

بِسْمِ اللَّهِ الرَّحْمَنِ الرَّحِيمِ



Immunotherapy of GI Cancers

S.H. Mirpour;MD.

Medical Oncologist,GUMS,1401/02/05

Why Does Cancer Occur?

- I now understand that cancer “happens” all the time in all of us
 - Most malignant cells are either cleared by the immune system, undergo DNA repair, or experience apoptosis
- That means that cancers that grow and need treatment, either:
 - Evade the immune system, or
 - Have broken DNA repair and apoptosis mechanisms

Select Key Validated Biomarkers and Associated Therapies Across GI Cancers

Tumor Type	Molecular Testing Commonly Used (Associated Targeted Agents)
Colorectal	BRAF V600E mut (encorafenib + cetuximab); HER2+ (trastuzumab + pertuzumab or lapatinib; trastuzumab deruxtecan); MSI-H/dMMR (pembrolizumab; nivolumab ± ipilimumab)
Gastric/GEJ/esophageal	PD-L1+ (nivolumab; pembrolizumab); HER2+ (trastuzumab; trastuzumab deruxtecan; pembrolizumab + trastuzumab/fluoropyrimidine- and platinum-containing CT)
Pancreatic	BRCA1/2 mut (olaparib)
Cholangiocarcinoma	FGFR2 mut (pemigatinib; infigratinib), IDH1 mut (ivosidenib)
Tumor agnostic	MSI-H/dMMR (pembrolizumab, dostarlimab); TMB-H (pembrolizumab); NTRK fusion (larotrectinib, entrectinib)

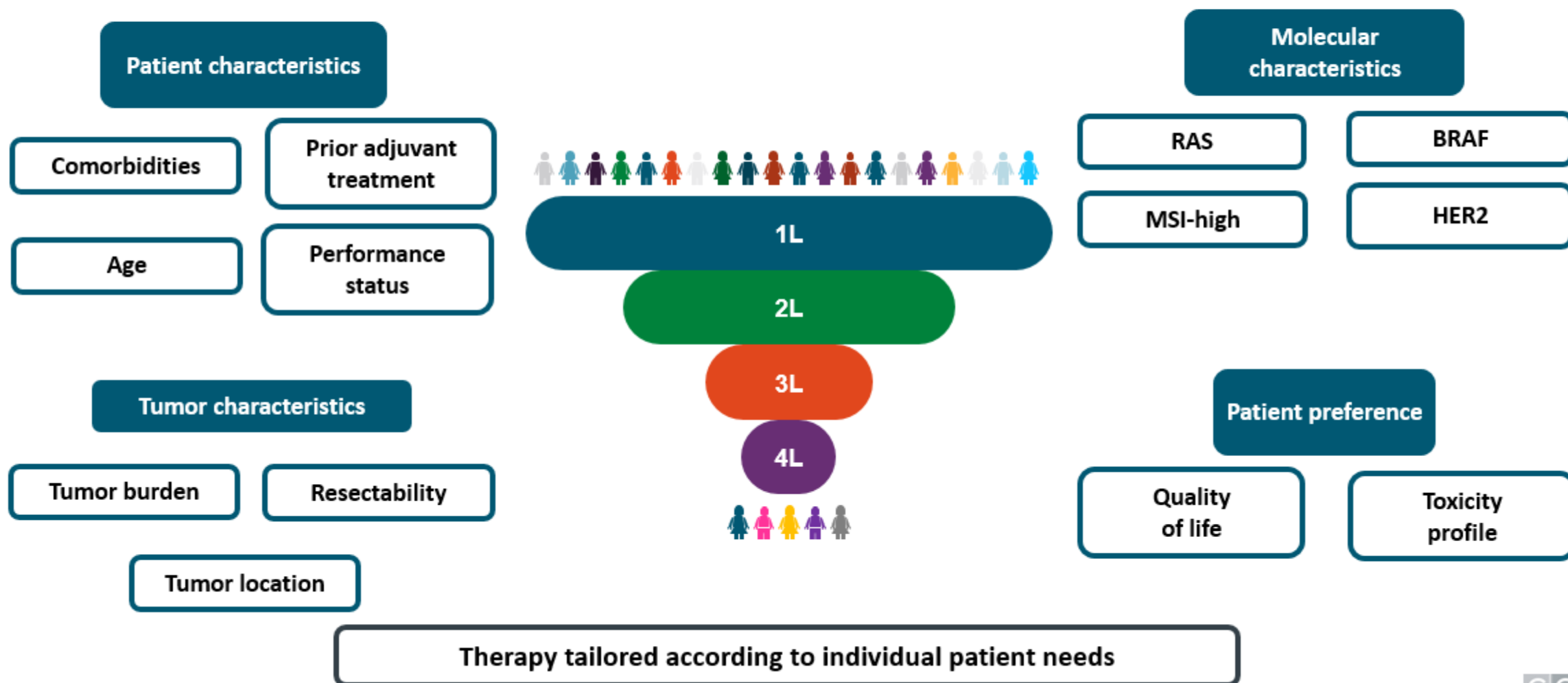
Biomarker-Driven Treatment of Advanced Colorectal Cancer



STRONGERTOGETHER

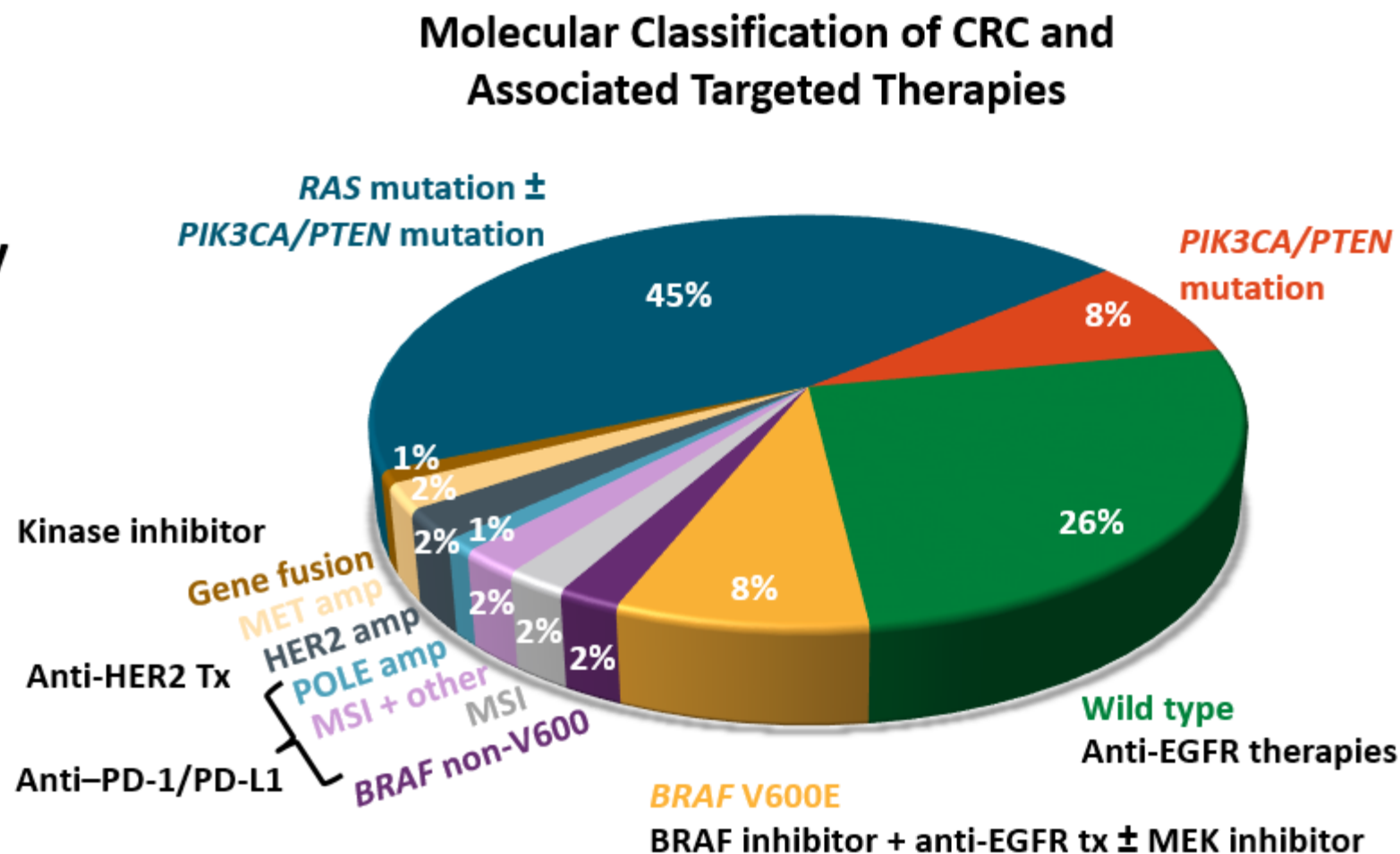


What Influences Treatment Choices in mCRC?



Biomarker Testing in CRC

- For all colon cancers:
 - MMR
 - Microsatellite stability
- Metastatic disease:
 - *KRAS*, *NRAS*, *BRAF*
 - *HER2* amplification
 - Panels: \pm fusion, broad NGS



General Treatment Algorithm for Metastatic CRC

	<i>BRAF/RAS wt</i>	<i>RAS mut</i>	<i>BRAF V600 mut</i>	<i>MSI-H/dMMR</i>	<i>NTRK fusion</i>	<i>HER2 amplification</i>
1L	<i>R side: CT + bev</i> <i>L side: CT + EGFRi or bev¹⁻⁴</i>	CT + bev ⁵⁻⁷	FOLFOXIRI + bev or Doublet + bev	Pembrolizumab	As with <i>BRAF/RAS wt</i>	
	Consider capecitabine + bev maintenance ⁹					
2L	CT as with <i>RAS mut</i> ; If bev: in 1L continue bev or change to EGFRi If EGFRi in 1L, bev	<i>If prior oxaliplatin:</i> irinotecan-based regimen + bev <i>If prior irinotecan:</i> oxaliplatin-based regimen + bev <i>If prior FOLFOXIRI:</i> regorafenib ¹¹ or TAS-102 ± bev ^{12,13}	Consider encorafenib + EGFRi i ¹⁴ ; otherwise, CT + bevacizumab	<i>If no prior PD-1i:</i> PD-1i ± CTLA-4i*; otherwise, CT/TT as with <i>BRAF/RAS wt</i>	<i>If no prior TRK inhibitor:</i> consider larotrectinib or entrectinib ^{15,16} ; otherwise, CT/TT as with <i>BRAF/RAS wt</i>	Consider trastuzumab (+ pertuzumab or lapatinib) or trastuzumab deruxtecan ¹⁷⁻¹⁹ ; otherwise, CT/TT as with <i>BRAF/RAS wt</i>
3L+	<i>If prior oxaliplatin- and irinotecan-based regimens:</i> regorafenib or TAS-102 ± bev			CT, chemotherapy regimens, including oxaliplatin- and/or irinotecan-based regimens (eg, FOLFOX, FOLFIRI, FOLFOXIRI, CAPOX). EGFRi, EGFR inhibitors, including cetuximab or panitumumab.		

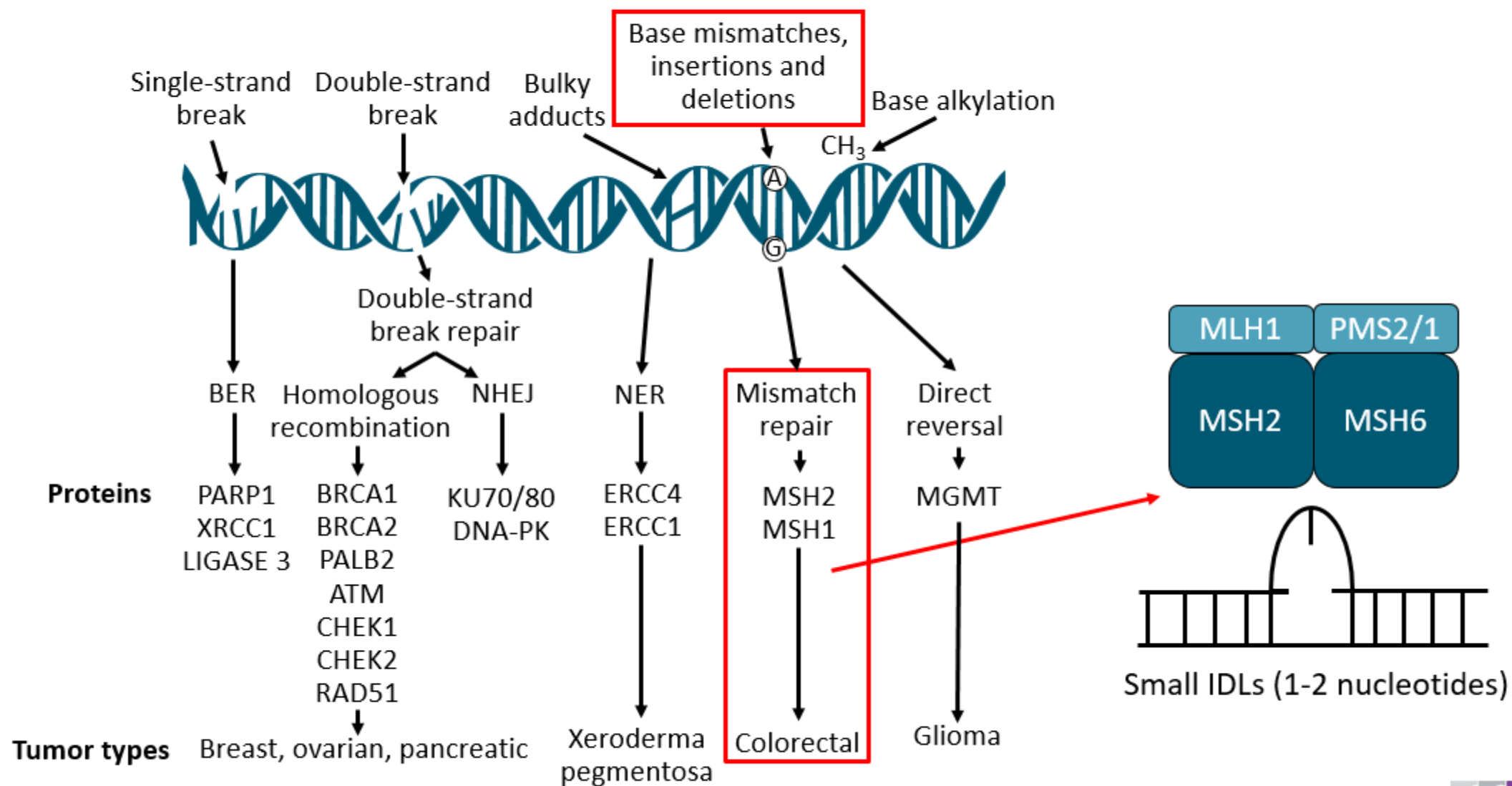
CT, chemotherapy regimens, including oxaliplatin- and/or irinotecan-based regimens (eg, FOLFOX, FOLFIRI, FOLFOXIRI, CAPOX). EGFRi, EGFR inhibitors, including cetuximab or panitumumab.

*If prior PD-1i monotherapy only, can consider PD-1i + CTLA-4i.

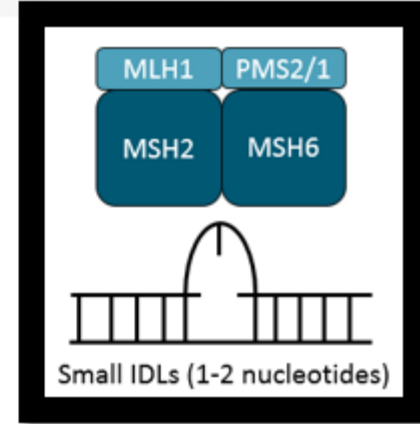
1. Tejpar. JAMA Oncol. 2017;3:194.
2. Venook. JAMA. 2017;317:2392.
3. Loupakis. NEJM. 2014;371:1609.
4. Cremolini. Lancet Oncology. 2020;21:497.
5. Parikh. Clin Cancer Res. 2019;25:2988.
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7. Van Cutsem. JCO. 2011;29:2011.
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11. Grothey. Lancet. 2013;381:303.
12. Mayer. NEJM. 2015;372:1909.
13. Pfeiffer. Lancet Oncology. 2020;21:412.
14. Tabernero. JCO. 2021;39:273.
15. Hong. Lancet Oncol. 2020;21:531.
16. Doebele. Lancet Oncol. 2020;21:271.
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18. Meric-Bernstam. Lancet Oncol. 2019;20:518.
19. Siena. Lancet Oncol. 2021;22:779.



DNA Repair Mechanisms

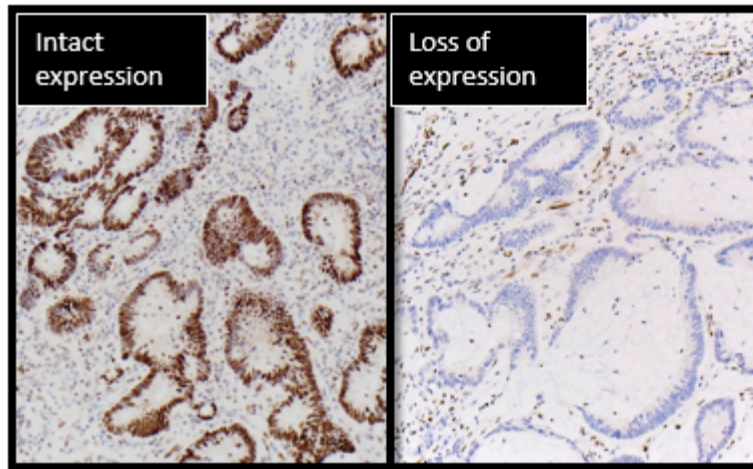


dMMR Testing: Methods



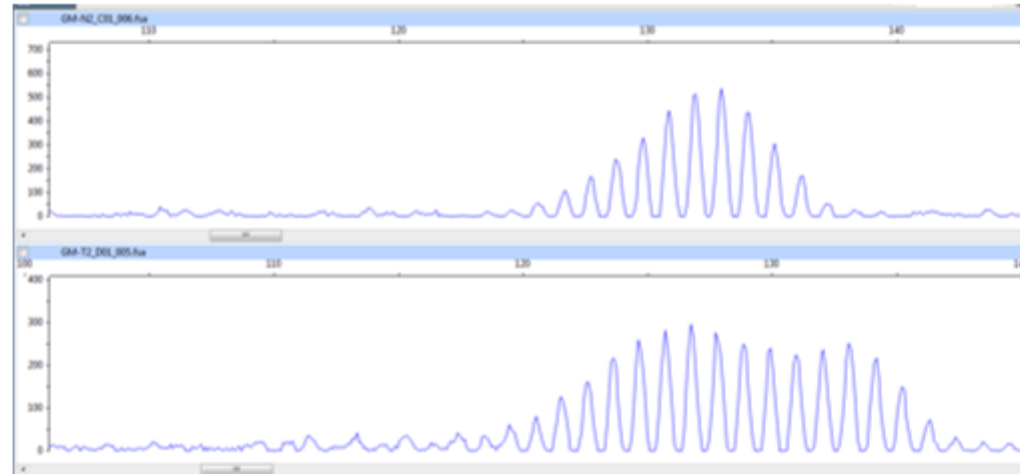
IHC

Complete loss of expression in one of the MMR proteins = MSI high



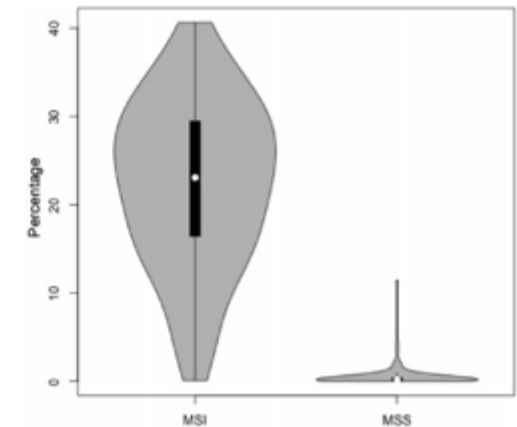
PCR

Panel of 5 or more microsatellites with allelic shift in 2 (> 30%) or more markers = MSI high



Next-Generation Sequencing

Differentiates microsatellite instable vs microsatellite stable samples



Interpreting MSI Status

- A loss (ie, “reduced” or “deficient”) in any of these proteins—MLH1, MSH2, PMS2, and MSH6—suggests potential MSI-H status and warrants further testing with PCR to confirm MSI-H
- Some facilities only test for PMS2 and MSH6; the intact presence of these 2 markers alone is insufficient to rule out MSI-H status
- If all 4 proteins are intact (or “expressed,” “retained,” “present”) tumor is microsatellite stable and unlikely to respond to checkpoint inhibitor monotherapy . . . most of the time

Microsatellite Testing Options

- IHC and PCR can be used to determine microsatellite status and are likely equally efficacious

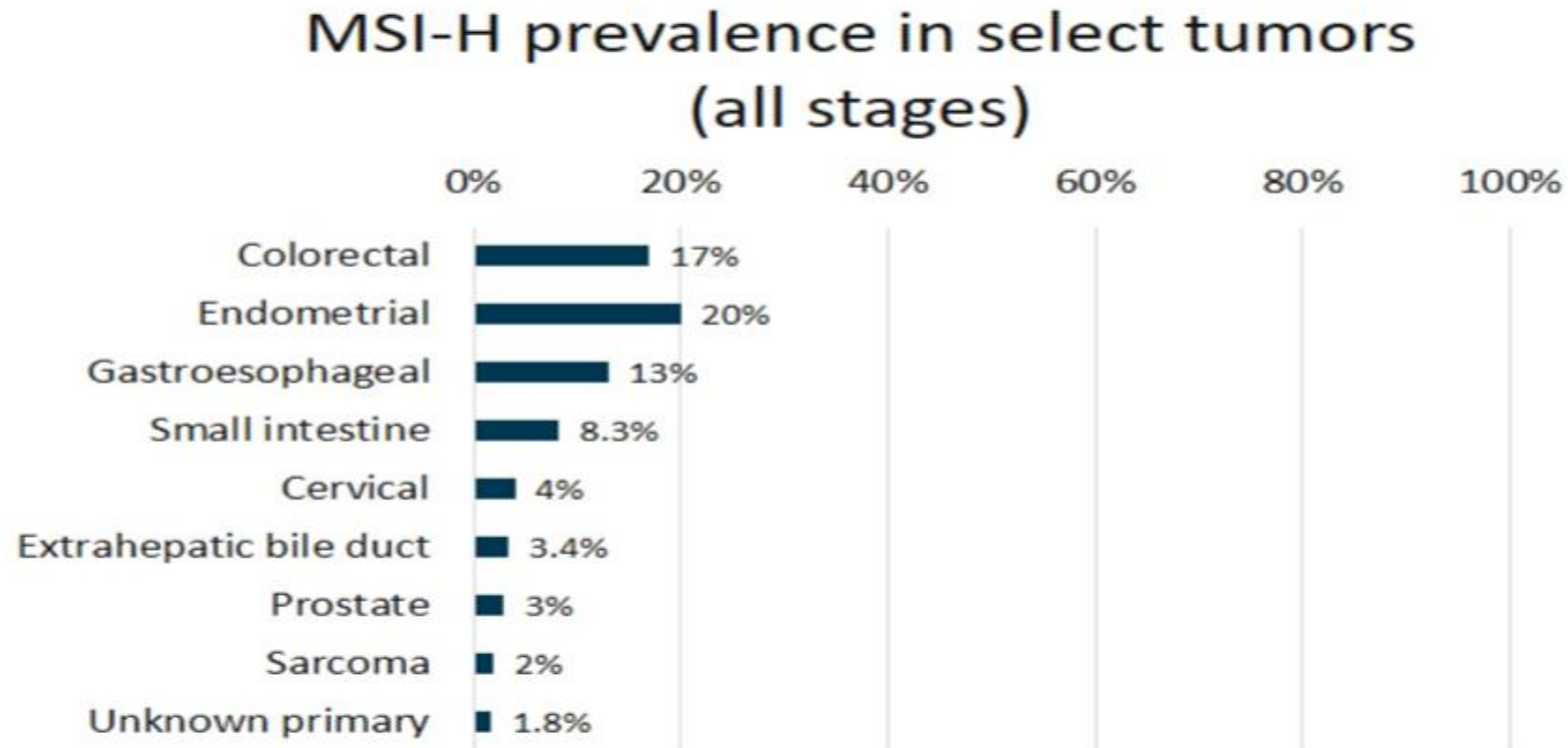
IHC

- Assesses for protein products of specific MMR genes (ie, *MLH1*, *MSH2*, *MSH6*, and *PMS2*)
- Classifies tumor as MSI or MSS
 - MSI:** loss of at least 1 of *MLH1*, *MSH2*, *MSH6*, and/or *PMS2* proteins, indicating tumor is MMR deficient
 - MSS:** all proteins are retained, indicating tumor is pMMR
- When test report reads "positive for expression," it means MSS and should not be misinterpreted as MSI

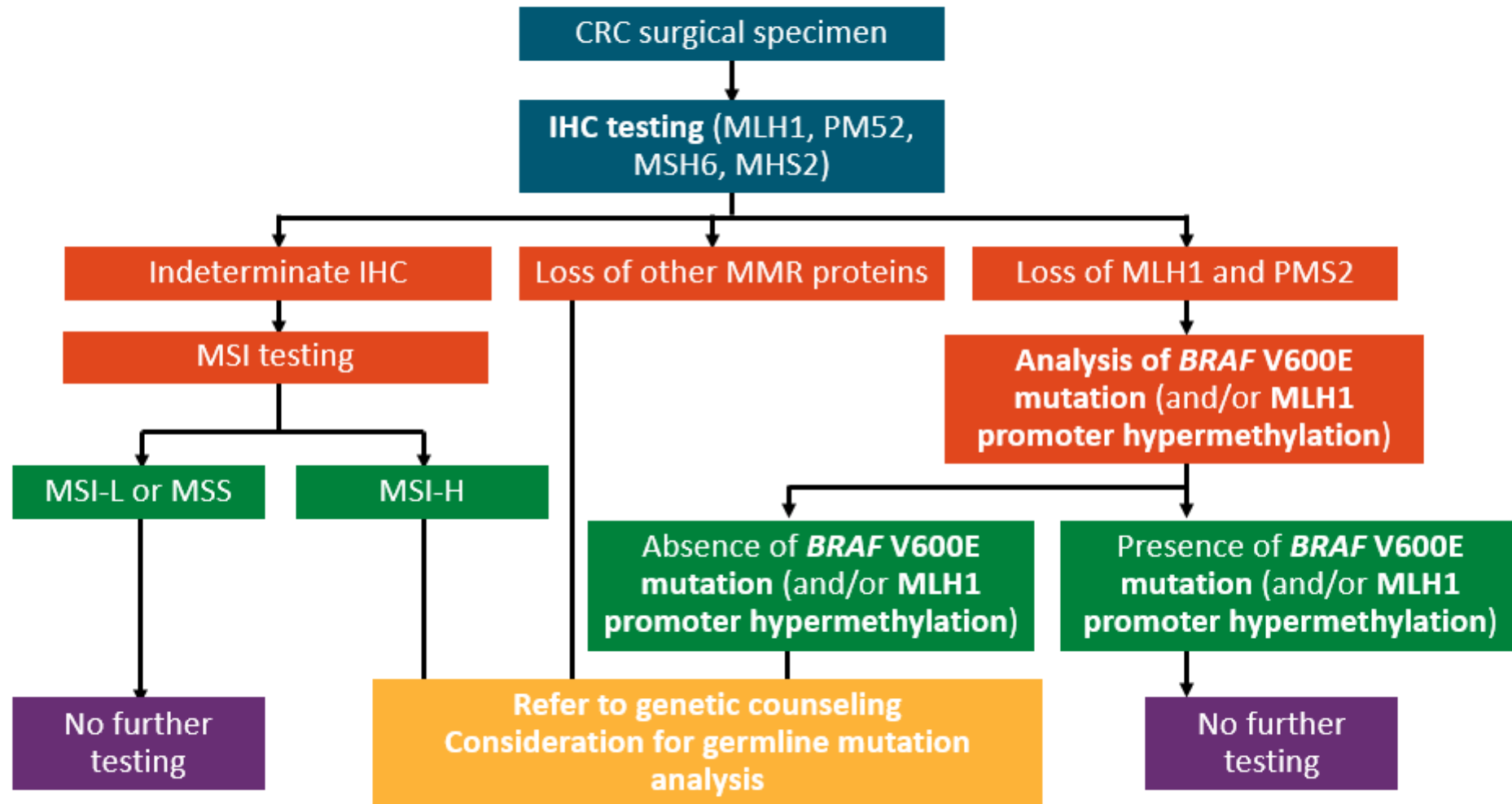
PCR

- Assesses for 5 microsatellite markers and compares them with normal DNA from same patient
 - 2 mononucleotides:** BAT25 and BAT26
 - 3 dinucleotides:** D2S123, D5S346, and D17S250
- Classifies tumor as high (MSI-H), low (MSI-L), or stable (MSS)
 - MSI-H:** ≥ 2 markers show instability
 - MSI-L:** 1 marker shows instability
 - MSS:** no markers show instability

Microsatellite Instability in Cancer



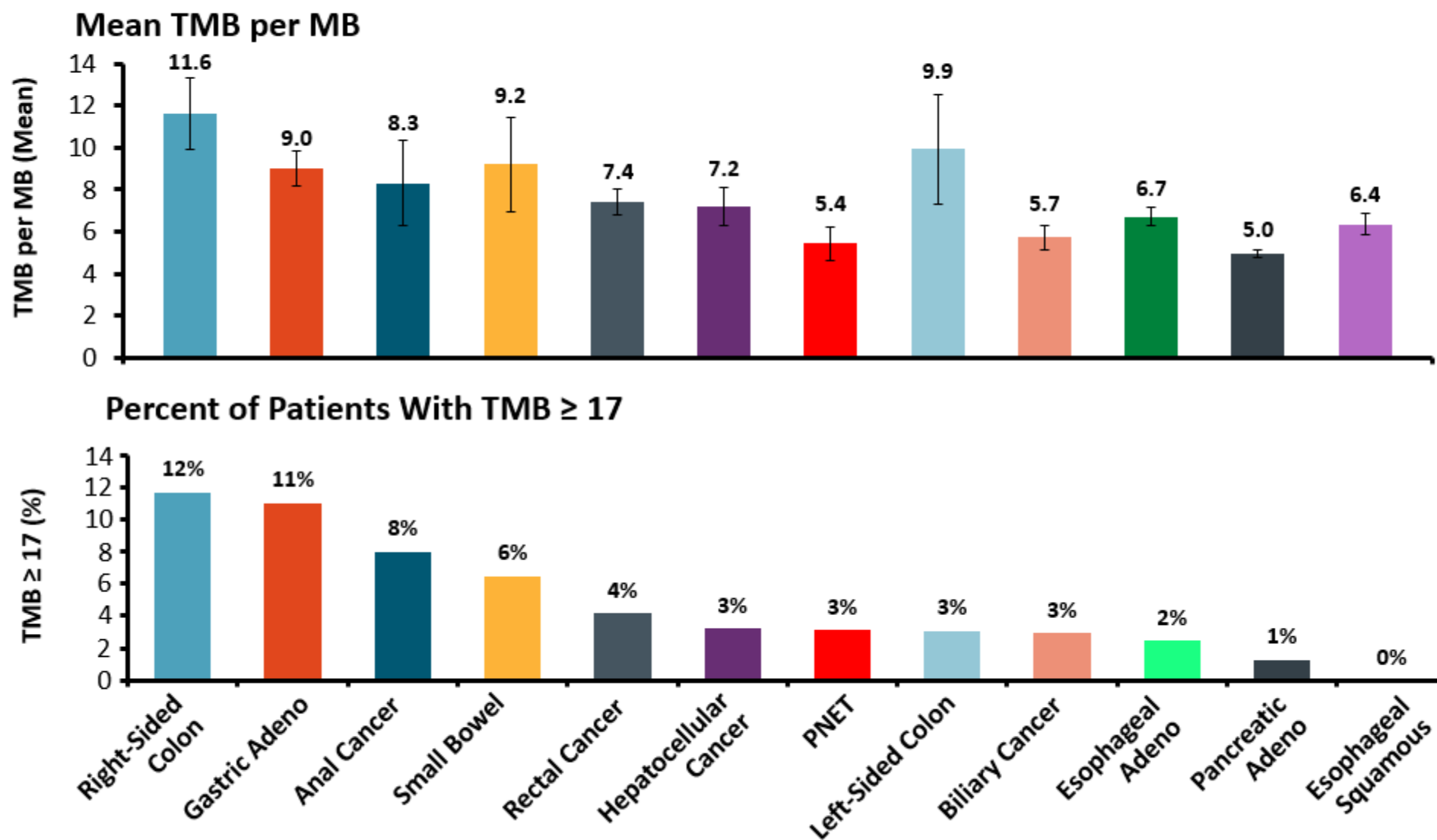
Algorithm for MSI Testing in CRC



Relationship Between MSI and TMB

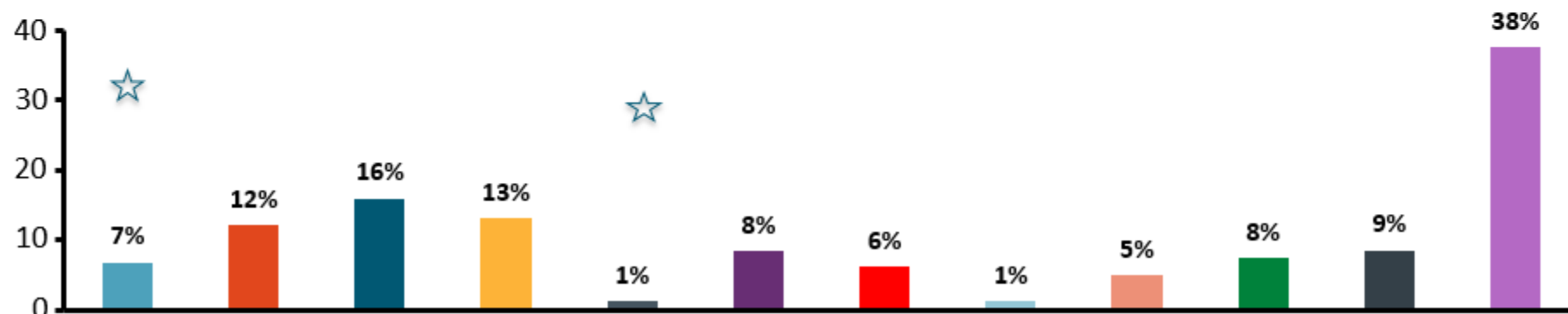


Tumor Mutation Burden Across GI Cancers

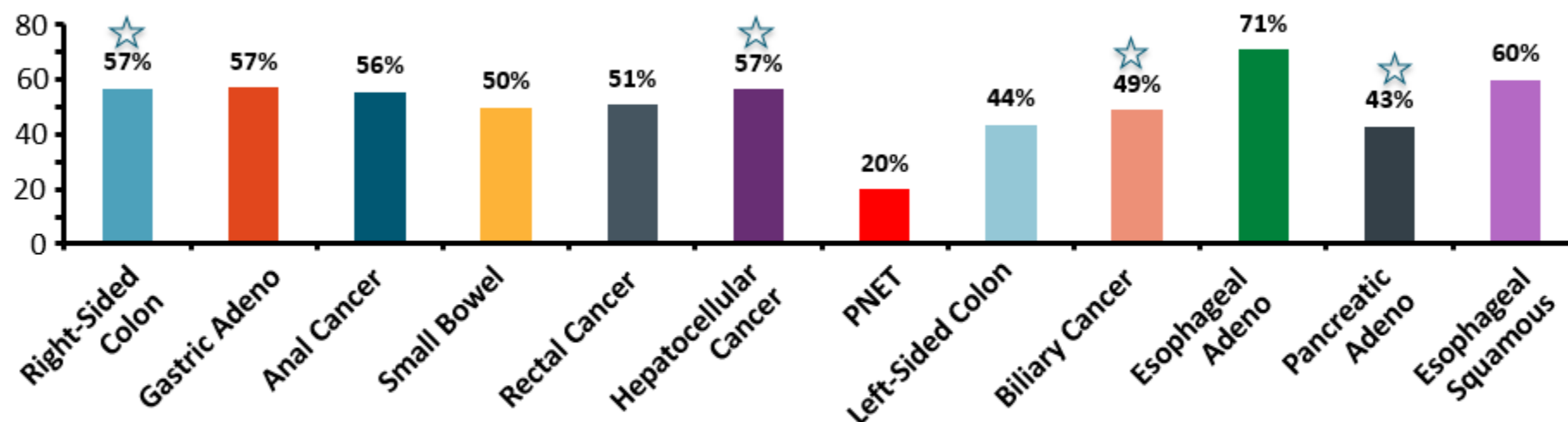


PD-1/PD-L1 Expression Across GI Cancers

PD-L1 Expression on Tumor Cells, %

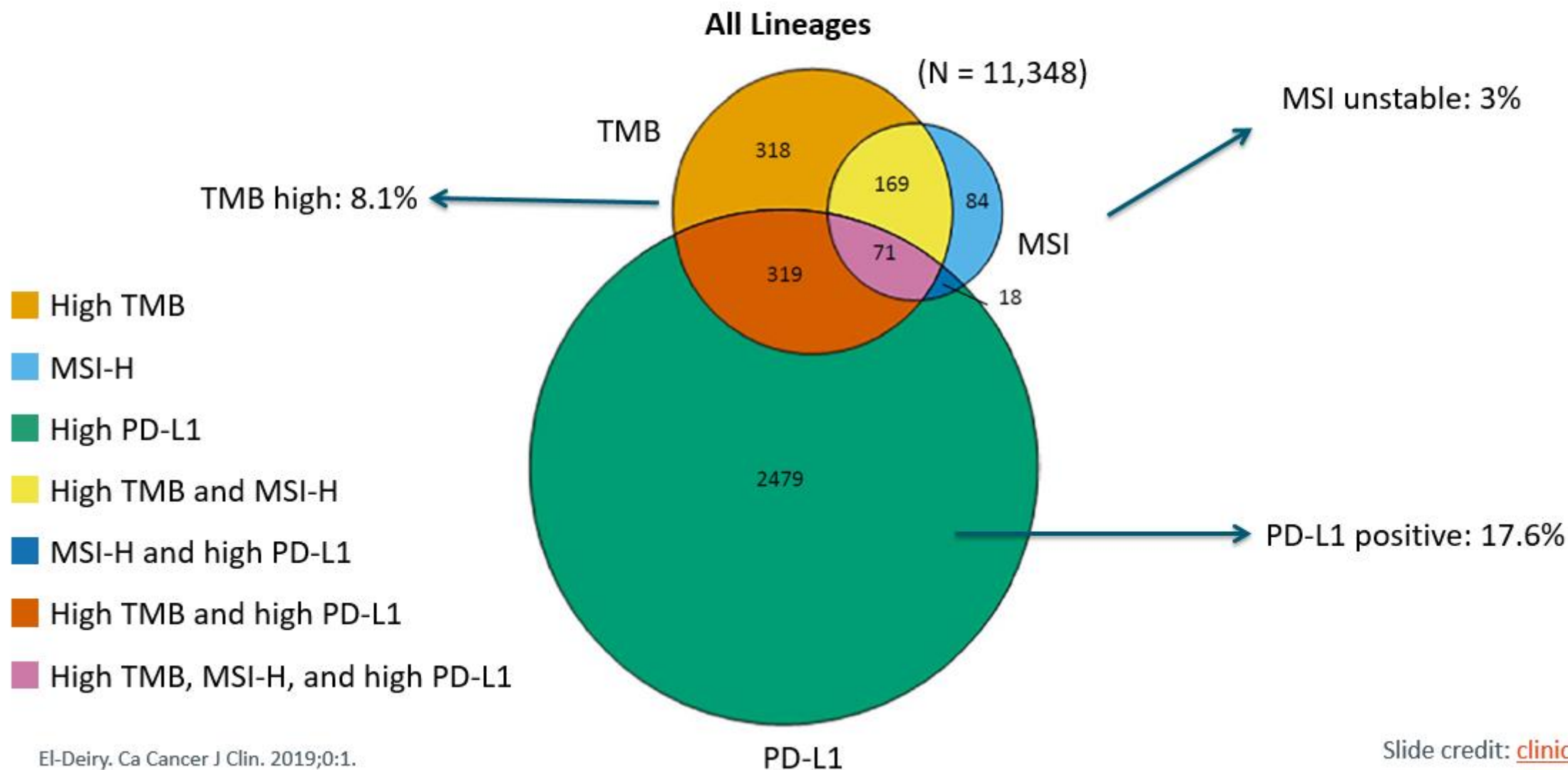


PD-1 Expression on TILs, %



★ = correlation of PD-1/PD-L1 expression with TMB ($P < .05$)

Relationship Between MSI, TMB, and PD-L1



MSI and TMB: Conclusions

- TMB varies significantly across MSI-H tumors
- Loss of expression of the MSH2/MSH6 heterodimer by IHC associated with higher mean TMB than loss of the MLH1/PMS2 heterodimer
- MSI-H CRC carries the highest TMB compared with MSI-H endometrial cancers and others MSI-H solid tumors, consistent with the different prevalence of MSH2/6 protein loss
- These data support gene-specific and disease site-specific heterogeneity among MSI-H tumors, which may partially explain the different responses seen to immune checkpoint inhibitor therapy

Current Indications for Immune Checkpoint Inhibitors in MSI-H/dMMR Cancer

- Pembrolizumab for **MSI-high/mismatch repair–deficient unresectable or metastatic CRC** with PD after a fluoropyrimidine, oxaliplatin, and irinotecan
- Pembrolizumab for **MSI-high/mismatch repair–deficient unresectable or metastatic solid tumors** with PD after prior treatment and no satisfactory alternative treatments
- Nivolumab ± ipilimumab for **MSI-high/mismatch repair–deficient metastatic CRC** with PD after a fluoropyrimidine, oxaliplatin, and irinotecan

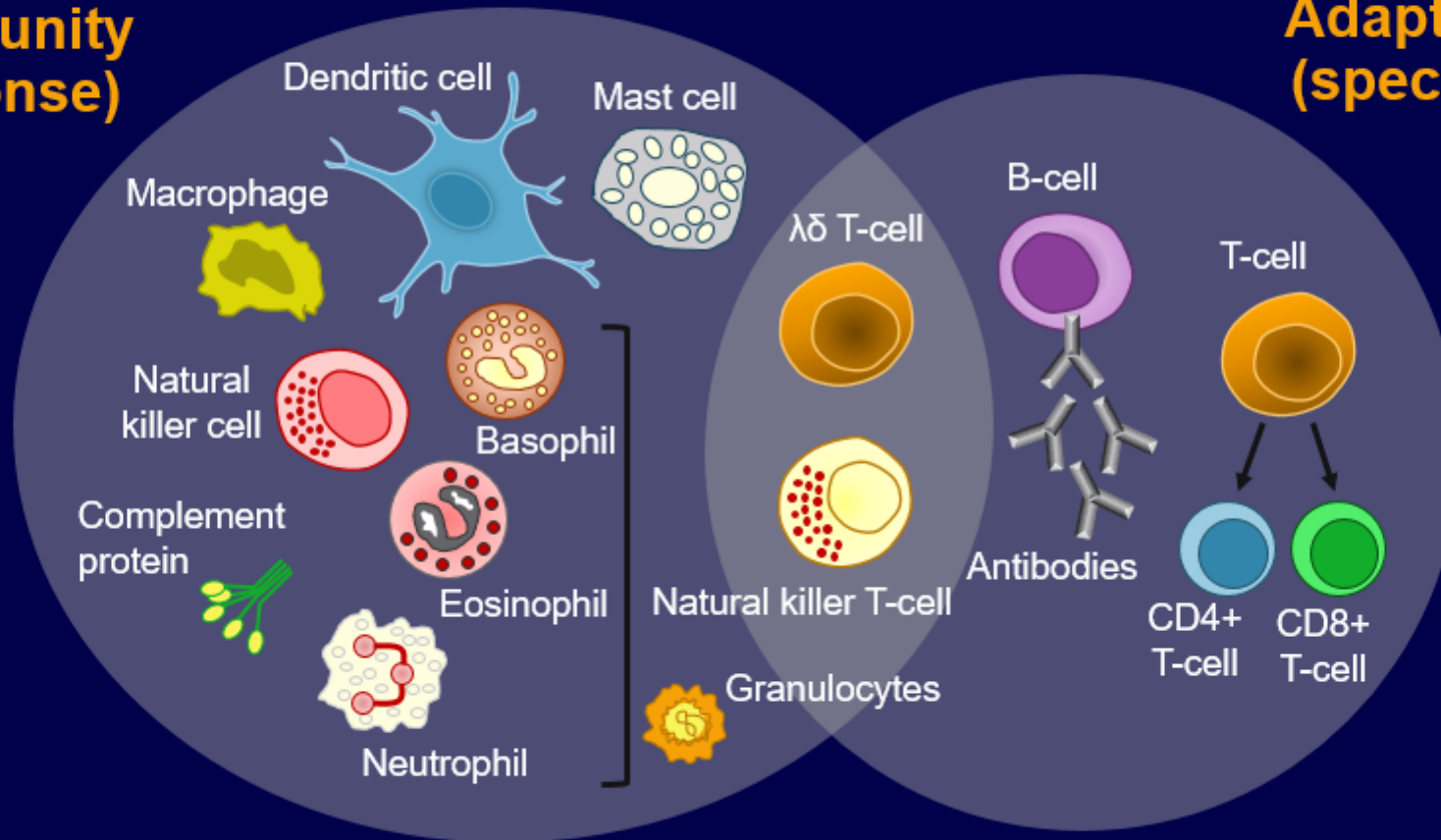
Tumor-Agnostic FDA Indications for Pembrolizumab in Solid Cancers

- Unresectable or metastatic **MSI-H or MMR deficient solid tumors** with PD after previous treatment with no satisfactory alternative treatment options
 - Assessed in several trials, including KEYNOTE-016, -164, -012, -028, -158, and -177
- Unresectable or metastatic **TMB-H (≥ 10 mut/Mb) solid tumors** with PD after previous treatment with no satisfactory alternative treatment options
 - KEYNOTE-158

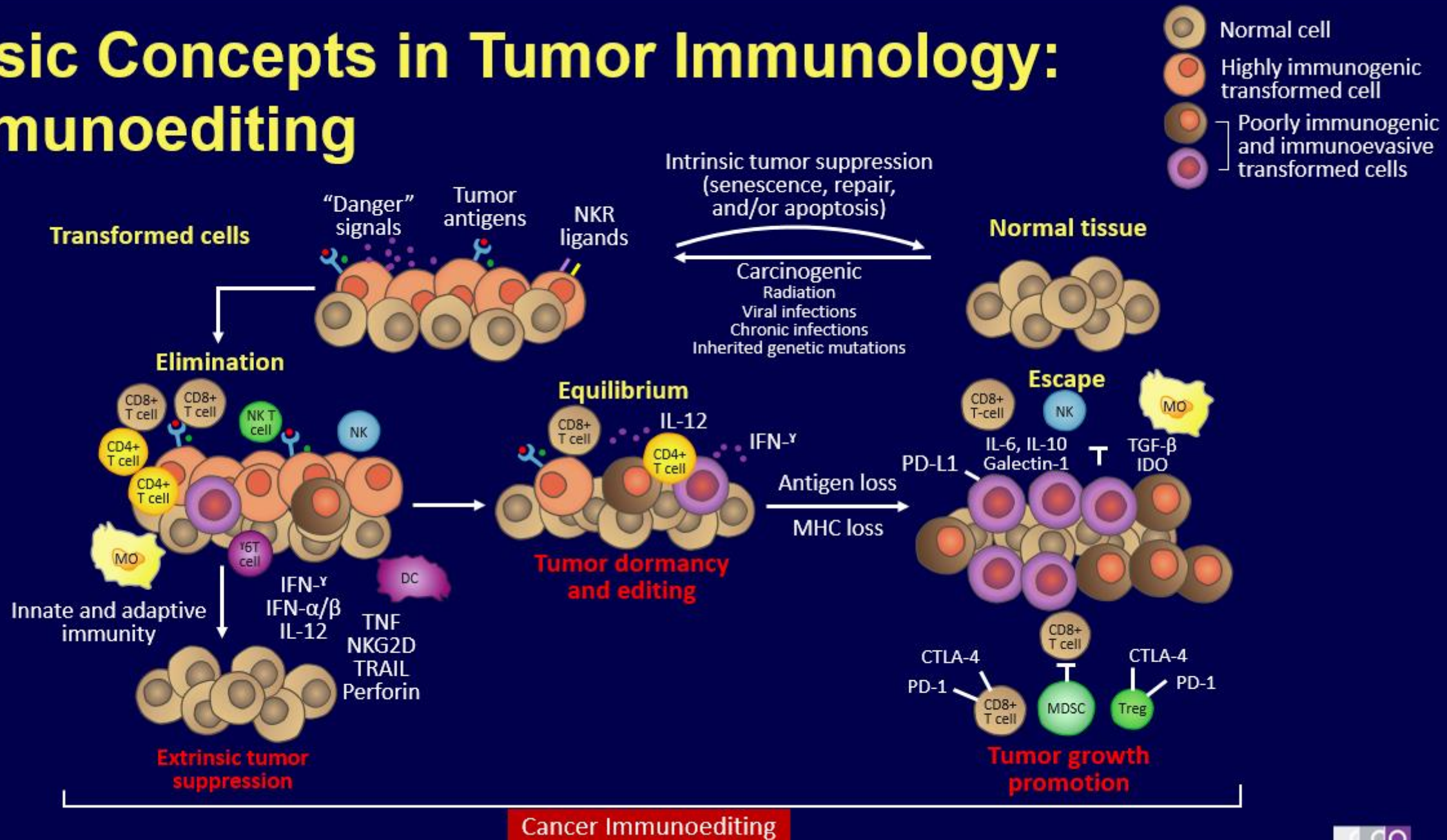
Accelerated approvals.

Immune System Function and Immune Response

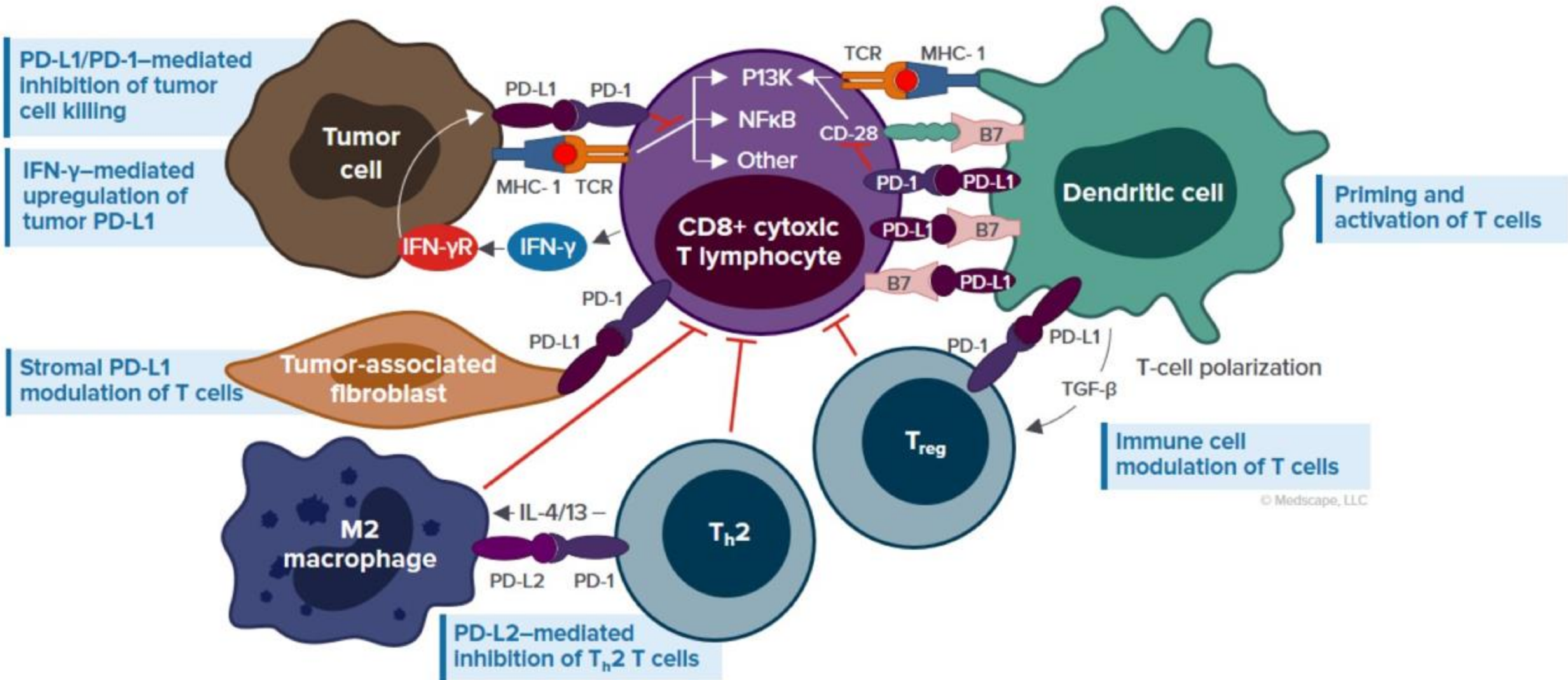
Innate Immunity (fast response)



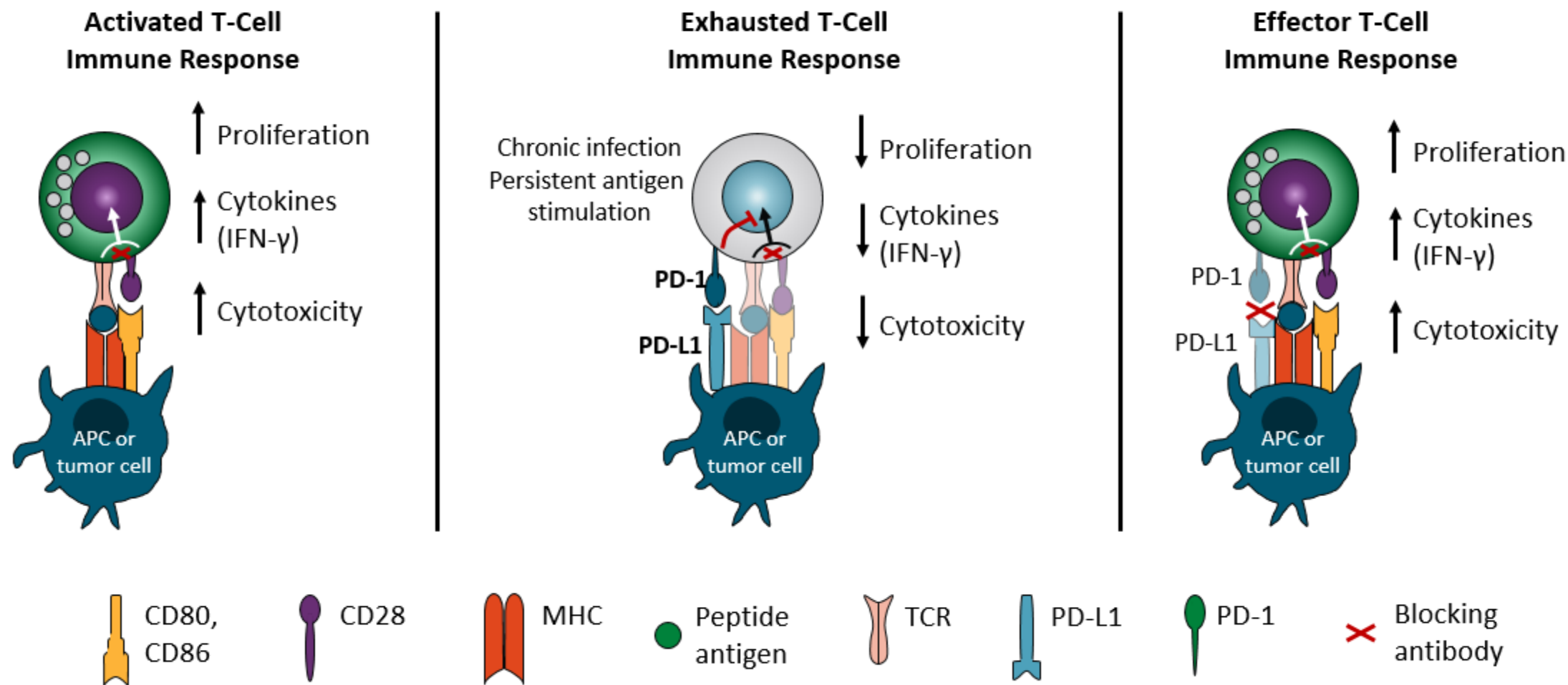
Basic Concepts in Tumor Immunology: Immunoediting



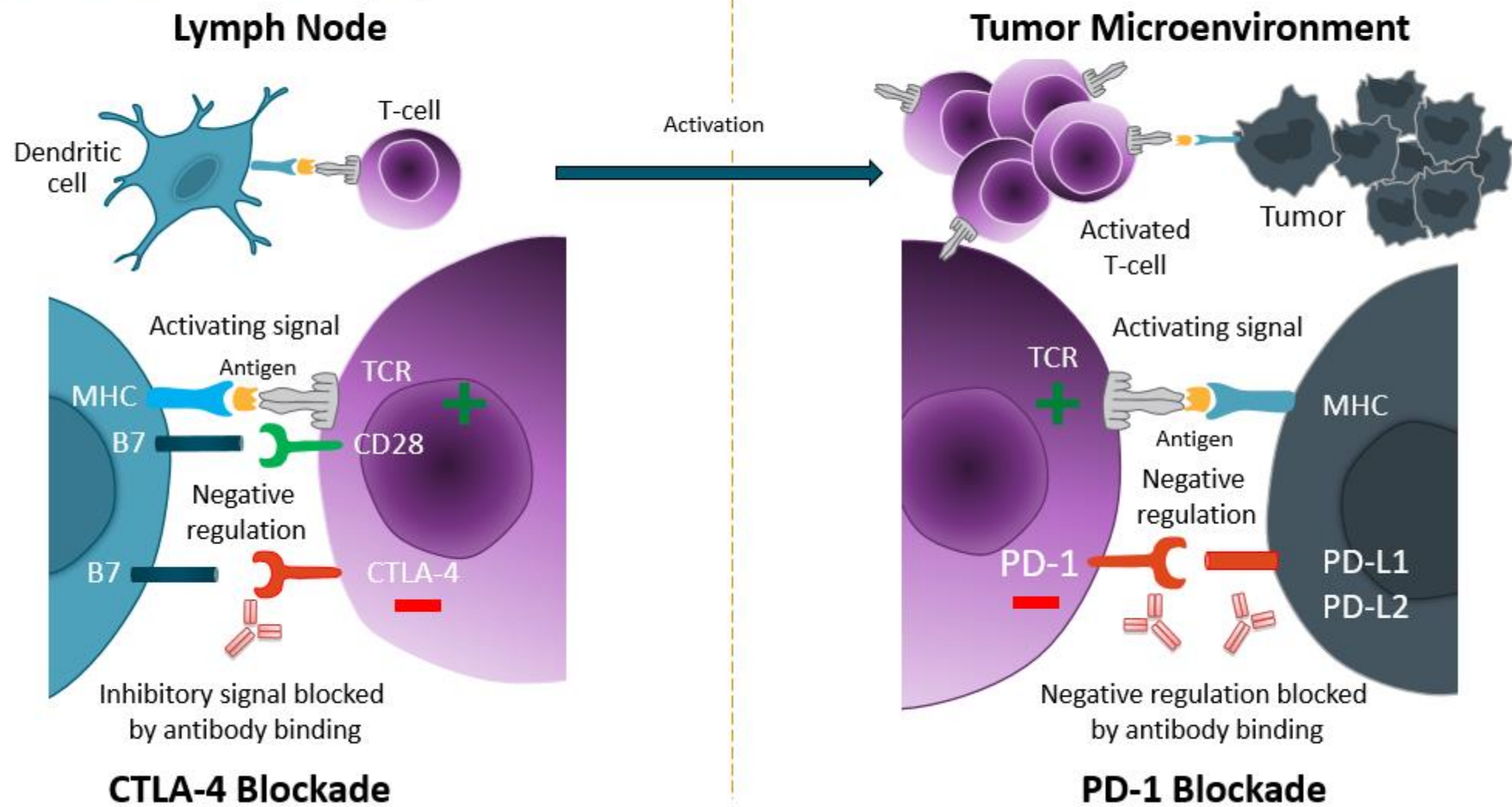
Immunologic Synapses Within Tumor Microenvironment



T-Cell Exhaustion and Reinvigoration



Biologic Rationale for Immune Checkpoint Inhibition as Cancer Therapy

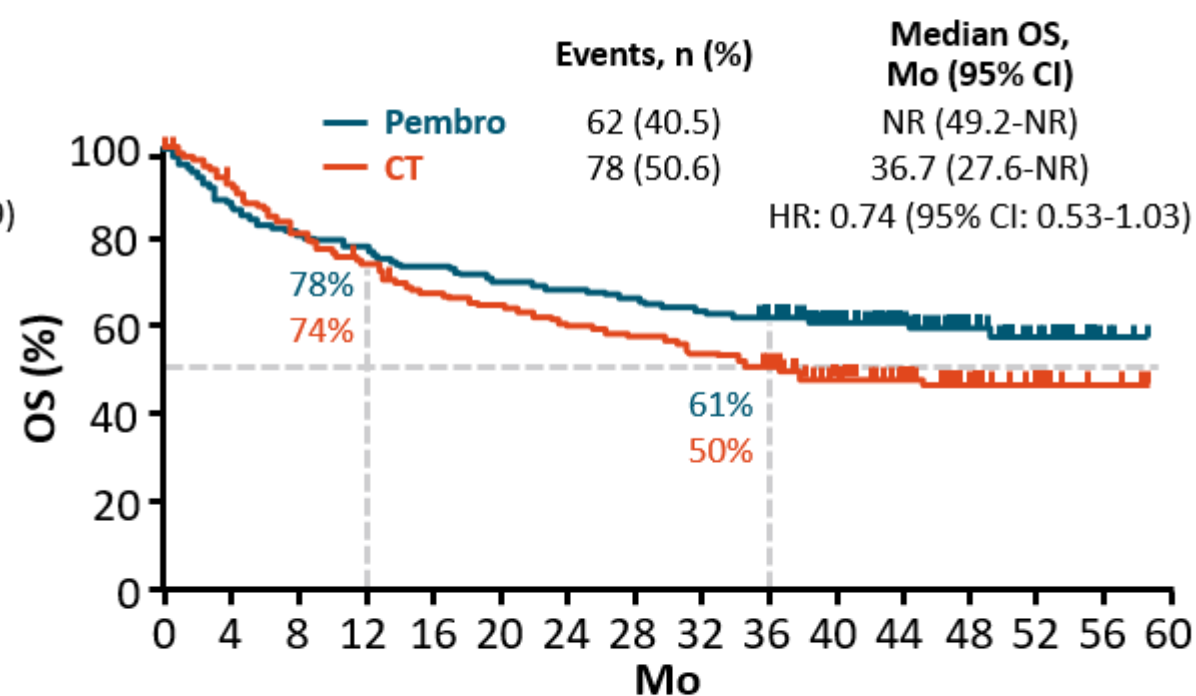
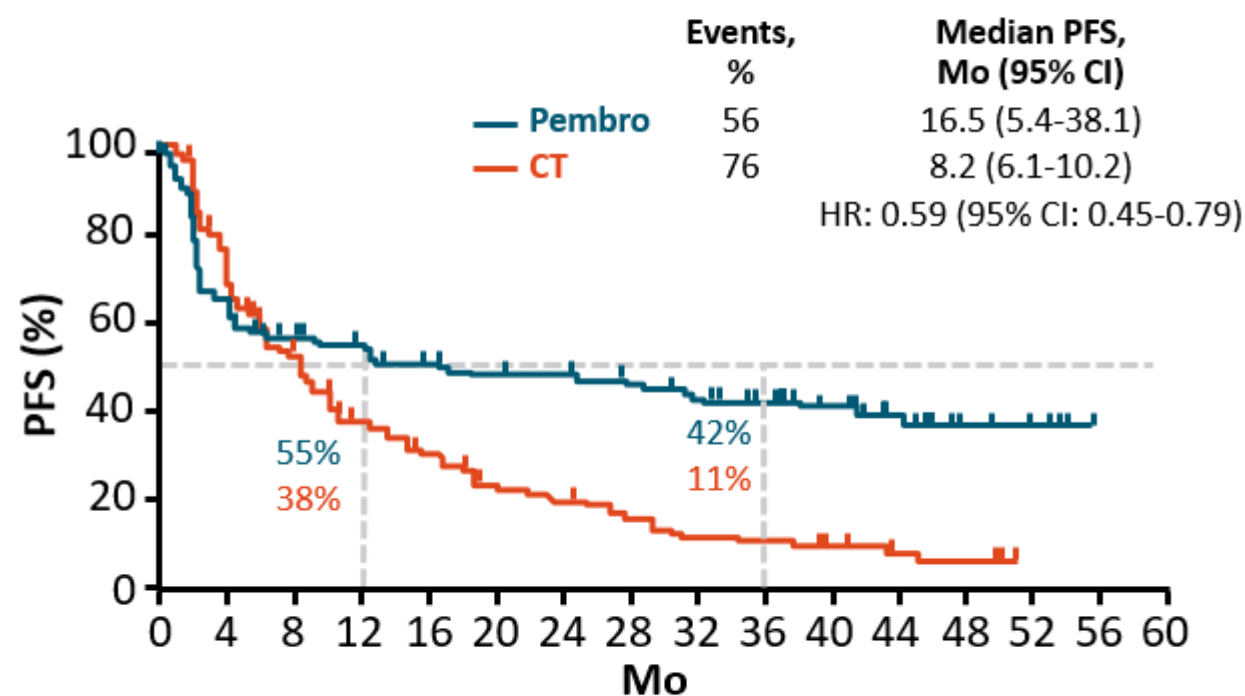


Tumor Agnostic and GI-Specific FDA Indications for MSI-H/dMMR Tumors

Treatment	Indication	Key Supporting Trials
Pembrolizumab	<ul style="list-style-type: none"> Unresectable/metastatic MSI-H or dMMR solid tumors with PD after previous treatment with no satisfactory alternative treatment options Unresectable/metastatic TMB-H (≥ 10 mut/Mb) solid tumors with PD after previous treatment with no satisfactory alternative treatment options Unresectable/metastatic MSI-H/dMMR CRC 	<ul style="list-style-type: none"> KEYNOTE-016, -164, -012, -028, -158, -177 KEYNOTE-158 (ORR: 18%-45% in non-CRC GI cancers) KEYNOTE-177 (see next)
Dostarlimab	<ul style="list-style-type: none"> Unresectable/metastatic dMMR solid tumors with PD after previous treatment with no satisfactory alternative treatment options 	<ul style="list-style-type: none"> GARNET (ORR: 36% in CRC and 43% in non-CRC GI cancers)
Nivolumab \pm ipilimumab	<ul style="list-style-type: none"> Metastatic MSI-H/dMMR CRC with PD after fluoropyrimidine, oxaliplatin, and irinotecan 	<ul style="list-style-type: none"> CheckMate-142 (ORR: 55% w/ nivo + ipi, 31% w/ nivo in recurrent CRC; 69% with nivo + ipi in mCRC with no prior treatment)

KEYNOTE-177: First-line Pembrolizumab vs Chemotherapy in MSI-H/dMMR Metastatic CRC

- Randomized, open-label phase III study of pembrolizumab vs CT* for patients with treatment-naïve MSI-H/dMMR mCRC (N = 307)



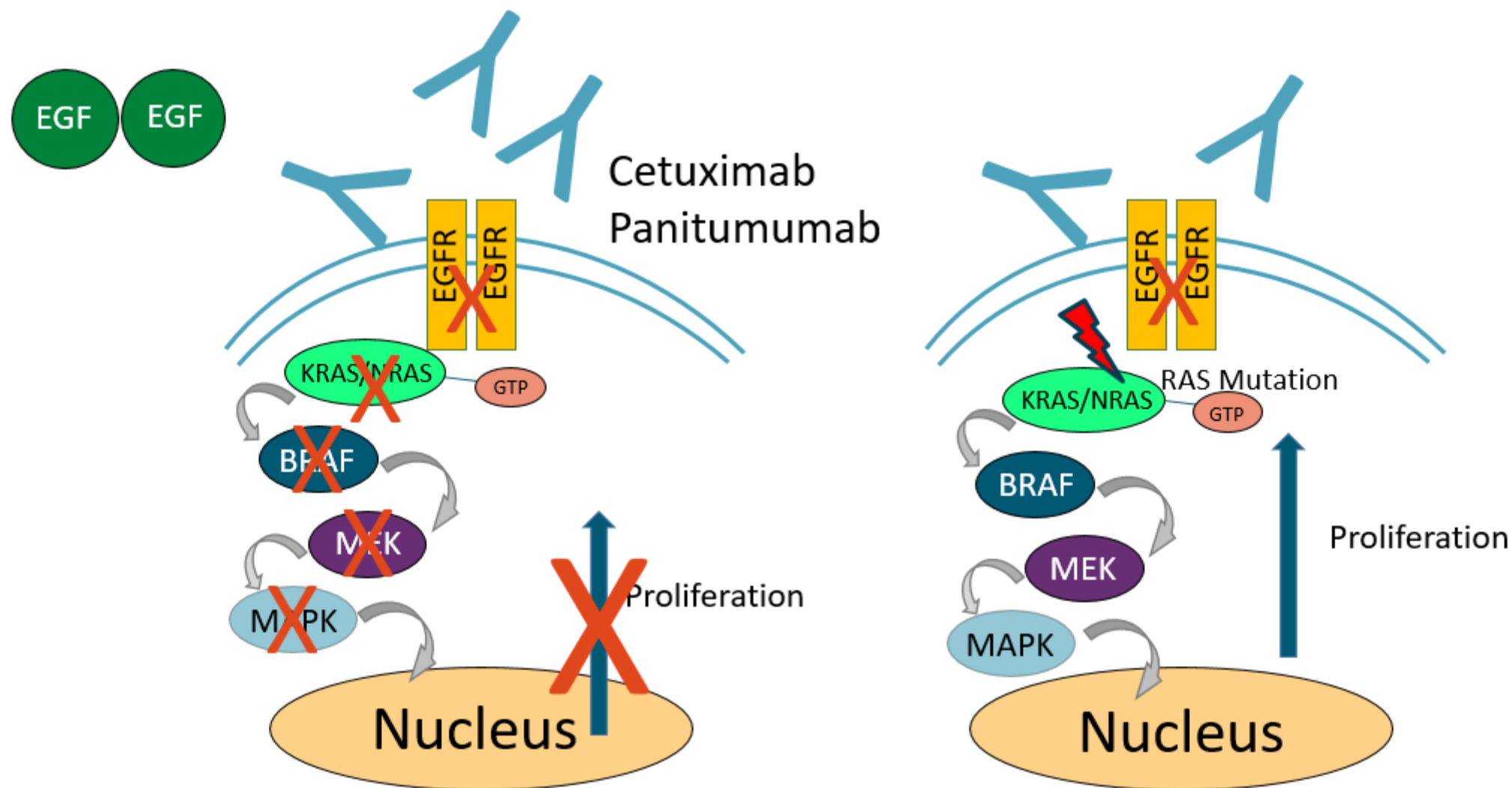
- ORR: pembrolizumab, 45%; CT, 33%

*mFOLFOX-6 ± bevacizumab or cetuximab or FOLFIRI ± bevacizumab or cetuximab.

***RAS* and *BRAF* Mutations: Impact on Therapy**



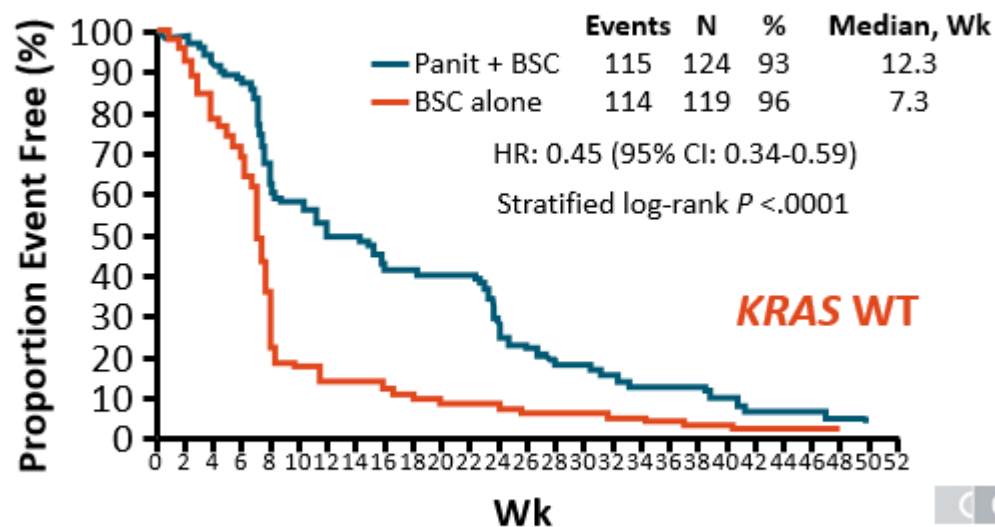
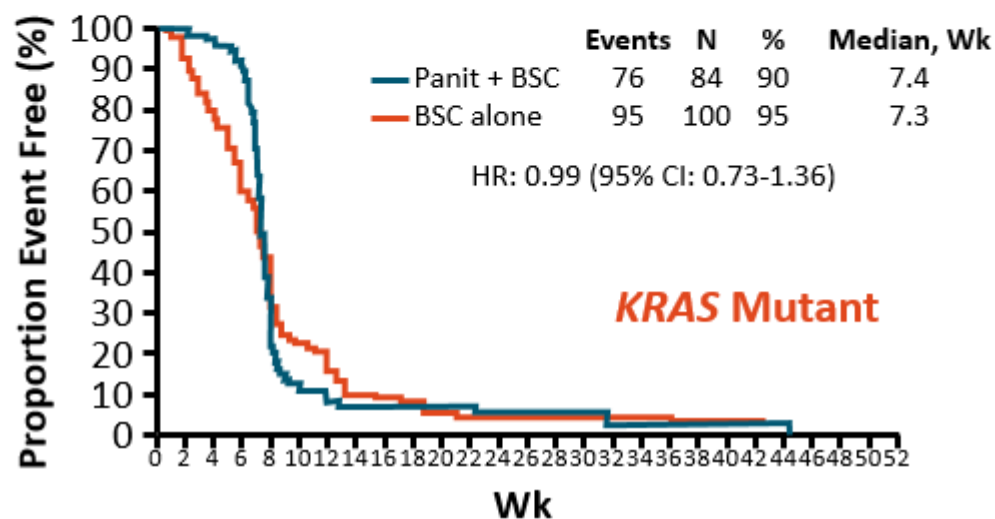
EGFR Inhibitors and Key Mutations in CRC



RAS Mutations (*KRAS*, *NRAS*, *HRAS*)

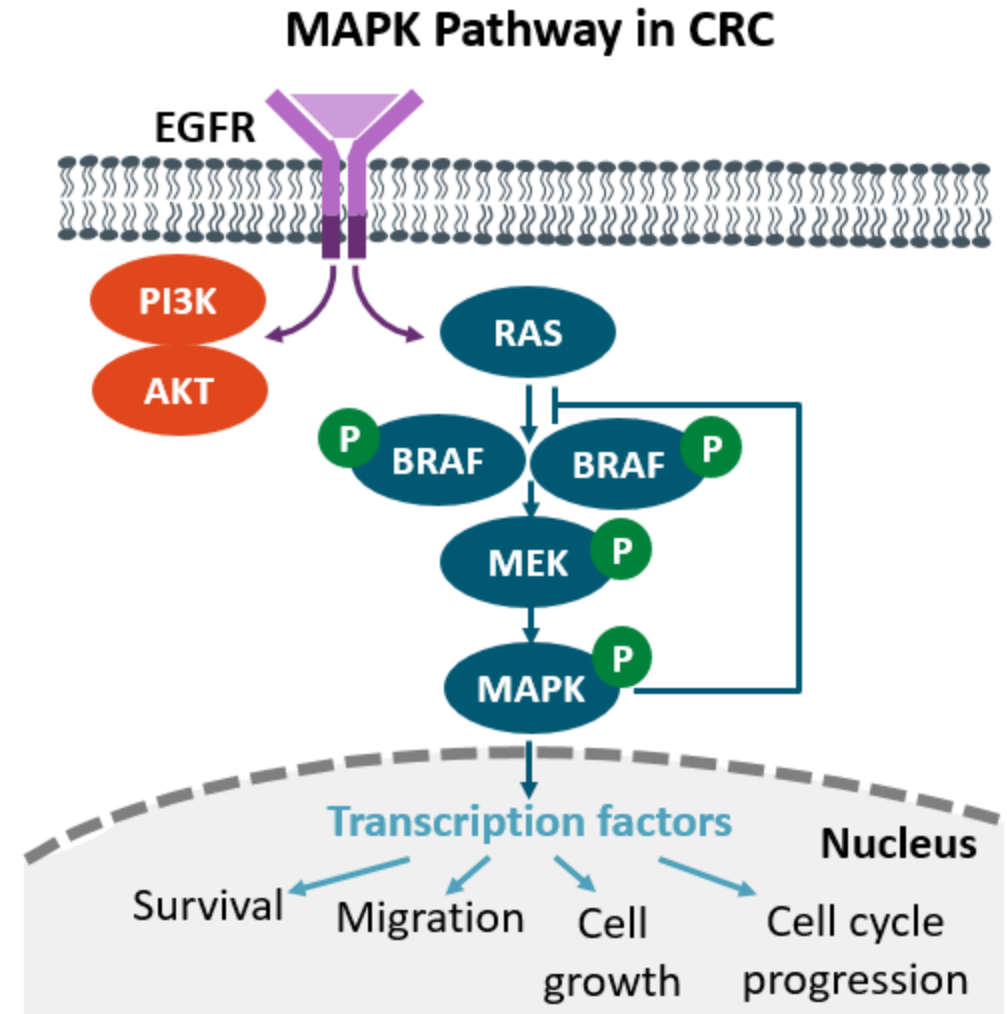
- Most frequently mutated oncogenes¹
 - 90% of pancreatic cancers, **45% of colon cancers**, 35% of lung cancers
 - *KRAS* most prevalent in these tumor types
- In CRC, RAS testing is required prior to anti-EGFR therapy (eg, cetuximab or panitumumab)
 - Patients with *KRAS* and *NRAS* mutations should not be treated with anti-EGFR therapy²⁻⁴
 - *HRAS* mutations are much less common (1.7%) but likely have the same negative predictive value

Panitumumab + BSC vs BSC¹



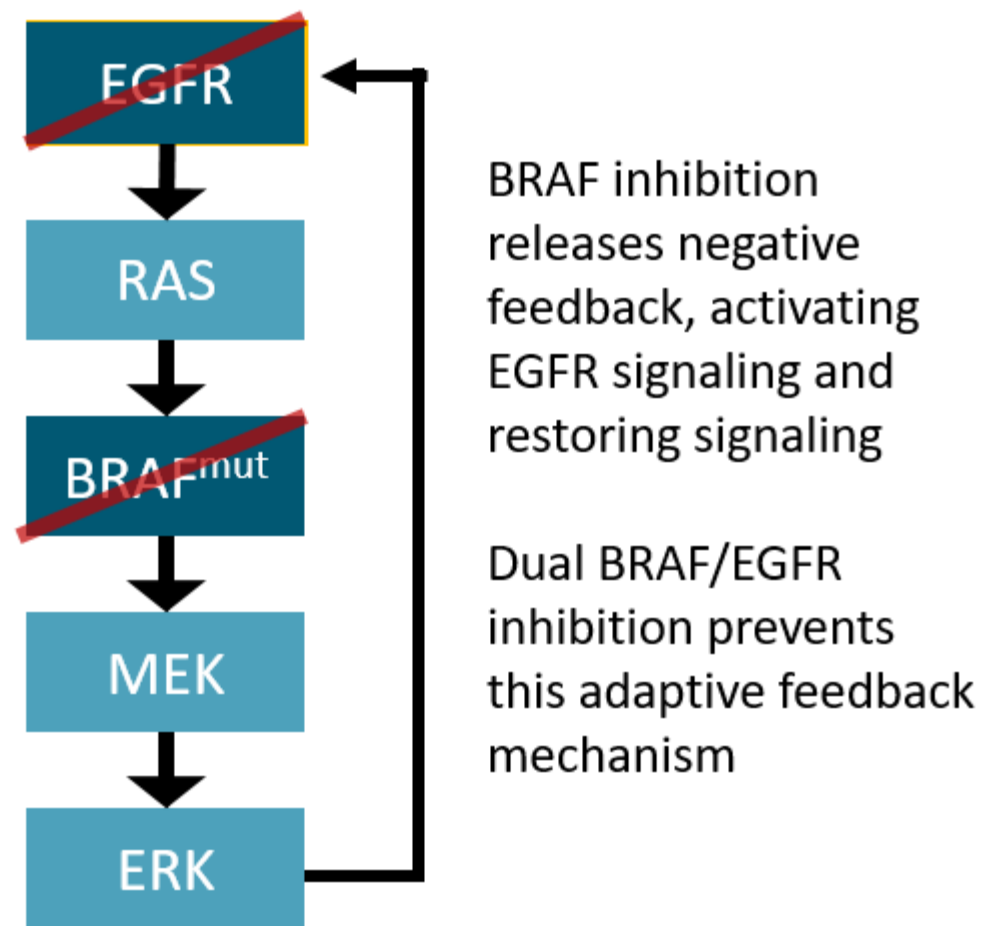
BRAF Mutations in CRC

- BRAF: primary effector of RAS signaling
- *BRAF* mutations
 - Occur most frequently in exon 15 (V600E)
 - Accounts for majority of *BRAF* mutations
 - **Found in ~10% of patients with CRC**
 - Mutually exclusive with *RAS* mutations



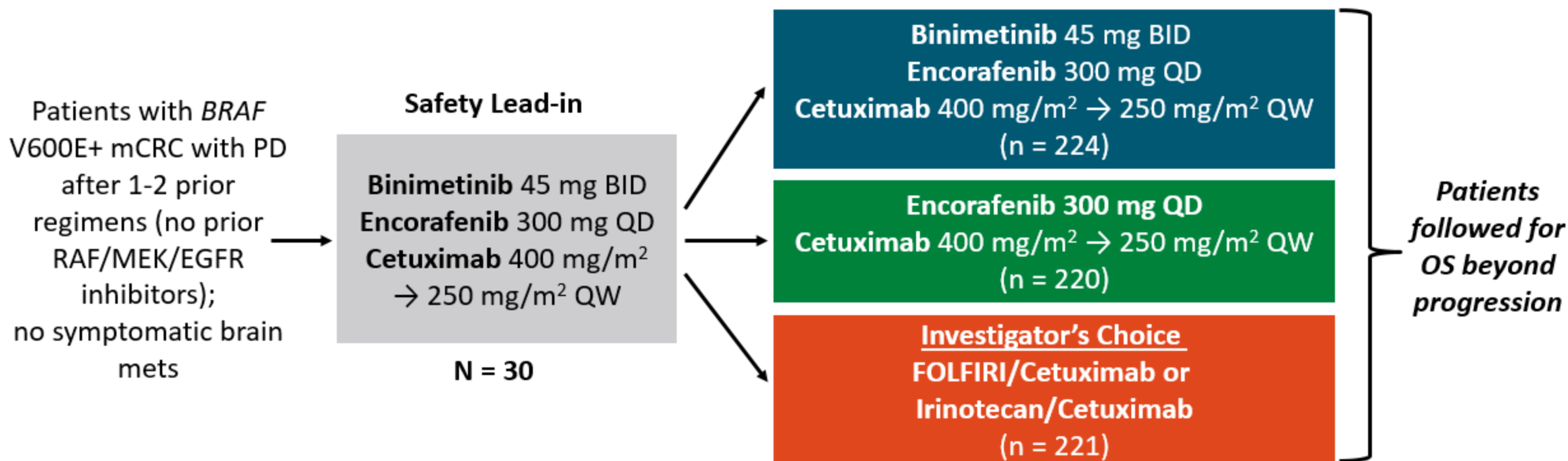
RAS and BRAF Mutations

- *KRAS*, *NRAS*, *HRAS* most frequently mutated oncogenes¹
 - 90% of pancreatic cancers, 45% of colon cancers
 - *KRAS* most prevalent in these tumor types
 - In CRC, RAS testing is required before anti-EGFR therapy; patients with *KRAS* and *NRAS* mutations should not receive anti-EGFR therapy²⁻⁴
- *BRAF* mutations occur most frequently in exon 15 (V600E)
 - Found in ~10% of patients with CRC



BEACON CRC: Encorafenib + Cetuximab ± Binimetinib for *BRAF* V600E–Mutant mCRC

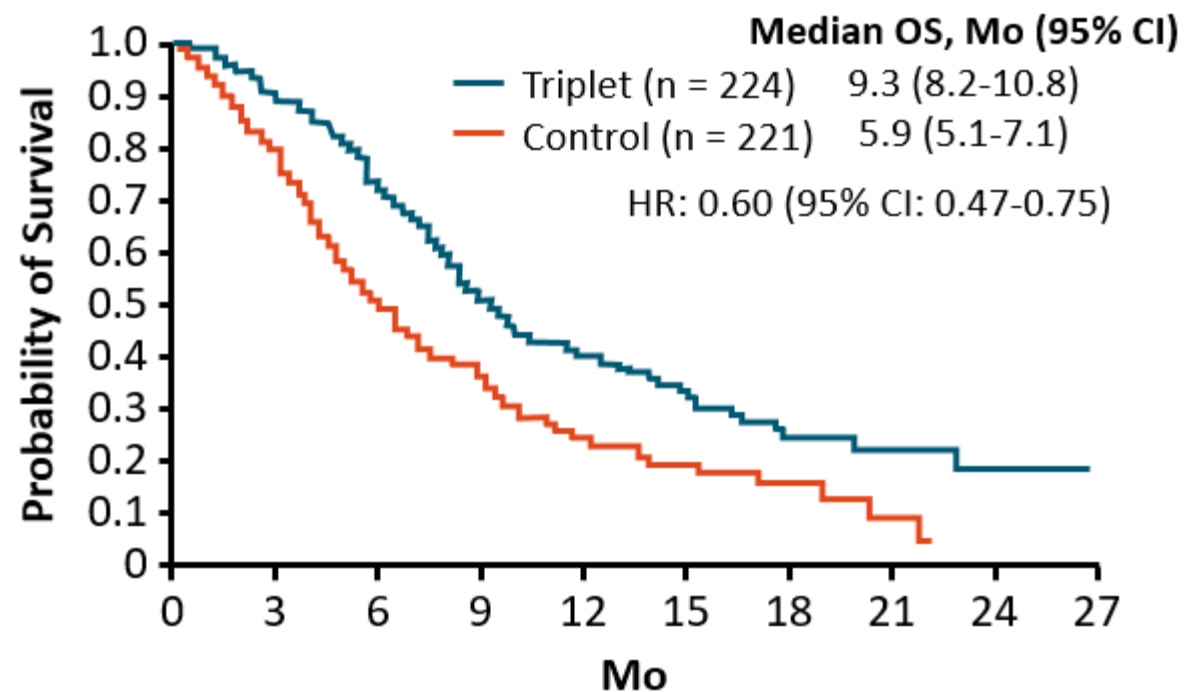
- A multicenter, randomized, open-label, 3-arm phase III trial



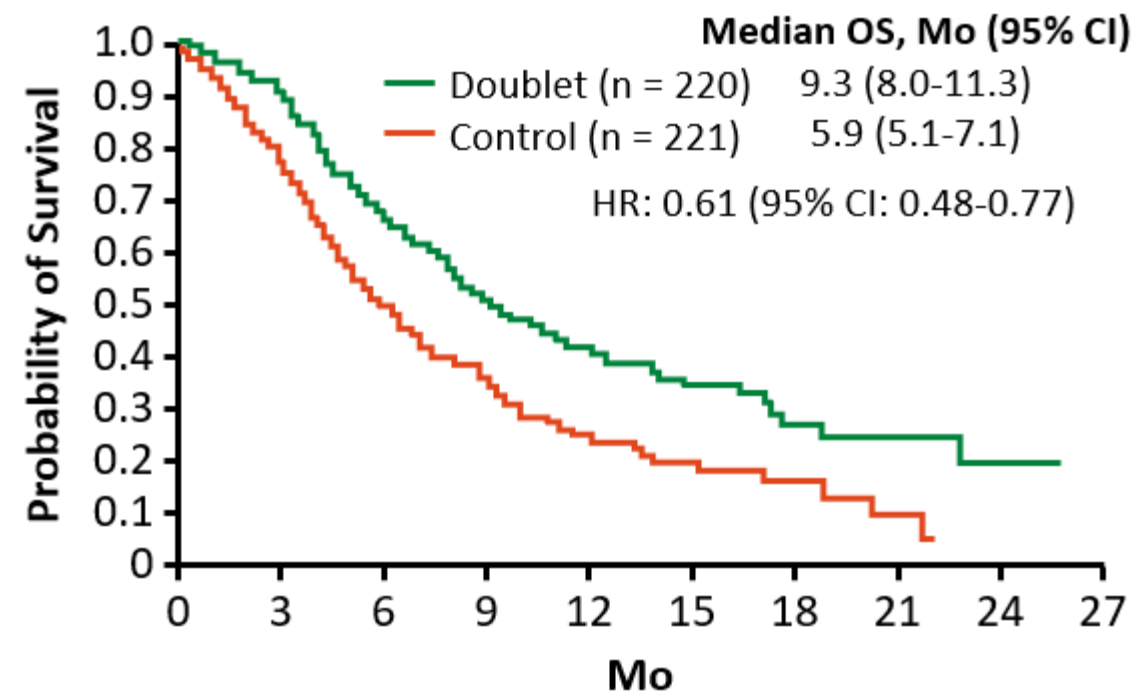
- Primary endpoints: OS and ORR for triplet vs control; secondary endpoints: OS and ORR for doublet vs control, triplet vs doublet; PFS; safety

BEACON CRC: OS and ORR

Triplet vs Control (Primary Endpoint)



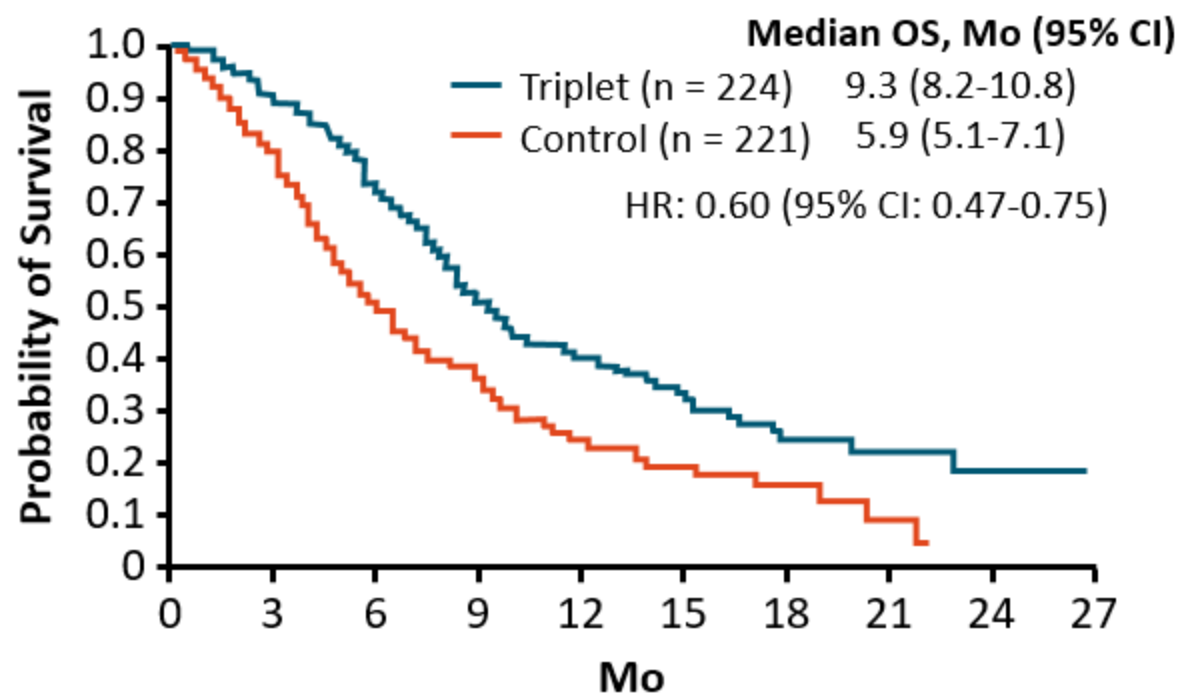
Doublet vs Control



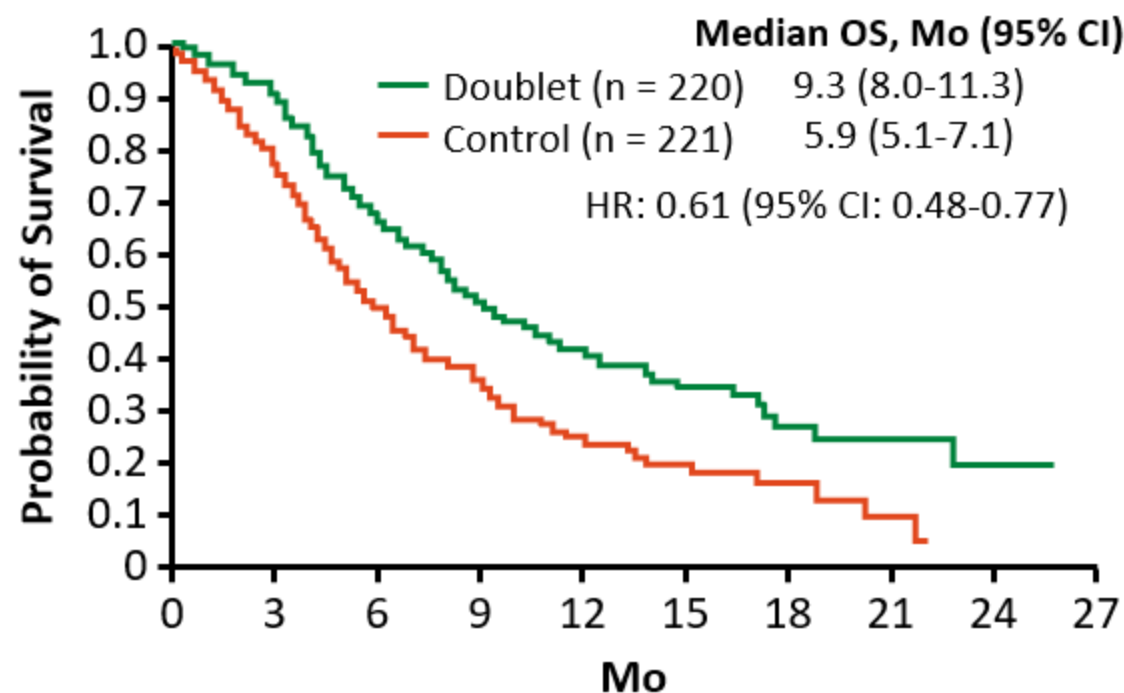
Confirmed Response by BICR	Triplet Regimen (n = 224)	Doublet Regimen (n = 220)	Control (n = 221)
ORR, % (95% CI)	27 (21-33)	20 (15-25)	2 (<1-5)
P value (vs control)	< .0001	< .0001	

BEACON CRC: OS and ORR

Triplet vs Control (Primary Endpoint)



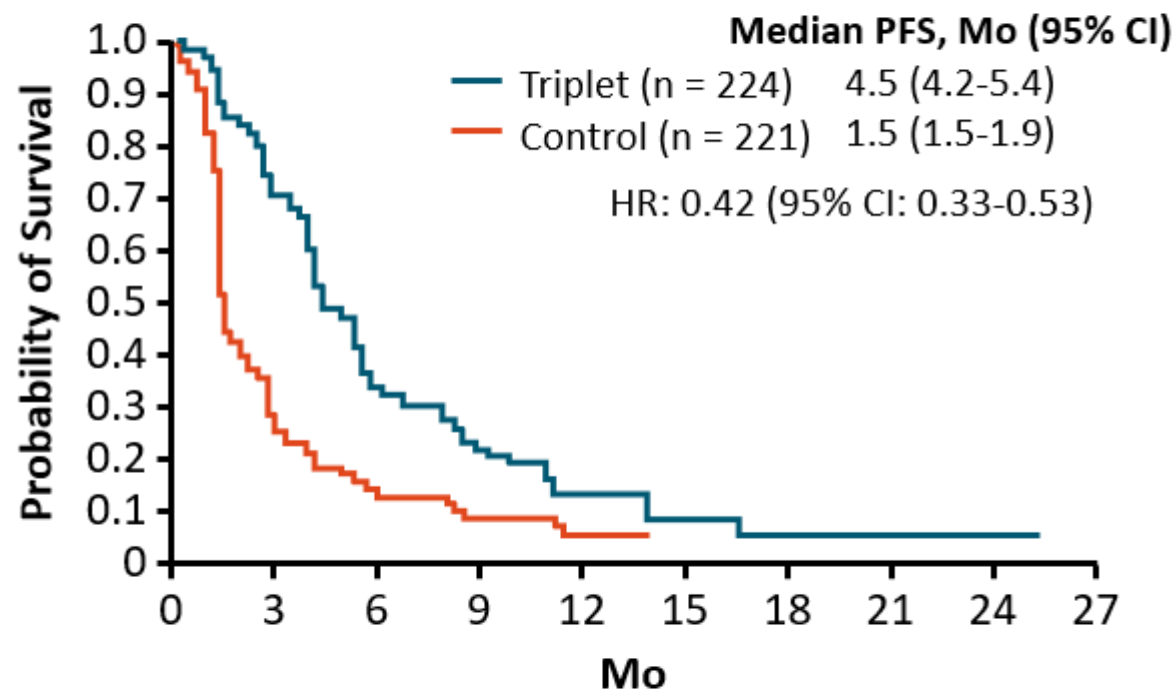
Doublet vs Control



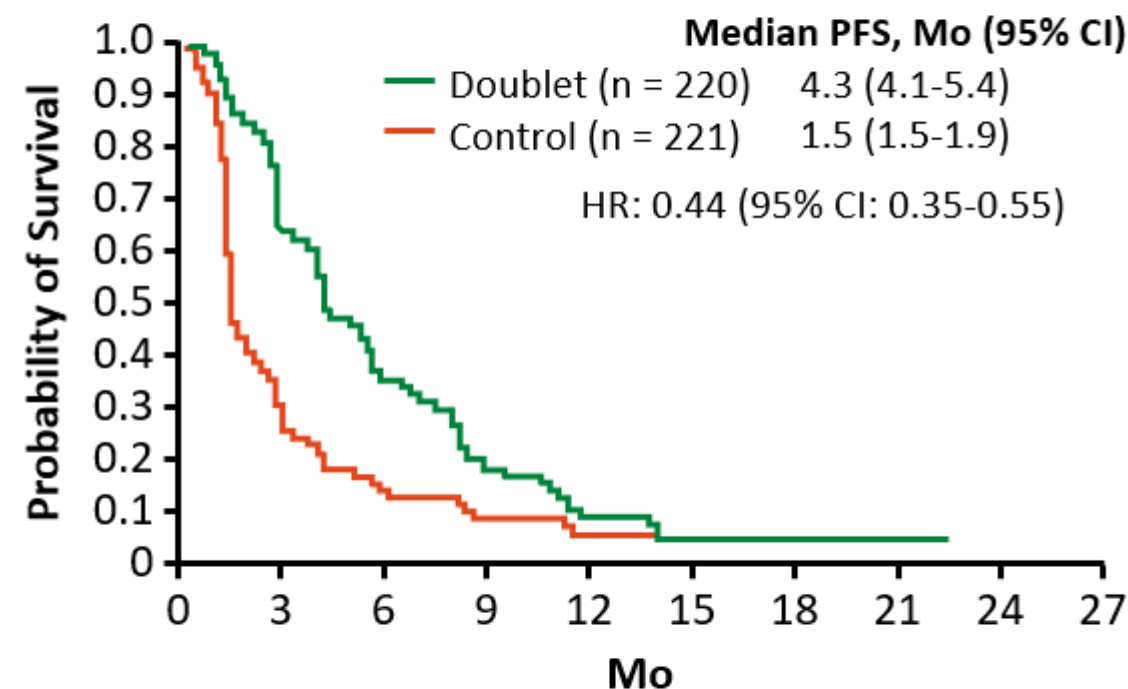
- FDA indication: encorafenib + cetuximab for *BRAF* V600E–mutated mCRC after previous systemic therapy

BEACON CRC: PFS (BICR)

Triplet vs Control (Primary Endpoint)



Doublet vs Control



- FDA/EMA indication: encorafenib + cetuximab for *BRAF* V600E-mutated mCRC after previous systemic therapy

Ongoing Trials and Select Recent Data With BRAF and KRAS G12C Inhibitors in Advanced CRC

Regimen	Phase	Treatment	Population	Key Findings
Recent Data				
KRYSTAL-1 (NCT03785249)	I/II	Adagrasib ± cetuximab	Advanced <i>KRAS</i> G12C CRC, previous therapy	<ul style="list-style-type: none"> ▪ A: ORR, 22%; DCR, 87% (n = 45) ▪ A + C: ORR, 43%; DCR, 100% (n = 28) ▪ Ongoing
Ongoing				
KRYSTAL-10 (NCT04793958)	III	Adagrasib + cetuximab vs CT	Advanced <i>KRAS</i> G12C CRC, previous 1L therapy	<ul style="list-style-type: none"> ▪ Ongoing
BREAKWATER (NCT04607421)	III	Encorafenib + cetuximab ± CT vs CT	<i>BRAF</i> V600E–mutant mCRC, no previous therapy	<ul style="list-style-type: none"> ▪ Ongoing

Additional Biomarkers: *HER2* Amplification



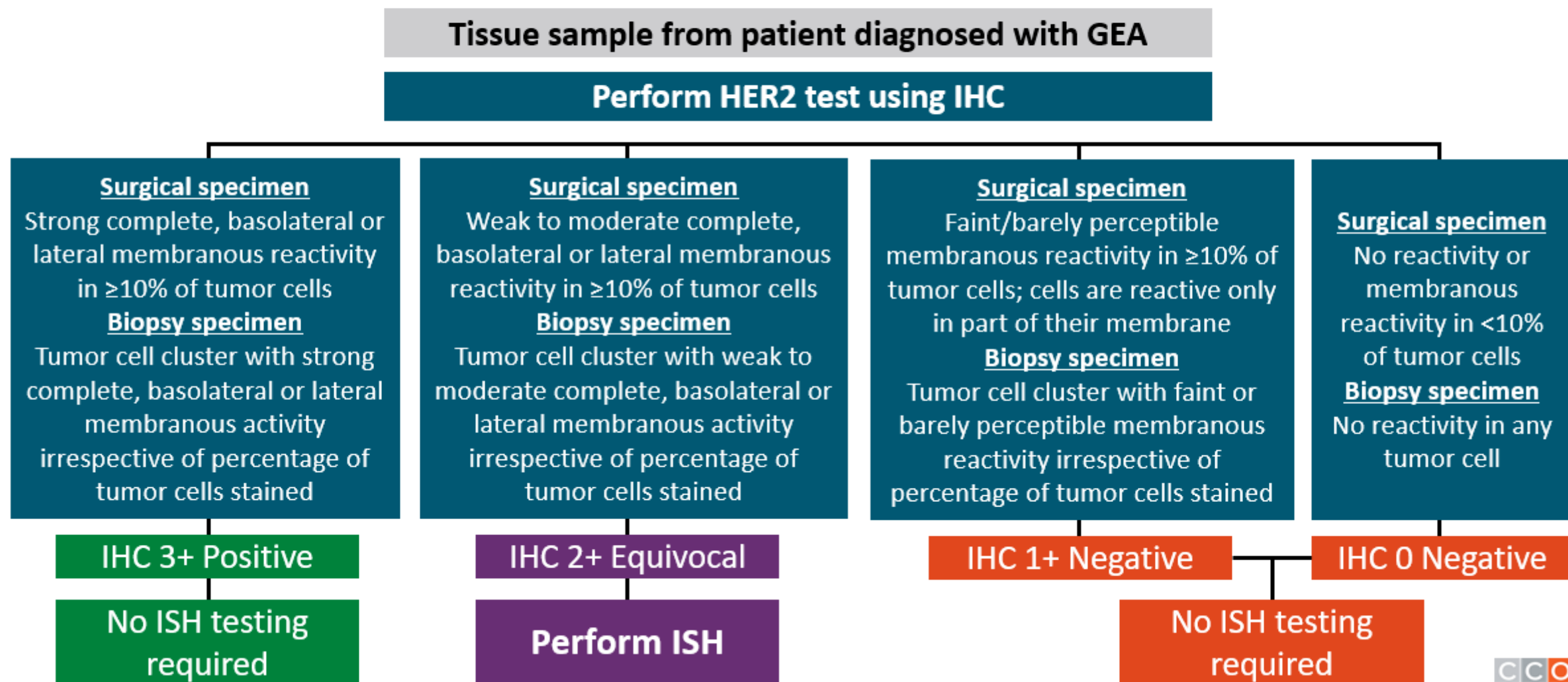
Genomic Alterations in *HER2/ERBB2* in Breast, Gastroesophageal, and Colorectal Cancers

Alterations, %	Breast	Gastro-esophageal	Colorectal
<i>HER2</i> amplification	18	16	2-5
<i>HER2</i> KD mutations	2	2	2
<i>HER2</i> ECD mutations	<1	--	<1

- Mutually exclusive with *RAS/BRAF* mutations
- *HER2* amplification enriched in *KRAS*, *NRAS*, *BRAF*, and *PIK3CA* WT tumors

Gastroesophageal Adenocarcinoma

Algorithm for HER2 Testing by IHC

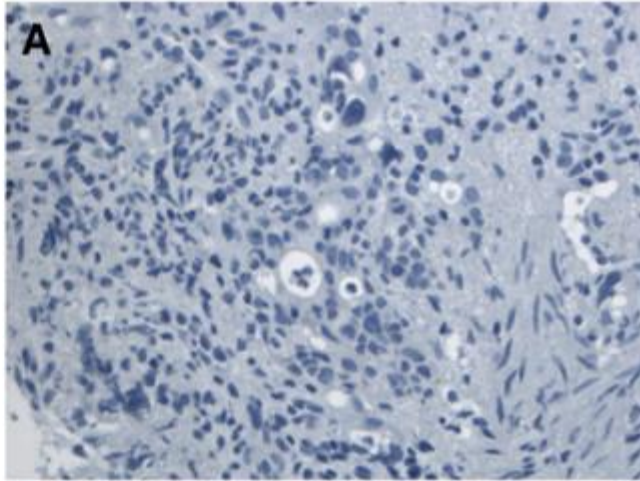


ASCO/CAP Guideline for HER2 Testing

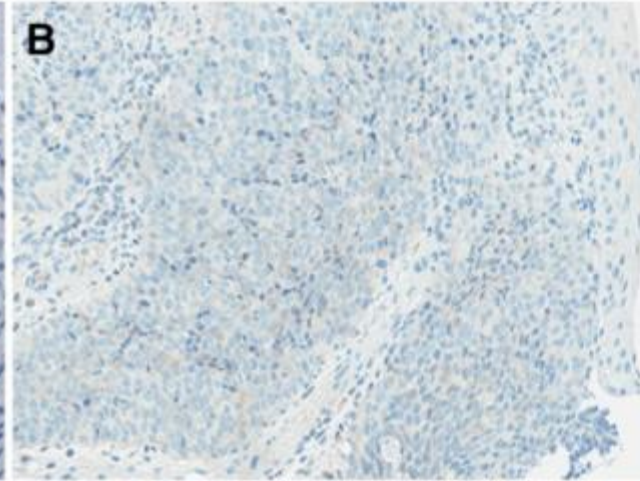
Group 1	Group 2	Group 3	Group 4	Group 5
Ratio ≥ 2	Ratio ≥ 2	Ratio < 2	Ratio ≥ 2	Ratio < 2
≥ 4 HER2 signals/cell	< 4 HER2 signals/cell	≥ 6 HER2 signals/cell	≥ 4 and < 6 HER2 signals/cell	≥ 4 HER2 signals/cell
Dual Probe Assay				
HER2 Positive		HER2 Negative		
Group 1		Group 2 AND concurrent IHC 0-1+ or 2+		
Group 2 AND concurrent IHC 3+		Group 3 AND concurrent IHC 0-1+		
Group 3 AND concurrent IHC 2+ or 3+		Group 4 AND concurrent IHC 0-1+ or 2+		
Group 4 AND concurrent IHC 3+		Group 5		

HER2 IHC in Representative Gastroesophageal Adenocarcinomas

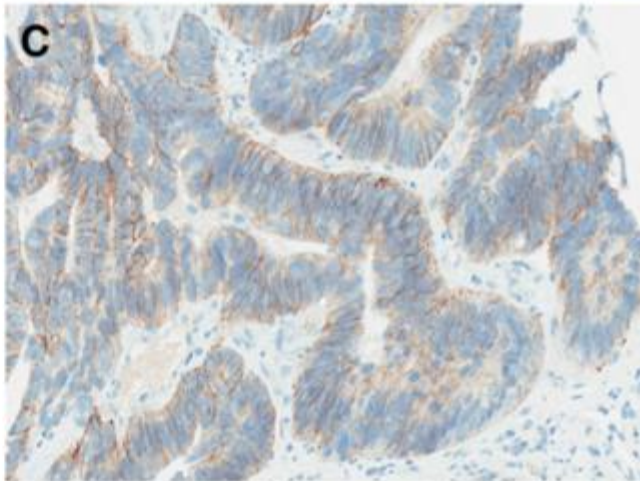
Negative 0



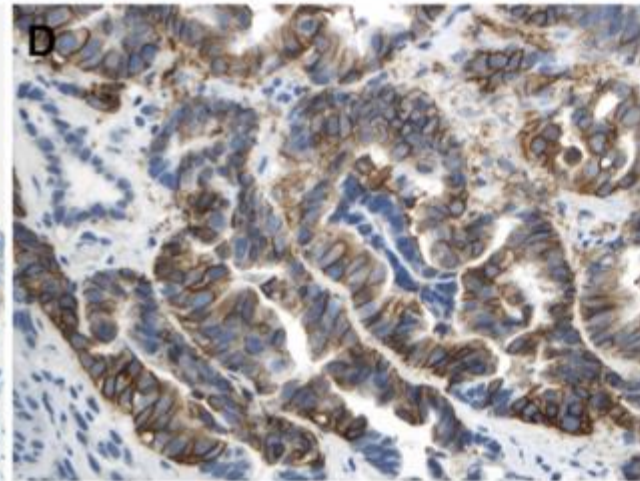
Negative 1+



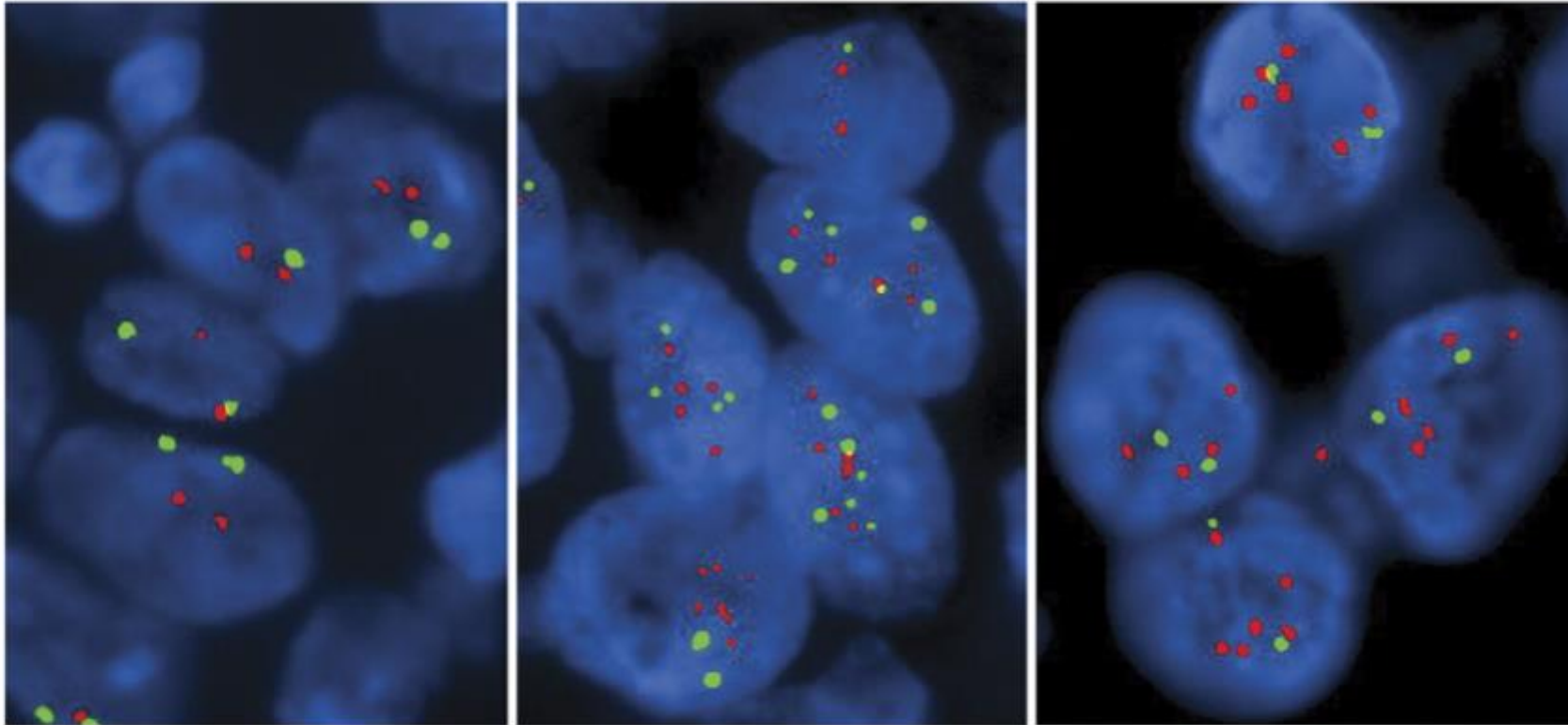
Equivocal 2+



Positive 3+



HER2 and CEP17 FISH in Representative Gastroesophageal Adenocarcinomas



$$\text{HER2 Ratio} = \text{HER2/CEP17} = \frac{1.9}{1.8} = 1.0$$

Not amplified

$$\frac{3.4}{2.7} = 1.3$$

Not amplified

$$\frac{5.2}{1.7} = 3.0$$

“Amplified”

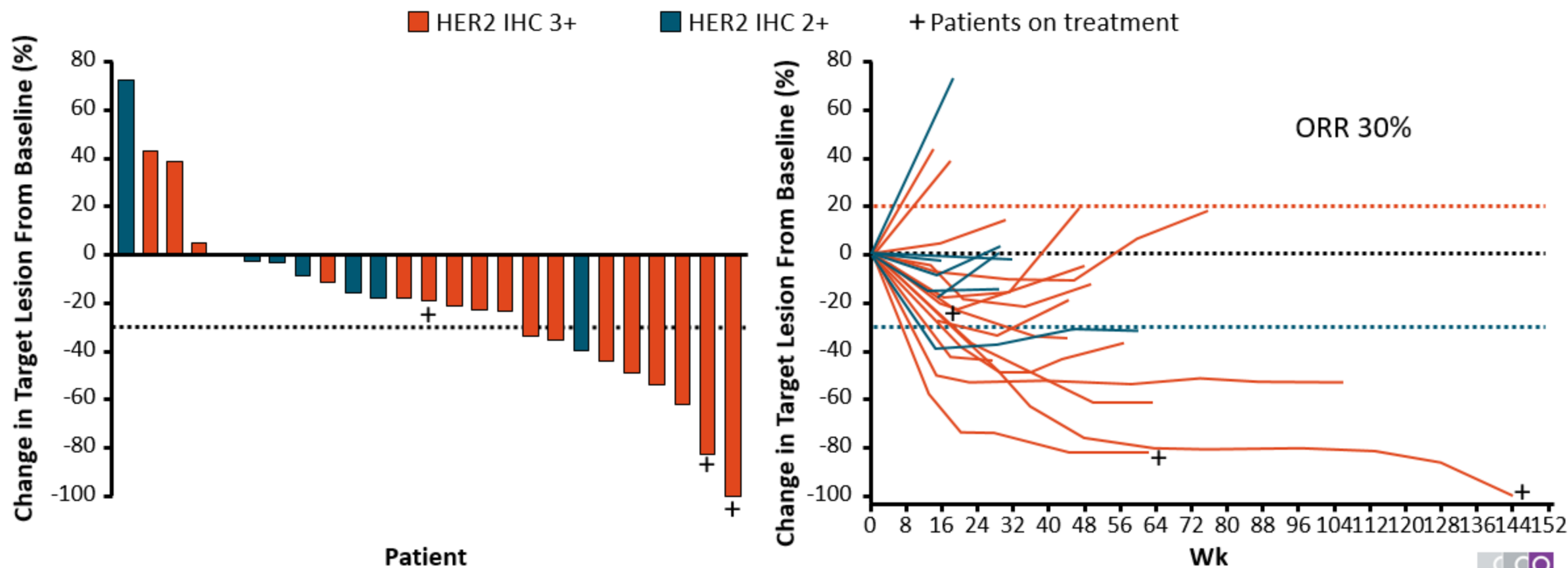
Recent Data With HER2-Targeted Regimens for Advanced CRC

Regimen	Key Trials	Key Findings	Ongoing Trials
Trastuzumab + lapatinib	HERACLES <ul style="list-style-type: none"> For HER2+ mCRC with PD after standard treatment (phase II, N = 32) 	<ul style="list-style-type: none"> ORR: 28% 	
Trastuzumab + pertuzumab	MyPathway <ul style="list-style-type: none"> For HER2+ mCRC refractory to standard treatment (phase II, N = 84) 	<ul style="list-style-type: none"> ORR: 31% 	<ul style="list-style-type: none"> Trastuzumab + pertuzumab vs CT (NCT03365882/S1613, ph II)
Tucatinib + trastuzumab	MOUNTAINEER <ul style="list-style-type: none"> For HER2+ mCRC that was previously treated (phase II, N = 26) 	<ul style="list-style-type: none"> ORR: 55% 	<ul style="list-style-type: none"> Ongoing as expanded pivotal trial (NCT03043313, ph II) Tucatinib + trastuzumab ± CT ± pembro (NCT04430738, ph I/II)
Trastuzumab deruxtecan	DESTINY-CRC01 <ul style="list-style-type: none"> For advanced CRC of varied HER2 expression with ≥2 prior regimens (phase II, N = 78) 	<ul style="list-style-type: none"> ORR: 45.3% if HER2+ 	<ul style="list-style-type: none"> DESTINY-CRC02 (NCT04744831, ph II)

- Others: neratinib, zanidatamab

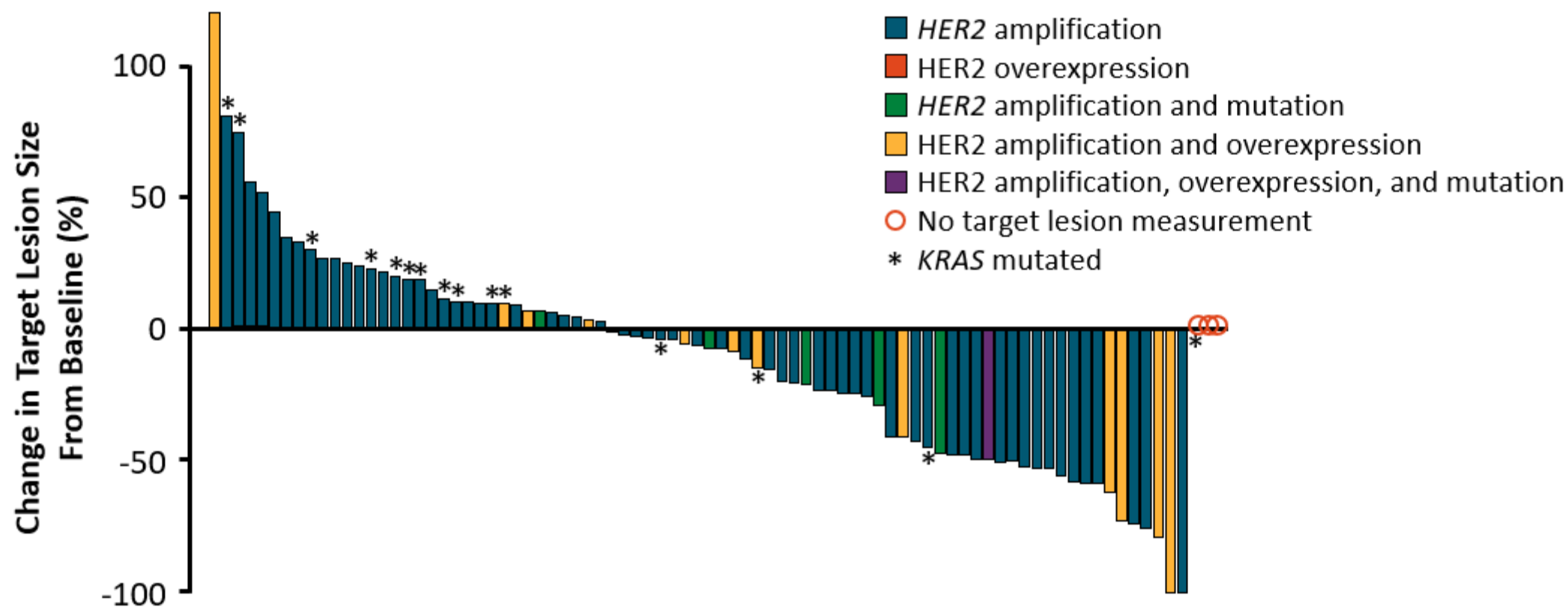
HERACLES: Trastuzumab + Lapatinib for Previously Treated mCRC

- Multicenter, open-label phase II trial of trastuzumab + lapatinib for patients with HER2+/KRAS exon 2 WT metastatic CRC; PD on/within 6 mo of approved standard treatment for CRC* (N = 27)



MyPathway: Trastuzumab + Pertuzumab for HER2+ Metastatic CRC

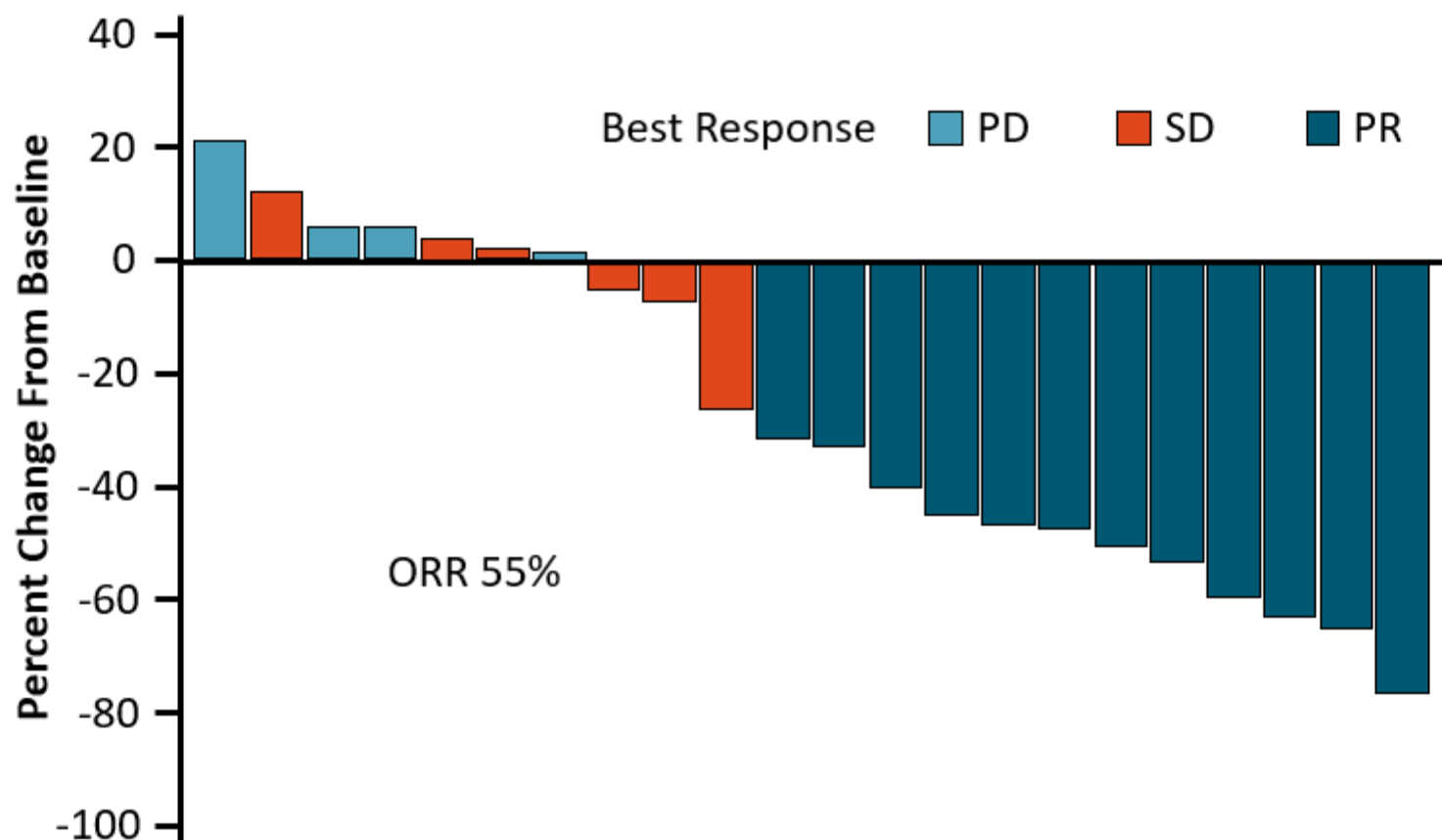
- Open-label phase IIa basket study (n = 84 with CRC)



- ORR: 30.9% in *KRAS* wild-type (n = 68)

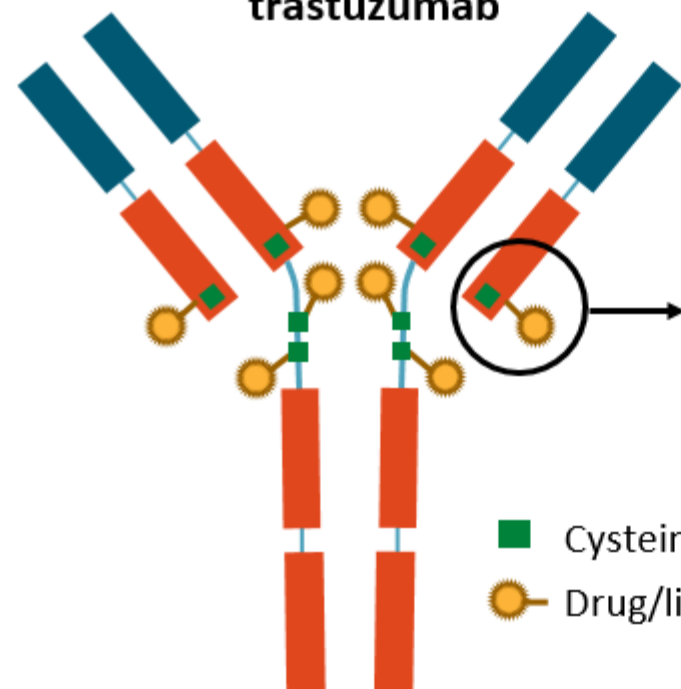
MOUNTAINEER: Tucatinib + Trastuzumab in *HER2*-Amplified mCRC

- Open-label, single-arm phase II study of tucatinib + trastuzumab for patients with previously treated, *RAS*-WT, *HER2*-amplified, metastatic or unresectable CRC (N = 26)

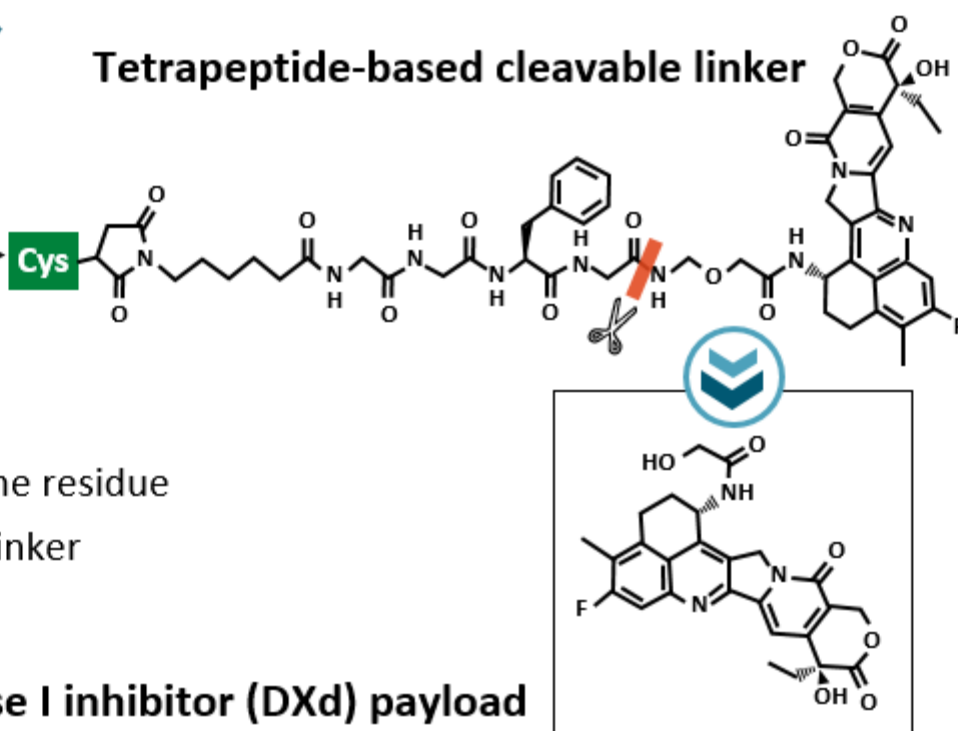


HER2-Targeted ADC: Trastuzumab Deruxtecan

Humanized anti-HER2 IgG1 mAb
with same AA sequence as
trastuzumab



Tetrapeptide-based cleavable linker

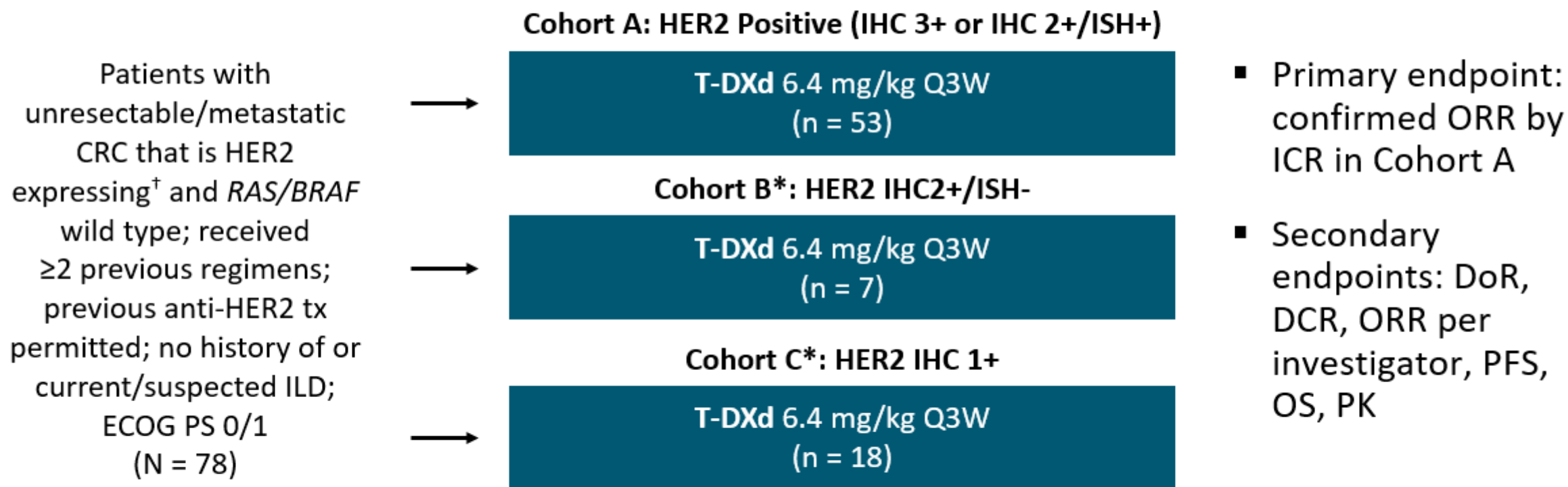


Topoisomerase I inhibitor (DXd) payload
(exatecan derivative)

- High drug:antibody ratio: ~8
- Stable linker-payload; tumor-selectable cleavable linker
- High potency, membrane-permeable payload with short systemic half-life; bystander killing effect
- FDA indicated for adult patients with locally advanced or metastatic HER2+ gastric or GEJ adenocarcinoma who have received a prior trastuzumab-based regimen

DESTINY-CRC01: Trastuzumab Deruxtecan for Patients With HER2-Expressing Metastatic Colorectal Cancer

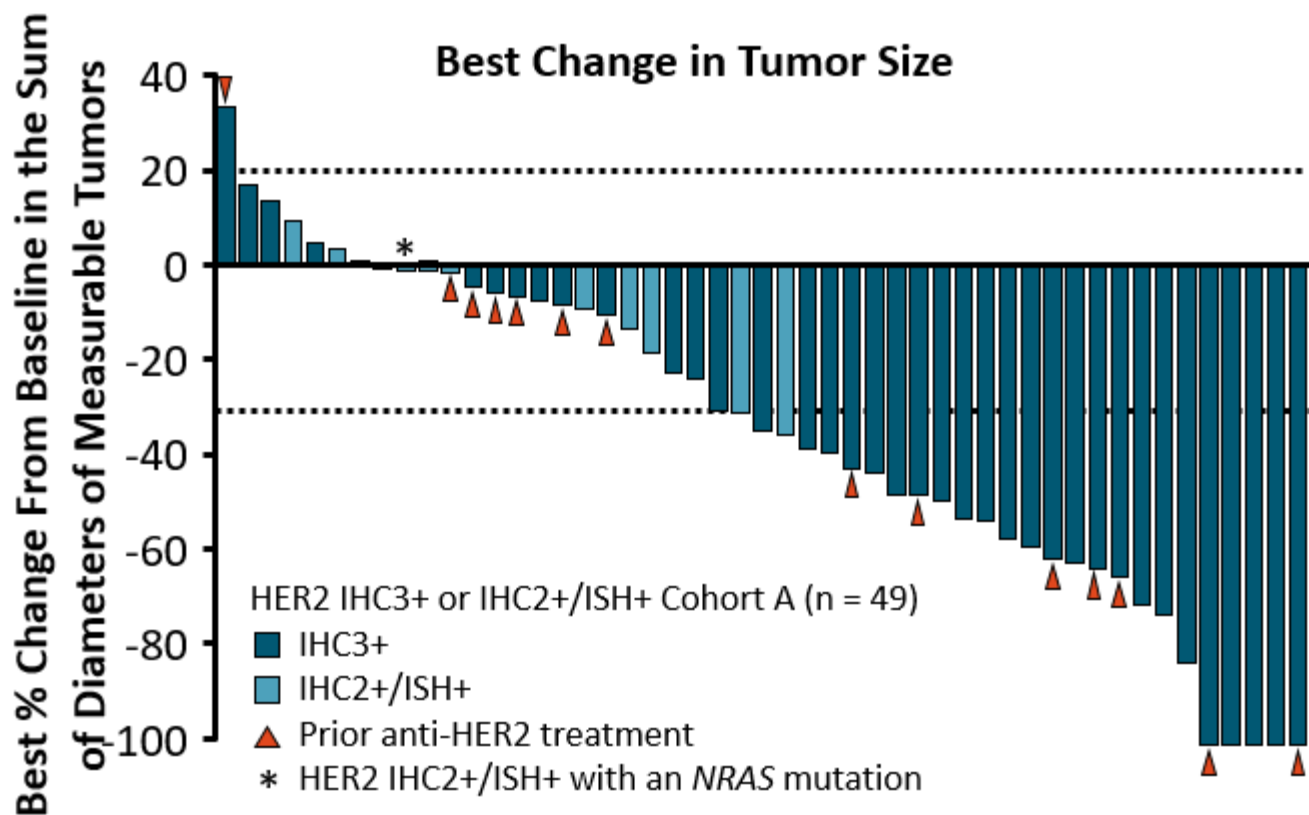
- International, open-label, multicohort phase II trial



*Cohorts B and C opened when futility monitoring performed after ≥20 patients enrolled on Cohort A had 12 wk of follow-up.

[†]Centrally confirmed.

DESTINY-CRC01: Best Change in Tumor Size and Response With Trastuzumab Deruxtecan in HER2+ CRC Cohort A



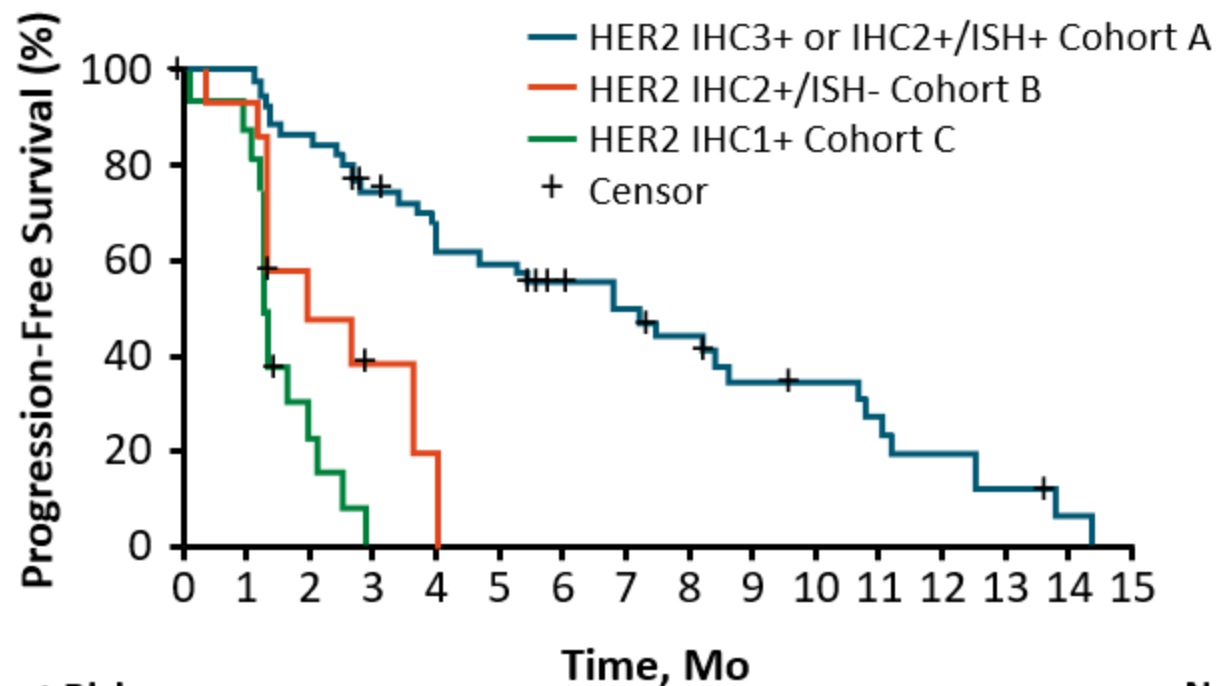
Response, n (%)	HER2+ Cohort A (n = 53)
Confirmed ORR by ICR (primary endpoint)	24 (45.3)
▪ CR	0
▪ PR	24 (45.3)
▪ SD	20 (37.7)
▪ PD	5 (9.4)
▪ NE	4 (7.5)*
DCR, % (95% CI)	83.0 (70.2-91.9)
Median DoR, mo (95% CI)	7.0 (5.8-9.5)

*Postbaseline scans missing.

- Tumor shrinkage generally detected by Mo 2 and sustained or deepened over time
- No confirmed responses by ICR in cohorts B and C

DESTINY-CRC01: PFS and OS by Cohort

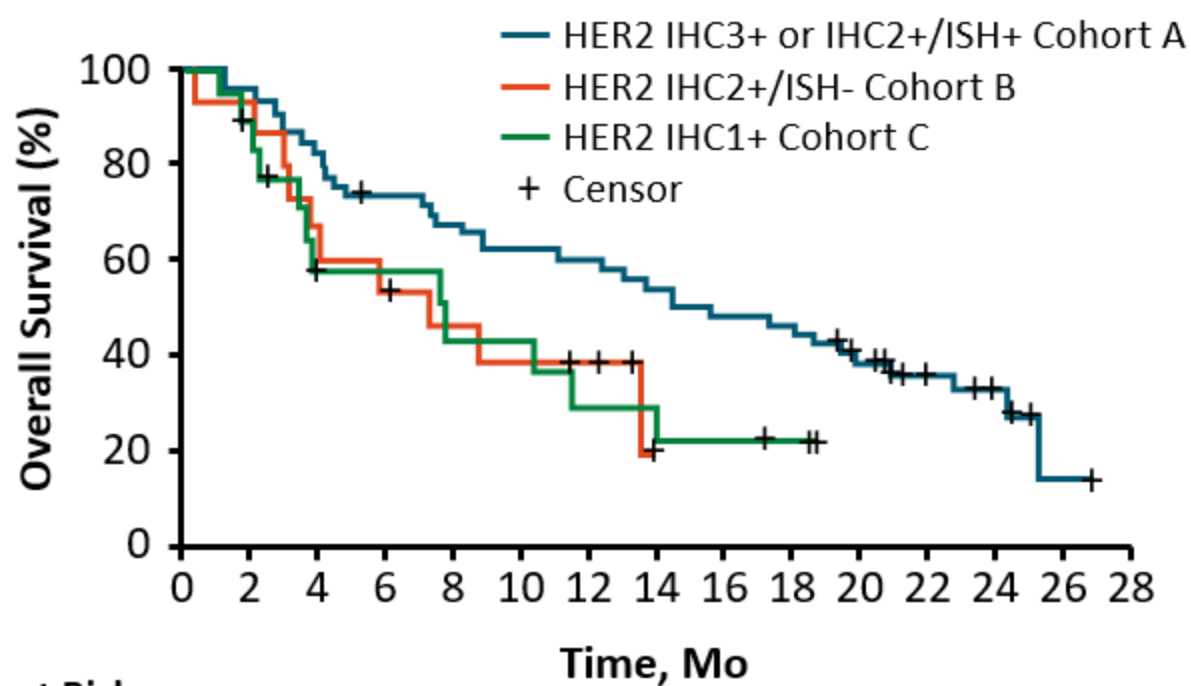
Progression-Free Survival



No. at Risk

Cohort A	53	51	44	36	33	27	22	18	15	10	9	7	5	3	1	0
Cohort B	15	14	4	4	1	0	0	0	0	0	0	0	0	0	0	0
Cohort C	18	15	4	1	0	0	0	0	0	0	0	0	0	0	0	0

Overall Survival



No. at Risk

Cohort A	53	51	44	36	35	32	31	28	25	24	18	12	6	1	0
Cohort B	15	14	10	8	6	5	4	0	0	0	0	0	0	0	0
Cohort C	18	15	8	8	6	6	4	3	3	2	0	0	0	0	0

Select Ongoing Biomarker-Based Trials in CRC

- Many ongoing biomarker-based studies

Study	Phase	Treatment	Population
BREAKWATER (NCT04607421)	III	Encorafenib + cetuximab ± CT vs CT	<i>BRAF</i> V600E-mutant mCRC
KRYSTAL-10 (NCT04793958)	III	Adagrasib + cetuximab vs CT	Advanced <i>KRAS</i> G12C CRC, previous 1L therapy
DESTINY-CRC02 (NCT04744831)	II	Trastuzumab deruxtecan	Advanced HER2+ CRC, prior therapy
S1613 (NCT03365882)	II	Trastuzumab + pertuzumab vs cetuximab + irinotecan	Advanced HER2+ CRC, ≤2 prior lines of therapy

NTRK Genes and TRK Proteins: Roles in Normal Biology and Cancer

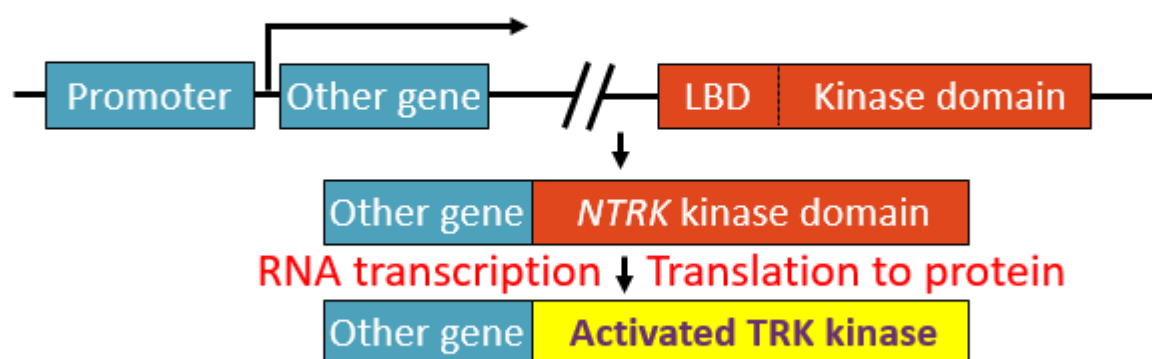
- **NTRK** genes and **TRK** receptors¹:

- In normal biology, expressed in neuronal tissue; roles in development, nervous system function via activation by neurotrophins
- Rarely expressed in normal nonneuronal or cancerous tissues

Receptor ²⁻⁴	Gene	Function
TRKA	NTRK1	Pain, thermoregulation
TRKB	NTRK2	Movement, memory, mood, appetite, weight
TRKC	NTRK3	Proprioception

- **NTRK** gene fusions and **TRK** chimeric fusion proteins¹:

- In-frame rearrangement of any *NTRK* gene links tyrosine kinase domain with upstream fusion partner to generate a chimeric RNA and TRK fusion protein
- Uncontrolled TRK kinase function results⁵

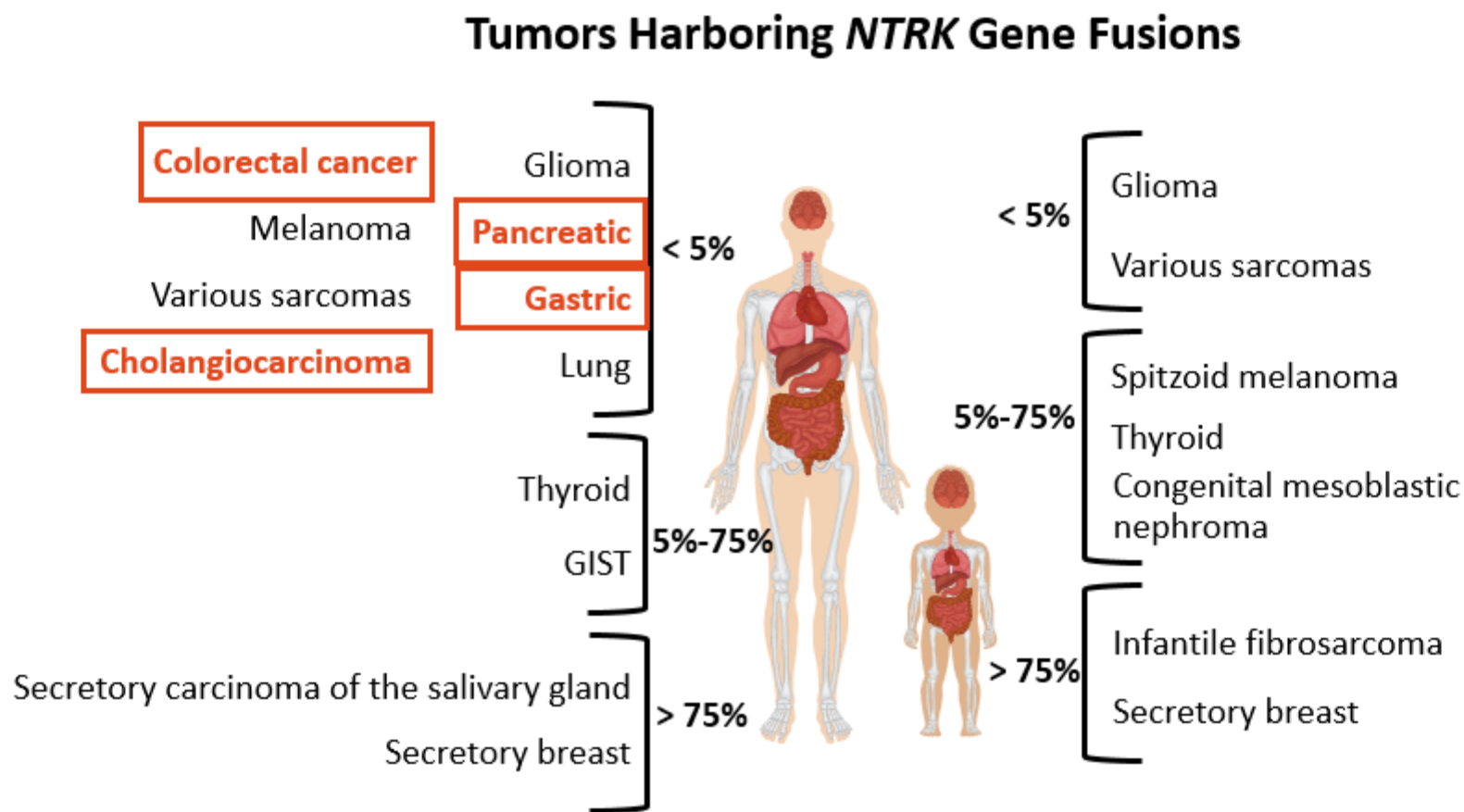


NTRK Gene Fusions in GI Cancers: Rare but Actionable

- *NTRK* gene fusions and TRK fusion chimeric proteins occur in <5% of GI cancers

Treatment	Tumor FDA Agnostic Indication
Larotrectinib	<ul style="list-style-type: none">▪ Adult/pediatric patients with solid tumors with a <i>NTRK</i> gene fusion without a known acquired resistance mutation who are either metastatic or not candidates for surgical resection due to likely severe morbidity and who have no satisfactory alternative treatments or PD following treatment
Entrectinib	<ul style="list-style-type: none">▪ Adult/pediatric patients (≥12 yr of age) with solid tumors with a <i>NTRK</i> gene fusion without a known acquired resistance mutation who are either metastatic or not candidates for surgical resection due to likely severe morbidity and who have no satisfactory alternative treatments PD following treatment

NTRK Gene Fusions in GI Cancers: Rare but Actionable



FDA-Approved TRK Inhibitors

Larotrectinib

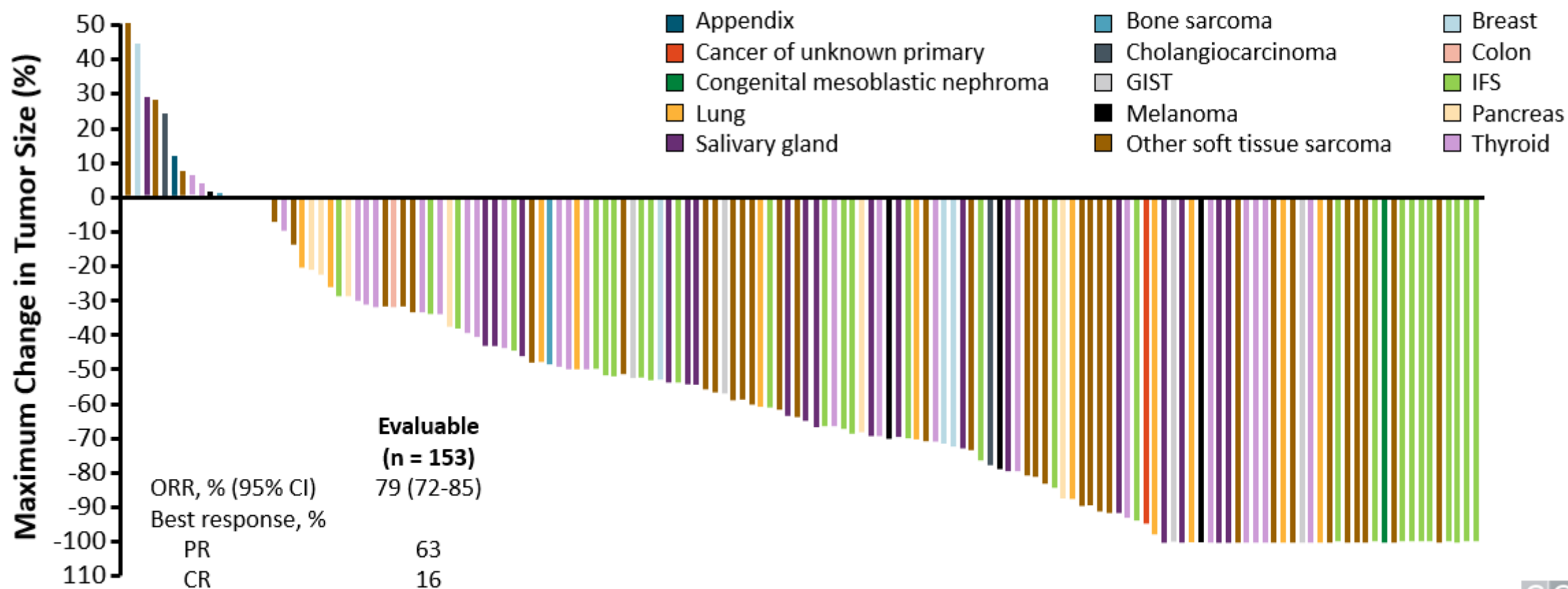
- Adult and pediatric patients with **solid tumors with a *NTRK* gene fusion** without a known acquired resistance mutation who are either **metastatic or not candidates for surgical resection** due to likely severe morbidity and who have **no satisfactory alternative treatments or whose cancer has progressed following standard treatment**
- Second tissue-agnostic approval (after pembrolizumab)

Entrectinib

- Adult and pediatric patients with **locally advanced or metastatic *NTRK* fusion–positive solid tumors** without a known acquired resistance mutation who are either metastatic or not candidates for surgical resection due to likely severe morbidity and **who have no satisfactory alternative treatments or whose cancer has progressed following standard treatment**
- Third tissue-agnostic approval

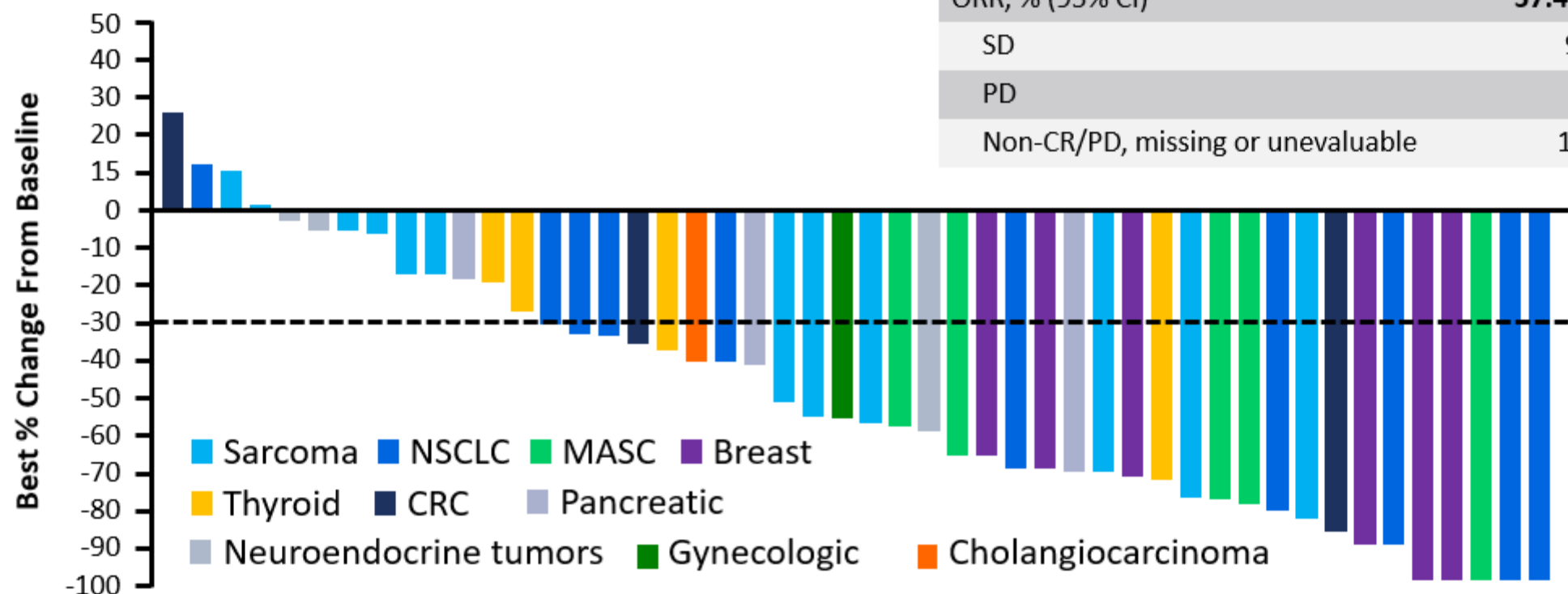
Larotrectinib: Antitumor Activity Across Tumor Types

- Analysis of 3 open-label trials (phase I, adults; phase I/II, children; phase II, adolescents/adults) assessing larotrectinib for treating advanced solid tumors with *NTRK* gene fusion (N = 159)



Entrectinib in *NTRK* Fusion+ Solid Tumors

- Analysis of 3 open-label trials (phase I or II trials in adults) assessing entrectinib for treating advanced solid tumors with *NTRK* gene fusion (N = 54)



Patients (N = 54)	
ORR, % (95% CI)	57.4 (43.2-70.8)
SD	9 (16.7)
PD	4 (7.4)
Non-CR/PD, missing or unevaluable	10 (18.5)

Results per Blinded Independent Central Review

Methods for Detection of *NTRK* Gene Fusions

	IHC	FISH	NGS
Advantages	<ul style="list-style-type: none"> ✓ Rapid results ✓ Detects transcribed and translated events only ✓ Low cost as single test 	<ul style="list-style-type: none"> ✓ Rapid results 	<ul style="list-style-type: none"> ✓ Potential for multiplexed testing ✓ Less depletion of tissue ✓ Fusion partner/position defined
Disadvantages	<ul style="list-style-type: none"> – Depletion of tissue – Fusion partner/position unknown – Less well-validated currently 	<ul style="list-style-type: none"> – Depletion of tissue – Fusion partner/position unknown – Can be difficult to interpret 	<ul style="list-style-type: none"> – Longer wait time for results – Cost

DNA- vs RNA-Based NGS Diagnostic Approaches

DNA-based NGS	RNA-based NGS
<ul style="list-style-type: none"> ✓ Potential for multiplexed assessment for multiple different fusion targets 	
<ul style="list-style-type: none"> – Detected fusions may or may not be expressed 	<ul style="list-style-type: none"> ✓ Detection of expressed transcripts
<ul style="list-style-type: none"> – Requires tiling through intronic regions 	<ul style="list-style-type: none"> ✓ Read directly through expressed transcripts
<ul style="list-style-type: none"> – Fusion positions inferred based on reads through genomic DNA 	<ul style="list-style-type: none"> ✓ Fusion positions are directly demonstrated

Select Ongoing Studies of Immune Checkpoint and TRK Inhibitors

Study	Phase	Treatment	Population
CheckMate 8HW (NCT04008030)	III	Nivolumab ± ipilimumab vs CT	mCRC with dMMR/MSI-H
NCT04895722	III	Pembrolizumab or pembrolizumab/quavonlimab	mCRC with dMMR/MSI-H
NAVIGATE (NCT02576431)	II	Larotrectinib	Advanced solid tumors with <i>NTRK</i> fusion
STARTRK-2 (NCT02568267)	II	Entrectinib	Advanced solid tumors with <i>NTRK</i> fusion or <i>ROS1</i> / <i>ALK</i> rearrangement
NCT03215511	I/II	Selitrectinib	Advanced solid tumors with <i>NTRK1/2/3</i> fusion and prior TRK inhibitor, no satisfactory treatment options
TRIDENT-1 (NCT03093116)	I/II	Repotrectinib	Patients aged ≥12 yr, advanced solid tumors with <i>NTRK1/2/3</i> , <i>ROS1</i> , or <i>ALK</i> fusion
NCT04094610	I/II	Repotrectinib	Children and young adults, advanced malignancies with <i>NTRK1/2/3</i> , <i>ROS1</i> , or <i>ALK</i> fusion

Conclusions: Clinical Relevance and Use of Biomarkers

- Current precision therapies with clinical activity:

Alteration	Targeted Therapy
<i>BRAF</i> V600E mutation	Dual EGFR + BRAF inhibition (\pm MEK inhibitor)
<i>HER2</i> amplification	Trastuzumab + lapatinib or pertuzumab
MSI-H/dMMR	Nivolumab \pm ipilimumab, pembrolizumab
<i>NTRK</i> fusion	Larotrectinib or entrectinib



powered by cea

Biomarker-Driven Treatment of Gastric, Esophageal, and Gastroesophageal Junction Cancers



STRONGERTOGETHER



Biomarker Testing in Gastroesophageal Cancers

- What test results must we have?
 - *HER2* amplification (IHC/ISH)
 - MSI-H (PCR/NGS) or dMMR (IHC)
 - TMB-H (NGS)
 - PD-L1 examination (IHC)
 - *NTRK* fusion (NGS)
 - Comprehensive genomic profiling (if enough tissue)
- When should testing occur, and for whom?
 - All newly diagnosed patients with metastatic disease

PD-L1 in Gastroesophageal Cancers

- 23% to 60% of gastric cancers are PD-L1+ (tumor cells + tumor-infiltrating immune cells)¹⁻³
- Serial sections: HE and IHC; clone 22 C3
- Minimum 100 tumor cells in IHC slide
- Clinically relevant cutoffs >1%, >10%, and ~20%
- Higher response to PD-1 inhibitors in advanced gastroesophageal cancers with higher PD-L1 levels; pembrolizumab indications based on specific levels^{4,5}

Assessing PD-L1 Levels (IHC Testing)

$$\text{Combined Positive Score} = \frac{\text{PD-L1 staining cells, n} \text{ (tumor cells, lymphocytes, macrophages)}}{\text{Tumor cells evaluated, n}} \times 100$$

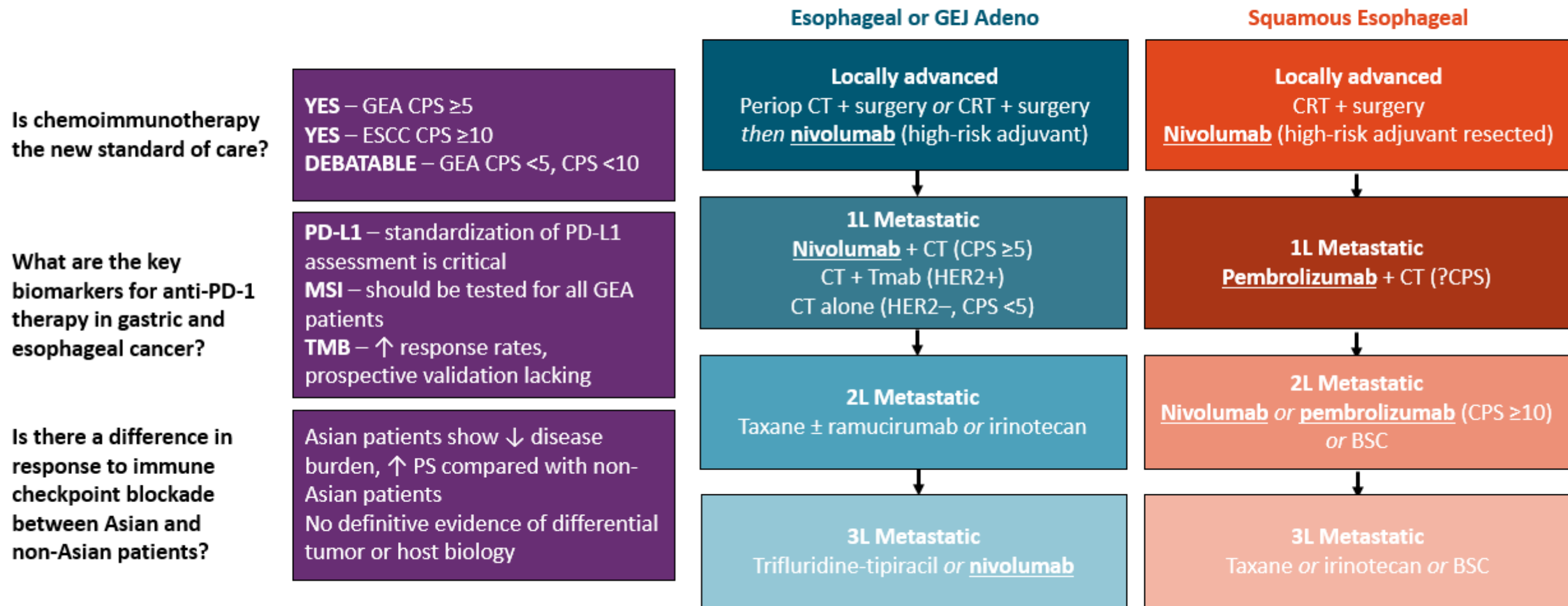
FDA-Approved Indications for Checkpoint Inhibitors in Advanced Gastroesophageal Cancers

Indication	Pembrolizumab	Nivolumab
Gastric	<ul style="list-style-type: none"> ▪ HER2+ locally advanced unresectable or metastatic gastric or GEJ adenocarcinoma (with trastuzumab and fluoropyrimidine- and platinum-containing CT) <u>as first-line therapy</u> ▪ PD-L1 CPS ≥ 1 recurrent locally advanced or metastatic gastric or GEJ adenocarcinoma <u>with PD on or after ≥ 2 previous therapies</u>, including fluoropyrimidine- and platinum-containing CT and, if appropriate, HER2-targeted therapy 	<ul style="list-style-type: none"> ▪ Advanced or metastatic gastric or GEJ cancer and esophageal adenocarcinoma with fluoropyrimidine- and platinum-containing CT
Esophageal	<ul style="list-style-type: none"> ▪ Locally advanced or metastatic esophageal or GEJ carcinoma: <ul style="list-style-type: none"> – With platinum- and fluoropyrimidine-based CT – For PD-L1 CPS ≥ 10 <i>squamous</i> carcinoma <u>after ≥ 1 line(s)</u> of systemic therapy 	<ul style="list-style-type: none"> ▪ Unresectable advanced, recurrent, or metastatic esophageal <i>squamous</i> cell carcinoma <u>after prior</u> fluoropyrimidine- and platinum-based CT ▪ See above
Tumor agnostic	<ul style="list-style-type: none"> ▪ MSI-H or MMR deficient or TMB-H (≥ 10 mut/Mb) unresectable or metastatic solid tumors <u>with PD after previous treatment</u> with no satisfactory alternative treatment options 	

Simplified First-line Treatment Algorithm for Advanced Gastroesophageal Adenocarcinomas

	No Biomarkers or HER2-	HER2+
Gastric	Fluoropyrimidine + platinum ± nivolumab (CPS ≥5; CheckMate 649)	Fluoropyrimidine + platinum + trastuzumab ± pembrolizumab (KEYNOTE-811)
Esophageal/ GEJ	Fluoropyrimidine + platinum ± nivolumab (CPS ≥5; CheckMate 649) Fluoropyrimidine + platinum ± pembrolizumab (CPS ≥10; KEYNOTE-590)	Fluoropyrimidine + platinum + trastuzumab ± pembrolizumab (KEYNOTE-811)

Standard of Care for Gastroesophageal Cancers in 2021

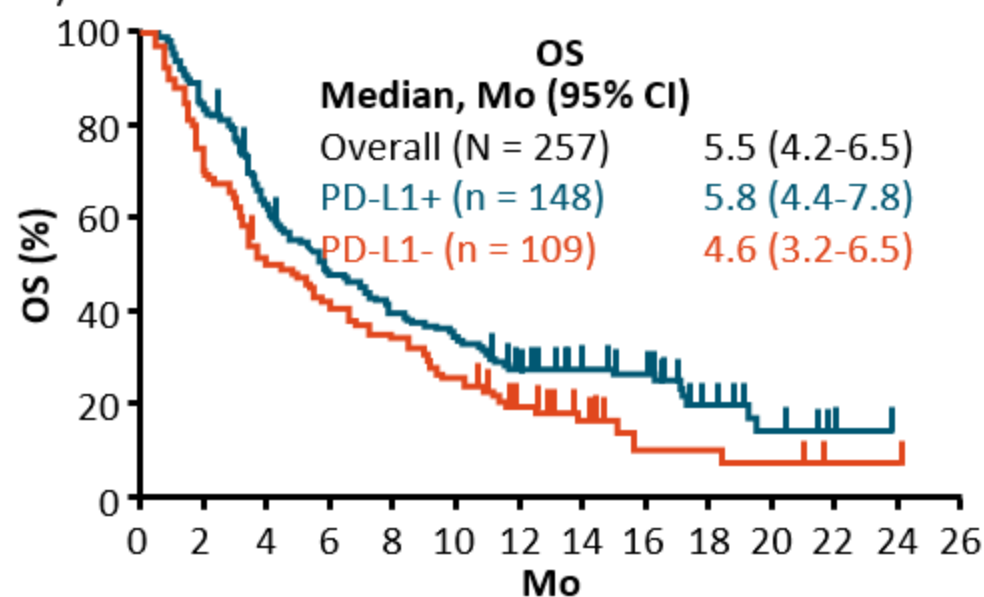
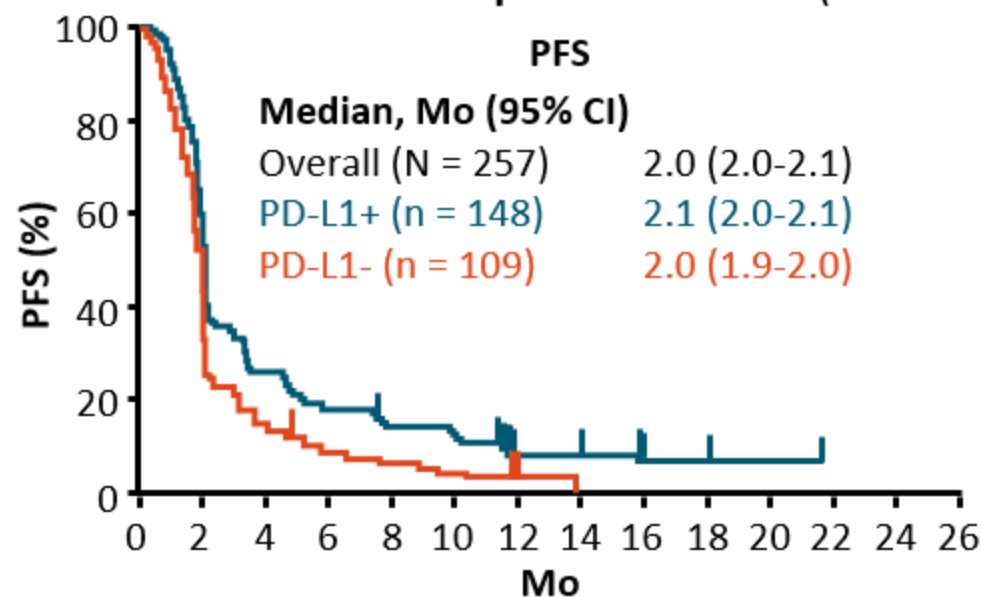


Additional FDA Indications for Immune Checkpoint Inhibitors in MSI-H/dMMR/PD-L1+ GI Cancers

- Pembrolizumab for **MSI-high/dMMR unresectable or metastatic CRC**
- Nivolumab ± ipilimumab for **MSI-high/dMMR metastatic CRC** with PD after fluoropyrimidine, oxaliplatin, and irinotecan
- Pembrolizumab for **PD-L1 CPS ≥ 1 recurrent locally advanced or metastatic gastric or GEJ adenocarcinoma** with PD on or after ≥ 2 previous therapies, including fluoropyrimidine- and platinum-containing CT and, if appropriate, HER2/neu-targeted therapy
- Pembrolizumab for **PD-L1 CPS ≥ 10 locally advanced or metastatic esophageal or GEJ squamous carcinoma** after ≥ 1 line(s) of systemic therapy

KEYNOTE-059: Pembrolizumab for Gastric/GEJ Adenocarcinoma With ≥ 2 Prior Lines of CT

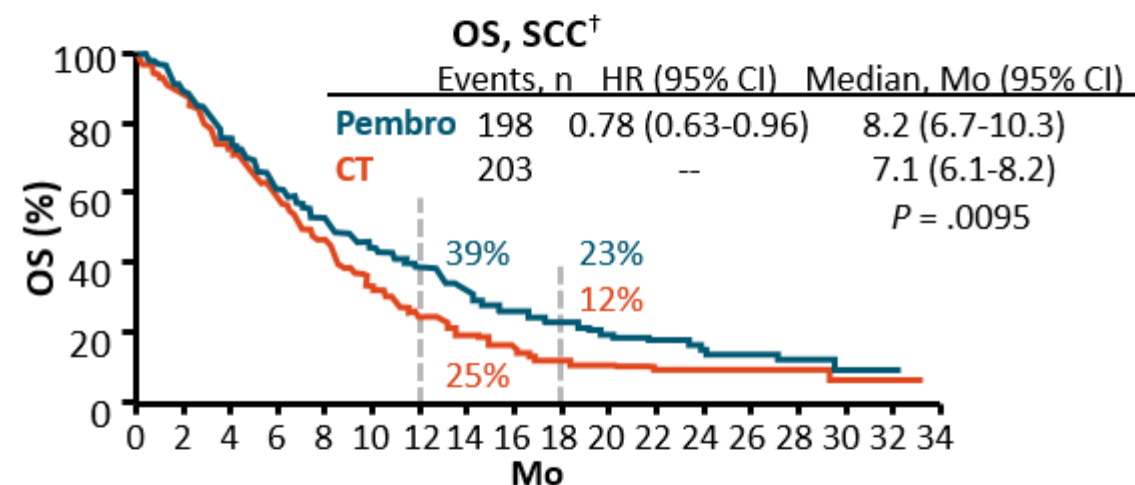
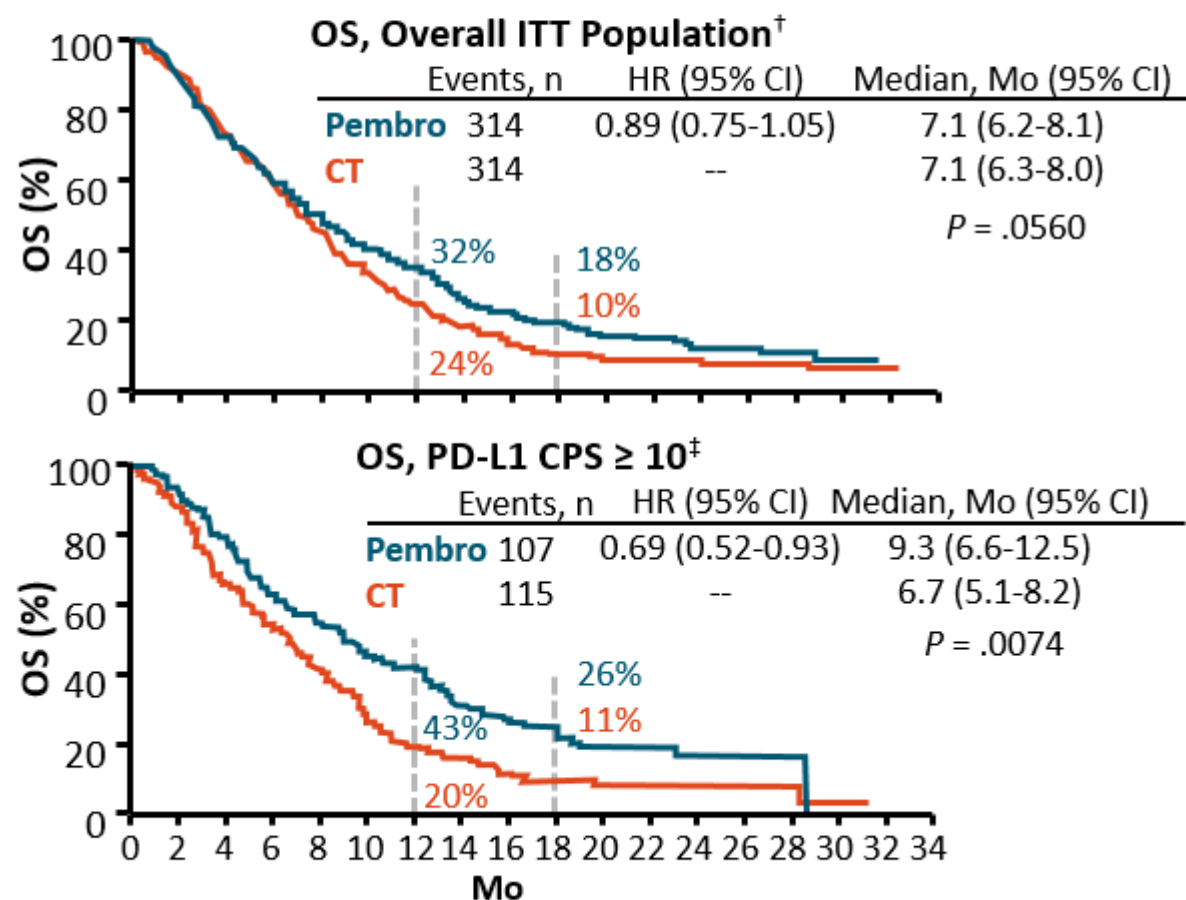
- Open-label phase II study of **pembrolizumab** for patients with recurrent or metastatic **gastric or GEJ adenocarcinoma** and ≥ 2 prior lines of CT (cohort 1, N = 259)



Outcome	Overall	PD-L1		MSI Status	
		Positive (n = 148)	Negative (n = 109)	High (n = 7)	Non-H (n = 167)
ORR, % (95% CI)	11.6 (8.0-16.1)	15.5 (10.1-22.4)	6.4 (2.6-12.8)	57.1	9.0
Median DoR, mo (95% CI)	8.4 (1.6+ to 17.3+)	16.3 (1.6+ to 17.3+)	6.9 (2.4 to 7.0+)	--	--

KEYNOTE-181: Second-Line Pembrolizumab vs Chemotherapy for Esophageal/GEJ Cancer

- Randomized phase III trial of **pembrolizumab** vs **investigator's choice CT*** for patients with advanced/metastatic **esophageal adenocarcinoma/SCC** or **GEJ adenocarcinoma** and **PD after 1L tx** (N = 628)

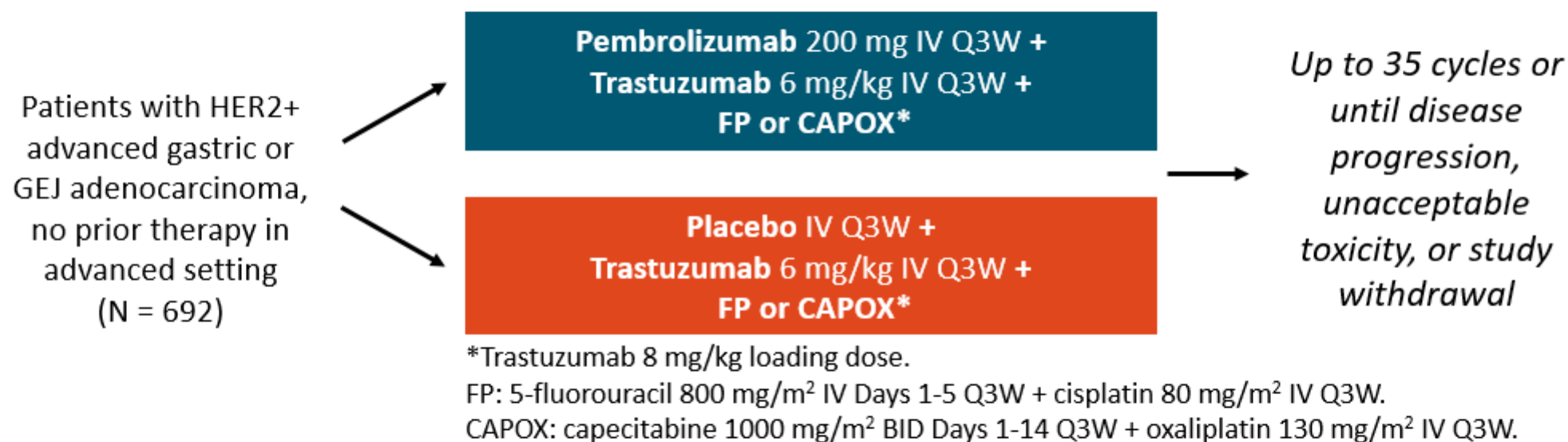


ORR, %	Pembrolizumab	Chemotherapy	P Value
Total population	13.1	6.7	.0037
PD-L1 CPS ≥ 10	21.5	6.1	.0006
SCC	16.7	7.4	.0022

*Paclitaxel or irinotecan. [†]Did not meet primary endpoint per statistical prespecifications. [‡]Met primary endpoint per statistical prespecifications.

KEYNOTE-811: 1L Pembrolizumab + Trastuzumab + Chemotherapy in HER2+ Metastatic Gastric/GEJ Cancer

- Randomized, double-blind, placebo-controlled phase III study



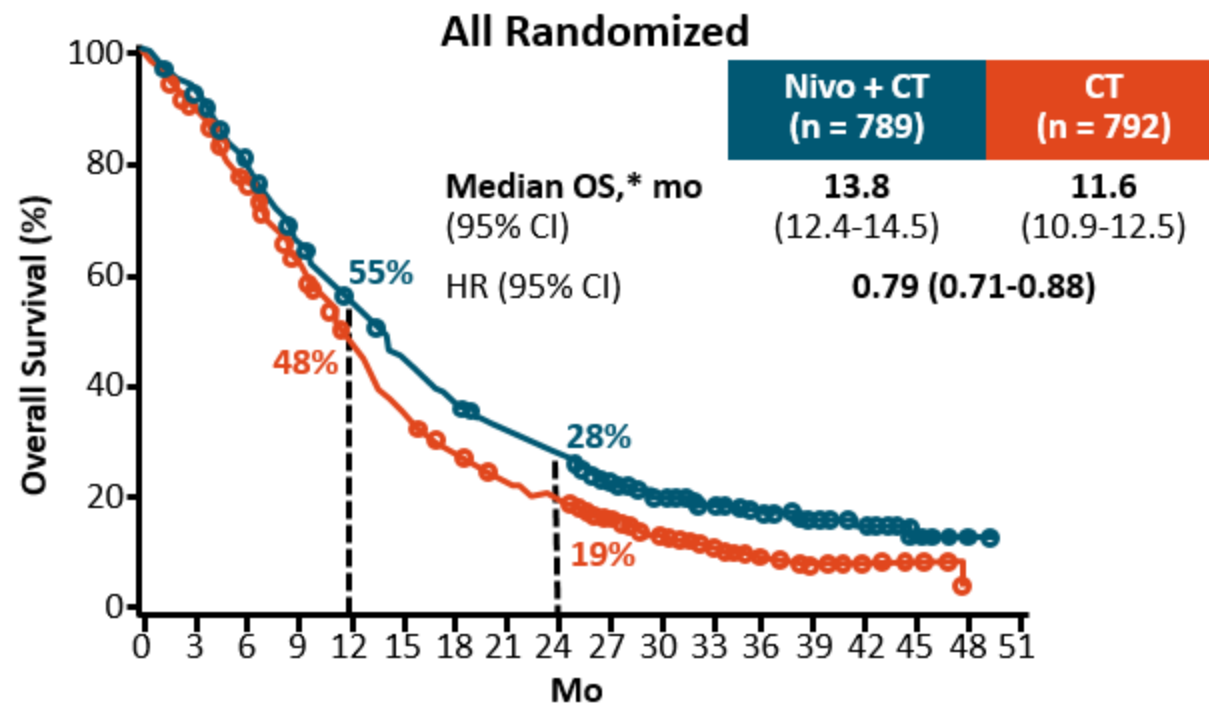
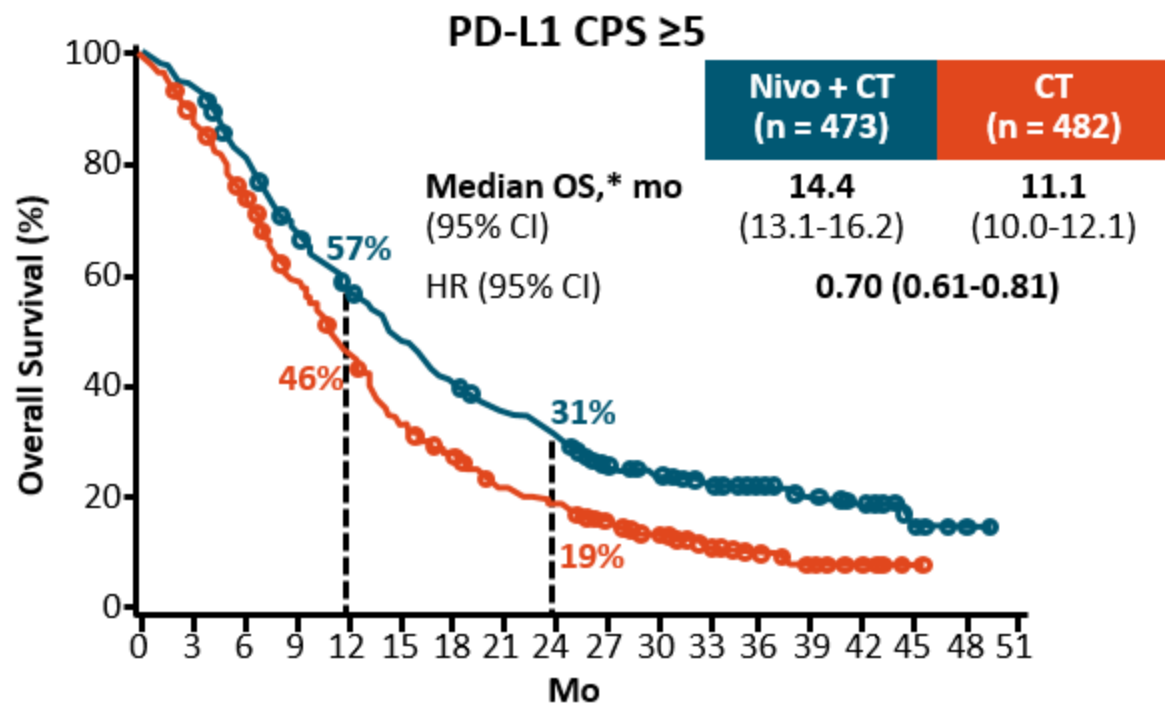
- Efficacy analysis: first 264 patients enrolled; safety analysis: 433 patients who received ≥ 1 dose of study medication
- Primary endpoints: OS, PFS per RECIST v1.1 by BICR; secondary endpoints: ORR and DoR per RECIST v1.1 by BICR, safety

KEYNOTE-811 Interim Analysis: Efficacy

Outcome	Pembrolizumab (n = 133)	Placebo (n = 131)
ORR, % (95% CI)	74.4 (66.2-81.6)	51.9 (43.0-60.7)
ORR difference*	22.7 (11.2-33.7); P = .00006	
DCR, % (95% CI)	96.2 (91.4-98.8)	89.3 (82.7-94.0)
Best response, n (%)		
▪ CR	15 (11)	4 (3)
▪ PR	84 (63)	64 (49)
▪ SD	29 (22)	49 (37)
▪ PD	5 (4)	7 (5)
▪ Not evaluable	0	2 (2)
▪ Not assessed	0	5 (4)
Duration of response [†]	(n = 99)	(n = 68)
▪ Median, mo (range)	10.6 (1.1+ to 16.5+)	9.5 (1.4+ to 15.4+)
▪ ≥6 mo duration, %	70.3	61.4
▪ ≥9 mo duration, %	58.4	51.1
Size reduction from baseline, n (%)	(n = 124)	(n = 122)
▪ Any decrease	97	90
▪ ≥80% decrease	32	15

CheckMate 649: First-line Nivolumab + Ipilimumab or CT vs CT in Gastroesophageal Cancer

- Randomized phase III trial of nivolumab + ipilimumab, **nivolumab + CT,*** or **CT*** for patients with previously untreated unresectable, advanced, or metastatic gastric/GEJ/esophageal adenocarcinoma (N = 1581)



	CPS ≥ 5 (n = 955)	CPS ≥ 1 (n = 1296)	All Randomized	CPS < 5 (n = 606)
HR for OS	0.70	0.76	0.79	0.94

*XELOX or FOLFOX.

KEYNOTE-590: First-line Pembrolizumab + Chemotherapy vs Chemotherapy for Esophageal/GEJ Cancer

- Randomized phase III trial of **pembrolizumab + CT*** vs **CT*** for previously untreated patients with locally advanced unresectable or metastatic EAC, ESCC, or GEJA (N = 749)

Outcome	All Patients			All Patients PD-L1 CPS ≥10			ESCC			ESCC PD-L1 CPS ≥10		
	Pembro + CT (n = 373)	CT (n = 376)	HR/ P Val	Pembro + CT (n = 186)	CT (n = 197)	HR/ P Val	Pembro + CT (n = 274)	CT (n = 274)	HR/ P Val	Pembro + CT (n = 143)	CT (n = 143)	HR/ P Val
Median OS, [†] mo	12.4	9.8	0.73/ <.0001	13.5	9.4	0.62/ <.0001	12.6	9.8	0.72/ .0006	13.9	8.8	0.57/ <.0001
Median PFS, [†] mo	6.3	5.8	0.65/ <.0001	7.5	5.5	0.51/ <.0001	6.3	5.8	0.65/ <.0001	--	--	--

	CPS ≥10 (n = 383)	All Randomized	CPS <10 (n = 347)
HR for OS	0.62	0.73	0.86

*5-FU + cisplatin. [†]Primary endpoint.

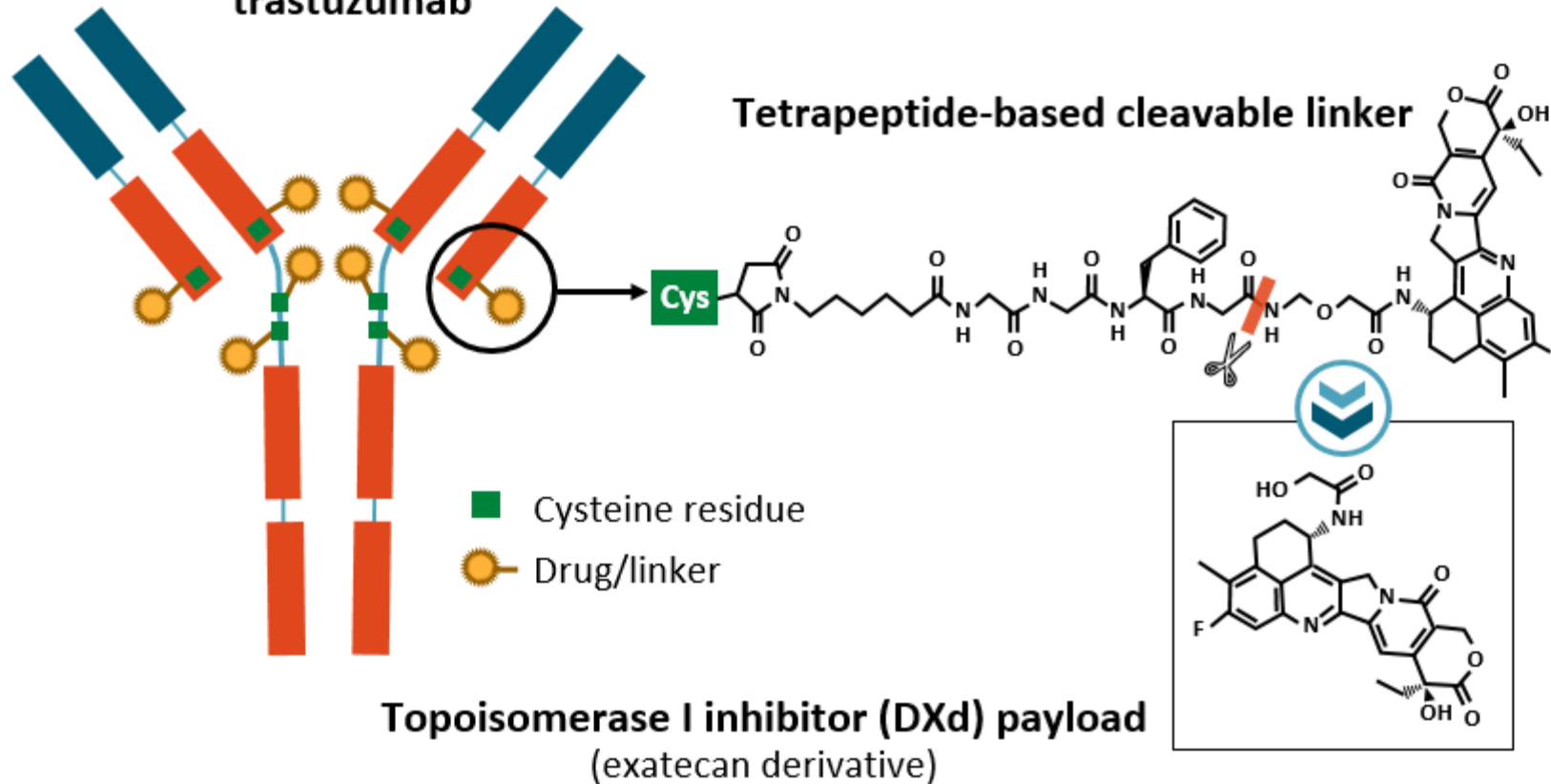
Sun. Lancet. 2021;398:759.



Slide credit: clinicaloptions.com

HER2-Targeted ADC: Trastuzumab Deruxtecan

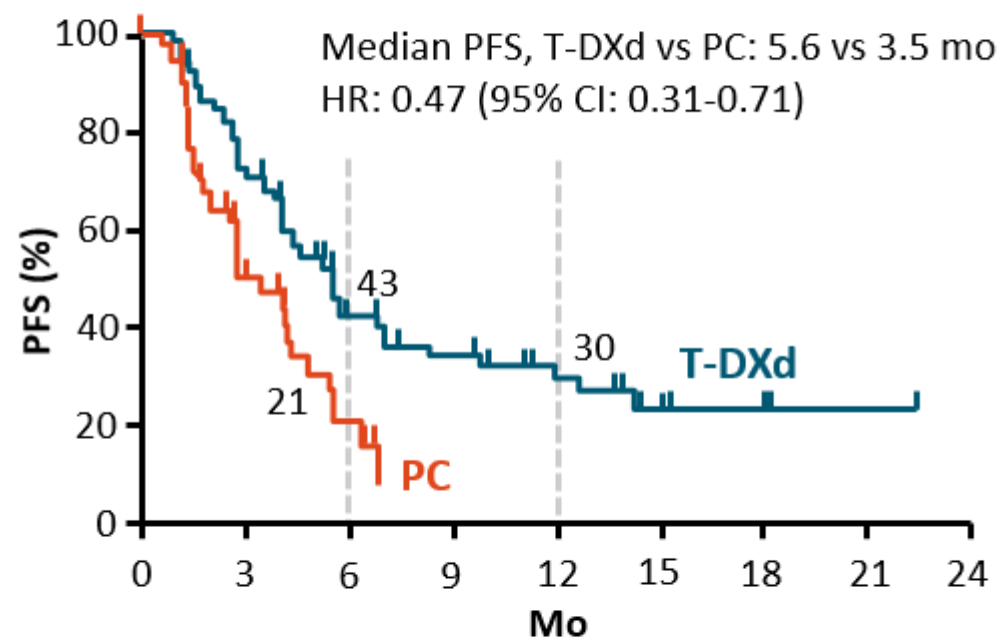
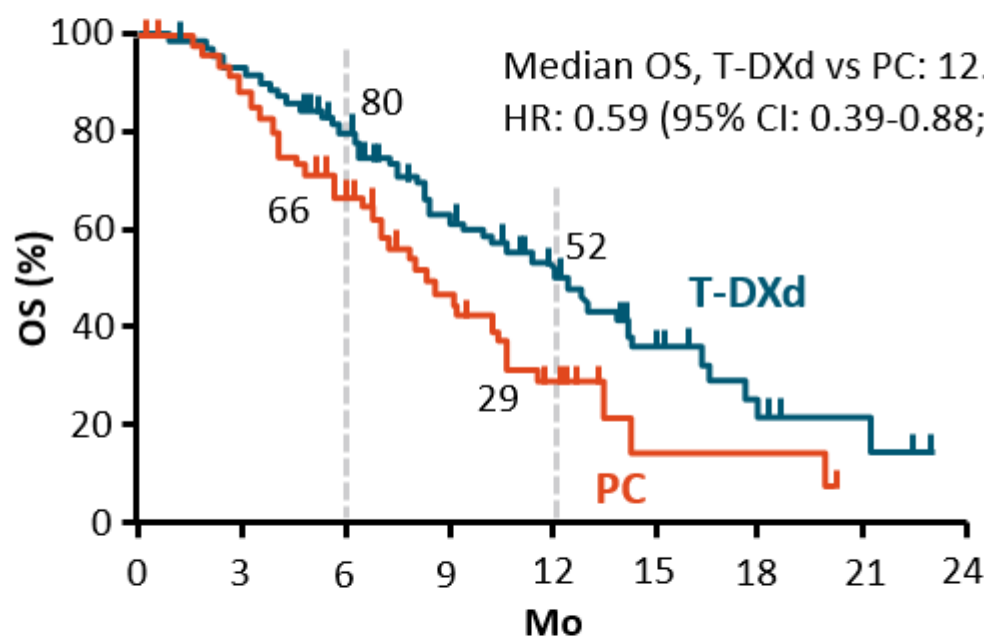
**Humanized anti-HER2 IgG1 mAb
with same AA sequence as
trastuzumab**



- High drug:antibody ratio: ~8
- Stable linker-payload
- Tumor-selectable cleavable linker
- High potency, membrane-permeable payload with short systemic half-life
- Bystander killing effect

DESTINY-Gastric01: Trastuzumab Deruxtecan in Previously Treated, HER2+ Gastric/GEJ Adenocarcinoma

- Open-label, randomized phase II study of T-DXd vs irinotecan or paclitaxel for pts with HER2+ locally advanced or metastatic gastric or GEJ cancer that progressed on ≥ 2 prior regimens (N = 188)



- ORR: T-DXd, 54%; CT, 14%
- FDA approved for pts with locally advanced or metastatic HER2+ gastric or gastroesophageal junction adenocarcinoma who have received a prior trastuzumab-based regimen

Trastuzumab Deruxtecan Approval

- HER2-targeted antibody-drug conjugate
 - mAb has same AA sequence as trastuzumab; topoisomerase I inhibitor payload (exatecan derivative)
- Indicated for adult patients with locally advanced or metastatic HER2-positive gastric or gastroesophageal junction adenocarcinoma who have received a prior trastuzumab-based regimen

Select Investigational Targeted Agents for Advanced Gastroesophageal Cancers

Agent	MoA	Key Trial	
Margetuximab	HER2-targeted antibody	CP-MGAH22-05: phase Ib/II study of margetuximab + pembrolizumab for previously treated advanced HER2+ GE cancers (N = 95)	ORR: 18%; median PFS: 3 mo; median OS: 13 mo
Zanidatamab	HER2-targeted bispecific antibody	Phase Ib/II study of zanidatamab ± CT for previously treated advanced HER2+ GE cancers (N = 52)	ORR: zan, 33%; zan + pac, 50%; zan + cape, 57%
Bemarituzumab	FGFR2b-targeted antibody	FIGHT: phase II study of first-line bemarituzumab vs placebo (both + mFOLFOX6) for advanced G/GEJ cancer with <i>FGFR2b</i> overexp/amp (N = 155)	Median OS, mo: bema, NR; placebo, 12.9 (HR: 0.58; <i>P</i> = .0268)
Zolbetuximab	CLDN18.2-targeted antibody	FAST: phase II study of first-line zolbetuximab + EOX vs EOX for advanced CLDN18.2+ G/GEJ cancer (N = 252)	Median OS, mo: zolb + EOX, 13; EOX, 8.3 (HR: 0.55; <i>P</i> < .0005)

Selected Ongoing Phase III Trials for Advanced Gastroesophageal Cancer

Trial	Regimen	Population	Phase
MOUNTAINEER-02 (NCT04499924)	Tucatinib + trastuzumab vs placebo (both with ramucirumab + paclitaxel)	2L+, GC/GEJC, HER2+	II/III
MAHOGANY (NCT04082364)	Margetuximab + PD-1 inhibitor ± CT or margetuximab + CT ± dual checkpoint inhibitor or trastuzumab + CT	1L, GC/GEJC, HER2+	II/III
DESTINY-Gastric04 (NCT04704934)	Trastuzumab deruxtecan vs ramucirumab + paclitaxel	2L+, GC/GEJC, HER2+	III
FORTITUDE-101 (NCT05052801)	Bemarituzumab + mFOLFOX6 vs placebo + mFOLFOX6	1L, GC/GEJC, FGFR2+	III
GLOW (NCT03653507)	Zolbetuximab + CAPOX vs placebo + CAPOX	1L, GC/GEJC, CLDN18.2+	III
SPOTLIGHT (NCT03504397)	Zolbetuximab + mFOLFOX6 vs placebo + mFOLFOX6	1L, GC/GEJC, CLDN18.2+	III

Biomarker-Driven Treatment of Pancreatic and Hepatobiliary Cancers

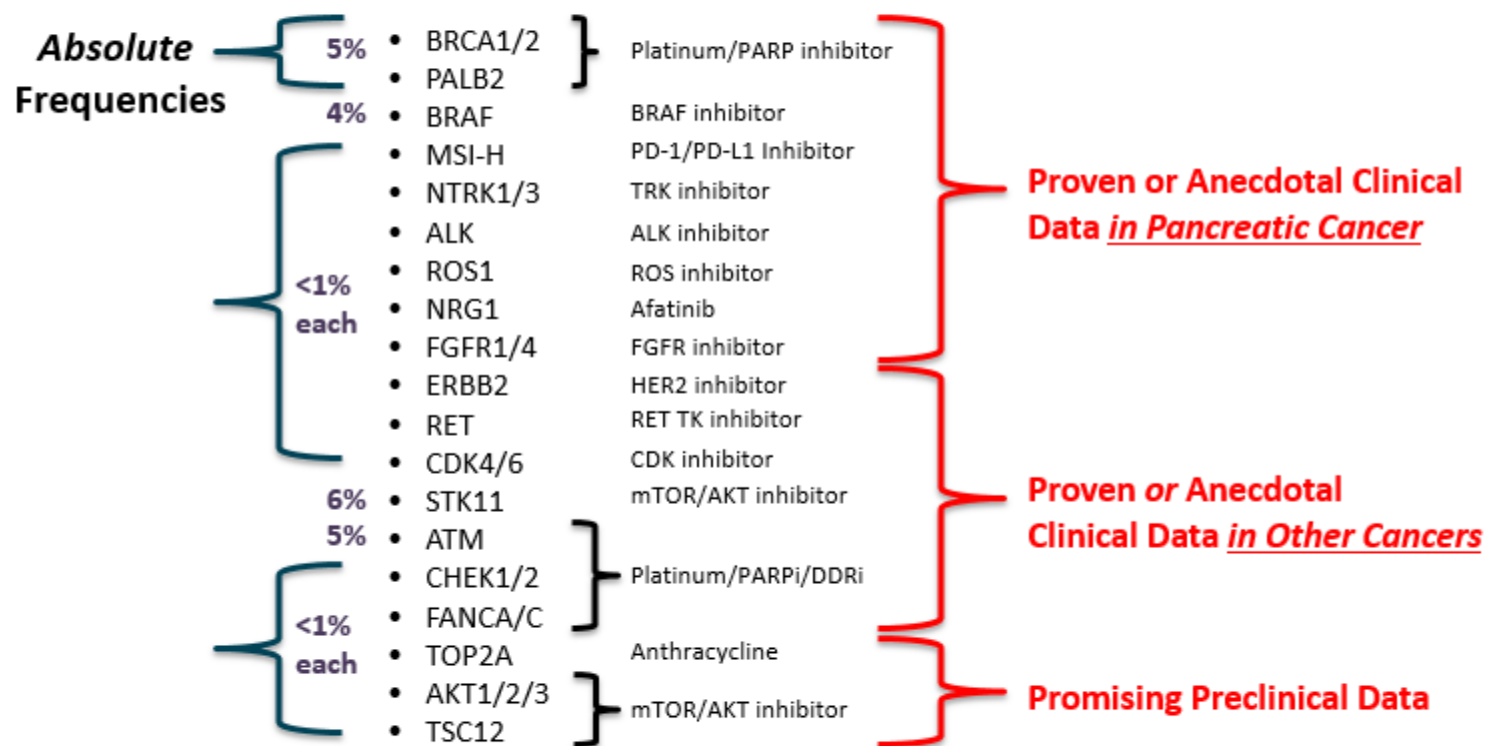


STRONGERTOGETHER



Pancreatic Cancers *DO* Harbor Actionable Mutations

- NGS efforts have consistently revealed that $\geq 25\%$ of pancreatic cancers have potentially highly actionable molecular biomarkers

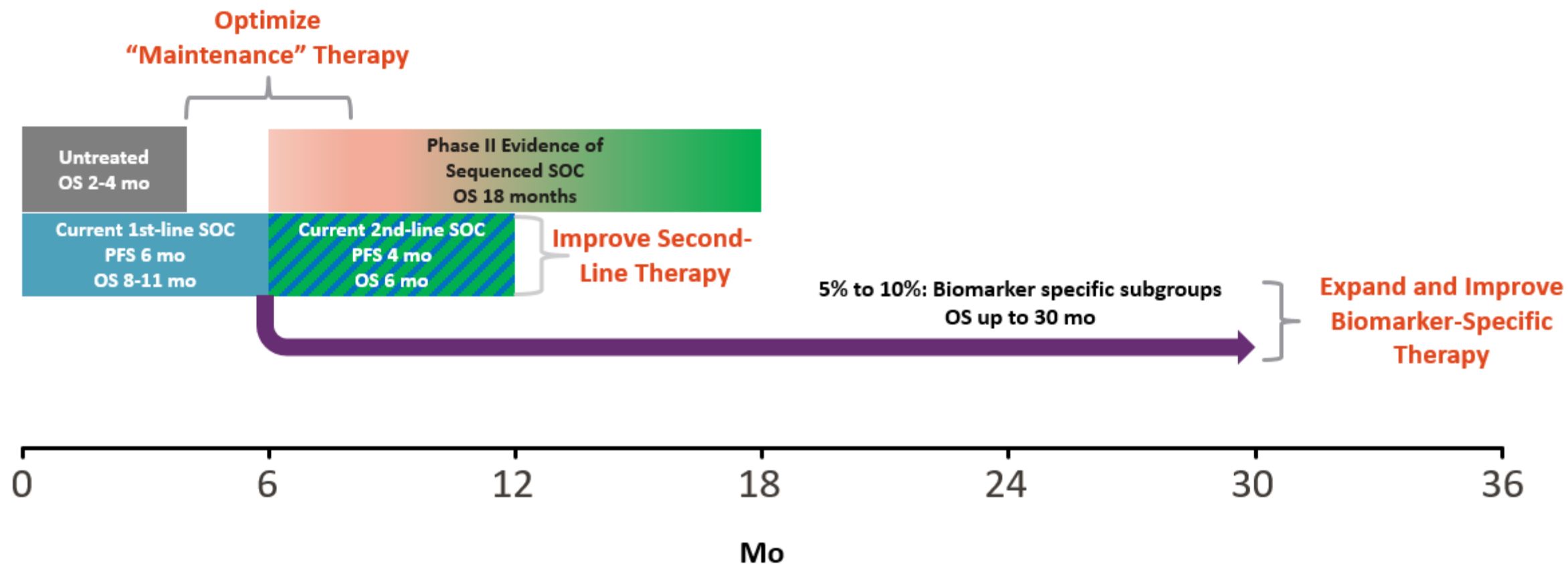


Singhi. Gastroenterology. 2019;156:2242. Pishvaian. Clin Cancer Res. 2018;24:5018. Heeke. JCO Precis Oncol. 2018;2018:10.1200/PO.17.00286. Aguire. Cancer Discov. 2018;8:1096. Witkiewicz. Nat Commun. 2015;6:6744. Lowery. Clin Cancer Res. 2017;23:6094. Waddell. Nature. 2015;518:495. Bailey. Nature. 2016;531:47. Biankin. Nature. 2012;491:399. Collisson. Nat Med. 2011;17:500. Pishvaian. Oncology (Williston Park). 2017;31:168. Jones. Clin Cancer Res. 2019;25:4674.

Pancreatic Cancer: Guideline Recommendations

- Germline testing recommended for any patient with PDAC
 - Multigene panel
- Tumor/somatic profiling recommended for all locally advanced/metastatic patients who are candidates for anticancer therapy to identify uncommon actionable mutations
 - Tissue testing optimal
 - cfDNA backup if insufficient tissue

Metastatic Pancreatic Cancer Survival



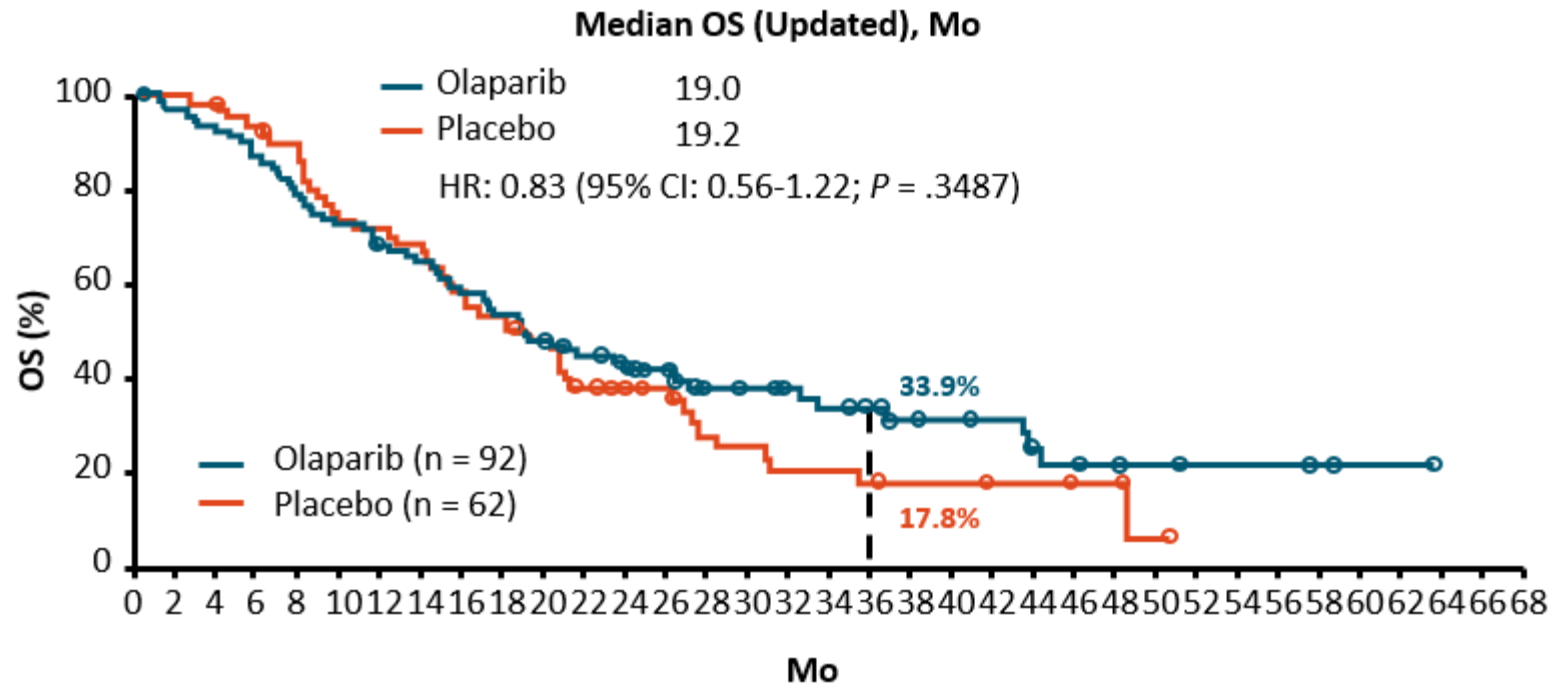
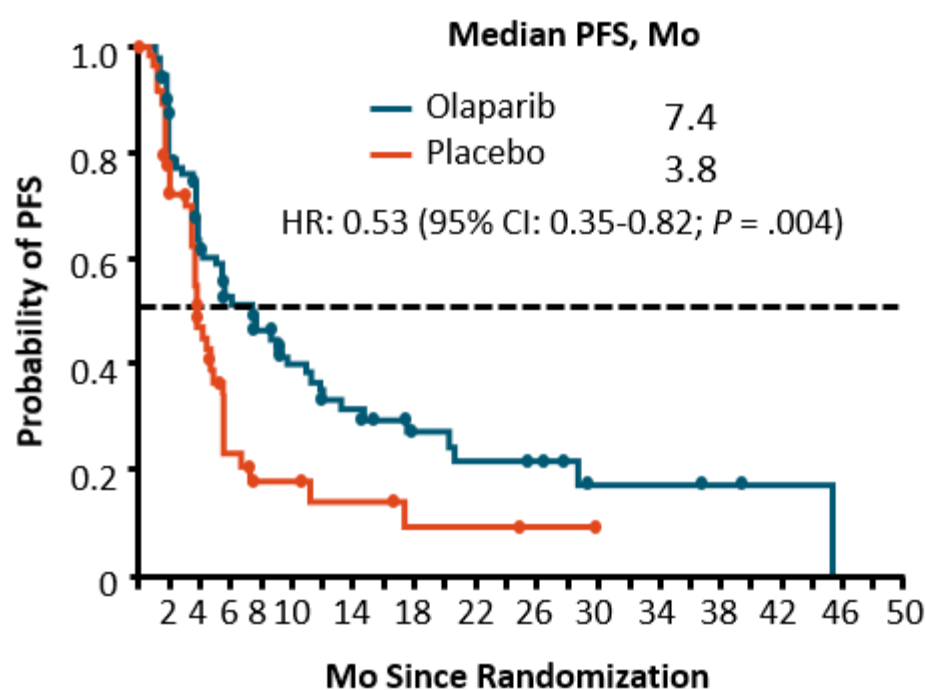
Germline and Somatic Testing in Pancreatic Cancer: Guidelines Recommendation

Germline testing is recommended for any patient with confirmed pancreatic cancer, using comprehensive gene panels for hereditary cancer syndromes

Tumor/somatic gene profiling is recommended for patients with locally advanced/metastatic pancreatic cancer (~80% of patients) who are candidates for anti-cancer therapy to identify uncommon but actionable mutations

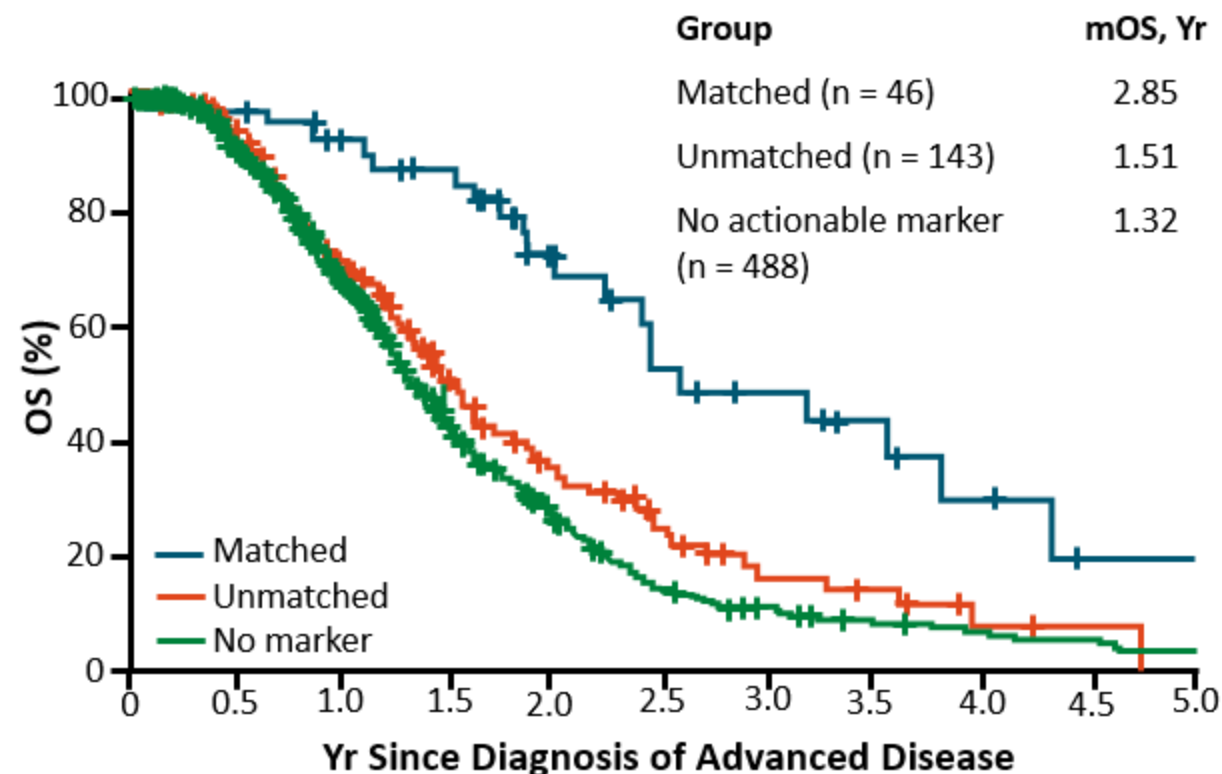
POLO: Maintenance Olaparib vs Placebo After First-line, Platinum-Based Therapy in Metastatic Pancreatic Cancer

- Randomized phase III trial of maintenance olaparib or placebo for patients with metastatic pancreatic cancer and deleterious/suspected deleterious *gBRCA1/2* mutation, ≥ 16 wk of first-line platinum-based therapy without progression (N = 154)



Gold Standard: Overall Survival Benefit

- 1028 pancreatic cancer patients
 - All underwent molecular profiling w/NGS
- 677 patients with outcome information
 - 189 with actionable findings
 - 46 received molecularly matched therapy
 - 143 received “unmatched” therapy
 - 488 with no actionable findings
- Overall survival
 - **Matched 1 y > unmatched**
 - **Matched 1.3 y > no actionable marker**



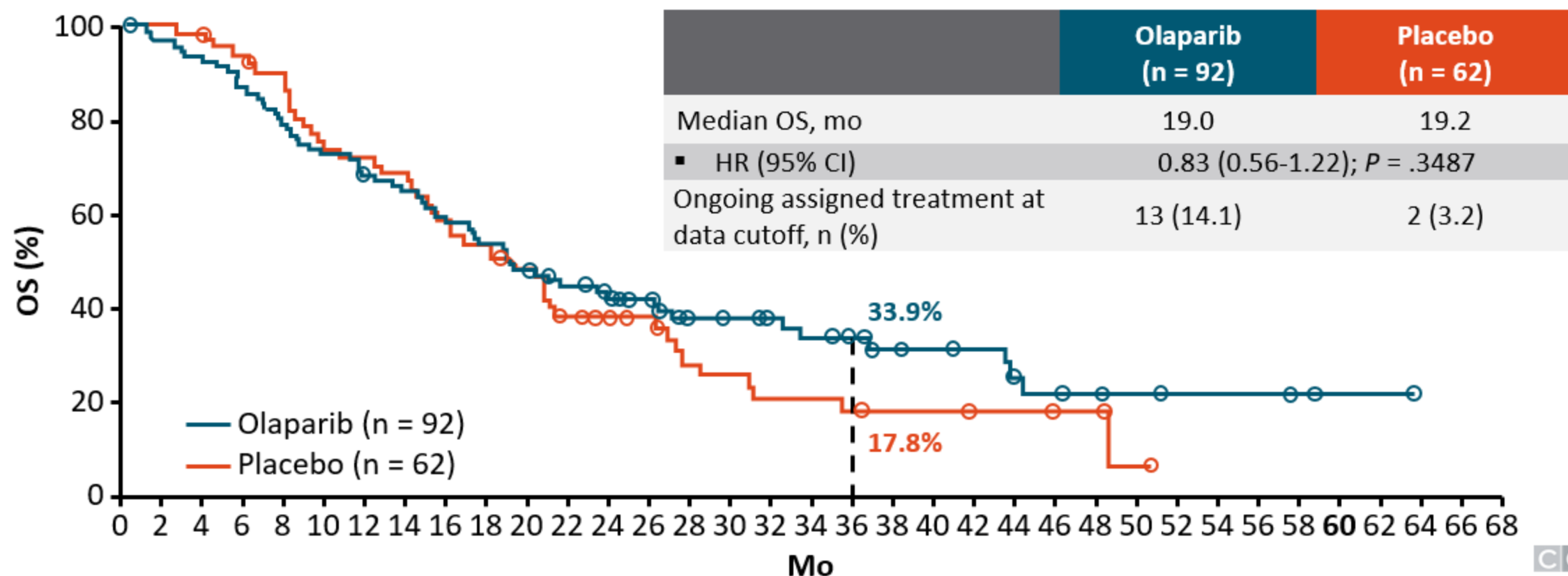
Comparison	HR (95% CI)
Matched vs unmatched (highly actionable)	0.42 (0.26-0.68); P = .0004
Matched vs no actionable marker	0.34 (0.22-0.53); P <.0001

cfDNA/ctDNA in Pancreatic Cancer

- Sensitivity is not too high^{1,2}
 - *KRAS* mutation detection: 30% to 75%
- But specificity is very high²
 - Concordance with *KRAS* mutations in tissue: 90% to 100%
- Can be used to track outcomes on serial samples
 - Decrease in mutated *KRAS* an early indicator of response to therapy³
- Might be useful as a prognostic and/or predictive marker³⁻⁵
 - Median OS: 3.2 mo for detectable vs 8.4 mo for undetectable ctDNA⁴

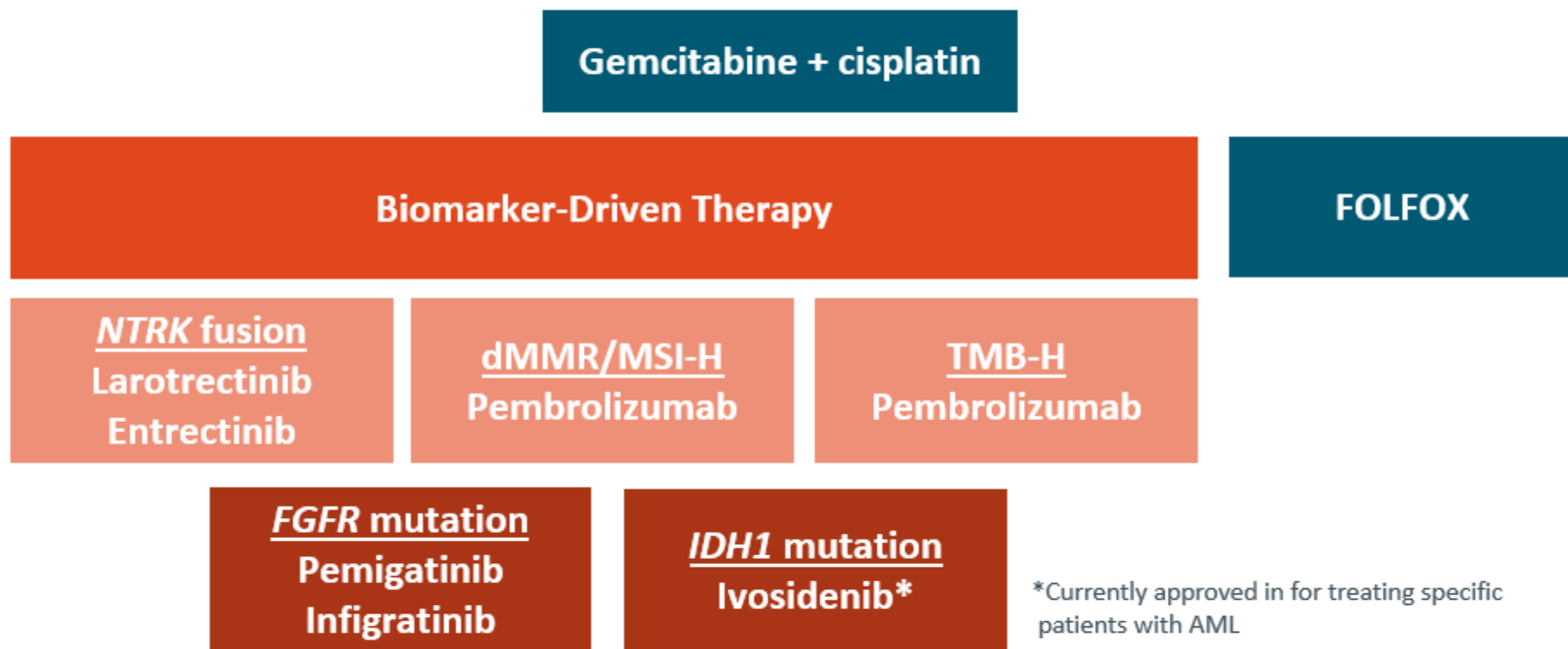
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Select Treatments for Advanced BTC

FDA-Approved Treatment Options



*Currently approved in for treating specific patients with AML

- Currently being assessed: gemcitabine + cisplatin + nab-paclitaxel or immunotherapy; TKIs + immunotherapy; Nal-IRI plus 5-FU/LV; trastuzumab (*HER2* amplified); dabrafenib + trametinib (*BRAF* V600E mutation)

Cholangiocarcinoma: Target-Rich Disease

Genetic Abnormality, n (%)	Intrahepatic CCA (n = 55)	Extrahepatic CCA (n = 20)	Odds Ratio	P Value	95% CI
<i>TP53</i>	16 (29.1)	9 (45)	0.506	.268	0.154-1.669
<i>KRAS</i>	13 (23.6)	8 (40)	0.469	.244	0.138-1.626
<i>ARID1A</i>	11 (20)	1 (5)	4.678	.164	0.599-214.83
<i>HER2</i>	1 (1.8)	5 (20)	0.058	.004	0.001-0.576
<i>PBRM1</i>	6 (10.9)	1 (5)	2.305	.667	0.252-112.51
<i>BAP1</i>	5 (9.1)	2 (10)	0.901	1	0.132-10.26
<i>FBXW7</i>	3 (5.5)	3 (15)	0.333	.333	0.041-2.719
<i>SMAD4</i>	2 (3.6)	5 (25)	0.333	.333	0.01-0.804
<i>IDH</i>	13 (23.6)	0	Infinite	.01	1.274-INF
MAP-ERK pathway	19 (34.5)	11 (55)	0.437	.121	0.133-1.389
mTOR pathway	14 (25.5)	8 (40)	0.517	.258	0.154-1.776
DNA repair pathway	9 (16.4)	8 (40)	0.299	.058	0.081-1.094
FGF pathway	7 (12.7)	1 (5)	2.741	.673	0.316-131.27
Chromatin modification pathway	18 (32.7)	3 (15)	2.724	.157	0.659-16.364

Targeted Approaches for BTC

Target	~Frequency in CC	Drug	Benefit	Status
MSI-H/dMMR	3%	Pembrolizumab	ORR: 40% ¹	Tumor agnostic approval
TMB >10 mut/Mb	2.4%	Pembrolizumab	ORR: 29% ¹	Tumor agnostic approval
<i>NTRK</i> fusion	1%	Larotrectinib	ORR: 75% ²	Tumor agnostic approval
<i>FGFR2</i> fusion	14% (intrahepatic)	Pemigatinib or infigratinib	ORR: 37% (pemigatinib) ³ ; 23% (infigratinib) ⁴	Cholangiocarcinoma approval
<i>IDH1</i> mutation	10%-20% (intrahepatic)	Ivosidenib	PFS HR: 0.37 ⁵	Cholangiocarcinoma approval
<i>BRAF</i> V600E	4%	Dabrafenib/trametinib	ORR: 41% (ROAR) ⁶	Open-label basket study
<i>HER2</i>	9% of BTC*†	Pertuzumab/trastuzumab	ORR: 23% (MyPathway) ⁷	Open-label basket study
<i>RET</i>	1%	Pralsetinib	Responses ⁸	2/2 PR in basket trial
<i>BRCA1/2</i> , DDR	20%*	PARP inhibitor	Responses reported	Case reports
ROS1	1%	Crizotinib	Response reported	Case reports

Most common in *extrahepatic or †GB.

1. Pembrolizumab PI. 2. Lenvatinib PI. 3. Pemigatinib PI. 4. Infigratinib PI. 5. Ivosidenib PI. 6. Subbiah. Lancet Oncol. 2020;21:1234.

7. Javle. Lancet Oncol. 2021;22:1290. 8. Subbiah. ASCO 2020. Abstr 109. Thornblade. Cancers (Basel). 2021;13:4062.

Select Ongoing Randomized Trials of Immunotherapy/Targeted Therapy for BTC

Trial	Regimens	Phase	Population
TOPAZ-1 (NCT03875235)	Gemcitabine + cisplatin ± durvalumab	III	Previously untreated unresectable/metastatic BTC (planned N = 757)
KEYNOTE-966 (NCT04003636)	Gemcitabine + cisplatin ± pembrolizumab	III	Previously untreated unresectable/metastatic BTC (planned N = 1048)
S1815 (NCT03768414)	Gemcitabine + cisplatin ± nab-paclitaxel	III	Previously untreated advanced BTC (planned N = 452)
IMbrave151 (NCT04677504)	Gemcitabine + cisplatin + atezolizumab ± bevacizumab	II	Previously untreated unresectable/metastatic BTC (planned N = 150)
HERIZON-BTC-01 (NCT04466891)	Zanidatamab	II	Previously treated advanced/metastatic HER2+ BTC (planned N = 100)
NCT04042831	Olaparib	II	Advanced BTC with DDR alterations (planned N = 36)
NCT03639935	Rucaparib + nivolumab	II	Previously treated advanced/metastatic BTC (planned N = 35)

Rationale for Immunotherapy in HCC (cont)

The liver is tolerogenic

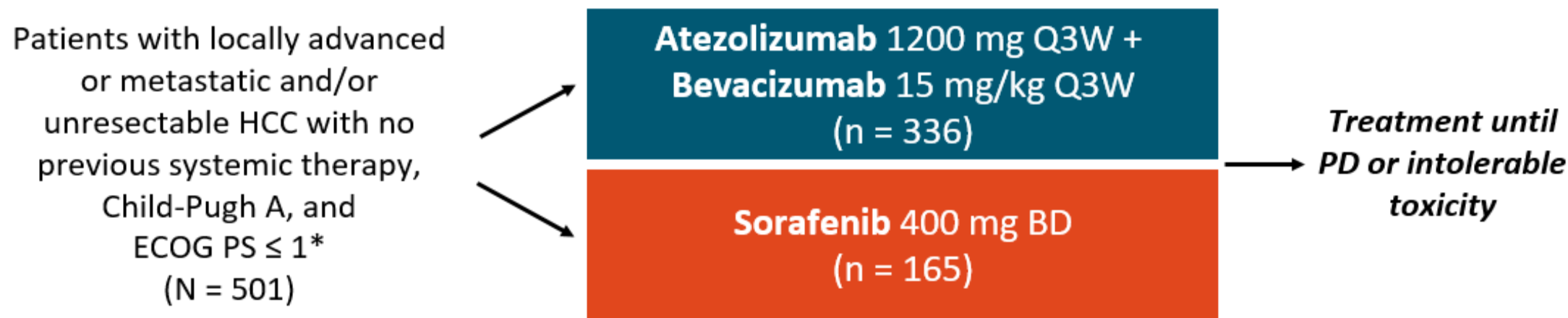
- Tolerates foreign antigens
- Increases risk of viral infection, which can lead to chronic liver damage
 - Decreases liver function
 - Increases cirrhosis
 - Increases cancer risk

HCC may be considered 2 concurrent diseases: cancer and liver disease

- Patients are frail and may not tolerate some therapies
- Effective but well tolerated treatment options are needed

IMbrave150: Atezolizumab + Bevacizumab vs Sorafenib for First-line Treatment of HCC

- Multicenter, randomized, open-label phase III trial^[1]
 - GO30140: randomized phase Ib study showed potential benefit of atezolizumab + bevacizumab for patients with advanced HCC (ORR 36%)^[2]



- Coprimary endpoints: OS and PFS

*Trial included subgroups of high-risk patients excluded from other contemporary phase III trials: ~ 40% had macrovascular invasion; specifically included patients with 50% hepatic involvement or main portal vein invasion or invasion of the portal vein branch contralateral to the primarily involved lobe.

Combining VEGF and PD-1/PD-L1 Inhibitors

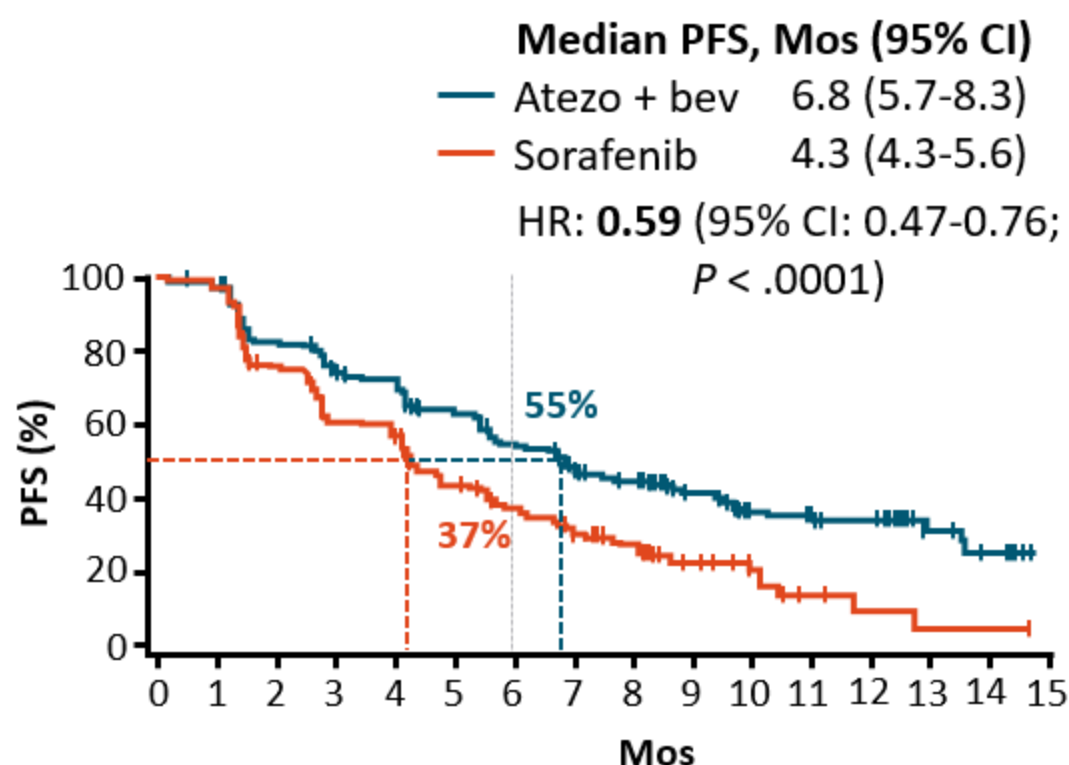
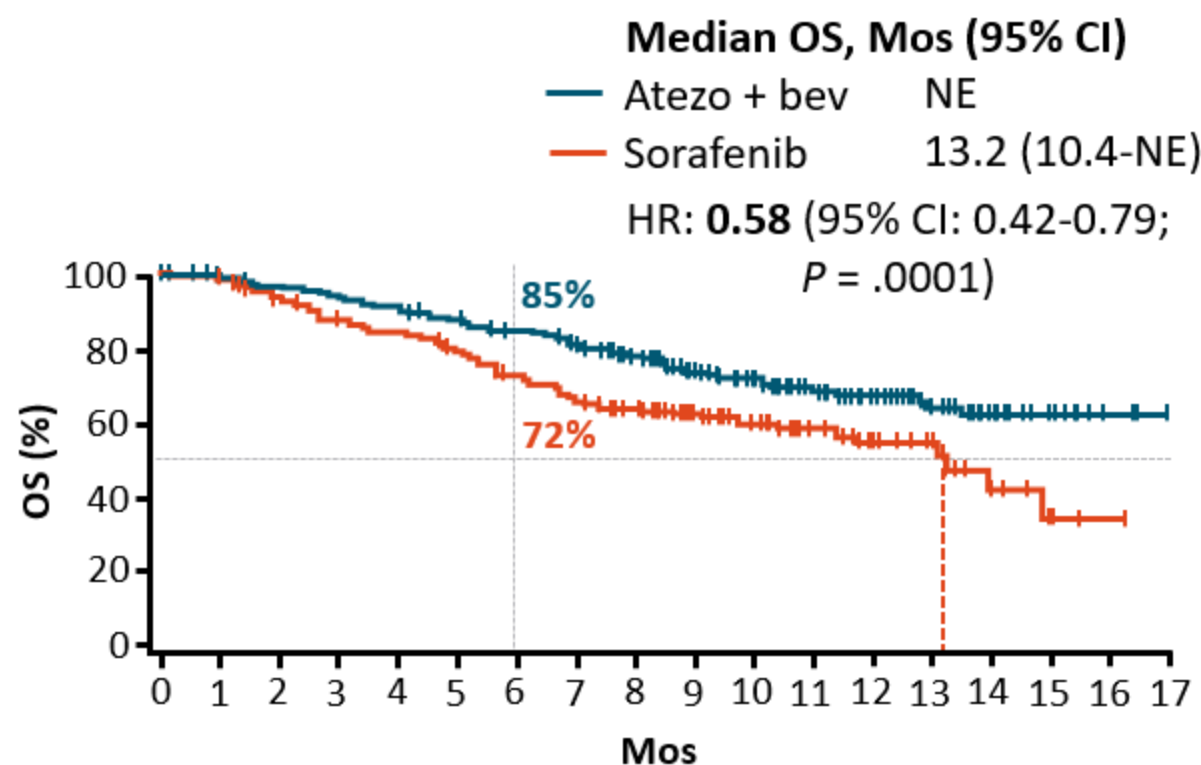
Atezolizumab (PD-L1 Inhibitor)

- Promotes T-cell activation by allowing B7.1 costimulation
- Restores anticancer immunity with activity further enhanced through VEGF-mediated immunomodulatory effects

Bevacizumab (VEGF Inhibitor)

- Promotes DC maturation
 - Normalizes tumor vasculature, increasing T-cell infiltration
 - Decreases activity of immunosuppressive cells (MDSCs, Tregs)
-
- **In combination, bevacizumab may enhance efficacy of atezolizumab by reversing VEGF-mediated immunosuppression to promote T-cell infiltration into tumor**

IMbrave150: OS, PFS, and Response

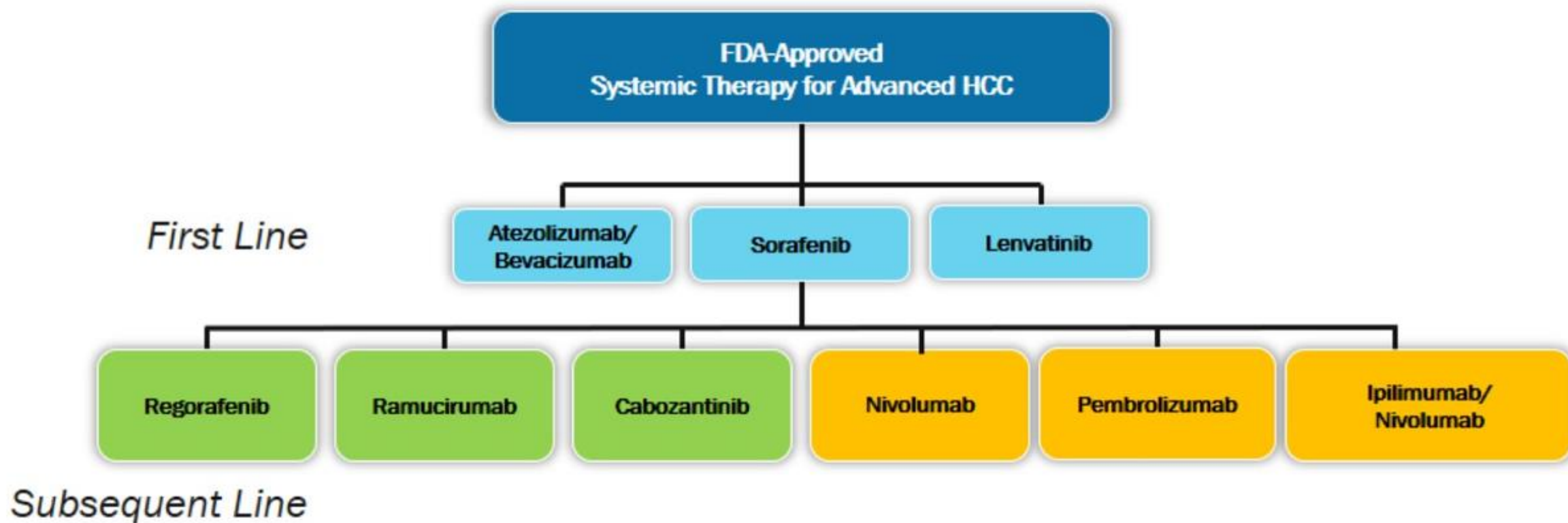


- ORR by HCC-specific modified RECIST with atezo + bev vs sorafenib: 33.2% vs 13.3%; CR rate, 10.2% vs 1.9%

Median follow-up: 8.6 mos.

Finn. NEJM. 2020;382:1894.

Treatment Landscape of Advanced HCC



NCCN Guidelines for HCC

First-line Systemic Therapy

Preferred regimens

- Atezolizumab-bevacizumab (Child-Pugh Class A only) (category 1)

Other recommended regimens

- Sorafenib (Child-Pugh Class A [category 1] or B7)
- Lenvatinib (Child-Pugh Class A only) (category 1)

Useful in certain circumstances

- Nivolumab (if ineligible for TKIs or other antiangiogenic agents) (category 2B)
- FOLFOX (category 2B)

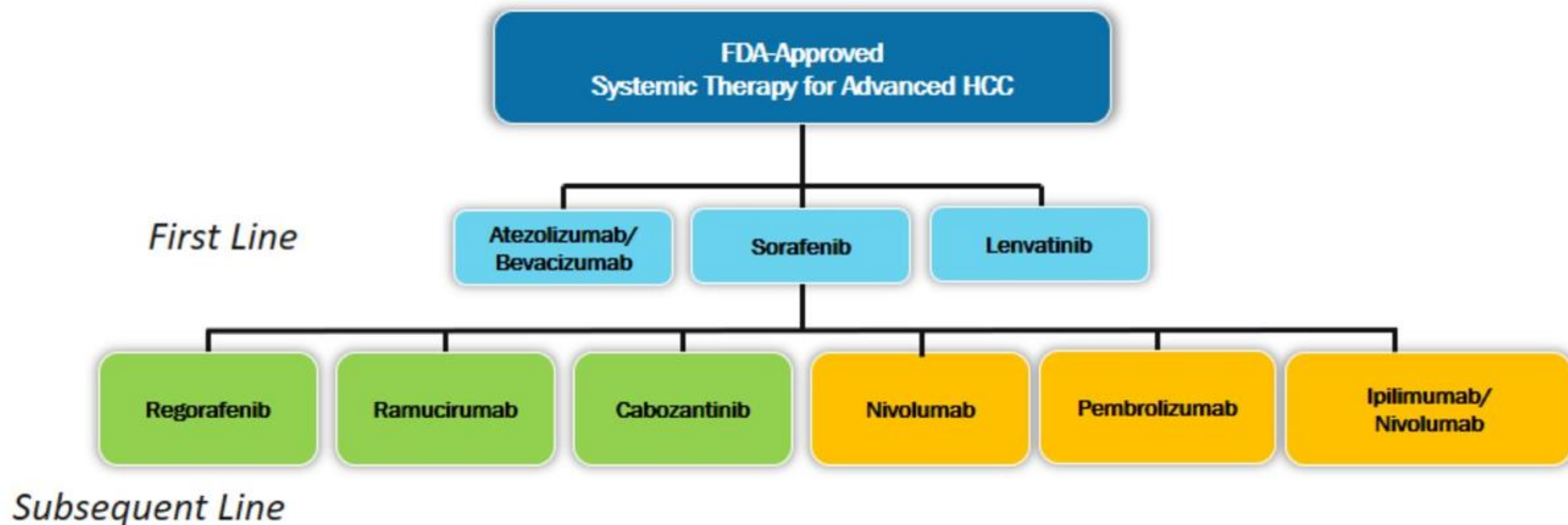
Subsequent-line therapy if disease progression

- Regorafenib (Child-Pugh Class A only) (category 1)
- Cabozantinib (Child-Pugh Class A only) (category 1)
- Ramucirumab (AFP \geq 400 ng/mL only) (category 1)
- Lenvatinib (Child-Pugh Class A only)
- Sorafenib (Child-Pugh Class A or B7)

Other recommended regimens

- Nivolumab (Child-Pugh Class A or B)
- Nivolumab-ipilimumab (Child-Pugh Class A only)
- Pembrolizumab (Child-Pugh Class A only) (category 2B)

Treatment Landscape of Advanced HCC



NCCN Guidelines. Hepatobiliary Cancers. V2.2021.

NCCN Guidelines for HCC

1L systemic therapy

Preferred regimens

- Atezolizumab-bevacizumab (Child-Pugh Class A only) (category 1)

Other recommended regimens

- Sorafenib (Child-Pugh Class A [category 1] or B7)
- Lenvatinib (Child-Pugh Class A only) (category 1)

Useful in certain circumstances

- Nivolumab (if ineligible for TKIs or other antiangiogenic agents) (Child-Pugh Class A or B; category 2B)
- FOLFOX (category 2B)

Subsequent-line therapy if disease progression

- Regorafenib (Child-Pugh Class A only) (category 1)
- Cabozantinib (Child-Pugh Class A only) (category 1)
- Ramucirumab (AFP \geq 400 ng/mL only) (category 1)
- Lenvatinib (Child-Pugh Class A only)
- Sorafenib (Child-Pugh Class A or B7)

Other recommended regimens

- Nivolumab (Child-Pugh Class B only)
- Nivolumab-ipilimumab (Child-Pugh Class A only)
- Pembrolizumab (Child-Pugh Class A only) (category 2B)
- Dostarlimab-gxly for MSI-H/dMMR tumors (category 2B)

Immunotherapy for Advanced HCC

Nivolumab

- PD-1 inhibitor
- FDA-approved (2017) as second-line treatment of HCC in patients previously receiving sorafenib^[a]
- Phase 1/2 CheckMate 040 study; ORR: 15%^[b]

Nivolumab + Ipilimumab

- Ipilimumab is a CTLA-4 blocking antibody
- FDA-approved (2020) as second-line treatment of HCC in patients previously on sorafenib^[c]
- Phase 1/2 CheckMate 040 study; ORR: 31%^[d]

Pembrolizumab

- PD-1 inhibitor
- FDA-approved (2018) as second-line treatment of HCC in patients previously receiving sorafenib^[e]
- Phase 1/2 study KEYNOTE-224: ORR 17%

a. Opdivo® (nivolumab). [PI]. 2018; b. El-Khoueiry AB, et al. *Lancet*. 2017;389:2492-2502; c. US FDA Drug Approvals. [fda.gov/drugs](https://www.fda.gov/drugs). Accessed March 21, 2020. d. Yau T, et al. *J Clin Oncol*. 2019;37(suppl): Abstract 4012. e. Keytruda® (pembrolizumab) [PI]. 2021; f. Zhu AX, et al. *Lancet Oncol*. 2018;19:940-952.

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