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(Article begins on next page)

Lymphovascular space invasion in endometrial carcinoma: a prognostic factor independent from molecular signature

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ABSTRACT

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- 34 Background: The 2020 ESGO/ESTRO/ESP guidelines stratify the prognosis of endometrial
- carcinoma (EC) patients combining The Cancer Genome ATLAS (TCGA) molecular signature and
- pathological factors, including lymphovascular space invasion (LVSI). However, little is known about
- 37 the prognostic independence of LVSI from molecular signature.
- 38 **Aim:** To assess whether the prognostic value of LVSI is independent from the TCGA signature.
- 39 Material and methods: A systematic review and meta-analysis was performed by searching 5
- 40 electronic databases from their inception to March 2021. All peer-reviewed studies reporting
- 41 assessing LVSI as a prognostic factor independent from the TCGA groups in EC were included.
- 42 Multivariate HRs with 95% confidence interval (CI) were pooled separately for overall survival (OS),
- 43 disease-specific survival (DSS) and disease-free survival (DFS). The absence of LVSI was
- considered as a reference. In DFS analyses, locoregional and distant recurrence were separately
- 45 considered for one study.
- Results: Six studies with 3,331 patients were included in the systematic review and three studies
- with 2,276 patients in the meta-analysis. LVSI showed a pooled multivariate HR of 1.818 (CI 95%,
- 48 1.378-2.399) for OS, 1.849 (CI 95%, 1.194-2.863) for DSS, 1.377 (CI 95%, 1.008-1.880) for DFS
- 49 excluding one study, 1.651 (CI 95%, 1.044-2.611) for DFS additionally considering locoregional
- recurrence from one study, and 1.684 (Cl 95%, 1.05-2.701) for DFS additionally considering distant
- recurrence from the same study.
- 52 Conclusion: LVSI has a prognostic value independent of TCGA signature, as well as age and
- 53 adjuvant treatment, increasing the risk of death of any cause, death due to EC and recurrent or
- 54 progressive disease by 1.5-2 times.
- **KEYWORDS:** cancer; tumor; endometrium; prognosis; treatment; risk assessment; PROMISE.

INTRODUCTION

- In 2020, endometrial carcinoma (EC) was the most common gynecological cancer in Europe, with
- an incidence of 31.2% and a 5-year prevalence of 34.1% [1]. In the last decades, EC has showed a
- 300% increase in the number of deaths despite a lower increase in incidence (80%) [1-10]. Such
- data seems due to an inaccurate risk stratification underlying patient management [2,3,10-16].
- In the attempt of improving the risk stratification system, in 2020, the European Society of
- 64 Gynaecological Oncology (ESGO), the European Society for Radiotherapy and Oncology (ESTRO)
- and the European Society of Pathology (ESP) drafted joint guidelines for the management of EC,
- 66 recommending the assessment of molecular signature as additional prognostic factor to be
- 67 integrated with classic pathological factors, such as histotype, International Federation of
- 68 Gynaecology and Obstetrics (FIGO) stage, FIGO grade, myometrial invasion, and lymphovascular
- space invasion (LVSI) [21]. In fact, in 2013, The Cancer Genome Atlas (TCGA) Research Network
- 70 identified four prognostic groups of ECs with a different molecular signature: POLE-
- 71 mutated/ultramutated (POLEmt), microsatellite-instable/hypermutated (MSI), copy-number-
- high/p53-mutated (p53mt), and no specific molecular profile (NSMP) [17]. Afterwards, The Proactive
- 73 Molecular Risk Classifier for Endometrial Cancer (ProMise), which adopted immunohistochemistry
- 74 (cheap, fast and widely available) as a surrogate of sequencing techniques, was demonstrated to
- be prognostically informative and more applicable in clinical practice [2,12,13].
- However, how the molecular signature can be integrated with classic pathological factors is still
- 77 under investigation. In particular, the histotype seems to have a crucial prognostic value
- 78 independently of the TCGA groups, with non-endometrioid carcinomas having a worse prognosis in
- each TCGA group [6]. On the other hand, deep myometrial invasion has shown to affect the risk of
- recurrence independently from the TCGA groups, but not the risk of death of any cause [11]. On the
- contrary, the prognostic independence of LVSI from the molecular signature has not been enough
- 82 investigated.
- The aim of this study was to evaluate whether the prognostic value of LVSI is independent from the
- TCGA group, through a systematic review and meta-analysis.

MATERIALS AND METHODS

Study protocol

- 88 All study steps were defined according to an *a priori* designed protocol and were independently
- 89 performed by two authors. In the case of disagreement, a discussion with a third author was adopted
- 90 as resolution method.
- The study was reported according to the Preferred Reporting Item for Systematic Reviews and Meta-
- 92 analyses (PRISMA) statement and checklist [23].

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Search strategy

- 95 MEDLINE, Web of Sciences, Google Scholar, Scopus and ClinicalTrial.gov were searched from their
- 96 inception to May 2021 through several combinations of the following text words: "endometr*";
- "carcinoma"; "cancer"; "tumor"; "neoplasia"; "malignancy"; "LVSI"; lymphovascular space invasion";
- 98 "lymph*"; "immunohistochem*"; "marker"; "TCGA"; "ATLAS"; "genome"; "ProMisE"; "Proactive
- 99 Molecular Risk Classifier"; "TransPORTEC"; "PORTEC"; "prognosis"; "survival"; "hypermutated";
- "mismatch repair"; "microsatellite instability"; "MMR"; "MSI"; "MLH1"; "MSH2"; "MSH6"; "PMS2";
- "EPCAM"; "ultramutated"; "POLE"; "copy number"; "TP53"; "tumor protein 53"; "p53"; "ESMO",
- "ESGO", "ESTRO", "ESP". All references from full-text screened studies were assessed for eligible
- 103 studies.

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Study selection

- Peer-reviewed studies with extractable data about LVSI as a prognostic factor independent of the
- 107 TCGA groups in EC were included. In detail, we selected all studies reporting multivariate hazard
- ratios (HR) for LVSI and assessing the TCGA group as a variable in the multivariate analysis. Case
- reports and reviews were excluded. In the cases of overlapping data between two studies (i.e. same
- period of enrollment, institution and/or results), only the study with higher sample size was
- 111 considered for the analysis.

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Data extraction

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- Data were extracted without modifications and in accordance with the PICO items [23]. "Population"
- of the systematic review and meta-analysis consisted of women with EC.
- "Intervention" (or risk factor) consisted of the presence of LVSI. For included studies that substratified
- LVSI, "substantial" LVSI was considered as presence of LVSI.
- "Comparator" consisted of the absence of LVSI. For included studies that substratified LVSI,
- "absent" and "mild" LVSI were considered as absence of LVSI.
- "Outcomes" consisted of overall survival (OS) as primary outcome, and disease-specific survival
- (DSS) and disease-free survival (DFS) as secondary outcomes. In particular, OS (or time to death)
- was defined as time from surgery until death of any cause. DSS (or time to death from disease) was
- defined as time from surgery until death due to EC. DFS (or time to recurrence or progression) was
- defined as time from surgery until there is evidence of recurrent or progressive disease (this was
- based on either clinical evidence of recurrence or imaging confirmation of recurrence) or if death
- from the disease occurred prior to the censoring date.

Risk of bias within studies evaluation

- 130 The risk of bias within studies was evaluated following The Methodological Index for Non-
- Randomized Studies (MINORS) [24]. Each included study was judged for 7 applicable domains
- related to risk of bias: 1) Aim (was the aim clear?); 2) Patient selection (were patients randomly or
- consecutively selected?); 3) Prospective collection of data (were data prospectively collected?); 4)
- Endpoints appropriate to the study (were OS, DSS and DFS assessed?); 5) Unbiased evaluation of
- the study endpoints (were endpoints assessed without bias?); 6) Follow-up time (was the follow-up
- at least 2 years?); 7) Lost to follow-up (were patients lost to follow-up less than 5% of the study
- 137 population?).
- Judgments for each domain were the following: "low risk", "unclear risk" or "high risk" of bias based
- on information was "reported and adequate", "not reported" and "reported but inadequate",
- 140 respectively.

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Data synthesis

- For our analyses, we considered multivariable OS, DSS and DFS analyses that assessed the prognostic value of LVSI and considered the TCGA group as a variable from the included studies.
- In particular, all included studies used cox proportional hazard models.
- 148 Pooled HRs between EC women with and without LVSI were calculated based on the above-
- 149 mentioned multivariable analyses, separately for death of any cause, death due to EC, and
- recurrence. The absence of LVSI was considered as a reference. HRs with 95% confidence interval
- (CI) were graphically reported as individual and pooled estimates on forest plots.
- 152 The random effect model of DerSimonian and Laird was used in the analyses.
- Statistical heterogeneity among studies was evaluated by using the inconsistency index I², as
- previously described [25-27]. In particular, heterogeneity was judged as: null for I²=0%, minimal for
- 155 $0 < l^2 < 25\%$, low for $25 \le l^2 < 50\%$, moderate for $50 \le l^2 < 75\%$ and high for $l^2 \ge 75\%$.
- 156 Comprehensive Meta-Analysis (Biostat,14 North Dean Street, Englewood, NJ 07631, USA) and
- Review Manager v. 5.4 (Copenhagen: The Nordic Cochrane Centre, Cochrane Collaboration, 2014)
- were used as software.

RESULTS

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Study selection

Electronic databases searches led to 12,558 studies. Of them, 4,186 remained after duplicates 162 removal and 72 after title screening. Abstract screening led to 17 studies, which were evaluated for 163 eligibility [3,13-15,28-40]. Of them, 11 were excluded because of the absence of multivariable 164 survival analysis for LVSI [13,30,32,34-36] or the absence of LVSI as a prognostic factor 165 [29,31,33,37,38]. Therefore, 6 studies with 3,331 EC women were included in the systematic review 166 [3,14,15,28,39,40]. Lastly, 2 studies were excluded from the meta-analysis due to overlapping 167 patient data [3,40] with another included study [39], while one study was excluded because 168 multivariate survival analysis was reported only for the NSMP group [14]. In conclusion, three studies 169 with 2,276 EC women were included in the meta-analysis [15,28,39] (Figure S1). 170

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Included studies and patients

- Each included study performed *ad hoc* analyses on samples from previous cohorts of EC patients.
- In particular, the cohort was retrospective for 4 studies [3,14,39,40], prospective for one study [28],
- and provided from a randomized controlled trial for the latter one [15].
- No included study adopted pathological criteria for patient selection, except for two studies [15,28]
- which only assessed endometrioid ECs (Table S1).
- 178 Our study population had a age which ranged from 41 to 90 years, and a mean follow-up which
- ranged from 60.6 to 131 months. ECs had endometrioid histotype in 85.7% of cases, International
- Federation of Gynecology and Obstetrics (FIGO) grade 3 in 27.3%, LVSI in 22.3%, and FIGO stage
- 181 I in 73.8%. Women underwent adjuvant treatment in 57.7% of cases (Table S2). Regarding TCGA
- groups, 6.8% of ECs were POLE-mt, 29.3% were MMR-d, 50% were NSMP, and 13.8% were p53-
- 183 abn (Table S3).

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Risk of bias within studies evaluation

- All included studies were at low risk of bias for each domain, except for the "Endpoints appropriate
- to the study" and "Unbiased evaluation of the study endpoints" domains.

In detail, the study by Stelloo et al. [15] was judged at unclear risk of bias for the "Endpoints appropriate to the study" because it did not consider DSS as a study endpoint and assessed separately locoregional and distant recurrence for DFS. On the other hand, the study by Kommoss et al. was judged at high risk of bias because multivariate survival analysis was reported only for the NSMP group [14].

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Data synthesis

Three out of six studies were included in the meta-analysis [15,28,39], as one study was excluded due to the high risk of bias in the "Unbiased evaluation of the study endpoints" domain [14], and two studies [3,40] were excluded due to overlapping data with another study [39]. In particular, all three studies were suitable for OS analysis, while 2 were suitable for DSS and DFS [3,14,28,39,40]. In detail, the study by Stelloo et al [15] was excluded from DSS analysis because it did not consider DSS as a study endpoint, while it was included in additional DFS analysis because it assessed separately locoregional and distant recurrence. Therefore, DFS analyses consisted of 3 separate analyses: one excluding the study by Stelloo et al, one including such study as locoregional recurrence, and one including it as distant recurrence.

Pooled HR of LVSI was:

- 1.818 (CI 95%, 1.378-2.399; p=0.0; I²=0%) for OS (Figure 1);
- 1.849 (CI 95%, 1.194-2.863; p=0.006; I² not calculable) for DSS (Figure 2);
- 1.377 (Cl 95%, 1.008-1.880; p=0.044; l² not calculable) for DFS excluding the study by Stelloo et al. [15] (Figure 3);
- 1.651 (Cl 95%, 1.044-2.611; p=0.032; l²=45.7%) for DFS including the study by Stelloo et al. [15] as locoregional recurrence (Figure 4);
- 1.684 (Cl 95%, 1.05-2.701; p=0.03; l²=51.5%) for DFS including the study by Stelloo et al.
 [15] as distant recurrence (Figure 5).

DISCUSSION

- This study shows that, in EC patients, LVSI has a prognostic value independent of TCGA groups,
- as well as age and adjuvant treatment. In particular, the presence of LVSI increased the risk of death
- of any cause, death due to EC and recurrent or progressive disease by 1.5-2 times. These findings
- 218 may be the first pooled data in the field.
- 219 Since President Obama's 2014 State of the Union address [41,42], Precision Medicine initiative has
- been essential in oncological studies [42,43]. Precision medicine appears as the most promising
- approach to cancer, recommending measures tailored to the individual, with the paradigm "the right
- therapy, at the right time, in the right patient..." [44]. In fact, rather than classifying tumors from the
- organs in which they originated, they may be classified based on the molecular signature [42].
- Among all cancers, this approach has provided the most promising results in EC, where TCGA
- findings have shown the potential to subvert the negative epidemiological trend observed in the last
- two decades and due to an inaccurate risk assessment [1-11,13,17].
- On these bases, in the attempt to improve and tailor the risk stratification of EC women, the 2020
- 228 ESGO/ESTRO/ESP guidelines for the management of patients with EC recommend to integrate
- classic pathological prognostic factors with TCGA molecular signature [21]. However, the allocation
- of a patient to a specific risk group based on such an integration should presuppose prognostic
- independence between factors [11]. To date, while the prognostic independence of molecular
- signature from pathological factors appears demonstrated [3,13-17,28-40], the opposite has not
- 233 been well-assessed yet. In detail, while data sustaining the at least partial prognostic independence
- of histotype and deep myometrial invasion have been recently provided [6,11], little is known about
- 235 LVSI, with only few studies assessing its impact in a specific TCGA group. In particular, while LVSI
- was not associated with an increased risk of tumor recurrence or progression [32] and death from
- disease [45] in POLE-mt ECs, it appeared as an independent predictor of poor outcome in the MSI
- group [46,47]. Lastly, in the study by Stelloo et al., LVSI was the only pathological prognostic factor
- independent of TCGA groups for recurrence and death of any cause [15].
- 240 Regarding the integrated risk stratification proposal, the 2020 ESGO/ESTRO/ESP guidelines
- 241 upgrade the risk group from low or intermediate to high-intermediate for stage I MMRd/NSMP
- 242 endometrioid ECs based on the presence of LVSI. Therefore, LVSI status would affect the
- recommendation of adjuvant treatment in these patients. On the contrary, no role has been attributed
- to LVSI in the risk stratification of POLE-mt and p53-mt ECs [21].
- Our results seem to support the use of LVSI as additional prognostic factor to be integrated with
- TCGA signature. The integration of molecular signature and pathological factors would ensure a
- 247 more tailored management of patients in accordance with the principles of the precision medicine.

On the one hand, this might reduce the under- and overtreatment of EC women observed in the last years [4,6,8,11]; on the other hand, it might reduce the risk of wrong conclusion from oncological trials as patients with different prognosis would not be lumped together based on histological examination alone.

However, in order to further confirm 2020 ESGO/ESTRO/ESP recommendations and to obtain a more and more tailored management of EC patients, additional studies are needed to separately assess the prognostic value of LVSI in each TCGA group. This is necessary as each TCGA group is differently affected from pathological factors, with the POLEmt group appearing as the group least affected [4]. Indeed, the impossibility to provide pooled data about LVSI prognostic independence separately in each TCGA group may be the major limitation of our study. In addition, there was only a small number of studies which were suitable for our analysis. Another point to be clarified is the criterion to categorize LVSI. In fact, in one of the included studies, as well as in the ESGO/ESTRO/ESP guidelines [21], LVSI was dichotomized as "absent/focal" vs "substantial" [15,21]; such system was also showed to work in cervical cancer [48]. By contrast, two of the included studies dichotomized LVSI as "absent" vs "present" [28,39]. Lastly, we could not assess the prognostic independence of LVSI from other histological factors, as only TCGA group, age and adjuvant treatment were considered in the multivariate analysis of all individual studies. These limitations highlight the great need for further studies in the field.

CONCLUSION

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- In EC patients, LVSI has a prognostic value independent of TCGA signature, as well as age and adjuvant treatment, increasing the risk of death of any cause, death due to EC and recurrent or progressive disease by 1.5-2 times.
- Further studies are necessary to confirm these findings and to assess the prognostic impact of LVSI separately in each TCGA group.

CONTRIBUTION AR and AT independently assessed electronic search, eligibility of the studies, inclusion criteria, risk of bias, data extraction and data analysis. DR, DN, MM, PC and AS contributed to the elaboration of methods for risk of bias assessment, data extraction and analysis. AR, AT, DN, LI, FZ, GFZ, AM and RS conceived the study; DR, FR, PC, AS and LI worked on the design of the study; AR, AT, DN, DR, FR and LC worked on the manuscript preparation; PC, LI, FZ, GFZ, AM and RS supervised the whole study. **CONFLICT OF INTEREST STATEMENT** Authors report no conflict of interest. **FUNDING INFORMATION** No financial support was received for this study.

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LEGENDS FOR TABLES AND FIGURES

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427 Figure 1. Forest plot of individual and pooled hazard ratios for death of any cause between 428 endometrial carcinoma patients with and without lymphovascular space invasion; multivariable 429 analyses including TCGA group as a variable were considered from included studies. The absence 430 of lymphovascular space invasion was considered as a reference. 431 432 Figure 2. Forest plot of individual and pooled hazard ratios for death due to endometrial carcinoma 433 between patients with and without lymphovascular space invasion; multivariable analyses including 434 TCGA group as a variable were considered from included studies. The absence of lymphovascular 435 436 space invasion was considered as a reference. 437 438 Figure 3. Forest plot of individual and pooled hazard ratios for recurrent or progressive disease between endometrial carcinoma patients with and without lymphovascular space invasion; 439 440 multivariable analyses including TCGA group as a variable were considered from included studies. The absence of lymphovascular space invasion was considered as a reference. The study by Stelloo 441 et al. [15] was not included. 442 443 Figure 4. Forest plot of individual and pooled hazard ratios for recurrent or progressive disease 444 445 between endometrial carcinoma patients with and without lymphovascular space invasion; 446 multivariable analyses including TCGA group as a variable were considered from included studies. 447 The absence of lymphovascular space invasion was considered as a reference. Locoregional 448 recurrence was considered for the study by Stelloo et al. [15]. 449 Supplementary Figure 1. Flow diagram of studies identified in the systematic review (Prisma 450 451 template [Preferred Reporting Item for Systematic Reviews and Meta-analyses]). 452 Supplementary Figure 2. a) Assessment of risk of bias. Summary of risk of bias for each study; 453

Plus sign: low risk of bias; minus sign: high risk of bias; guestion mark: unclear risk of bias; b) Risk

of bias graph about each risk of bias item presented as percentages across all included studies.

Supplementary Table 1. Characteristics of the studies included in the qualitative synthesis. 456 FIGO: International Federation of Gynaecology and Obstetrics 457 **RCT:** Randomized controlled trial 458 459 **OS:** overall survival **DFS:** disease-free survival 460 DSS: disease-specific survival 461 462 *: DFS was assessed as locoregional recurrence-free survival and distant recurrence-free survival 463 Supplementary Table 2. Characteristics of patients from studies included in the qualitative 464 465 synthesis. FIGO: International Federation of Gynecology and Obstetrics 466 LVSI: Lymphovascular space invasion 467 *: In the Stelloo et al. study, LVSI was assessed as absent/mild or substancial 468 469 470 Supplementary Table 3. Distribution of TCGA groups in the studies included in qualitative synthesis. 471 472 Supplementary Table 4. List of variables considered at multivariable analyses in the studies 473 474 included in the qualitative synthesis. FIGO: International Federation of Gynaecology and Obstetrics 475 L1CAM: L1 Cell Adhesion Molecule 476