

The many uses of p53 IHC in Gynecological Pathology

PRESENTED BY

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Martin Köbel reported no relevant financial relationships

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YOUR CELL PHONES**



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The Journal of Pathology: Clinical Research

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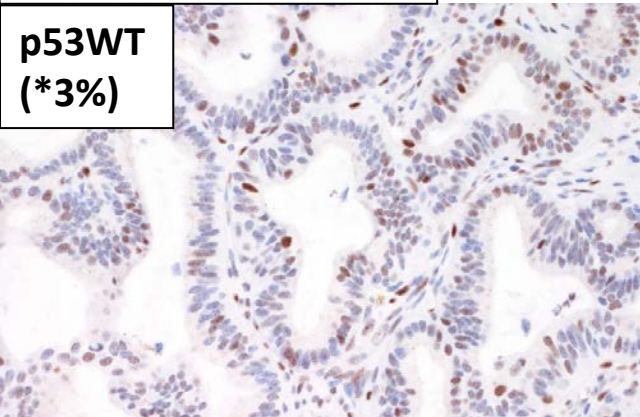
Original Article

Optimized p53 immunohistochemistry is an accurate predictor of *TP53* mutation in ovarian carcinoma

Martin Köbel,^{1†} Anna M Piskorz,^{2†} Sandra Lee,¹ Shuhong Lui,¹ Cecile LePage,^{3,4} Francesco Marass,² Nitzan Rosenfeld,² Anne-Marie Mes Masson^{3,4} and James D Brenton^{2*}

The normal Wild-type pattern

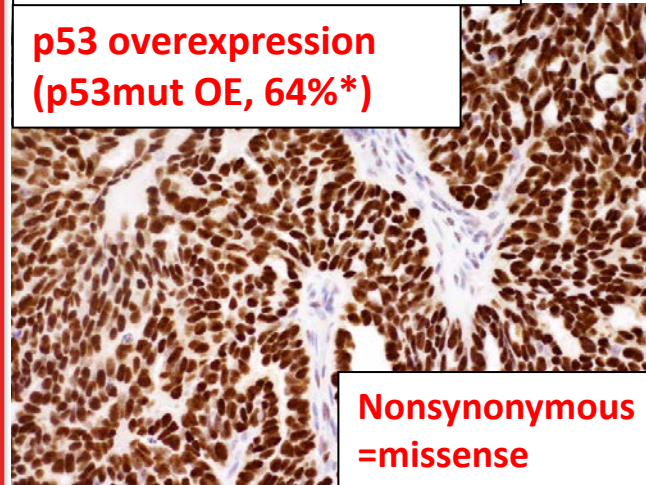
p53WT
(*3%)



No *TP53* mutation (rare exceptions)

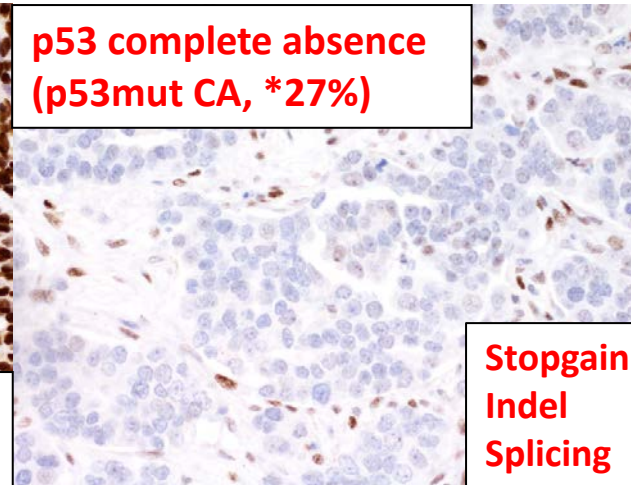
3 Mutant Patterns:

p53 overexpression
(p53mut OE, 64%*)



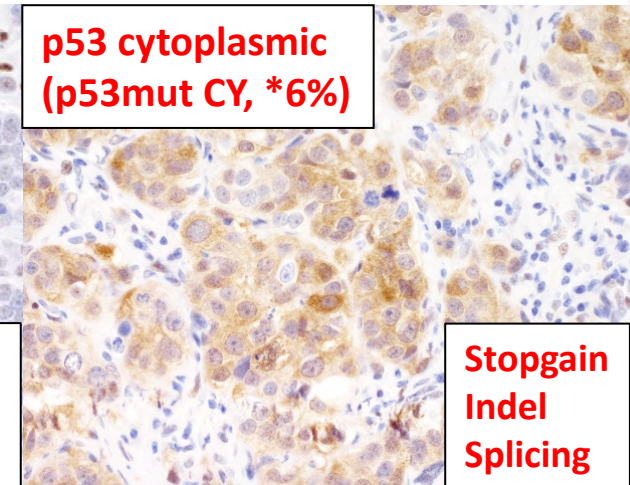
Nonsynonymous
=missense

p53 complete absence
(p53mut CA, *27%)

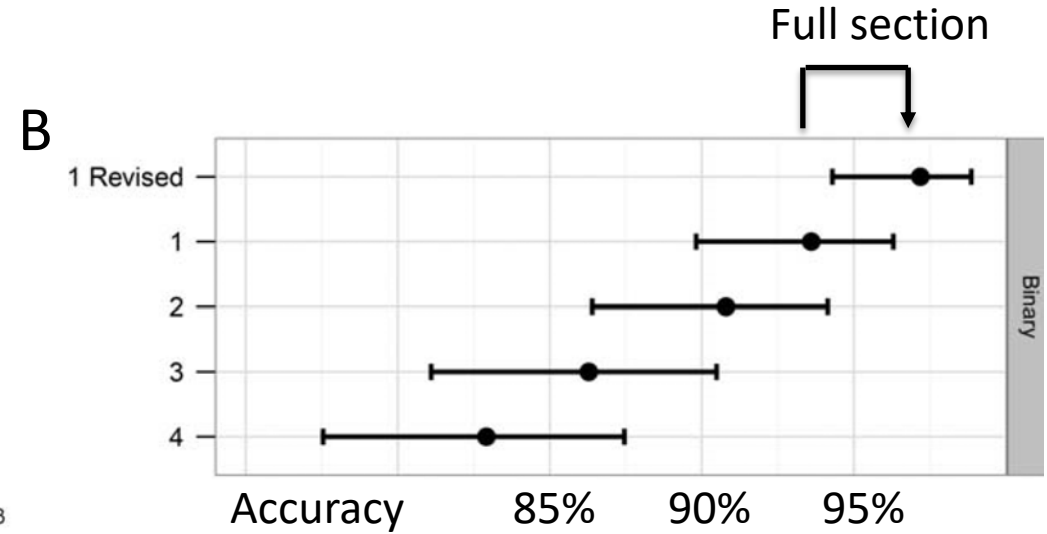
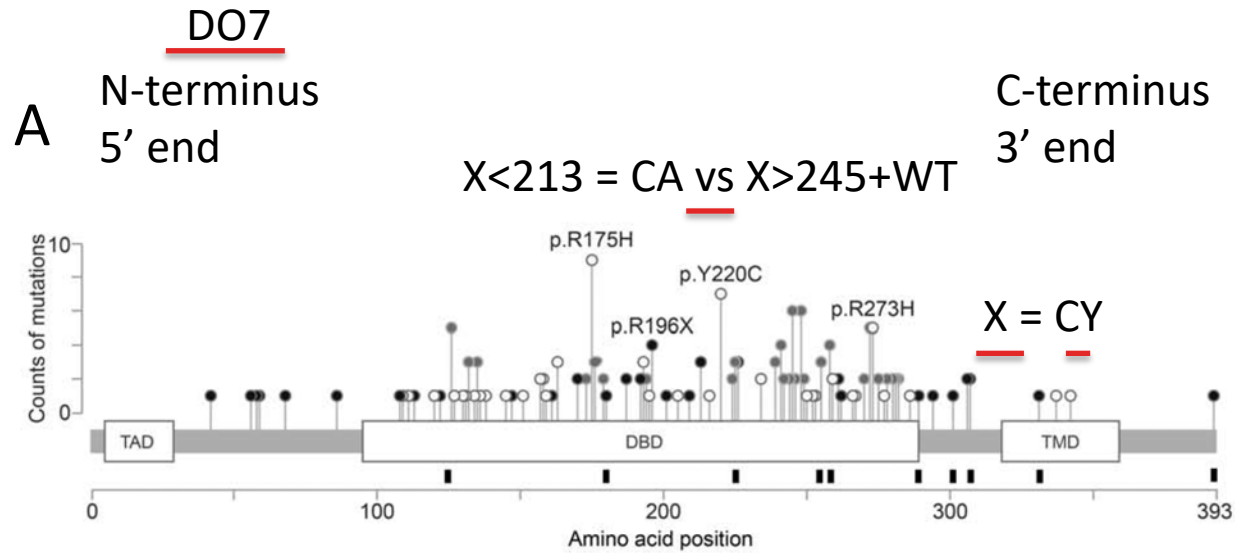


Stopgain
Indel
Splicing

p53 cytoplasmic
(p53mut CY, *6%)



Stopgain
Indel
Splicing

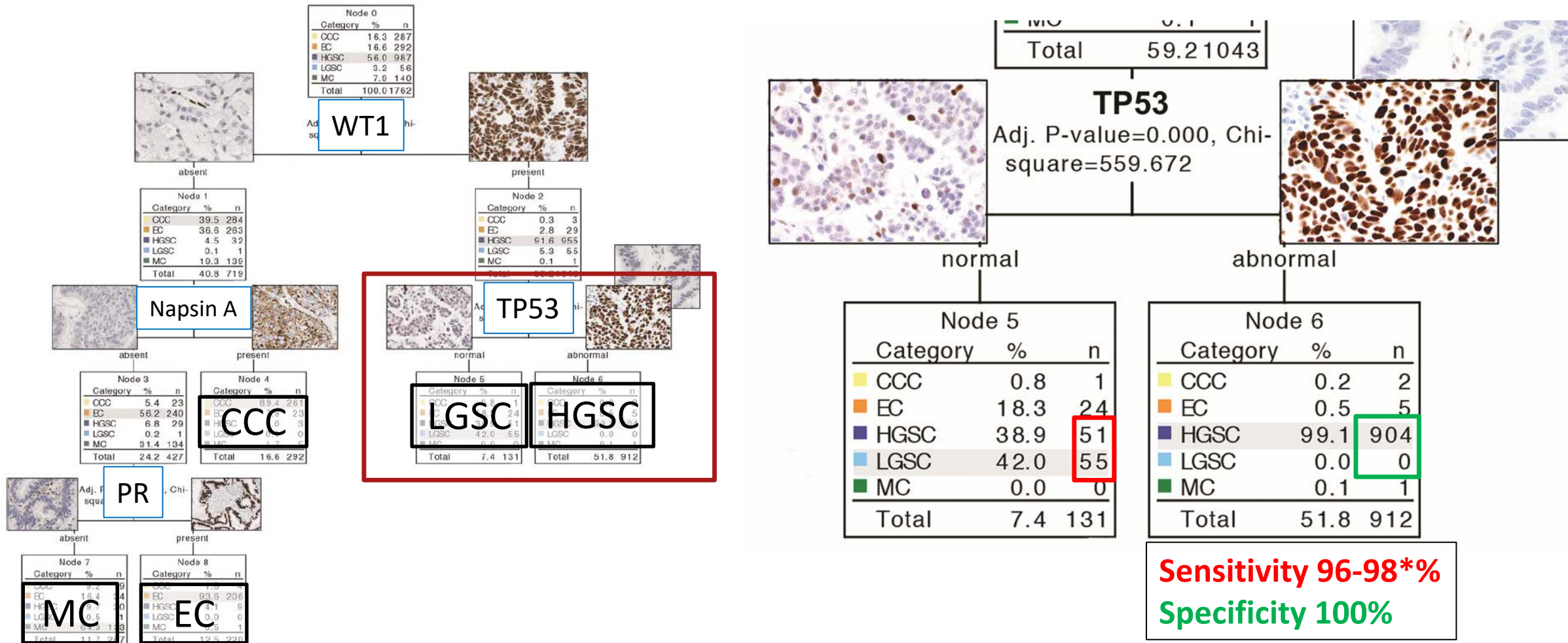


C

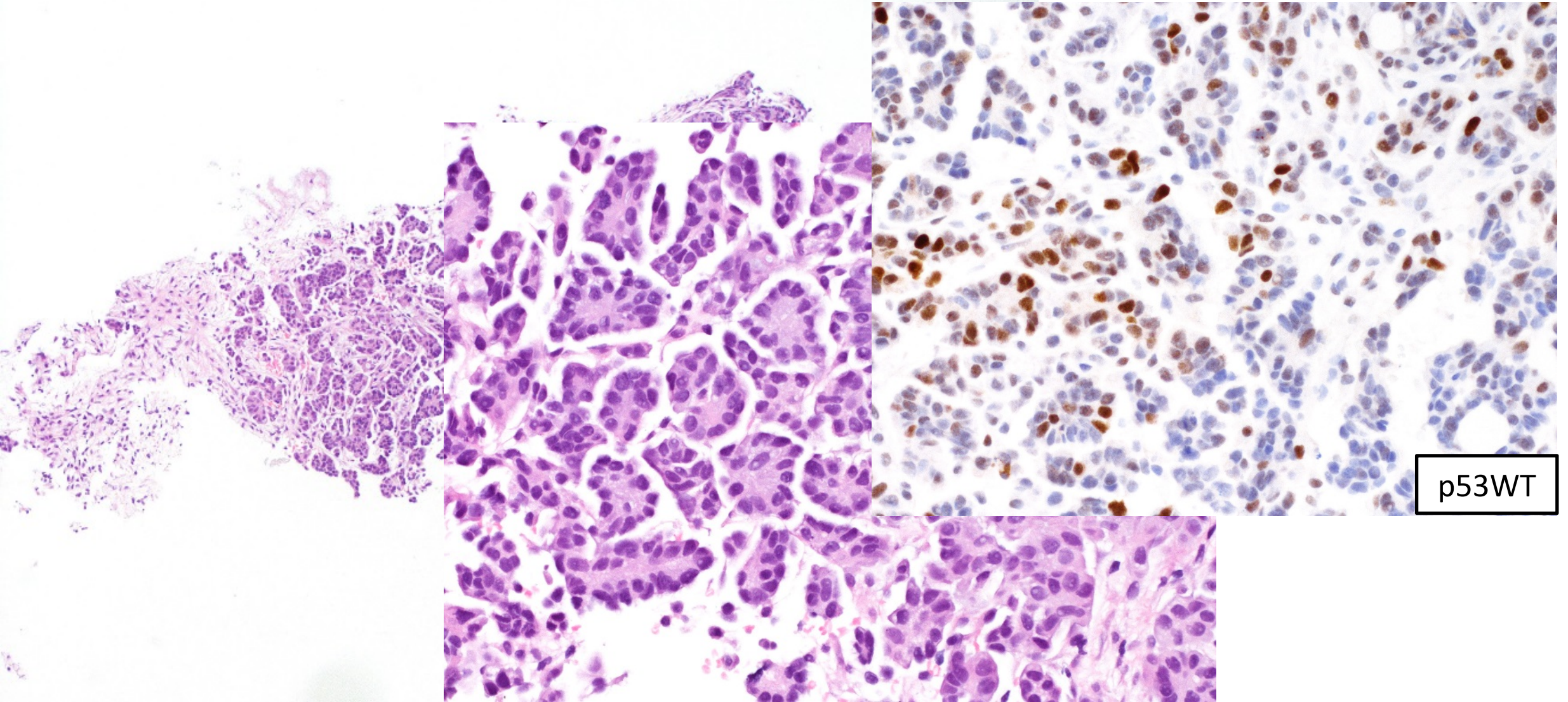
IHC	Nonsynonymous	Indel	Stopgain	Splicing	NDM	Total
OE	115	2	2	2	0	121
CA	0	16	13	12	0	41
CY	0	2	2	0	0	4
WT	0	4	0	3	76	83
Total	115	24	17	17	76	249

Sensitivity 96%
Specificity 100%

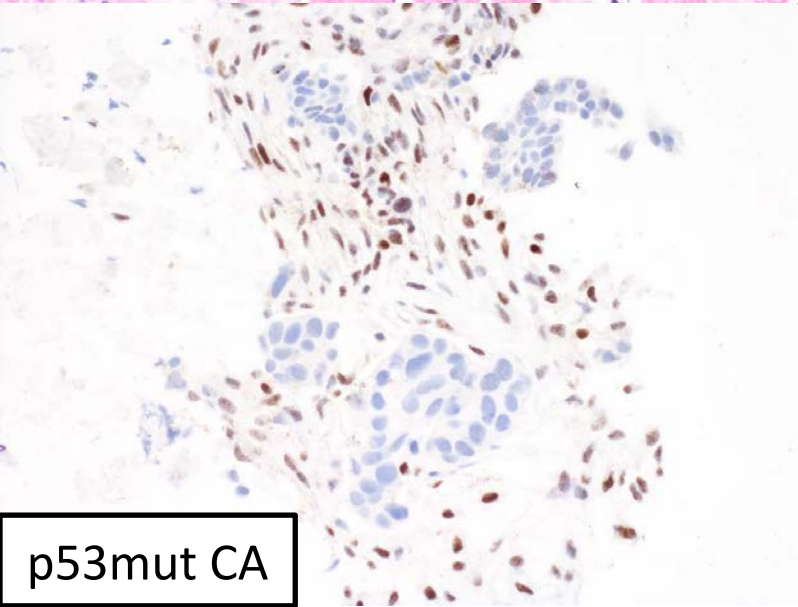
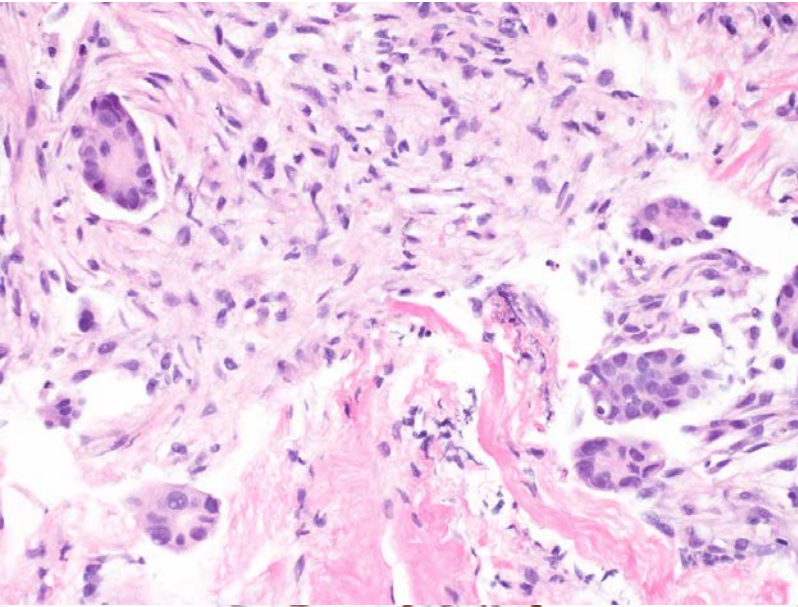
1. Distinction of tubo-ovarian high-grade serous from ovarian/peritoneal low-grade serous carcinoma



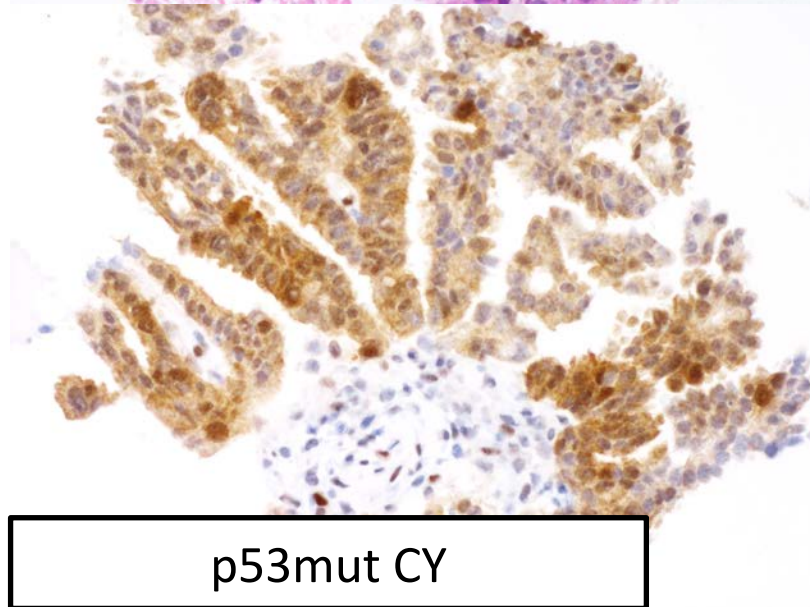
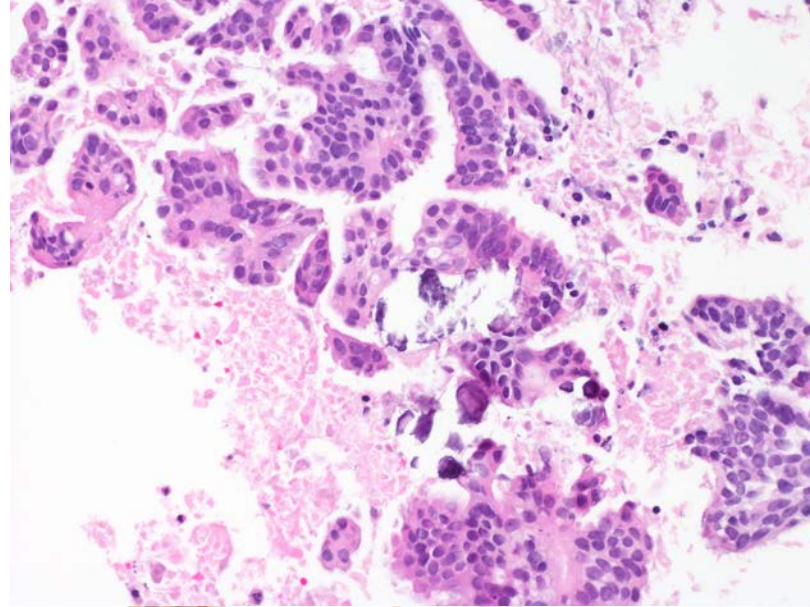
1.a) Confirmation of LGSC



1.b) The danger of underreliance on p53 IHC



p53mut CA

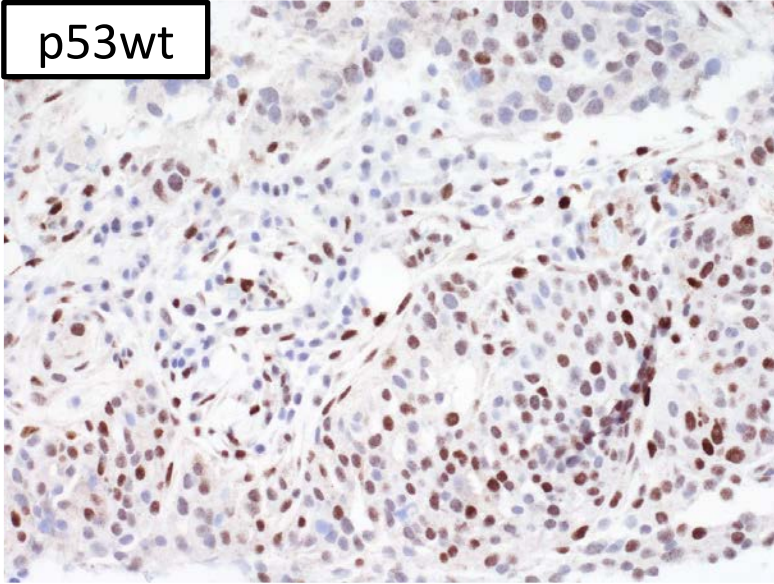


p53mut CY

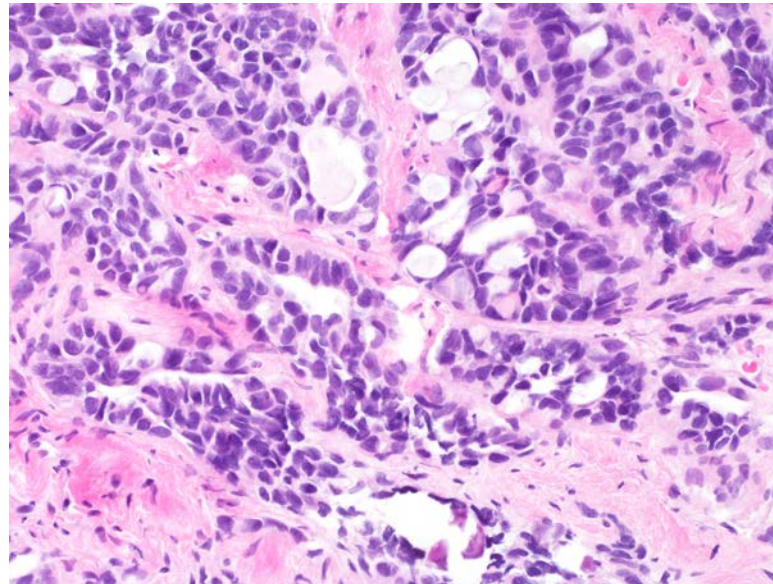
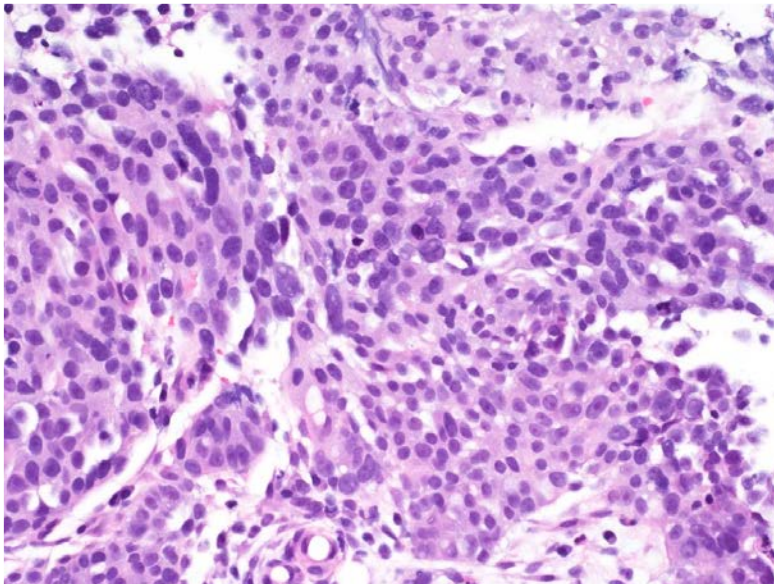
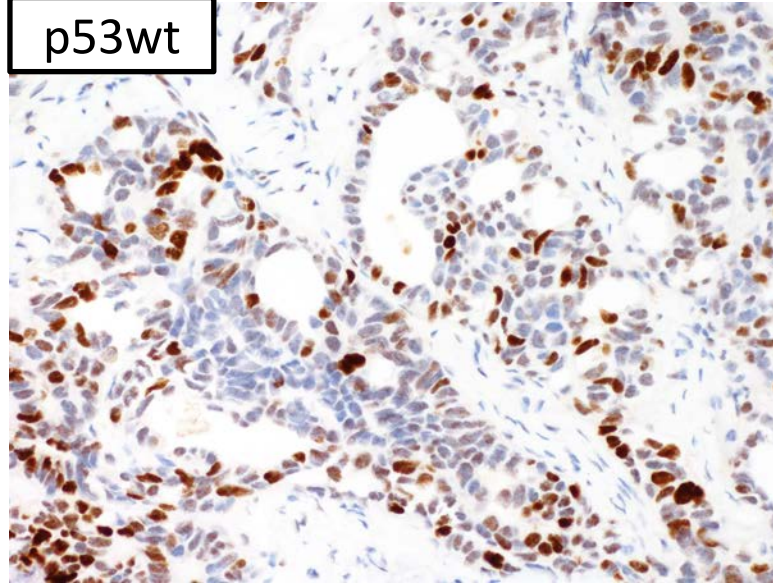
- Some HGSC can mimic LGSC on preoperative biopsy: micropapillary architecture and moderate nuclear atypia
- Confirm with p53 IHC

1.c) Avoid the sensitivity trap

p53wt



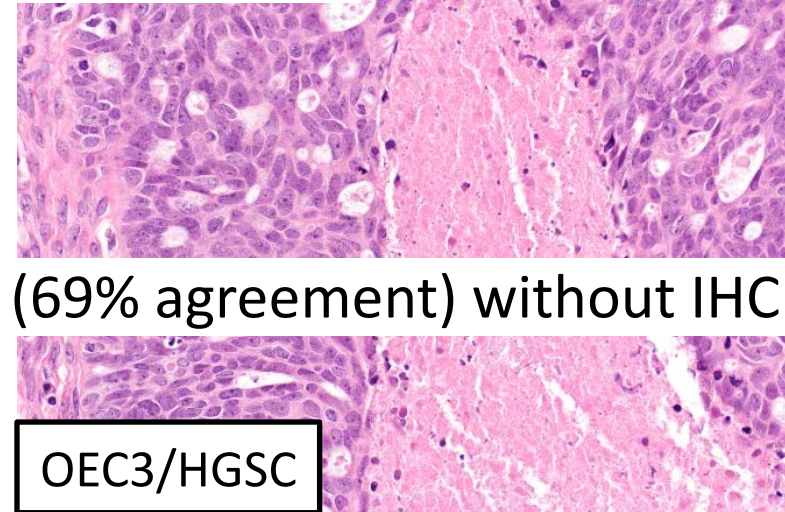
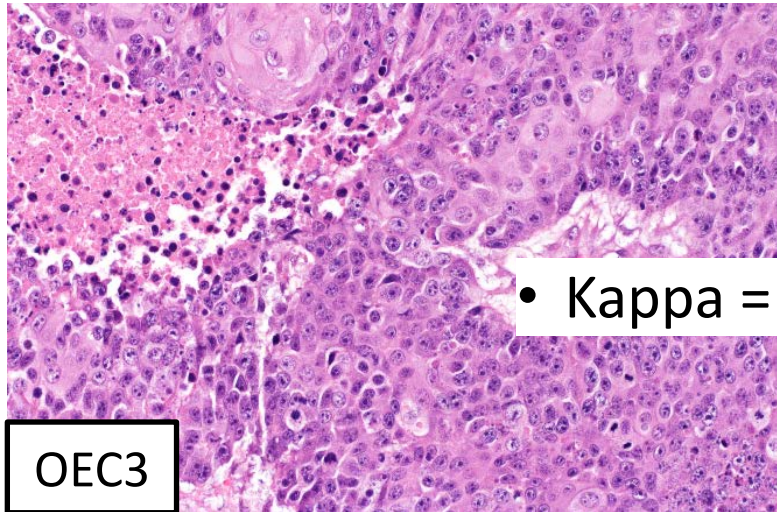
p53wt



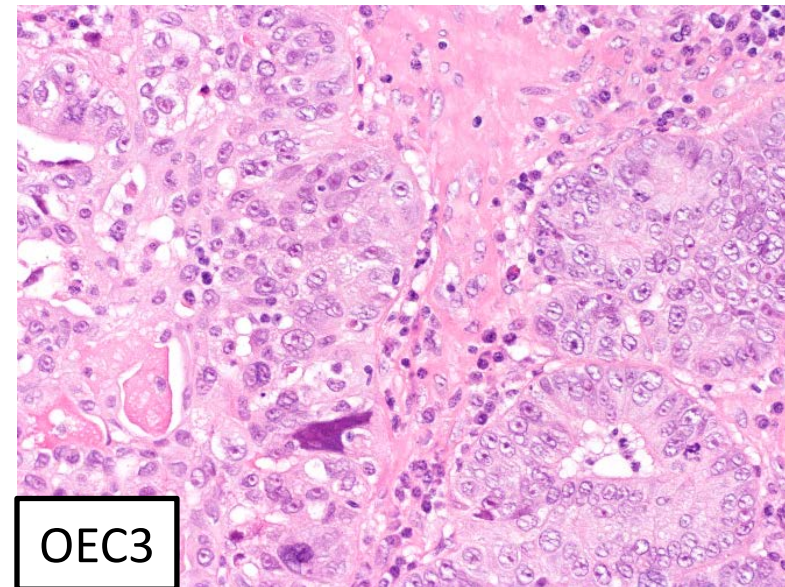
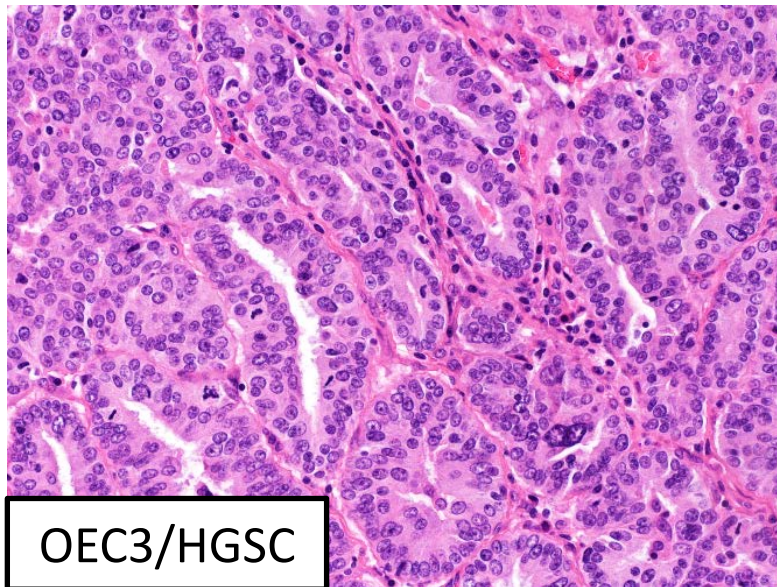
- ~2% of tubo-ovarian carcinoma harbor a *TP53* mutation that results in expression of a non functional protein in wild type pattern.
- These mutation can either be C-terminal truncating or splice site mutation.
- A tumor with p53wt can still be a tubo-ovarian high-grade serous carcinoma – morphological correlation is required.

2. Differential diagnosis of ovarian endometrioid grade 3 versus HGSC

Interobserver Reproducibility between OEC3 and HGSC



- Kappa = 0.40 (69% agreement) without IHC

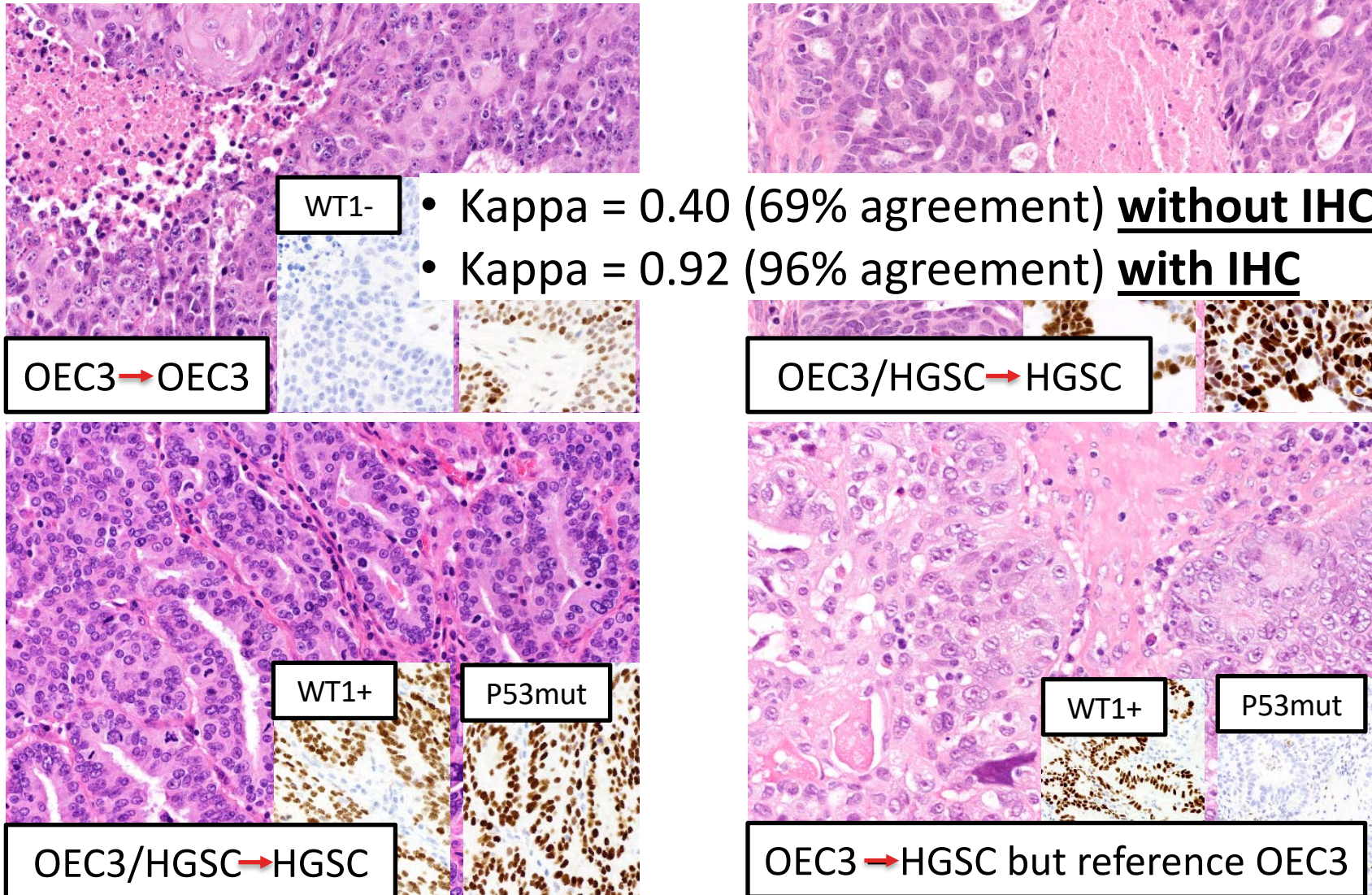


Fist line ancillary markers assessing tubo-ovarian carcinoma

Histotype	PAX8, present	WT1, present	TP53, mutant-type	Napsin A, present	PR, present
HGSC	95%	97%	94-98%	1%	37-42%
LGSC	87-100%	98-100%	<u>0</u>	0	59-60%
EC	<u>82%</u>	10-14%	14-15%	3-8%	81-85%
CCC	95%	<u>1%</u>	11-12%	92%	5-7%
MC	39-47%	<u>0-1%</u>	61-66%	0-3%	<u>0-4%</u>

2. Differential diagnosis of ovarian endometrioid grade 3 versus HGSC

Interobserver Reproducibility between OEC3 and HGSC



The figure displays four panels of histological images comparing Ovarian Endometrioid Grade 3 (OEC3) and High-Grade Serous Carcinoma (HGSC). Each panel includes a main H&E stained image and smaller inset images showing immunohistochemical (IHC) staining for WT1 and P53. The panels are labeled as follows:

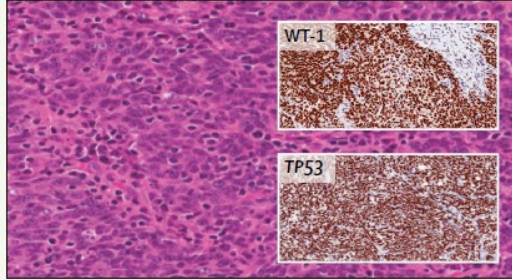
- Top Left:** OEC3 → OEC3. IHC markers: WT1- (negative).
- Top Right:** OEC3/HGSC → HGSC. IHC markers: WT1+ (positive), P53mut (mutated).
- Bottom Left:** OEC3/HGSC → HGSC. IHC markers: WT1+ (positive), P53mut (mutated).
- Bottom Right:** OEC3 → HGSC but reference OEC3. IHC markers: WT1+ (positive), P53mut (mutated).

Summary statistics for interobserver reproducibility:

- Kappa = 0.40 (69% agreement) **without IHC**
- Kappa = 0.92 (96% agreement) **with IHC**

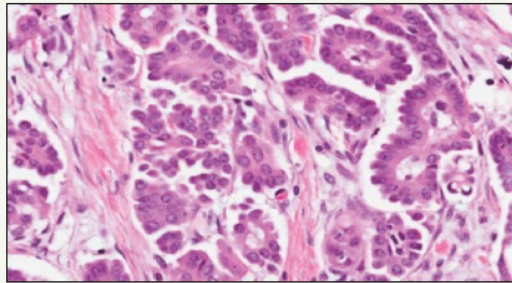
2.b) WT1/p53 to confirm HGSC: rare pitfalls

HGSOc

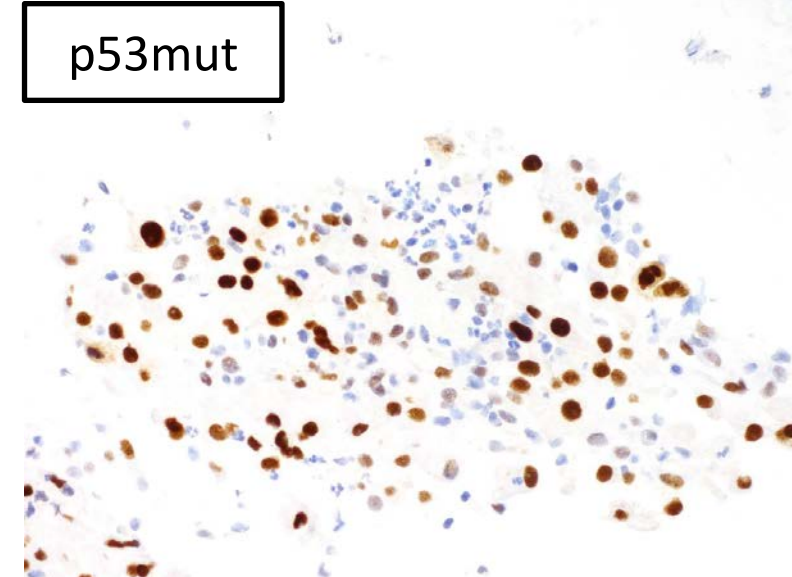
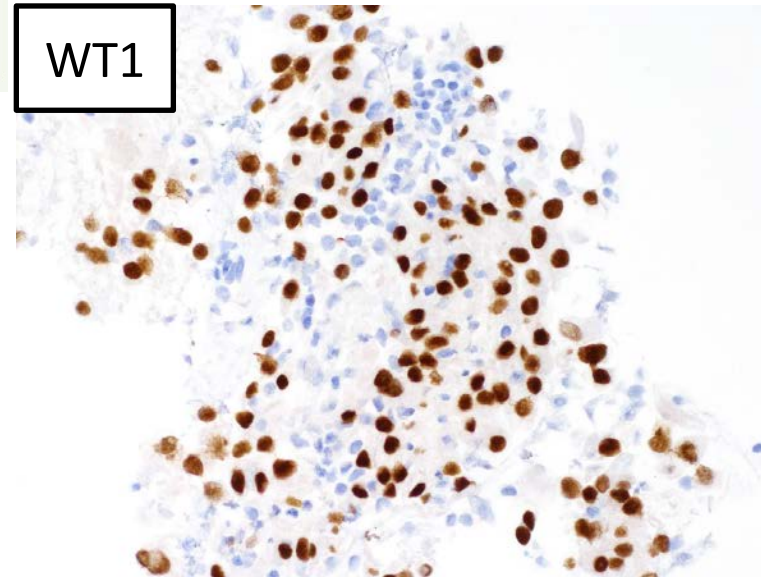
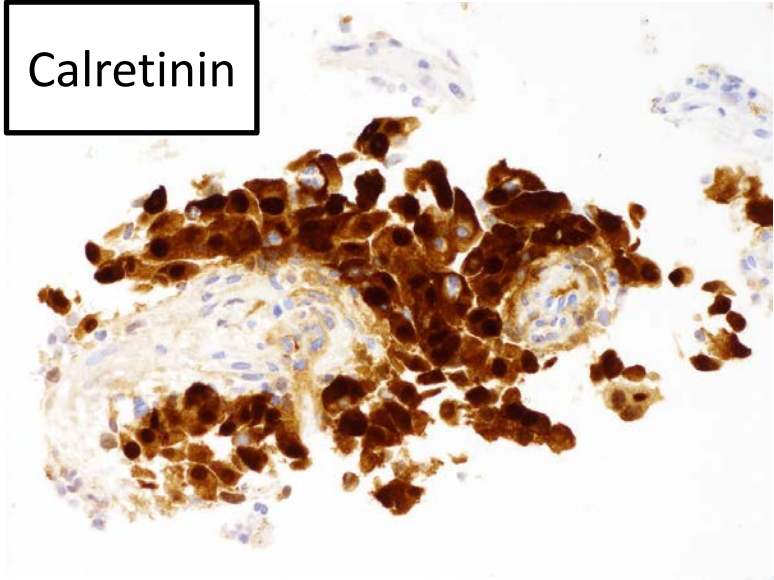
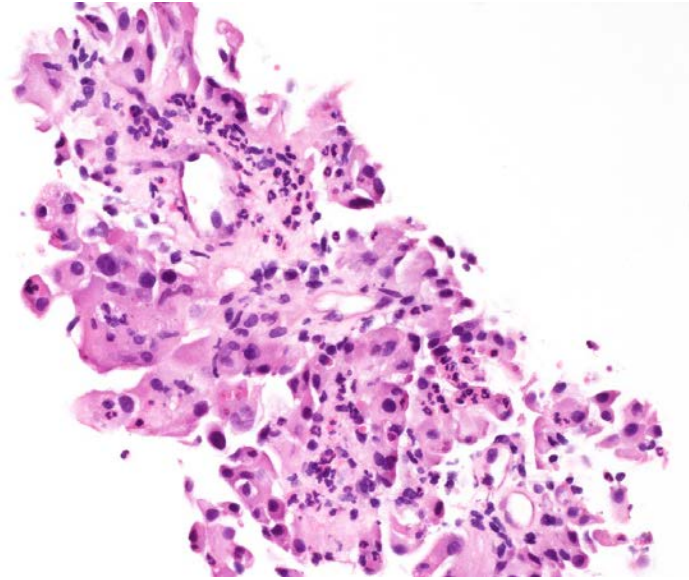


- Highly aggressive tumours
- Papillary or solid growth pattern
- Tumour cells with atypical, large irregular nuclei
- High proliferative rate
- Initial chemosensitivity with subsequent acquisition of increasing resistance
- Key targets: *TP53*, *BRCA1* and 2, and *HRR* genes

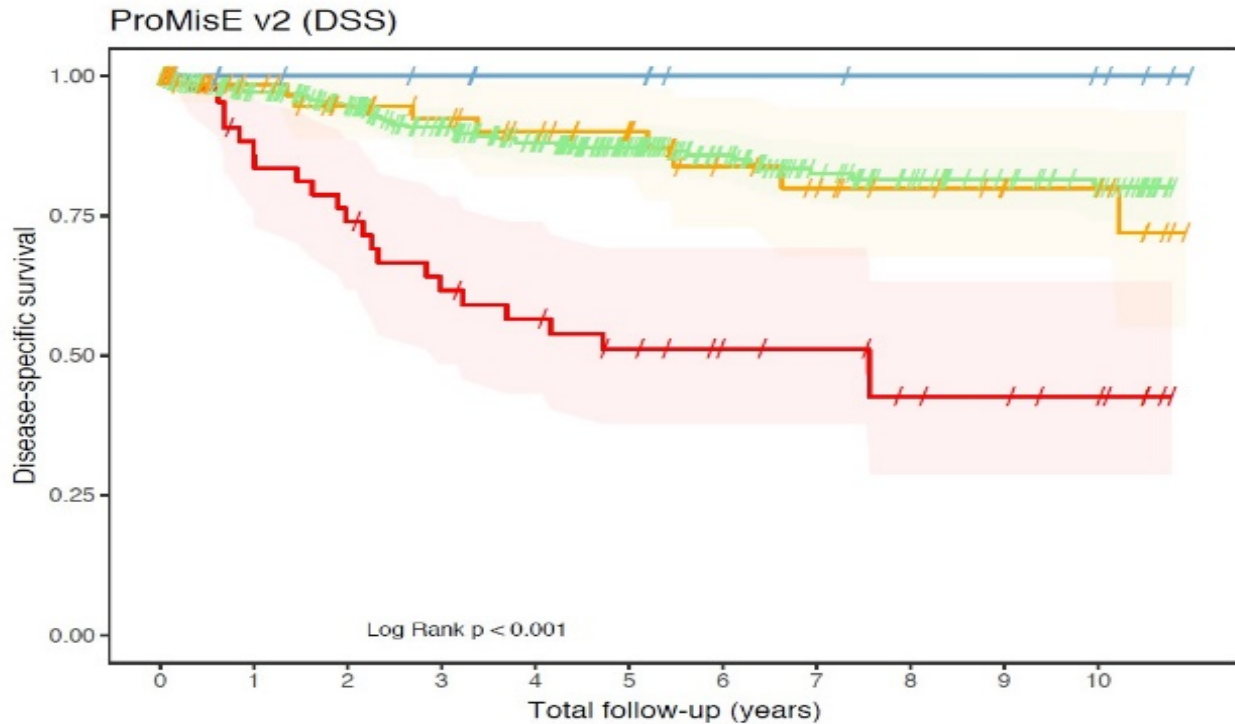
LGSOc



- Indolent behaviour
- Micro-papillary pattern
- Tumour cells with small uniform nuclei
- Low proliferative rate
- Relative chemoresistance
- Key targets: *BRAF*, *KRAS*, *NRAS*, and *PIK3CA*



3. Molecular subtype of ovarian endometrioid carcinoma

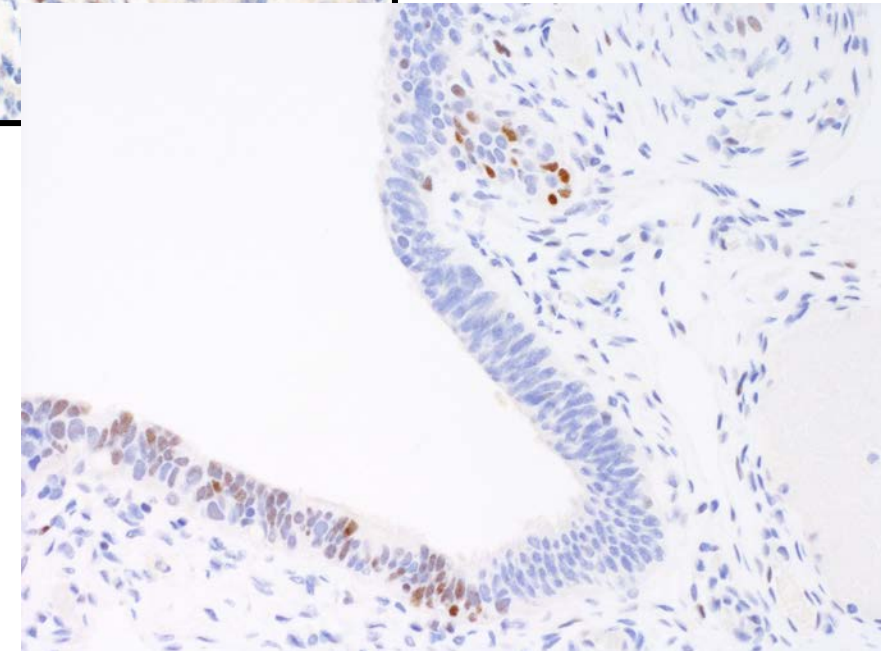
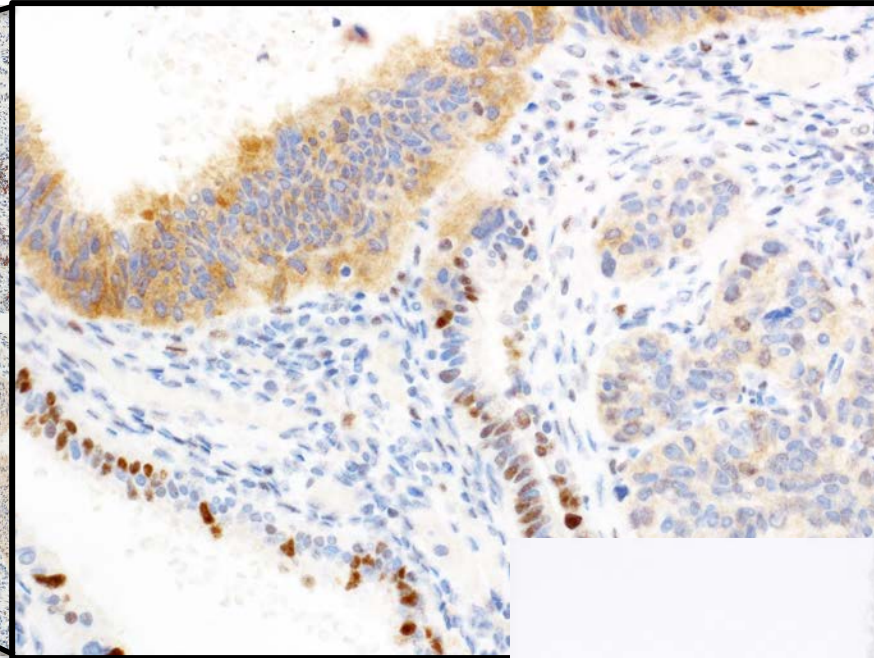
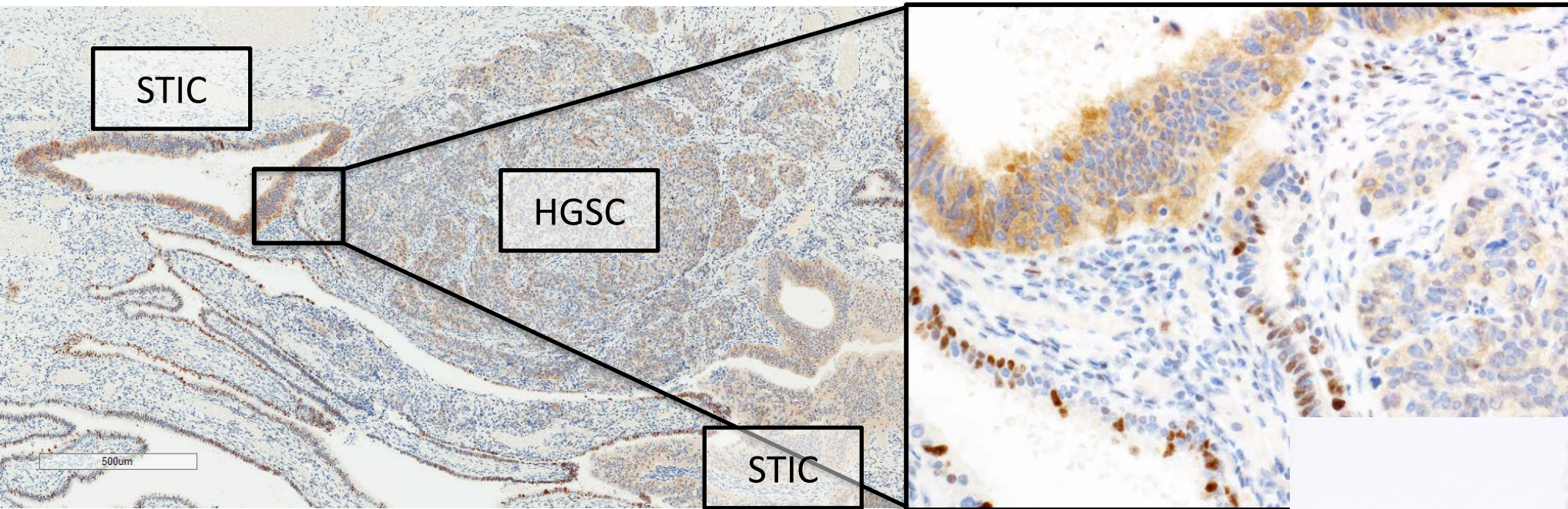


	0	1	2	3	4	5	6	7	8	9	10
MMRd	67	54	46	42	37	31	23	20	16	13	12
POLE	16	13	12	11	9	9	6	6	5	5	4
p53abn	45	36	31	25	22	18	14	13	9	8	6
p53wt	360	310	281	239	207	154	115	90	75	63	54

Numbers at risk

	Ovarian endometrioid carcinoma N=511	Endometrial carcinoma N=920
POLE	17 (3.3%)	84 (9.1%)
MMRd	70 (13.7%)	232 (25.2%)
NSMP	375 (73.4%)	430 (46.7%)
P53 mut	49 (9.6%)	166 (18.0%)

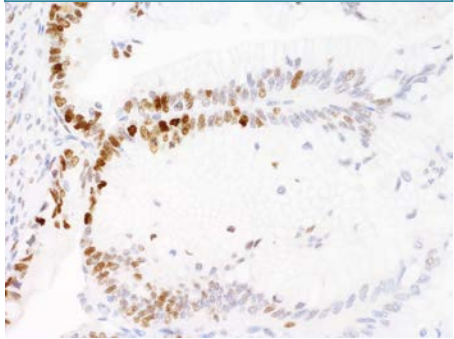
3. Confirming precursor lesion of HGSC



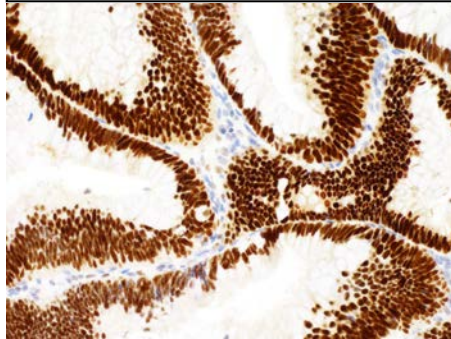
Uncommon patterns can only be appreciated with optimized immunohistochemistry

3. p53 IHC in ovarian mucinous tumors

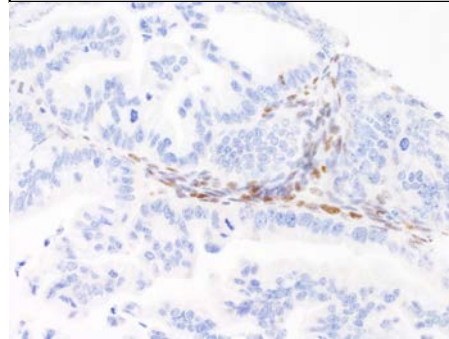
Wild-type pattern



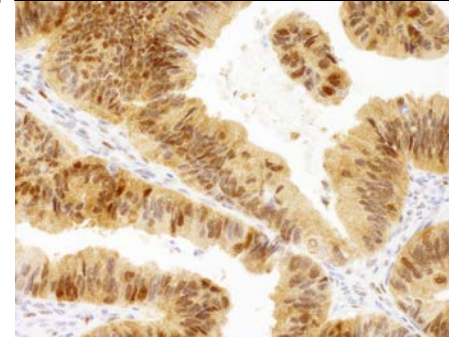
p53mut OE



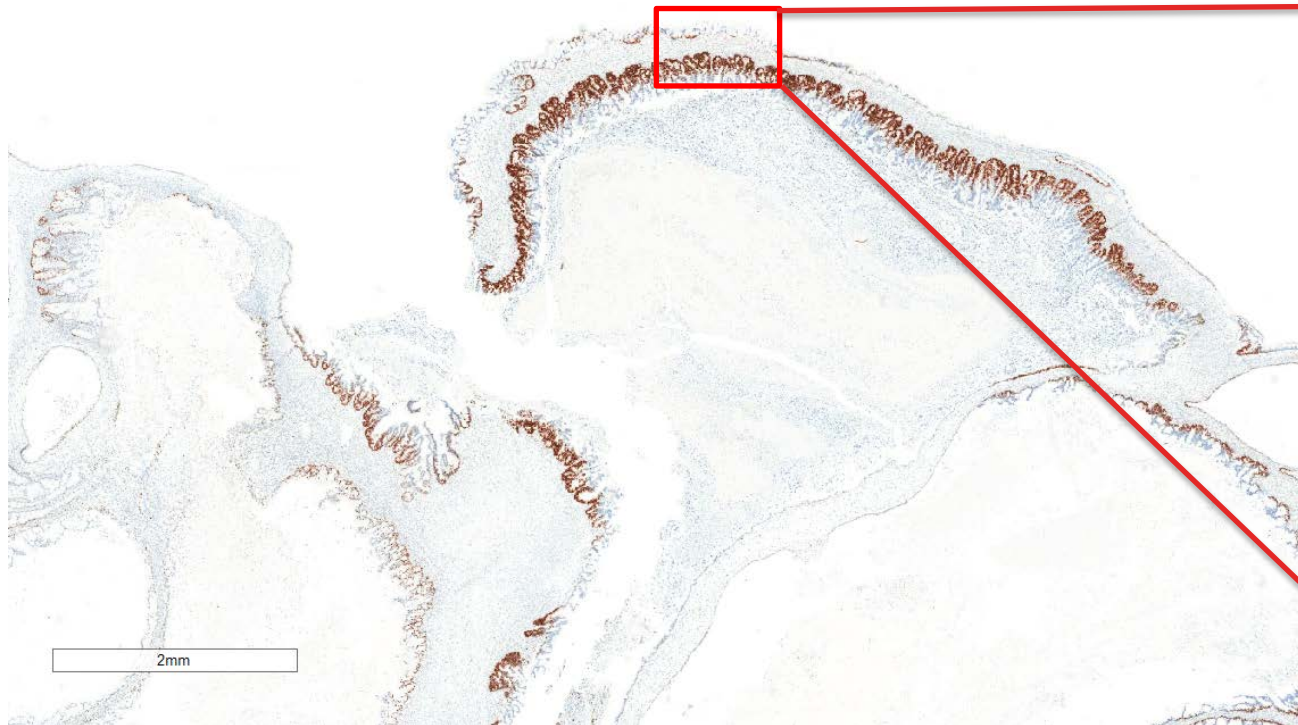
p53mut CA



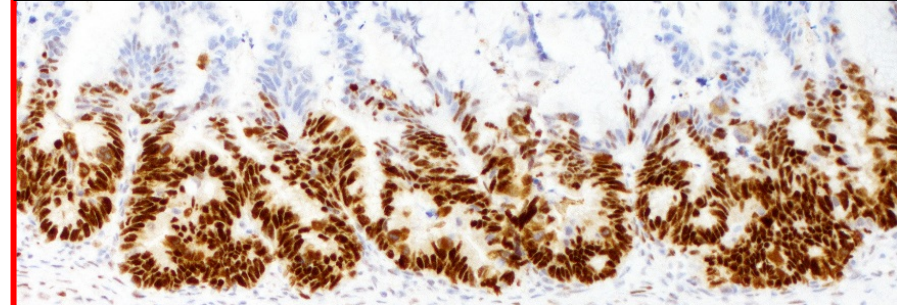
p53mut CY



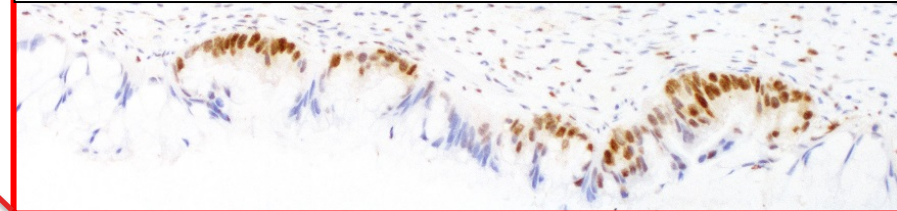
The 4 patterns of p53 IHC also exist in ovarian mucinous tumors but ...



p53mut OE with terminal differentiation

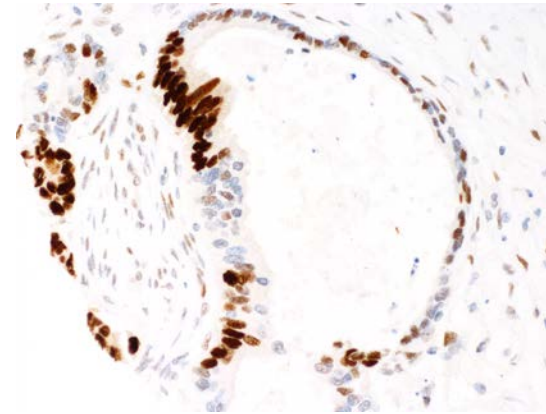
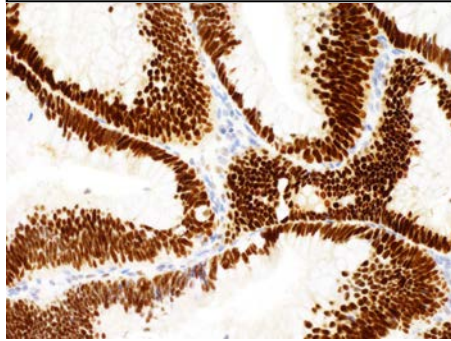


Below wild type = subclonal



3. p53mut OE cut-off in ovarian mucinous tumors

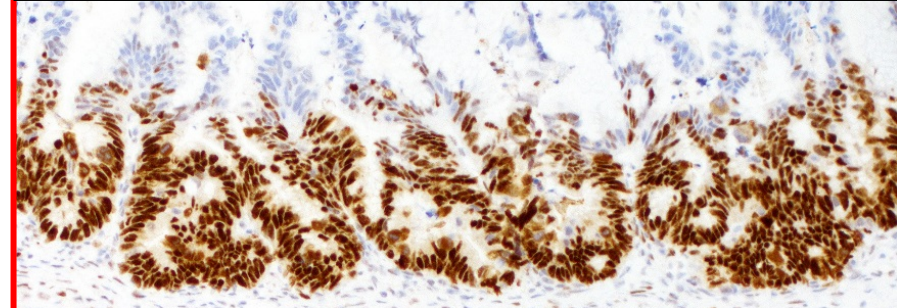
p53mut OE



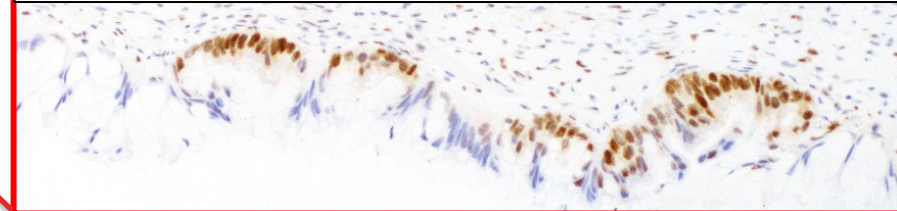
12 consecutive strongly staining cells (“p53 signature”) – lowest threshold for **p53mut OE** in mucinous tumors



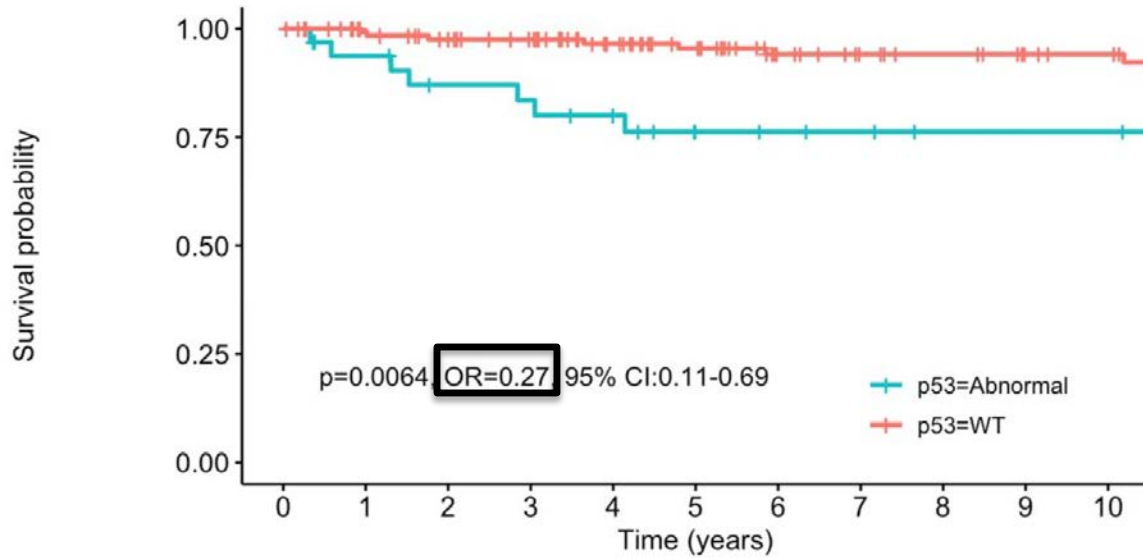
p53mut OE with terminal differentiation



Below wild type = subclonal



Combined MBOT OS

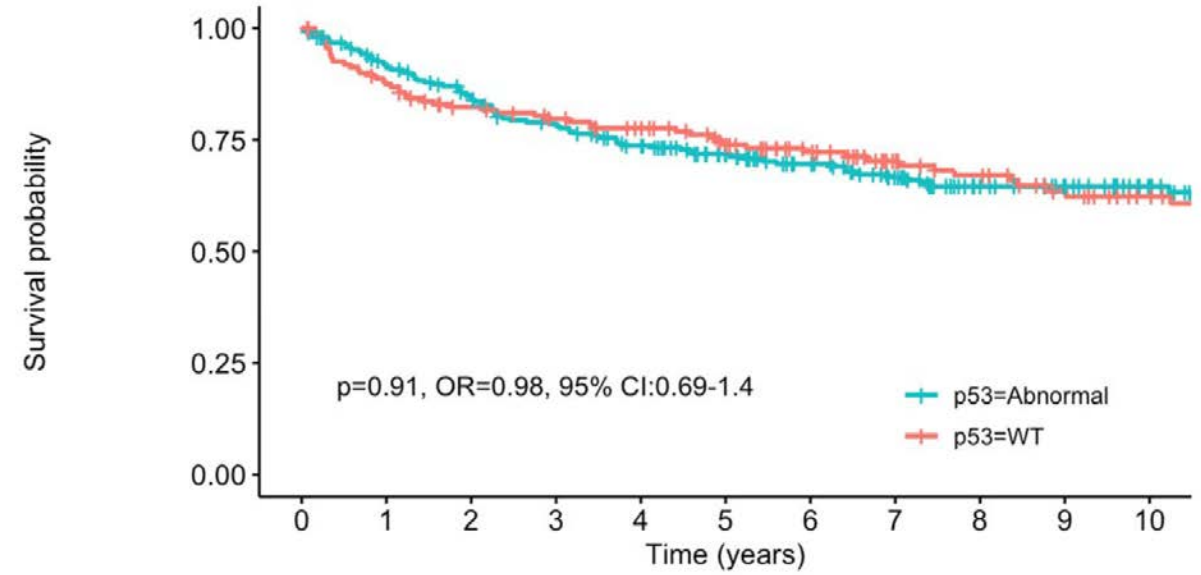


Number at risk

p53=Abnormal	33	29	25	24	22	16	15	14	12	12	12
p53=WT	137	126	119	112	96	84	66	60	57	52	50
	0	1	2	3	4	5	6	7	8	9	10

Time (years)

Combined MOC OS



Number at risk

p53=Abnormal	255	223	199	182	163	140	122	102	79	70	54
p53=WT	162	139	125	117	109	95	82	68	61	51	42
	0	1	2	3	4	5	6	7	8	9	10

Time (years)

4. Endometrial carcinoma histotypes/molecular subtype as per WHO

Histotype/Grade:

1. Endometrial endometrioid carcinoma, grade 1 (50%)
2. Endometrial endometrioid carcinoma, grade 2 (15%)
3. Endometrial endometrioid carcinoma, grade 3 (13%)
4. Dedifferentiated/undifferentiated (from endometrioid, <1%)
5. Endometrial serous carcinoma (10%)
6. Clear cell carcinoma (3%)
7. Carcinosarcoma (7%)
8. Neuroendocrine carcinoma (rare)

Molecular subtype:

1. NSMP (32-59%)
2. MMRd (26-33%)
3. POLEmut (6-13%)
4. p53mut (9-22%)

Journal of Pathology

J Pathol 2018; **245**: 249–250

Published online 16 April 2018 in Wiley Online Library
(wileyonlinelibrary.com) DOI: 10.1002/path.5068

Letter in response to: McAlpine J, Leon-Castillo A, Bosse T.
The rise of a novel classification system for endometrial
carcinoma; integration of molecular subclasses. *J Pathol* 2018;
244: 538–549

Received 16 February 2018; Accepted 1 March 2018

**Implementation or integration of molecular
subclasses?**

LETTER TO THE EDITOR

We have difficulty supporting their proposal because we don't interpret the cited evidence in the same way. *Kappa* values of moderate agreement strength are quoted



Histotype/Grade:

1. Endometrial endometrioid carcinoma, grade 1 (0%)
2. Endometrial endometrioid carcinoma, grade 2 (low)
3. Endometrial endometrioid carcinoma, grade 3 (25%)
4. Dedifferentiated/undifferentiated (from endometrioid, 0%)

5. Endometrial serous carcinoma (100%)
6. Clear cell carcinoma (59%)
7. Carcinosarcoma (90%)
8. Neuroendocrine carcinoma (Common)

Molecular subtype:

1. NSMP (32-59%)
2. MMRd (26-33%)
3. POLEmut (6-13%)
4. p53mut (9-22%)

Histotype/Grade:

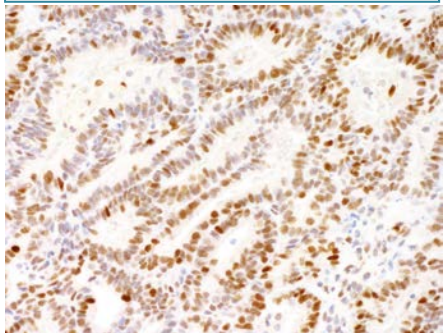
1. Endometrial endometrioid carcinoma, grade 1 (25%)
2. Endometrial endometrioid carcinoma, grade 2 (25%)
3. Endometrial endometrioid carcinoma, grade 3 (45%)
4. Dedifferentiated/undifferentiated (from endometrioid, 53%)
5. Endometrial serous carcinoma (0%)
6. Clear cell carcinoma (0%)
7. Carcinosarcoma (from endometrioid, 4%)
8. Neuroendocrine carcinoma (44%)

Molecular subtype:

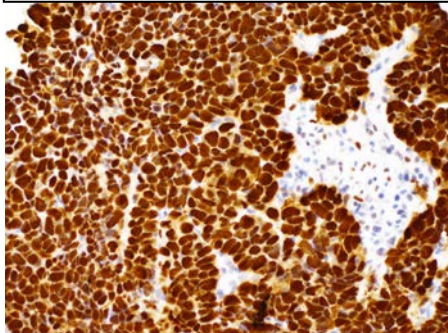
1. NSMP (32-59%)
2. MMRd (26-33%)
3. POLEmut (6-13%)
4. p53mut (9-22%)

Endometrial carcinoma: high interassay & observer IHC agreement

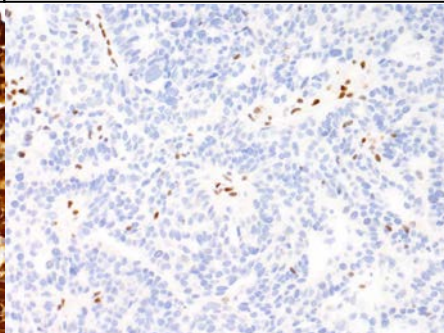
Wild-type pattern



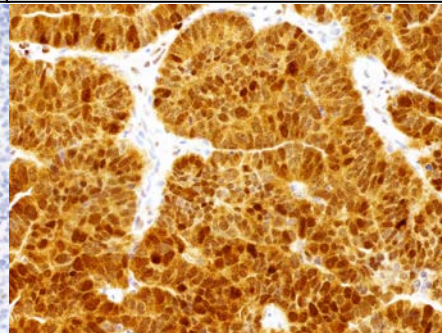
Mut-OE



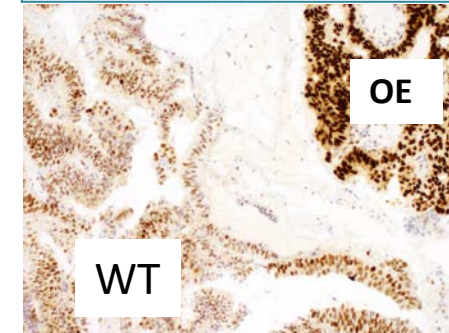
Mut-CA



Mut-CY



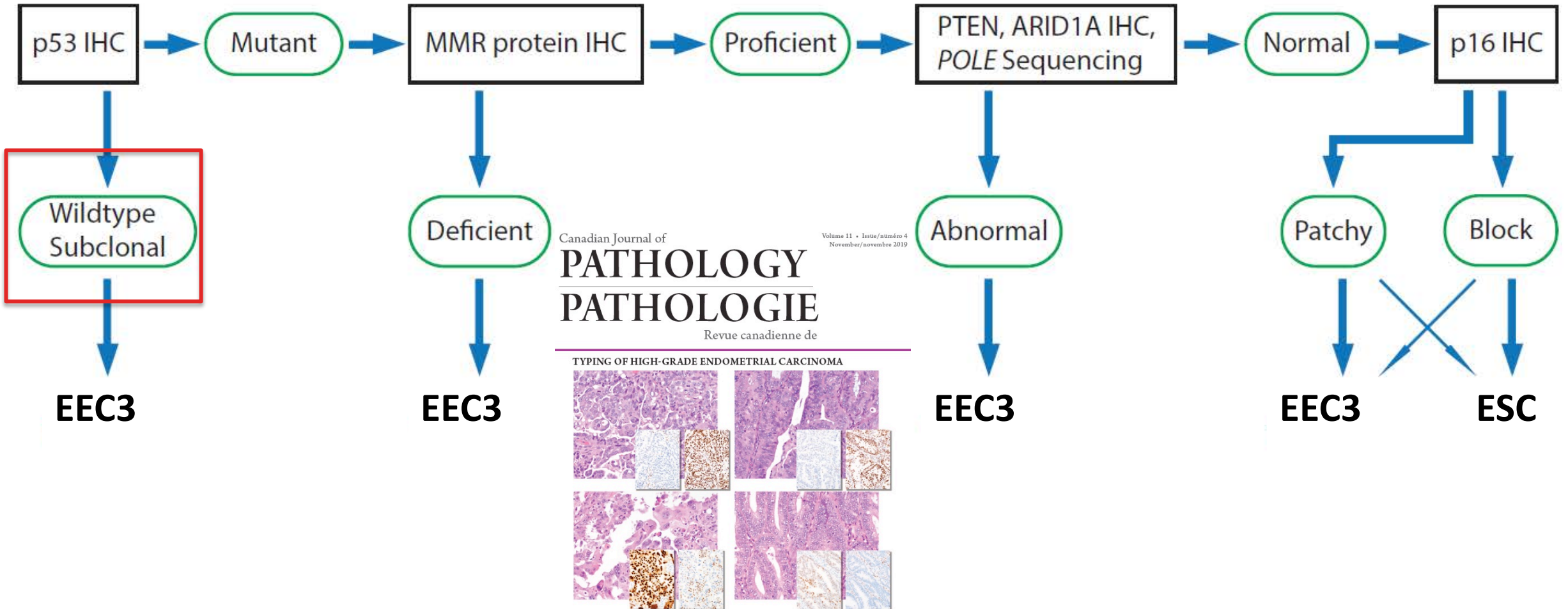
Subclonal

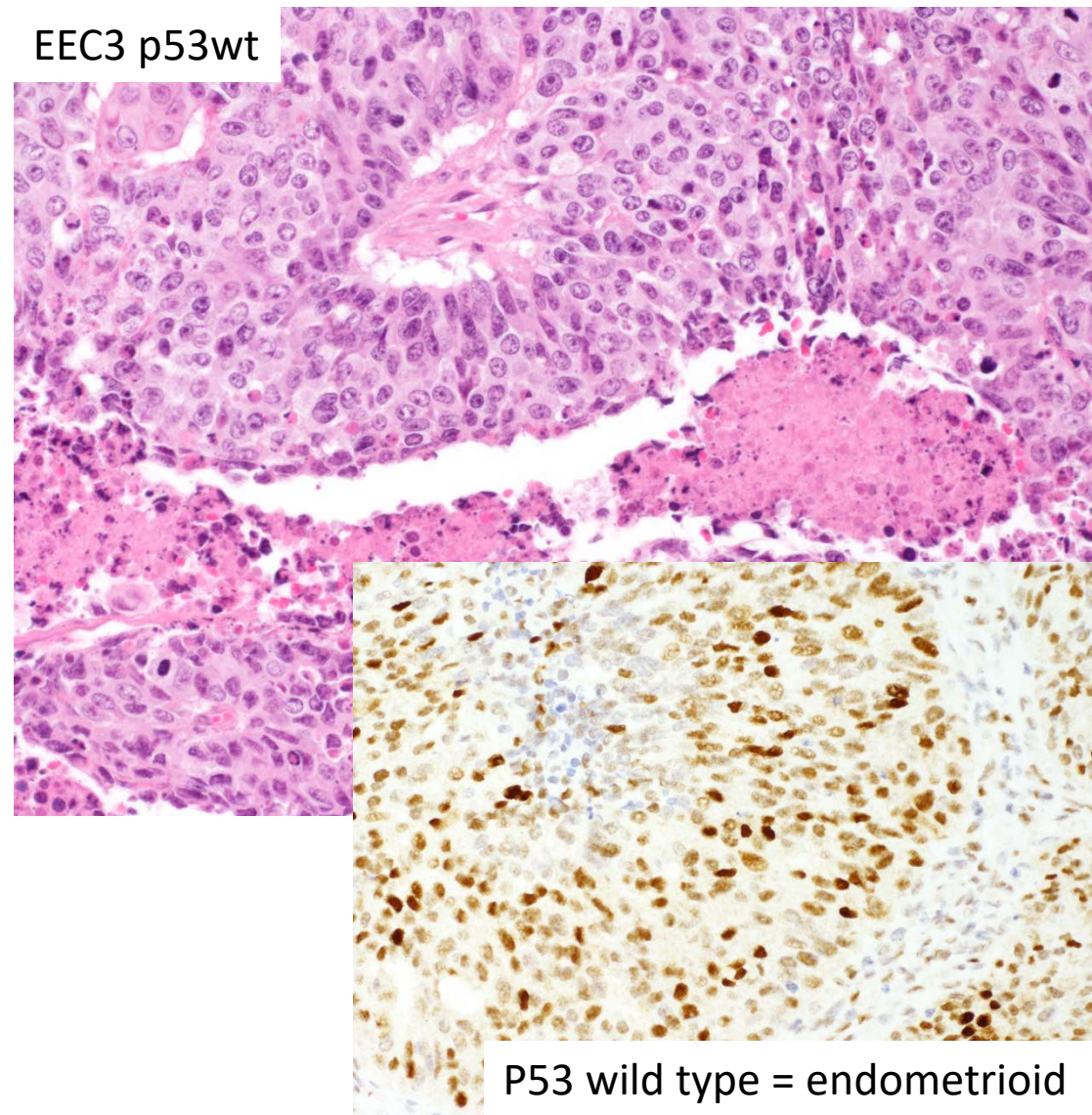
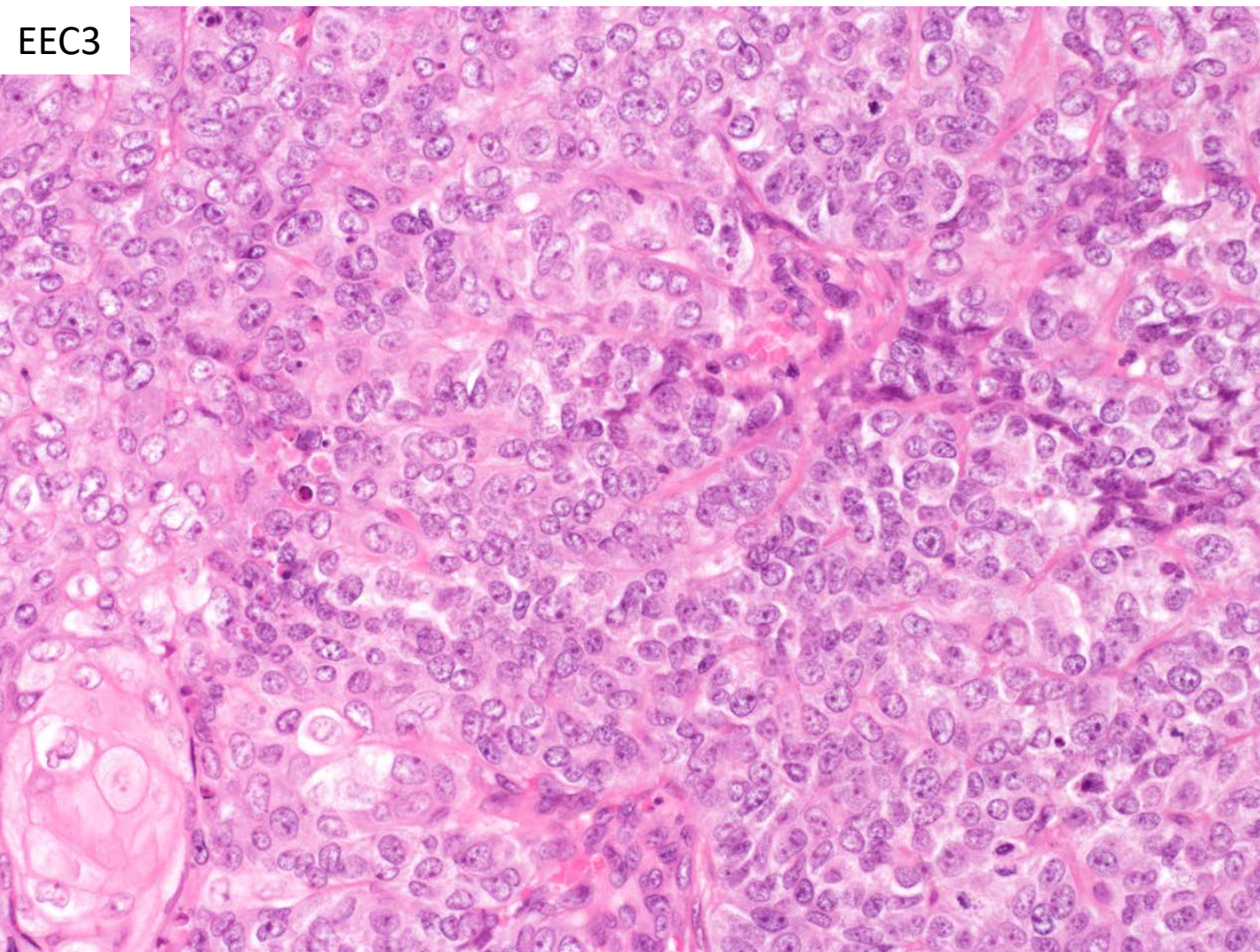


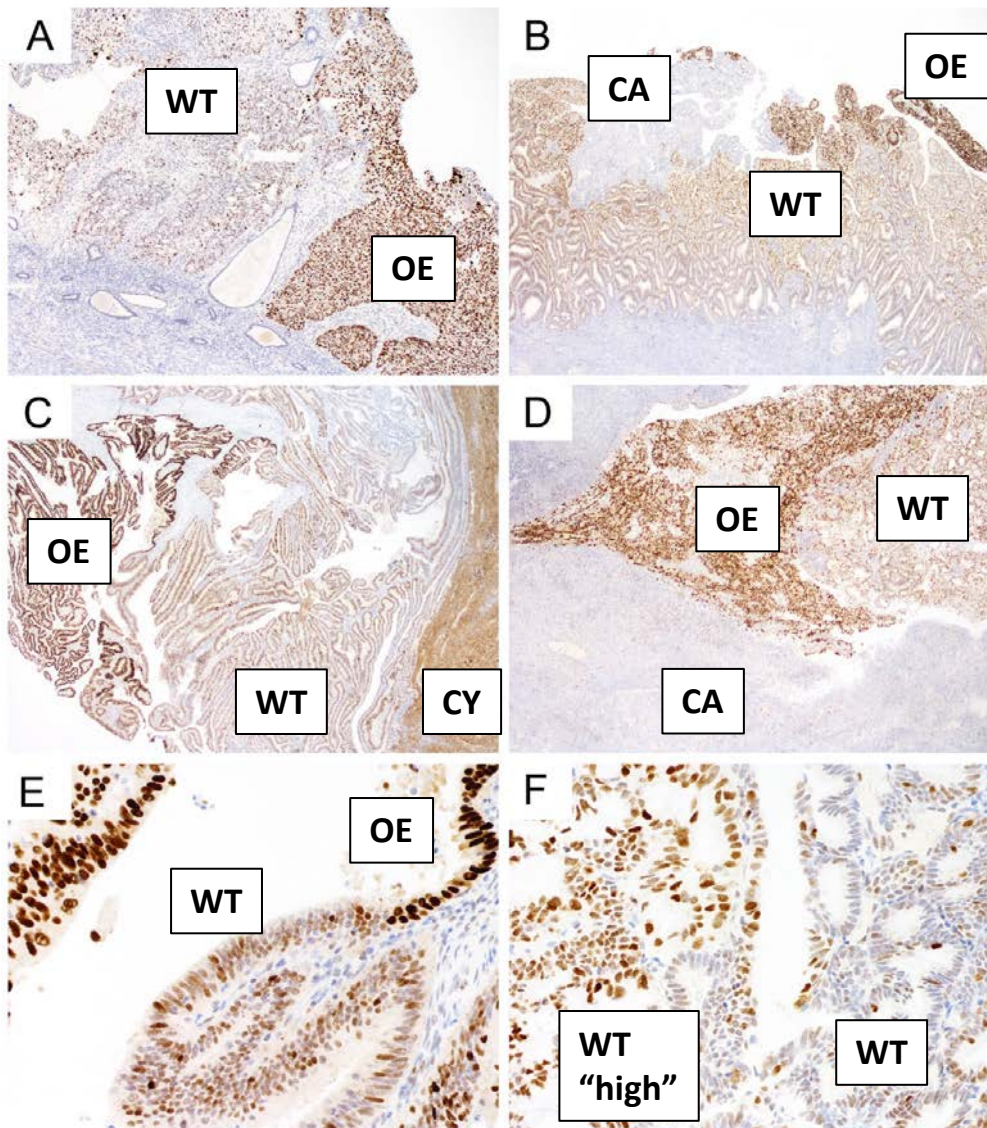
		CENTRAL p53 IHC					
		WT	Mut-OE	Mut-CA	Mut-CY	Subclonal	TOTAL
LOCAL p53 IHC	WT	67	1	0	1	3	72
	Mut-OE	2	68	1	1	0	72
	Mut-CA	1	0	14	1	0	16
	Mut-CY	0	0	0	1	0	1
	Subclonal	0	0	0	0	3	3
TOTAL		70	69	15	4	6	164

		TP53 mutation	
		Present	Absent
IHC	Mut	85	4
	WT	2	32
Sensitivity: 97.70% (95% CI 91.94% to 99.7%)			
Specificity: 88.89% (95% CI 73.94% to 96.89%)			
Accuracy*: 95.12% (95% CI 89.68% to 98.19%)			

4. Histotype distinction of endometrial endometrioid carcinoma grade 3 (EEC3) from serous (ESC)







P53 subclonal

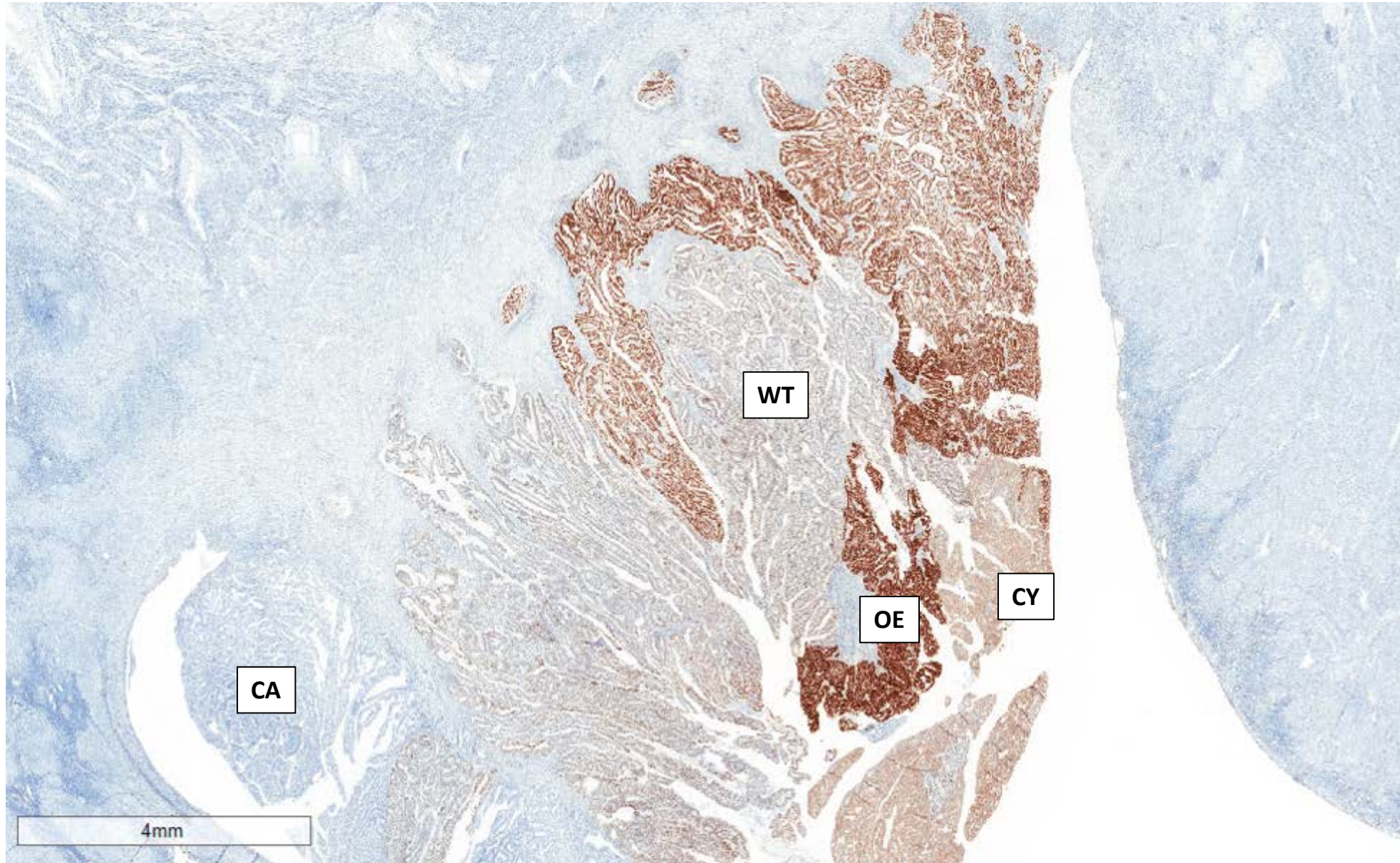
Definition: combination of wild type pattern (WT) with mutant patterns (OE, CA, CY)

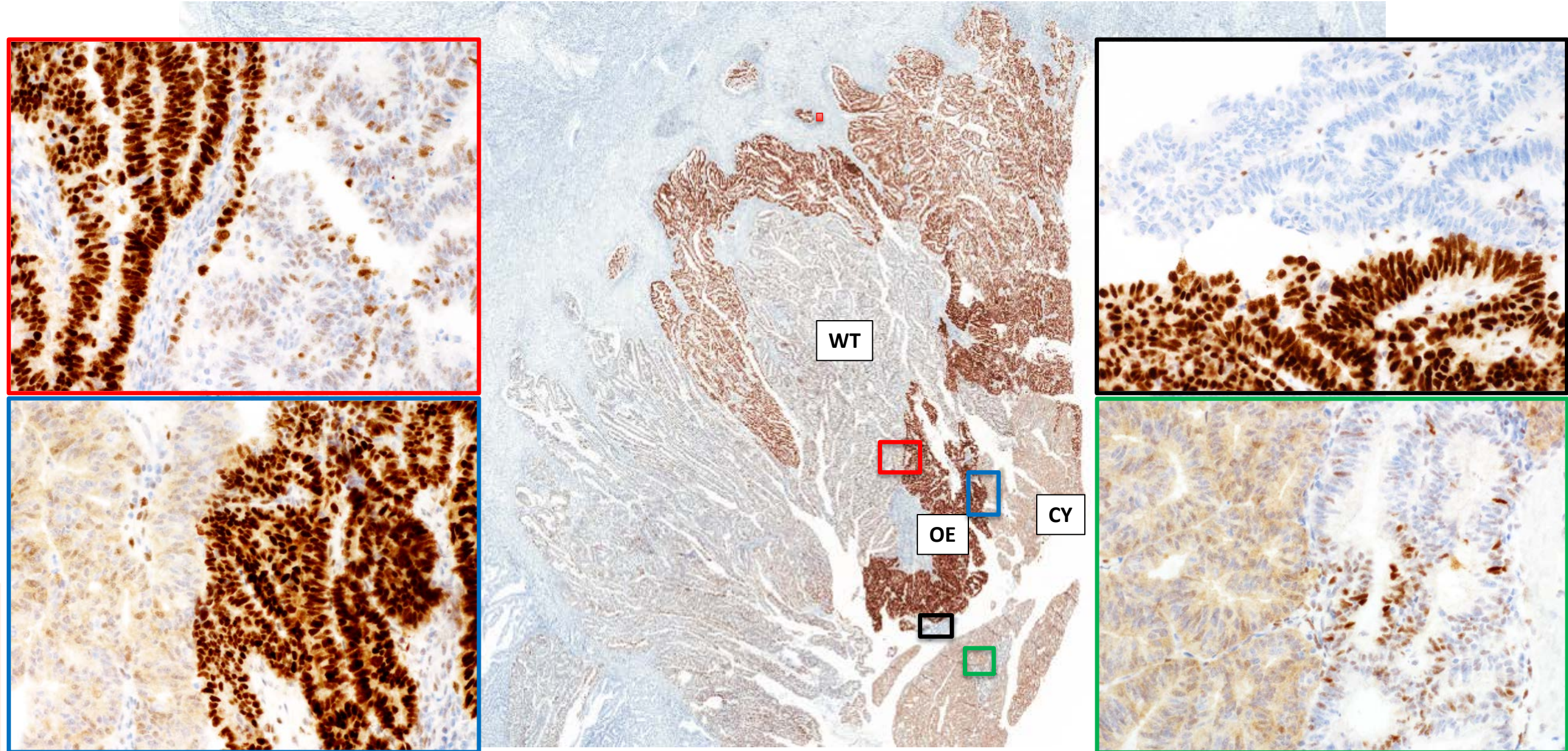
Frequency: ~5%

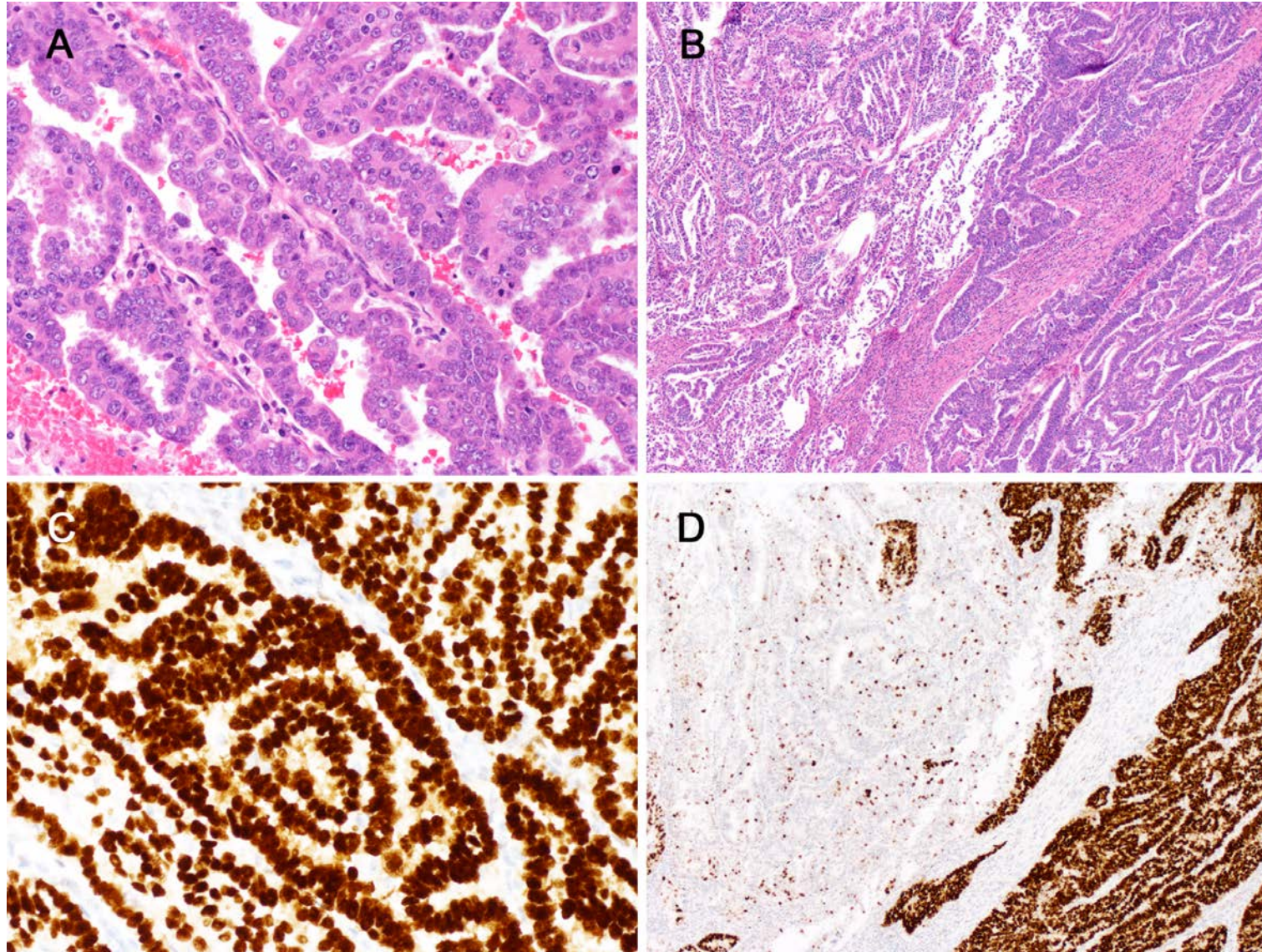
Interpretation:

Confirms endometrioid histotype

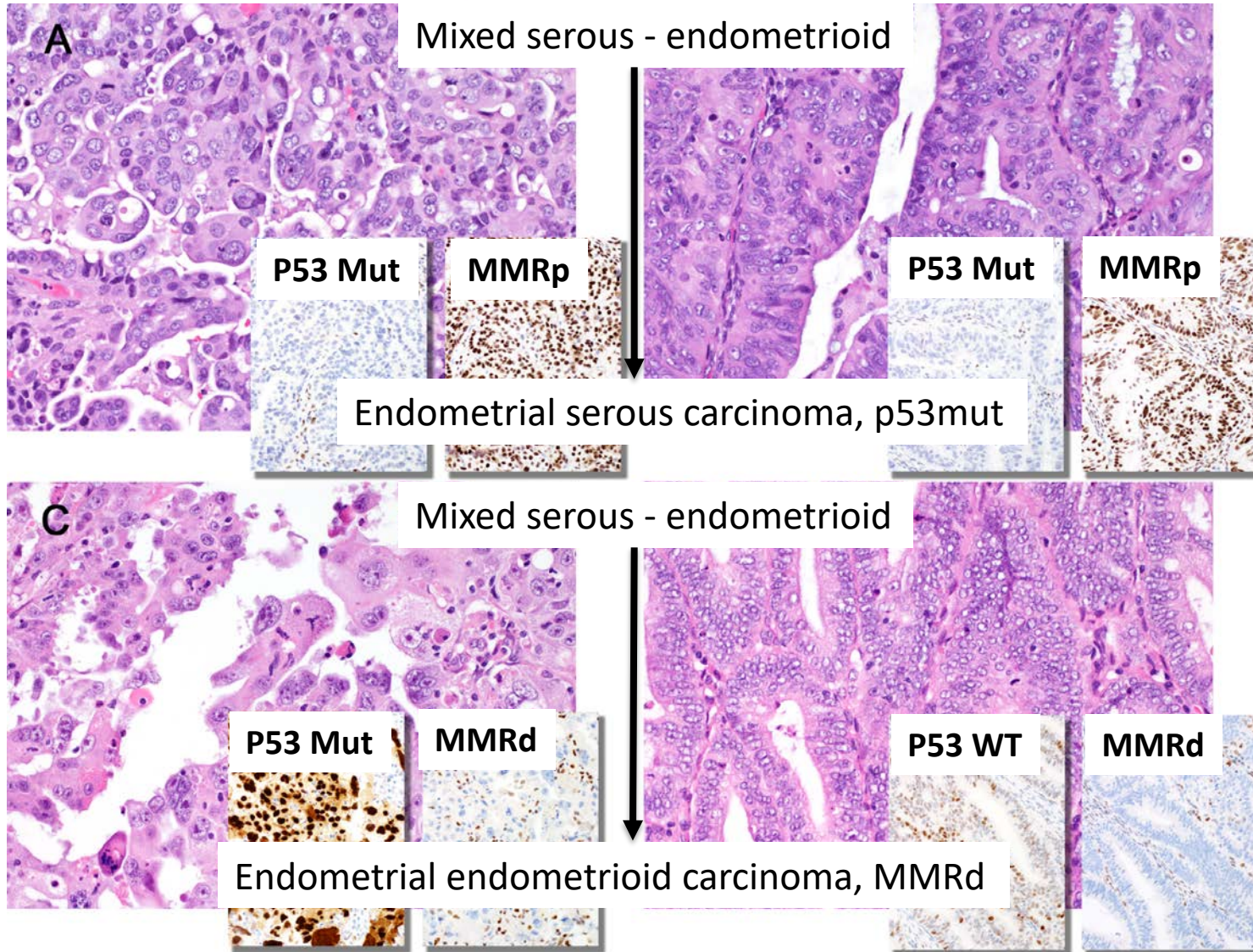
Majority associated with MMRd or POLEmut







4. Subclonal ~ mixed morphology



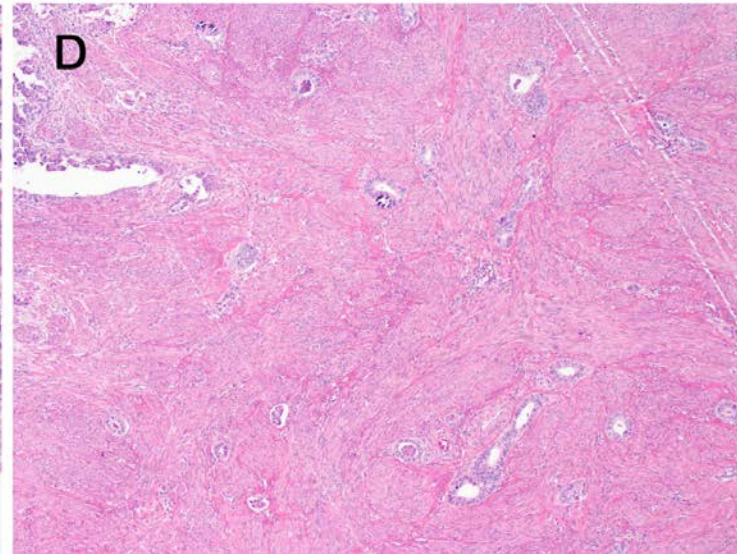
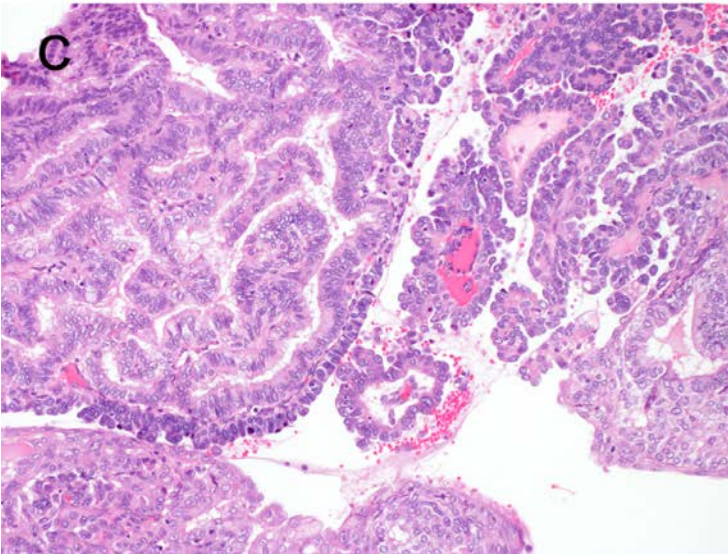
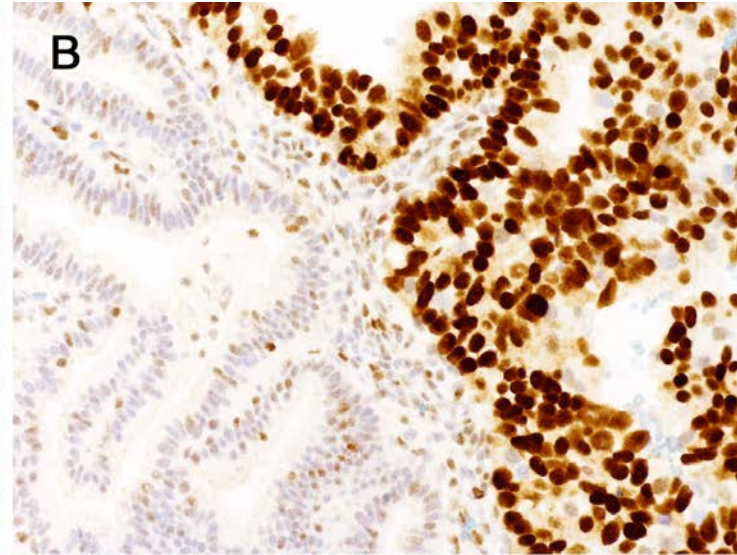
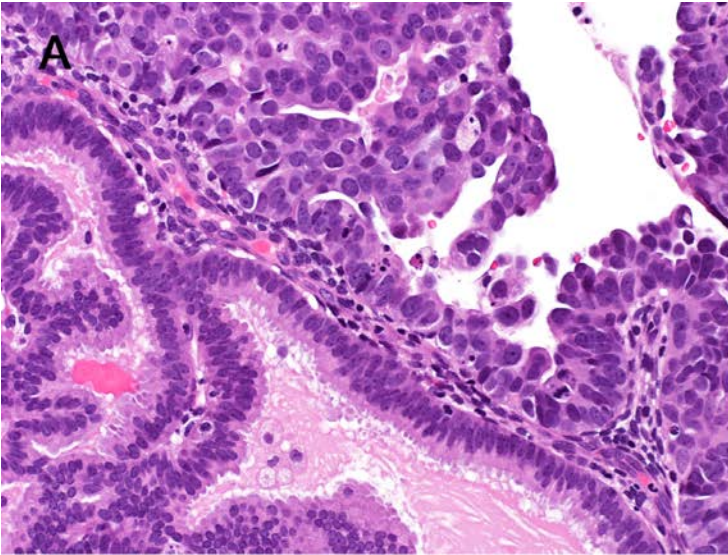
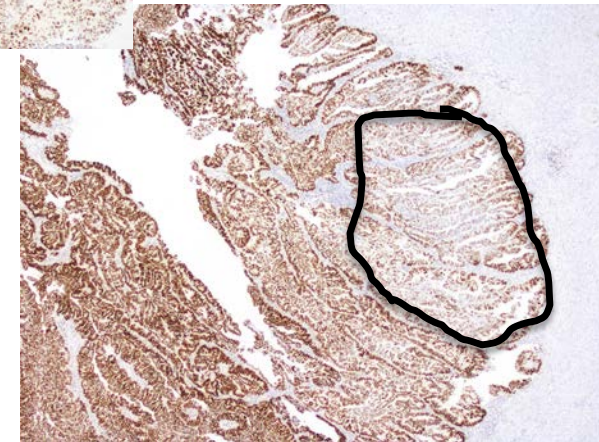
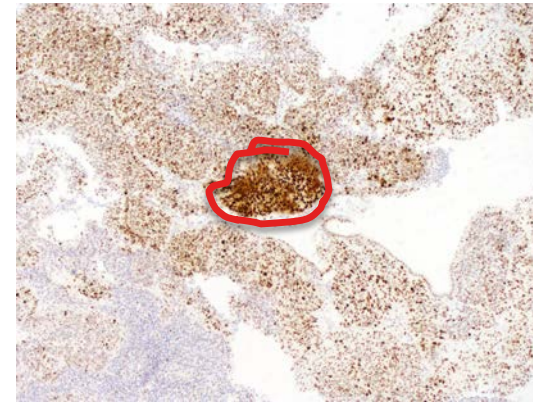
Mixed endometrial carcinoma can be assigned to a single histotype

Represent intratumoral heterogeneity (morphological mimicry) probably due to acquisition of non-founder ('passenger?') mutations

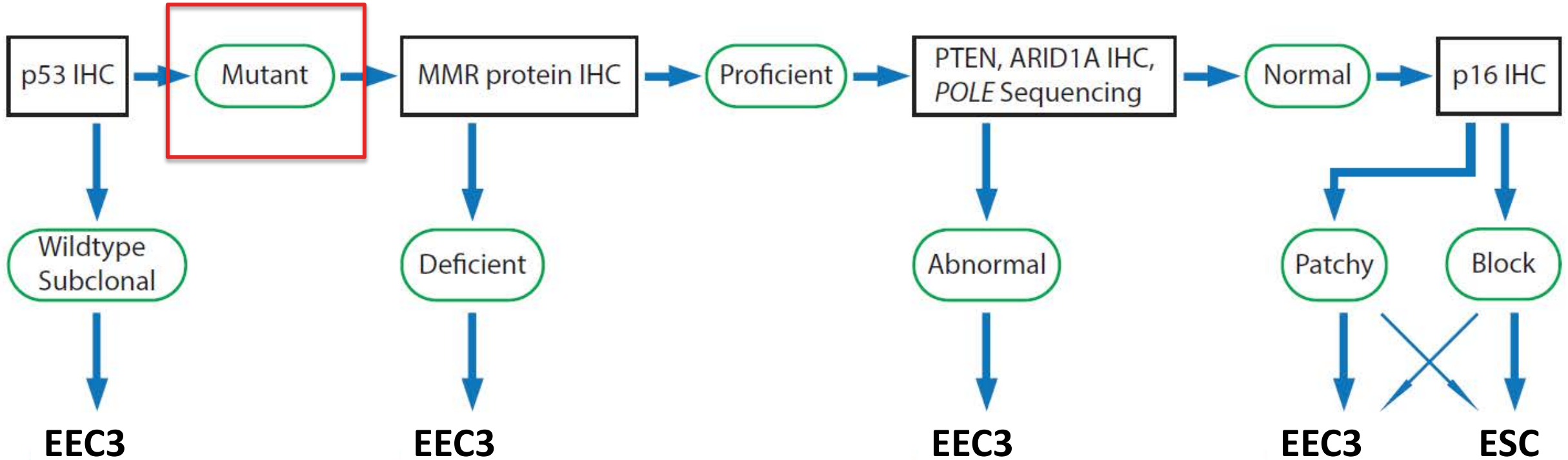
4. Subclonal MMRp, no POLE mutation

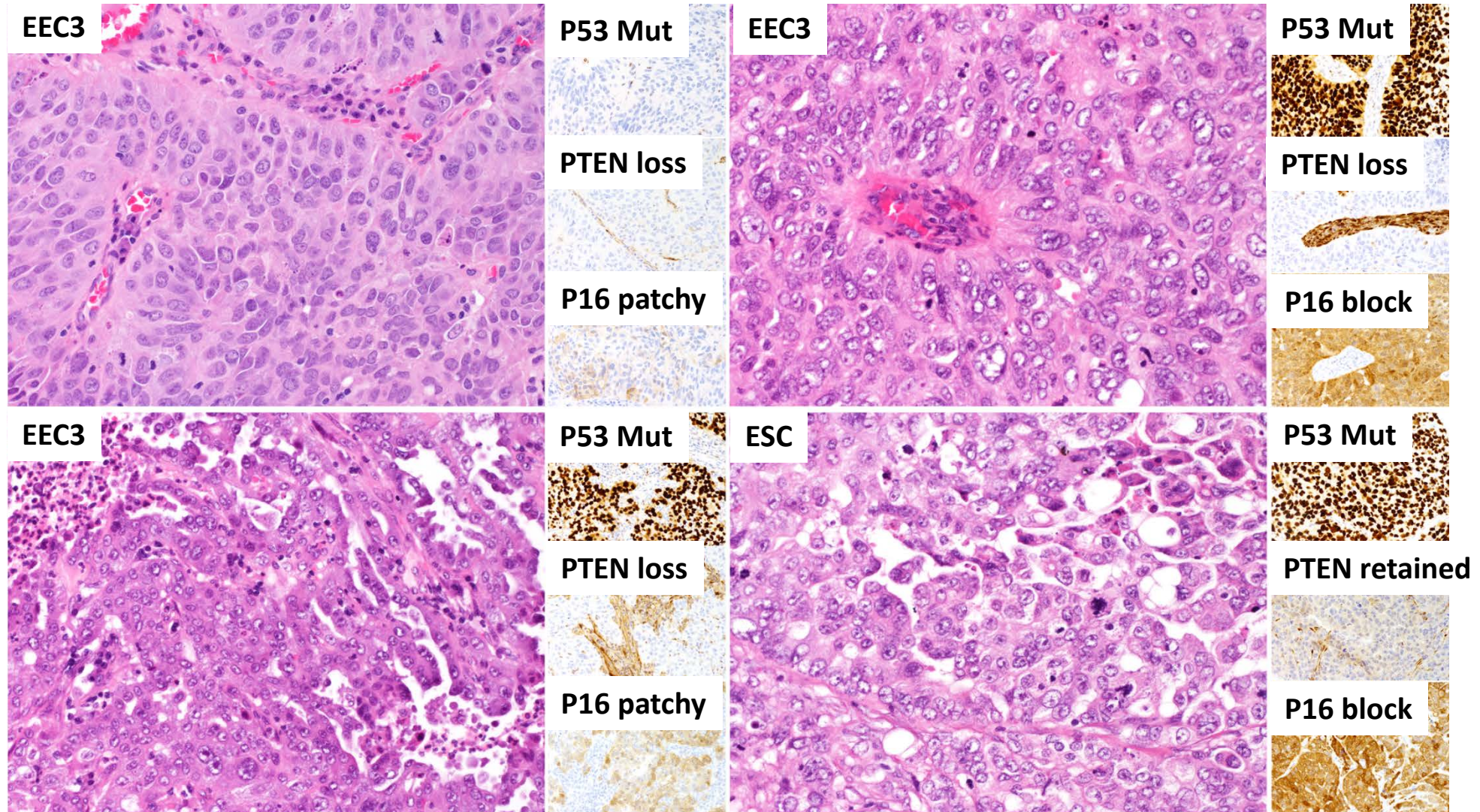
1. Exclude collision tumor of ESC with EEC1 (left)

Single tumor MMRp, no POLE mutation:
Uncommon (~2%) but prognostic uncertain
What amount of subclonal p53 is relevant?
Example below 2% **OE** versus 5% WT?



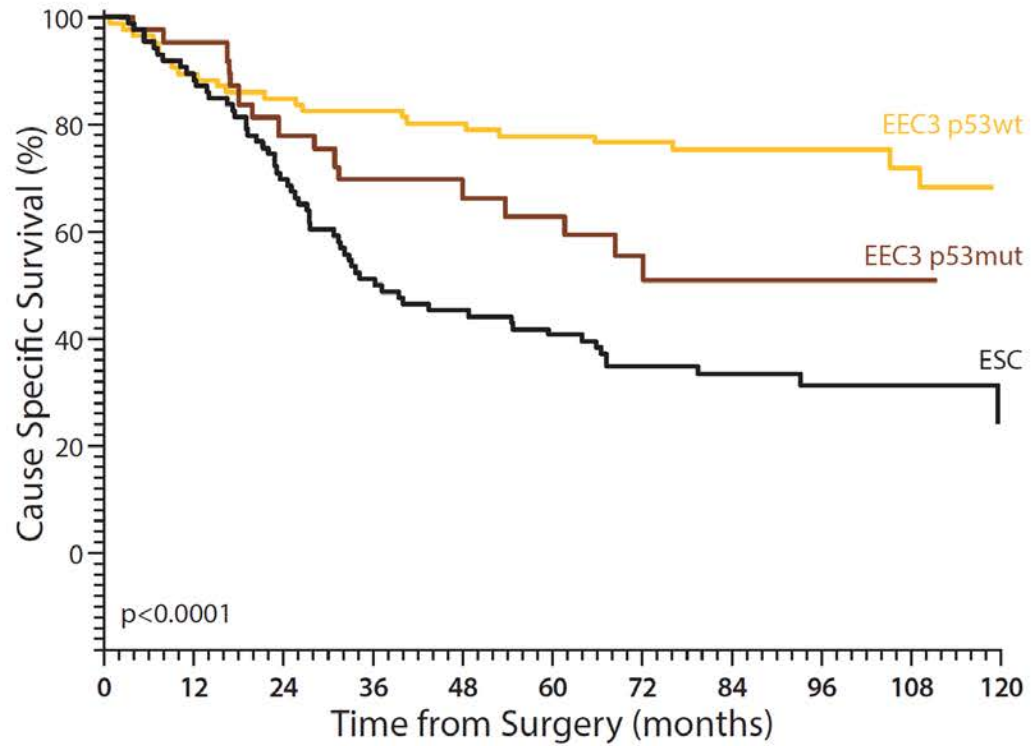
4. Histotype distinction of endometrial endometrioid carcinoma grade 3 (EEC3) from serous (ESC)



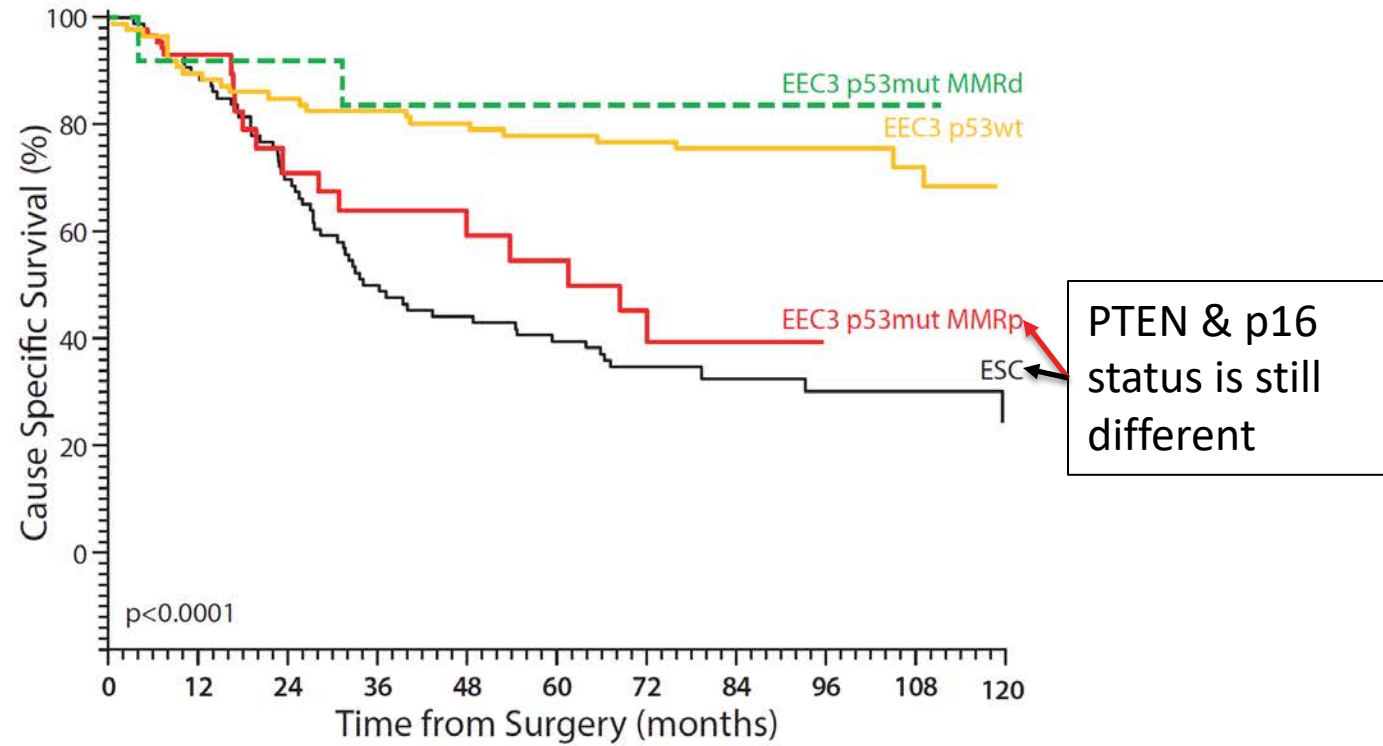


4. Subclonal p53 often associated with MMRd (“double classifier”)

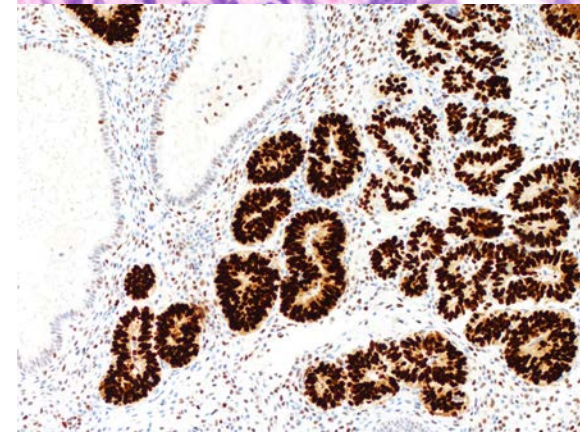
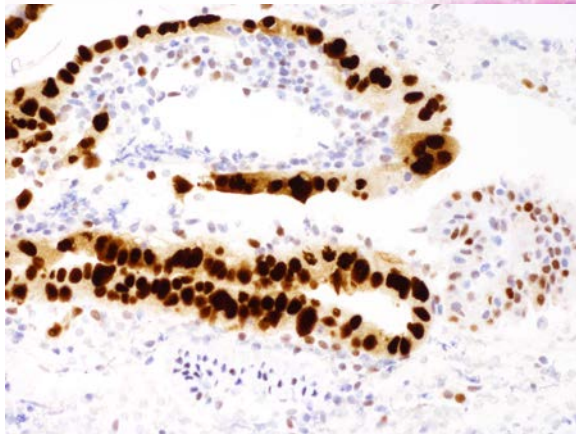
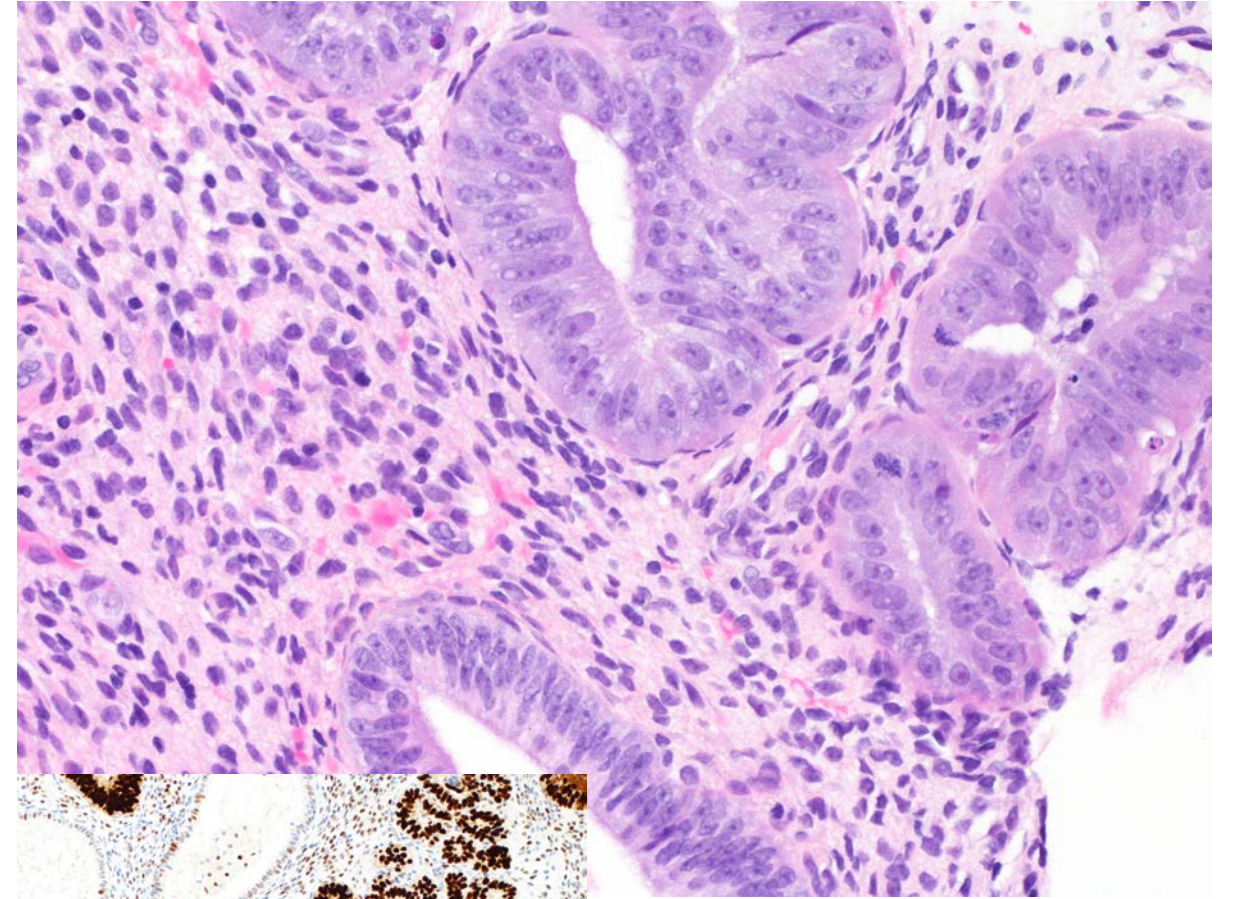
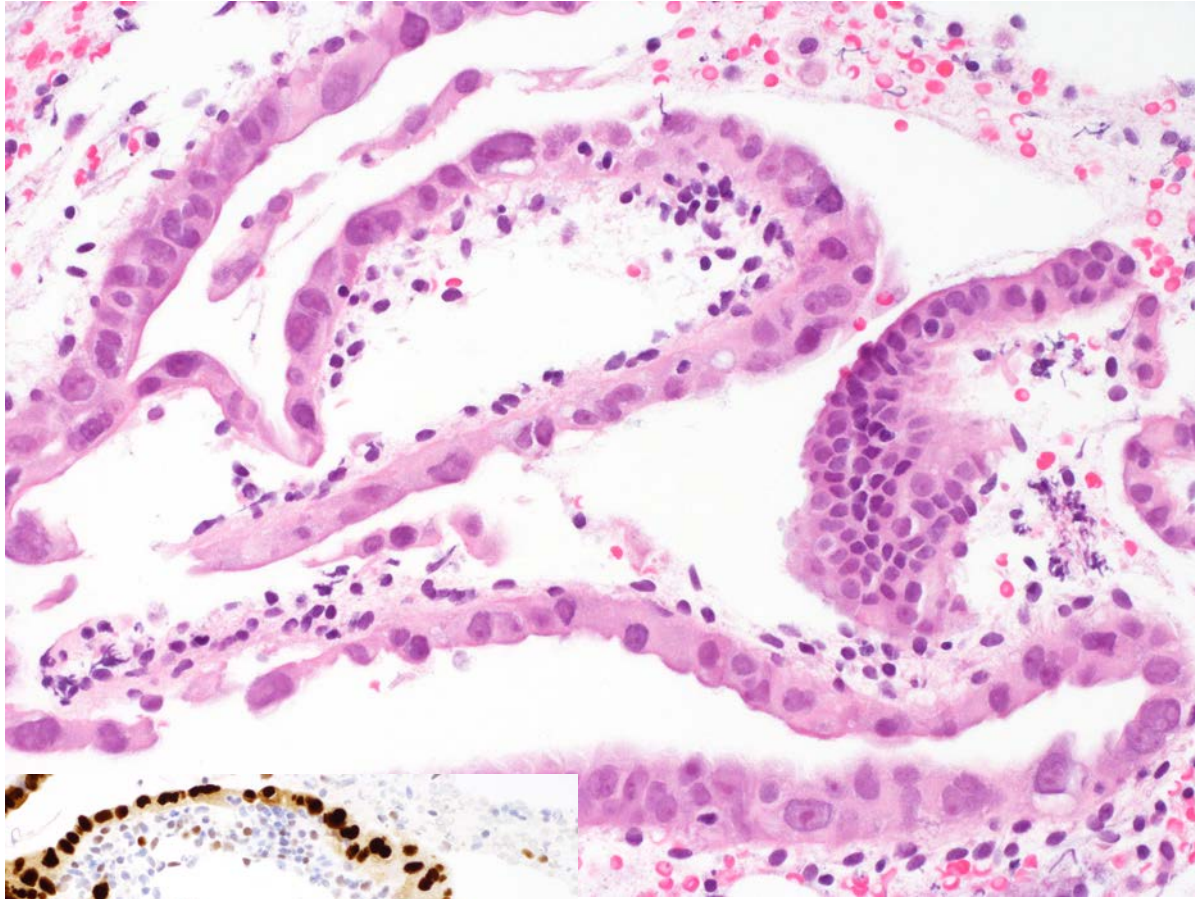
A



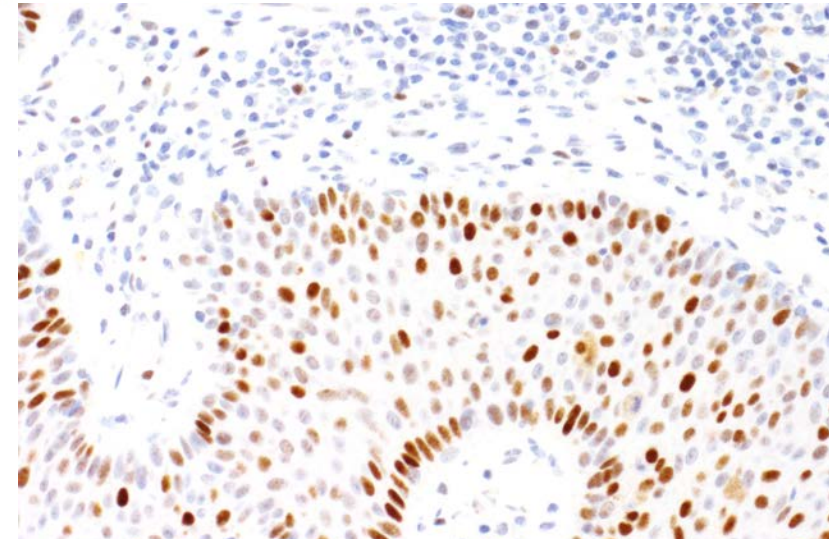
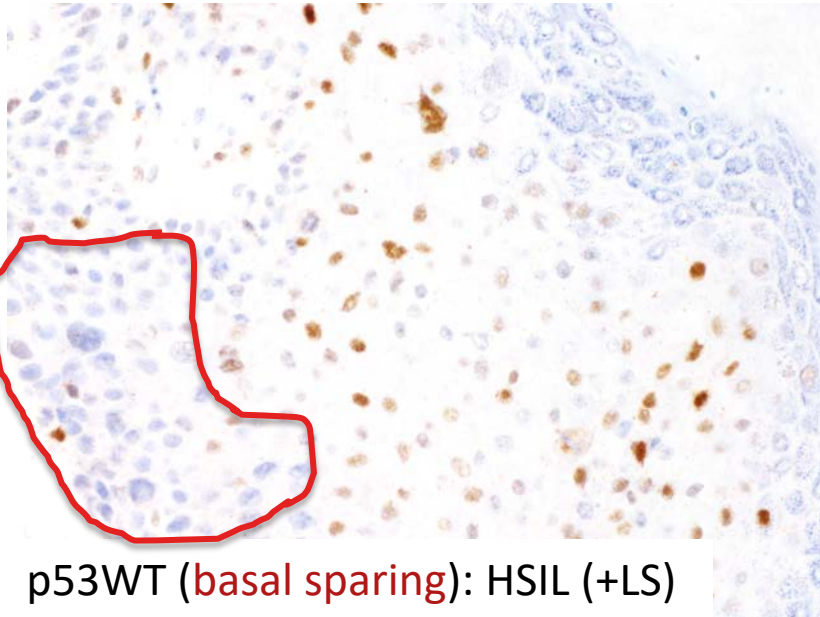
B



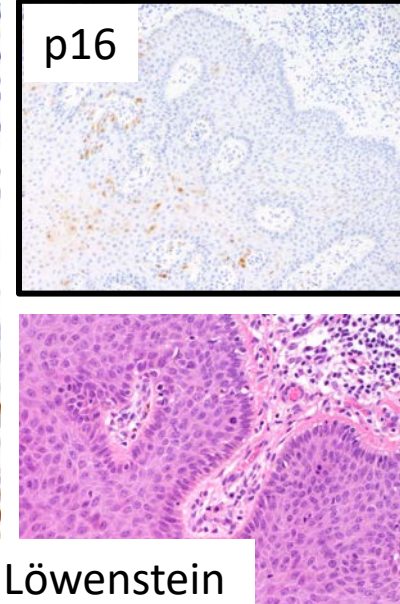
EEC3 MMRd and p53mut (double classifier) can be classified as MMRd



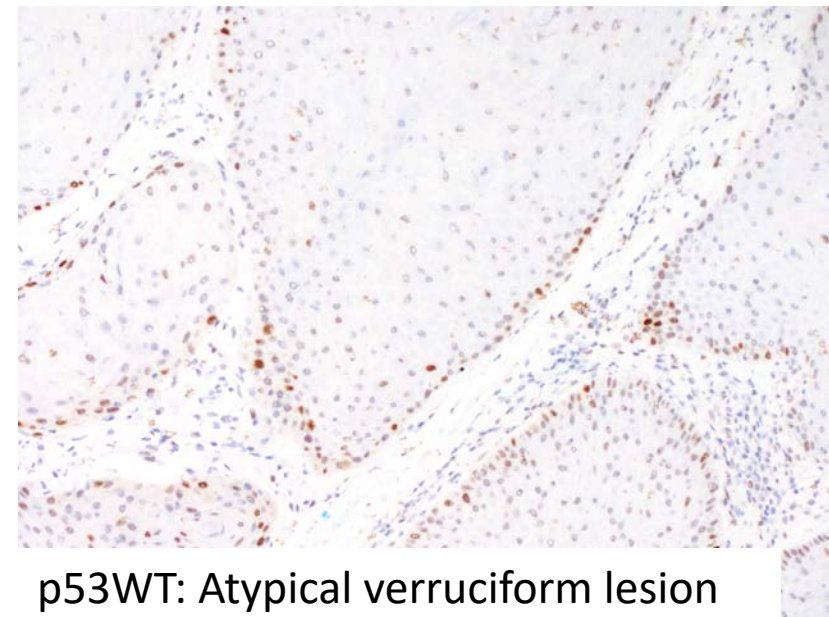
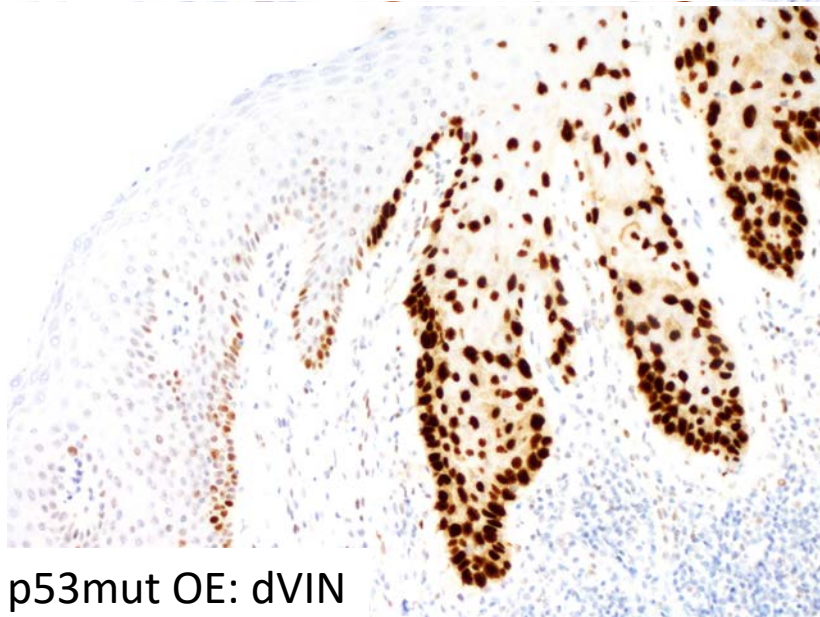
p16



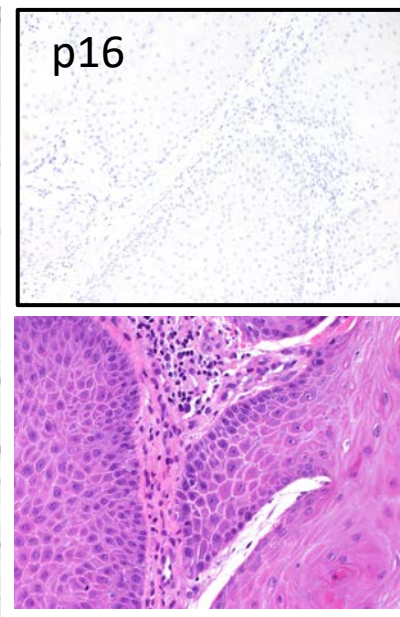
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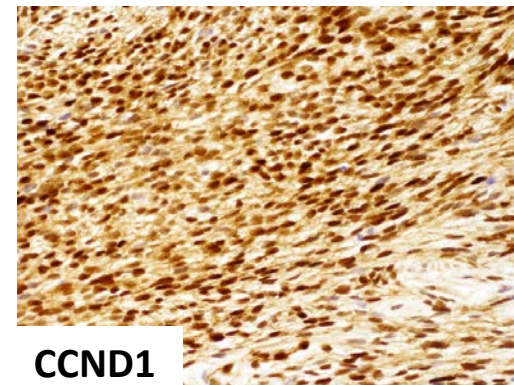
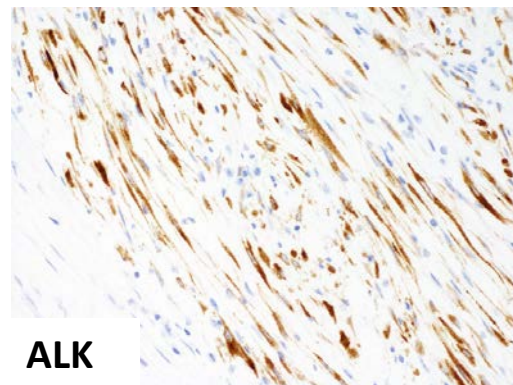
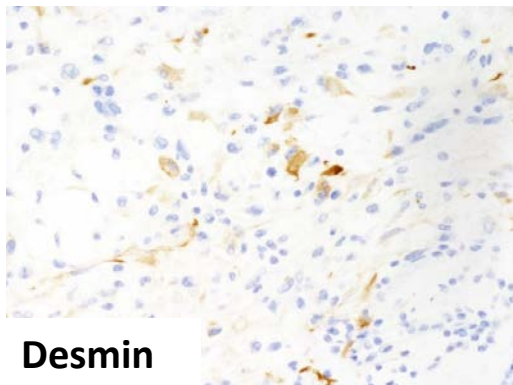
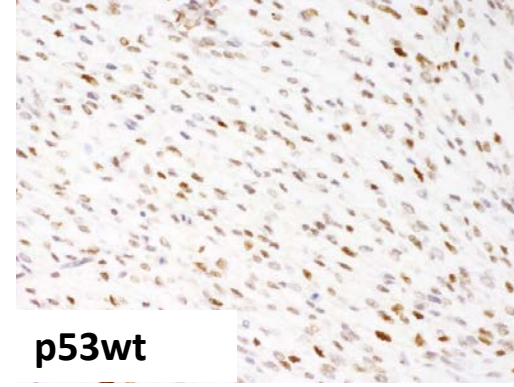
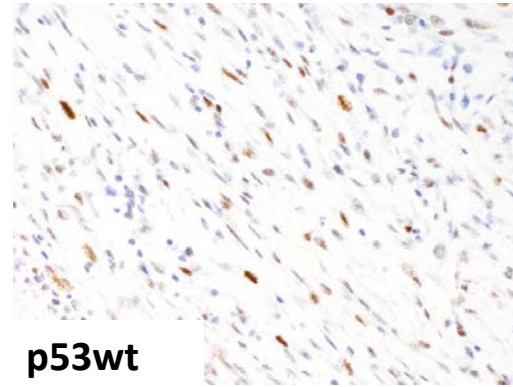
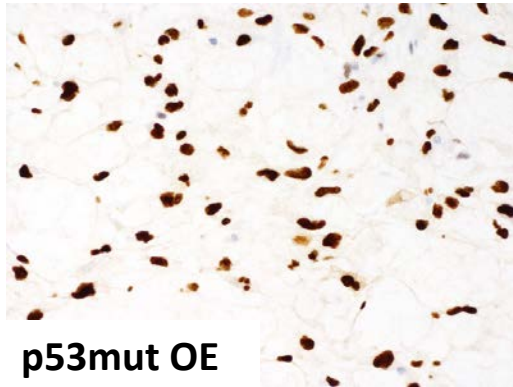
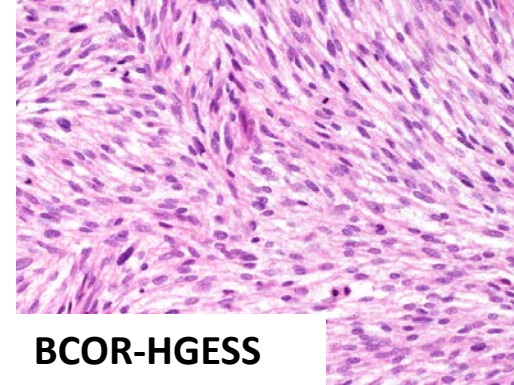
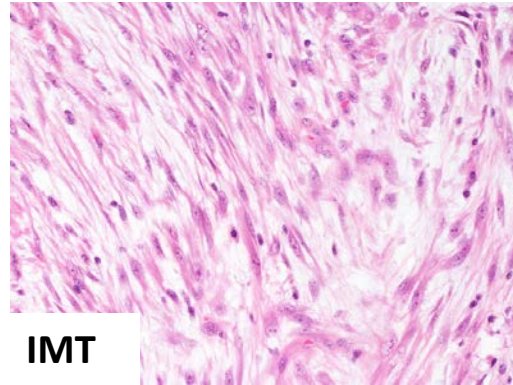
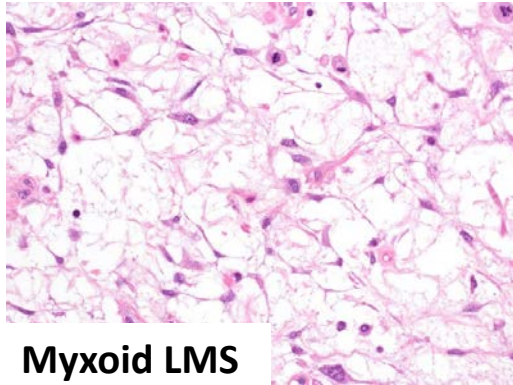
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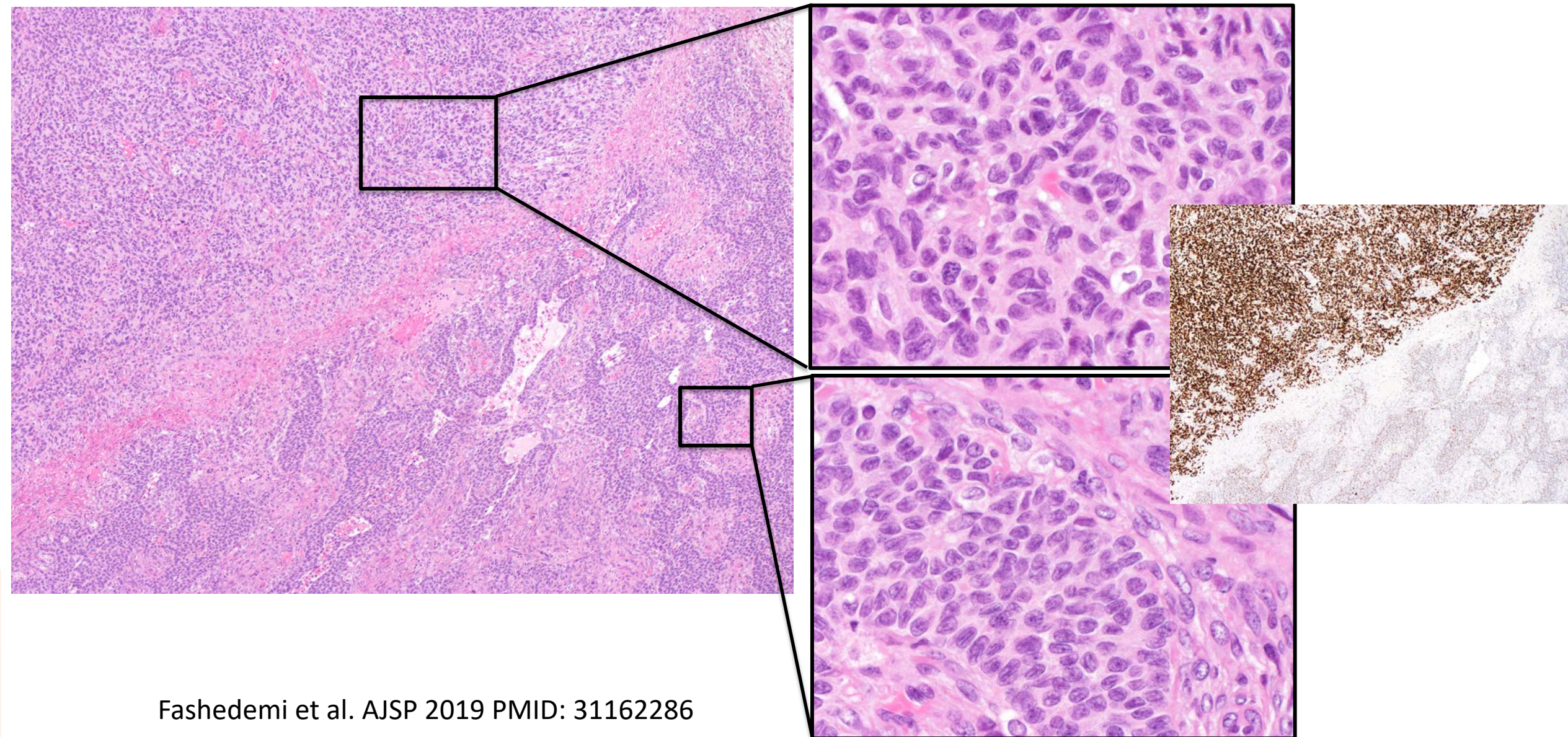
p16



8. Triaging of uterine mesenchymal neoplasms to translocation testing

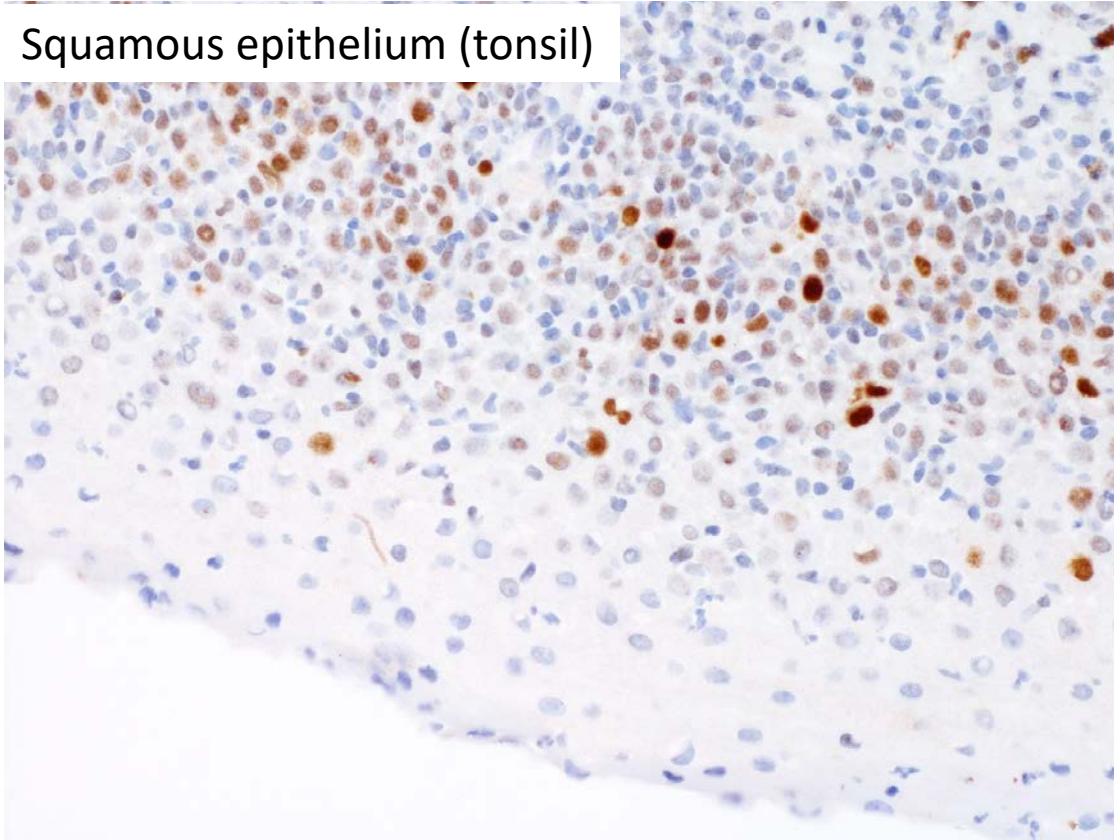


P53 wild type cases may be referred to molecular translocation testing

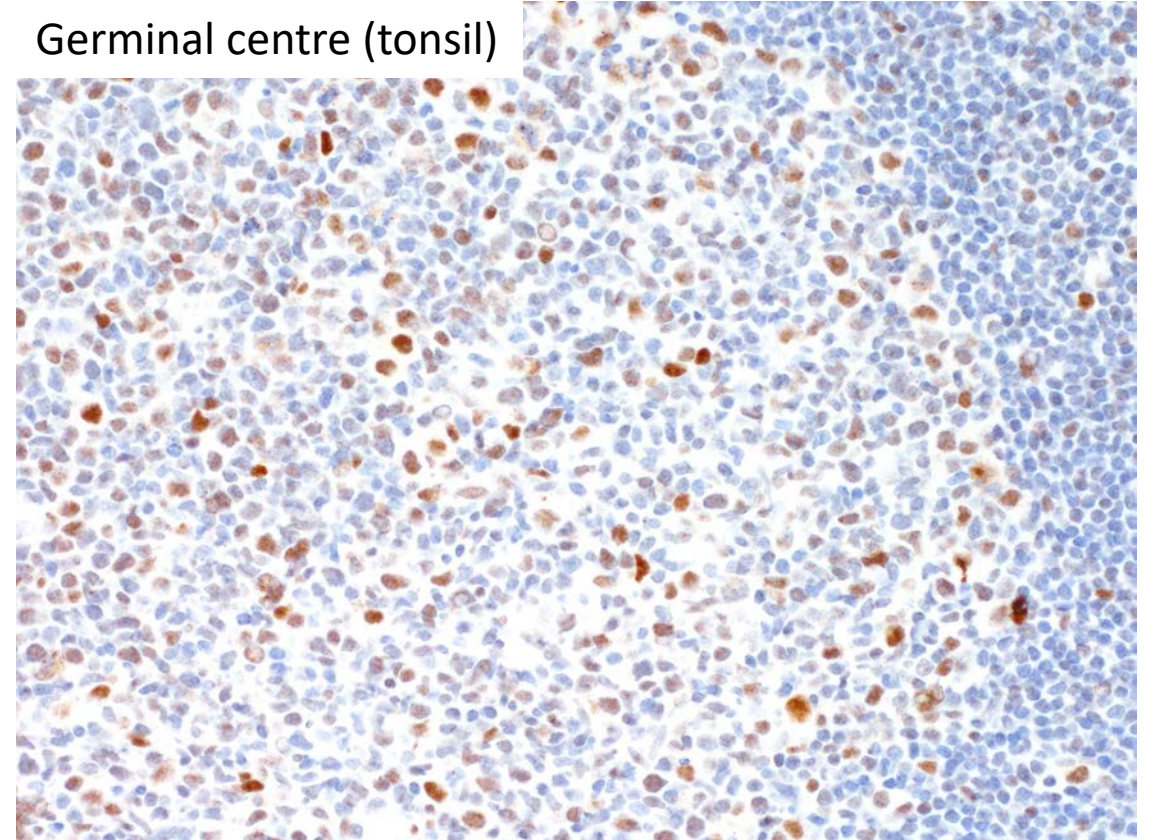


1. Distinction of LGSC from HGSC
2. Distinction of OEC from HGSC
 - a) Assigning molecular subtype of OEC
3. Confirmation of HGSC precursors
4. Assessing risk in MBOT
5. Distinction of EEC3 from ESC
 - a) Assigning molecular subtype of EEC
6. Confirming of ESC precursors
7. Prognostication in squamous vulvar lesions
8. Triaging of uterine mesenchymal neoplasms to translocation testing

Squamous epithelium (tonsil)



Germinal centre (tonsil)



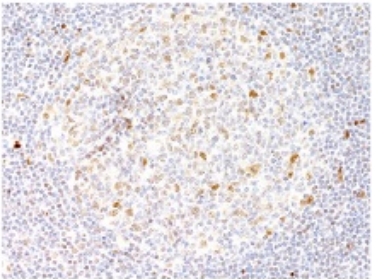
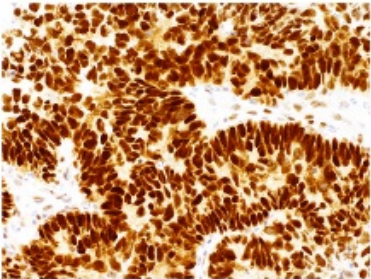
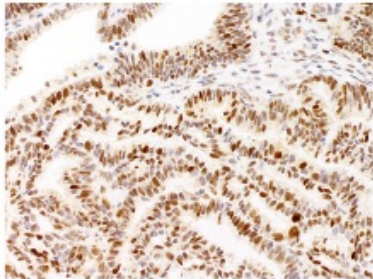
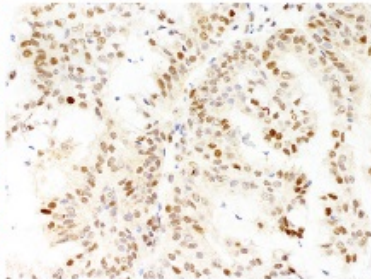
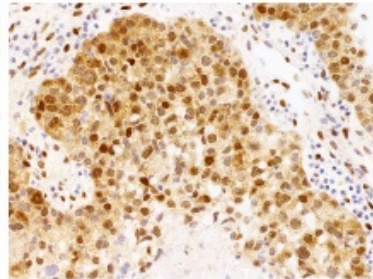
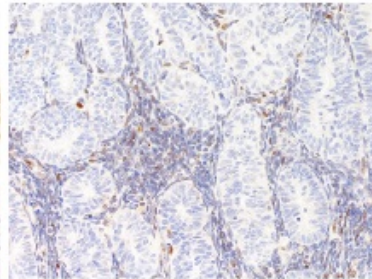
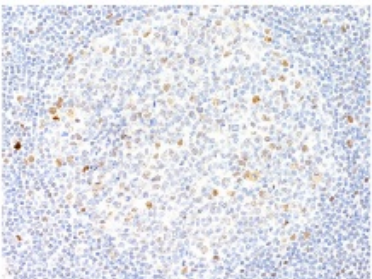
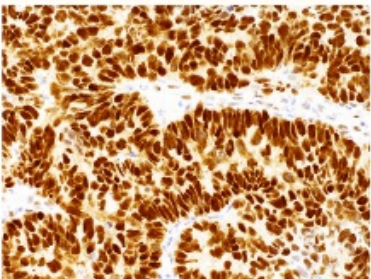
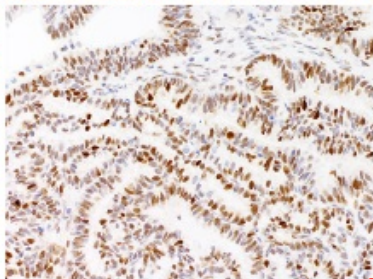
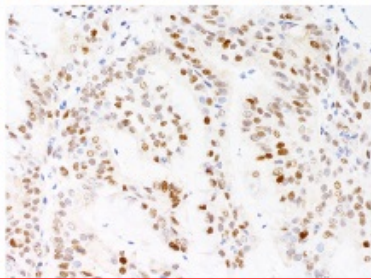
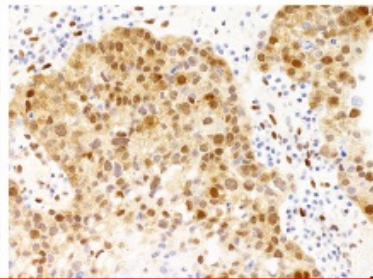
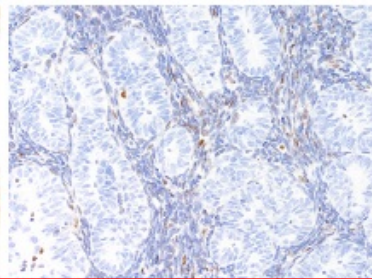
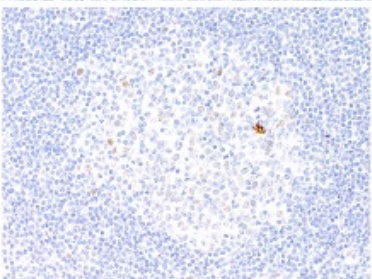
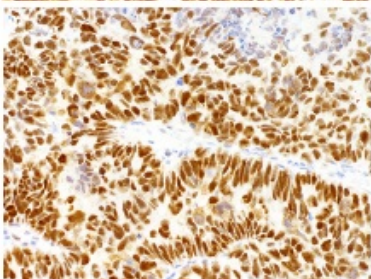
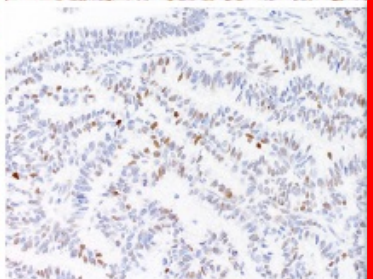
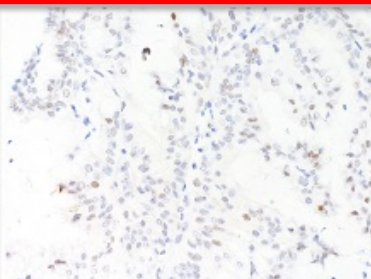
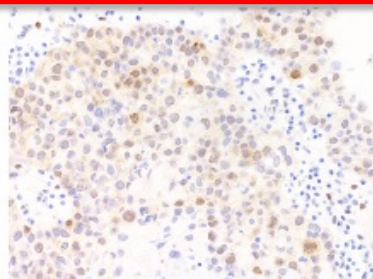
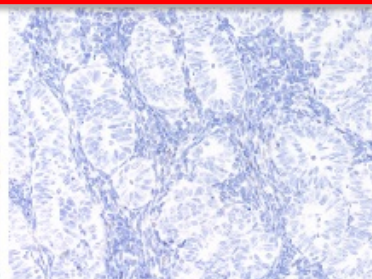
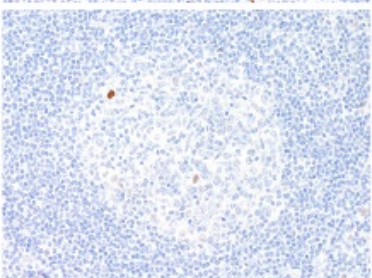
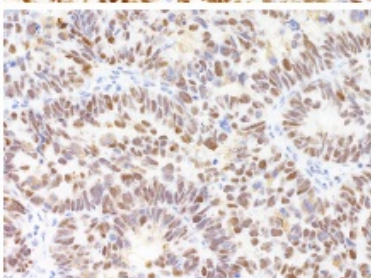
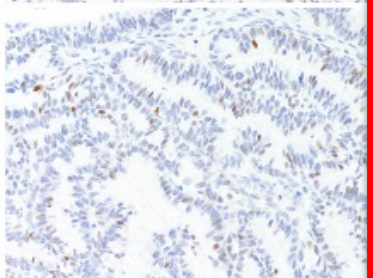
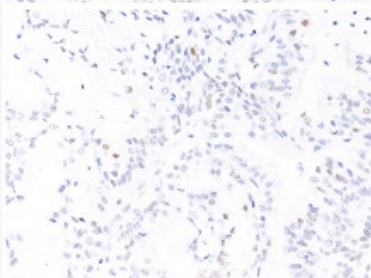
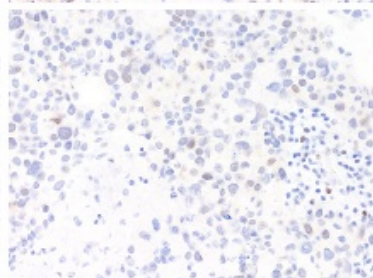

DO7 clone on Dako Omins

Detailed protocol: Singh et al. J Pathol PMID: 31829441

Email: mkoebel@ucalgary.ca

<https://www.nordiqc.org/>

<https://www.cpqa.ca/>

	Tonsil: Wild-type	Overexpression	Wild-type	Wild-type	Cytoplasmic	Complete absence
#1						
#2						
#3						
#4						

Köbel et al.
IJGP 2019
PMID: 29517499

THANK YOU!



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