

Predictive and Prognostic Immunohistochemistry in the Evaluation of Colorectal Cancer

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No relevant disclosures or
conflicts of interest

Objectives

- Discuss interpretation and clinical significance of immunohistochemical biomarkers in colorectal cancer
 - Mismatch repair (MMR) proficiency and deficiency
 - Checkpoint inhibitor therapy (PD-L1)
 - Her2neu amplification

Immunohistochemical Biomarkers

- **Mismatch repair (MMR) proficiency and deficiency**
- PD-L1
- Her2neu amplification

Mismatch Repair Proteins

- Correct single base mismatches in DNA microsatellites
- MLH1, PMS2, MSH2, MSH6
- Presence of all 4 indicates microsatellite stable (mismatch repair proficient - pMMR)*
- Staining loss of 1 or more indicates microsatellite instability (mismatch repair deficient - dMMR)

* Like all rules, there are exceptions. Stay tuned.

MLH1

PMS2

MSH2

MSH6

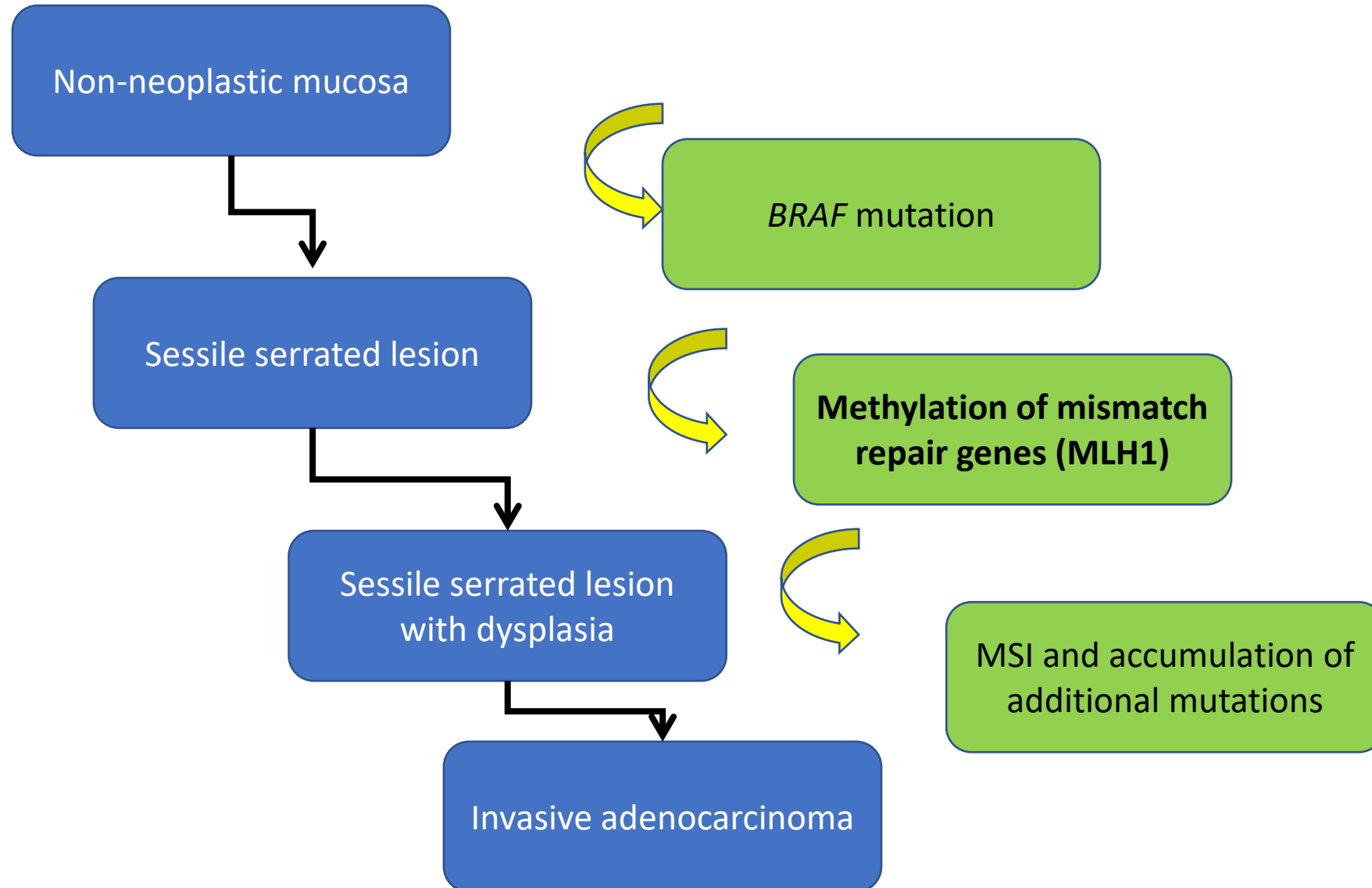
Lynch Syndrome

- Autosomal dominant
- Germline mutations in 1 of the 4 mismatch repair genes or in *EPCAM*
- Universal screening advocated by Centers for Disease Control and Prevention, National Comprehensive Cancer Network, American College of Gastroenterology, American Society of Clinical Oncology

Screening for Lynch Syndrome

- Why do we do it?
 - To enter patients and their families into monitoring programs
 - To plan appropriate surgery
 - Subtotal colectomy
 - Only possible when the biopsy is tested

Sporadic MSI Cancers and the Serrated Neoplastic Pathway



Identifying Sporadic MSI Colon Cancers

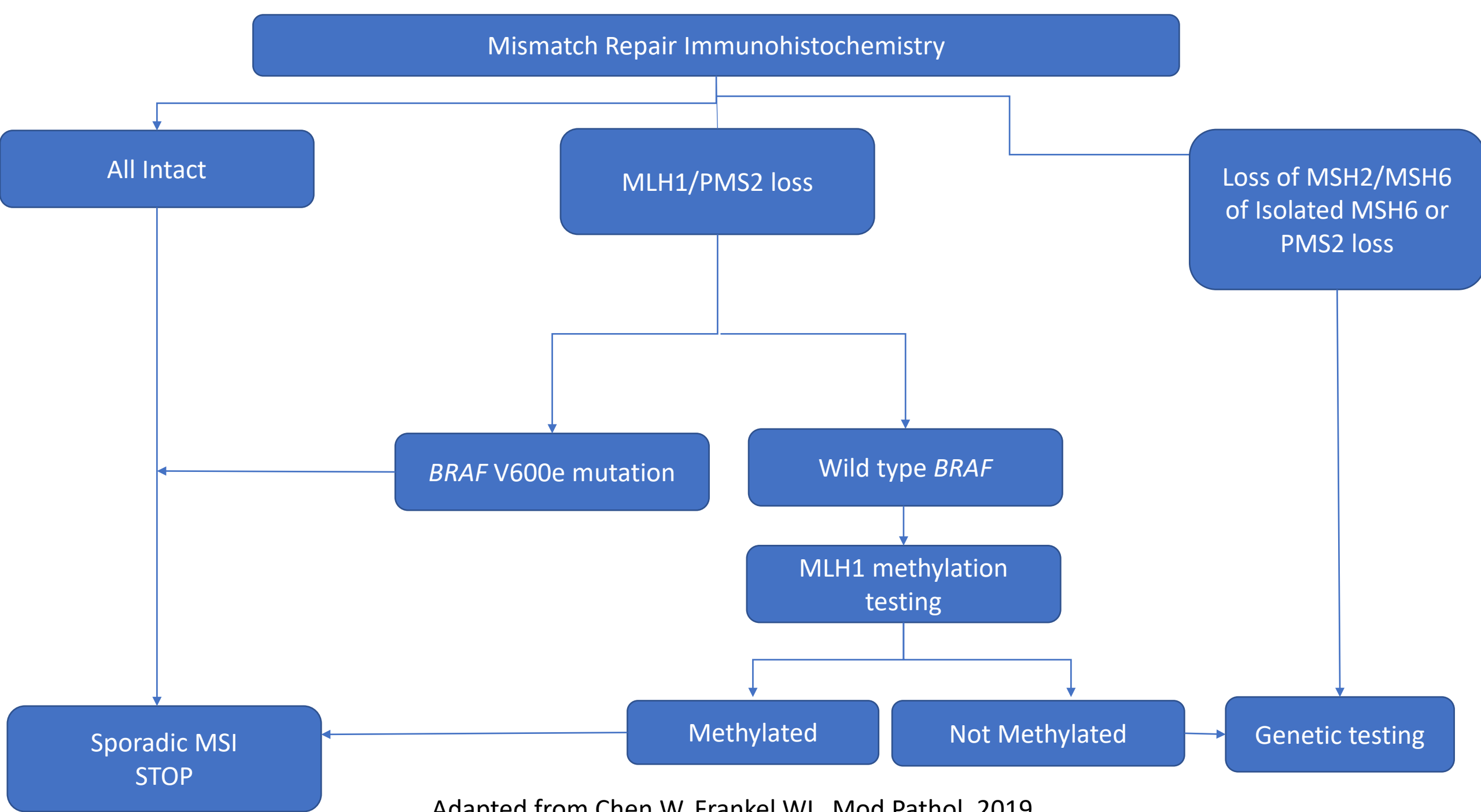
- Why do we do it?
 - Overall better prognosis
 - Poor response to Fluorouracil-based therapy
 - Benefit from immune checkpoint inhibitors

Screening for Mismatch Repair Deficiency: How Do We Do It?

- Most institutions perform immunohistochemistry (IHC) on biopsy samples
- Discordant results between IHC and PCR rare
- PCR is warranted when IHC results are equivocal
- MSI status is included in some next generation sequencing panels

Hissong E, et al. Mod Pathol. 2018

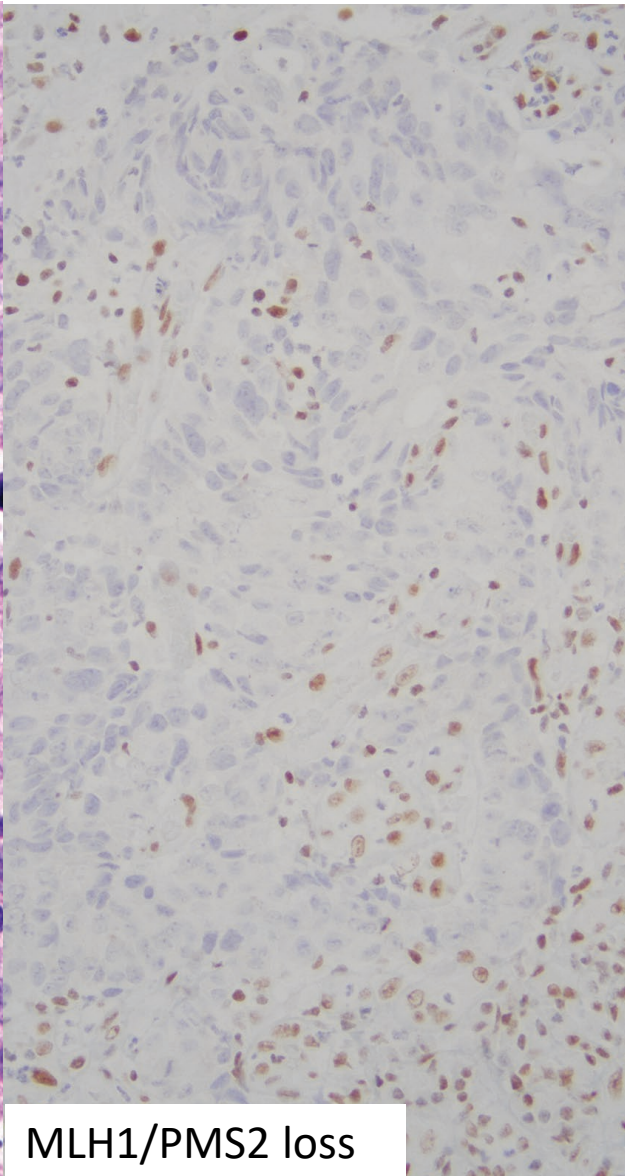
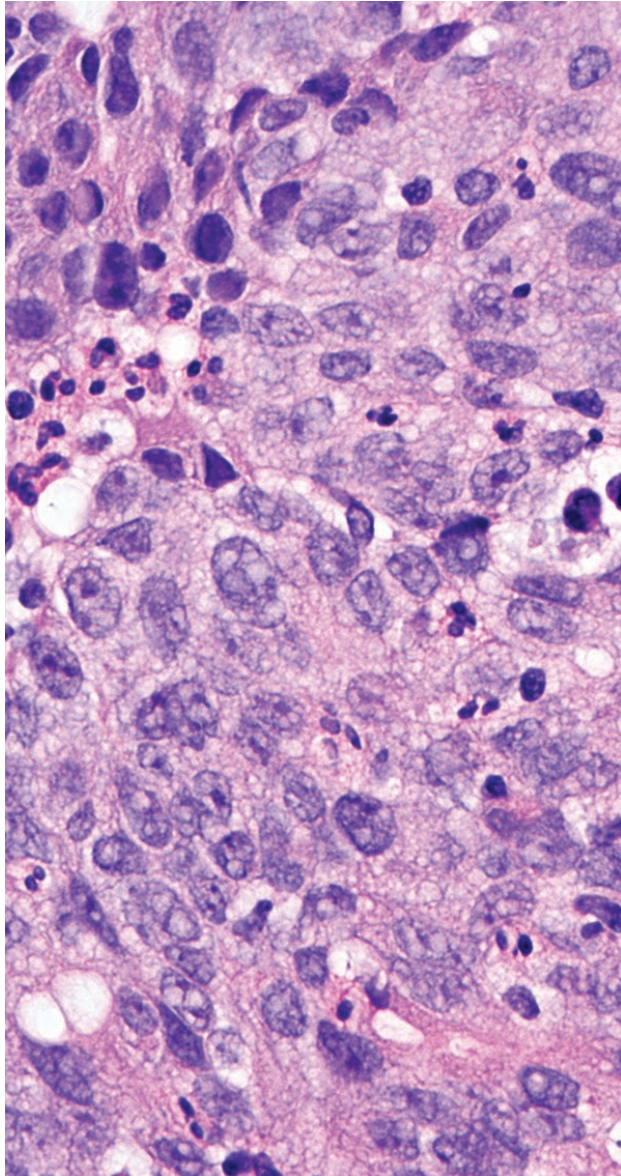
Hechtman JF, et al. Mod Pathol. 2020 May;33(5):871-879



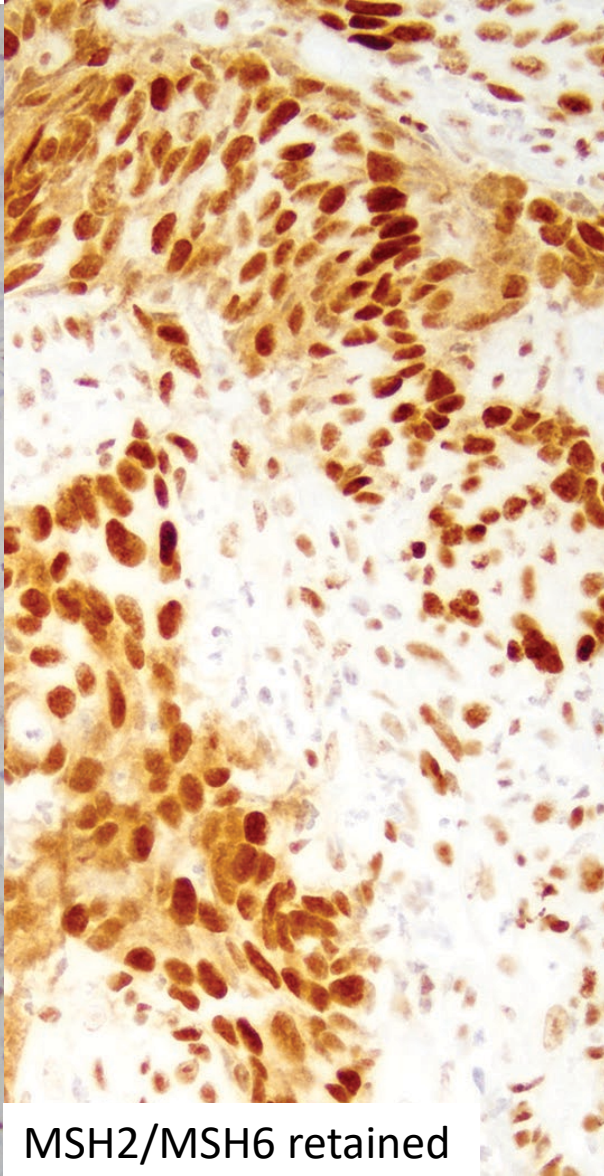
Adapted from Chen W, Frankel WL. Mod Pathol. 2019.

MMR Protein
Immunohistochemistry
Interpretation and Pitfalls

Mismatch Repair Protein Immunohistochemistry



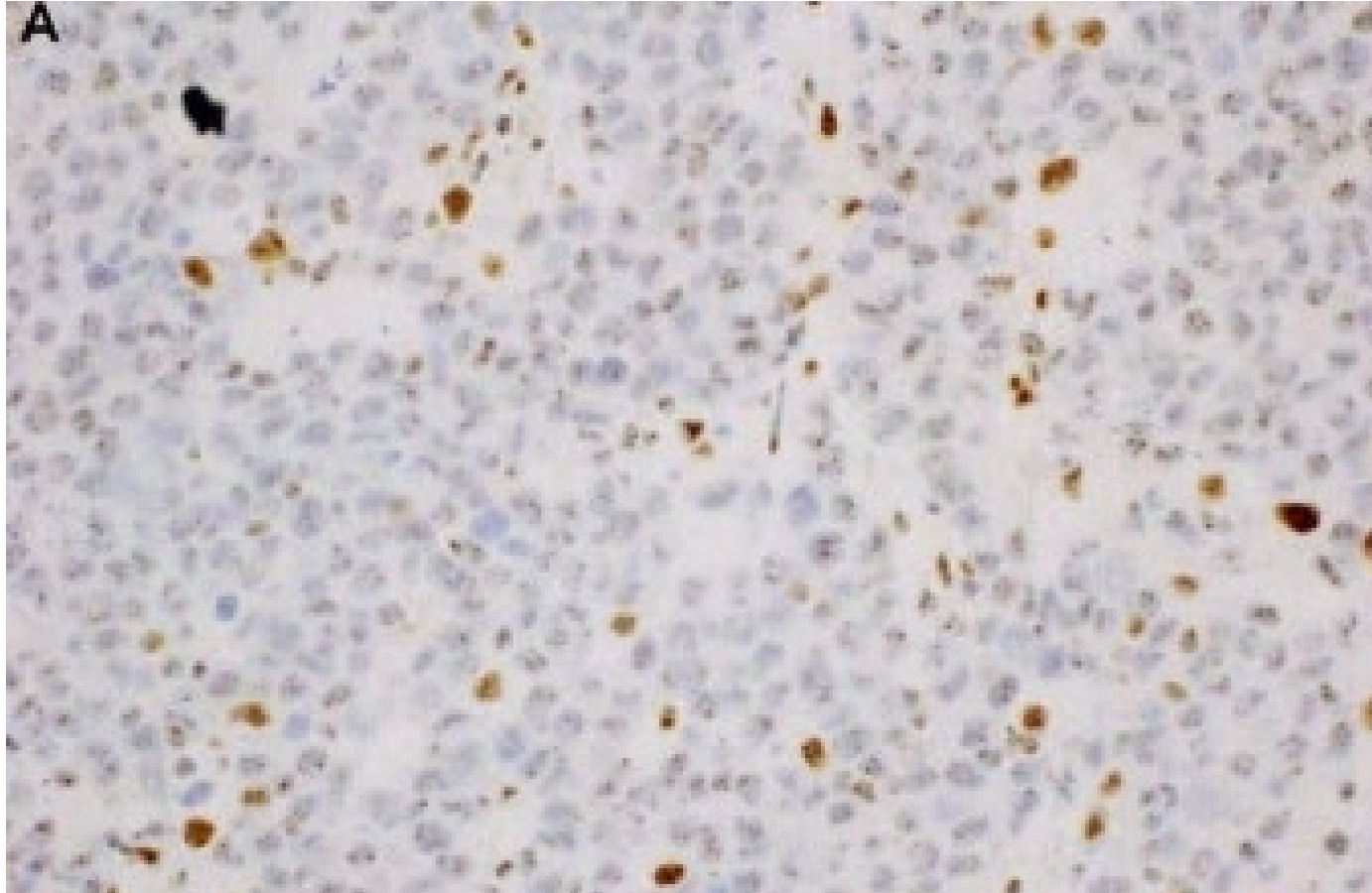
MLH1/PMS2 loss



MSH2/MSH6 retained

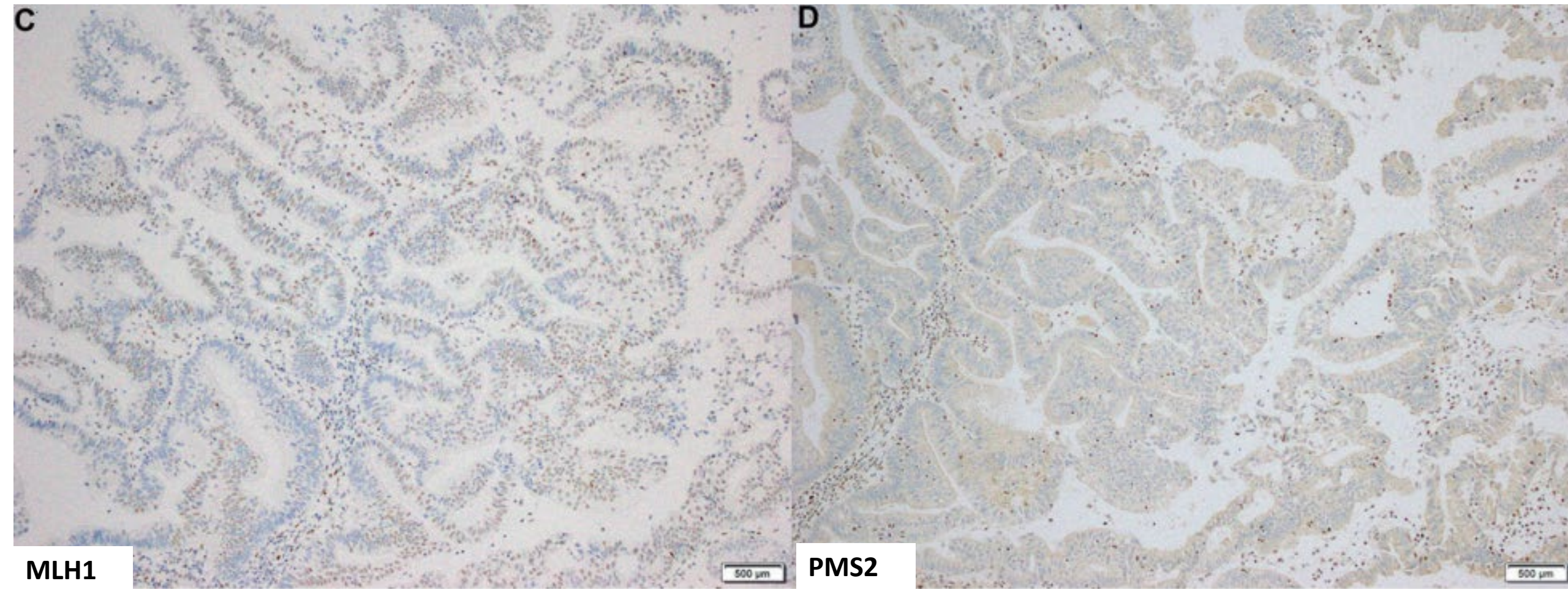
- Intact staining should be AT LEAST as strong as the internal control
 - Any proliferating cell
- No official threshold for proportion of cells staining
 - Some experts endorse $\geq 5\%$

“Discordant” IHC



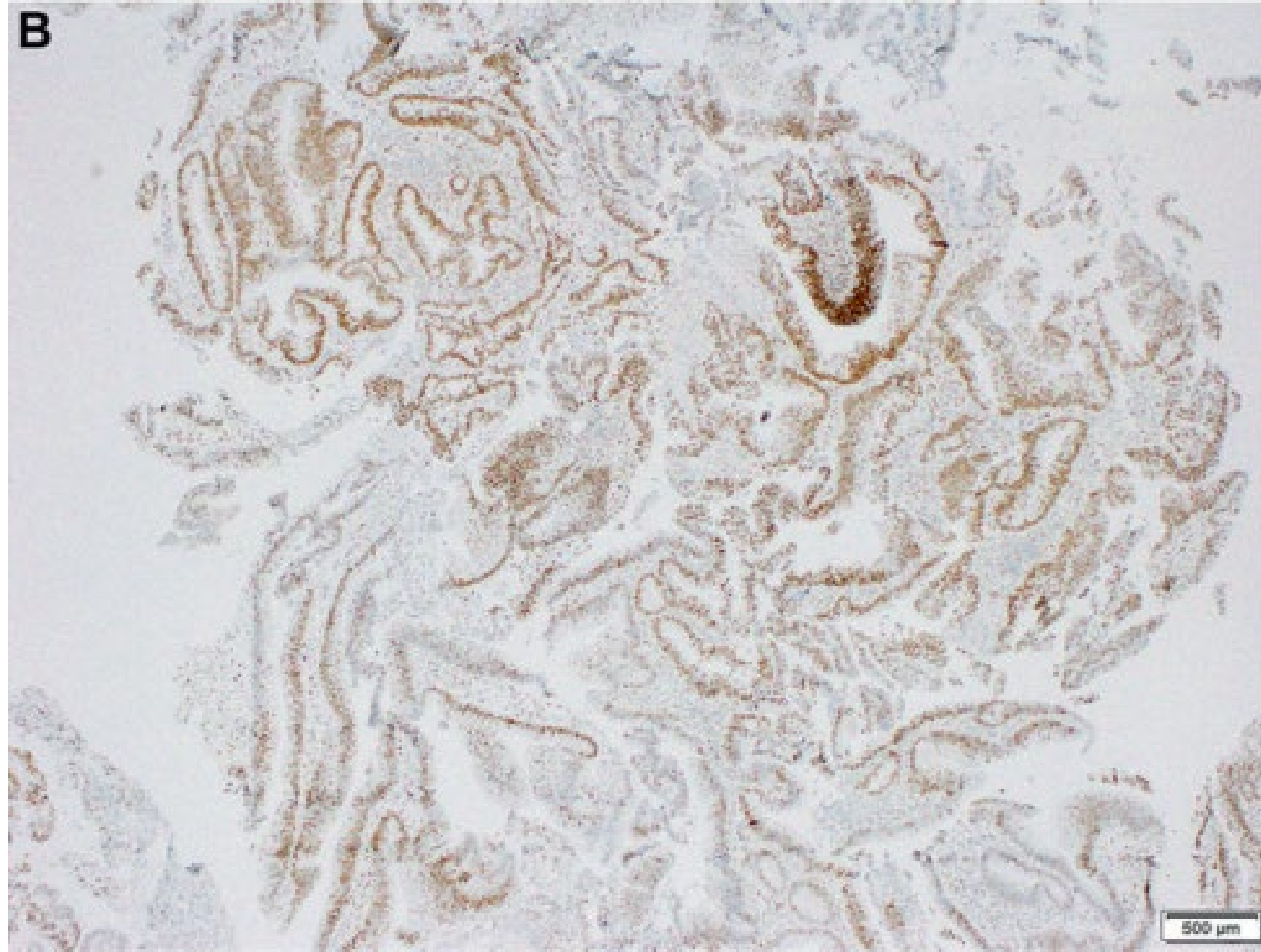
Dot-like MLH1 staining may be seen in cases with germline, somatic, or promotor hypermethylation

“Discordant” IHC



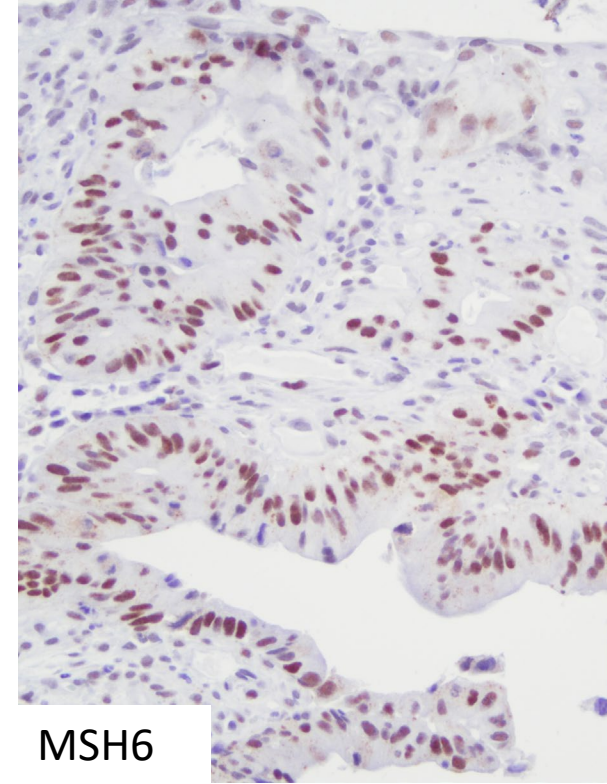
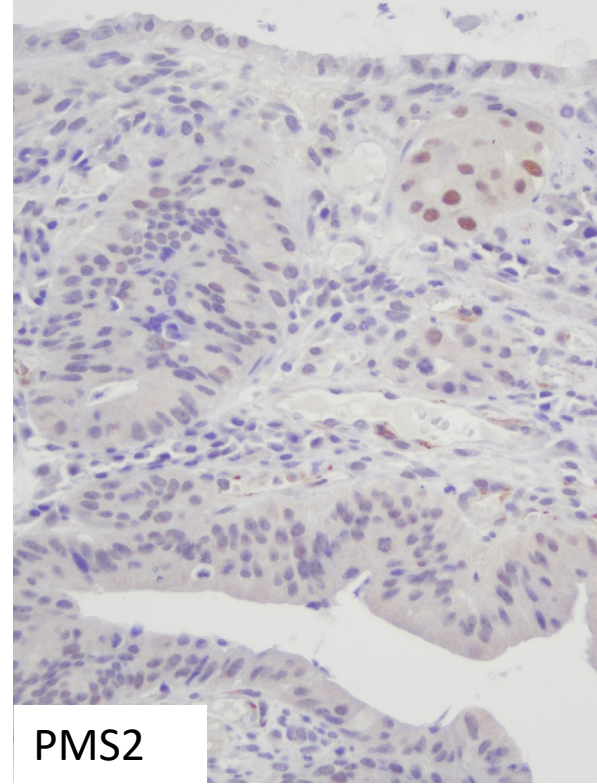
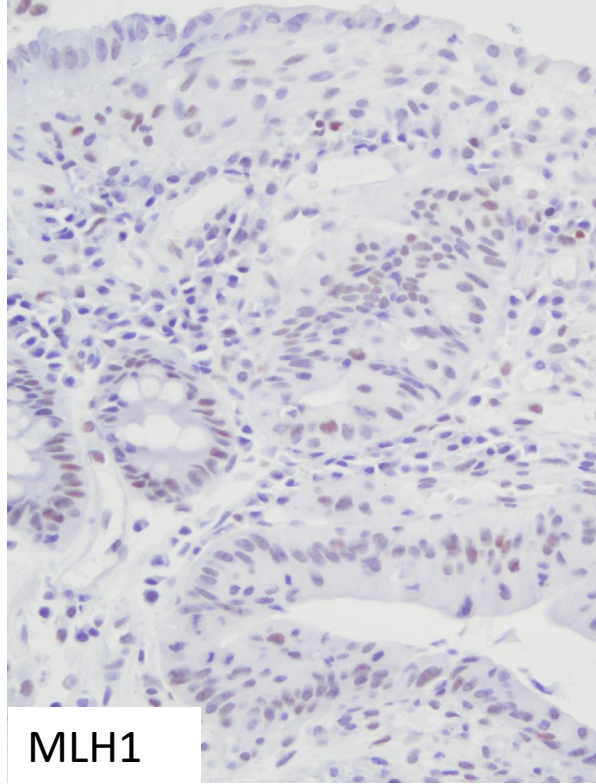
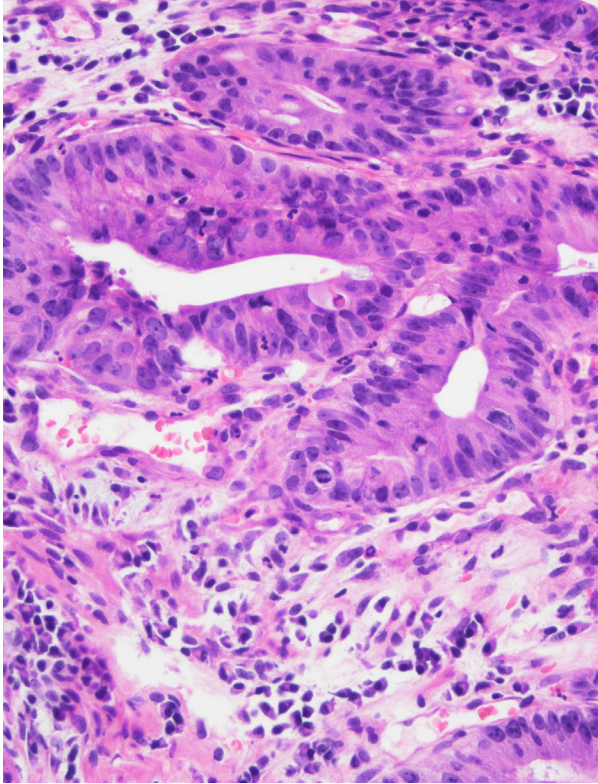
Weak MLH1 staining and PMS2 loss in a case with MLH1 promoter hypermethylation

Discordant IHC

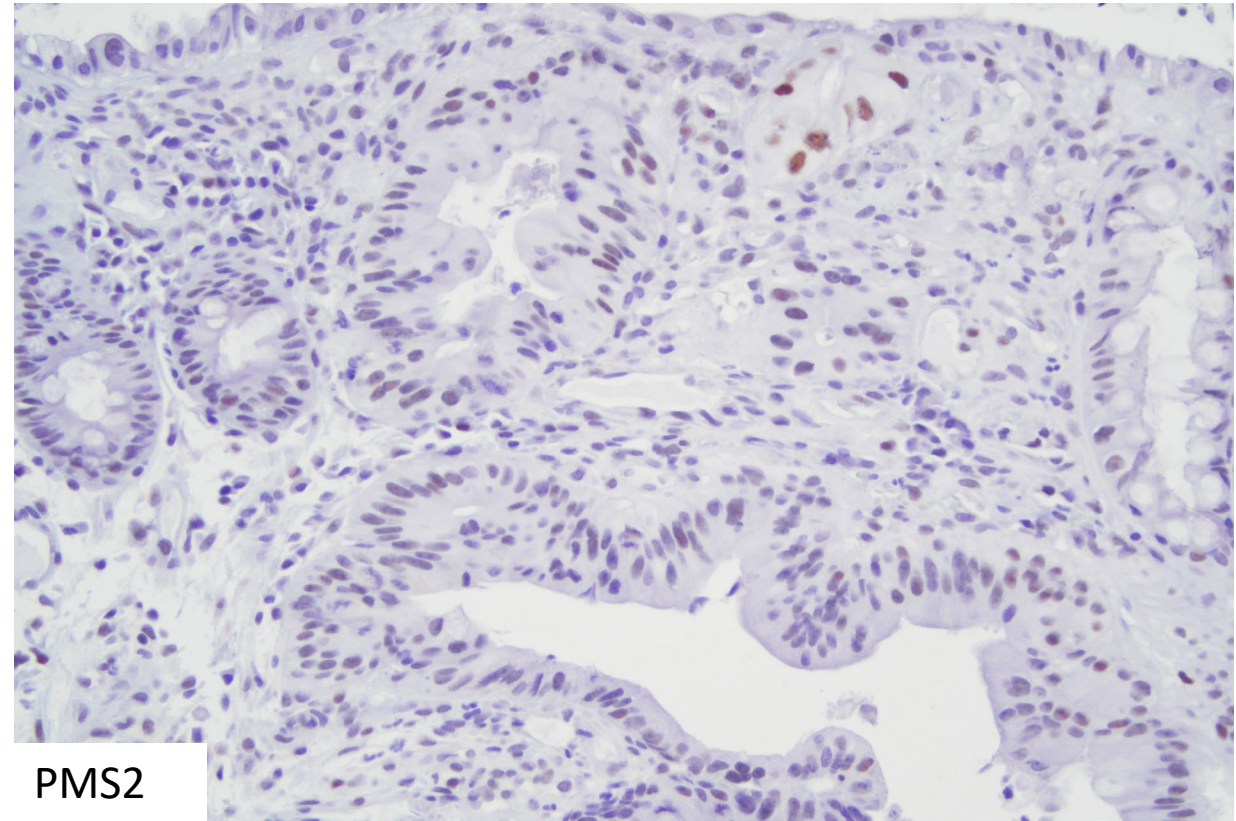
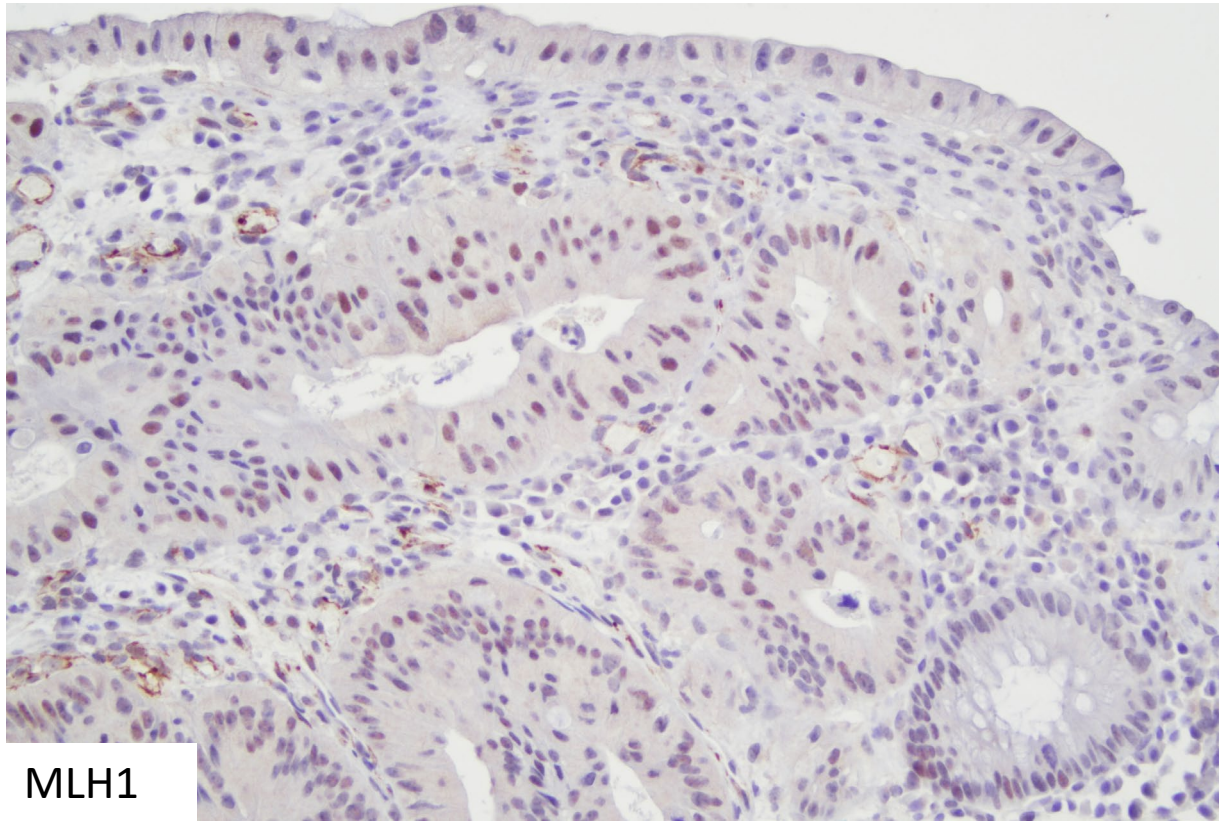


Retained MSH6 staining in a case with two somatic *MSH6* mutations

A Case From My Signout...



Repeat Stains



We could probably convince ourselves of retained staining in the tumor cells, but the control is still pretty weak

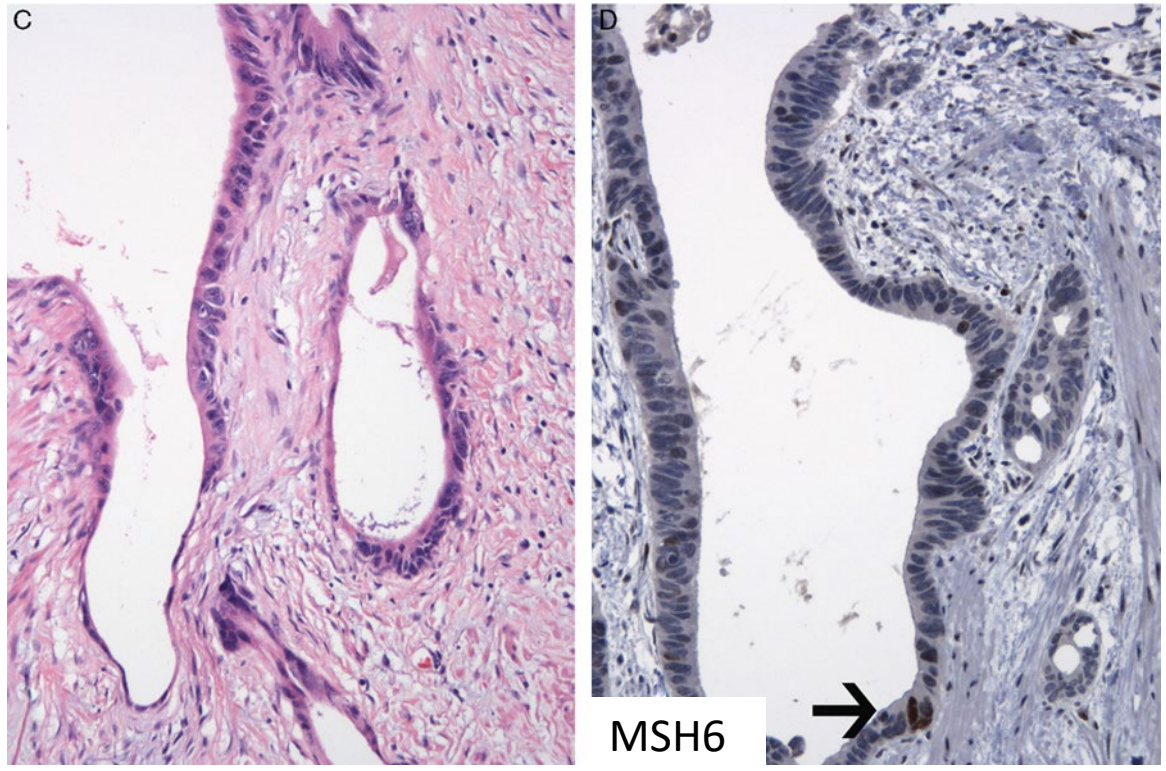
Referred for PCR → MSS

Summary

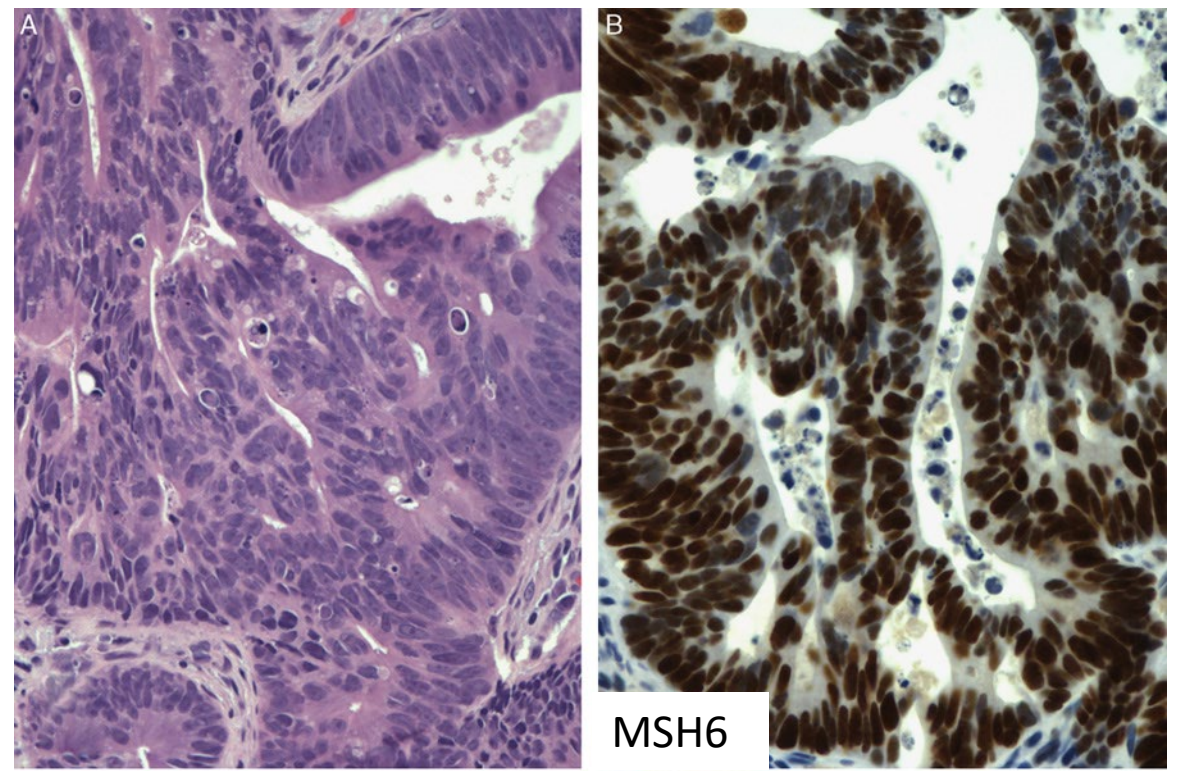
- Staining must be AT LEAST as strong as the internal control
- DO NOT interpret cases without a valid internal control
- Mutated mismatch repair proteins may be immunoreactive but this is rare
- Most discrepancies result from use of outdated criteria for interpreting IHC and can be solved by further testing (i.e. PCR)

One last pitfall... Post-Neoadjuvant Therapy

Resection



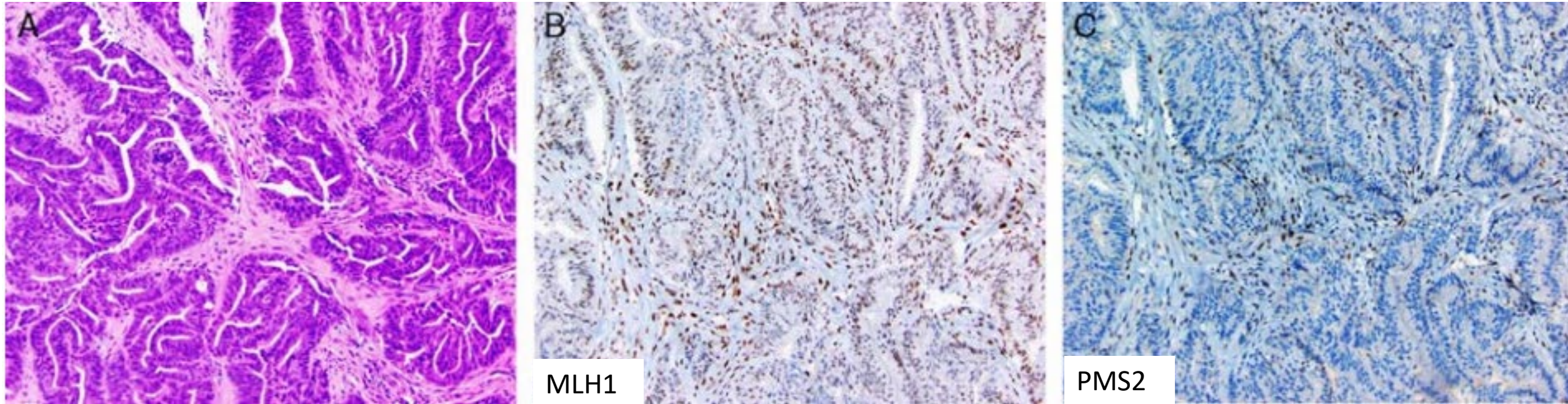
Biopsy



MMR Protein Immunohistochemistry

Some unusual patterns to know about

Isolated PMS2 Loss Without a Germline *PMS2* Mutation



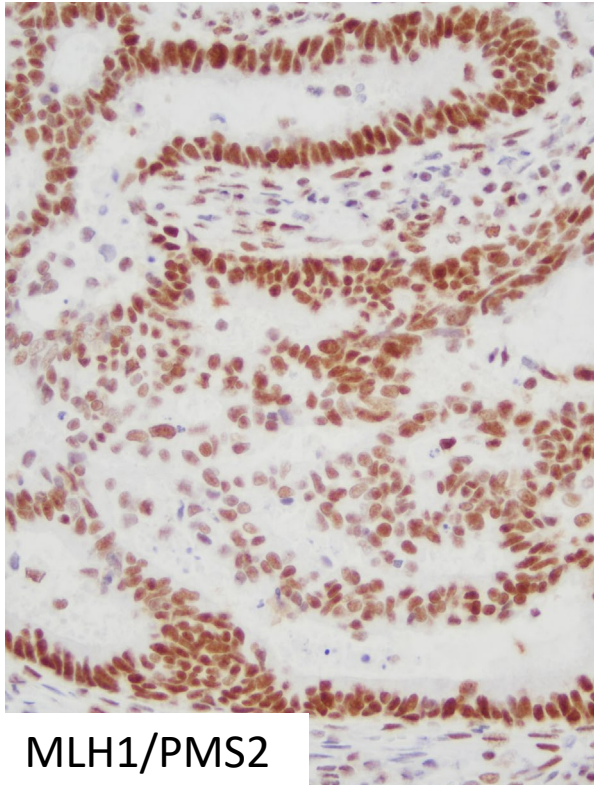
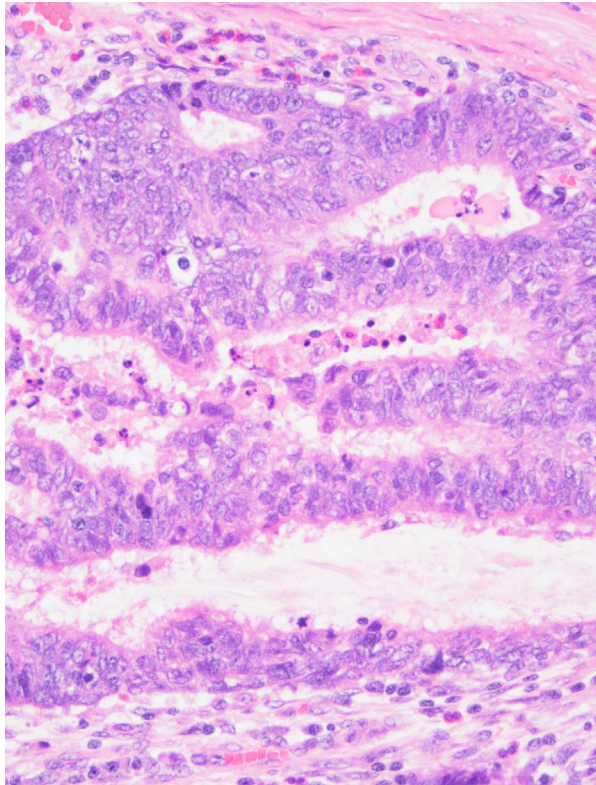
Dudley B, et al. Am J Surg Pathol. 2015 Aug;39(8):1114-20.

- Germline pathogenic variants in *MLH1*
- Bottom line: This pattern is dMMR
 - If *PMS2* is normal, *MLH1* should be tested

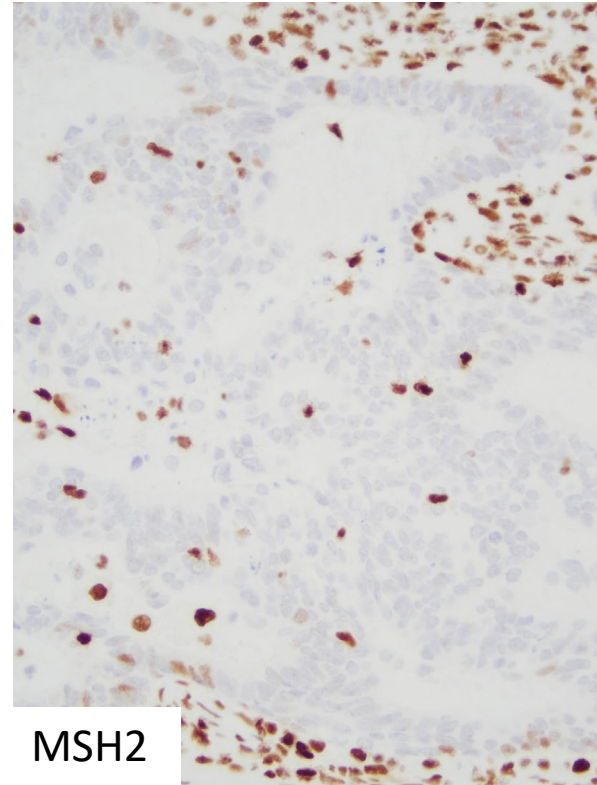
Dudley B, et al. Am J Surg Pathol. 2015 Aug;39(8):1114-20.

Rosty C, et al. BMJ Open. 2016 Feb 19;6(2):e010293.

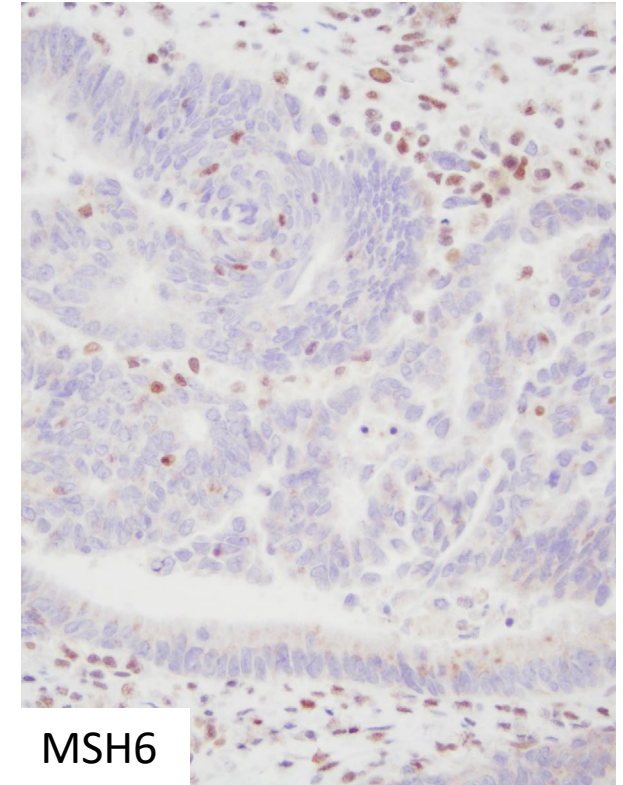
Another Case From My Signout...



MLH1/PMS2

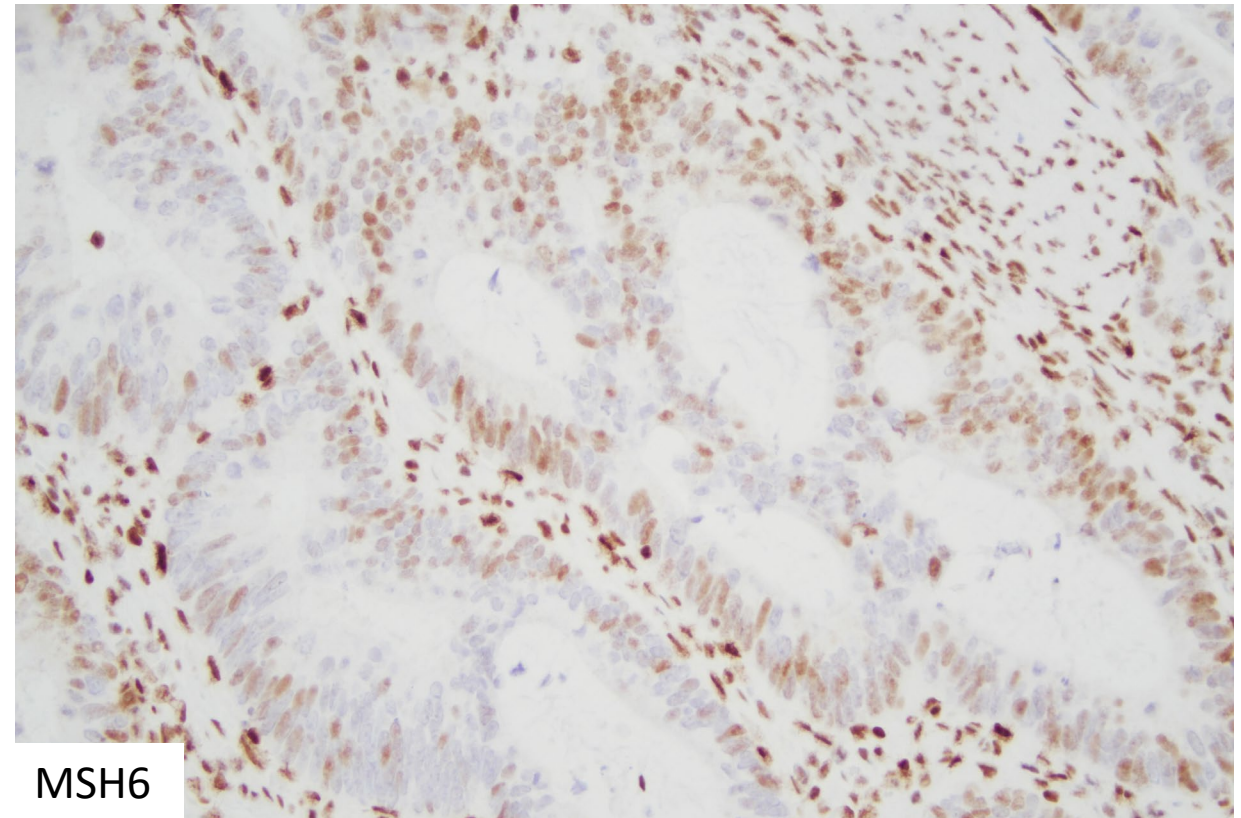
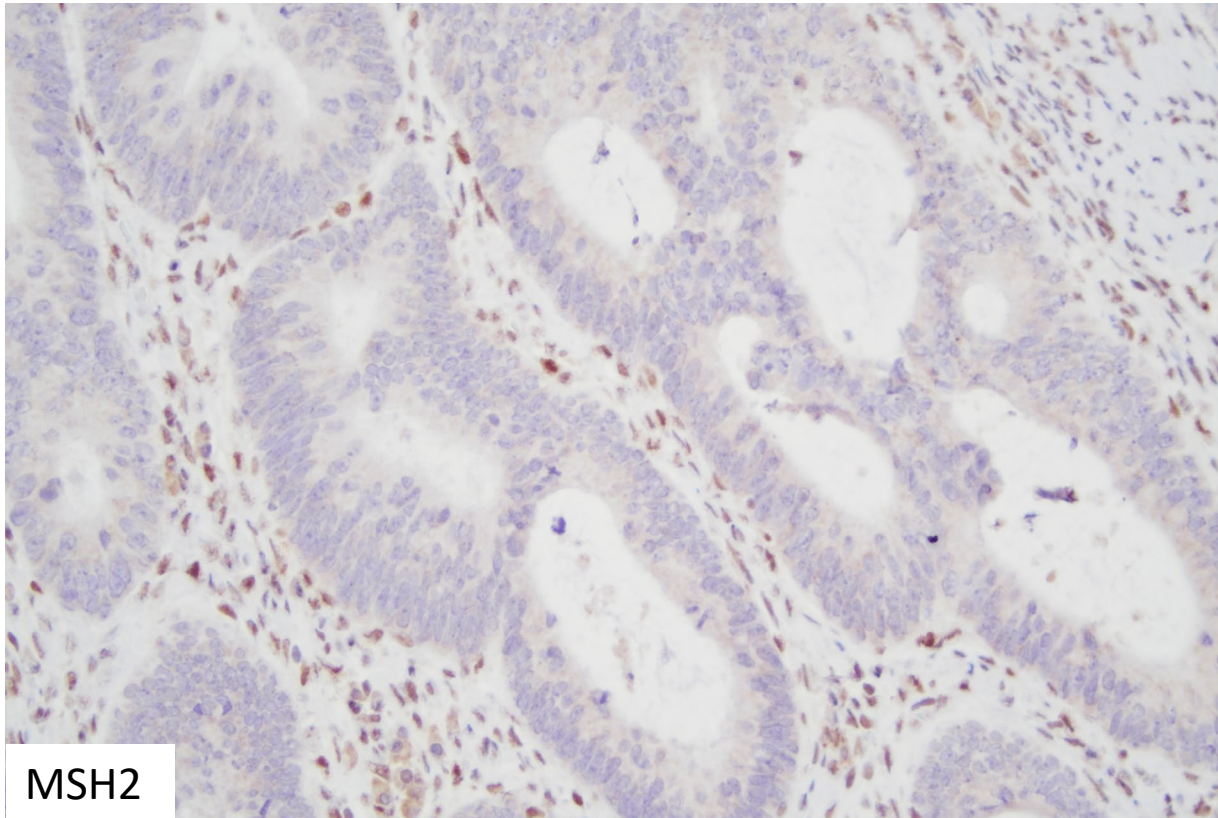


MSH2



MSH6

Same Case, Different Area

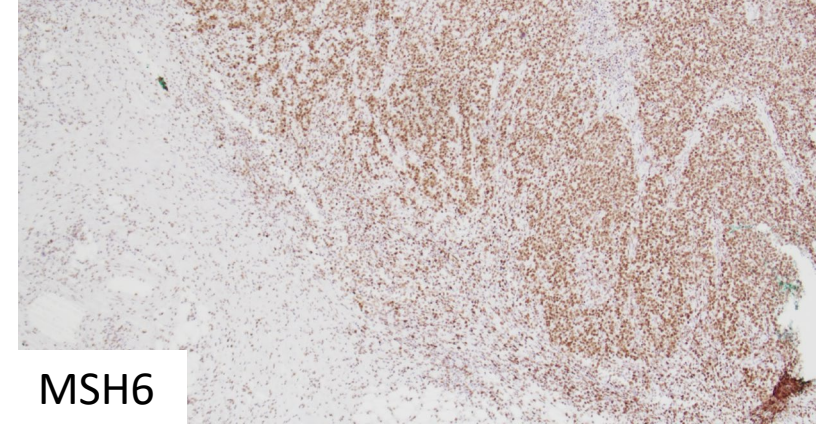
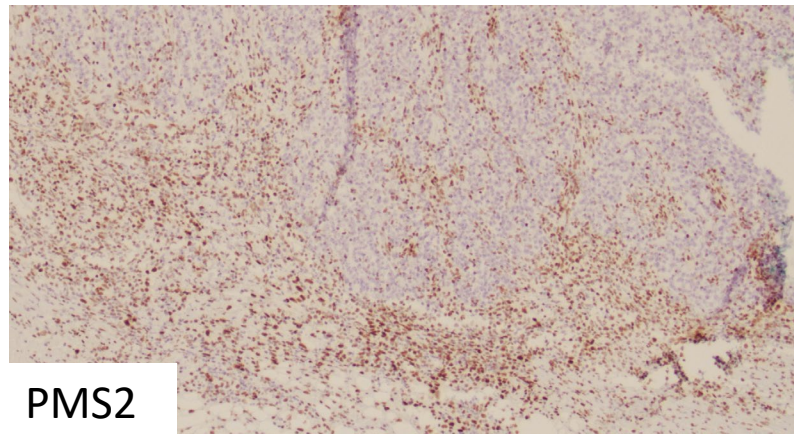
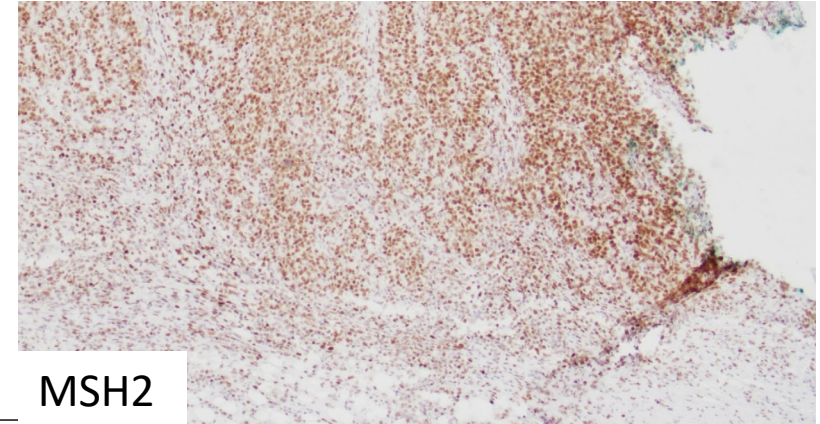
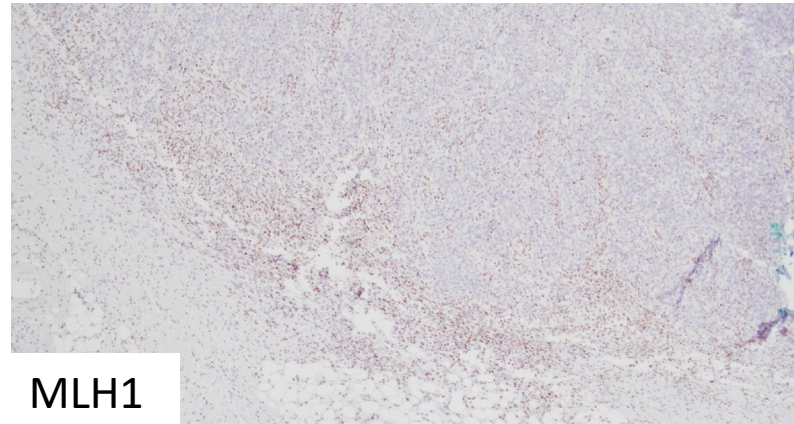
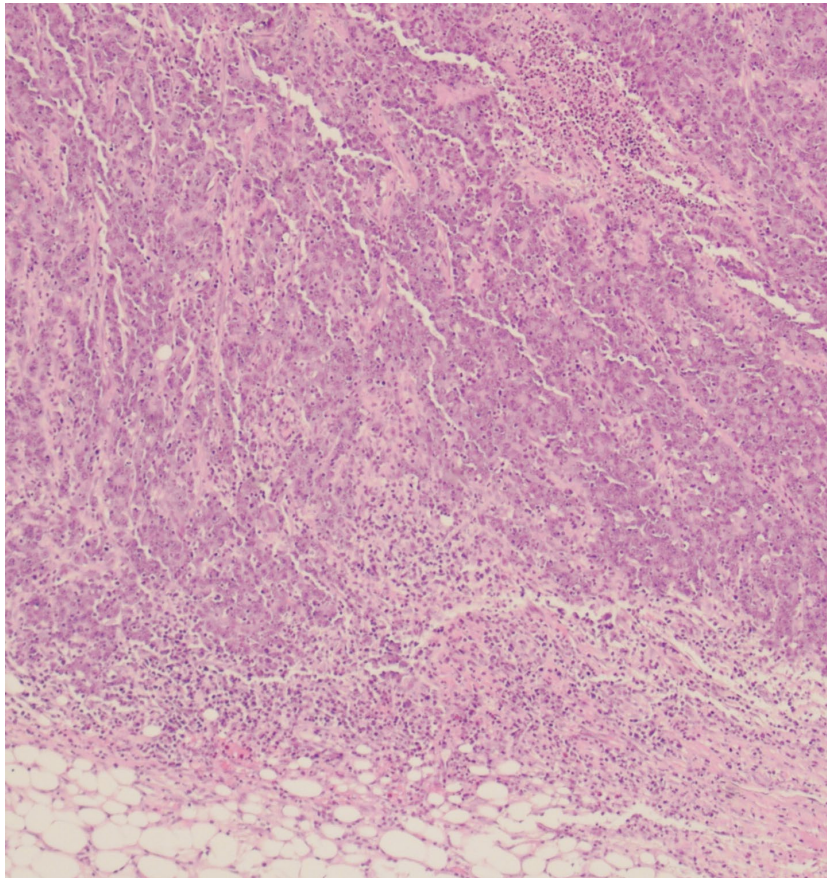


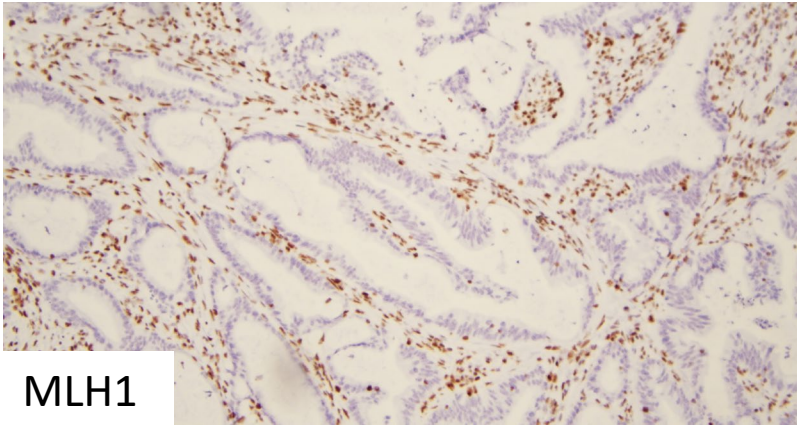
Isolated MSH2 Loss

- These cases have been found to have germline or somatic mutations in *MSH2*
 - Mechanism of MSH6 retention unclear
 - Bottom line: This pattern is dMMR

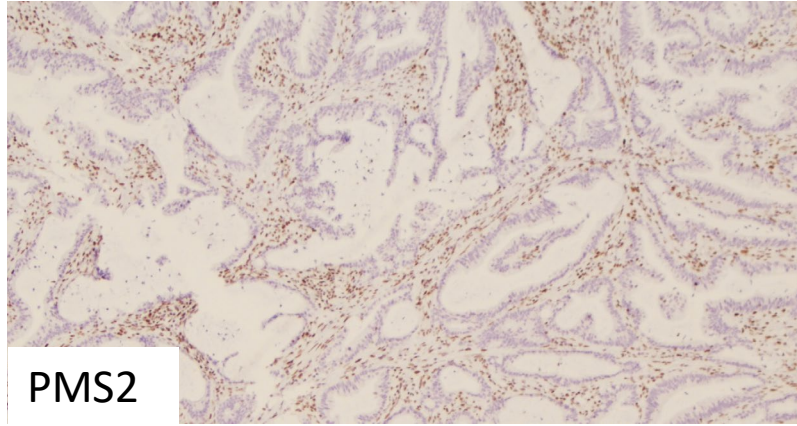
Another Case From My Signout

Tumor #1: Medullary carcinoma

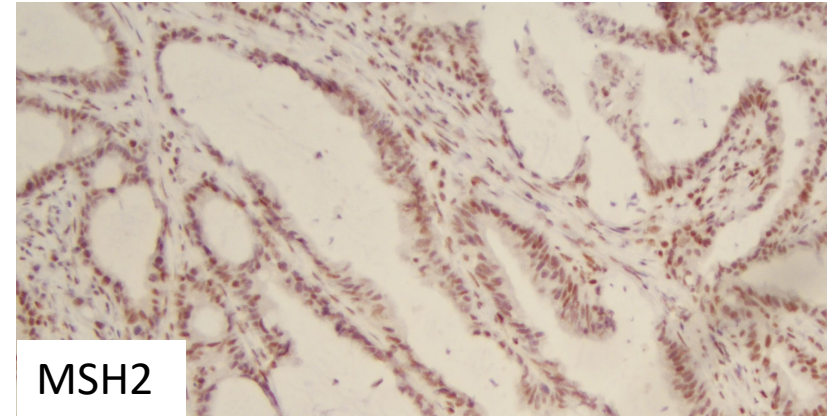




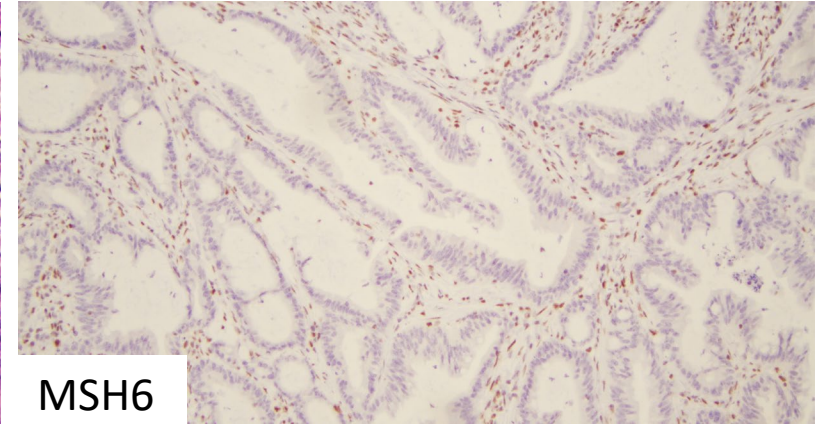
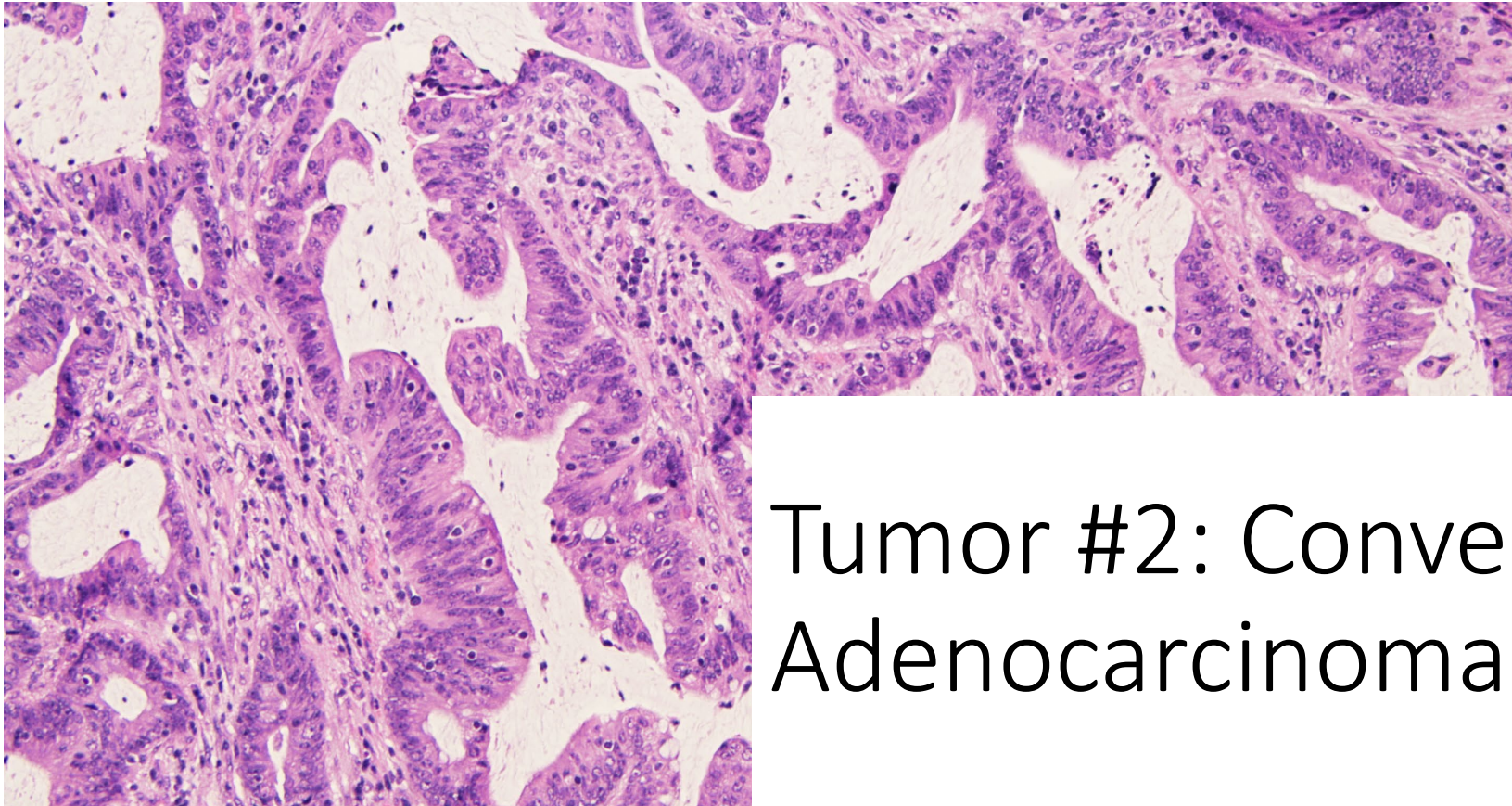
MLH1



PMS2



MSH2



MSH6

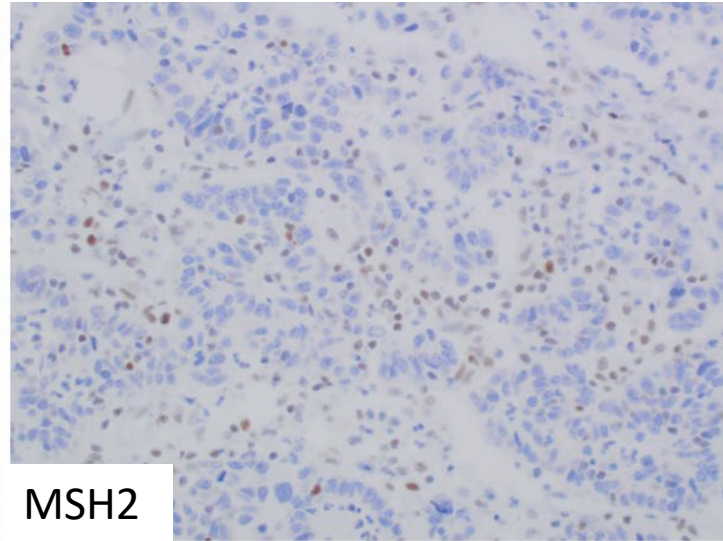
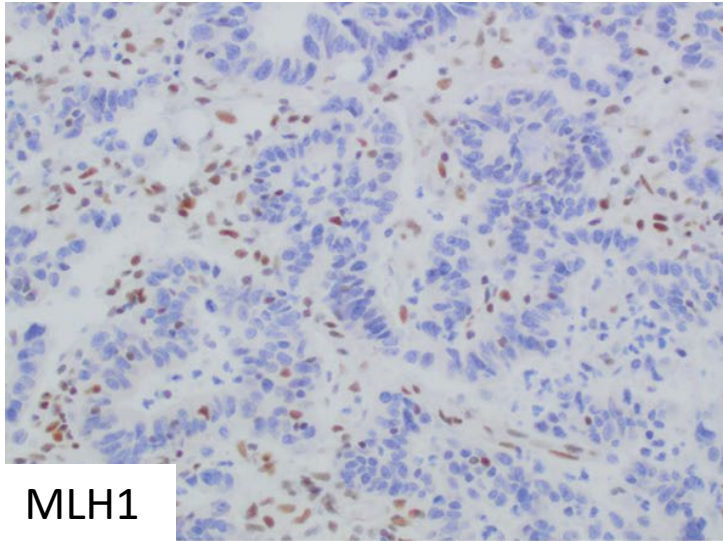
Tumor #2: Conventional
Adenocarcinoma

Loss of MLH1 and/or PMS2, and minimal or absent MSH6

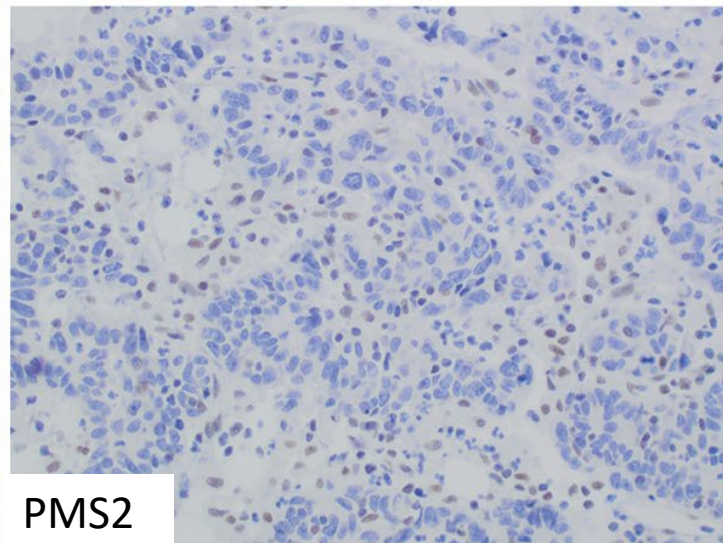
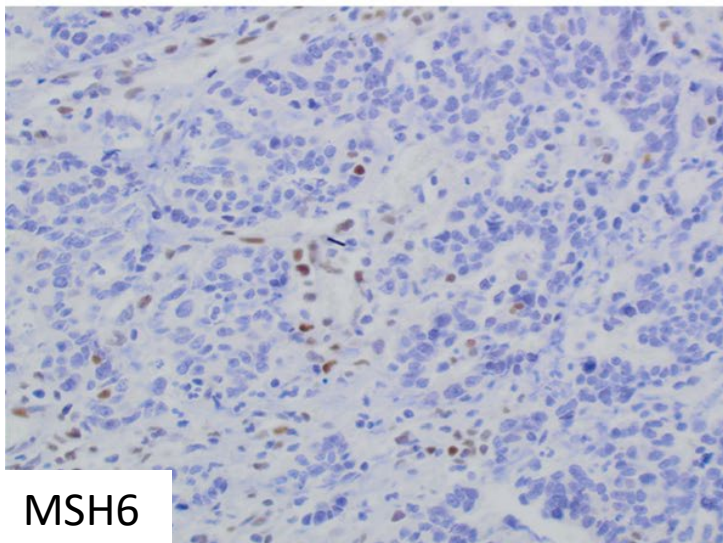
- Hypermethylation of *MLH1* causes MLH1/PMS2 loss
- Tumor subsequently develops somatic mutation in *MSH6*

Shia J, et al. Mod Pathol. 2013 Jan;26(1):131-8.

Null Pattern



- Germline mutation in *MSH2*
- Hypermethylation of *MLH1*



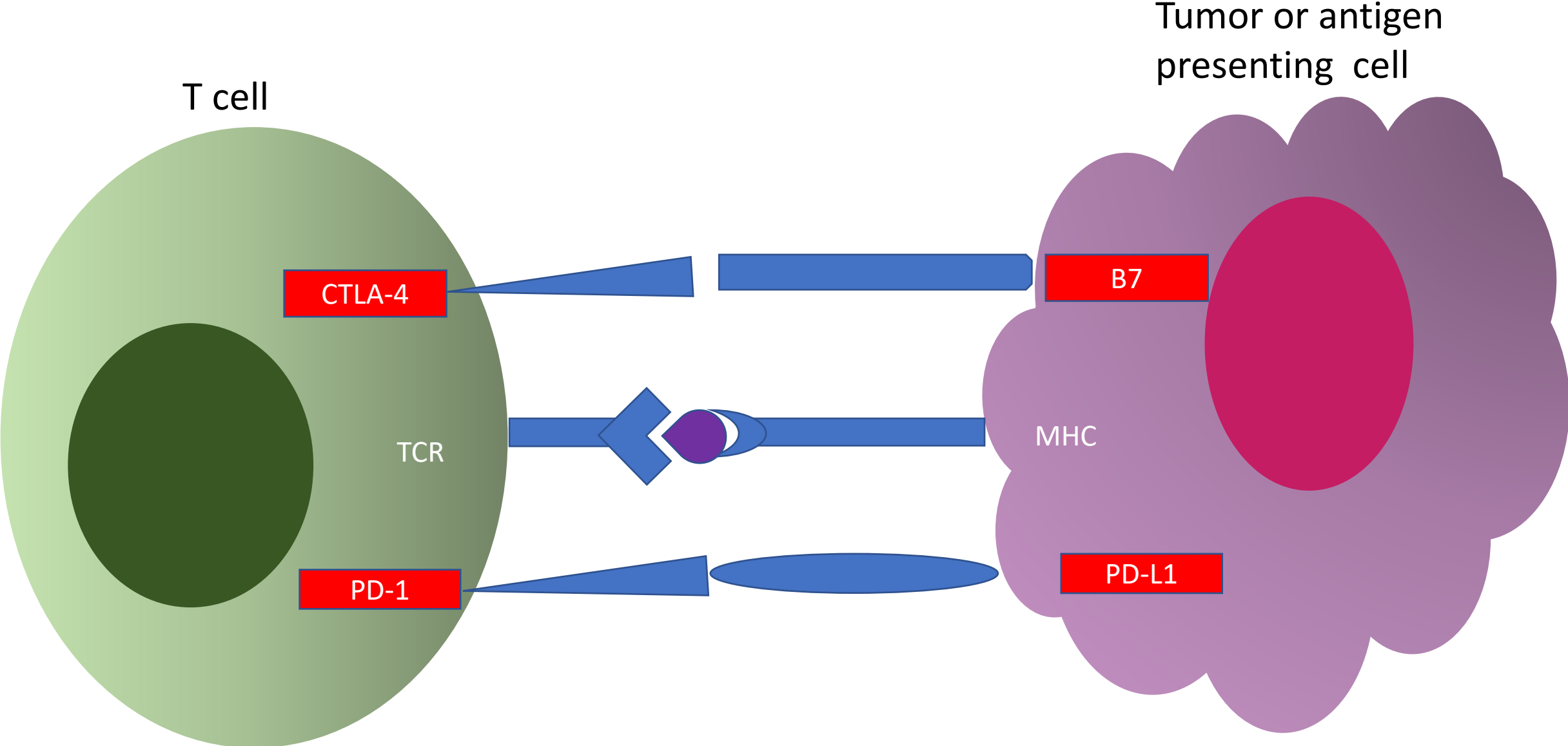
Take Home Points

- Universal Screening for MSI
 - Identifies families with Lynch syndrome
 - Is prognostic and predictive of response to some therapies for colorectal cancer patients
- Immunohistochemistry is sufficient in the vast majority of cases
 - It's all about the internal control
- Have a low threshold for repeating stains when non-standard patterns occur
- PCR or NGS for confirmation or equivocal results

Immunohistochemical Biomarkers

- Mismatch repair (MMR) proficiency and deficiency
- **Checkpoint inhibitor therapy (PDL1)**
- Her2neu amplification

Immune Checkpoint Function



Immune Checkpoint Inhibition

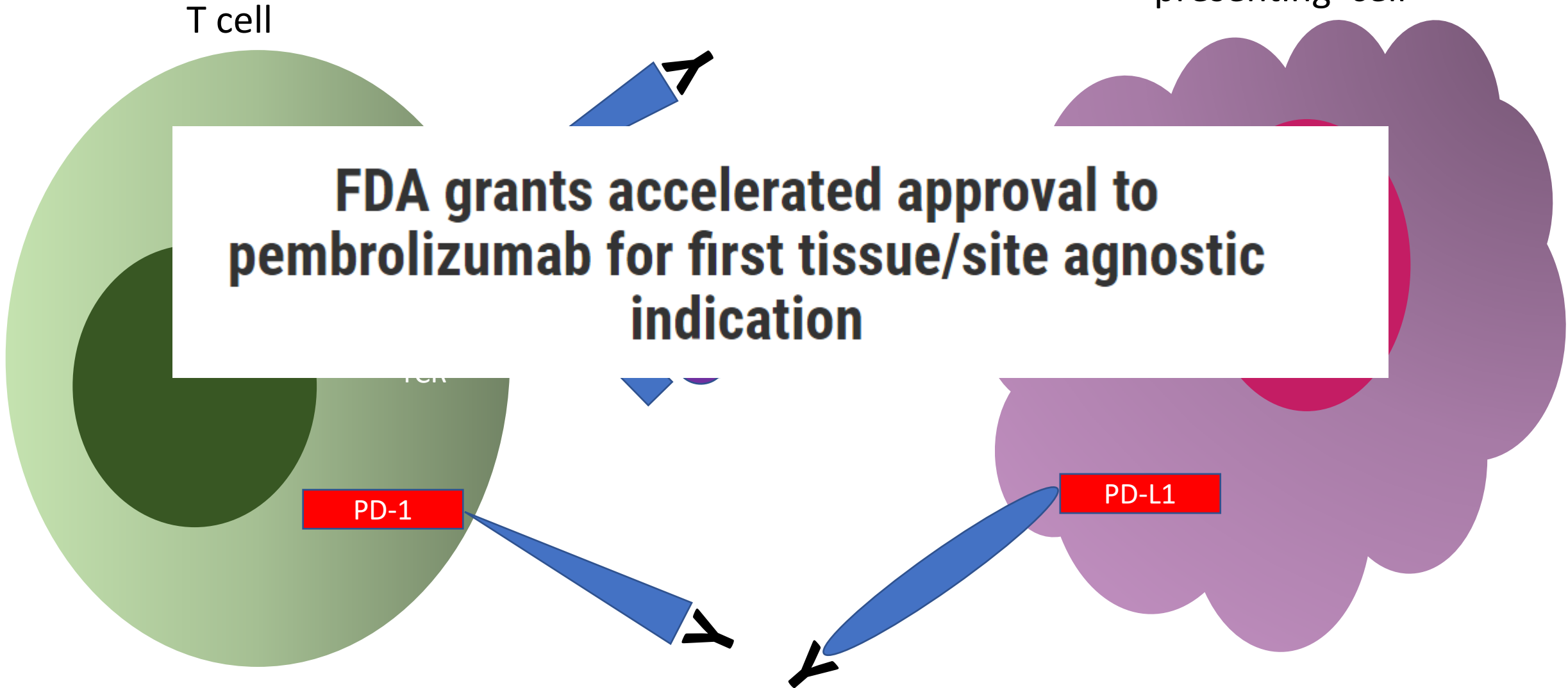
T cell

Tumor or antigen
presenting cell

**FDA grants accelerated approval to
pembrolizumab for first tissue/site agnostic
indication**

PD-1

PD-L1



PD-L1 Immunohistochemistry: Gastric and Gastroesophageal Adenocarcinoma

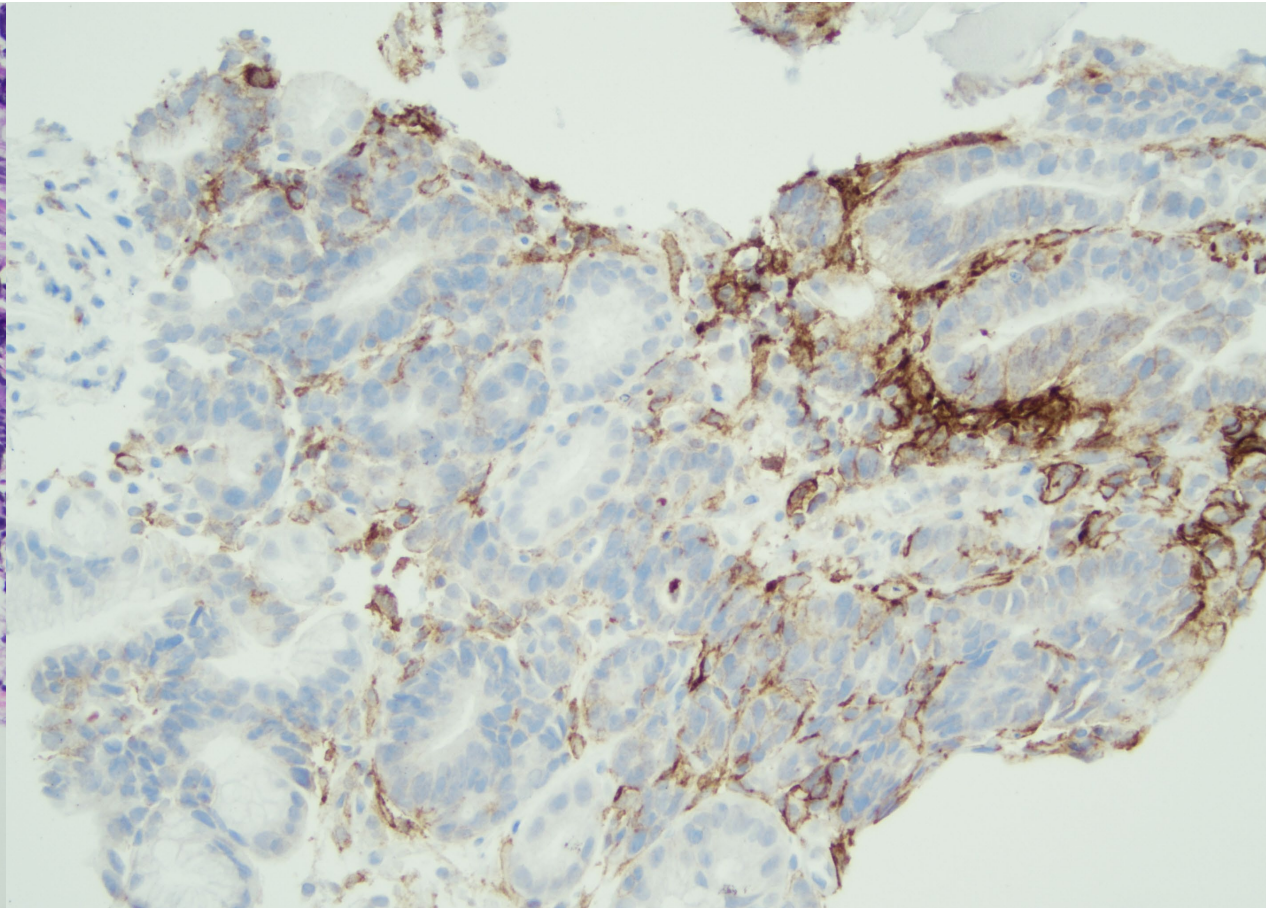
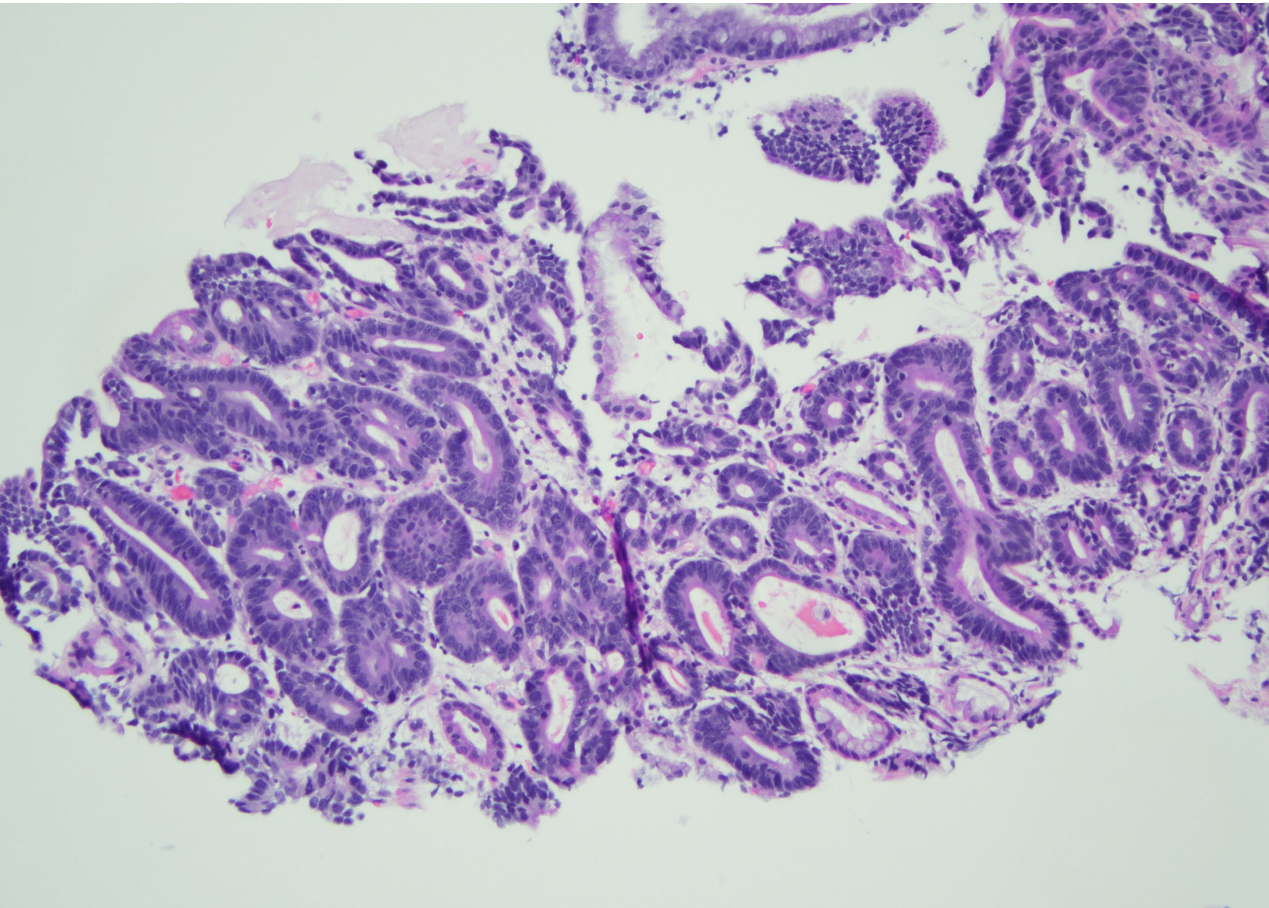
$$\text{Combined positive score (CPS)} = \frac{\text{PD-L1 positive tumor AND immune cells}}{\text{Total viable tumor cells}}$$

- Used for upper gastrointestinal cancers
- Cutoff >1 determines response to checkpoint inhibitor therapy
- Cutoff >10 recently proposed as more predictive of response to checkpoint inhibitor monotherapy

Kulangara K, et al. Arch Pathol Lab Med. 2019 Mar;143(3):330-337.

Wainberg ZA, et al. Clin Cancer Res. 2021 Apr 1;27(7):1923-1931.

CPS: PD-L1 Stain



Checkpoint Inhibitor Therapy in Colon Cancer

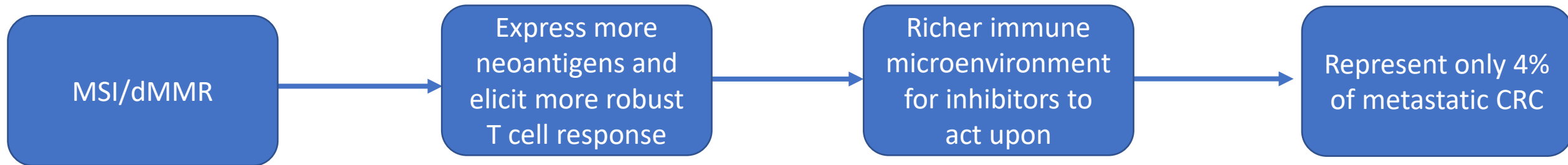
Phase II Study	Agents	Response Rate		
		dMMR CRC	dMMR non-CRC	pMMR CRC
KEYNOTE 016	pembrolizumab	40%	78%	0%!!!!
CheckMate 142	nivolumab	31%	N/A	N/A
	Nivolumab + ipilimumab	55%	N/A	N/A

- Phase III trials ongoing

Le DT, et al. N Engl J Med. 2015;372:2509-2520.

Overman MJ, et al. Lancet Oncol. 2017 Sep;18(9):1182-1191.

Checkpoint Inhibitor Therapy in Colon Cancer



What about mismatch repair proficient (pMMR) CRC?

- Some may harbor mutations that stimulate T cell response (*TP53*, *KRAS*)
- Combination of traditional chemo and radiotherapy may expose tumor antigens that potentiates the effect of checkpoint inhibition
- Trials are ongoing

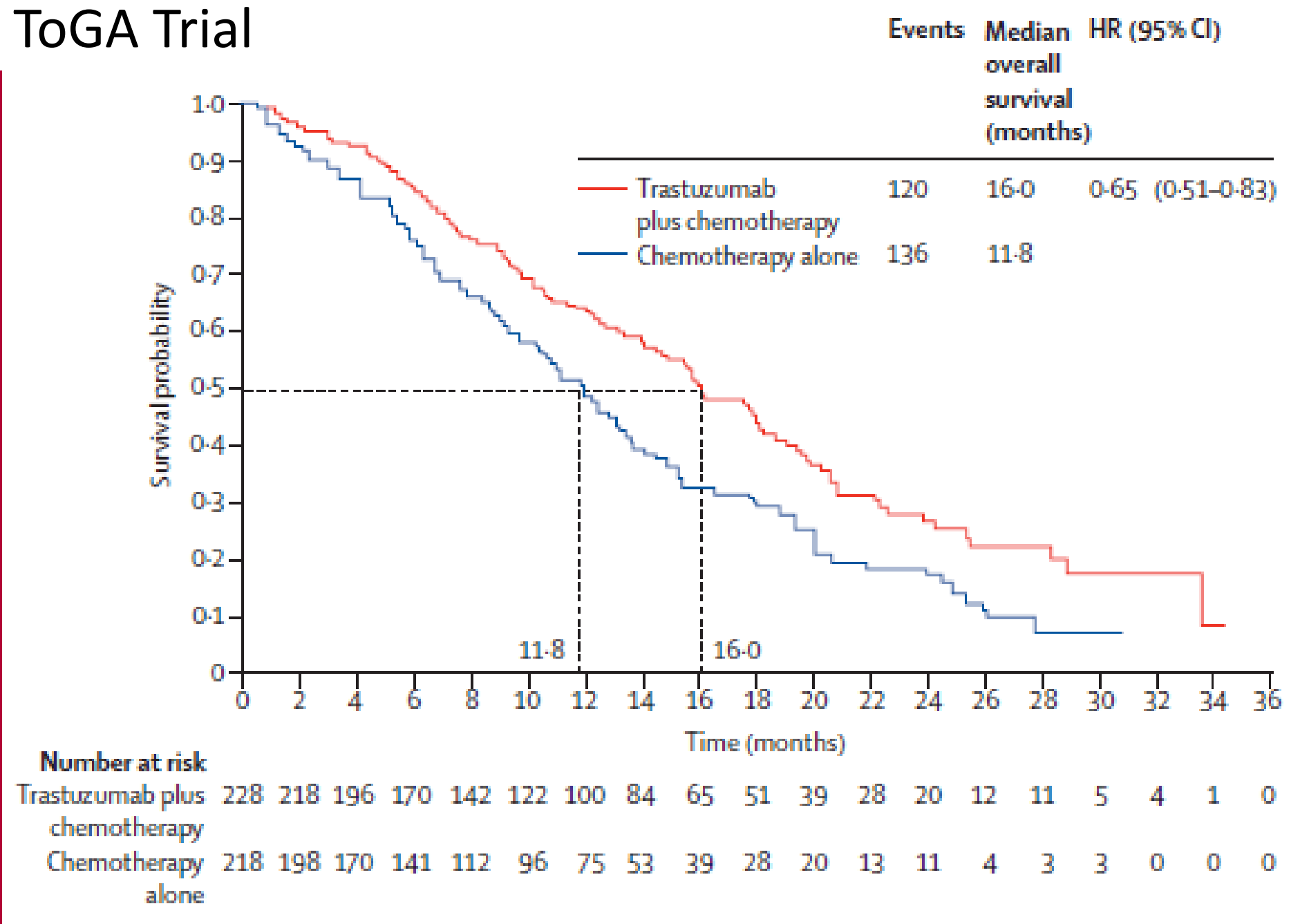
Take-Home Points

- Currently, the best marker of checkpoint inhibitor therapy response for colorectal cancer is dMMR

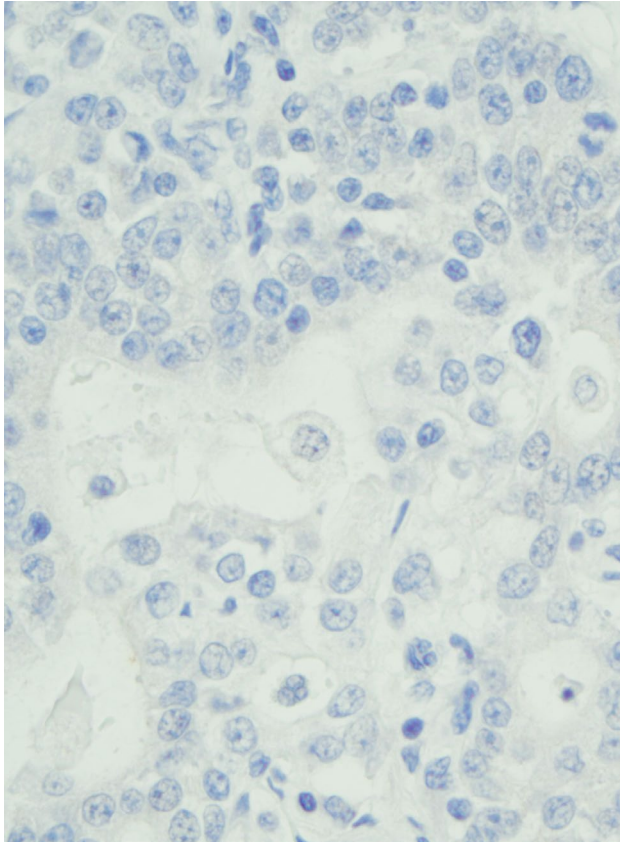
Immunohistochemical Biomarkers

- Mismatch repair (MMR) proficiency and deficiency
- PDL1
- **Her2neu amplification**

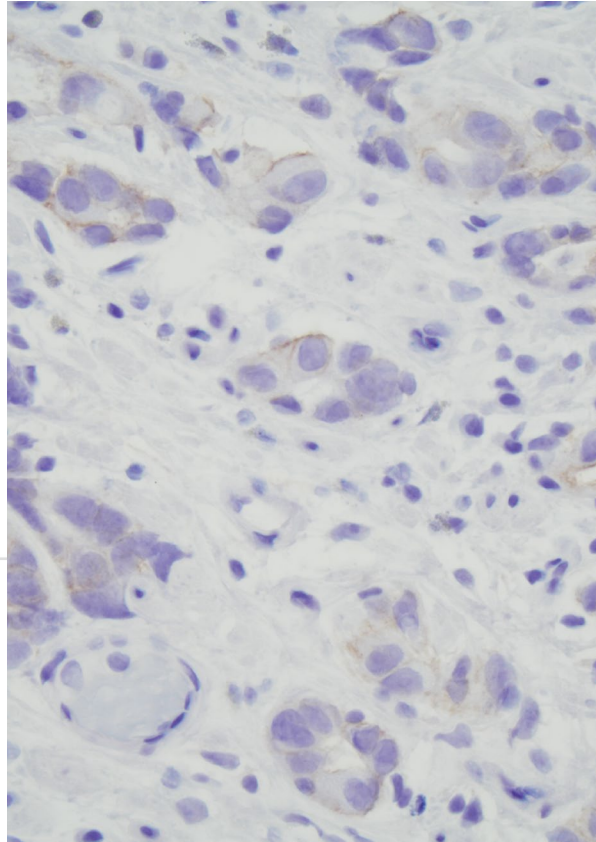
ToGA Trial



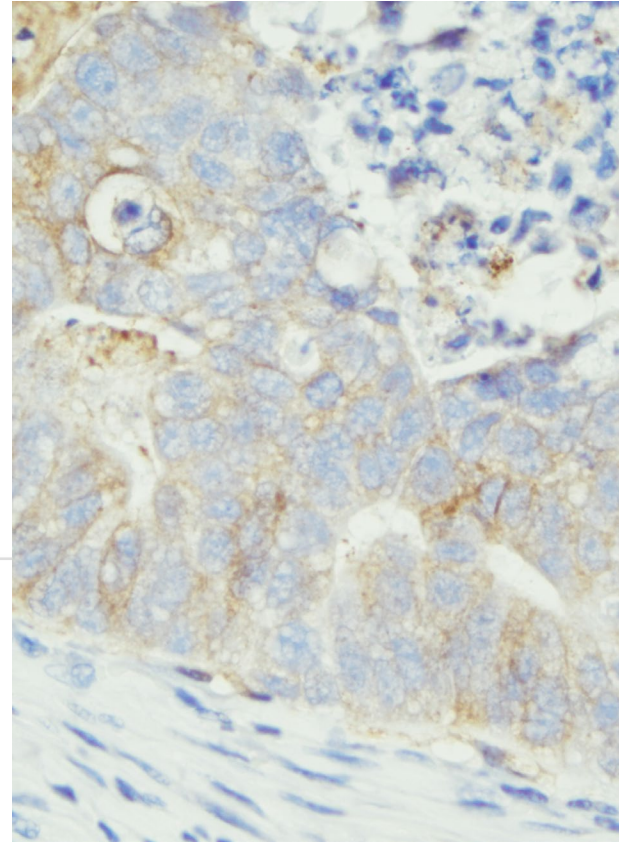
Gastric and Gastroesophageal Junction Cancer HER2 Scoring



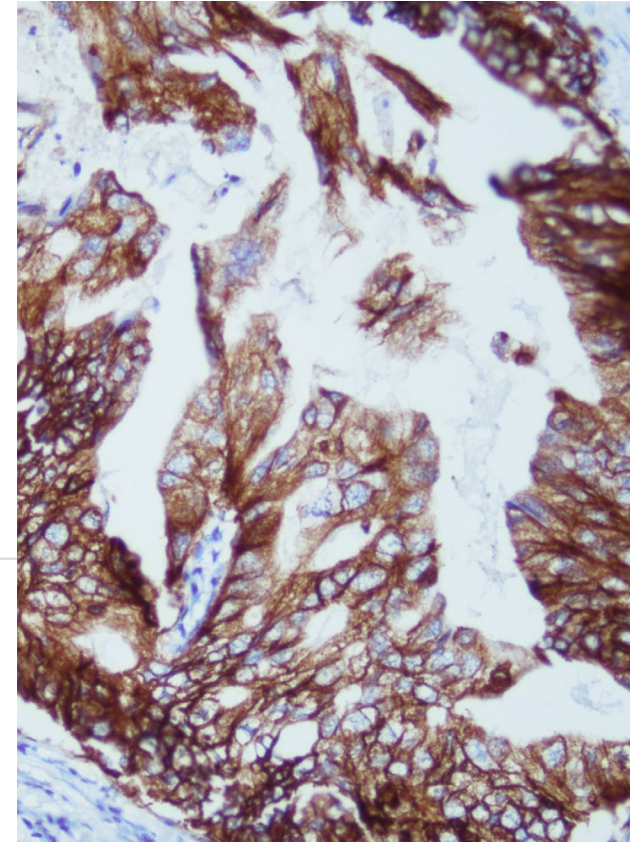
0: No staining



1+: Barely perceptible

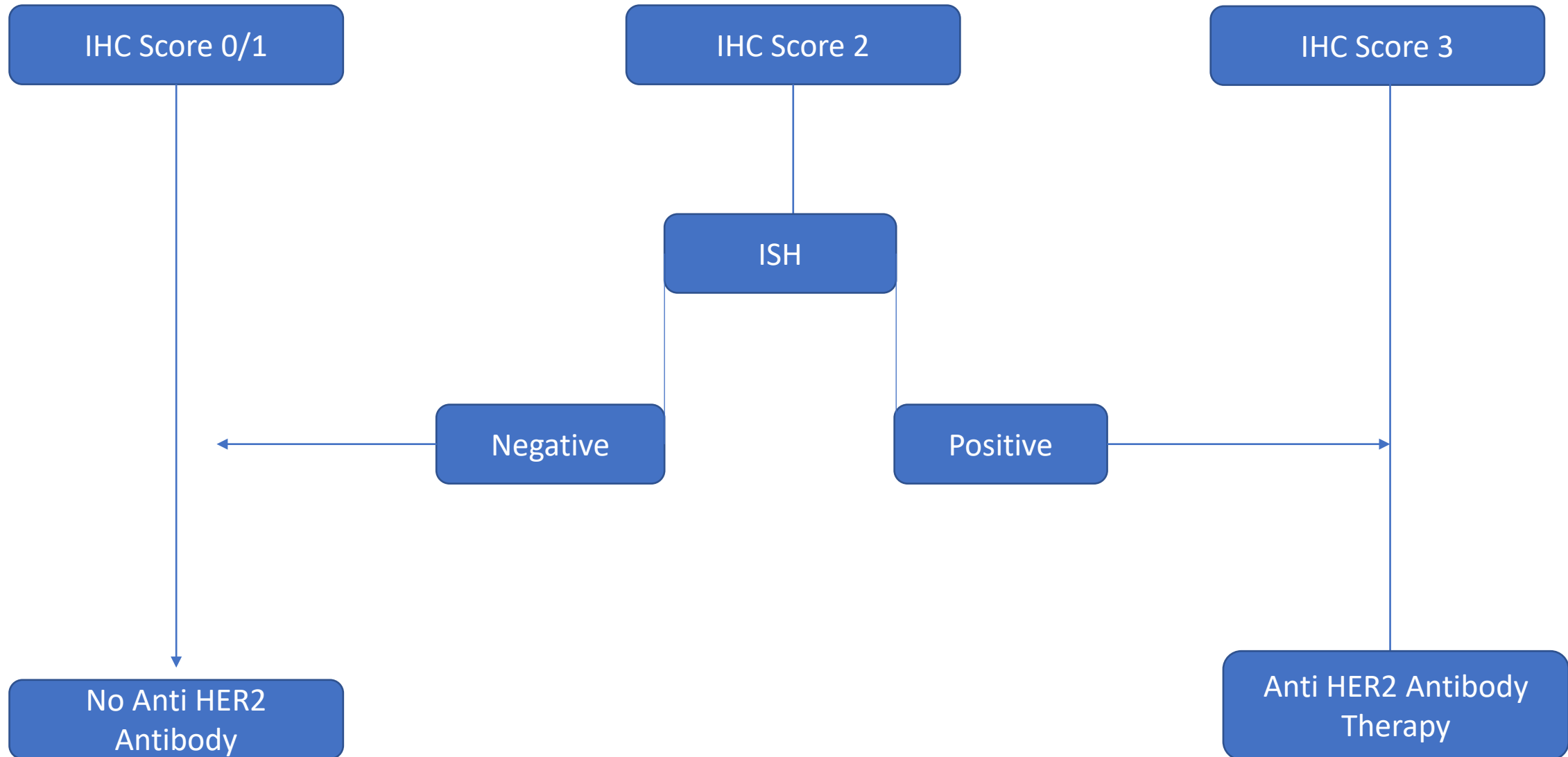


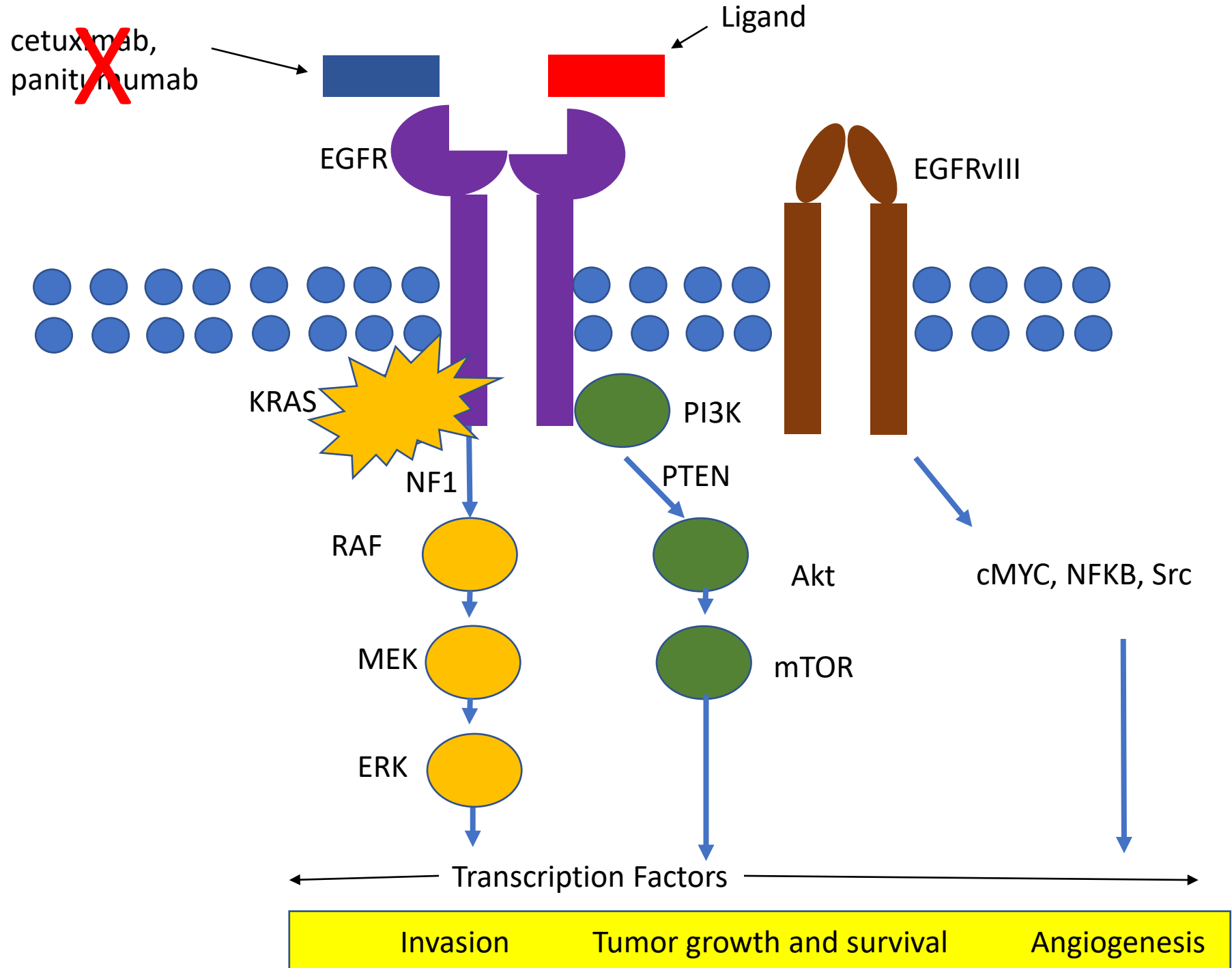
2+: Weak to moderate

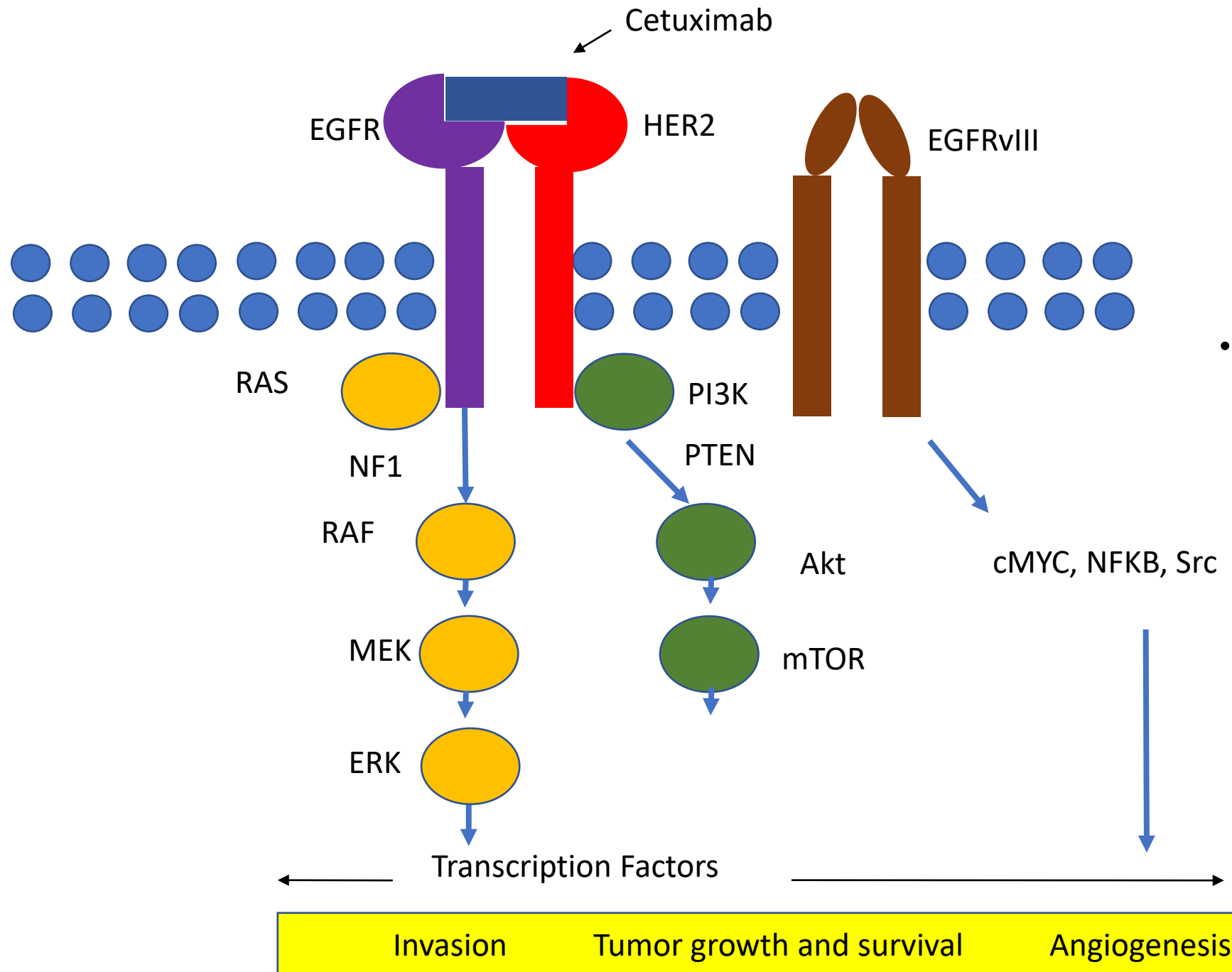


3+: Strong

Gastric and Gastroesophageal Junction Her2 Expression Testing Algorithm







- **HER2 is amplified in approximately 1-8% of colorectal**

HERACLES (HER2 Amplification for Colo-rectal cancer Enhanced Stratification)

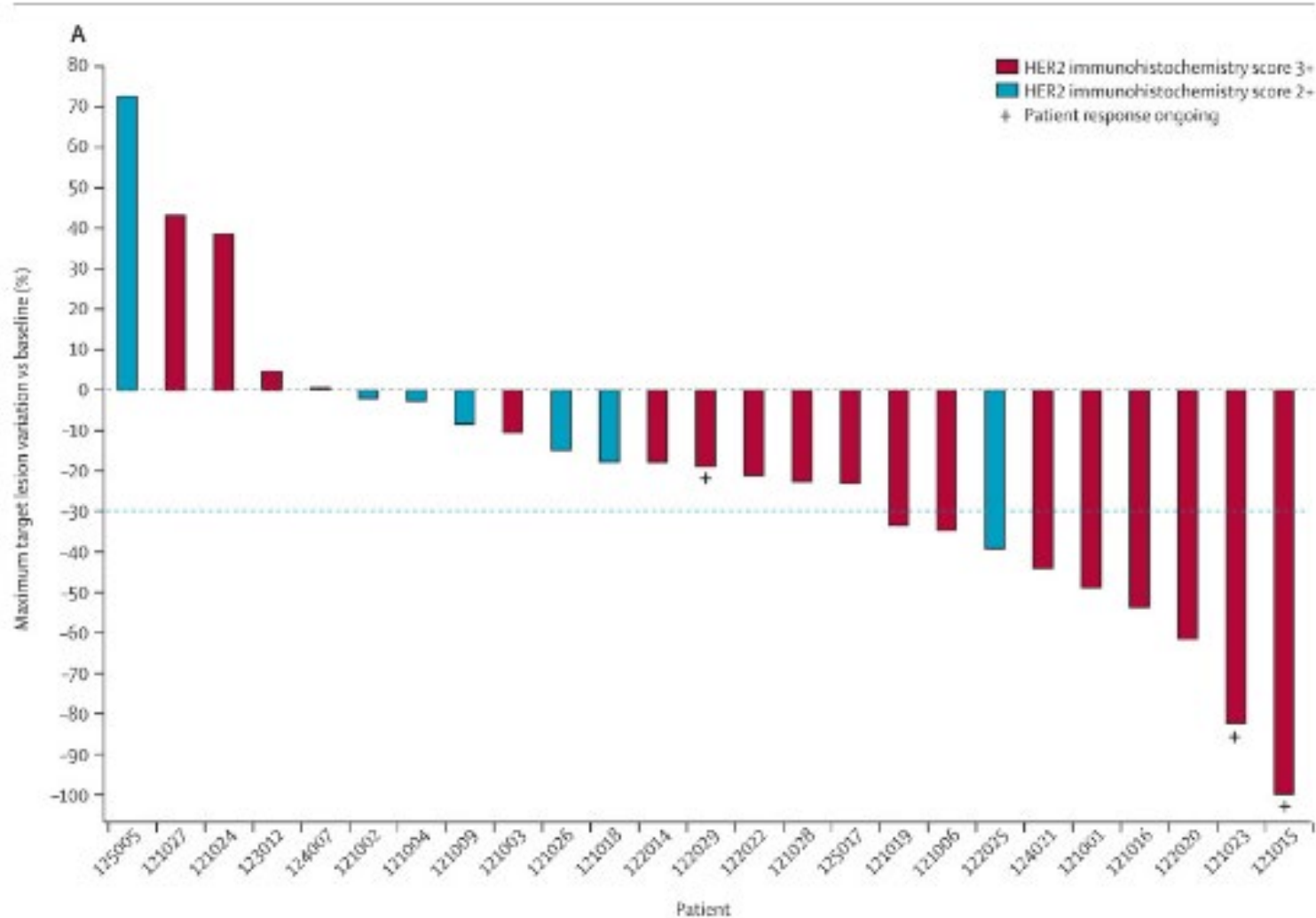
- Patients with metastatic colorectal cancer
- Had progressed on standard therapies (inclusive of cetuximab and panitumumab)
- Trastuzumab – anti-Her2 antibody
- Lapatinib – dual tyrosine kinase inhibitor that blocks HER2/neu and EGFR

HERACLES (HER2 Amplification for Colo-rectal cancer Enhanced Stratification)

Score	Staining Pattern
0	No staining
1+	Faint segmental or granular staining
2+	Moderate circumferential, basolateral, or lateral staining
3+	Intense circumferential, basolateral, or lateral staining

- HER2 positivity
 - 3+ immunostaining in >50% of cells
 - 3+ in ≥ 10 , but <50% of cells and positive by FISH
 - 2+ in >50% of cells and positive by FISH

HERACLES Results



- 27 patients
 - 1 complete response
 - 7 partial response
 - 12 stable disease
 - Duration of response at least 16 weeks

MyPathway

- HER2 positivity was defined as any of the following:
- HER2 IHC 3+ in >10 percent of cells
- *HER2*:CEP17 ratio ≥ 2.0 or *HER2* count >6 per cell
- Increased *HER2* gene copy number by molecular methods
- *HER2* activating mutations
- 57 HER2-positive advanced colorectal cancer patients
- Eighteen patients achieved an objective response, including one complete response and 17 partial responses; seven additional patients had stable disease for greater than four months

Take-Home Points

- Mismatch repair proteins
 - Control is the KEY
- Checkpoint inhibitor (PDL1)
 - Mismatch repair currently the best marker
- Her2neu amplification
 - Promising target
 - No uniformly agreed upon IHC criteria
 - Criteria may differ from upper gastrointestinal tract
 - May employ multiple modalities in conjunction with IHC



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