Assessing the prognostic value of PAX2 and PTEN in endometrial carcinogenesis

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Abstract

In order to avoid the consequences of over- and under-treatment of endometrial hyperplasia, diagnostic accuracy and progression risk assessment must be improved. The aim of this study was to assess whether PAX2 or PTEN expression could predict progression-free survival in endometrial intraepithelial neoplasia (EIN) and endometrial endometrioid carcinoma (EEC). Immunohistochemistry for detection of PAX2 and PTEN was performed on 348 endometrial samples; 75 proliferative endometrium (PE), 36 EIN and 237 EEC. Cases classified as PTEN null (1 or more glands negatively stained) were more prevalent in EEC than in PE and EIN (64% EEC vs 11% PE/EIN). A progressive decrease in PAX2 expression was observed from PE to EIN to EEC. Long-term clinical follow-up (6–310 months, median: 126) was available for 62 PE cases, all 36 EIN cases and 178 EEC cases. No patients with PE demonstrated progression to EIN or EEC. Progression of disease was observed in 10 (28%) EIN patients. These patients had significantly lower PAX2 expression than those that regressed (P = 0.005). Progression-free survival analysis revealed that EIN patients with a high-risk PAX2 expression score (H-score ≤ 75) had a higher probability of progression of disease in comparison to those with a low-risk score (H-score >75). PAX2 expression was not prognostic in EEC nor was PTEN status of prognostic value in either EIN or EEC. PAX2 expression analysis by means of H-score has prognostic potential for the identification of high-risk progression cases in EIN but needs to be validated in a larger cohort.

Key Words
- endometrial endometrioid carcinoma
- endometrial intraepithelial neoplasia (EIN)
- PAX2
- PTEN
- prognostic biomarkers

Introduction

Endometrial cancer is currently the most frequent form of gynaecologic cancer in developed countries (http://globocan.iarc.fr/Pages/fact_sheets_population.aspx, Accessed: 28 February 2018). Endometrial endometrioid cancer (EEC) is often preceded by a premalignant lesion termed endometrial hyperplasia (Bokhman 1983). These lesions have been previously categorised by the WHO94 system according to the degree of glandular complexity and nuclear atypia; however, the disadvantages of its use in the clinic has been greatly debated citing its poor
reproducibility and inadequate predictive capability (Skov et al. 1997, Kendall et al. 1998, Bergeron et al. 1999). The newer EIN classification system, recently adopted by the World Health Organisation (WHO) in its 2014 publication (Kurman et al. 2014), demonstrates improved reproducibility and clinical predictive ability compared to the older scheme (Baak et al. 2005a, Hecht et al. 2005, Mutter & Group TIENW 2000, Usubutun et al. 2012). The EIN classification distinguishes two forms of endometrial hyperplasia: benign reactive hyperplasia, which is a polyclonal, diffuse lesion that is a result of hormone imbalance and oestrogen hyper-stimulation, and endometrial intraepithelial neoplasia (EIN), defined as a monoclonal, neoplastic lesion that has a high risk of progression to EEC (Mutter & Group TIENW 2000).

Of note, the EIN system more clearly distinguishes high progression risk categories from low risk compared to the WHO94 system, especially when EIN is diagnosed by computerised morphometric analysis or D-score (Baak et al. 2005a).

With the ability to better distinguish between high-risk and low-risk lesions; a clearer prognosis can result in improved allocation of the most appropriate treatment. Despite an EIN diagnosis predisposing to an increased risk of EEC, some patients will never develop EEC, and likewise some rare non-EIN cases can develop EEC. For premenopausal women wishing to preserve fertility and high-risk surgery patients, hysterectomy may be considered as over-treatment. However, as the chances of concurrent carcinoma is high and D-score diagnosed EIN are reported to have a 45-fold risk of progressing to EEC (Baak et al. 2005a, Kurman et al. 2014), progestin treatments and monitoring may result in under-treatment. It would therefore be beneficial to be able to ascertain which cases are likely to progress to reduce the likelihood of over and under-treatment, particularly for these patients. Currently, the EIN classification alone is unable to 100% accurately predict which endometrial cases will progress to cancer. To achieve optimal management of patients, discrimination between lesions that should be handled differently is crucial. There is therefore precedence for the establishment of accurate prognosticators to distinguish between low and high progression risk EIN.

Considerable research has been conducted searching for prognostic biomarkers to predict progression amongst EIN and EEC cases. One of the most commonly investigated biomarkers implicated in endometrial carcinogenesis is PTEN and of recent interest PAX2. The tumour suppressor gene PTEN (10q23.31) is found inactivated in up to 83% of EEC (Mutter et al. 2000). It encodes the PTEN protein, a lipid phosphatase which functions as a negative regulator of the PI3K/AKT pathway (Kurose et al. 2001). The PAX2 gene (10q24.31) encodes PAX2 (paired-box protein 2), a transcription factor active in embryogenesis and promotion of cell proliferation (Shang 2007). Oestrogen- and tamoxifen-induced hypomethylation of the PAX2 promoter has been demonstrated in endometrial carcinogenesis (Wu et al. 2005). Several immunohistochemical studies have shown that loss of expression of both PTEN and PAX2 increases from normal endometrium to endometrial hyperplasia to EEC (Mutter et al. 2000, Baak et al. 2005b, Monte et al. 2010, Pavlakis et al. 2010, Allison et al. 2012). Furthermore, combined loss of PTEN and PAX2 in endometrial glands has been described as an early event and is suggested to promote neoplastic transformation, although they are activated independently (Mutter et al. 2000, Monte et al. 2010, Allison et al. 2012). Notably, in one study of 103 hyperplasias, no morphometrically high-risk (i.e., D-score <1) EIN lesions with PTEN-positive glands developed cancer in a long follow-up, contrasting to a high cancer incidence in PTEN negative, D-score <1 cases (Baak et al. 2005b). Thus, in EIN, D-score and PTEN might have additional prognostic value. This suggests a combinatorial approach to deduce prognosis could be promising. PAX2 expression has been reported to be diminished in EEC (Strissel et al. 2008, Monte et al. 2010, Allison et al. 2012) but one study has demonstrated increased expression in EEC (Kahraman et al. 2012). No studies have, to the best of our knowledge, yet commented on whether PAX2 is associated with progression-free survival in EEC. Therefore, a thorough prognostic evaluation of the additive prognostic value of PAX2 would be beneficial.

In our investigation, we used strictly optimised and standardized immunohistochemistry to observe expression patterns of PTEN and PAX2 in proliferative endometrium (PE), EIN and EEC. We aimed to assess whether PAX2 and PTEN expression has the potential to predict progression-free survival in a large EEC study population with long-term follow-up, in addition to its potential to predict progression of EIN to EEC. We aimed to review their potential as prognostic biomarkers in order to improve low-risk and high-risk progression categorisation so as to avoid the risks of over- and under-treatment of EIN and EEC.
Materials and methods

Study population

This study was approved by the Regional Ethics Committee of Health West Norway (2010/2464). None of the patients included in this study were required to provide written informed consent to participate. All insights in a patient’s journal were monitored electronically, and all, except the treating physician, were required to state the reason why they needed to read that patient’s journal. This log was always open for the patient to view. A total of 348 cases were selected from a vast database, which includes 5494 patients each with one or more endometrial histology samples from the archives of the Department of Pathology, Stavanger University Hospital, Stavanger, Norway, dating from 1977 to 2004 (Steinbakk et al. 2009). Cases were carefully and independently reviewed by two pathologists (J B, E G) using strict microscopic criteria and classified as PE, EIN or endometrioid type endometrial carcinoma (EEC). In case of discrepancies in diagnosis between the pathologists, a consensus meeting with a multi-head microscope was used to decide on a final diagnosis. In total, 75 cases with the diagnosis PE, 36 cases with EIN and 237 with ECC were included in the current study (Table 1). Formalin-fixed paraffin-embedded pipeile/curettage/hysterectomy samples were available for all these cases. All endometrial carcinoma samples were available as tissue microarrays (TMA) created as described by Steinbakk et al. (2009). Histologic sections 4μm thick underwent standard haematoxylin and eosin (H&E) staining. All endometrial cancer samples were classified according to FIGO stage (Table 1) whilst all EIN samples were classified according to WHO14 guidelines. Inclusion criteria for retrospective analysis required follow-up of at least 6 months. Follow-up data were available for 62 of the 75 PE cases (median: 131, range: 6–202 months), all 36 EIN cases (median: 128, range: 6–310) and 178 EEC cases (median: 138, range: 13–275) and 178 EEC cases (median: 138, range: 13–275) and 178 EEC cases (median: 138, range: 13–275). Follow-up data were available for 62 of the 75 PE cases (median: 131, range: 6–202 months), all 36 EIN cases (median: 128, range: 6–310).

Immunohistochemistry

The protocols for H&E staining and immunohistochemistry were performed as outlined previously (Baak et al. 2005b).

Histologic sections were cut to 4μm adjacent to H&E sections. Sections were mounted on silanised slides (Dako) and dried overnight at 37°C. The slides were incubated at 60°C for 1 h and then deparaffinised in xylene followed by rehydration in decreasing concentrations of alcohol. For PAX2 and PTEN, antigen retrieval was performed in the following conditions: 120°C, 1.9 bar in 10mmol/L TRIS/1 mmol/L EDTA (pH 9.0) for 2 min then cooled for 15 min. Immunostaining was performed using an automated stainer from Dako with Tris-buffered saline (S1968) 0.05% Tween 20 (pH 7.6) as a wash buffer. Endogenous peroxidase activity was blocked with a peroxidase blocking reagent (S2001, Dako) for 10 min. The slides were then incubated for 30 min with a PTEN monoclonal antibody (1:300, clone 6h2.1, Cascade Biosciences, Winchester, MA, USA) or PAX2 polyclonal antibody (1:200, cat. No 71-60000, Invitrogen). A peroxidase/DAB system (ChemMate Envision Kit, Dako) was used to visualise the immune-complex; slides were incubated in Envision/HRP secondary antibodies for 30 min and thereafter 10 min in DAB-chromogen. Sections were then counterstained with haematoxylin, dehydrated and mounted. Immunostaining of adjacent normal endometrial tissue and positive immunostaining of endometrial stroma, for PTEN, were considered as internal controls.

Evaluation of immunohistochemical results

For each sample with successful staining a single area, between 2 × 2 mm and 5 × 5 mm in size, was selected for evaluation of PTEN and PAX2. For cases diagnosed as PE, a representative area was chosen by one of the pathologists (J B, E G) and scored by two independent observers (A E V, E J). For cases diagnosed with EIN, an experienced pathologist (J B) marked an area of the section with the most high-graded lesion for evaluation. For endometrioid endometrial cancer (EEC) samples, the TMA’s contained the most poorly differentiated, well-preserved tumour areas and therefore each individual TMA was considered as a single measurable area. Two samples were taken from each EEC case for TMA and their mean score was used.

Table 1 Overview of the number of cases in each diagnostic category with successful PTEN and PAX2 staining.

<table>
<thead>
<tr>
<th></th>
<th>PE* (N=75)</th>
<th>EIN* (N=36)</th>
<th>EEC* (FIGO stage) (N=237)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of cases with successful staining according to diagnostic category</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>PTEN</td>
<td>71/75</td>
<td>35/36</td>
<td>23/31 86/122 35/39 8/9 11/14 8/9 0/1 5/6 1/6 177/237</td>
</tr>
<tr>
<td>PAX2</td>
<td>75/75</td>
<td>36/36</td>
<td>25/31 89/122 36/39 8/9 11/14 8/9 0/1 5/6 1/6 183/237</td>
</tr>
</tbody>
</table>

*Proliferative endometrium; †endometrial intraepithelial neoplasia; ‡endometrioid endometrial carcinoma.
for analysis. Two trained observers (A E V, E J), blinded to the outcome, independently evaluated and scored each case using a light microscope (objective 40×, field diameter 450μm at specimen level). PTEN staining was determined as positive or negative by presence or absence of nuclear or cytoplasmic staining respectively; stromal PTEN staining was used as a positive control. Sections containing one or more PTEN-null glands within the marked area were scored as negative (J B), as previously described (Steinbak et al. 2009). Sections stained with PAX2 were scored using the H-score method. Glandular nuclei could be assigned a score of 0–3 according to staining intensity (0 = negative, 1 = weak positive, 2 = positive, 3 = strongly positive). The observers would evaluate the marked area and deduce the percentage of glandular nuclei at each staining level. Each staining score was then multiplied with the respective percentage, and these were then added together to result in an H-score that ranged from 0 to 300. If PAX2 H-score between the two observers differed by more than 50 a consensus meeting with a multi-head microscope was used to decide on a final score. If the PAX2 score differed by less than 50 points the average of the two scores determined the final score.

Computerised image analysis

The computerised morphometric analysis termed D-score was performed, as previously described (Baak et al. 2001), on H&E-stained sections using the QPRODIT system (Leica). Lesions scored as D-score <1 were defined as high-risk for progression whilst those scored as D-score ≥1 were deemed as low risk. D-score was performed on 35 of 36 EIN cases, where one lesion did not meet the size criteria.

Statistical analysis

Descriptive statistics and cross tables were used to control database quality, to identify any unlikely values or combinations. Regression (no change in diagnosis for PE cases) or progression of endometrial disease was used as the measured endpoints. Regression for EIN and EEC was defined as cured, no recurrence at final follow-up. For cases with an initial diagnosis of PE, progression was defined by a diagnosis of EIN, EEC or EEC-related death in the follow-up. For cases with an initial diagnosis of EIN, progression was defined by a diagnosis of EEC or EEC-related death in the follow-up. For the purposes of this study, EIN cases that persisted in the follow-up, with a lower D-score compared to the original sample, were grouped together with progression cases. For cases with an initial diagnosis of EEC, progression was defined by a diagnosis of higher stage disease or EEC-related death in the follow-up; patients diagnosed with higher stage disease gave the same results as those who succumbed to EEC and were therefore grouped together in this study. Patients who were either lost to follow-up or suffered death from other non-EEC-related causes were censored to last known follow-up date as alive, no evidence of disease. SPSS (SPSS) for Windows, version 23.0 was used for all analyses. A one-way ANOVA was used to observe differences between the three initial diagnostic categories, and an independent sample t-test was used to compare progression and regression patients within each category, according to age, follow-up and PAX2. A chi-square test was used to test for significant differences between progression and regression patients within each category. For survival analysis, Kaplan–Meier survival curves were created and significant differences between groups were tested using the log-rank test. The endpoint was progression-free survival. Data were considered significant at P<0.05.

Results

Age distribution of patients varied between the groups. For patients diagnosed with PE the mean age was 38 (range 25–50), for EIN 46 years (range 24–87) and for EEC 66 years (range 32–92) (Table 2). The lesion of interest was lost in one case for EIN and four cases for PE for PTEN staining (Table 1). Of the 237 EEC samples, PAX2 and

Table 2  Mean age, mean PAX2 score and PTEN status according to endometrial diagnostic category.

<table>
<thead>
<tr>
<th></th>
<th>PE *</th>
<th>EIN §</th>
<th>EEC ‡</th>
</tr>
</thead>
<tbody>
<tr>
<td>N</td>
<td>75</td>
<td>36</td>
<td>183</td>
</tr>
<tr>
<td>Mean age (range)</td>
<td>38 (25–50)§</td>
<td>46 (24–87)§</td>
<td>66 (32–92)§</td>
</tr>
<tr>
<td>Mean PAX2 score (range)</td>
<td>157 (0–245)§</td>
<td>123 (0–300)§</td>
<td>55 (0–215)§</td>
</tr>
<tr>
<td>PTEN % cases</td>
<td>89% pos (63/71)</td>
<td>89% pos (31/35)</td>
<td>34% pos (61/177)</td>
</tr>
<tr>
<td></td>
<td>11% neg (8/71)</td>
<td>11% neg (4/35)</td>
<td>66% neg (116/177)</td>
</tr>
</tbody>
</table>

*Proliferative endometrium; † endometrial intraepithelial neoplasia; ‡ endometrioid endometrial carcinoma; § significant between all groups: One-Way ANOVA, P<0.05.
PTEN staining was successful in 183 (183/237, 77.2%) and 177 (177/237, 74.7%) cases, respectively (Table 1).

Endometrial samples demonstrating a loss of PTEN glandular expression could be readily identified; glands with no PTEN staining were distinct from glands with strong PTEN staining and surrounding stroma (Fig. 1). In PE and EIN, 89% were categorised as PTEN positive and 11% were PTEN negative (Table 2). Whereas in EEC, 34% were classified as PTEN-positive and 66% as PTEN negative (Table 2). No difference in PTEN status was observed between different FIGO stages of EEC (data not shown).

Whilst staining for PTEN allowed for distinct definition of PTEN-null glands, PAX2 exhibited a nuclear staining pattern with variable intensity (Fig. 1). A decreasing trend in PAX2 expression was observed with increasing severity of diagnosis where mean scores decreased from 157 for PE to 123 for EIN to 55 for EEC, which was found to be significant (PE vs EIN: \( P=0.017 \), EIN vs EEC: \( P<0.001 \), PE vs EEC: \( P<0.001 \)) (Fig. 2 and Table 2). There was no observable difference in PAX2 expression between FIGO stages of EEC (data not shown). Scoring of PAX2 had good reproducibility, where 22.4% (41/183) of EEC cases demonstrated conflict between the two observers and a consensus discussion was required.

For 62 of 75 PE cases with known follow-up, none demonstrated progression. Follow-up criteria were met for all 36 patients with an initial EIN diagnosis whilst 178 EEC patients met follow-up criteria. Of the 178 EEC patients with follow-up, all had successful PAX2 staining and 172 had successful PTEN staining.

Of the 36 EIN patients, 26 demonstrated regression of disease, whilst seven patients were diagnosed with EEC. The remaining three patients were diagnosed as persistent EIN at final follow-up. Of note, although still diagnosed with EIN in the follow-up, all three patients had a lower D-score in the follow-up sample in addition to the observation that the glands were more tightly packed and crowded with less overall stroma over a wider area in direct comparison to the original lesion. As a result, these patients were grouped together with the seven progression cases in this study and henceforth these ten patients will be referred to as progression cases. These ten patients were found to have a significantly lower average PAX2 expression score in the original EIN sample in comparison to patients that regressed (\( P=0.005 \)) (Fig. 3). In EIN progression cases, 40% were PTEN negative in the initial sample, whereas 20% of EIN regression cases were PTEN negative; however, this difference was not significant (Table 3). For patients with an initial diagnosis of EEC, no difference was observed between average PAX2 expression scores or PTEN status of patients with or without progression (Table 3). EIN progression risk scores by the D-score method were calculated for 35 of 36 EIN cases. Patients with an initial diagnosis of EIN who later progressed had a significantly lower D-score than cases that demonstrated regression (\( P=0.013 \)) (Table 3).

**Figure 1** Example of an EIN lesion immunostained for the detection of (A and B) PAX2 or (C and D) PTEN protein. The D-score was –0.5. (A) Intensity of PAX2 staining was observed to vary across the lesion. Of note, glands with weak expression and glands with strong expression tended to group within the lesion. (B) A close-up of glands indicated by the box in (A) compares a gland with low expression (top) with a gland with high PAX2 expression (bottom). PAX2 expression could also vary within a single gland with some nuclei strongly expressing PAX2 neighbouring nuclei that did not. (C) This lesion was defined as PTEN-null due to the presence of one or more PTEN-null glands (null glands indicated by absence of PTEN staining). PTEN-negative staining was very distinct from PTEN-positive staining with neighbouring glands showing contrasting expression in some areas. (D) A close-up of the glands indicated by the box in (C) emphasising that PTEN staining was very distinct readily distinguishing PTEN-positive (bottom) glands from PTEN-negative glands (top). Scale bar represents 100 µm. A full colour version of this figure is available at https://doi.org/10.1530/ERC-18-0106.
Age at the time of initial diagnosis was also evaluated for any potential differences between progression and regression cases in EIN and EEC. Only for the EIN category was a significant difference observed between the age of patients that demonstrated regression (mean 42 years) compared to progression (mean 57 years) (P=0.029) (Table 3).

Expanding on the observation that EIN progression patients had a lower mean PAX2 expression score compared to regression patients, a risk threshold was established for PAX2 in order to group patients into progression risk categories. For the purpose of this study, a threshold score of 75 was used to identify patients with low risk of progression: PAX2 >75, and patients with a high risk of progression: PAX2 ≤75. This threshold was based on box plot analysis where 75% of all EIN patients that progressed had a PAX2 score equal to or below 75 (Fig. 3) and where ROC analysis revealed that specificity decreased beyond this threshold (data not shown). The threshold defined 4% of PE cases as high-risk (n=75), 31% of EIN (n=36) and 68% of EEC (n=183). None of the patients with PE developed EIN or EEC. For patients with an initial EIN diagnosis, follow-up revealed that 73% (8/11) of patients with a high-risk PAX2 score progressed and only 8% (2/25) of patients with a low-risk PAX2 score progressed (Table 4). To note, at final follow-up, no EIN patients were dead due to EEC-related or unrelated causes (0/36). For patients in the EEC category where follow-up was available (n=178), 10% (12/121) of EEC patients with a high-risk score demonstrated progression whilst 14% (8/57)
of EEC patients with a low-risk score demonstrated progression (Table 4).

For risk analysis according to PTEN status, we defined PTEN negative as high risk of progression and PTEN positive as low risk of progression due to the observation that PTEN loss occurs at a higher percentage in EEC in comparison to PE and EIN. For patients with an initial diagnosis of EIN, we observed progression in 44% (4/9) of high-risk cases and 23% (6/26) of low-risk cases (Table 4). Whereas for patients with an initial diagnosis of EEC, we observed progression in 13% (3/61) of high-risk cases and 11% (12/111) of low-risk cases (Table 4).

To assess whether the PAX2 threshold or PTEN status had any significant prognostic value in EIN or EEC, we utilised Kaplan–Meier curve analysis to assess progression-free survival. Analysis revealed that high-risk EIN patients classified by PAX2 expression score had a significant lower progression-free survival than low-risk patients (Fig. 4). However, for patients with EEC, there was no difference in progression-free survival between patients with PAX2 low-risk or high-risk status (Table 4). Analysis of PTEN status showed no significant difference in progression-free survival for EIN and EEC patients (Table 4). We also analysed the PAX2 risk threshold in combination with PTEN status and D-score but no additional prognostic value (from PTEN) was observed in either EIN or EEC (Table 4).

Table 4  Overview of the proportion of patients that demonstrated disease progression in either EIN or EEC according to risk classification defined by PAX2 score, PTEN status and/or D-score.

<table>
<thead>
<tr>
<th>Biomarker</th>
<th>Risk status</th>
<th>Percent EIN* patients</th>
<th>Survival analysis</th>
<th>Percent EEC‡ patients</th>
<th>Survival analysis</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>that progressed†</td>
<td>(Kaplan–Meier, log rank)</td>
<td></td>
<td>(Kaplan–Meier, log rank)</td>
</tr>
<tr>
<td>PAX2</td>
<td>≤75</td>
<td>High</td>
<td>73% (8/11)</td>
<td>P=0.001</td>
<td>10% (12/121)</td>
</tr>
<tr>
<td></td>
<td>&gt;75</td>
<td>Low</td>
<td>8% (2/25)</td>
<td></td>
<td>14% (8/57)</td>
</tr>
<tr>
<td>PTEN</td>
<td>Negative</td>
<td>High</td>
<td>44% (4/9)</td>
<td>P=0.268</td>
<td>13% (8/61)</td>
</tr>
<tr>
<td></td>
<td>Positive</td>
<td>Low</td>
<td>23% (6/26)</td>
<td></td>
<td>11% (12/111)</td>
</tr>
<tr>
<td>D-score</td>
<td>&lt;1</td>
<td>High</td>
<td>50% (6/12)</td>
<td>P=0.094</td>
<td>N/A</td>
</tr>
<tr>
<td></td>
<td>≥1</td>
<td>Low</td>
<td>17% (4/23)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>PAX2 PTEN in four risk groups</td>
<td>≤75</td>
<td>High</td>
<td>67% (4/6)</td>
<td>P=0.012</td>
<td>11% (9/81)</td>
</tr>
<tr>
<td></td>
<td>≤75</td>
<td>Positive</td>
<td>80% (4/5)</td>
<td></td>
<td>8% (3/36)</td>
</tr>
<tr>
<td></td>
<td>&gt;75</td>
<td>Uncertain</td>
<td>0% (0/3)</td>
<td></td>
<td>1% (3/30)</td>
</tr>
<tr>
<td></td>
<td>&gt;75</td>
<td>Negative</td>
<td>10% (2/21)</td>
<td></td>
<td>20% (5/25)</td>
</tr>
<tr>
<td>PAX2 D-score in four risk groups</td>
<td>≤75</td>
<td>High</td>
<td>100% (5/5)</td>
<td>P=0.009</td>
<td>N/A</td>
</tr>
<tr>
<td></td>
<td>≤75</td>
<td>&lt;1</td>
<td>50% (3/6)</td>
<td></td>
<td></td>
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<tr>
<td></td>
<td>≥1</td>
<td>≥1</td>
<td>17% (1/6)</td>
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<tr>
<td></td>
<td>&gt;75</td>
<td>&lt;1</td>
<td>6% (1/18)</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>&gt;75</td>
<td>≥1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>PTEN and D-score in four risk groups</td>
<td>≤75</td>
<td>High</td>
<td>67% (4/6)</td>
<td>P=0.388</td>
<td>N/A</td>
</tr>
<tr>
<td></td>
<td>≤75</td>
<td>&lt;1</td>
<td>0% (0/3)</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>≥1</td>
<td>&lt;1</td>
<td>33% (2/6)</td>
<td></td>
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</table>
*Endometrial intraepithelial neoplasia; †progression event for EIN was defined as a diagnosis of persistent EIN with more poorly differentiated glands or diagnosis of EEC in the follow-up; ‡endometrioid endometrial carcinoma; §progression event for EEC was defined as higher stage EEC or death due to EEC.
Discussion

In this study, we have used strictly optimised and standardised immunohistochemistry to observe expression patterns of PTEN and PAX2 in PE, EIN and EEC. A higher proportion of PTEN-negative cases were observed in EEC than in EIN and PE. Furthermore, PAX2 expression decreased progressively from PE to EIN to EEC.

PTEN loss was detected in 11% of PE samples, which is similar to results found in three separate studies (Feng et al. 2012, Lee et al. 2012, Yang et al. 2015). In EEC, PTEN loss occurred in 66% of samples which is similar to findings in literature where loss has been observed to range from 55 to 70.2% (Kanamori et al. 2002, Uegaki et al. 2005, Mackay et al. 2010, Monte et al. 2012, Lee et al. 2012, Yang et al. 2015). For 11% of EIN cases, PTEN loss was observed, equal to that of PE, which is far lower than what has been observed in the literature with 44% and 63% observed in two separate studies where the EIN classification was used (Mutter et al. 2001, Monte et al. 2010), and in studies where the WHO94 scheme has been used, PTEN loss has been observed at 42.9–71% in the category atypical hyperplasia (Lacey et al. 2008, Feng et al. 2012, Lee et al. 2012). The discrepancy here between our observation and the literature could be due to our relatively small sample size for EIN.

In addition, no association between PTEN status and risk of progression in patients diagnosed with EIN or EEC was found. Previously, PTEN status has been shown to have a low predictive value in predicting clinical outcome of EIN, however, in combination with D-score, its prognostic value is increased (Baak et al. 2005b). It is important to note that the sample number of EIN patients that progressed was very small in both their and our studies (n=7). These findings would need to be validated in a larger population. Supporting our preliminary findings, Lacey et al. (2008) demonstrated in a large sample group that PTEN status was not associated with progression risk of endometrial hyperplasia.

With regards to PTEN status in EEC, several studies report that there is an association with PTEN status and progression in EEC but even here there are discrepancies. Some studies state that a PTEN-positive status is associated with a favourable prognosis (Kanamori et al. 2002, Uegaki et al. 2005) and that with higher stages of EEC, PTEN loss increases (Daniilidou et al. 2013), but two studies suggest that a PTEN-negative status is associated with a favourable prognosis (Mackay et al. 2010, Akiyama-Abe et al. 2013). The present study found no association between PTEN status and progression-free survival in EEC or any difference between PTEN status and FIGO stage. In combination, our findings and those of the literature makes PTEN a challenging prognostic biomarker for both EIN and EEC.

In the present study, a progressive decrease in PAX2 expression from PE to EIN to EEC was observed, which is also demonstrated in two separate studies (Monte et al. 2010, Allison et al. 2012). This suggests that loss of PAX2 expression is an early event of endometrial carcinogenesis and clonal expansion of these glands. For the purposes of this study, the H-score method was used to assess PAX2 expression in endometrial samples, whereas other studies have scored based on the percentage of PAX2 null glands in a sample (Monte et al. 2010) or grouped patients based on complete loss (0% cells staining), partial loss (1–75% cells staining) or minimal to no loss (76–100% cells staining) of PAX2 (Allison et al. 2012) or the number of positive cells per 1000 epithelial cells or tumour cells for EEC cases (Kahraman et al. 2012). Of interest, Kahraman et al. (2012) reported an increase in PAX2 expression with increasing severity of diagnosis despite the use of two different scoring methods including one employed by Monte et al. (2010) who had previously reported the opposite observation akin to our findings. Discrepancies here could be due to different immunohistochemistry methods. Whilst Monte et al. (2010) has used the same antibody utilised in the present study, Kahraman et al. (2012) made use of another. On closer comparison...
Kahraman et al. (2012) observed PAX2 staining in the stroma, an observation neither we nor several other studies (Monte et al. 2010, Allison et al. 2012, Joinet et al. 2015, Ørbo et al. 2015) have observed.

In this study, we present a method and threshold that could prove promising for identification of high-risk EIN lesions that will progress to EEC. EIN lesions with a PAX2 expression H-score below or equal to 75 (high risk) had significantly lower progression-free survival than those with an H-score above 75 (low risk). Furthermore, more EIN patients with high-risk PAX2 showed progression (73%; 8/11) compared to low-risk patients (8%; 2/25). This is the first time, to our knowledge, that PAX2 expression has been associated with progression in EIN in a population with long-term follow-up. We acknowledge that this study uses a small EIN sample number and therefore would need to be validated in a large patient cohort; however, we would like to emphasise that use of PAX2 for risk categorisation of EIN holds promise and should be investigated further. To be noted, a previous study demonstrated no association with PAX2 expression and progression of endometrial hyperplasia, however, this study had a very short follow-up of 2–6 months (Upson et al. 2012). It is recognised that endometrial carcinogenesis is a long process and development of EEC from normal endometrium can take 30–40 years where progression of EIN to EEC can take as long as 10 years. Although our study is limited by the small EIN sample size, the follow-up of these patients is long. Unlike for EIN, the PAX2 expression threshold was not prognostic in EEC, where patients with a high-risk score progressed similarly to those with a low-risk score (13% vs 11%). Furthermore, no additional benefit was demonstrated of using PTEN status in combination with PAX2 to determine progression in either EIN or EEC.

A challenge in comparing literature, and what could account for the variations seen between our observations and previous studies, is the use of different immunohistochemistry methods, observed staining patterns and scoring protocols. Some have observed staining in both stroma and glands whilst others only observe nuclear staining of the endometrial glands. Interpretation of immunostained tissue can be a significant limiting factor for accurate diagnostic and prognostic evaluation. By utilising digital image analysis, this variability and error due to subjective interpretation of staining patterns can be reduced. Immunostaining of PAX2 has been shown to be technically robust and its distinct nuclear staining is not difficult to interpret (Monte et al. 2010, Quick et al. 2012, Ørbo et al. 2015).

In the current study, reproducibility of scoring was high. A benefit of PAX2 staining is adjacent normal endometrial tissue functions well as an internal control (Quick et al. 2012).

In conclusion, decreased PAX2 expression is associated with clinical progression of EIN lesions. Thus, evaluation of PAX2 expression in EIN lesions could have significant impact on treatment decisions. To be able to distinguish between lesions that ought to be treated differently could significantly reduce the number of patients that are over and under treated and further investigation is therefore warranted.

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

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