



The 62nd Annual Congress of Taiwan Association of Obstetrics and Gynecology

# Assessment of Mismatch Repair Deficiency and Associated Clinicopathologic Significance

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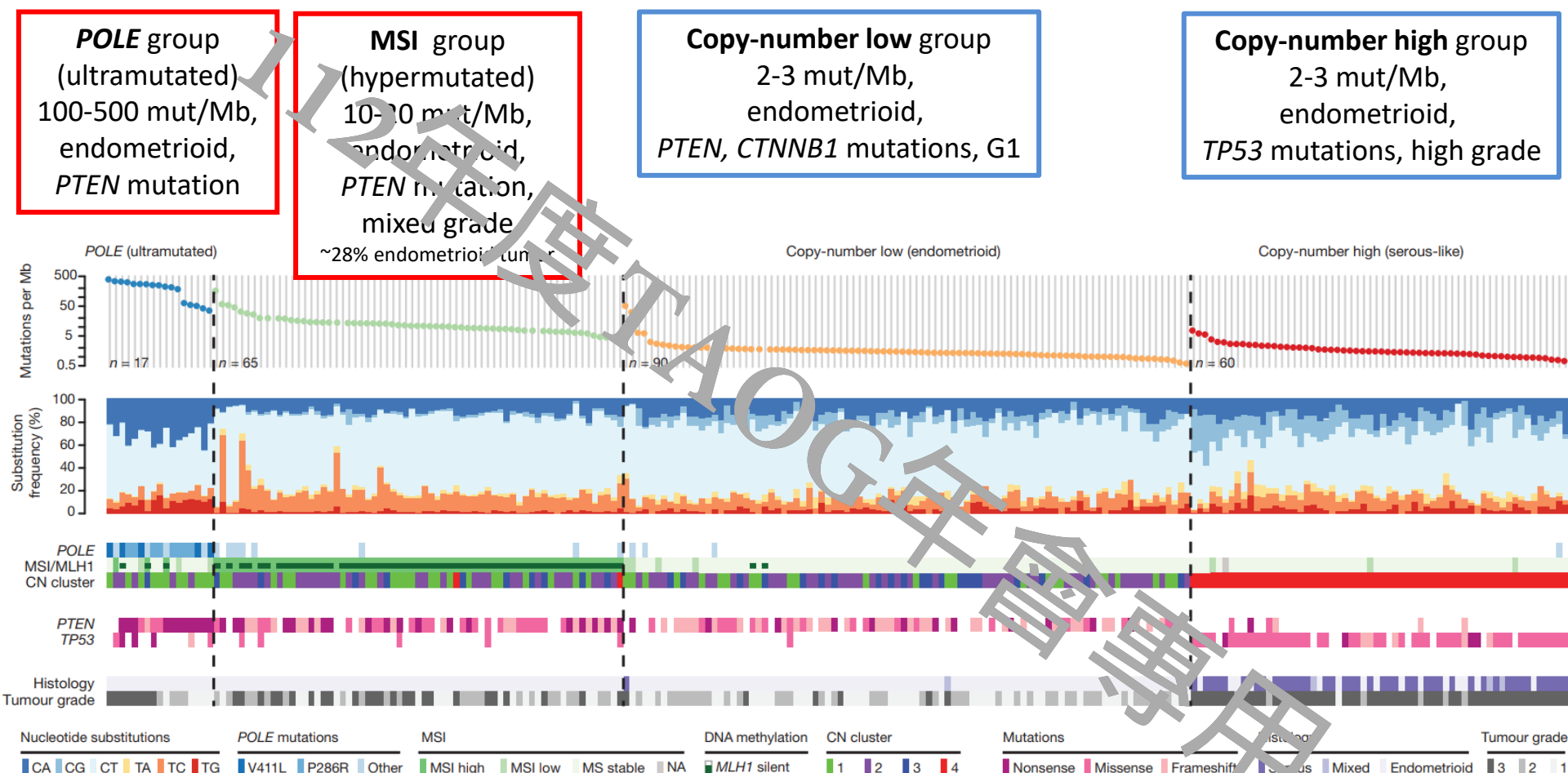
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# Overview

- May. 2017 Pembrolizumab (PD-1 inhibitor) received accelerated FDA approval for Solid MSI-H or dMMR tumor that has progressed following prior treatment
- Jun 2018 Cervical cancer with PD-L1 /2nd line
- Apr 2021 Dostarlimab dMMR endometrial cancer/2nd line (GARNET)
- Jul 2021 Pembrolizumab + lenvatinib endometrial cancer /2nd line (KEYNOTE-775)
- Aug 2021 Dostarlimab dMMR solid tumors /2nd line (GARNET)
- Mar 2022 Pembrolizumab dMMR endometrial cancer/2nd line (KEYNOTE-158)
- MSI-H or dMMR tumors tend to be hypermutated, release more neoantigens that stimulate an immune response, and may be more susceptible to immunotherapeutic strategies

# Mutation Spectra Across Endometrial Carcinomas

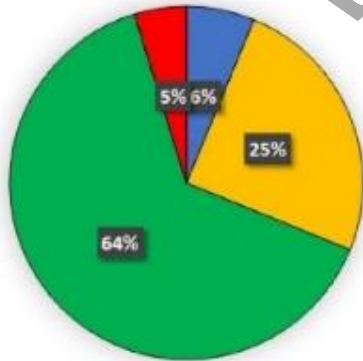


**High genome instability  
immunogenic**

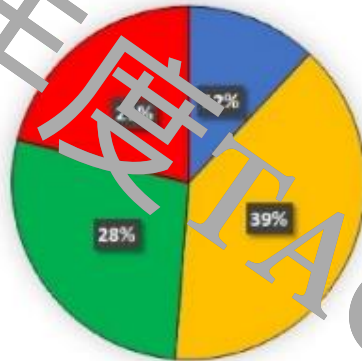
**Hypo-mutated, low expression of immune-related  
biomarkers**

# Molecular Classifications of Endometrial Carcinomas

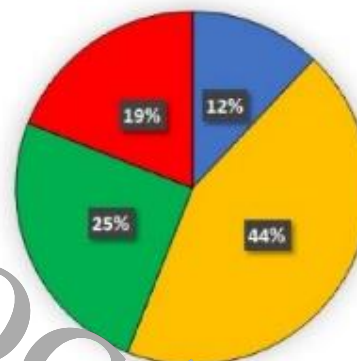
Low-grade endometrioid



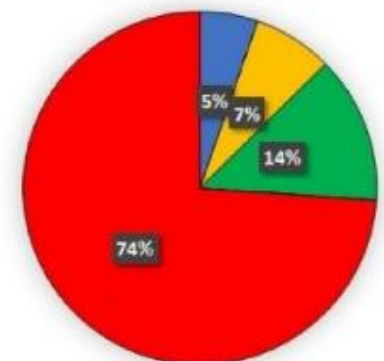
High-grade endometrioid



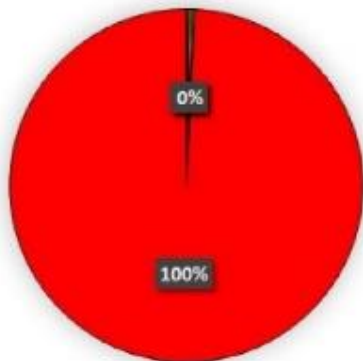
Un-/Dedifferentiated



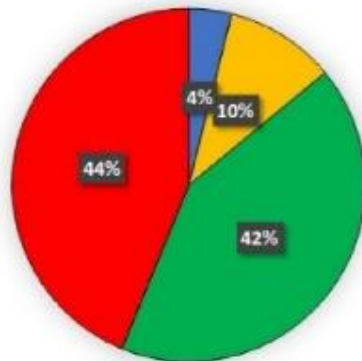
Carcinosarcoma



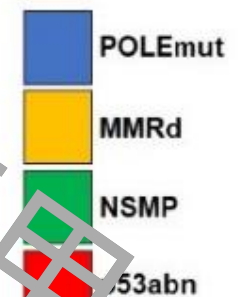
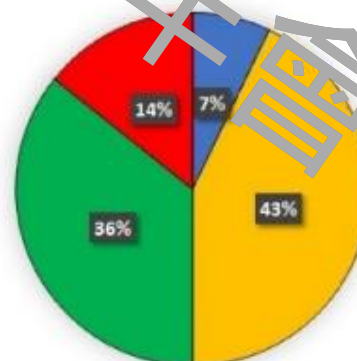
Serous



Clear cell

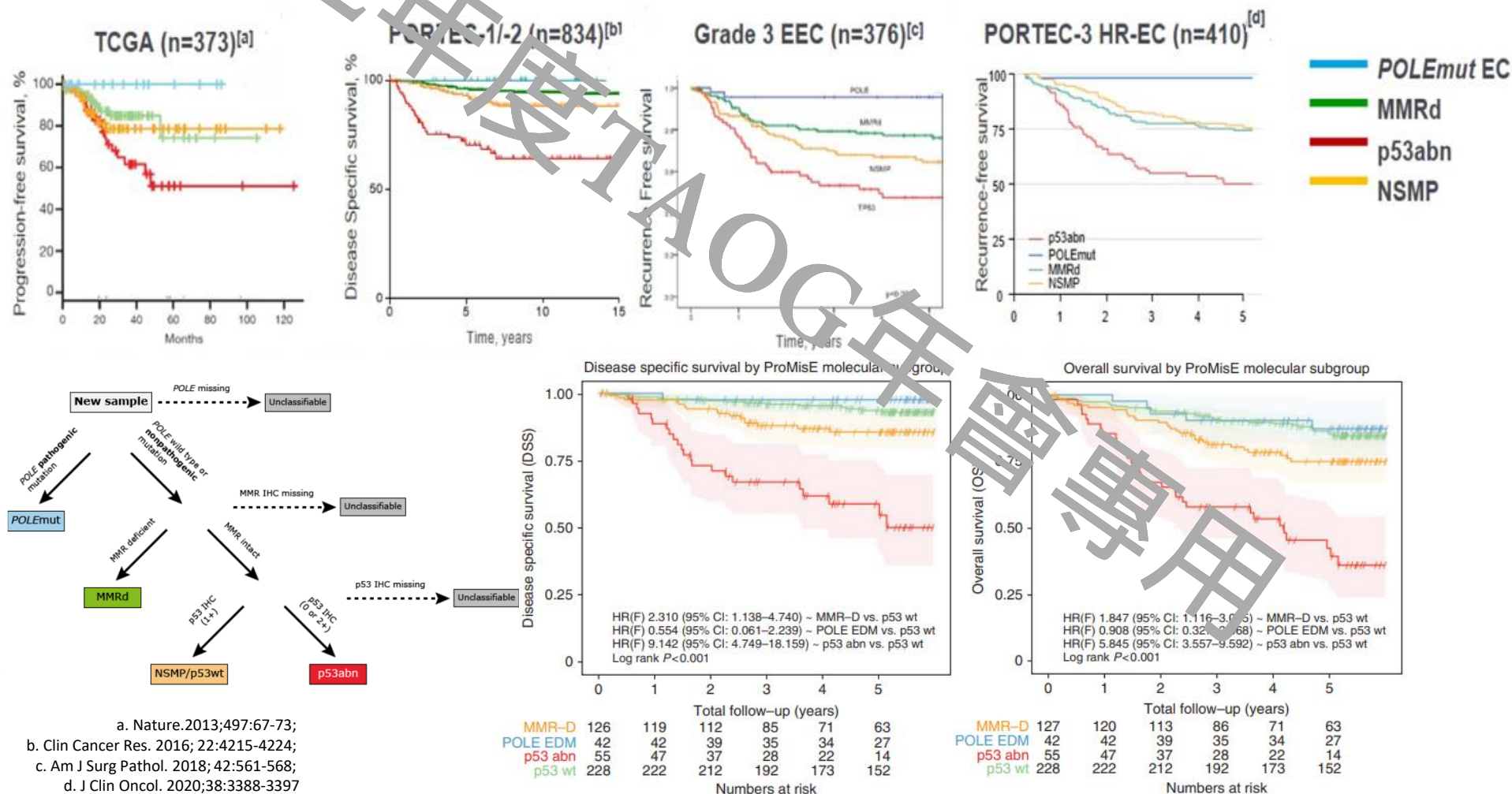


Neuroendocrine



# High prognostic value of molecular characterization of endometrial cancer

- Best survival for *POLE*mut (ultramutated) EC
- ProMisE (Proactive Molecular Risk Classifier for Endometrial Cancer)





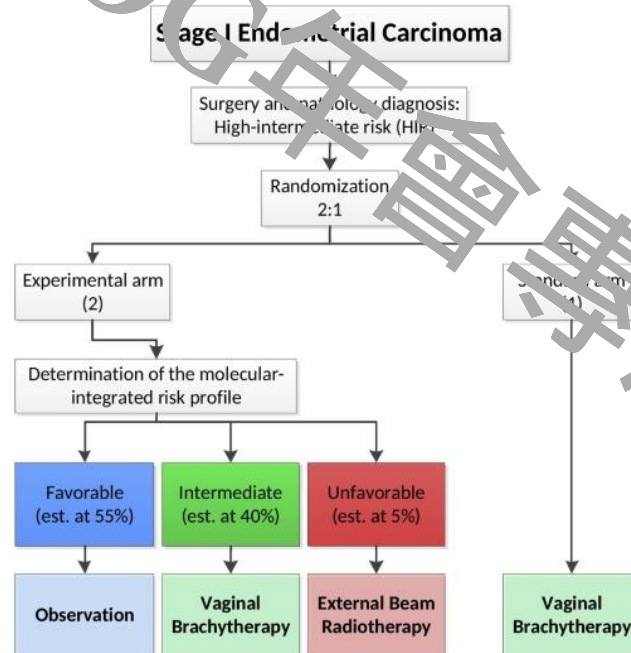
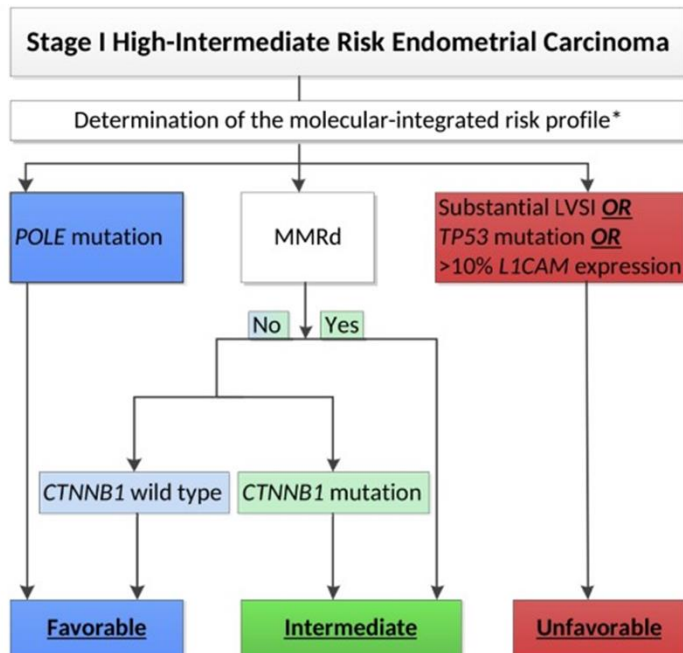
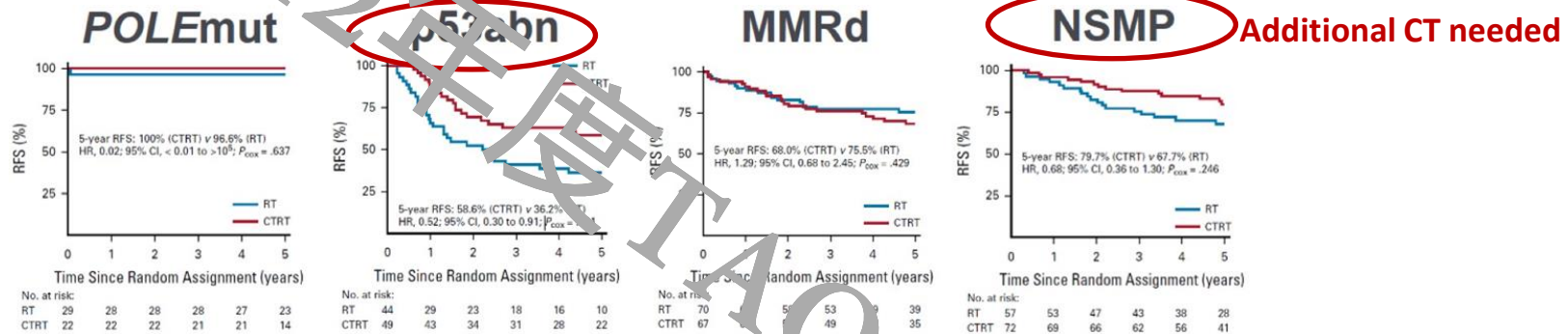
# 2020 ESGO/ESTRO/ESP Guidelines

## *Molecular Classifications of Endometrial Carcinomas*

Risk Group	Molecular Classification Unknown	Molecular Classification Known
Low	<ul style="list-style-type: none"> <li>➤ Stage IA endometrioid + low-grade + LVSI negative or focal</li> </ul>	<ul style="list-style-type: none"> <li>➤ Stage I-II <b>POLEmut</b> endometrial carcinoma, no residual disease</li> <li>➤ Stage IA <b>dMMR/NSMP</b> endometrioid carcinoma + low grade + LVSI negative or focal</li> </ul>
Intermediate	<ul style="list-style-type: none"> <li>➤ Stage IB endometrioid + low-grade + LVSI negative or focal</li> <li>➤ Stage IA endometrioid + high-grade + LVSI negative or focal</li> <li>➤ Stage IA nonendometrioid (serous; clear cell; undifferentiated carcinoma; carcinosarcoma, mixed) without myometrial invasion</li> </ul>	<ul style="list-style-type: none"> <li>➤ Stage IB <b>dMMR/NSMP</b> endometrioid carcinoma + low grade + LVSI negative or focal</li> <li>➤ Stage IA <b>MMRd/NSMP</b> endometrioid carcinoma + high grade + LVSI negative or focal</li> <li>➤ Stage IA <b>p53abn</b> or nonendometrioid (serous; clear cell; undifferentiated carcinoma; carcinosarcoma, mixed) without myometrial invasion</li> </ul>
High-intermediate	<ul style="list-style-type: none"> <li>➤ Stage I endometrioid + substantial LVSI regardless of grade and depth of invasion</li> <li>➤ Stage IB endometrioid high grade regardless of LVSI status</li> <li>➤ Stage II</li> </ul>	<ul style="list-style-type: none"> <li>➤ Stage I <b>dMMR/NSMP</b> endometrioid carcinoma + substantial LVSI regardless of grade and depth of invasion</li> <li>➤ Stage IB <b>dMMR/NSMP</b> endometrioid carcinoma high grade regardless of LVSI status</li> <li>➤ Stage II <b>dMMR/NSMP</b> endometrioid carcinoma</li> </ul>
High	<ul style="list-style-type: none"> <li>➤ Stage III-IVA with no residual disease</li> <li>➤ Stage I-IVA nonendometrioid (serous; clear cell; undifferentiated carcinoma; carcinosarcoma, mixed) with myometrial invasion and with no residual disease</li> </ul>	<ul style="list-style-type: none"> <li>➤ Stage III-IVA <b>dMMR/NSMP</b> endometrioid carcinoma with no residual disease</li> <li>➤ Stage I-IVA <b>p53abn</b> endometrial carcinoma with myometrial invasion, with no residual disease</li> <li>➤ Stage I-IVA <b>NSMP/dMMR</b> serous; undifferentiated carcinoma; carcinosarcoma with myometrial invasion, with no residual disease</li> </ul>
Advanced metastatic	<ul style="list-style-type: none"> <li>➤ Stage III-IVA with residual disease</li> <li>➤ Stage IVB</li> </ul>	<ul style="list-style-type: none"> <li>➤ Stage III-IVA with residual disease of any molecular type</li> <li>➤ Stage IVB of any molecular type</li> </ul>

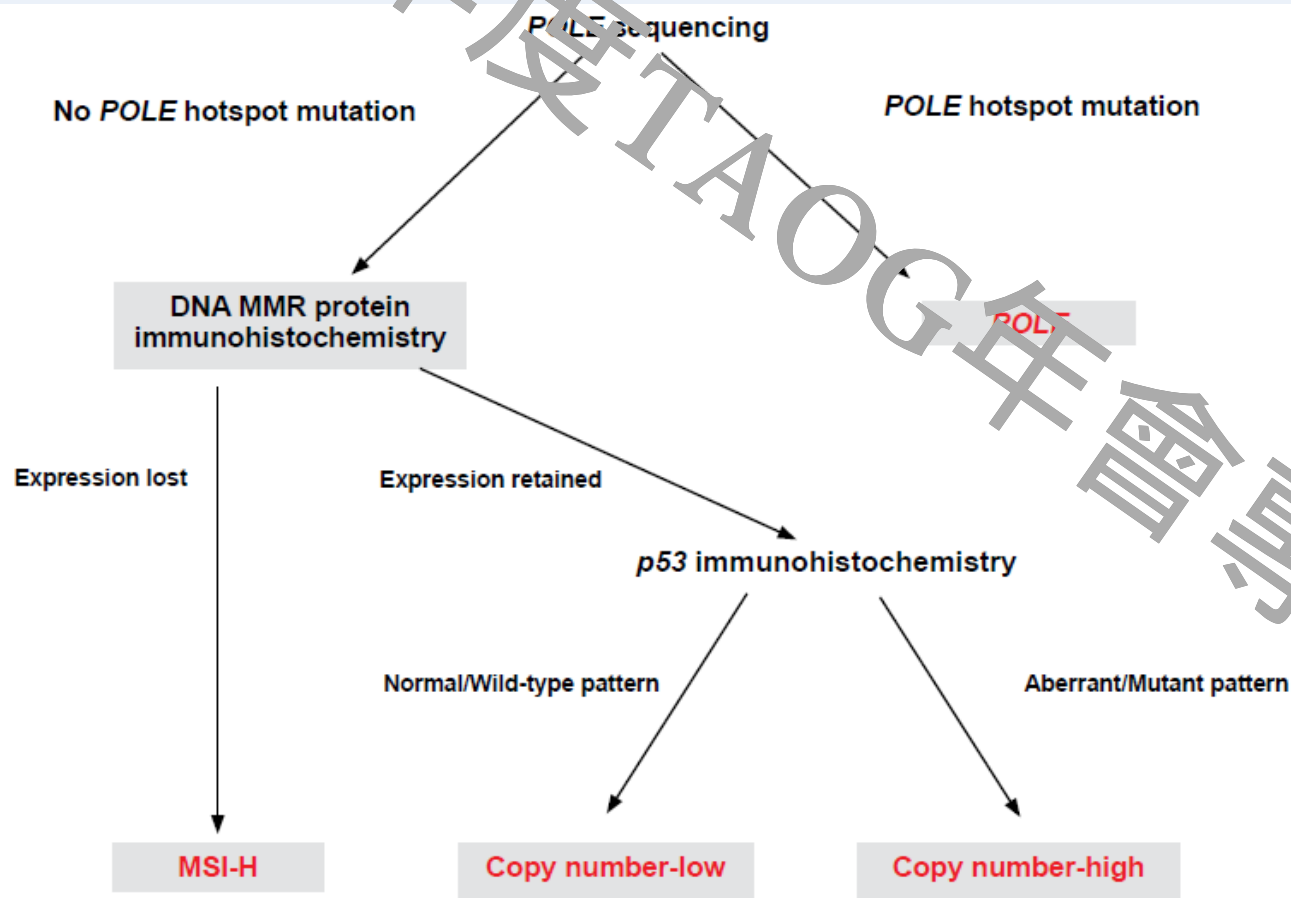
# Molecular typing-guided treatment

- Predictive potential for adjuvant platinum-based treatment in PORTEC-3
- Ongoing PORTEC-4a<sub>NCT03469674</sub>, CANSTAMP<sub>NCT04159155</sub>, TAPER<sub>NCT04705649</sub>.....



# Principle of molecular analysis

- Recommend **universal testing for MMR proteins** (MSI testing if results equivocal)
- Assess promoter methylation in MLH1 loss (epigenetic mechanism)
- Consider NTRK gene fusion testing for metastatic or recurrent EC
- Consider TMB testing through validated and/or FDA-approved assays





# Updated 2023 FIGO Staging of Endometrial Cancer

## Define substages

- In all stages, grade, histological type and LVSI must be recorded. If available and feasible, molecular classification testing (POLEmut, MMRd, NSMP, p53abn) is encouraged in all patients for prognostic risk-group stratification
- addition of “m” for molecular classification and a subscript for subtype
- Non-aggressive histological types: G1/2 EECs; aggressive histological types: G3 EEC, serous, clear cell, undifferentiated, mixed, mesonephric-like, gastrointestinal mucinous type carcinomas and carcinosarcomas.
- **Disease upstaging or downstaging if “m” classification of p53abn or POLEmut in stages I and II (IICm<sub>p53abn</sub> or IAM<sub>POLEmut</sub>).**

TABLE 2 FIGO endometrial cancer stage with molecular classification.<sup>a</sup>

Stage designation	Molecular findings in patients with early endometrial cancer (stages I and II after surgical staging)
Stage IAM <sub>POLEmut</sub>	POLEmut endometrial carcinoma, confined to the uterine corpus or with cervical extension, regardless of the degree of LVSI or histological type <i>Stage II (FIGO 2009)</i>
Stage IICm <sub>p53abn</sub>	p53abn endometrial carcinoma confined to the uterine corpus with any myometrial invasion, with or without cervical invasion, and regardless of the degree of LVSI or histological type <i>Stage I (FIGO 2009)</i>

# Updated 2023 FIGO Staging

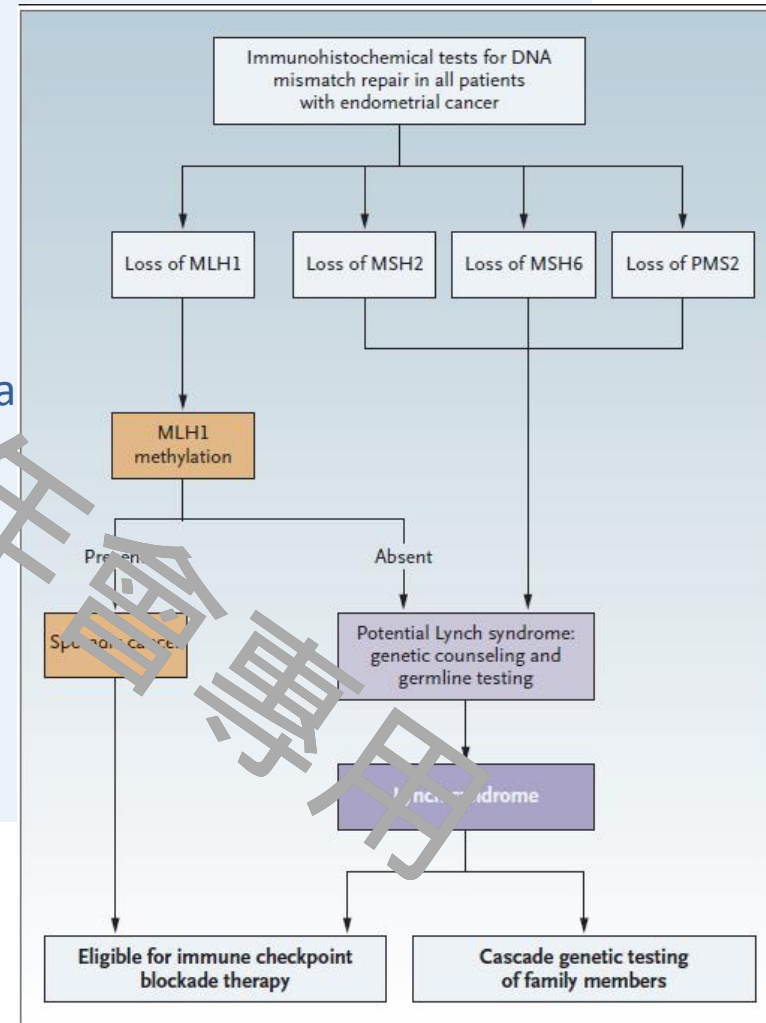
TABLE 1 2023 FIGO staging of cancer of the endometrium.<sup>a,b</sup>

Stage	Description
Stage I	Confined to the uterine corpus and ovary <sup>c</sup>
IA	Disease limited to the endometrium OR non-aggressive histological type, i.e. low-grade endometrioid, with invasion of less than half of myometrium with no or focal lymphovascular space involvement (LVSI) OR good prognosis disease
IA1	Non-aggressive histological type limited to an endometrial polyp OR confined to the endometrium
IA2	Non-aggressive histological types involving less than half of the myometrium with no or focal LVSI
IA3	Low-grade endometrioid carcinomas limited to the uterus and ovary <sup>c</sup>
IB	Non-aggressive histological types with invasion of half or more of the myometrium, and with no or focal LVSI <sup>d</sup>
IC	Aggressive histological types limited to a polyp or confined to the endometrium
Stage II	Invasion of cervical stroma with extra-uterine extension OR with substantial LVSI OR aggressive histological types with myometrial invasion
IIA	Invasion of the cervical stroma of non-aggressive histological types
IIB	Substantial LVSI <sup>d</sup> of non-aggressive histological types
IIC	Aggressive histological types <sup>e</sup> with any myometrial involvement
Stage III	Local and/or regional spread of the tumor of any histological subtype
IIIA	Invasion of uterine serosa, adnexa, or both by direct extension or metastasis
IIIA1	Spread to ovary or fallopian tube (except when meeting criteria A3 criteria)
IIIA2	Involvement of uterine subserosa or spread through the uterine serosa
IIIB	Metastasis or direct spread to the vagina and/or to the parametria or pelvic peritoneum
IIIB1	Metastasis or direct spread to the vagina and/or the parametria
IIIB2	Metastasis to the pelvic peritoneum
IIIC	Metastasis to the pelvic or para-aortic lymph nodes or both <sup>f</sup>
IIIC1	Metastasis to the pelvic lymph nodes
IIIC1i	Micrometastasis
IIIC1ii	Macrometastasis
IIIC2	Metastasis to para-aortic lymph nodes up to the renal vessels, with or without metastasis to the pelvic lymph nodes
IIIC2i	Micrometastasis
IIIC2ii	Macrometastasis
Stage IV	Spread to the bladder mucosa and/or intestinal mucosa and/or distance metastasis
IVA	Invasion of the bladder mucosa and/or the intestinal/bowel mucosa
IVB	Abdominal peritoneal metastasis beyond the pelvis
IVC	Distant metastasis, including metastasis to any extra- or intra-abdominal lymph nodes above the renal vessels, lungs, liver, brain, or bone

# Why testing for MMR/MSI relevant in endometrial and ovarian carcinoma?

## Recommended testing for MMR status/MSI in all EC, irrespective of age

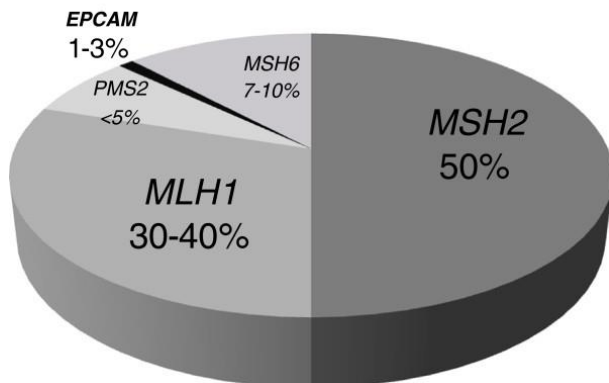
- Diagnostic, MMRd/MSI considered a marker for endometrioid-type EC
- Pre-screening to identify patients at higher risk for Lynch syndrome
  - 3-5% of all endometrial cancer
  - About 10% of MMRd/MSI-H endometrial carcinoma
  - EC often precedes subsequent ca (sentinel cancers)
- Prognostic, as identified by TCGA classification
  - Hypermuted MMRd/MSI category
- Predictive testing for MMRd tumors eligible for immune checkpoint inhibitors



# Lynch syndrome

- Autosomal dominant inherited, hereditary cancer syndrome  
Prevalence? 1 in 600 to 1 in 3,000 individuals
- Germline mutation in 1 of 4 MMR genes (MLH1, MSH2, MSH6, PMS2) or the epithelial cell adhesion molecule (EPCAM)
- Different risks for cancers  
Colorectum (18–61%), endometrium (16–61%), ovaries(5-10%), stomach, small bowel, bile duct, pancreas, and upper urinary tract
  - 2% to 6% (2.3 %) of endometrial cancers (5–9 % in <50 years)
  - 2.2% in women with colorectal cancer
- Screening and identification of patients by clinical characteristics Amsterdam criteria II (3-2-1 rule) and revised Bethesda criteria (2004)

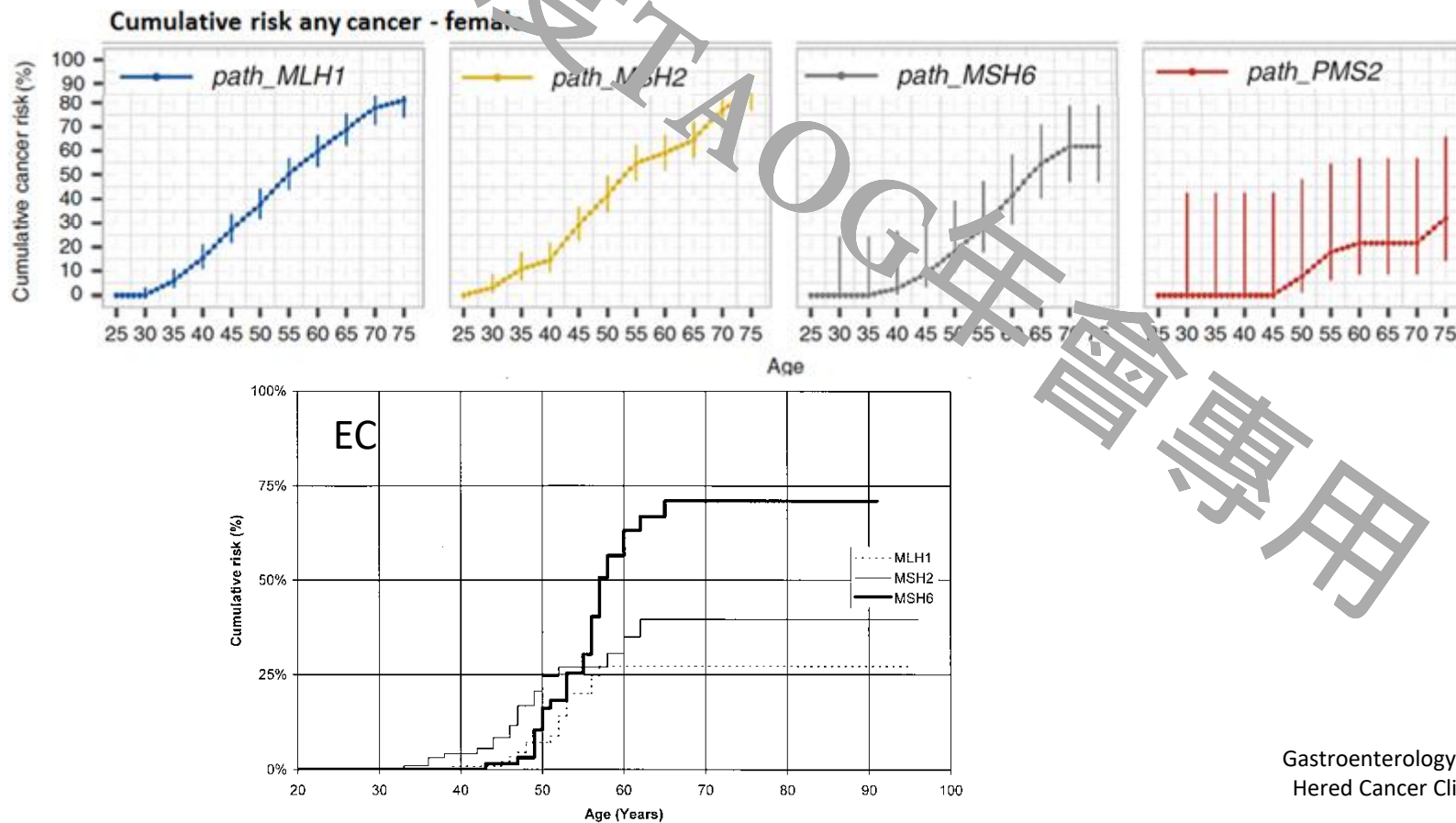
LYNCH SYNDROME MUTATIONS



MLH1 and MSH2 mutations ~ 90% (NCBI)

# Lynch syndrome

- Highest cancer risks in path\_MLH1 and path\_MSH2 carriers
- Penetrance for path\_MSH6 variants lower but females had high risks for gynecological cancers
- Higher risk for EC for MSH6 mutation carriers than MLH1 and MSH2

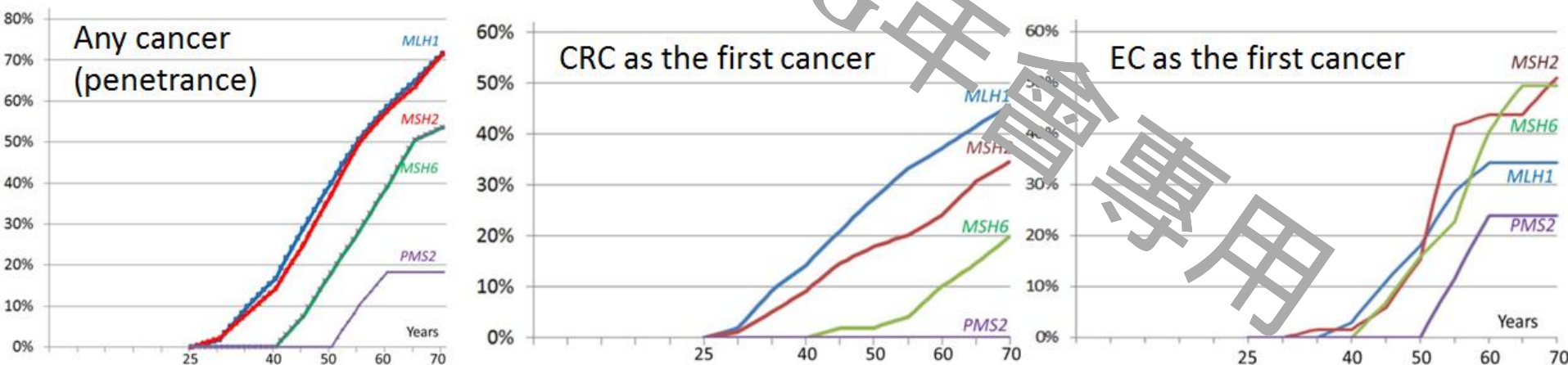




# Lynch syndrome

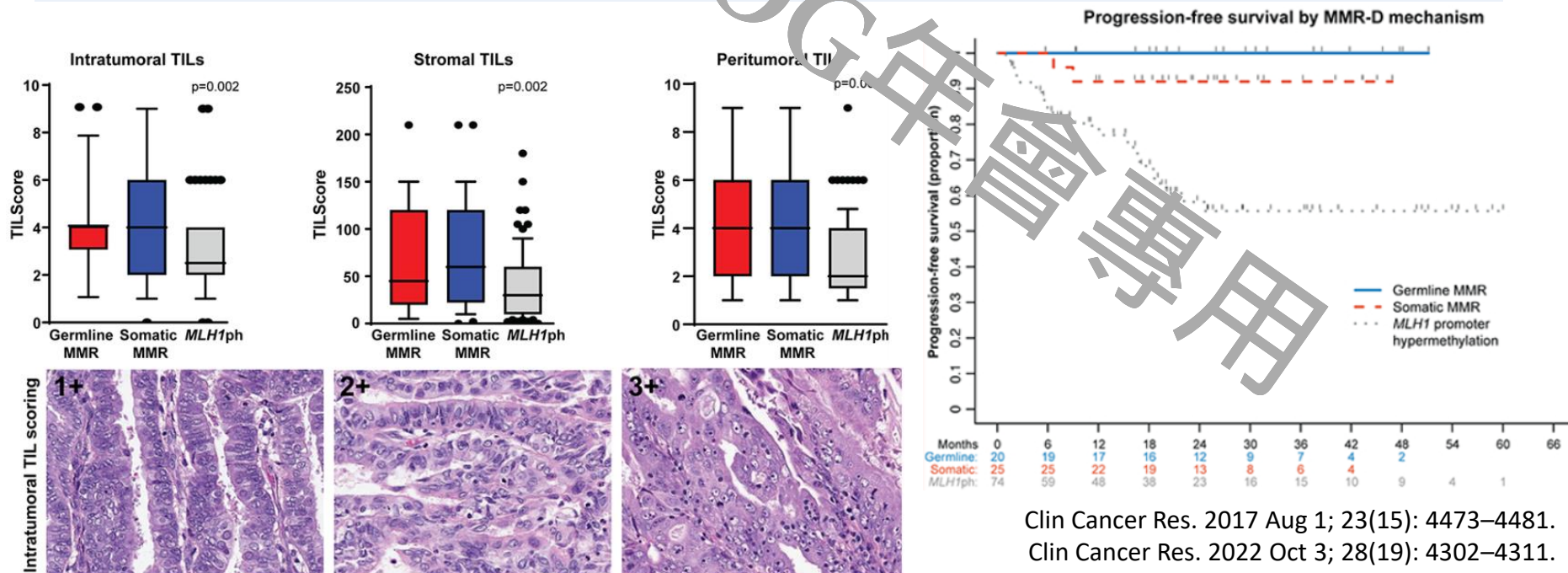
- Age of cancer onset varies among specific mutated genes, types of mutations
- Cumulative incidences for EC at 70 years:  
34%, 51%, 49%, and 24% for MLH1, MSH2, MSH6, and PMS2 mut
- Cumulative incidences for OC at 70 years:  
11%, 15%, 0%, and 0% for MLH1, MSH2, MSH6, and PMS2 mut
- Surveillance for EC in general start at the age of 35 years  
Ryan et al suggest gyn surveillance from age 30 (MSH2 mut), 35 (MLH1 mut), 40 (MSH6 mut)

Calculated cumulative incidences by age and mutated gene



# Hereditary or sporadic origin matters?

- Increased immune response in MSI-H tumors with increased immune cell infiltration and PD-L1 positive cells
- Equivalent immune response in MSI-H EC with sporadic or inherited Lynch syndrome origins?
- Increased CD8+ cells and activated CTLs in stroma: LS > sporadic MSI-H cases
- TIL score: germline/somatic MMR mutations > MLH1ph ECs
- MLH1ph EC was associated with inferior PFS



Clin Cancer Res. 2017 Aug 1; 23(15): 4473–4481.

Clin Cancer Res. 2022 Oct 3; 28(19): 4302–4311.

# Microsatellite Instability (MSI)

## ■ Microsatellites, short tandem repeats

Repetitive DNA sequences, 1-6 bases, in both coding and noncoding regions along the genome, particularly sensitive to DNA mismatching errors during DNA replication or damage

## ■ MSI A condition of genetic hypermutability

- Clustering of mutations in microsatellites consisting of repeat length alterations, phenotypic evidence of defective DNA mismatch repair
- A marker of dMMR, characterizes a hypermutable state
- Assessed with :
  - (i) Defective expression of MMR proteins as determined by IHC
  - (ii) Molecular tests, including PCR-based tests and NGS approaches

# DNA Mismatch Repair, MMR

## ■ DNA MMR

Restore DNA mismatching errors, single base mismatches or short indels

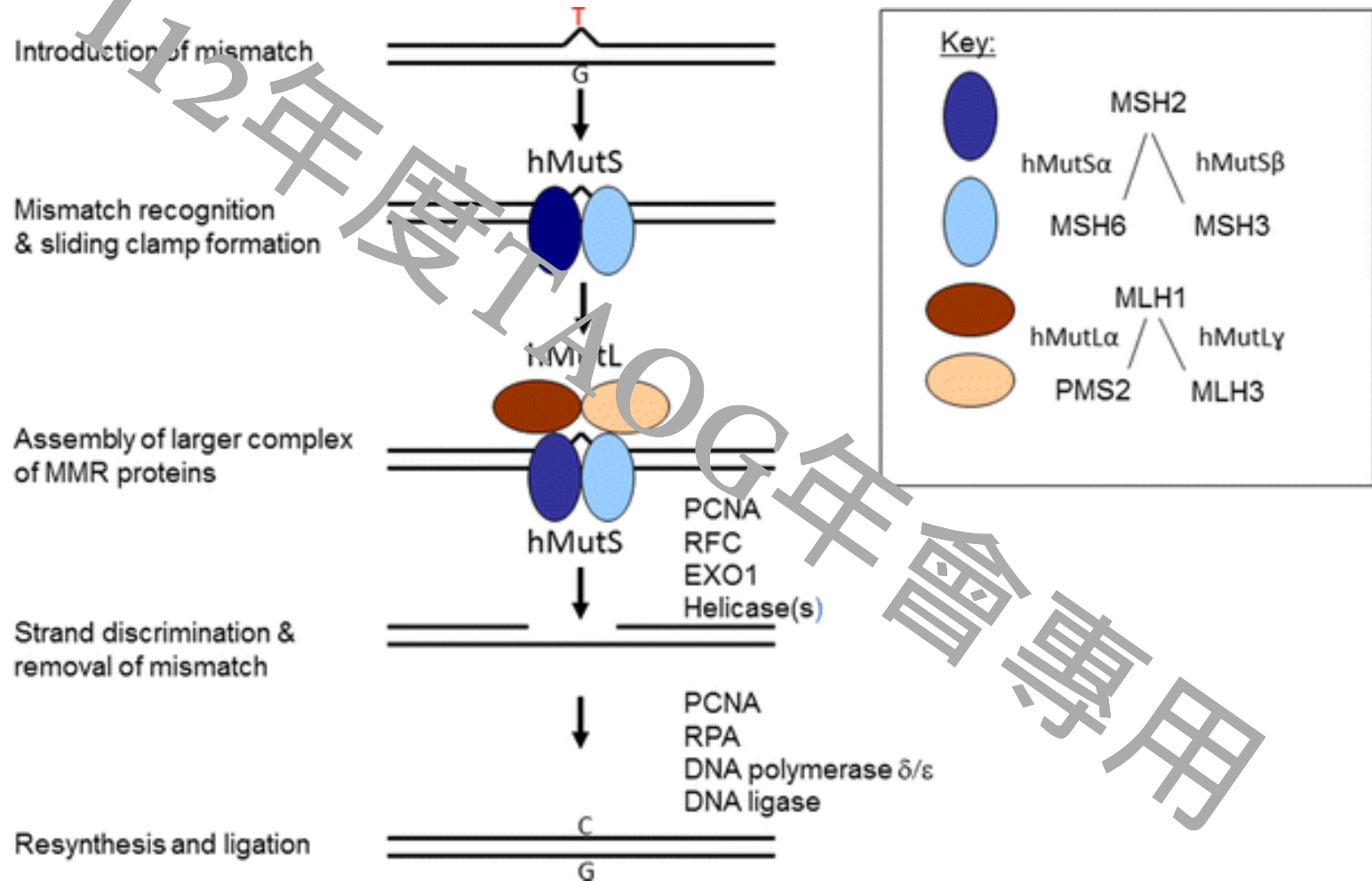
- Critical genes include MLH1, MSH2, MSH6 and PMS2
- Function in heterodimers, MLH1-PMS2 and MSH2-MSH6.
- Germline, somatic mutations or epigenetic silencing of one of these genes results in a defective MMR (dMMR)

## ■ MSI-H/MMRd tumor

A tumor that accumulates thousand of mutations, particularly clustered in microsatellites, consisting of repeat length alterations, result in MSI

Heterodimer	Components	Type of mismatches repaired	Predominant type of MSI (defective gene)
hMutSa	MSH2 + MSH6	Single-base mismatches, ins/del loops	MSI-H (MSH2 or MSH6)
hMutSβ	MSH2 + MSH3	Ins/del loops	MSI-L/EMAST (MSH3)
hMutLa	MLH1 + PMS2	Single-base mismatches, ins/del loops	MSI-H (MLH1 or PMS2)
hMutLβ	MLH1 + PMS1	?	?
hMutLy	MLH1 + MLH3	Single-base mismatches, small loops	MSI-L/EMAST or MSI-H (MLH3)

# DNA Mismatch Repair, MMR



PCNA, proliferating cell nuclear antigen; RFC, replication factor C;  
EXO1, exonuclease 1; RPA, replication protein A  
Familial Cancer 15, 385–393 (2016).



# MSI status defined by IHC

- IHC for 4 MMR proteins required in cancer type belonging to spectrum of Lynch syndrome [colorectal, endometrial, small intestine, urothelial, gliomas/glioblastomas and sebaceous gland]

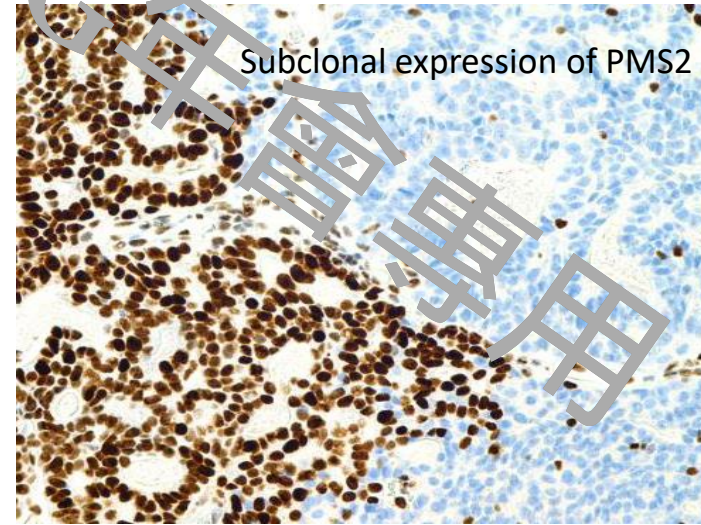
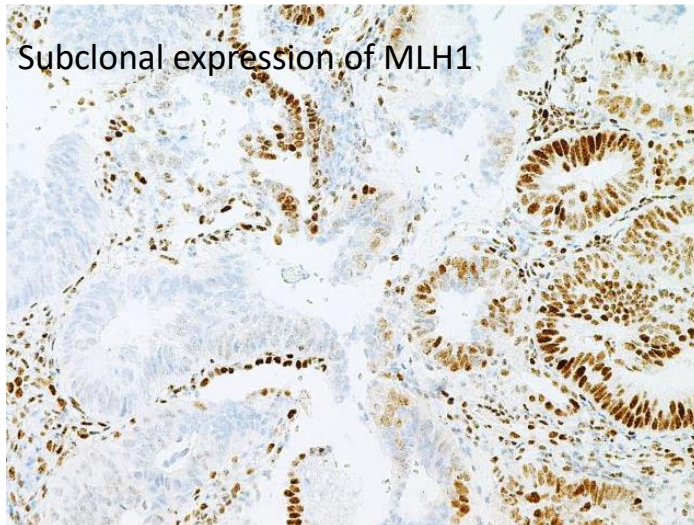
MMR genes mutation interfere with dimerization → Heterodimers degradation

→ Loss of both obligatory and secondary proteins

- Mutations in MLH1 → IHC loss of both MLH1 and PMS2
- Mutations in MSH2 → IHC loss of both MSH2 and MSH6
- PMS2 antibody detects MLH1 or PMS2 abnormalities
- MSH6 antibody detects MSH2 or MSH6 abnormality
- Advantages
  - Perform IHC on biopsies or surgical specimens?
- Pitfalls
  - False negative due to tissue fixation, aberrant staining patterns
  - Missense mutation with catalytically inactive but antigenically intact mutant protein
  - Lack of PMS2 or MSH6 substituted by other secondary proteins (MSH3, MLH3, PMS1)

# Common problems in MMR IHC interpretation

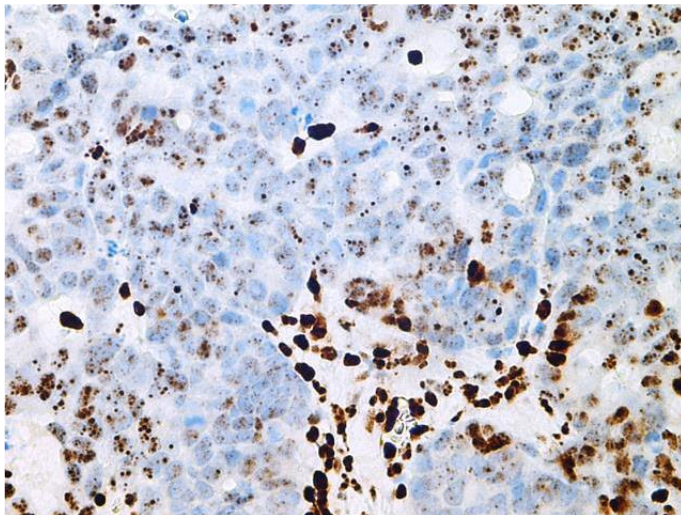
- Suboptimal fixation
- Defective or equivocal staining: very weak or focal expression seen in the presence of MMRd; unknown diagnostic, and clinical implications of this pattern
- Subclonal expression (focal expression loss): normal staining must be seen as internal control; occur as acquired defect during tumor progression, such as MLH1 promoter methylation; “passenger mutation” in MSH6 gene



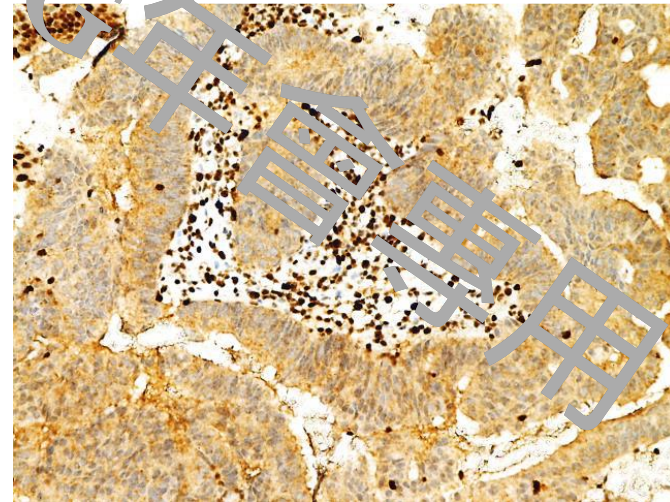
# Common problems in MMR IHC interpretation

- Punctate nuclear expression pattern in some case of MLH1 loss: erroneously interpreted as retained/normal expression
- Cytoplasmic/membranous staining should be reported as abnormal
- Others: 3 or more proteins loss, MMR IHC/MSI or MMR IHC/genetic testing discordancy...

Punctate nuclear staining for MLH1



Cytoplasmic staining for PMS2





# MSI status defined by IHC

- ESMO recommendations
- The first test of choice is IHC
- Use all 4 MMR proteins. Whether testing all 4 antibodies simultaneously or in a sequential manner, i.e. using two-antibody screening followed by reflex IHC for the appropriate partner protein
- Screen for MMRd by testing for PMS2 and MSH6: cost-effective but equivalent accuracy to testing for all 4 proteins ?
- Move to MSI-PCR whenever there is any doubt in IHC interpretation

**Table 5** Selected literature reports on patterns of immunohistochemical staining for MLH1, MSH2, MSH6, and PMS2 in extraintestinal neoplasms

Tumor site	Reference	Total no.	Abnormal mismatch repair protein immunohistochemical staining pattern							All intact
			MLH1/ PMS2	MLH1- only	PMS2- only	MSH2/ MSH6	MSH2- only	MSH6- only	Other patterns	
Skin sebaceous tumor	Orta <i>et al</i> <sup>39</sup>	27	2	0	1	8	0	1	0	15
Gynecologic tract	Modica <i>et al</i> <sup>39</sup>	85	23	0	6	6	1	9	1 with MLH1/MSH2/ MSH6, 2 with MLH1/ PMS2/MSH6	37
Gynecologic tract	Garg <i>et al</i> <sup>40</sup>	71	19	0	0	9	0	4	0	39
Gynecologic tract	Backes <i>et al</i> <sup>41</sup>	140	24	0	0	4	0	2	0	110
Ampulla of Vater	Agaram <i>et al</i> <sup>42</sup>	54	1	0	0	0	0	2	0	51

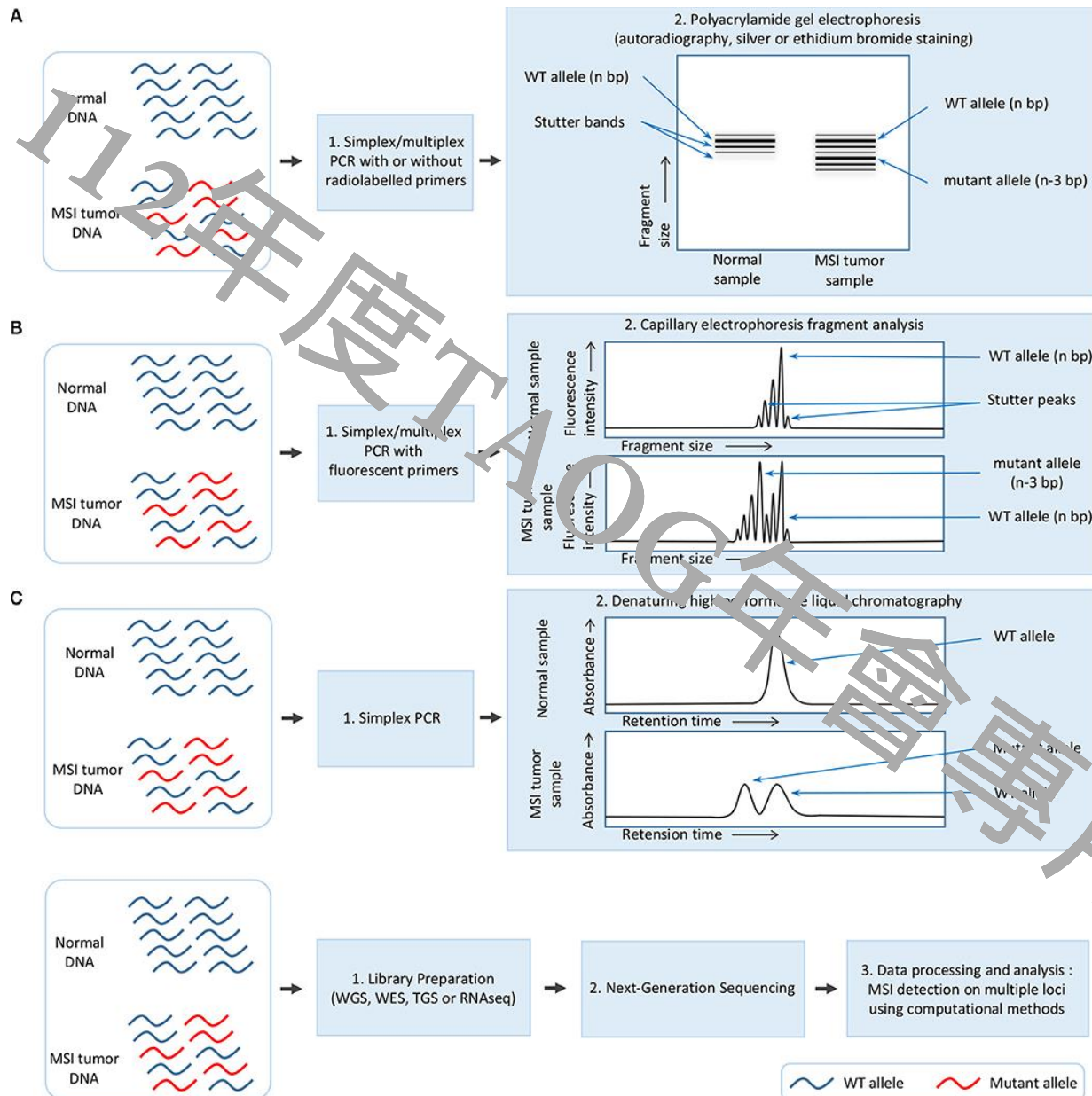
# MSI status defined by PCR-based testing

- MSI-PCR molecular testing indicated in indeterminate IHC results or in case of loss of only one heterodimer subunit
- PCR amplification of microsatellite markers
  - Panel 1 (Bethesda/NCI): BAT-25, BAT-26, D5S346, D2S123 and D17S250
  - Panel 2: BAT-25, BAT-26, NR-21, NR-24, NR-27 (Sen 95.6%, Spe 100% ).
- MSI: 2+ mononucleotide markers show repeats length alteration

The terms MSI-H ( $\geq 2$ ) or MSI-L (1 marker) should be abandoned and MSI-L should be included with MSS tumors
- NGS-based MSI testing
  - Couple MSI analysis with the determination of TMB
  - Larger set of microsatellites
  - Identify other targetable alterations other than immunotherapy
    - MSIplus panel, ColonCore Panel, smMIP panel



# MSI status defined by PCR-based testing



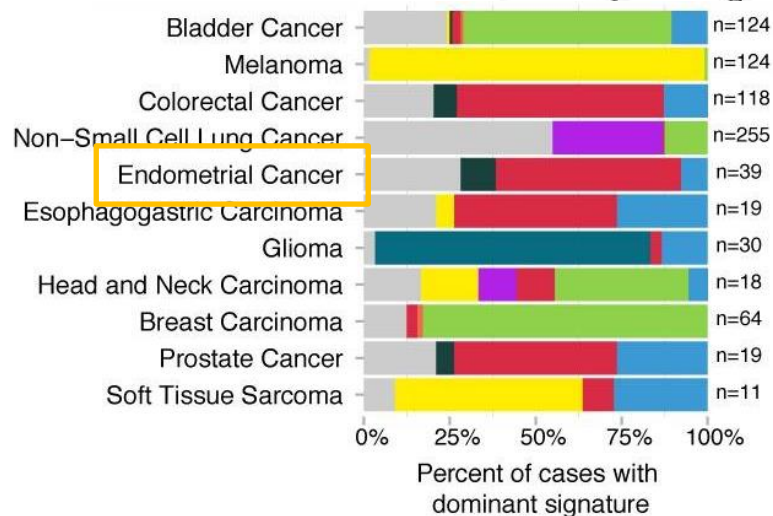
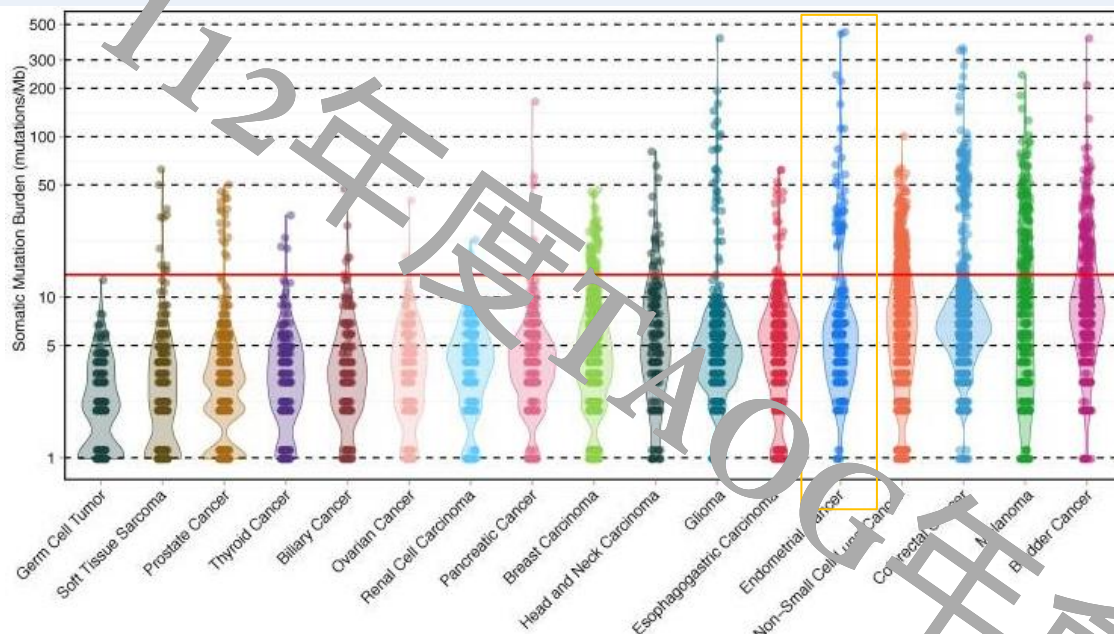
# Testing Algorithm for MSI

- Better diagnostic performance of molecular assays in CRC than in EC
- In EC, lower sensitivity for PCR assay (67%) and NGS (75%)
- Discrepant IHC and molecular MSI in samples with loss of MSH6 expression

	Case number	Tumor type	Tumor cell %	IHC result	IHC pattern	PCR										NGS										IHC																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
						MSI-H	MSI-L	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI	MSI

# Significant mutational burden in endometrial cancer

- Microsatellite stable (MSS) in 70-75%; microsatellite instability high (MSI-H) in 25-30%



## Mutational signatures (MSK-IMPACT)

### Signature

- Aging
- APOBEC
- BRCA1/2
- MMR
- Smoking
- TMZ
- POLE
- UV
- Other



# Significant mutational burden in endometrial cancer

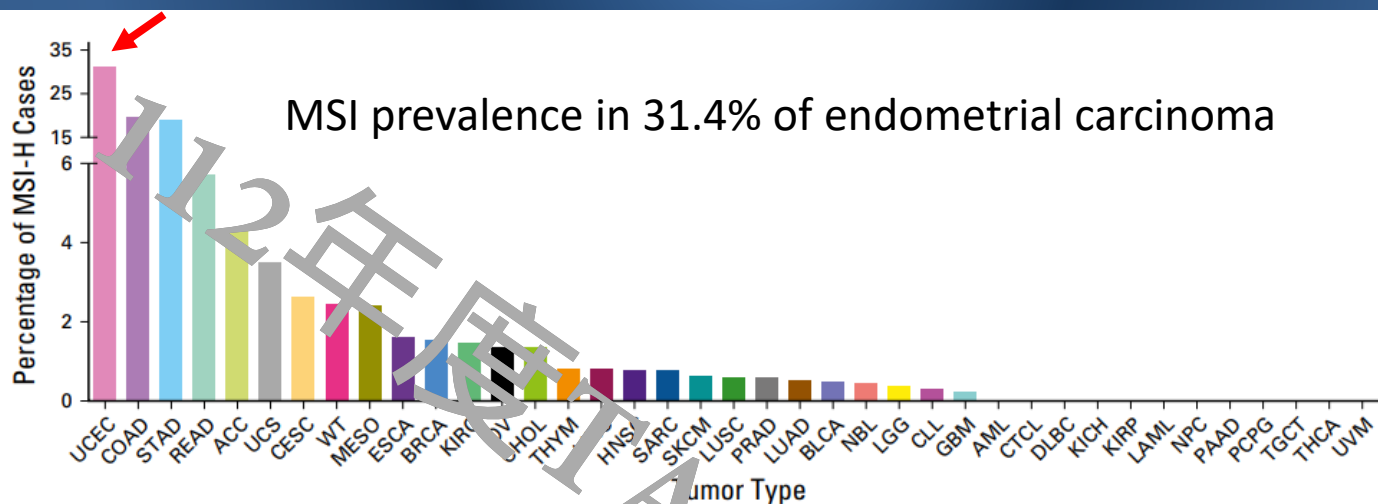
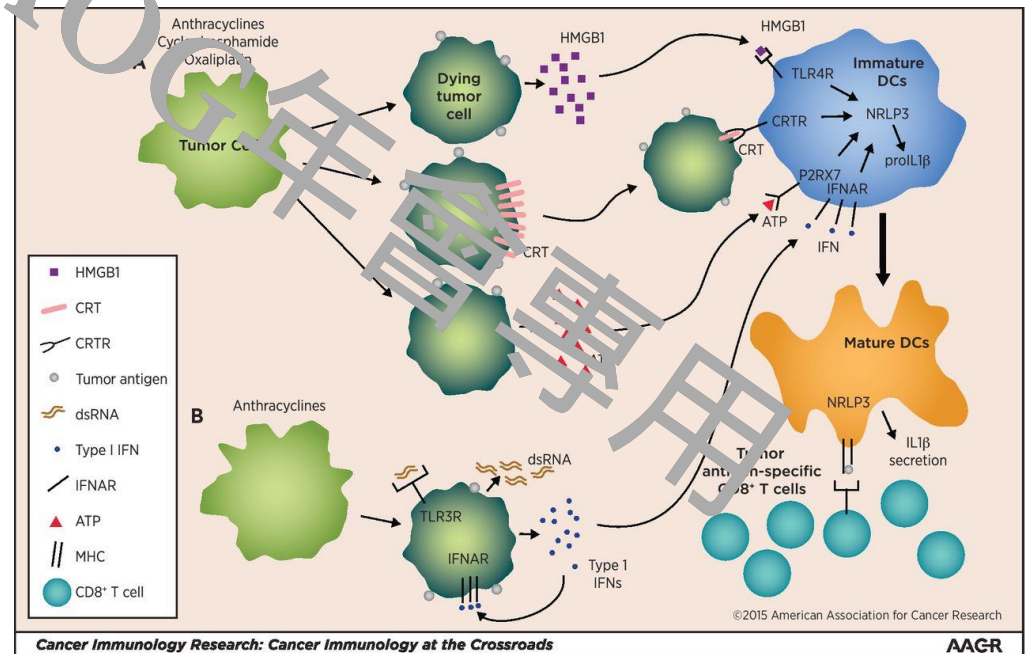
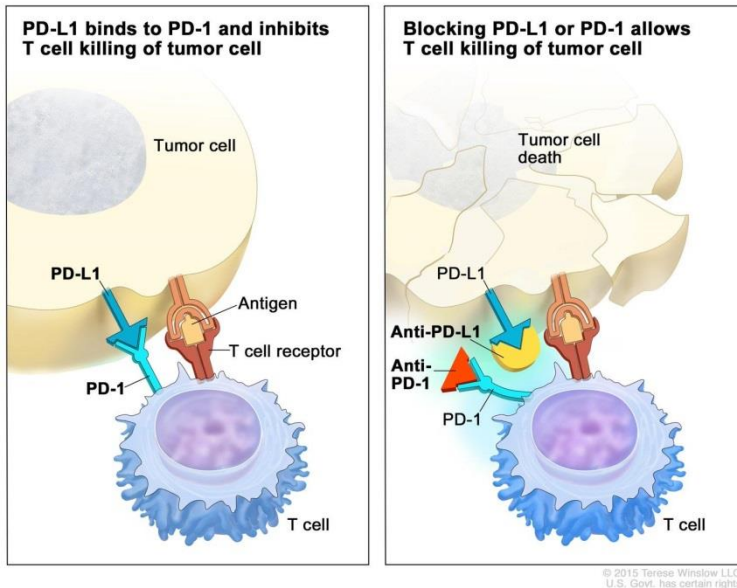


Table 2. Recommendation for MSI testing in different cancer types and in the framework of immunotherapy

Cancer type	MSI prevalence (all stages)	Specific MSI-associated histotype, if any
<b>Sporadic tumour types belonging to the spectrum of Lynch syndrome that can be tested using IHC and MSI-PCR or NGS (testing is indicated for stage IV cancers, whose MSI prevalence is lower than that of earlier stages)</b>		
Colorectal	17%	Medullary, mucinous, poorly differentiated, neuroendocrine
Endometrial	20%	Lower uterine segment-located, undifferentiated/dedifferentiated, mixed morphology, tumours showing high levels of tumour-infiltrating lymphocytes/lymphoid stroma
Gastric-oesophageal	13%	Adenocarcinoma (MSI up to 39% in case of carcinoma with lymphoid stroma, and absent in oesophageal squamous cell carcinoma)
Small intestine	8.3%	Including duodenum and ampulla of Vater
Ovarian	3.5%–10%	Endometrioid, clear cell
Glioblastoma	6%–13%	
<b>All common or rare tumour types not belonging to the spectrum of Lynch syndrome with low prevalence of MSI and little data available on the reliability of IHC and MSI-PCR, to be tested using NGS</b>		
Unknown primary	1.8%	
Cervical	4%	
Extrahepatic bile duct	3.4%	
Pancreatic	1%–7%	Medullary, IPMN-associated, periampullary (when the origin from ampulla, terminal bile duct or pancreatic duct is uncertain)
Prostate	3%	
Non-small-cell lung cancer	< 1%	
Head and neck	<1%	
Melanoma	NS	1% uveal melanoma
Sarcomas	2%	Uterine, peritoneal and retroperitoneal
Anal	NS	
Kidney	NS	

# Rationale for Immunotherapy-Based Combinations

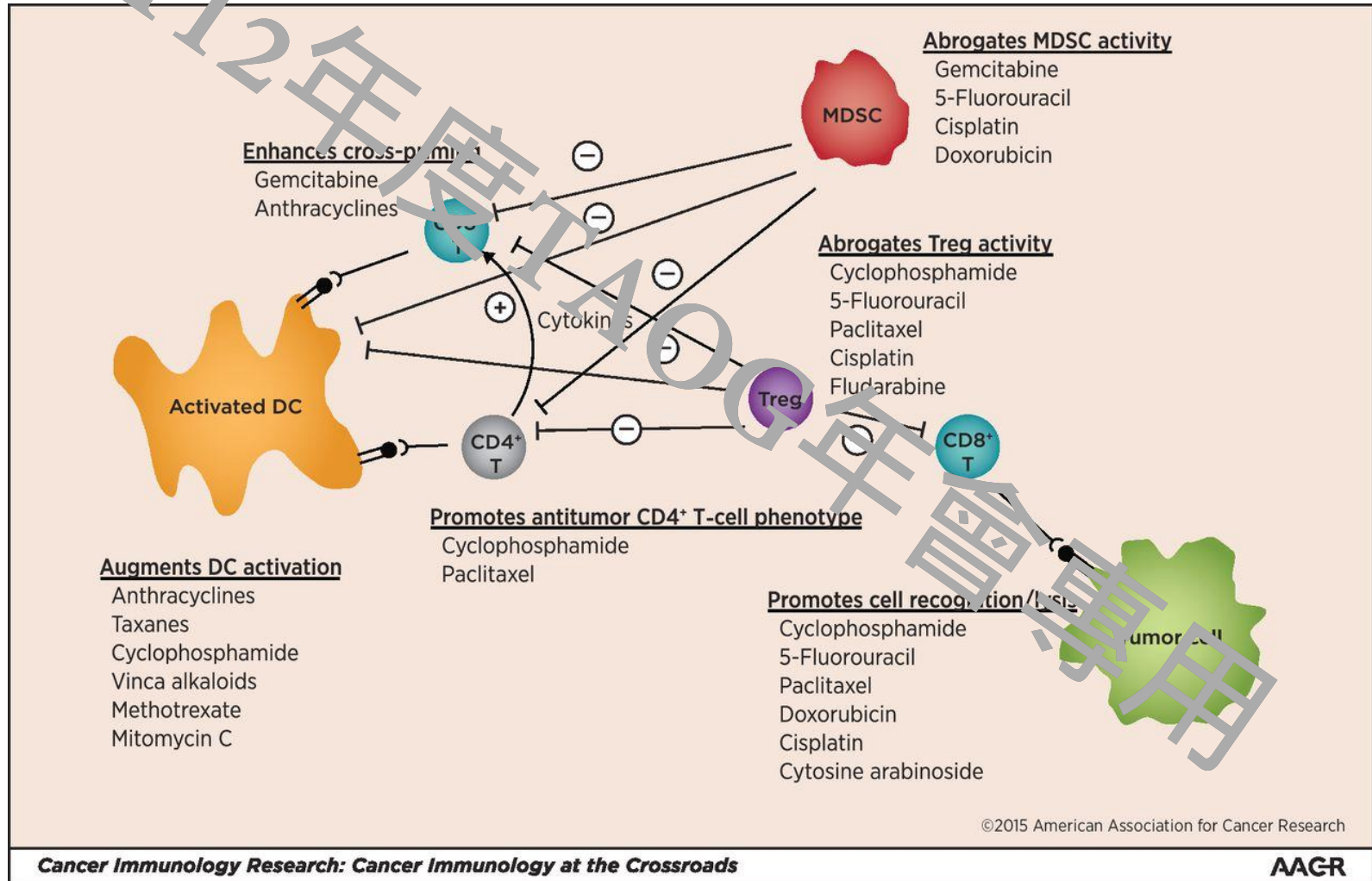
- Checkpoint immunotherapy mechanism of action
- Immune checkpoint inhibitors (ICIs) block interaction of PD-L1 with PD-1; allow T cells to kill tumor cells
- Immunogenic effects of chemotherapy
- Taxanes: T-cell priming, DC activation, MDSCs depletion
- Platinum-based agents: Downregulation of PD-L1/L2 on DC, Induction of immunogenic cancer cell death





# Rationale for Immunotherapy-Based Combinations

## *Immunogenic effects of chemotherapy*



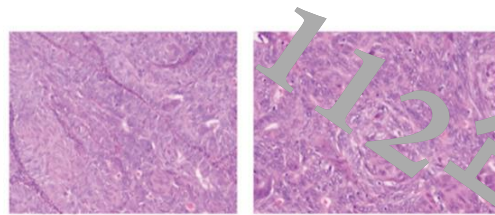
# NCCN Recommended Systemic Treatments for Endometrial Carcinoma

Primary or Adjuvant Therapy (Stage I-IV)	
Chemoradiation Therapy	Systemic Therapy
<u>Preferred Regimens</u> • Cisplatin plus RT followed by carboplatin/paclitaxel <sup>1,2</sup>	<u>Preferred Regimens</u> • Carboplatin/paclitaxel <sup>3</sup> • Carboplatin/paclitaxel/pembrolizumab (for stage III-IV tumors, except for carcinosarcoma) (category 1) <sup>a,b,4</sup> <b>NRG-GY018</b> • Carboplatin/paclitaxel/dostarlimab-gxly (for stage III-IV tumors) (category 1) <sup>b,c,5</sup> <b>RUBY (ENGOT-EN6; GOG-3031)</b> • Carboplatin/paclitaxel/trastuzumab (for stage III/IV HER2-positive uterine serous carcinoma) <sup>d,e,6</sup> • Carboplatin/paclitaxel/trastuzumab (for stage III/IV HER2-positive carcinosarcoma) (category 2B) <sup>d,e,6</sup>

# NCCN Recommended Systemic Treatments for Endometrial Carcinoma

RECURRENT DISEASE <sup>f,g</sup>	
First-Line Therapy for Recurrent Disease <sup>h</sup>	Second-Line or Subsequent Therapy
<p><b>Preferred</b></p> <ul style="list-style-type: none"> <li>• Carboplatin/paclitaxel (category 1 for carcinosarcoma)<sup>i,3</sup></li> <li>• Carboplatin/paclitaxel/pembrolizumab (except for carcinosarcoma) (category 1)<sup>b,4</sup></li> <li>• Carboplatin/paclitaxel/dostarlimab-gxly (category 1)<sup>b,5</sup></li> <li>• Carboplatin/paclitaxel/trastuzumab<sup>e</sup> (for HER2-positive uterine serous carcinoma)<sup>d,6</sup></li> <li>• Carboplatin/paclitaxel/trastuzumab<sup>e</sup> (for HER2-positive carcinosarcoma) (category 2B)<sup>d,6</sup></li> </ul> <p><b>Other Recommended Regimens</b></p> <ul style="list-style-type: none"> <li>• Carboplatin/docetaxel<sup>j</sup></li> <li>• Carboplatin/paclitaxel/bevacizumab<sup>k,7,8</sup></li> </ul> <p><b>Useful in Certain Circumstances</b> (Biomarker directed: after prior platinum-based therapy including neoadjuvant and adjuvant)</p> <ul style="list-style-type: none"> <li>• Lenvatinib/pembrolizumab (category 1) for mismatch repair proficient (pMMR) tumors<sup>b,9</sup></li> <li>• Pembrolizumab<sup>b</sup> for TMB-H<sup>l,10</sup> or MSI-H/dMMR<sup>m</sup> tumors<sup>11</sup></li> <li>• Dostarlimab-gxly for dMMR/MSI-H tumors<sup>b,12</sup></li> </ul>	<p><b>Other Recommended Regimens</b></p> <ul style="list-style-type: none"> <li>• Cisplatin/doxorubicin<sup>13</sup></li> <li>• Cisplatin/doxorubicin/paclitaxel<sup>n,13</sup></li> <li>• Cisplatin</li> <li>• Carboplatin</li> <li>• Doxorubicin</li> <li>• Liposomal doxorubicin</li> <li>• Paclitaxel<sup>14</sup></li> <li>• Albumin-bound paclitaxel<sup>o</sup></li> <li>• Topotecan</li> <li>• Bevacizumab<sup>k,p,15</sup></li> <li>• Temsirolimus<sup>16</sup></li> <li>• Cabozantinib</li> <li>• Docetaxel<sup>f</sup> (category 2B)</li> <li>• Ifosfamide (for carcinosarcoma)</li> <li>• Ifosfamide/paclitaxel (for carcinosarcoma)<sup>17</sup></li> <li>• Cisplatin/ifosfamide (for carcinosarcoma)</li> </ul> <p><b>Useful in Certain Circumstances</b> (Biomarker directed therapy)</p> <ul style="list-style-type: none"> <li>• Lenvatinib/pembrolizumab (category 1) for mismatch repair proficient (pMMR) tumors<sup>b,9</sup></li> <li>• Pembrolizumab<sup>b</sup> for TMB-H<sup>l,8</sup> or MSI-H/dMMR<sup>m</sup> tumors<sup>m,11</sup></li> <li>• Dostarlimab-gxly for dMMR/MSI-H tumors<sup>b,12</sup></li> <li>• Larotrectinib or entrectinib for NTRK gene fusion-positive tumors (category 2B)</li> <li>• Avelumab for dMMR/MSI-H tumors<sup>b</sup></li> <li>• Nivolumab for dMMR/MSI-H tumors<sup>b,18</sup></li> </ul>

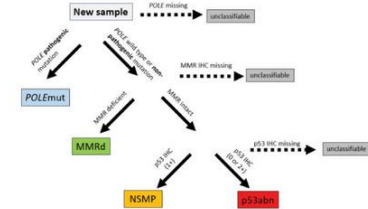
# Molecular subtype-specific adjuvant therapy



Endometrial Biopsy or Curettage

## Pathology + Molecular Classification

Impact surgical decision /staging



### POLEmut

Early ~90%  
Advanced ~10%

De-escalation e.g., surgery only

? De-escalation  
? Radiation only

**Clinicopathologic and molecular parameters; e.g., LVI, grade, myoinvasion, L1CAM do not add prognostic or predictive stratification**

**Adjuvant Clinical Trials**

- PORTEC-4a (de-escalation)
- RAINBO Blue (de-escalation)
- TAPER (de-escalation)

### MMRd

Adjuvant Radiation

? No additional benefit from chemotherapy

Loss of MSH6, MSH2, PMS2

Loss of MLH1

Not Methylated

Methylated

Hereditary Cancer Referral

Consider ICB

? Less response to ICB  
? Pembro + Lenvat

**Further stratified by MLH1 and immune profile?**

**Adjuvant Clinical Trials**

- NRG GY 020 (+ICB; pembrolizumab)
- RAINBO Green (+ICB; durvalumab)
- ADELE Trial (+ICB; tislelizumab)

### NSMP

Additional Stratification

- LVI
- Grade
- ER/PR
- L1CAM?
- CTAD/B1?
- Immune e.g., CD8?

Highly Favourable

- De-escalation
- Vault brachy
- Endocrine therapy

Unfavourable

- Chemotherapy
- Pembro + Lenvat

**Clinicopathologic parameters do add prognostic and possible predictive stratification**

**Adjuvant Clinical Trials**

- RAINBO Orange (+endocrine therapy)
- TAPER (de-escalation)

### p53abn

Stage IA  
No myoinvasion

Stage IA with myoinvasion/Stage IB+

? Surgery only  
? Vault brachy  
? ChemoRT

Significant benefit from chemotherapy

HRD (20-25%) PARPi

HER2 (20-25%) Anti-HER2 Rx

CCNE1 (20-30%) Wee1-i

Pembro + Lenvat (50% response in serous)

**Aggressive tumors regardless of grade and histotype**

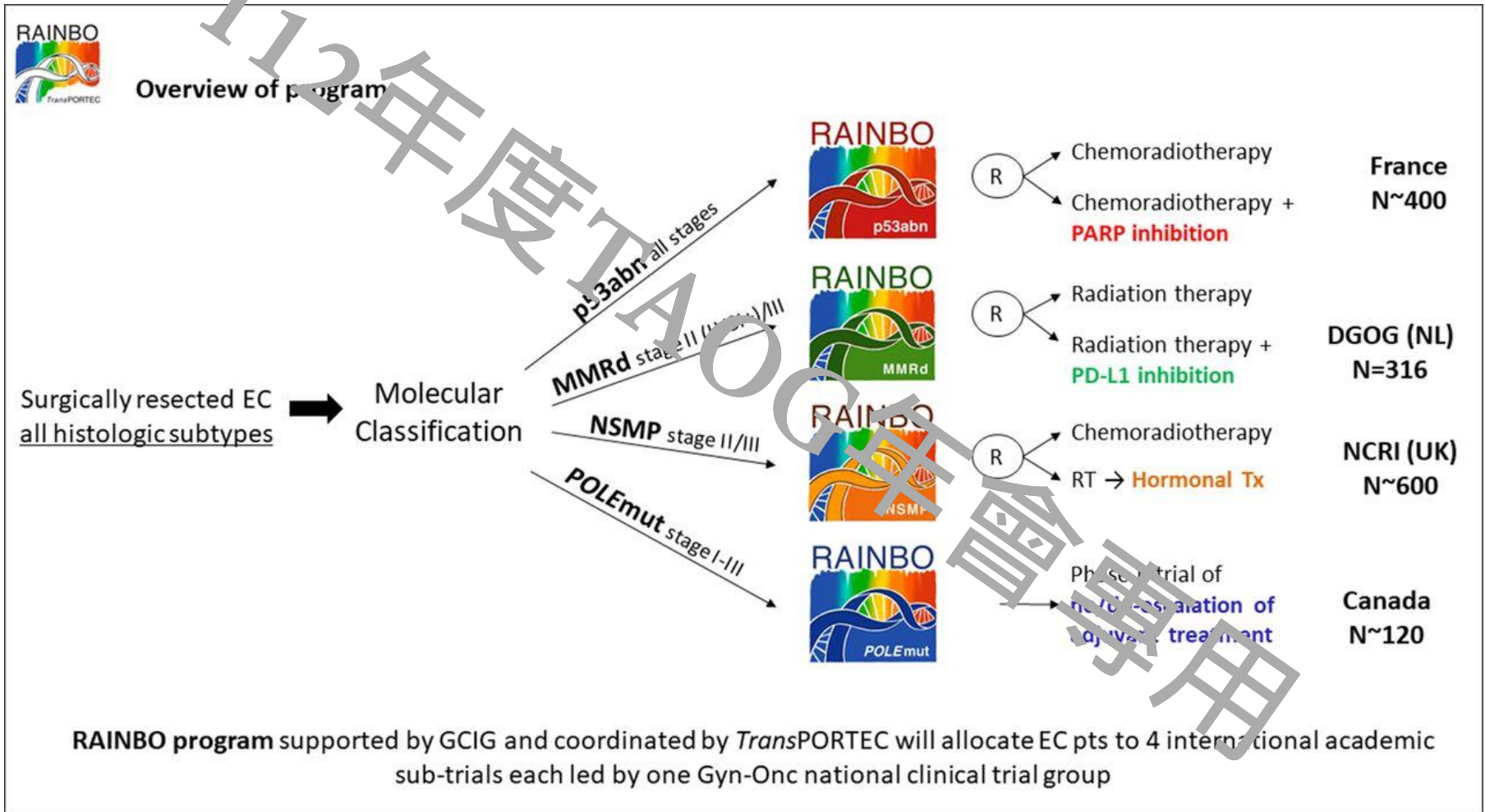
**Clinicopathologic parameters do add prognostic and possible predictive stratification**

**Adjuvant Clinical Trials**

- RAINBO Red (+PARPi; olaparib)
- CAMP STAMP (+PARPi; niraparib)
- NRG GY 026 (+HER2; trastuzumab/pertuzumab)



# TransPORTEC RAINBO Umbrella Trial





# MMR protein loss by IHC, PCR-based or NGS based MSI analysis?

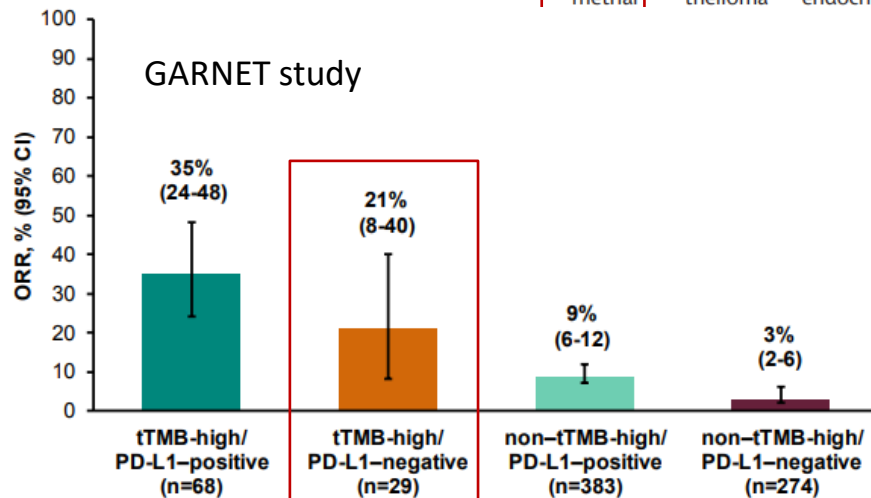
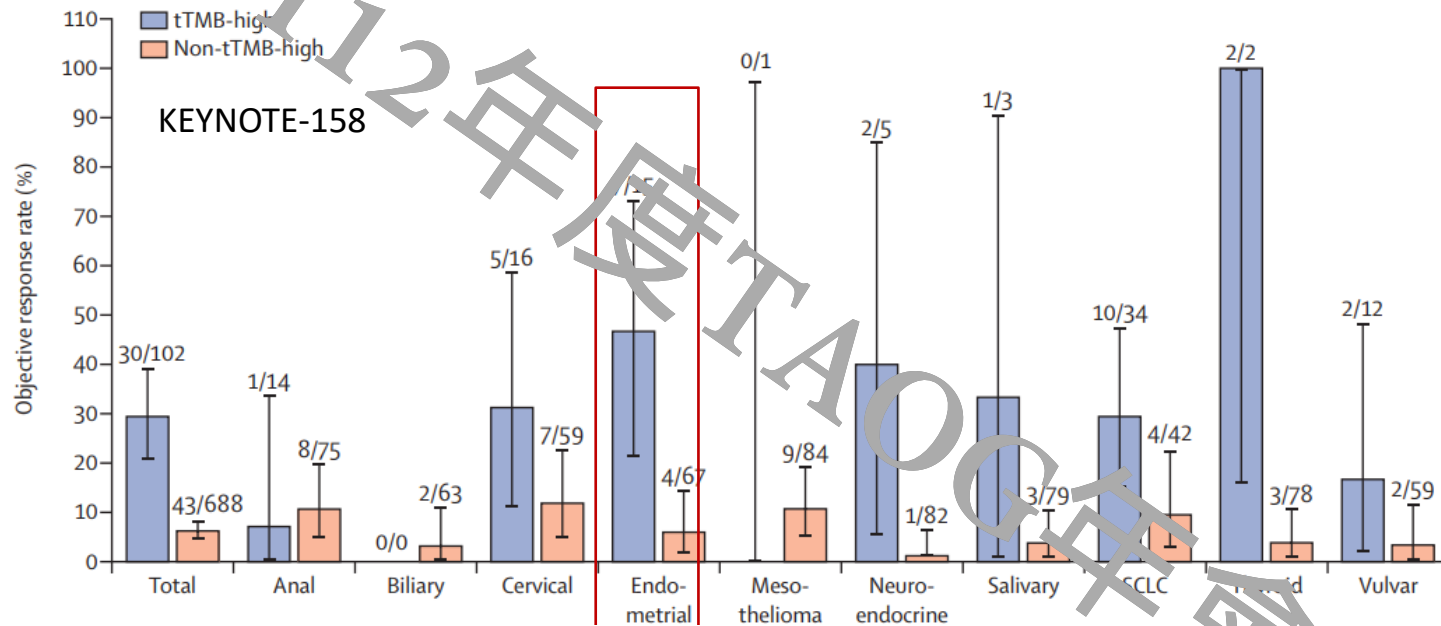
- MSI-NGS discrepancies in non-CRC cancers may due to other involved loci not measured by MSI-PCR
- Different MSI levels between EC and CRC patients with Lynch syndrome
  - MSI lower in EC patients
  - MSI lower in EC patients from MSH6 mutation carriers
  - More MSH6 mutations in EC patients (x5)
  - MSI-L or MSS status in EC with MSH6 deficiency range from 29% to 50%

Table 3. Results of MSI Analyses in Tumors of *MSH6* Mutation Carriers

Tumor	MSI high	MSI low	Microsatellite stable	Total
Colorectal carcinoma (%)	18 (86)	3 (14)	0	21
Endometrial carcinoma (%)	11 (69)	4 (25)	1 (6)	16
Transitional cell carcinoma (%)	5 (71)	2 (29)	0 (0)	7
Ovarian carcinoma	2	0	0	2
Breast carcinoma	1	0	0	1
Stomach carcinoma	0	0	1	1
Adenocarcinoma of the cervix	0	1	0	1
Total	35	9	5	49

# TMB as biomarker beyond dMMR/MSI-H

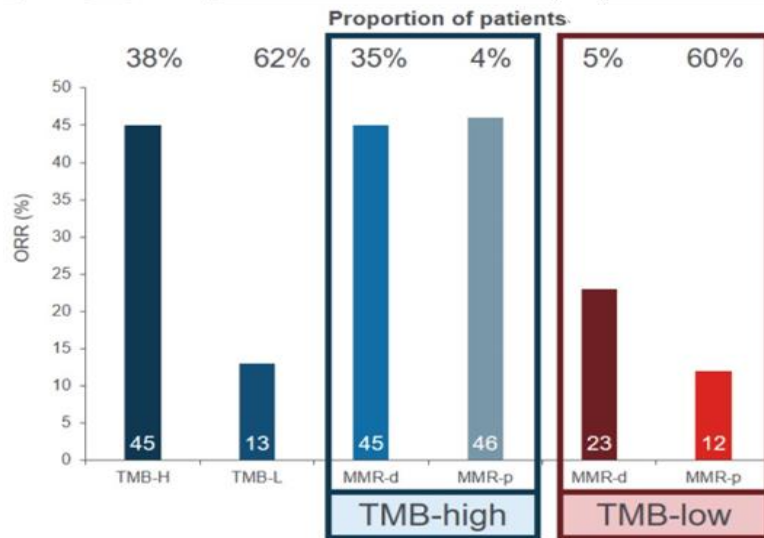
## TMB predicted outcomes with pembrolizumab irrespective of PD-L1 expression



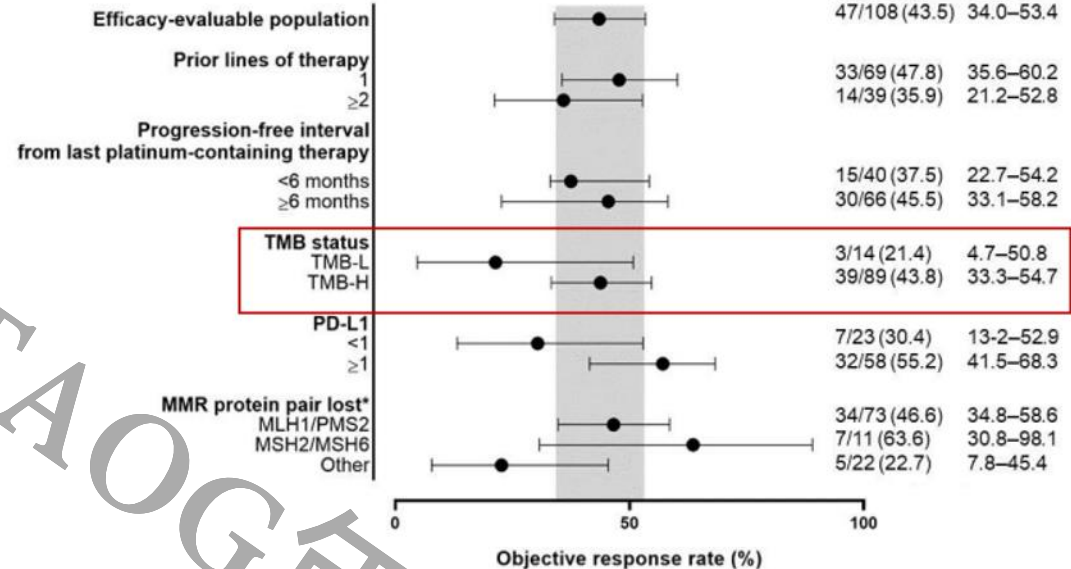
# TMB as biomarker beyond dMMR/MSI-H

## GARNET study

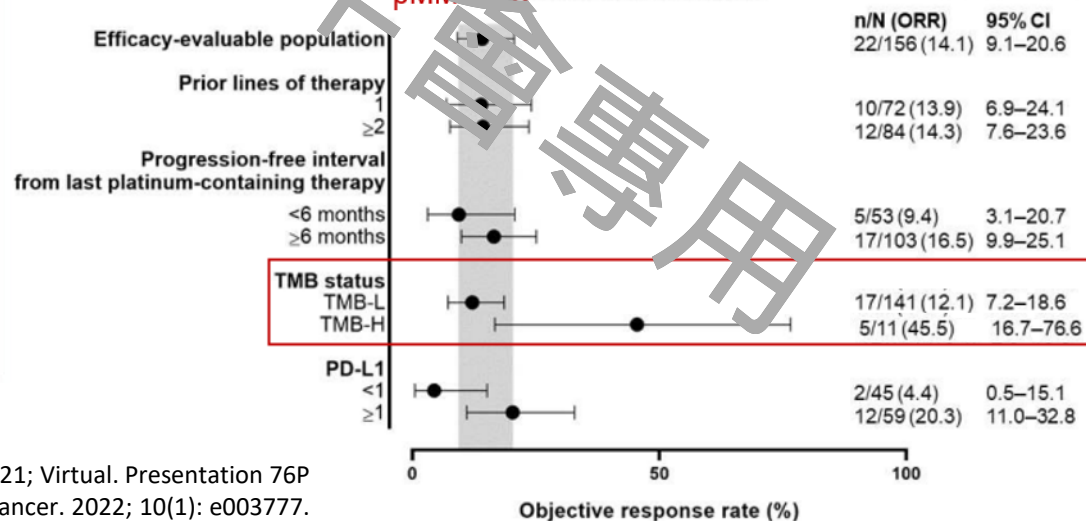
- Patients with MMR-d had high response rates
- But TMB-high tumors were associated with high responses irrespective of MMR status



## dMMR EC A1 subgroup analysis

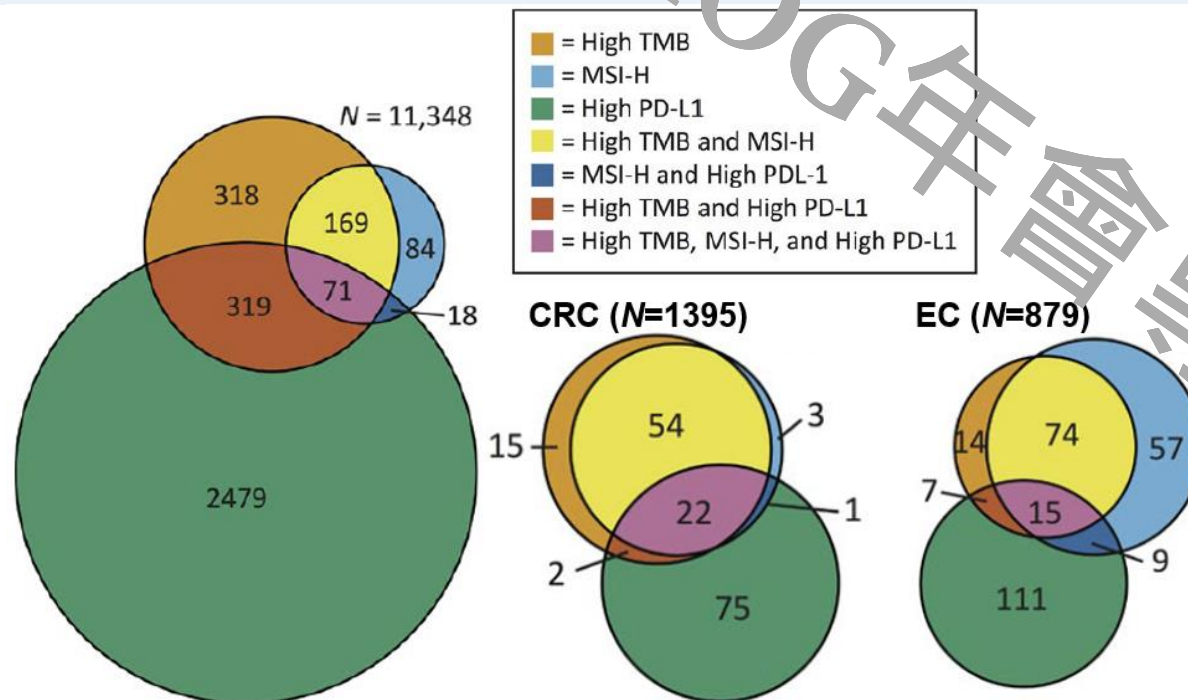


## pMMR EC A2 subgroup analysis



# TMB as a surrogate for PCR- and NGS based MSI assays?

- Elevated TMB associated with other etiologies e.g. POLE exonuclease-domain mutations
- 30% of MSI-H tumors were TMB-Low (<17 mut/Mb)
- Higher discrepant TMB and MSI-H rates in non-CRC
- 95% concordance in CRCs but only 57% MSI-H EC were TMB-H
- Discrepancy observed in OV (24%), NE (57%), and CC (33%)



# FDA Approved in vitro diagnostic companion diagnostic device

Diagnostic Name (Manufacturer)	Indication - Sample Type	Drug Trade Name (Generic) NDA / BLA	Biomarker(s)	Biomarker(s) (Details)	Clearance / Grant Date)
PD-L1 IHC 22C3 pharmDx (Dako North America, Inc.)	Cervical Cancer - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	PD-L1	PD-L1 protein expression	<a href="#">P150013/S009</a> (06/12/2018)
Ventana MMR Rx Dx Panel (Ventana Medical Systems, Inc.)	Endometrial Carcinoma (EC) - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a> in combination with Lenvima (lenvatinib) <a href="#">NDA 20694</a>	proficient mismatch repair (pMMR) proteins	MLH1, PMS2, MSH2 and MSH6	<a href="#">P210001/S002</a> (06/16/2022)
PD-L1 IHC 22C3 pharmDx (Dako North America, Inc.)	Esophageal Squamous Cell Carcinoma (ESCC) - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	PD-L1	PD-L1 protein expression	<a href="#">P150013/S016</a> (07/30/2019)
PD-L1 IHC 22C3 pharmDx (Dako North America, Inc.)	Head and Neck Squamous Cell Carcinoma (HNSCC) - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	PD-L1	PD-L1 protein expression	<a href="#">P150013/S014</a> (06/10/2019)
PD-L1 IHC 22C3 pharmDx (Dako North America, Inc.)	Non-Small Cell Lung Cancer (NSCLC) - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	PD-L1	PD-L1 protein expression	<a href="#">P150013</a> (02/02/2015)
Ventana MMR Rx Dx Panel (Ventana Medical Systems, Inc.)	Solid Tumors	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	deficient mismatch repair (dMMR) proteins	MLH1, PMS2, MSH2 and MSH6	<a href="#">P210001/S002</a> (06/16/2022)
FoundationOne CDx (Foundation Medicine, Inc.)	Solid Tumors - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	TMB	TMB ≥ 10 mutations per megabase	<a href="#">P170019/S016</a> (06/16/2020)
FoundationOne CDx (Foundation Medicine, Inc.)	Solid Tumors - Tissue	Keytruda (pembrolizumab) <a href="#">BLA 125514</a>	MSI-High	Microsatellite instability-High (MSI-H)	<a href="#">P170019/S029</a> (02/18/2022)



# FDA Approved in vitro diagnostic companion diagnostic device

Diagnostic Name (Manufacturer)	Indication - Sample type	Drug Trade Name (Generic) NDA / BLA	Biomarker(s)	Biomarker(s) (Details)	(Approval / Clearance / Grant Date)
Ventana MMR RxDx Panel (Ventana Medical Systems, Inc.)	Endometrial Carcinoma (EC) Tissue	Jemperli (dostarlimab-gxly) <a href="#">NDA 761174</a>	deficient mismatch repair (dMMR) proteins	MLH1, PMS2, MSH2 and MSH6	<a href="#">P200019</a> (04/22/2021)
Ventana MMR RxDx Panel (Ventana Medical Systems, Inc.)	Solid Tumors	Jemperli (dostarlimab-gxly) <a href="#">NDA 761174</a>	deficient mismatch repair (dMMR) proteins	MLH1, PMS2, MSH2 and MSH6	<a href="#">P210001</a> (08/17/2021)

# Summary of guidelines for patients considered immune checkpoint inhibitors

# ESMO recommendations for MSI testing

**Table 1. Summary table of recommendations for MSI testing in the framework of immunotherapy and comments from the ESMO TR and PM WG consensus panel**

## **Recommendation A: immunohistochemistry**

Coefficient of agreement: strong (8.7)

*Main comment: MMR proteins form heterodimers; for correct IHC interpretation, the consensus panel highlights that mutations in MLH1 are associated with IHC loss of both MLH1 and PMS2, while mutations in MSH2 are associated with IHC loss of both MSH2 and MSH6. There exist isolated losses of PMS2, MSH2 or MSH6, this strengthening the recommendation to use all four antibodies.*

The first test of choice is IHC, using antibodies recognising the four MMR proteins: MLH1, MSH2, MSH6 and PMS2.

## **Recommendation B: polymerase chain reaction**

Coefficient of agreement: strong (8.6)

*Main comment: both the suggested panels have been and are being used to assess MSI in clinical trials. Molecular tests guarantee the highest values of specificity and sensitivity in MSI testing.*

In case of doubt of IHC, confirmatory molecular analysis is mandatory. The first-line of molecular analysis is represented by PCR. It can be carried out using two possible panels: (i) a panel with two mononucleotide (BAT-25 and BAT-26) and three dinucleotide (D5S346, D2S123 and D17S250) repeats and (ii) a panel with five poly-A mononucleotide repeats (BAT-25, BAT-26, NR-21, NR-24, NR-27). The five poly-A panel is the recommended panel given its higher sensitivity and specificity.

## **Recommendation C: next-generation sequencing**

Coefficient of agreement: very strong (9.0)

*Main comment: NGS should be carried out only in selected centres devoted to these techniques.*

NGS represents another type of molecular tests to assess MSI. Its main advantages are represented by the possibilities of coupling MSI analysis with the determination of tumour mutational burden (TMB).

# 2022 CAP (College of American Pathologists) guidelines

Guideline Statement	Strength of Recommendation
1. For patients with CRC being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-IHC and/or MSI by PCR for the detection of DNA mismatch repair defects. Although MMR-IHC or MSI by PCR are preferred, pathologists may use a validated MSI by NGS assay for the detection of DNA mismatch repair defects Note: MSI by NGS assay must be validated against MMR-IHC or MSI by PCR and must show equivalency	Strong Recommendation
2. For patients with gastroesophageal and small bowel cancer being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-IHC and/or MSI by PCR over MSI by NGS for the detection of DNA mismatch repair defects Note: This recommendation does not include esophageal squamous cell carcinoma	Strong Recommendation
3. For patients with endometrial cancer being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-IHC over MSI by PCR or NGS for the detection of DNA mismatch repair defects	Strong Recommendation <b>Certainty of Evidence: Low</b>
4. For patients with cancer types other than CRC, GEA, small bowel, and endometrial being considered for immune checkpoint inhibitor therapy, pathologists should test for DNA mismatch repair, although the optimal approach for the detection of mismatch repair defects has not been established Note: Assays must be adequately validated for the specific cancer type being tested with careful consideration of performance characteristics of MMR-IHC and MSI by NGS or PCR for the detection of DNA mismatch repair defects	Conditional Recommendation
5. For all cancer patients being considered for immune checkpoint inhibitor therapy based on defective mismatch repair, pathologists should not use TMB as a surrogate for the detection of DNA mismatch repair defects. If a tumor is identified as TMB-High, pathologists may perform IHC and/or MSI by PCR to determine if high TMB is secondary to mismatch repair deficiency	Strong Recommendation <b>Certainty of Evidence: Low</b>
6. For cancer patients being considered for immune checkpoint inhibitor therapy, if a mismatch repair deficiency consistent with Lynch syndrome is identified in the tumor, pathologists should communicate this finding to the treating physician	Strong Recommendation

Table 4. Number of Studies by Outcome

Test	Rec 1 CRC	Rec 2 GEA and SI	Rec 3 EC	Rec 4 Other Cancer	Rec 5 TMB	Rec 6 LS
MMR-IHC diagnostic test characteristics	2 PCS 8 RCS	2 RCS	3 RCS	2 PCS 2 RCS	0	0
MMR-IHC status concordance with germline testing	1 PCS 8 RCS	0	2 RCS	0	0	1 PCS 9 RCS
MSI-PCR diagnostic test characteristics	2 PCS 7 RCS	0	3 RCS	1 RCS	0	0
MSI-PCR status concordance with germline testing	1 PCS 2 RCS	0	1 RCS	0	0	3 PCS 10 RCS
MMR-IHC and MSI-PCR status concordance	6 PCS 16 RCS	4 RCS	2 PCS 7 RCS	2 RCS	0	0
MSI-NGS diagnostic test characteristics	1 PCS 5 RCS	0	1 PCS 2 RCS	5 RCS	0	0
MSI-NGS and MMR-IHC status concordance	1 PCS 1 RCS	1 PCS	2 PCS	1 PCS 1 RCS	0	0
MSI-NGS and MSI-PCR status concordance	0	0	0	1 RCS	0	0
TMB diagnostic test characteristics	0	0	0	0	1 RCS	0
TMB and MMR-IHC status concordance	0	0	0	0	2 RCS	0
TMB and MSI-NGS status concordance	0	0	0	0	1 PCS 2 RCS	0
Association between LS prevalence and MMR MSI status	0	0	0	0	0	5 PCS 10 RCS



# Endorsement of CAP guidelines by ASCO

- Consider other potential information via NGS testing in decision making.  
Beyond MSI detection,
- Detection of HER2 amplification, high TMB due to non-MSI mechanisms

**Recommendation 1.** For patients with CRC being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-immunohistochemistry (IHC) and/or microsatellite instability (MSI) by polymerase chain reaction (PCR) for the detection of DNA MMR defects. Although MMR-IHC or MSI by PCR is preferred, pathologists may use a validated MSI by next-generation sequencing (NGS) assay for the detection of DNA MMR defects. *Note: MSI by NGS assay must be validated against MMR-IHC or MSI by PCR and must show equivalency.* (Strong recommendation)

**Recommendation 2.** For patients with gastroesophageal and small bowel cancer, being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-IHC and/or MSI by PCR over MSI by NGS for the detection of DNA MMR defects. *Note: This recommendation does not include esophageal squamous cell carcinoma.* (Strong recommendation)

**Recommendation 3.** For patients with endometrial cancer, being considered for immune checkpoint inhibitor therapy, pathologists should use MMR-IHC over MSI by PCR or NGS for the detection of DNA MMR defects. (Strong recommendation)

**Recommendation 4.** For patients with cancer types other than CRC, gastroesophageal adenocarcinoma, small bowel, and endometrial being considered for immune checkpoint inhibitor therapy, pathologists should test for DNA MMR although the optimal approach for the detection of MMR defects has not been established. *Note: Assays must be adequately validated for the specific cancer type being tested with careful consideration of performance characteristics of MMR-IHC and MSI by NGS or PCR for the detection of DNA MMR defects.* (Conditional recommendation)

**Recommendation 5.** For all cancer patients being considered for immune checkpoint inhibitor therapy based on defective MMR, pathologists should not use tumor mutation burden (TMB) as a surrogate for the detection of DNA MMR defects. If a tumor is identified as TMB-high, pathologists may perform IHC and/or MSI by PCR to determine if high TMB is secondary to MMR deficiency. (Strong recommendation)

**Recommendation 6.** For cancer patients being considered for immune checkpoint inhibitor therapy, if a MMR deficiency consistent with Lynch syndrome is identified in the tumor, pathologists should communicate this finding to the treating physician. (Strong recommendation)





Thank you for your attention