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## Endometrial Carcinoma Diagnosis: Use of FIGO Grading and Genomic Subcategories in Clinical Practice: Recommendations of the International Society of Gynecological Pathologists

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### SUMMARY

In this review, we sought to address two important issues in the diagnosis of endometrial carcinoma: how to grade endometrial endometrioid carcinomas (EECs) and how to incorporate the four genomic subcategories of endometrial carcinoma, as identified through The Cancer Genome Atlas, into clinical practice. The current FIGO grading scheme provides prognostic information that can be used to guide the extent of surgery and use of adjuvant chemotherapy or radiation therapy. We recommend moving toward a binary scheme to grade EECs by considering International Federation of Gynecology and Obstetrics (FIGO)-defined grade 1 and 2 tumors as “low grade” and grade 3 tumors as “high grade”. The current evidence base does not support the use of a three-tiered grading system, although this is considered standard by FIGO, the American College of Obstetricians and Gynecologists, and the College of American Pathologists. As for the

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four genomic subtypes of endometrial carcinoma (copy number low/p53 wild-type, copy number high/p53 abnormal, polymerase E (POLE) mutant, and mismatch repair deficient), which only recently have been identified, there is accumulating evidence showing these categories can be reproducibly diagnosed and accurately assessed based on biopsy/curettage specimens as well as hysterectomy specimens. Furthermore, this subclassification system can be adapted for current clinical practice and is of prognostic significance independent of conventional variables used for risk assessment in patients with endometrial carcinoma (e.g., stage). It is too soon to recommend the routine use of genomic classification in this setting; however, with further evidence, this system may become the basis for the subclassification of all endometrial carcinomas, supplanting (partially or completely) histotype and grade. These recommendations were developed from the International Society of Gynecological Pathologists Endometrial Carcinoma project.

## Keywords

Endometrial cancer; FIGO grade; The Cancer Genome Atlas; TCGA; genomic subtype

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## Introduction

This review focuses on the grading of endometrial endometrioid carcinoma (EEC) and the emerging molecular classification of endometrial carcinoma. In this review, we will discuss the pertinent literature and make recommendations as appropriate. At the end of each section, we will identify unresolved issues for which there are currently insufficient data to make recommendations, with the hope that these issues will be investigated in future research. These recommendations were developed by the authors as part of the International Society of Gynecological Pathologists Endometrial Carcinoma Project.

### Topic 1: How to apply the International Federation of Gynecology and Obstetrics (FIGO) grading system for EEC

According to current practice standards, EECs are assigned a FIGO grade based on the degree of glandular differentiation. Grade 1 tumors exhibit  $\leq 5\%$  solid non-glandular, non-squamous growth; grade 2 tumors from 6–50%; and grade 3 tumors  $>50\%$ .<sup>1–3</sup> The presence of marked cytologic atypia increases the grade one level. Since mucinous adenocarcinomas of the endometrium are closely related to endometrioid carcinomas, it is reasonable to use FIGO grade for those carcinomas as well. However, FIGO grading should NOT be used when endometrioid or mucinous differentiation is in doubt or cannot be established. All of the other endometrial tumor types carry an intrinsic tumor grade (i.e., serous, clear cell and undifferentiated carcinomas and carcinosarcomas are high grade). Strategies for grading mixed epithelial tumors will be discussed later in this review and in another review in this issue.

In clinical practice, determining FIGO grade is not always clear-cut. Based on personal experience and anecdotes accrued during participation in the Gynecologic Oncologic Group Pathology section, it appears that many pathologists consider tight, small microacini with barely visible lumens as solid growth, although FIGO grading rules do not specifically discuss this and some pathologists characterize such patterns as “glandular” (Figure 1). For

the purposes of grading, we endorse that a confluent microacinar pattern constitutes “solid” growth, although this is not evidence-based. As stated in the FIGO criteria, squamous differentiation should be discounted as evidence of solid growth, but there are inevitable problems with grading tumors containing solid growth that resembles immature squamous epithelium and tumors that feature transitions between non-keratinizing squamous epithelium and spindle cell change. It is reasonable to adjudicate these types of cases by paying attention to the nuclear grade, first in the glandular component and then, if that approach is not informative, in the solid component. We designate a tumor as FIGO grade 3 if the solid areas resemble poorly differentiated non-keratinizing squamous cell carcinoma.

Another common problem in grading concerns the degree and extent of nuclear atypia that is sufficient to upgrade a tumor from one FIGO category to another.<sup>4,5</sup> The philosophy underlying our approach to this problem is that discordance between architectural grade and nuclear grade should be uncommon in endometrioid adenocarcinomas. The first step is to ensure the nuclear features are sufficiently atypical. An easy guideline is to ask yourself, “Is this focus easily appreciated on scanning or intermediate power examination?” and “Is the atypia so bad that I would consider it grade 3 on a 3-point scale?” If the answers are “yes,” you should sample the tumor extensively to determine whether the finding is limited only to a few glands. If the change is diffuse, it is advisable to at least question whether part or all of the tumor could be a serous carcinoma or a clear cell carcinoma, instead of an endometrioid carcinoma.<sup>6</sup> It is acceptable to move such a tumor from one FIGO grade to another only after determining that the tumor is indeed endometrioid throughout. A mixed epithelial carcinoma with an endometrioid component should be diagnosed when the cytologically atypical area is determined to be serous or clear cell on further study. One should not upgrade endometrioid carcinomas containing only a few glands with atypical nuclei, especially if it took a prolonged search at high-power magnification to recognize them. We endorse the severe nuclear atypia qualification described by Zaino et al.,<sup>4</sup> i.e., that severe nuclear atypia in the majority of cells (>50%) is required to upgrade a grade 1 or grade 2 EEC. “Upgrading” a case based on nuclear features is only rarely prudent, since in most cases, tumors are upgraded inappropriately (nuclear atypia is mild-moderate and diffuse, or severe and only focally found) or are not endometrioid at all.

Classification and regression tree (CART) statistical analyses have demonstrated that after tumor stage, the next most informative prognostic division in EEC is between high-grade (grade 3) and low-grade (grade 1/2) tumors.<sup>7,8</sup> While it is reported that there is a small but statistically significant difference in survival of up to 5% between clinically low-stage grade 1 and grade 2 EECs,<sup>9</sup> this has not been consistently demonstrated. Therefore, the key consideration in tumor grading at hysterectomy should be to identify the presence of grade 3 EEC or any other high-grade components that may adversely affect patient prognosis, and not in trying to distinguish between grade 1 and 2 EEC. We have designated this system as “binary FIGO”, which entails combining grade 1 and grade 2 tumors into a low-grade category and grade 3 tumors into a high-grade category.<sup>10,11</sup> Said another way, EECs with low nuclear grade and harboring 50% solid tumor components would be considered “low-grade EEC” (Figure 1). This is in keeping with current risk assessment/treatment guidelines, such as those of the National Comprehensive Cancer Network (NCCN), European Society of Medical Oncology (ESMO),<sup>12</sup> the Mayo Clinic,<sup>13</sup> and Memorial Sloan Kettering Cancer

Center (MSK) in which grade 1 and grade 2 EECs are managed the same way. Although NCCN (NCCN guideline version 1.2018) still recognizes 3 EEC grades, the staging guidelines for grades 1 and 2 are very similar. Differences in the risk of lymph node metastasis in grades 1 and 2 EEC are most pronounced when decisions for staging are undertaken by evaluating biopsy or curettage material only. As will be discussed subsequently, the distinction between grades 1 and 2 is unimportant in the setting of planned lymphadenectomy or sentinel lymph node mapping, and distinguishing between these grades does not stratify patients who have had comprehensive surgical staging into different risk categories.

A literature review of various proposed binary grading schemes (summarized in Table 1 and a recent review on grading EEC)<sup>1</sup> reveals that EECs can readily be divided into either a low- or high-risk prognostic group based on a number of factors associated with increased tumor aggressiveness; the different grading systems show only slight disagreement over the distribution of a small number of gray-area cases between the two main prognostic groups. All of the proposed two-tiered grading systems have been shown to be equal or superior to the current three-tiered FIGO system in terms of interobserver variability kappa score, and the binary FIGO system, in the studies in which it was assessed, consistently exceeded the three-tiered FIGO system. The binary FIGO system also performs as well as or better in terms of prognostication, compared to other binary grading schemes, and has the added advantage of being based on the currently used FIGO grading system (i.e., practicing pathologists are already familiar with the system's criteria). An assessment of the potential clinical impact of the adoption of a binary EEC grading system is complicated by variations in clinical patient management, as well as the absence of a widely accepted consensus on the indications for extensive surgical staging in patients with a preoperative diagnosis of low-grade EEC. Soslow et al. recently analyzed a departmental database of clinical stage 1 EECs (n=1544) in order to better understand relationships between preoperative tumor grade, depth of myometrial invasion, and the risk of positive sentinel lymph nodes. The analysis demonstrated that while grade 1 tumors were associated with a significantly lower rate of lymph node metastases compared with grade 2 tumors overall (6.6% vs 11.6%; p=0.003), the difference disappeared after adjusting for depth of myometrial invasion (Table 2).<sup>1</sup> Those findings further endorse the view that preoperative tumor grade alone should not be used to assess the indication for extensive surgical staging. The use of preoperative tumor grade alone to select patients for lymphadenectomy is clearly not supported by the evidence and does not reflect the complex interplay between the various pathologic parameters. In summary, we propose that it would be appropriate to adopt a binary FIGO system for grading EEC by combining the grade 1 and 2 categories into a single low-grade category in biopsy or curettages when comprehensive surgical staging is planned and in hysterectomy specimens. For patients desiring a fertility-sparing therapeutic approach, it will continue to be necessary to distinguish grades 1 and 2 based on currently used clinical criteria for conservative, hormonal therapy whereby grade 1 tumors can be considered for this approach and grade 2 tumors will generally not (NCCN guideline version 1.2018). We also propose to retain the severe nuclear atypia qualification, based on the work of Zaino et al.,<sup>4</sup> whereby severe nuclear atypia in the majority of cells (>50%) in an architecturally low-grade EEC

would lead to a high-grade designation. This proposed system should be beneficial in terms of simplification and diagnostic reproducibility.

### Recommendations

- Discuss individual clinician's preferences for moving to a binary grading scheme, while emphasizing its potential benefits
- As a bridge towards uniform adoption of the binary FIGO grading scheme, it is acceptable to report both the binary grade and the FIGO grade (i.e., low-grade EEC [FIGO grade 2])
- Going forward, we recommend the adoption of the binary FIGO grading system, when appropriate. Existing reporting datasets/synoptic templates/checklists for reporting endometrial carcinoma will need to be revised and updated

### Unresolved Issues

- Anticipating the future incorporation of The Cancer Genome Atlas (TCGA) classification system into endometrial cancer diagnosis and prognostication, it remains uncertain whether grading will remain an important clinical discriminator for tumors, especially in the POLE category.

## Topic 2: How to synthesize genomically defined types of endometrial carcinoma with their morphological, immunophenotypic, and clinical characteristics

For the purposes of this review, FIGO grade 3 endometrioid, serous, and clear cell carcinomas will be considered "high-grade endometrial carcinomas" (HGECs). Undifferentiated and de-differentiated carcinomas and carcinosarcomas are dealt with elsewhere in this review series.

Various studies have shown that an HGEC diagnosis is not highly reproducible.<sup>20,21</sup> This implies that: 1) diagnostic criteria are insufficiently detailed for practical use; 2) and/or diagnostic criteria were developed empirically and may not be evidence based; 3) and/or uniform diagnostic criteria are not utilized in practice. This lack of reproducibility is significant in that the reported clinicopathological and molecular data concerning HGEC may not be readily interpreted because of differences in case series based on inconsistent diagnoses. Examples of the latter problem are the lack of consensus regarding whether grade 3 endometrioid carcinomas are "type II endometrial carcinomas" and whether they are as aggressive as serous carcinomas.

There have been a number of attempts to refine diagnostic criteria using evidence-based approaches. Among other groups, the MSK group used immunohistochemical and clinical evidence to describe gland-forming serous carcinomas that are distinct from FIGO grade 1 and 2 endometrioid carcinomas<sup>4,22</sup> (Figure 2); the MD Anderson group used supporting clinical evidence to separate undifferentiated carcinomas from grade 3 endometrioid carcinomas<sup>23,24</sup>; the Massachusetts General Hospital group used supporting clinical

evidence to separate corded and hyalinised endometrioid carcinomas (CHECs) from carcinosarcomas<sup>25</sup>; and the Vancouver group used a genomically based classifier to refine diagnostic criteria that more precisely separate grade 3 endometrioid carcinoma with a papillary/villoglandular or solid architectural pattern from serous carcinomas with similar architecture.<sup>26</sup> These latter criteria also have clinical relevance.

Publication of the TCGA endometrial carcinoma dataset has provided pathologists a unique opportunity to further refine diagnostic criteria for serous and grade 3 endometrioid carcinomas.<sup>27</sup> These data do not include information about clear cell carcinoma, undifferentiated/dedifferentiated carcinoma, or carcinosarcoma.

Unlike the Bokhman “type I vs type II” classification of endometrial carcinoma, TCGA data indicated four genomic types of endometrial carcinoma. These subcategories of endometrial carcinoma were identified based on their genomic architecture. They include the following: polymerase E (POLE) mutated/ultramutated; microsatellite instability-high (MSI-H)/hypermutated; copy number low (conceptually similar to “type I” endometrial carcinoma); and copy number high (presence of many genomic amplifications and/or deletions), also known as “the serous-like group”, which is similar to “type II” endometrial carcinoma. The copy number low group has the lowest mutation burden, while the hypermutated and ultramutated groups have, on average, 10- and 100-fold more mutations, respectively; the higher mutation burdens in these latter two groups are secondary to defects in mismatch repair and POLE-mediated DNA repair, respectively. Note that mutations in *TP53* were a distinguishing feature, when comparing copy number high versus copy number low groups (Figure 3), once POLE and MSI tumors were excluded.

The overwhelming majority of low-grade (FIGO grade 1 and 2) endometrioid carcinomas map to the copy number low and MSI-H categories. Grade 3 endometrioid carcinomas are highly heterogeneous, as they are found in every genomic category, and are least represented in the copy number low group. Endometrioid carcinomas mostly have an endometrioid genomic profile, with or without *TP53* mutation/high copy number alterations, while serous carcinomas have *TP53* mutations/high copy number alterations without additional mutations characteristic of endometrioid carcinomas.

Assigning a tumor to the serous category requires the presence of a *TP53* mutation/abnormal “mutation pattern” p53 immunostaining, the absence of endometrioid-type mutations (although this point is debatable, as some low-grade endometrioid carcinomas with *PTEN* mutations acquire *TP53* mutations during tumor progression, along with “serous” morphology), and the presence of high copy number alterations. All serous carcinomas map to the copy number high, “serous-like” category, as do approximately one quarter of grade 3 endometrioid carcinomas. Although TCGA data are incomplete, they were sufficient to generate a Kaplan-Meier survival curve showing that POLE carcinomas have outstanding clinical outcomes irrespective of grade and perhaps stage. MSI-H and copy number low tumors have intermediate clinical risk, and the serous-like group, unsurprisingly, has the worst outcomes.

The first approach to extracting this information for practical diagnostic use was to describe the morphology of endometrioid carcinomas in each group. Although there are many characteristic features of tumors from the POLE<sup>28,29</sup> (peritumoral lymphocytes, tumor infiltrating lymphocytes, and intratumoral heterogeneity, tumor giant cells), MSI-H<sup>30,31</sup> (peritumoral lymphocytes, tumor infiltrating lymphocytes, and intratumoral heterogeneity), and serous-like groups (diffuse high nuclear grade), there is a general feeling that these features are incomplete, partly overlapping, and may not be able to be applied reproducibly. A particular problem that is not adequately addressed with a morphology-only approach is the existence of morphologically ambiguous tumors that reside in the POLE, MSI-H, and serous-like categories (but not the copy number low category),<sup>32</sup> and many POLE endometrioid carcinomas have components or subclones that are histologically similar to serous carcinomas. Given the particular heterogeneity of grade 3 endometrioid carcinomas, we were faced with the following questions and answers:

- Are serous-like grade 3 endometrioid carcinomas, as genetically defined, really endometrioid based on morphological features/light microscopic examination?
  - In some cases, they can show unequivocally endometrioid morphological features, e.g., squamous differentiation.<sup>33</sup> Many are histologically ambiguous, as mentioned above, making the distinction between serous-like grade 3 endometrioid and serous carcinomas difficult or impossible; this also raises the question of whether there is a justification for attempting to separate these tumors.
- Are serous-like grade 3 endometrioid carcinomas clinically interchangeable with serous carcinomas?
  - *TP53*-mutated grade 3 endometrioid carcinomas, compared with those lacking *TP53* mutation, recur more frequently (T. Bosse et al, Am J Surg Pathol 2018, in press). Detailed analyses indicate that *TP53*-mutated grade 3 endometrioid carcinomas (i.e., serous-like grade 3 endometrioid carcinomas) have a high-risk profile, with only rare exceptions. Although there are few such cases, it appears that *TP53*-mutated grade 3 endometrioid carcinomas from the POLE category retain the clinically low-risk profile of POLE endometrioid carcinomas that lack a *TP53* mutation. There are sparse data suggesting that serous-like grade 3 endometrioid carcinomas metastasize to the peritoneum less frequently than serous carcinomas, but this requires validation. Currently, there is no compelling evidence to support that serous-like grade 3 endometrioid carcinomas are significantly clinically different than serous carcinomas, and unless there is improvement in diagnostic reproducibility for these two categories, it is difficult to envision such a distinction being a variable that can be used to guide patient treatment decisions.
- Can p53 immunohistochemistry alone be used to identify those grade 3 endometrioid carcinomas that belong to the copy number high group?

- No. As aberrant p53 staining is not confined to tumors in the serous-like category, additional ancillary tests would be necessary to exclude a *TP53*-mutated POLE or MSI-H grade 3 endometrioid carcinoma. Of interest, many *TP53* mutations found in the POLE and MSI-H categories involve portions of the *TP53* gene that are not mutated in serous carcinomas,<sup>34</sup> leading to imperfect correlation between *TP53* mutation and aberrant p53 staining in these groups of tumors.
- Is there a better assay than p53 immunohistochemistry to determine which grade 3 endometrioid carcinomas are serous-like?
  - p53 immunohistochemistry was compared to *TP53* mutational analysis and fluorescent in situ hybridization (FISH) for copy number analysis and found to be equivalent in identifying the copy number high serous/serous-like group<sup>34</sup> (see below). Since aberrant p53 immunostaining is not synonymous with a copy number high tumor, other methods may prove to be superior surrogate markers of this genomic subtype. Aneuploidy, which is intuitively a good candidate marker of copy number high tumor, frequently was found to be present within the MSI-H and POLE categories, so testing to exclude MSI-H and POLE is, at this time, necessary prior to separation of copy number high and low tumors.<sup>35</sup>

We used TCGA data to extract surrogate markers for the genome-wide analyses used to classify endometrial cancers into one of the four genomic categories, and then validated the classifier in an independent set of cases; this classifier, referred to as the ProMisE (Proactive Molecular Risk Classifier for Endometrial Cancer) classifier, includes mismatch repair (MMR) immunostaining (rather than MSI analysis, as was used in TCGA), sequencing of *POLE* exonuclease domains (rather than quantification of mutations genome wide) and p53 immunohistochemistry (rather than quantification of copy number alterations genome wide or *TP53* mutational analysis), and was able to recreate the four prognostically significant genomic subsets of endometrial carcinoma<sup>34,36</sup> (Table 3). An identical strategy independently used by the group from Leiden was applied to clinical trial cases and also demonstrated that the four molecular subtypes are of prognostic significance.<sup>37</sup>

## PROPOSAL

Based on this information, we propose a multimodality classification system of HGECs that separates the four genomically defined groups of HGEC using only the following assays (Table 3 and Figure 4): POLE mutational analysis and immunohistochemistry for p53 and the DNA mismatch repair markers PMS2 and MSH6. As noted previously, this algorithm has been validated in three independent case series, including in Post Operative Radiation Therapy in Endometrial Carcinoma (PORTEC) clinical trial cases.<sup>34,36,37</sup>

This multimodality molecular classification system may not be feasible in many pathology laboratories, especially in resource-restricted areas of the world. In resource-rich regions of the world, the main impediment to making this proposal the standard of care is not only buy-in from our clinical colleagues and the availability of POLE mutational testing, but also the



recognition by regulatory authorities that this is a crucial, medically valid, and necessary assay (and in the US, this means being worthy of reimbursement by insurance companies).

For laboratories that do not have the means to perform *POLE* mutational analysis, it will still be possible to identify the MMR-deficient (MMR-D) and p53 abnormal groups via immunostaining with a three-marker panel (PMS2, MSH6, and p53). The approximate 8% of endometrial cancers in the *POLE* group will not be identified, and separating these frequently histologically high-grade and FIGO stage I carcinomas from serous carcinomas will remain a problem. At this time, we do not know if the favorable prognosis of the histologically high-grade *POLE*-mutant endometrial cancers reflects indolent biology or a good response to treatment, and until this is resolved through clinical trials, conventional therapy, based on current risk assessment (e.g., ESMO guidelines), will continue to be implemented for these tumors, and their prognosis will remain excellent.

There will be obstacles to applying the ProMisE algorithm not only because of complexities involving implementation of *POLE* mutational analysis, but also because of some resistance to performing universal DNA MMR protein immunohistochemistry. The clinical argument for universal MMR testing is compelling, and this approach is recommended in the NCCN guidelines (NCCN guideline version 1.2018) for Lynch syndrome triage and in another review in this issue. There are other benefits to universal testing, as it may be used as an ancillary diagnostic test (i.e., to help exclude a serous carcinoma or to help assign endometrial rather than cervical primary site) and, in the setting of recurrent endometrial carcinoma, it provides therapeutic prediction for the use of immunotherapy.

What about clear cell carcinomas? Work from the MSK group indicates that endometrial clear cell carcinomas are just as heterogeneous as grade 3 endometrioid carcinomas. There are *POLE* hotspot mutated, MSI-H, copy number low, and copy number high (serous-like) clear cell carcinomas, and those in the copy number low and serous-like clusters have significantly worse clinical outcomes compared to the others.<sup>38</sup> Once this work is confirmed by other groups, we would provisionally propose that clear cell carcinomas be classified based on the same multimodality system as that of grade 3 endometrioid carcinomas.

## Related and unresolved issues

### Terminology

Current treatment paradigms rely on the classic parameters of tumor histotype, especially the distinction between endometrioid versus non-endometrioid histology, and grade (i.e., FIGO grade as applied to endometrioid-type tumors). Therefore, any recommended terminology using the genomic classifier must continue to clearly include these parameters for the foreseeable future, although unpublished data from MSK and recently published data from Vancouver<sup>34,36</sup> indicate that prognosis is independent of histotype (in a multivariate model), and instead depends on stage, grade, and genomic classification. We would therefore recommend that reporting of the genomic classifier be clearly noted, as well as the analyses used to assign the classification. For example:

Endometrioid endometrial carcinoma, FIGO grade 3. Genomic classifier group: Copy number high (p53 abnormal by immunostaining)

Or

Mixed endometrioid and serous carcinoma. Genomic classifier group: POLE (POLE exonuclease mutation identified by sequencing)

### When to utilize the ProMisE algorithm

Universal MMR testing should be performed on biopsy/curettage material in patients desiring fertility-sparing conservative medical management, since the recognition of Lynch Syndrome in such cases would encourage prophylactic hysterectomy and bilateral salpingo-oophorectomy at the appropriate age. These tests should also be considered in biopsy/curettage material when treatment planning depends on genomic classification. This is not yet a standard practice, although selected gynecological pathology teams are currently using this approach. It should also be noted that antigen preservation may be superior in biopsies/curettings compared to hysterectomy specimens, and that molecular classification based on the biopsy does accurately reflect the findings in the subsequent hysterectomy specimen.

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### Tumor grading

Tumor grading in the context of molecular subtype is a complex issue. FIGO grading is presumably only appropriate for tumors that are within the MSI-H/MMR-D or copy number low (also referred to as tumors with no specific molecular profile [NSMP], as they lack POLE mutation, MMR deficiency, or abnormal p53 immunostaining) groups, as these most clearly correspond to the “classical” endometrioid tumors. FIGO grading in these two groups may provide the prognostic information that one expects from older, histology-based clinical studies, and would allow them to be provisionally placed into low- and high-risk groups, although this remains to be studied. On the other hand, tumors in the copy number high/p53 abnormal group (“serous-like”) clearly belong in the “high-risk” group, in which no histologic grading is appropriate. There is likely no value in assigning grade to tumors in the POLE group, for which studies indicate a survival rate approaching 100%. A significant proportion of these tumors morphologically would be considered high grade and could potentially be treated with unnecessary lymphadenectomy and chemotherapy. The intersection of genomic classification, surgical staging, and delivery of adjuvant therapy is not yet resolved.

### Intratumoral heterogeneity

Some researchers strongly believe that endometrial carcinomas are more heterogeneous than their ovarian counterparts (i.e., more likely to contain true mixed tumors). Van Esterik et al. demonstrated that this morphological heterogeneity does not extend to the level of genomic abnormalities, i.e., genomic subtype is uniform throughout.<sup>41</sup>

Many tumors assigned to the MMR-D/MSI-H and POLE categories demonstrate significant intratumoral morphological heterogeneity, while tumors in these groups and the p53

abnormal/copy number high cluster also typically show morphological ambiguity<sup>42,43</sup> that gives rise to diagnostic problems.<sup>32</sup> So where does intratumoral heterogeneity stop and mixed epithelial carcinoma begin? Given emerging data<sup>44</sup> indicating that most mixed epithelial carcinomas represent different morphological expressions of one clonal tumor, one is left with the impression that mixed epithelial carcinomas may not be diagnosed with any frequency in the future, apart from some exceptional situations, e.g., grade 1 EEC giving rise to undifferentiated carcinoma. Also, what do we make of the assertion that any type II component of a mixed epithelial carcinoma places the tumor in a clinically high-risk category? It is probable that most “mixed epithelial carcinomas” with poor clinical outcomes represent heterogeneous or morphologically ambiguous p53 abnormal/copy number high tumors. Other examples, however, likely represent heterogeneous or ambiguous POLE or MSI-H tumors whose clinical behavior is baseline or favorable despite the histologically high-grade appearance of one portion of the tumor. These concepts need to be developed further and validated with good clinical, morphological, and genomic data. For the time being, it is reasonable to continue to follow the ISGP recommendation that grading mixed epithelial carcinomas (such as mixed endometrioid and serous or clear cell carcinoma) be based on the highest grade component (discussed in another review in this issue). However, it also must be recognized this practice may not be clinically informative in MSI-H and POLE tumors.

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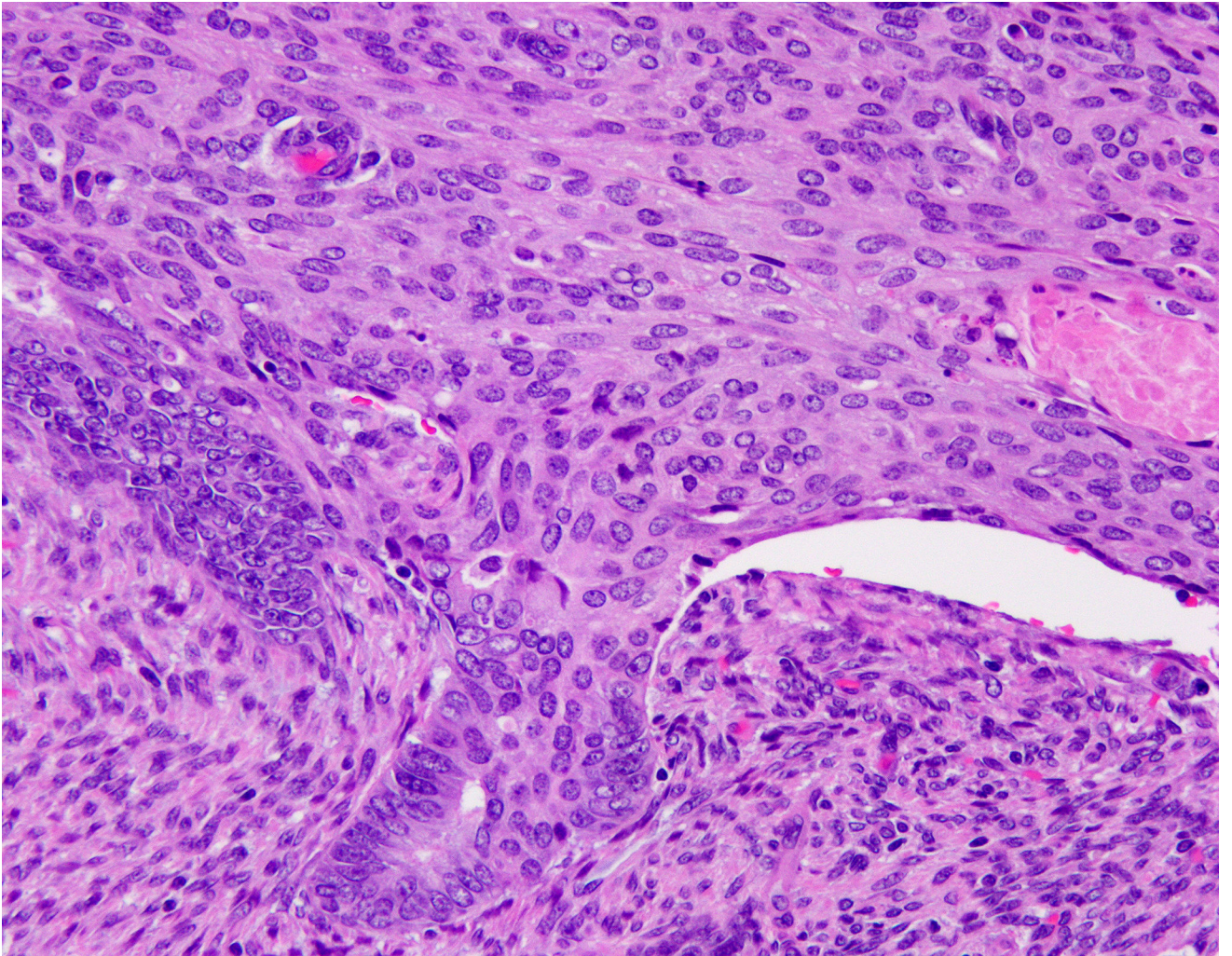
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**Summary/Recommendations**

- TCGA classification separates prognostically favorable from unfavorable types of FIGO grade 3 endometrioid carcinoma. Furthermore, it separates serous carcinoma from hitherto unknown types of clinically low-grade endometrial carcinomas that can morphologically mimic serous carcinoma to a substantial degree (e.g., POLE, MMR-D). The classification is powerfully prognostically relevant and is independently associated with clinical outcomes, unlike histological subtype assignment based on examination of hematoxylin and eosin slides alone.
- In laboratories undertaking genomic classification, this should be reported in conjunction with grade and histotype.

**Unresolved issues**

- Since the practical application of TCGA classification relies on universal POLE gene sequencing and universal immunohistochemistry for p53 and two or four of the DNA MMR markers, reimbursement and financial issues need to be considered. We do not expect significant problems with implementation once regulatory authorities understand the potentially enormous clinical impact of these assays.
- Once regulatory and financial issues are resolved, the most efficient workflow for accomplishing the ancillary tests, interpreting them and incorporating them into a diagnostic report that synthesizes morphology, POLE genotype and immunophenotype will need to be defined.
- As accumulating data already suggest modifications to the “high-intermediate risk category” of uterus-confined endometrial carcinoma based on ProMisE/TCGA cluster assignment, clinical trials to validate its usefulness for clinical applications will be needed.
- It is uncertain whether it will continue to be necessary to distinguish serous-like (p53 abnormal/copy number high) FIGO grade 3 endometrioid, serous, and clear cell carcinomas.
- Although there are data indicating that the ProMisE classifier reliably assigns matched tumors from preoperative and hysterectomy specimens to the same genomic group,<sup>39,40</sup> it is uncertain whether preoperative ProMisE/TCGA classification (using biopsy or curettage material) can be used to triage patients to different types of staging surgery. For example, patients with POLE-mutant tumors might not need comprehensive surgical staging. Similarly, patients with serous and serous-like carcinomas may not benefit from comprehensive surgical staging, since nearly all will require adjuvant chemotherapy, and it is possible that comprehensive surgical staging may only be applicable to the remaining categories.

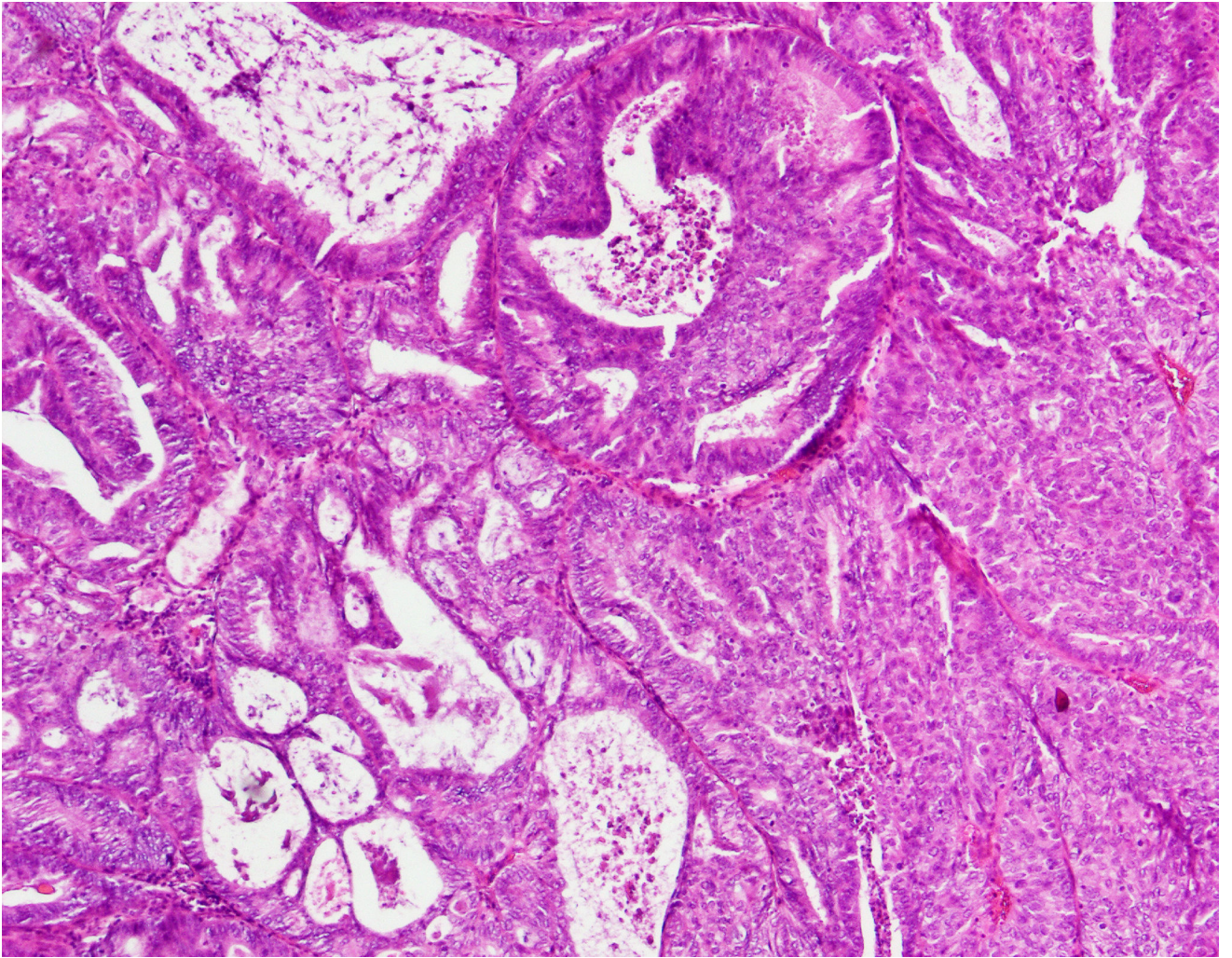


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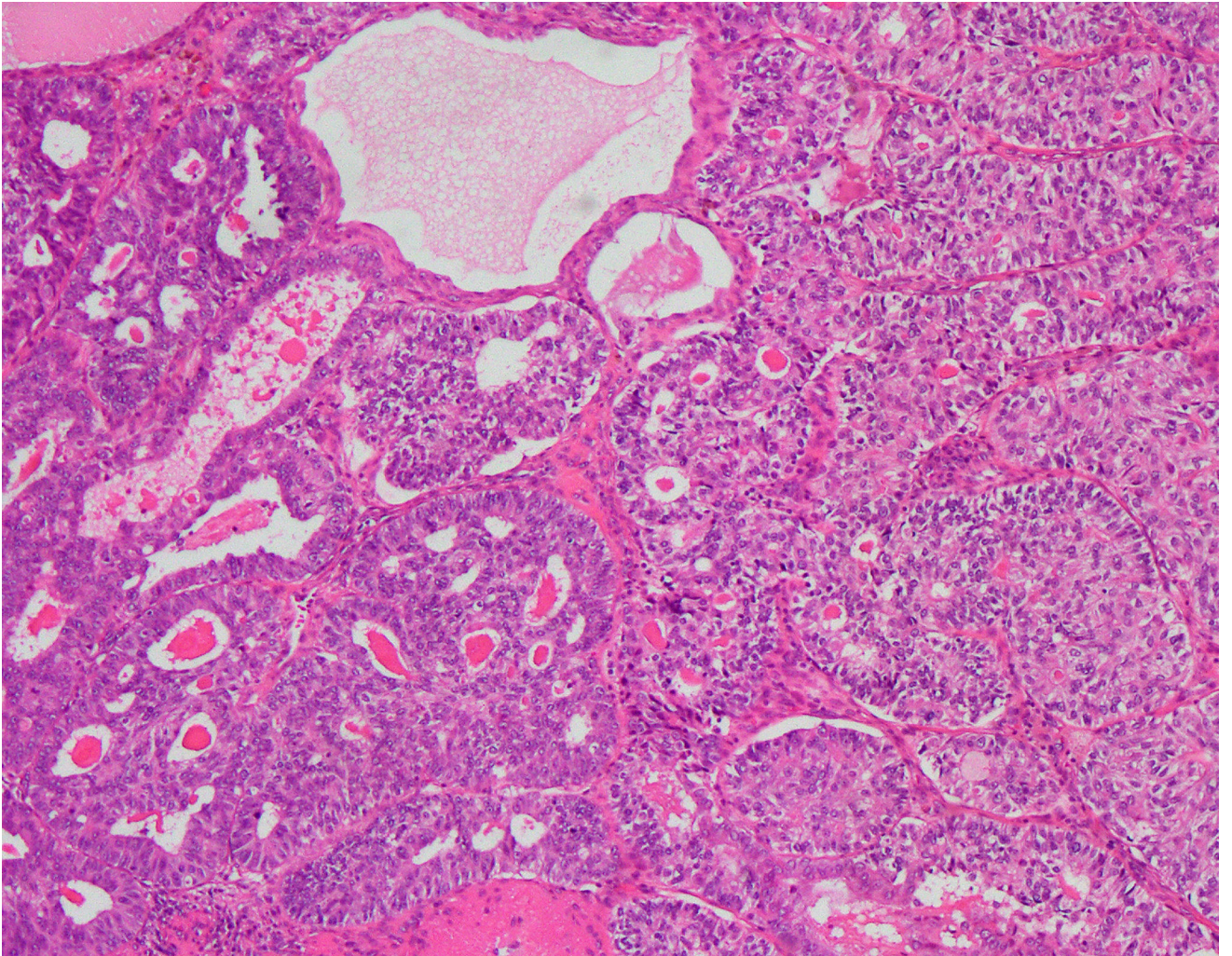
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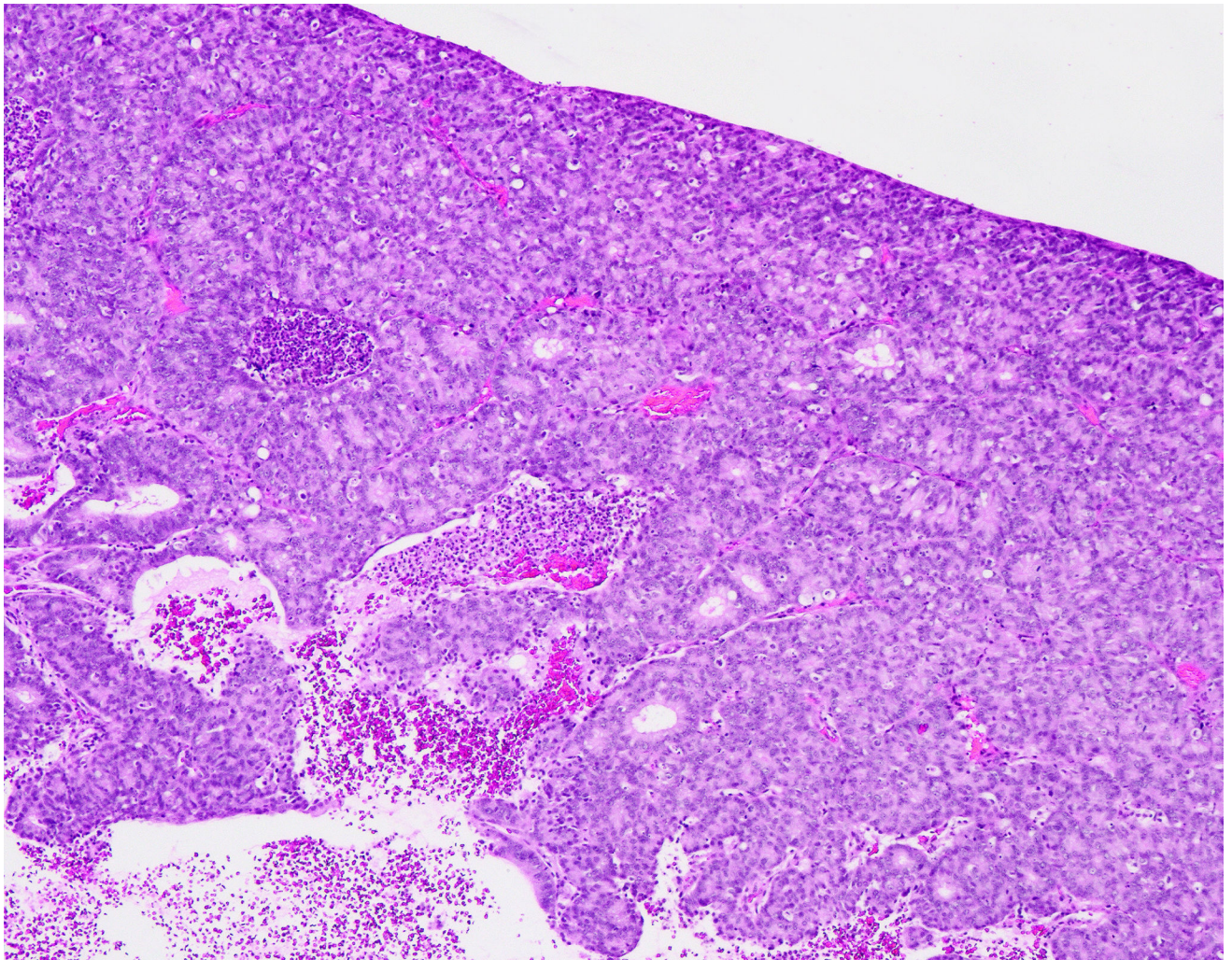


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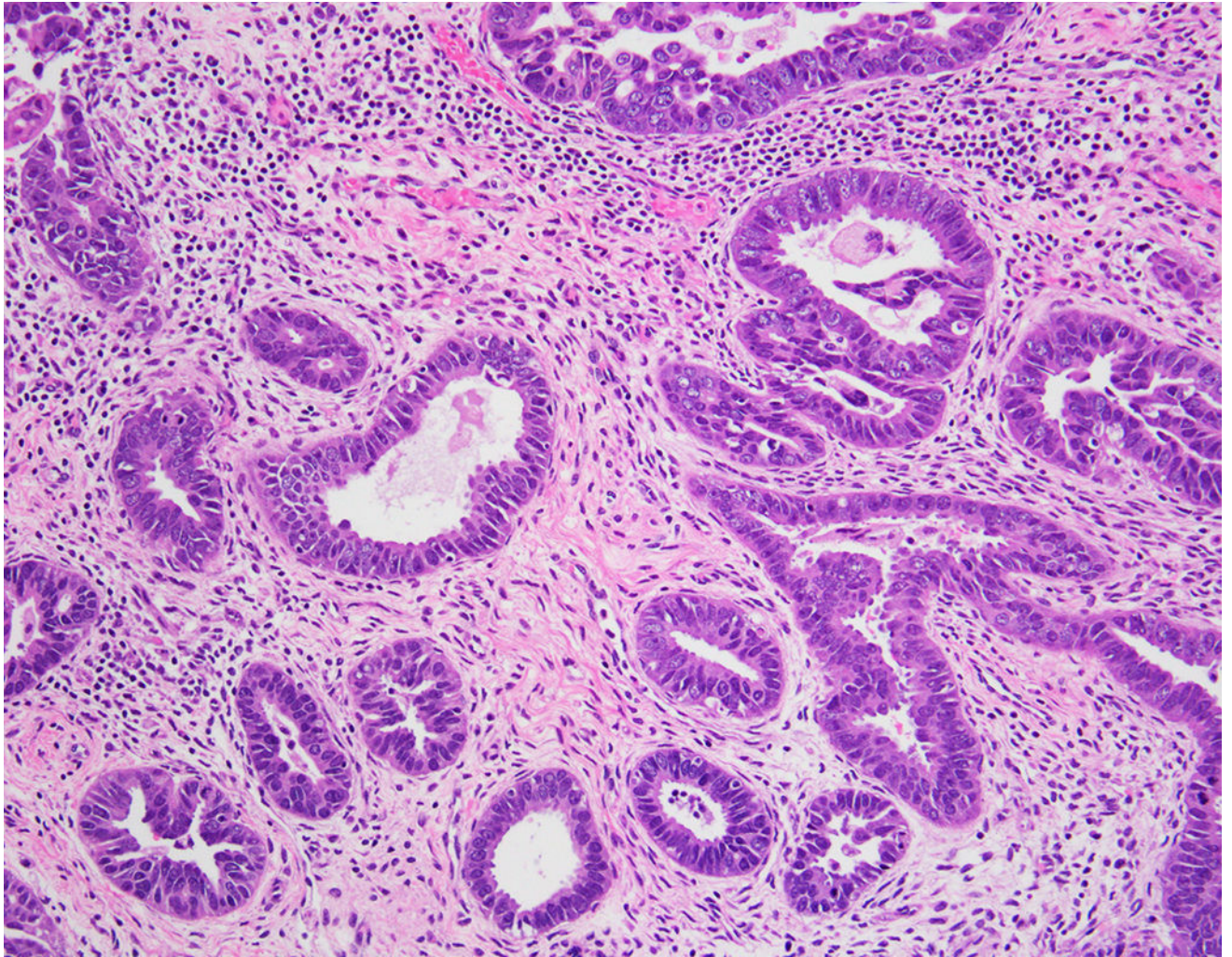
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**Figure 1.**

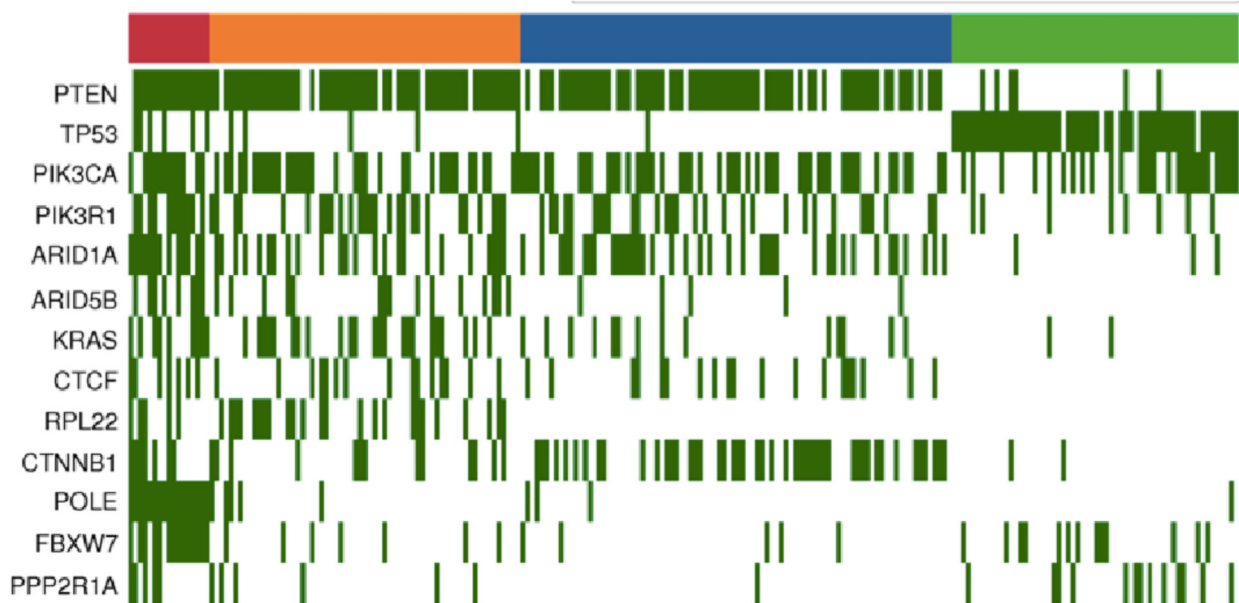
Examples of low- and high-grade endometrial endometrioid carcinomas (EEC): A) Low-grade EEC (International Federation of Gynecology and Obstetrics [FIGO] grade 1) with extensive squamous differentiation that does not qualify as “solid” for the purposes of grading; B) Low-grade EEC (FIGO grade 2) with <50% solid non-squamous growth; C) Low-grade EEC (FIGO grade 2) with <50% solid non-squamous growth; and D) High-grade EEC with a microacinar growth pattern that qualifies as “solid growth.” The presence of microacini should not be considered “glandular” for the purposes of assigning binary or FIGO grade.



**Figure 2. Serous carcinoma with a glandular pattern**

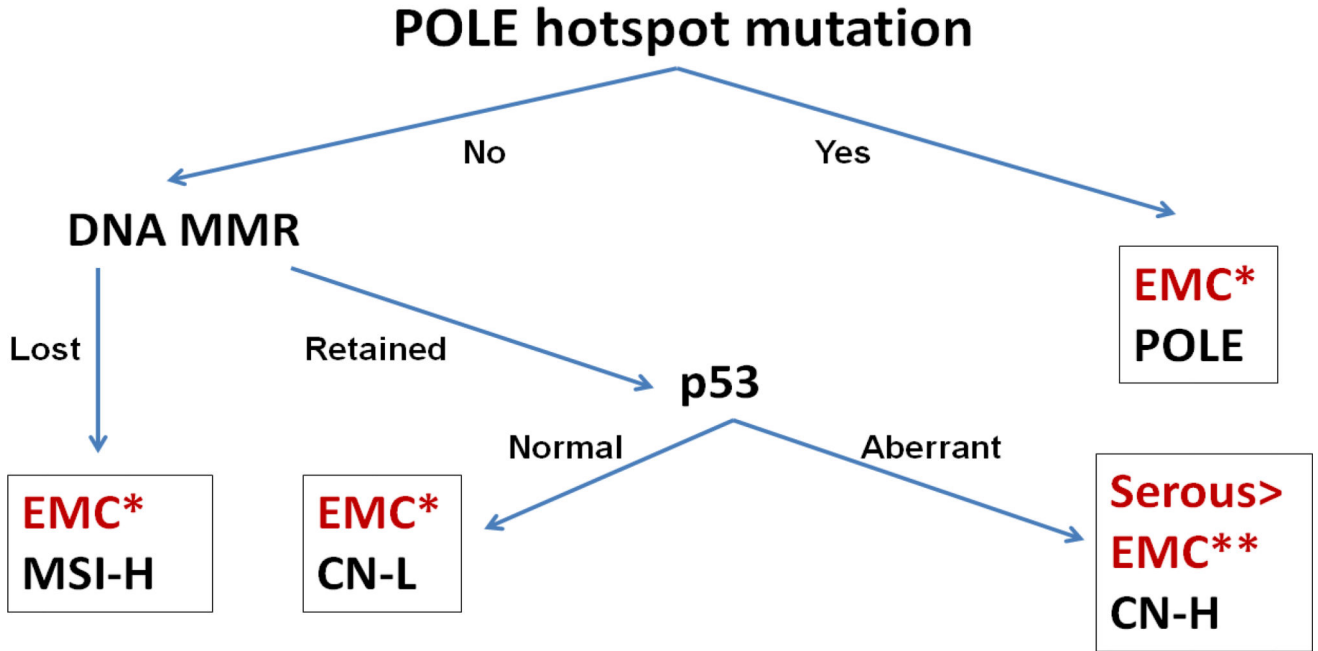
Serous differentiation can be recognized at intermediate-power magnification by the presence of cells with high nuclear-to-cytoplasmic ratios and at high power by the presence of enlarged and atypical nuclei. Confirmatory endometrioid features are lacking. The diagnosis can be confirmed with a p53 immunohistochemical stain, which shows mutation-type staining in at least 90% of serous carcinomas. Although mutation-type p53 staining can be present in endometrioid carcinomas, almost all are International Federation of Gynecology and Obstetrics (FIGO) grade 3 carcinomas, which glandular serous carcinomas do not resemble.

# TCGA Mutational Data by Clusters



**Figure 3. The Cancer Genome Atlas (TCGA) mutational data arranged by genomic clusters**  
 Data from TCGA showing that *TP53* mutation analysis is a sensitive and specific method for distinguishing between copy number high and copy number low tumors. Note however that *TP53* mutations are also seen in POLE and microsatellite instability-high (MSI-H) tumors.

# Harnessing TCGA for diagnosis (multimodality algorithm)



**Figure 4. Harnessing The Cancer Genome Atlas (TCGA) data for clinical diagnosis**  
 EMC: endometrial carcinoma; MSI-H: microsatellite instability high; CN-L: copy number low (type I EMC); POLE: polymerase E hotspot mutant tumor; CN-H: copy number high (serous or serous-like endometrial carcinoma)  
 \*EMC: This may also be applicable to clear cell carcinomas. Most cases can be separated with review of hematoxylin and eosin slides; \*\*EMC: This assay does not distinguish CN-H grade 3 endometrioid carcinoma or CN-H clear cell carcinoma from serous carcinoma

**Table 1**

Interobserver and intraobserver variability of different grading systems

<b>Grading system</b>	<b>Interobserver variability*</b>	<b>Intraobserver variability*</b>
FIGO Grade 1–3 <sup>5,10,14–19</sup>	0.41–0.65	0.66–0.73
Binary FIGO <sup>10,11</sup>	0.58–0.71	0.9
Lax <sup>10,11,14</sup>	0.39–0.75	0.67–0.79
Taylor <sup>15</sup>	0.90 (biopsy); 0.97 (hysterectomy)	
Alkushi <sup>10</sup>	0.76	0.81

FIGO, International Federation of Gynecology and Obstetrics

\* kappa statistics

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**Table 2**

Rate of lymph node metastases in clinical stage 1 endometrial carcinoma in terms of preoperative tumor grade and depth of myometrial invasion (DMI) on hysterectomy: apparent differences in nodal positivity rates between grades is lost when adjusted for DMI

	<b>Preoperative Grade 1 (n=828)</b>	<b>Preoperative Grade 2 (n=480)</b>	<b><i>P</i> (Grade 1 vs 2)</b>	<b>Preoperative Grade 3 (n=236)</b>
<b>Total</b>	55 (6.6%)	55 (11.5%)	0.0036	33 (14%)
<b>No myometrial invasion</b>	3/404 (0.7%)	1/147 (0.7%)	1	0/52 (0%)
<b>&lt;50% myometrial invasion</b>	23/329 (7%)	26/232 (11.2%)	0.095	10/119 (8.4%)
<b>&gt;50% myometrial invasion</b>	29/95 (30.5%)	28/101 (27.7%)	0.75	23/70 (32.9%)

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**Table 3**

The four molecular subtypes of endometrial carcinoma

<b>Mutation profile</b>	<b>Surrogate marker (genetic)</b>	<b>Surrogate marker (IHC)</b>
Ultramutated	<i>POLE</i> mutation	none available
Hypermutated	MSI-H	MMR-D
Copy-number high	<i>TP53</i> mutation	p53 abn
Copy-number low	absence of other markers	absence of other markers (p53 wt)

MSI-H, microsatellite instability (high); MMR-D, mismatch repair deficient; abn, abnormal; wt, wild type

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