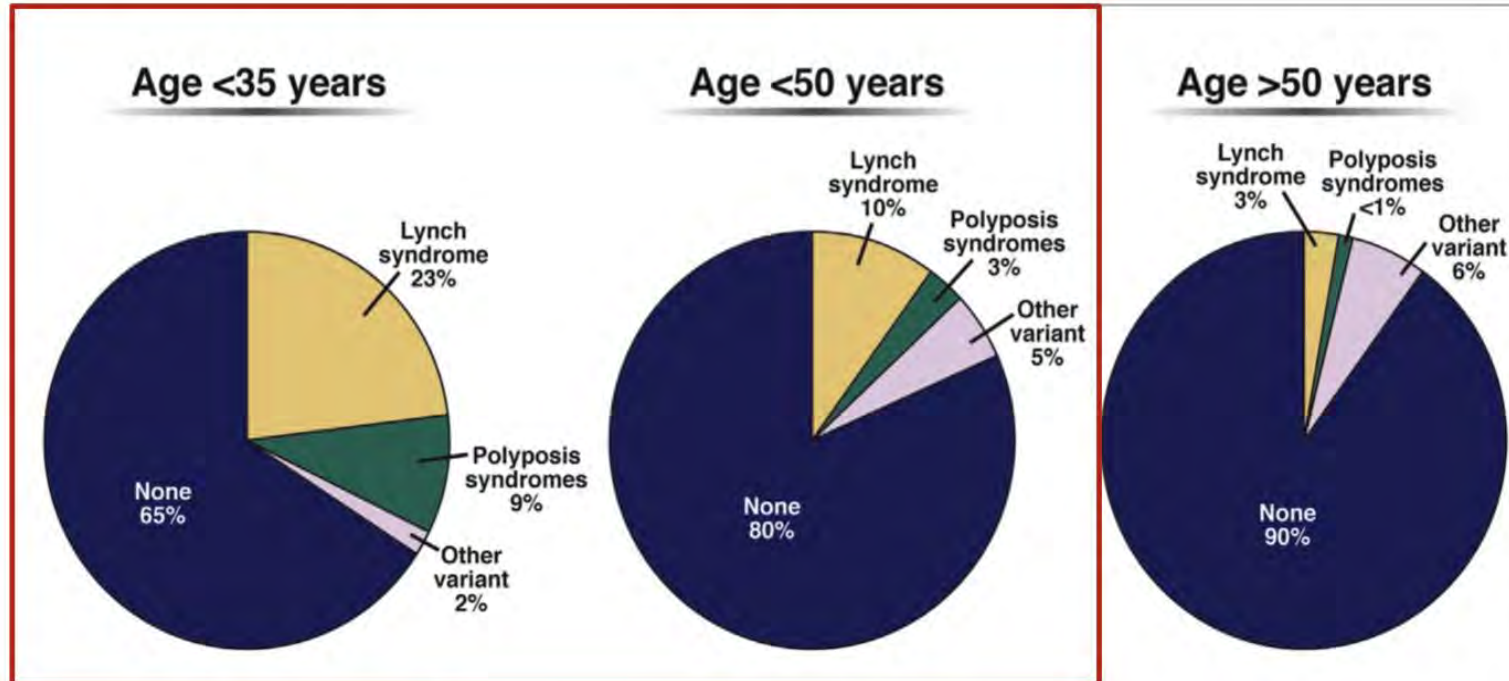
Making Cancer History®

# AGENDA

1. *Biology*
2. *Treatment*
3. *Circulating tumor DNA & Minimal Residual Disease*
4. *Conclusions*

# KEY BIOLOGIC ASPECTS OF YOCRC

# Genes of YOCRC



**~70-80% unknown**



**Sporadic YOCRC!**

Lynch syndrome	Polyposis syndromes	Other pathogenic variants	
		High penetrance	Moderate/low penetrance
<i>MLH1</i>	<i>APC</i>	<i>BRCA1</i>	<i>CHEK2</i>
<i>MSH2</i>	<i>MUTYH</i>	<i>BRCA2</i>	<i>ATM</i>
<i>MSH6</i>	<i>SMAD4</i>	<i>TP53</i>	<i>NBN</i>
<i>PMS2</i>	<i>BMPR1A</i>	<i>PALB2</i>	<i>BARD1</i>
	<i>PTEN</i>	<i>CDKN2A</i>	<i>BRIP1</i>
	<i>POLE</i>		

# MDACC Dataset + AACR GENIE

>36,000 pts

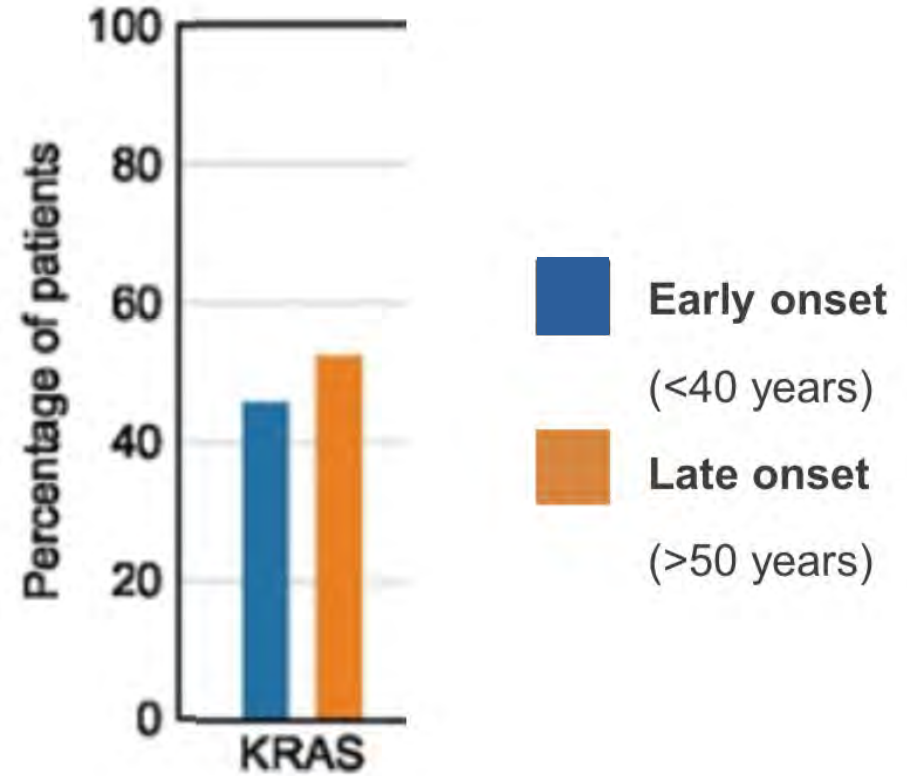
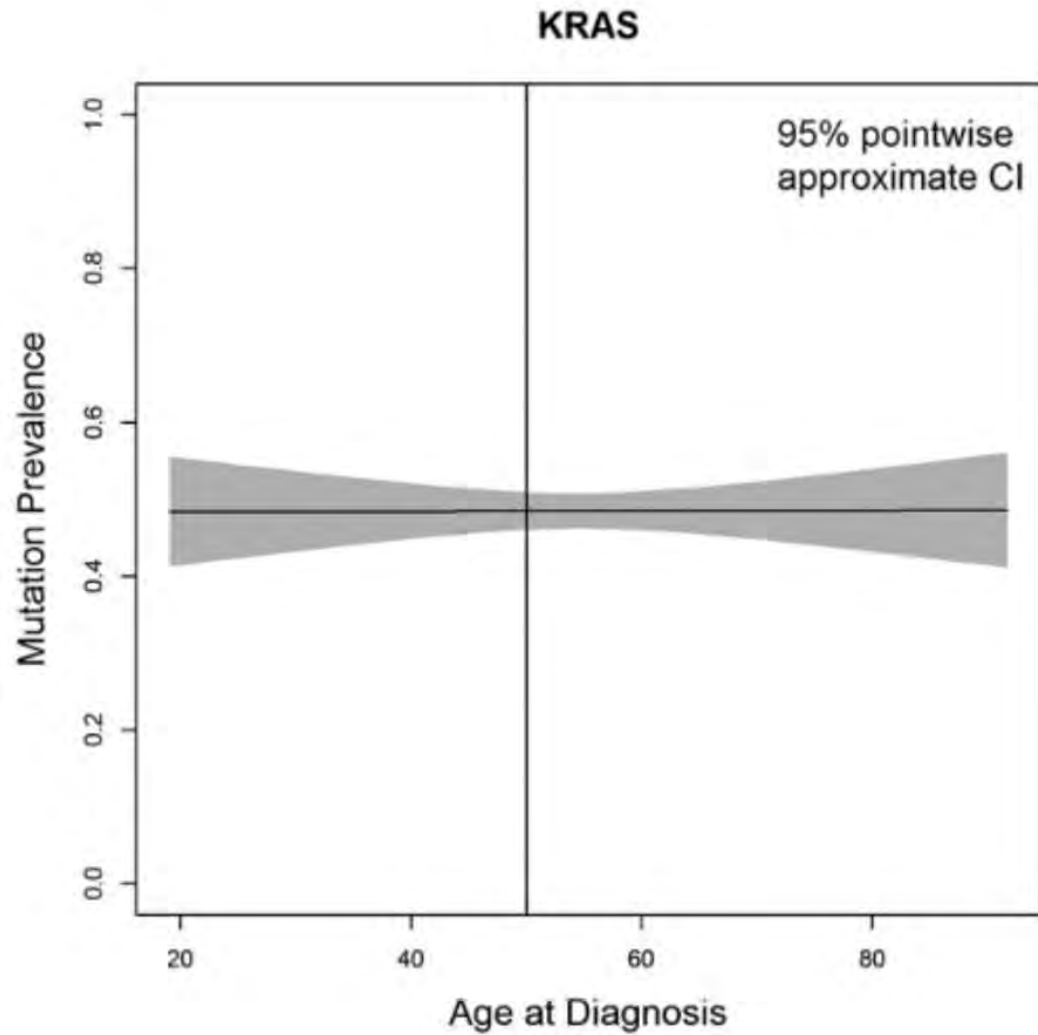
	MDACC Molecular Cohort	MDACC Tumor Registry Cohort	AACR Project GENIE Cohort	CMS Cohort
Patient Information	<ul style="list-style-type: none"> <li>N=1877</li> <li>Seen at MDACC from January 1, 2012 to September 1, 2016</li> </ul>	<ul style="list-style-type: none"> <li>N=32507</li> <li>Seen at MDACC from January 1, 1980 to present</li> </ul>	<ul style="list-style-type: none"> <li>N=1868</li> <li>Excluded patients from MDACC to prevent duplication of data</li> </ul>	<ul style="list-style-type: none"> <li>Total N=626</li> <li>N=448 from TCGA</li> <li>N=178 from MDACC</li> </ul>
Clinical Data	<ul style="list-style-type: none"> <li>Baseline clinical and pathologic characteristics</li> </ul>	<ul style="list-style-type: none"> <li>Baseline clinical and pathologic characteristics</li> </ul>	<ul style="list-style-type: none"> <li>Limited clinical and pathologic characteristics</li> </ul>	<ul style="list-style-type: none"> <li>Limited clinical and pathologic characteristics</li> </ul>
Molecular Data	<ul style="list-style-type: none"> <li>Mutational data available from 46- or 50-gene CLIA next-generation sequencing panel</li> </ul>	<ul style="list-style-type: none"> <li>Unavailable</li> </ul>	<ul style="list-style-type: none"> <li>Mutation data available from AACR Project GENIE database, which includes a mixture of next-generation sequencing platforms</li> </ul>	<ul style="list-style-type: none"> <li>RNA expression data.</li> <li>For TCGA patients, data were publicly available.</li> <li>For MDACC patients, data were obtained with Affymetrix RNA expression arrays.</li> </ul>
Cancer Stage(s)	<ul style="list-style-type: none"> <li>Stage IV</li> </ul>	<ul style="list-style-type: none"> <li>Stages I-IV</li> </ul>	<ul style="list-style-type: none"> <li>Majority stage IV</li> </ul>	<ul style="list-style-type: none"> <li>Stages I-IV</li> </ul>
Additional Data	<ul style="list-style-type: none"> <li>Comorbid predisposing condition information available for patients &lt; 50 years</li> </ul>			<ul style="list-style-type: none"> <li>Classification by CMS subtype</li> </ul>

# Foundation Medicine

18,218 total patients



- 1,420 patients under the age of 40
- 3,248 between 40 and 49
- 13,550 age 50 and older

# No significant difference in *KRAS*, *NRAS* mutations

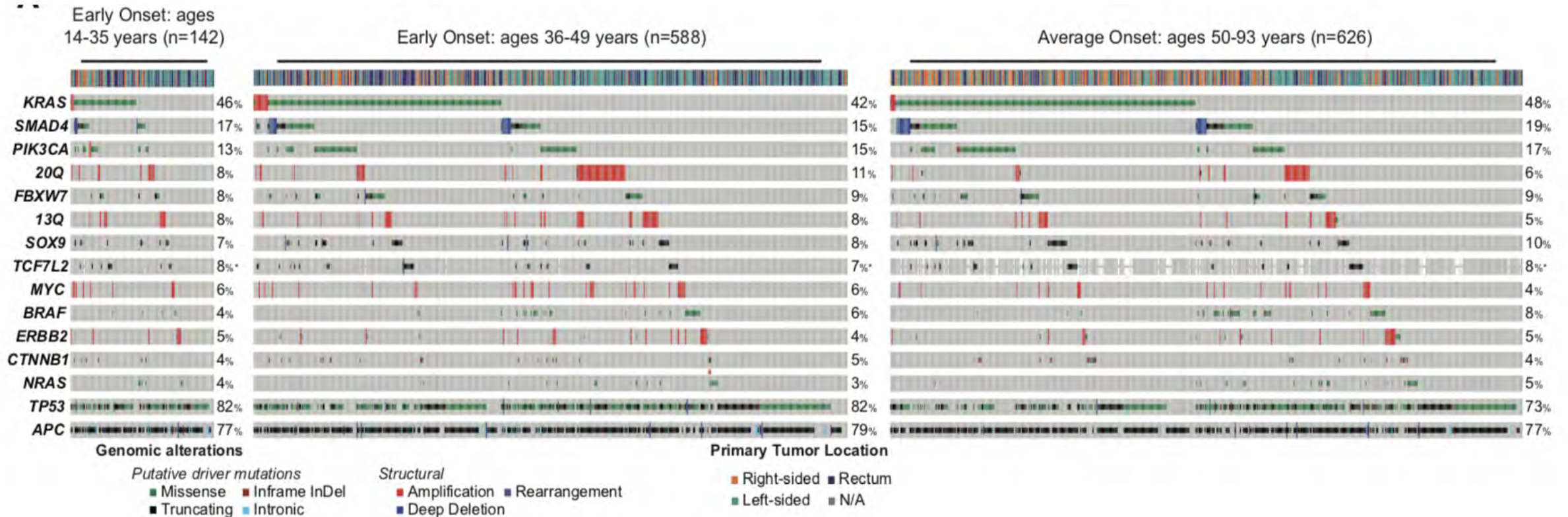


# Foundation One molecular testing → *CTNNB1*, *TP53*

**Table 1.** Significant alterations and alterations in genes of interest between cohorts using false discovery rate (FDR) in MSS colorectal cancer (CRC) and MSI-H colorectal cancer

Alteration rates in the MSS cohort			
Gene	Rate observed in under 40 group (%)	Rate observed in 50 and over group (%)	FDR
 <i>TP53</i>	82.3	76.7	1.56E-05
<i>APC</i>	65.8	79.7	4.84E-26
<i>KRAS</i>	45.6	52.4	1.56E-05
<i>PIK3CA</i>	14.1	17.5	0.002959601
 <i>CTNNB1</i>	4	2.7	0.013488987
<i>BRAF</i>	5.2	7.7	0.002067048
<i>FAM123B</i>	2	6.8	1.35E-12
<i>NRAS</i>	3.7	4.6	0.171847712

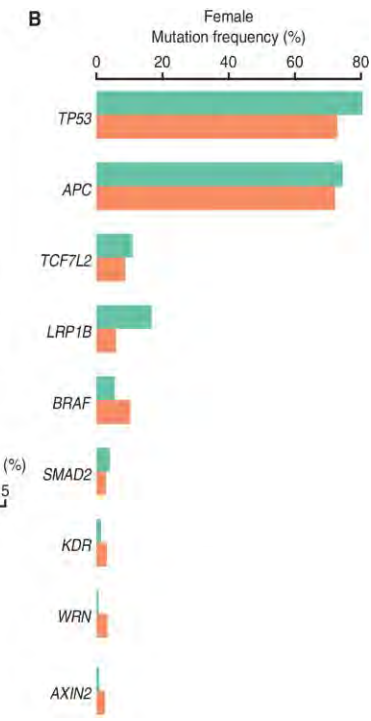
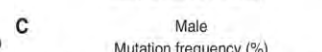
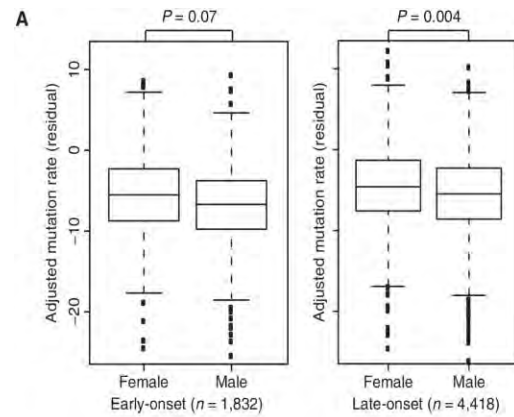
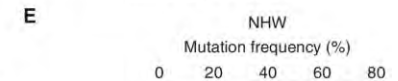
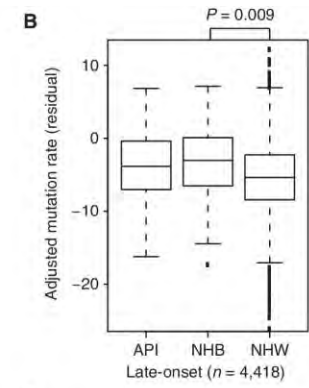
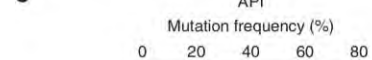
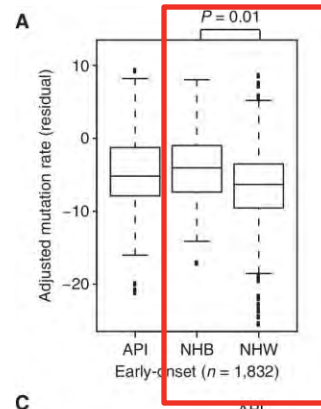
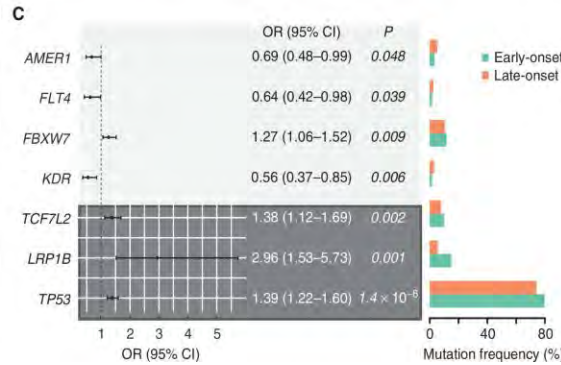
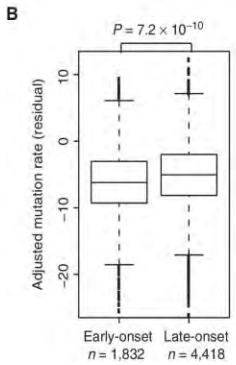
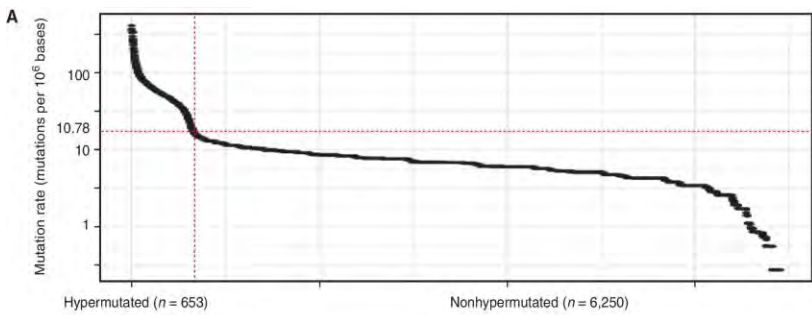
# Real world data @MSKCC: No major genomic differences between YO CRC and average onset CRC



Cercek et al, JNCI 21'



# YOCRC Biology differs based on race and sex; n=6903 pts (NHW, NHB, API)



Young patients (n=1832) with sporadic CRC had significantly higher odds of presenting with nonsilent mutations in **TP53, LRP1B, TCF7L2, and FBXW7**

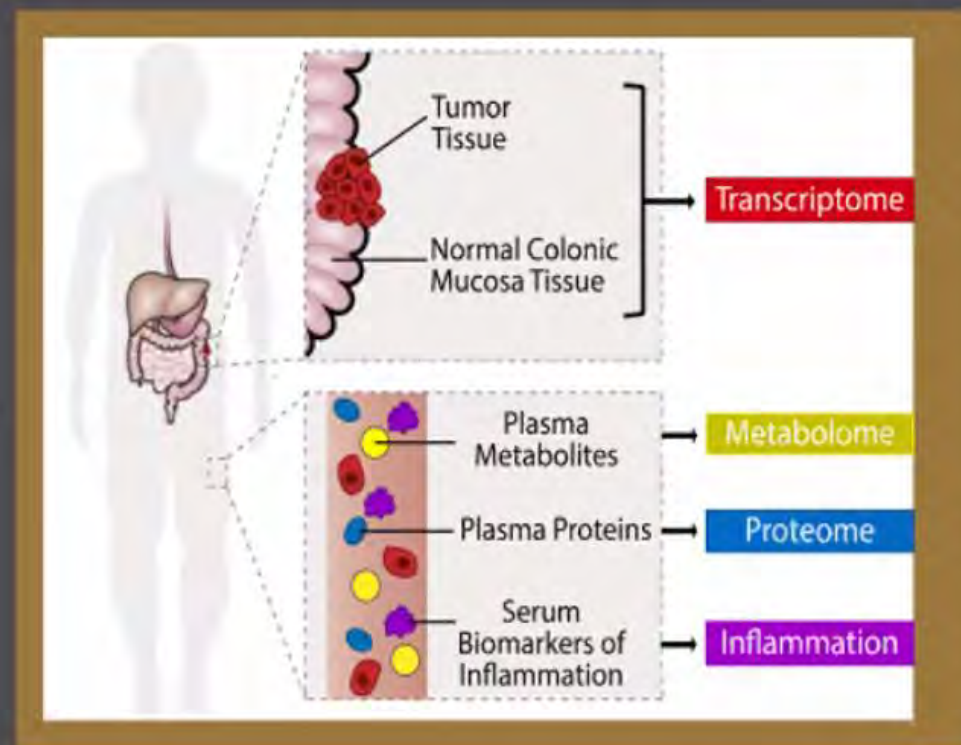
**NHBs, but not APIs, with early-onset sporadic colorectal cancer had higher adjusted tumor mutation rates versus NHWs.**

Differences for FLT4, FBXW7, RNF43, LRP1B, APC, PIK3CA, and ATRX mutation rates between **racial/ethnic groups** and EP300, KRAS, AXIN2, WRN, BRAF, and LRP1B mutation rates **by sex.**

## Deregulated redox homeostasis is a distinct molecular hallmark of early-onset sporadic CRC

Integrated, multi-omics analysis implicate perturbations in:

- *NRF2*-mediated oxidative stress response,
- glutathione metabolism, and
- the *CXCL12-CXCR4* signaling axis, as a molecular phenotype distinct to early-onset sporadic CRC.

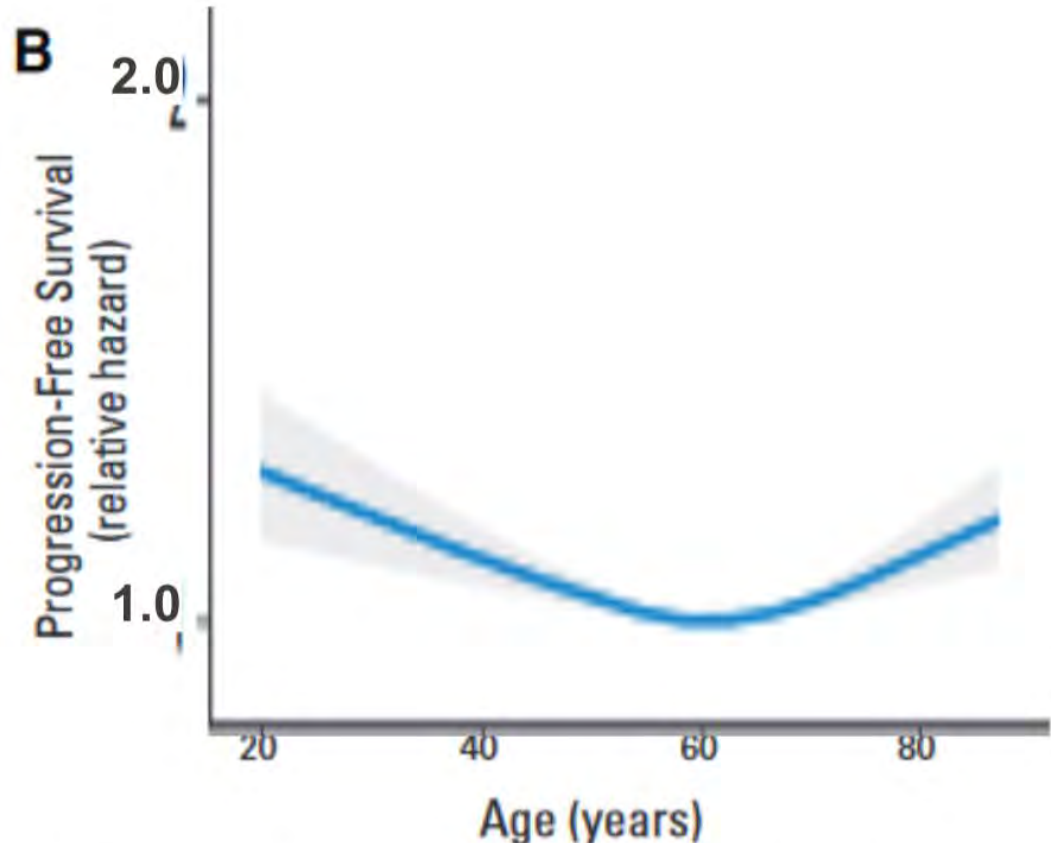
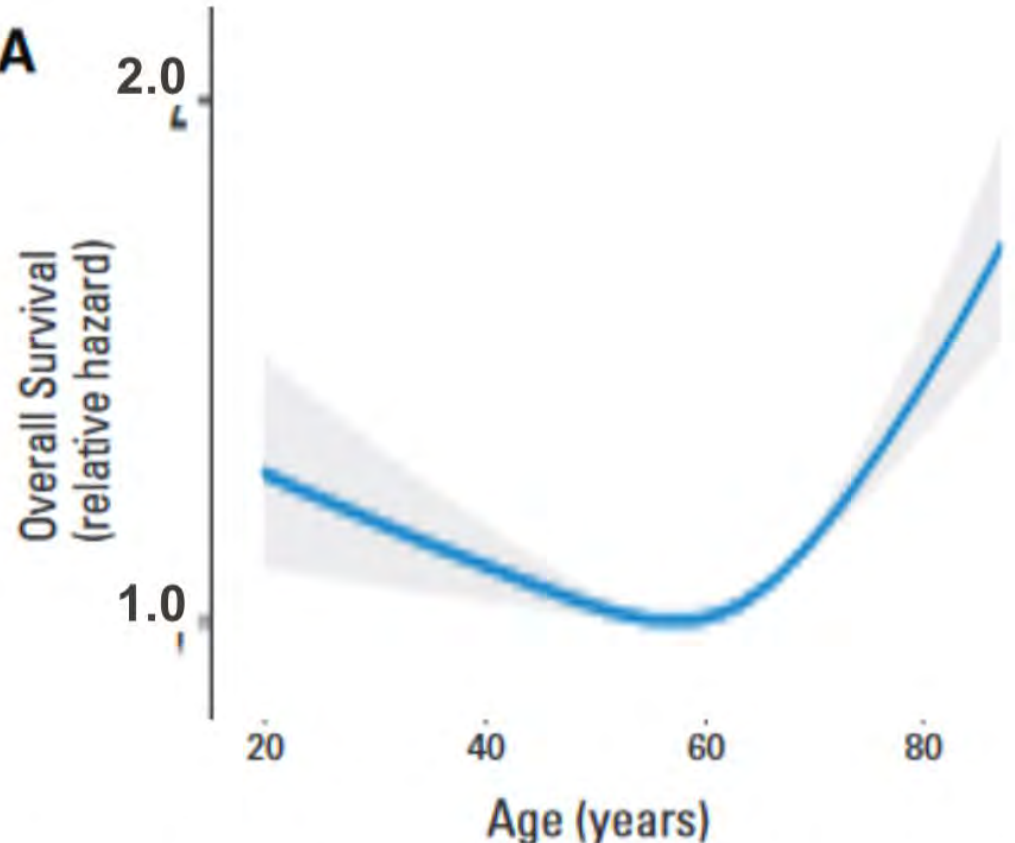


Holowatyj et al. *Gastroenterology*. 2020.

*Imbalance in  
glutathione  
metabolism*

# TREATMENT FOR YOCRC & NEW DEVELOPMENTS

# Overall survival and progression-free survival from diagnosis of mCRC is worse for EOCRC patients



20,003 patients from 24 first line studies of mCRC (ARCAD database)

# YOCRC: Warrants more 'aggressive' therapy?...

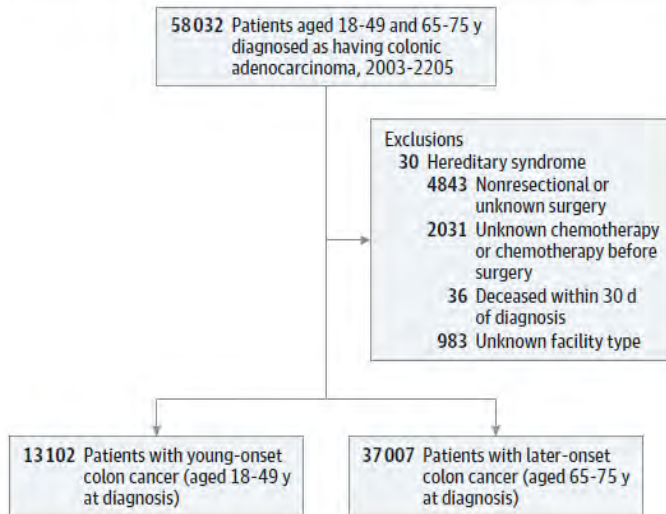
## More chemotherapy must be better.

Original Investigation

### Overtreatment of Young Adults With Colon Cancer More Intense Treatments With Unmatched Survival Gains

Peter J. Kneuert, MD; George J. Chang, MD, MS; Chung-Yuan Hu, MPH, PhD; Miguel A. Rodriguez-Bigas, MD; Cathy Eng, MD; Eduardo Vilar, MD, PhD; John M. Skibber, MD; Barry W. Feig, MD; Janice N. Cormier, MD, MPH; Y. Nancy You, MD, MHSc

Figure 1. Young Adults (Ages 18-49 Years at Diagnosis) vs Older Adults (Ages 65-75 Years at Diagnosis) With Colon Cancers



The patients were treated between January 1, 2003, and December 31, 2005, and were reported to the National Cancer Data Base.

## More Surgery must be better.

The prognostic impact of *RAS* on overall survival following liver resection in early versus late-onset colorectal cancer patients

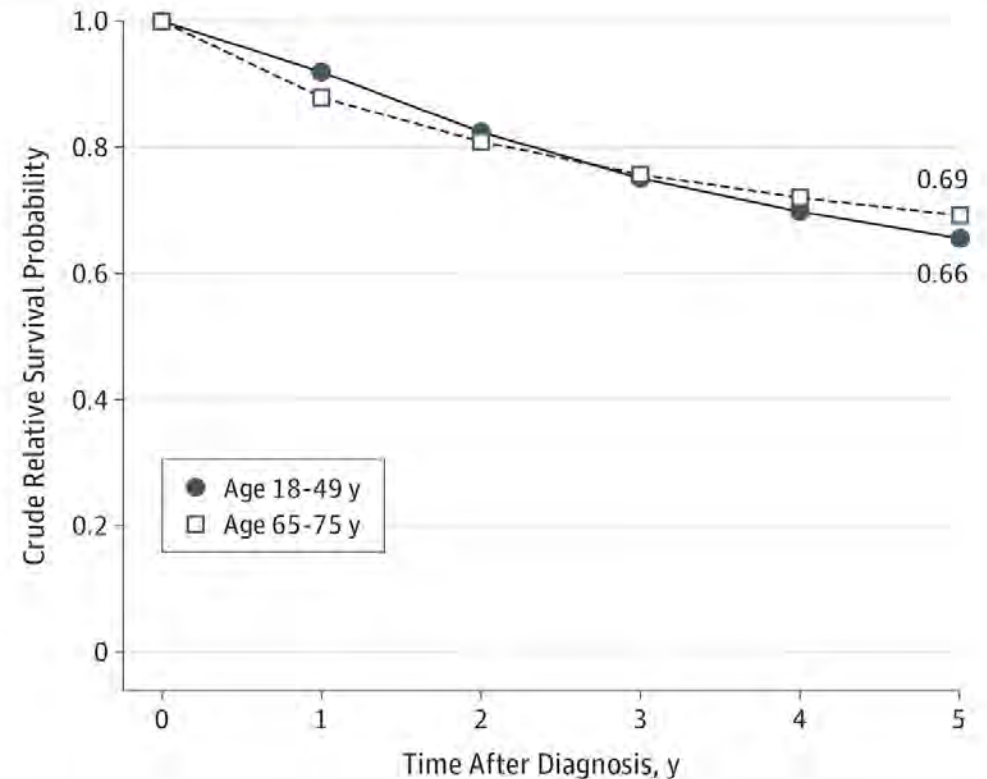
Alexandre A. Jácome<sup>1</sup>, Timothy J. Vreeland<sup>2</sup>, Benny Johnson<sup>1</sup>, Yoshikuni Kawaguchi<sup>2</sup>, Steven H. Wei<sup>2</sup>, Y. Nancy You<sup>2,3</sup>, Eduardo Vilar<sup>1,4</sup>, Jean-Nicolas Vauthey<sup>2</sup> and Cathy Eng<sup>1,5</sup>

Characteristic	Early-onset (n = 192)	Late-onset (n = 381)	P value
Median age at diagnosis (range), y	42 (22-49)	59 (50-81)	<0.001
Sex			
Male	99 (52)	236 (62)	0.019
Female	93 (48)	145 (38)	
RAS status			
Mutated	77 (40)	178 (47)	0.154
Wild-type	115 (60)	203 (53)	
BRAF status			
Mutated	5 (3)	4 (1)	0.294
Wild-type	163 (97)	293 (99)	
MSI status			
MSS	150 (98)	204 (97)	0.739
MSI-H	3 (2)	6 (3)	
Tumour location			
Ascending colon	29 (15)	93 (25)	0.012
Transverse colon	8 (4)	16 (4)	0.527
Descending colon	9 (5)	32 (8)	0.122
Rectosigmoid	146 (76)	240 (63)	0.001
Sidedness			
Right	37 (19)	109 (29)	0.015
Left	155 (81)	272 (71)	
CEA level > 10 ng/mL			
Yes	40 (22)	101 (27)	0.213
No	143 (78)	271 (73)	
Bilobar disease			
Yes	39 (21)	28 (26)	0.389
No	149 (79)	82 (75)	
≥2 liver lesions			
Yes	92 (48)	182 (49)	1
No	98 (52)	193 (51)	

# More intensive chemotherapy for YOCRC – did not translate to survival benefits

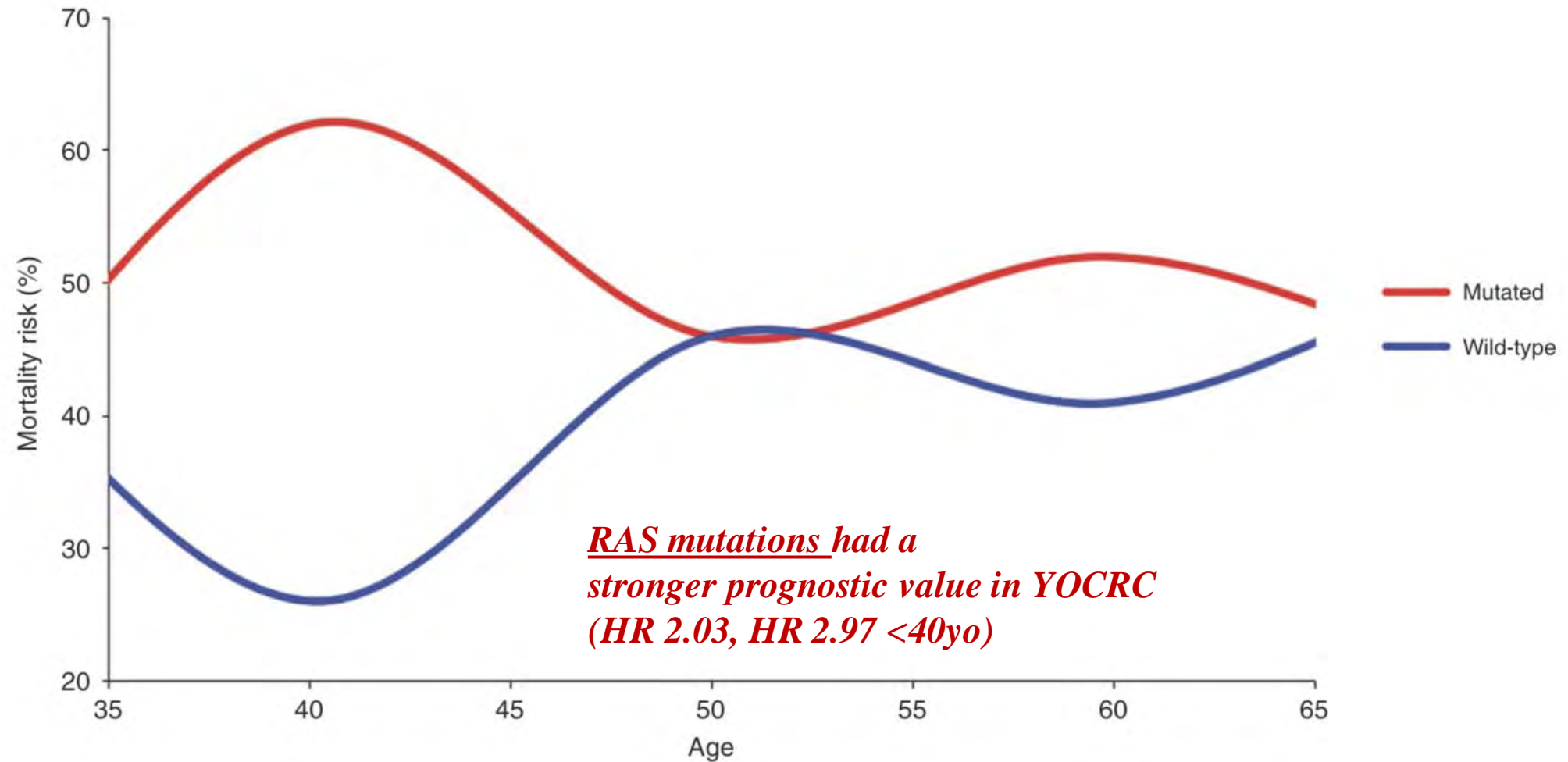
Patients Who Received Chemotherapy	Any Chemotherapy, No. (%)	Odds Ratio for Receiving Chemotherapy (95% CI)	Multiagent Regimens, No. (%)	Odds Ratio for Receiving Multiagent Regimen (95% CI)
<b>Stage I</b>				
Ages 65-75 y (n = 8991)	162 (1.8)	1 [Reference]	52 (43.0)	1 [Reference]
Ages 18-49 y (n = 1926)	109 (5.7)	2.88 (2.21-3.77)	43 (48.3)	1.38 (0.71-2.68)
<b>Stage II Overall</b>				
Ages 65-75 y (n = 11 011)	2748 (25.0)	1 [Reference]	773 (41.7)	1 [Reference]
Ages 18-49 y (n = 3083)	1732 (56.2)	3.93 (3.58-4.31)	670 (54.9)	1.71 (1.48-1.97)
<b>Stage II Low Risk</b>				
Ages 65-75 y (n = 4822)	923 (19.1)	1 [Reference]	313 (39.6)	1 [Reference]
Ages 18-49 y (n = 1636)	826 (50.5)	4.22 (3.70-4.81)	388 (52.5)	1.67 (1.34-2.09)
<b>Stage II High Risk</b>				
Ages 65-75 y (n = 6189)	1825 (29.5)	1 [Reference]	677 (42.7)	1 [Reference]
Ages 18-49 y (n = 1447)	906 (62.6)	3.69 (3.23-4.20)	454 (57.0)	1.77 (1.46-2.14)
<b>Stage III</b>				
Ages 65-75 y (n = 11 202)	8175 (73.0)	1 [Reference]	4209 (59.4)	1 [Reference]
Ages 18-49 y (n = 4780)	4132 (86.4)	2.42 (2.18-2.68)	2590 (71.5)	1.75 (1.58-1.93)
<b>Stage IV</b>				
Ages 65-75 y (n = 5803)	3652 (62.9)	1 [Reference]	2567 (80.4)	1 [Reference]
Ages 18-49 y (n = 3313)	2710 (81.8)	2.74 (2.44-3.07)	2136 (88.6)	1.90 (1.60-2.26)

Figure 2. Crude Relative Survival of Young Adults (Ages 18-49 Years at Diagnosis) vs Older Adults (Ages 65-75 Years at Diagnosis) With Colon Cancers

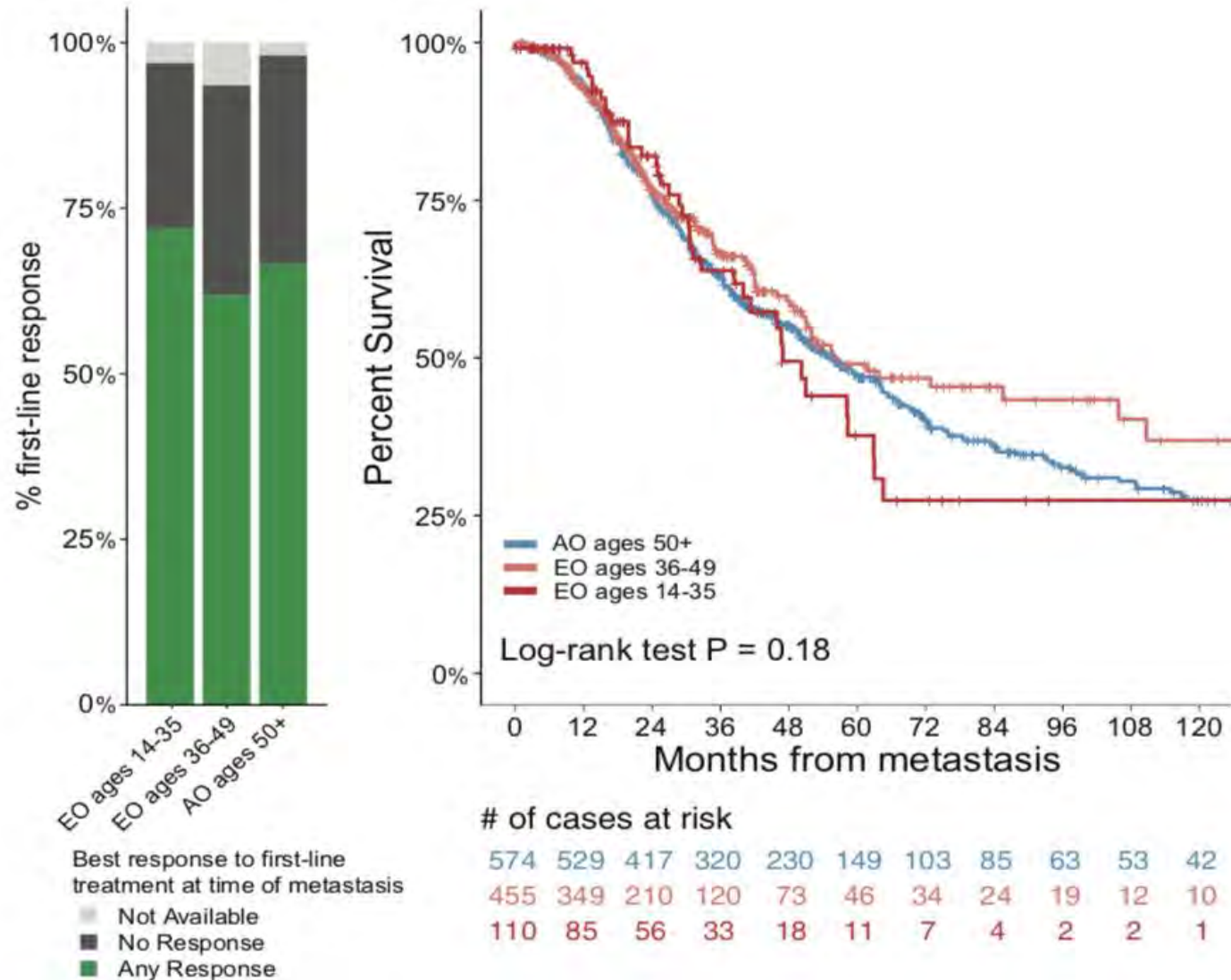


The unadjusted survival analysis showed slightly inferior 5-year relative survival for the young adults (0.66 vs 0.69,  $P < .001$ ).

# Surgical Outcomes in mCRC: Prognostic impact of *RAS* mutation status CLM in YOCRC



# Sporadic YOCRC with very similar response to 1L chemotherapy & survival as average age population



*Cercek et al JNCI 21'*



# FOLFOXIRI in mCRC

Triplet/bev vs doublets/bev in mCRC

Pooled analysis; n = 1697

Primary Endpoint: OS

	Triplet/bev	Doublet/bev*	p
ORR (%)	<b>64.5</b>	53.6	p < 0.001
Median OS(mos)	<b>28.9</b>	24.5	p < 0.001
Median PFS (mos)	<b>12.2</b>	9.9	p < 0.001
5-year OS (%)	<b>22.3%</b>	10.7%	P < 0.001

\*70% FOLFOX/bev; 30% FOLFIRI/bev

Triplet/bev vs doublets/bev in mCRC

Secondary analyses of survival in resected pts

Trial / Endpoint	Triplet/bev	Doublet/bev*	HR (95% CI)
OLIVIA / RFS (mos)	<b>17.1</b>	8.1	0.31 (0.12 – 0.75)
Cremoloni et al, OS (mos)	<b>64</b>	52.6	0.79 (0.5 – 1.24)

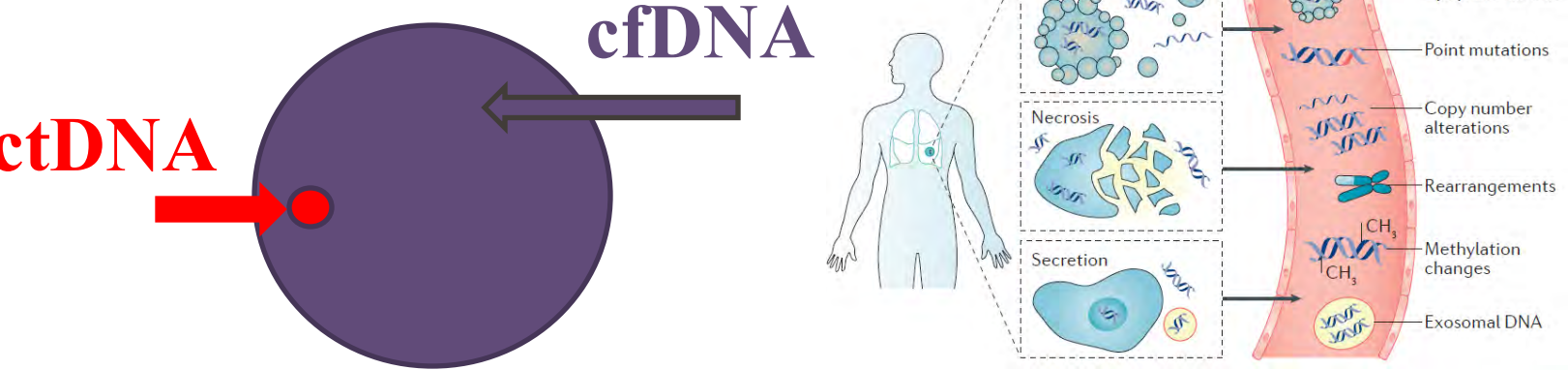
# How I treat a patient with Advanced Young Onset CRC

1. Establish **goals** of therapy & highlight **supportive services** (oncofertility, social work, integrative medicine, supportive care)
2. All pts: **Precision** : **Expanded molecular profiling** - tissue NGS or ctDNA testing
3. **Early** surgical consultation in stage IV with potential resectable liver or lung metastases; careful use of triplet chemotherapy for conversion (FOLFOXIRI/BEV)
4. **1L** - FOLFOX / FOLFIRI +/- BEV or anti-EGFR (if left sided colon cancer, RAS wt) ; Immunotherapy if Lynch syndrome/MSI-H ; 3 drug chemo if very symptomatic → *Clinical trial enrollment*
5. **2L** – opposite chemotherapy backbone + appropriate biologic / targets (HER2 amp; BRAFV600E; KRAS G12C) / → *Clinical trial enrollment*
6. **3L** – TAS-102/BEV or Regorafenib → *Clinical trial enrollment*

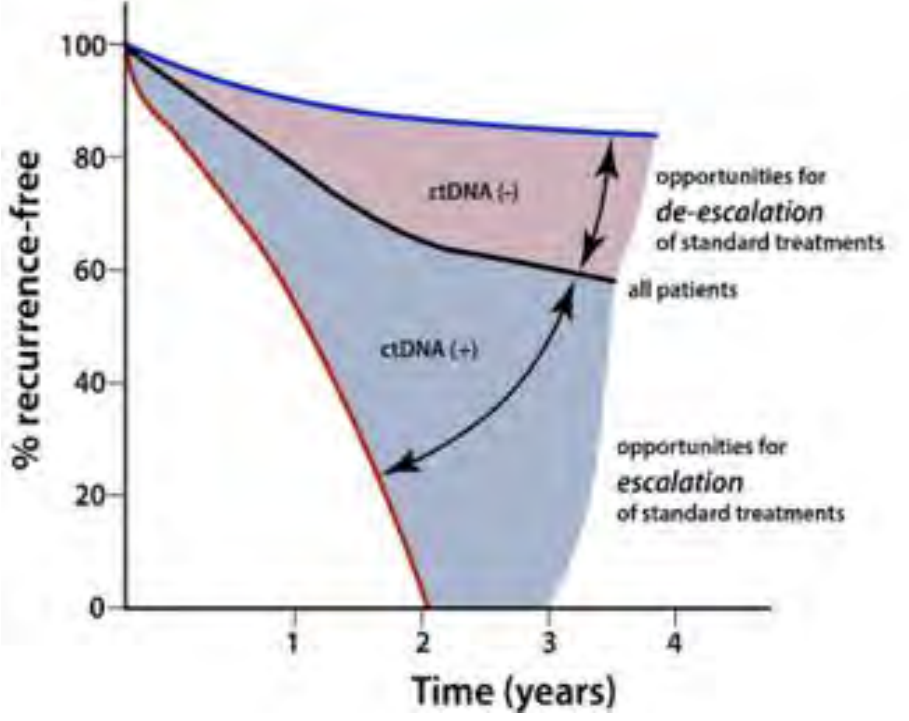
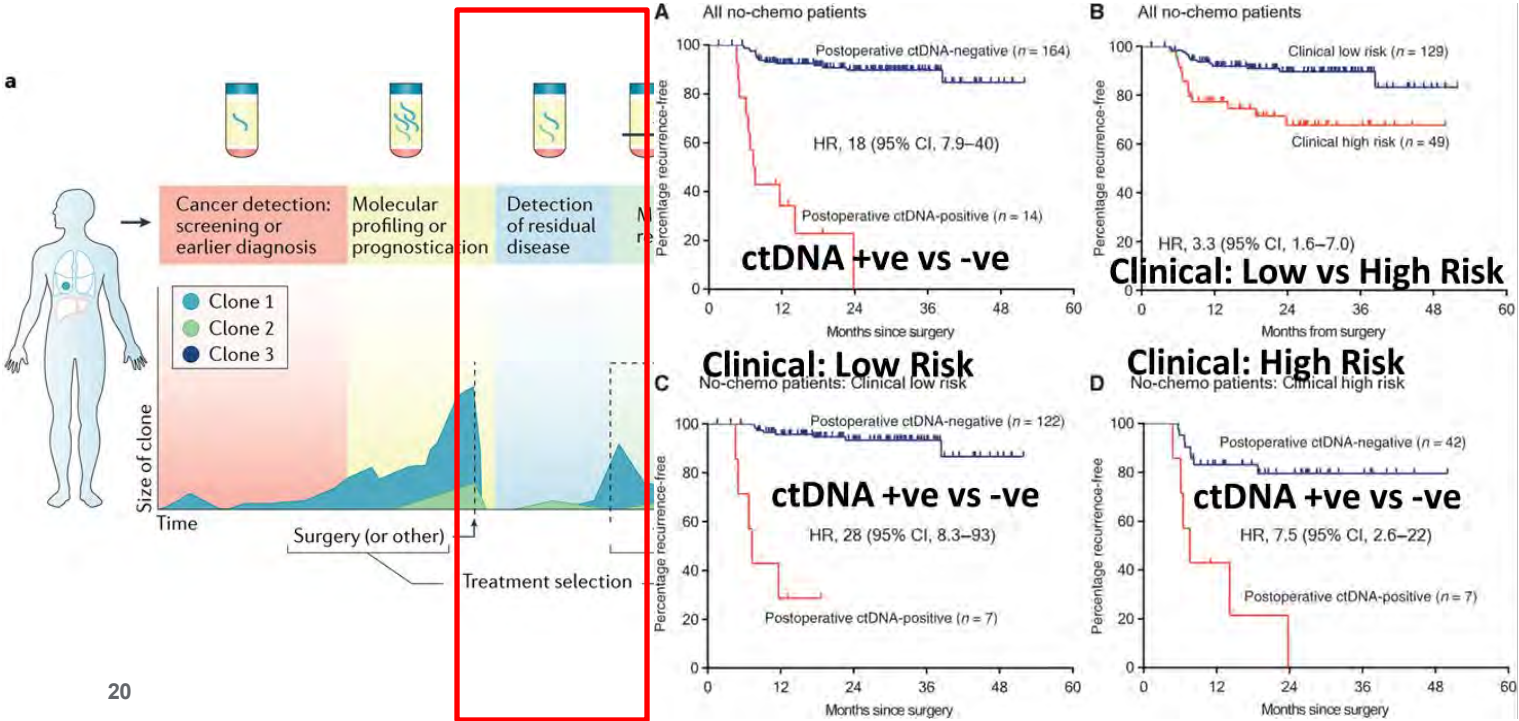
# Paradigm Shifts in CRC with impact for Young Onset

- 1. Expanded NGS/ctDNA testing** → Key biologic subgroups w/ FDA approved targeted therapies available:
  - *MSI-H / BRAFV600E / RAS/RAF wt / HER2 amplification / NTRK fusions*
- 2. 'Watch and Wait'** in rectal cancer for cCR after upfront chemotherapy/radiation
- 3. Emergence of ctDNA – the future:** guided/personalization of therapy in **stage II and III** CRC to avoid toxicity from oxaliplatin based chemotherapy (DYNAMIC, ongoing COBRA & CIRCULATE-US)
- 4. Immunotherapy alone for dMMR/MSI-H** locally advanced rectal cancer- (avoidance of chemotherapy/radiation/surgery) (n=12; 100% cCR) – Caveat: **applies to only a small subset of pts**
- 5. Clinical Trial/Emerging targets** – *KRAS G12C combo*; neoantigen vaccine approaches, TCR therapy; **MRD trials**

# Circulating tumor DNA (ctDNA) & Minimal Residual Disease (MRD) in Colorectal Cancer



- ctDNA can be detected in the blood following release from tumor cells,
  - Fragment size: cfDNA ~167 bp; ctDNA ~20-30 bp shorter
  - “real time” analysis;  $t_{1/2} \sim 2 - 3$  hours
- cfDNA described > 70 years ago;*  
*ctDNA described > 40 years ago*

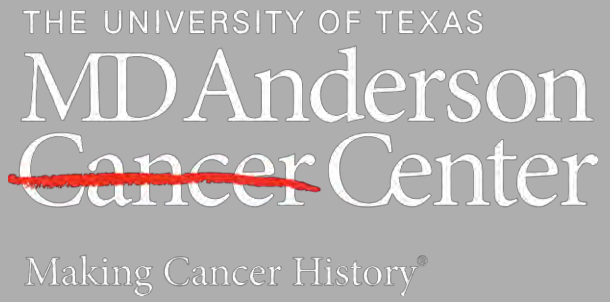


## Protocols at MDACC: Targeting CRC MRD with novel approaches

Intervention (INTERCEPT Study Lead)	Setting
COBRA NRG (Dr. Van Morris)	Stage II, immediate post op
CIRCULATE US– NRG / SWOG (Dr. Arvind Dasari)	Stage III / ctDNA+ stage II, immediate post op
BioNTech RNA vaccine (Dr. Van Morris / Dr. Scott Kopetz)	Stage II, III, immediate post op
Chemo de-escalation (Dr. Timothy Newhook)	Stage IV, immediately post op
KRAS vaccine (Dr. Shubham Pant)	Any stage
Cetuximab + NK Cell therapy (Dr. Pia Morelli)	Any stage
Lifestyle Bootcamp (Dr. Alisha Bent)	Any stage
TAS-102 (Dr. Arvind Dasari / Dr. Alisha Bent)	Any stage
CXCR1/2 inhibitor + anti-PD-1 (Dr. Benny Johnson)	Any stage

# Conclusions

- No major differences in frequency of traditional mutations of interest (*KRAS*, *NRAS*, *BRAF*, HER2 amplification, etc) in YOCRC.
- There are clues to underlying **genomic differences** noted in sporadic YOCRC biology – that may be related to **ethnicity & sex**.
- YOCRC treatment should incorporate **precision** regarding **key molecular drivers, goals** of therapy – essentially **personalized** for each patient.
- **Clinical trial enrollment** should be a part of the cancer journey for **all patients** with YOCRC.
- Novel de-escalation/escalation & monitoring strategies utilizing **circulating tumor DNA (ctDNA)** have relevant implications for YOCRC treatment & survivorship.



***Thank you!***  
***Any questions? – [bjohnson6@mdanderson.org](mailto:bjohnson6@mdanderson.org)***  
***@benjohnson1112***