

Endometrial cancer: redefining the molecular-targeted approach

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Abstract Endometrial cancer (EC) is the most frequent gynecologic malignancy in the world. Metastatic and recurrent disease confers a worse prognosis, and the side effects of the current cytotoxic agents are the main cause of treatment disruption. Recently, the genetic alterations that facilitate the start, development and progression of EC have been elucidated, reclassifying the disease in distinct subtypes with different mechanisms of carcinogenesis. Targeted therapy aims to interfere specifically these mechanisms causing less toxicity, therefore opening new perspectives for a tailored treatment and improvement of response and survival rates for heavily treated recurrent disease. Treatment with hormone therapy was not addressed in this review because it is an extensively discussed issue and would divert the discussion about molecular-targeted therapy. The purpose of this paper was to review the available literature data regarding the main genetic abnormalities related to the carcinogenesis and evaluate the safety and efficacy of the molecular-targeted agents in the treatment of metastatic and recurrent EC.

Keywords Cancer · Endometrium · Endometrial cancer · Chemotherapy · Targeted therapy · Molecular targets

Abbreviations

EC Endometrial cancer
GOG Gynecologic Oncology Group
HNPCC Hereditary non-polyposis colorectal carcinoma

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Introduction

Among the gynecologic malignancy, endometrial cancer (EC) is the most frequent, with 52,630 estimated new cases and 8590 deaths worldwide for 2014. The lifetime risk for developing this malignancy is approximately 2.7 %, based on 2008–2010 data [1]. Almost 90 % of the cases of EC are sporadic, whereas the remaining are hereditary [2].

Since 1983, according to the Bokhman [3] hypothesis, the sporadic endometrial tumorigenesis is broadly divided into type 1 and type 2. This dualistic model defines these cancers in terms of both histology and clinical behavior. Type 1 tumors are more frequent, approximately 70–80 %. They follow the estrogen-related pathway, arising from a background of unopposed estrogenic stimulation, associated with endometrial hyperplasia, expressing estrogen (ER) and progesterone receptors (PR). Most of the type 1 are low grade and, histologically, shows endometrioid differentiation. Mucinous adenocarcinomas express ER and/or PR, are low grade and are also considered type 1 carcinomas. In general, type 1 EC has a favorable behavior and good response to treatment with hormone therapy in the context of metastatic and recurrent disease.

The frequency of the sporadic type 2 carcinomas ranges from 10 to 20 %. They follow the estrogen-unrelated pathway and arise from the background of an atrophic endometrium. They are usually high-grade carcinomas of non-endometrioid differentiation and have a negative or weakly positive for expression of ER and PR. Type 2 carcinomas occur at an older age than the type 1 tumors [4].

In the last decade, the increasing molecular knowledge of the tumorigenesis pathways has distinguished the EC in different subtypes by genetics alterations. The Cancer Genome Atlas Research Network (TCGA) has reported an important integrated genomic, transcriptomic and

proteomic characterization of 373 EC (307 endometrioid, 53 serous and 13 mixed endometrioid and serous). Considering mutations, copy number aberrations and microsatellite instability status, EC was reclassified into four novel categories that could have immediate therapeutic application: POLE ultramutated, microsatellite instability (MSI) hypermutated, copy number low and copy number high. The POLE subtype occurred in 10 % of endometrioid tumors and was characterized by ultrahigh somatic mutation frequency, MSI and newly identified hotspot mutations in the exonuclease domain. All of the endometrial tumors were examined for mRNA expression ($n = 333$), protein expression ($n = 293$), miRNA expression ($n = 367$) and DNA methylation ($n = 373$) [5].

In this genomic data, mutations in PI3K/AKT pathway were more frequent in EC than in any other tumor type studied by TCGA. The endometrioid EC frequently presented POLE mutation, MSI (40 %), KRAS and CTNNB1 activation signaling, showing many characteristics in common with colorectal carcinoma. And similar to basal-like breast cancer and high-grade serous ovarian carcinoma (HGSOCs), the uterine serous carcinomas presented high frequency of non-silent *TP53* and somatic copy number alterations. However, HGSOCs and basal-like breast cancers did not demonstrate high frequency of *PIK3CA*, *FBXW7*, *PPP2R1A* and *ARID1A* mutations as shown in uterine serous carcinomas [5].

One important point of this analysis was that the molecular characterization demonstrated that approximately 25 % of tumors classified as high-grade endometrioid by pathologists have a molecular phenotype similar to uterine serous carcinoma. The TCGA has important limitations: Clear cell histology and carcinosarcoma were not evaluated, and high-grade endometrioid carcinoma classified as being of copy number high subtype might have included mixed tumors in which only endometrioid component was sampled.

Finally, in this context of different sets of genes and molecular alterations, referred as a 'multi-step carcinogenesis,' several novel treatment regimens are being developed based on drugs targeting specifically these pathways, achieving therapeutic effects with fewer side effects [6, 7].

The type I endometrial carcinomas (endometrioid)

The temporal sequence of the large number of mutations that promote the carcinogenesis of type I EC is not known. The most common genetic alterations include microsatellite instability, *PTEN* and β -catenin mutation. There is also evidence for different gene profiles of *EGFR*, *FGFR2* and mTOR pathway [8, 9].

The mismatch repair system (MRS) is composed by the genes *MLH* and *MSH*. The *MLH1* inactivation is the

most common mechanism of tumorigenesis in the endometrium, accomplished by hypermethylation of CpG islands in the promoter, the process of epigenetic silencing. *MLH1* is important in repair of short segments (2–4 bases). Since simple repeat sequences are unstable in cells with MSI, the observed mutation may be secondary to the MSI derived from defective *MLH1* expression, differently to colon cancer, where MSI is due to mutations in the *MSH2*, *MLH1* and *MSH6* genes. Inherited or somatically acquired mutations of *MSH6* are also common in patients with MSI EC [10–12].

MSI has been found in a substantial fraction of sporadic endometrial tumors, but data on whether these endometrial tumors differ from their MSI counterparts in clinical characteristics, prognostic variables and survival are still lacking [9, 13]. Basil et al. [14] demonstrated that only 13 % of MSI (+) endometrial tumors presented with advanced stage III or IV disease compared with 28 % of MSI (–) tumors. This clinical difference regarding stage has not been demonstrated previously in EC. They also suggested that MSI (–) was found to be associated with more aggressive histological subtypes by univariate analysis, but no difference in overall survival (OS) was observed. These findings were different from previous series involving other types of cancer, in which MSI (+) was associated with good behaviors and better prognosis, acting as a marker of aggressiveness [10–12, 15].

Hereditary non-polyposis colorectal carcinoma (HNPCC) is a well-characterized autosomal dominant cancer syndrome associated with tumor MSI, and EC is the second most common malignancy found in these patients [7, 16]. MSI is more frequent in endometrioid than in non-endometrioid tumors [10]. Approximately 12–24 % of sporadic colorectal cancers have MSI positivity, and it is associated with favorable prognosis and improved survival [17, 18]. Otherwise, sporadic endometrioid tumors are positive for MSI in 30 % of the patients [15].

Currently, by non-inferiority analysis, the GOG 209 study has established carboplatin plus paclitaxel as first-line chemotherapy regimen for patients with recurrent EC, demonstrating a better toxicity profile compared with the three-drug regimen paclitaxel, doxorubicin and cisplatin [19]. Prospective trials evaluating the interaction between tumor MSI and response to platinum compound therapy in women with advanced or recurrent EC are necessary to clarify the relationship between mismatch repair and response to chemotherapy.

The *PTEN* gene, located at chromosome 10q23, encodes a protein (phosphatase and tensin homolog, PTEN) with tyrosine kinase function and behaves as a tumor suppressor gene. The inactivation of the *PTEN* tumor suppressor gene is implicated in the carcinogenesis of a large range of human sporadic. This is the most frequent genetic alteration or abnormality (qualquer um dos dois) in endometrioid adenocarcinoma, present in 83 % of the cases. Numerous

mechanisms of inactivation and the pathway downstream that lead to the cancer phenotype have been described. These alterations seem to be conserved in a given histological subtype of adenocarcinoma irrespective of the primary site. For example, endometrioid adenocarcinomas of the ovary and endometrium have similar inactivation patterns associated. On the other hand, the consequences of altered *PTEN* expressions are different in various tissues [20, 21].

PTEN acts as a gatekeeper for initiation of carcinogenesis in the endometrium from a normal background state, and additional *PTEN* damage accumulates in the transition from premalignant to malignant disease. Thus, immunohistochemically detected *PTEN* loss expression is an informative biomarker for endometrial neoplasm, including precancerous lesions [22].

One of the most important signaling pathways involved in gynecologic carcinogenesis is the PI3K/AKT/mTOR pathway. Eighty percent of the cases of endometrioid EC have one or more somatic alterations affecting this pathway. Thus, amplifications, mutations and translocations, resulting in aberrant activation of this pathway, occur more frequently than in any other pathway [23, 24].

The primary negative regulator of the PI3K pathway is *PTEN*. Therefore, *PTEN* loss may lead to aberrant cell growth and an escape from apoptosis. *PIK3CA*, another gene often mutated in various types of cancer, may also hold a role in the alteration of the phosphatidylinositol 3-kinase (PI3K)/Akt pathway in EC. *PIK3CA* mutations appear in 25–36 % of endometrioid EC and in 15 % of non-endometrioid EC, often coinciding with *PTEN* mutations [25].

One important study explored whether mutations of the PI3K pathway, apart from *PIK3CA* and *PTEN*, were present in EC [26]. It has been reported by several groups a mutation rate of up to 20 % in *PI3K1*, which is significantly higher than any other lineage, therefore demonstrating a selective target in EC [25, 27, 28]. The *PI3K1* gene encodes the PI3K regulatory subunit p85a. Several of these mutations are known to phosphorylate AKT, thus activating the downstream signaling pathway. The *PI3K2* has also been established as a novel cancer gene. The mutation rate for *PI3K2* has been reported in up to 5 % of EC, and several mutations have shown to exhibit gain of function [25, 27]. Shoji et al. [29] detected the presence of *AKT1* mutations in 2 % of ECs tissue samples, which did not demonstrate other mutations in *PIK3CA*, *PTEN* or *KRAS*. The authors suggested that *AKT1* mutation might be mutually exclusive with other PI3K–AKT activating alterations.

The RAS–RAF–MAPK is another important pathway in a variety of human cancers, interacting with the PI3K pathway through the RAS proteins. This interaction may suggest a cooperation between the two pathways in order to determine functional outcomes. Somatic mutations of the

KRAS gene are found in 18–28 % of endometrioid EC [30, 31].

Constitutive activating mutations in *KRAS* are more frequently found in tumors with MSI, suggesting that both events may occur simultaneously before clonal expansion [32]. In EC, *KRAS* mutations can coexist with mutations in *PIK3CA*, *PIK3R1* and *PTEN*, suggesting that *KRAS* mutations are not functionally redundant with PI3K pathway mutations [33].

The most common molecular alterations in tumor cells leading to disruption of β -catenin degradation are mutations that inactivate APC or activate β -catenin itself [34]. These alterations produce an accumulation of cytoplasmic β -catenin that translocates into the nucleus, interacts with members of the lymphoid enhancer factor-1/T cell factor and activates the transcription of various genes, such as *cyclin D1* and *MYC*. The APC/ β -catenin signaling pathway has mainly been studied in the type 1 of EC. In these tumors, mutations of β -catenin occurred in up to 17 % of cases [35]. E-cadherin is a transmembrane protein with an intracellular domain that connects to the actin cytoskeleton through a complex with cytoplasmic catenin. Decreased expression of E-cadherin is associated with a loss of cell–cell cohesive forces and has been shown to precede tumor cell motility. The mainway that APC protein downregulates β -catenin levels is by cooperating with glycogen 76 synthase kinase 3 β (GSK-3 β), inducing phosphorylation of serine–threonine residues coded in exon 3 of the β -catenin gene and its degradation through the ubiquitin–proteasome pathway [36, 37]. The β -catenin alterations are almost exclusive and do not coexist with MSI, *PTEN* and *KRAS* mutations. When altered, β -catenin expression changes are usually seen throughout all tumor cells, its changes are present in some premalignant lesions [36]. This suggests that β -catenin mutation is an early step of endometrial tumorigenesis that is clonally represented in all tumor cells. On the other hand, alterations in β -catenin activity probably contribute to later tumor progression targeting the *cyclin D1* gene [38, 39].

Finally, some preclinical studies have addressed great attention in exploring the alterations in the fibroblast growth factor receptor 2 (FGFR2) in EC [40–42]. FGFR2 arose as a possible target for therapeutic approaches, enabling the development of the FGFR inhibitors as novel molecular agents for the treatment of several types of tumors. FGFR2 belongs to fibroblast growth factor receptor (FGFR) tyrosine kinase family that comprises four different transmembrane receptor tyrosine kinases (FGFR1–FGFR4) and their alternative spliced isoforms [42]. They activate downstream pathways such as RAS–RAF–MAPK and differentially respond to FGF ligands. Depending on the type of the cell, *FGFR2* has been shown to play crucial role in many physiological and pathological processes and

is considered as either oncogene or tumor suppressor gene. Several types of molecular alterations have been described, including gene overexpression and point mutation [43].

The type 2 endometrial carcinoma (non-endometrioid)

The background of the non-endometrioid EC (type 2) is fairly different. They are comprised of the high-grade papillary serous and clear cell carcinomas. These tumors arise in relatively older women and are not usually preceded by a history of unopposed estrogen exposure, but rather from a background of atrophic endometrium. They perform an aggressive clinical course, a greater propensity for early spreading and a worse prognosis than the more common endometrioid adenocarcinomas [44, 45].

The most common genetic alteration in type 2 EC occurs in *p53* gene. This tumor suppressor is located on chromosome 17 and is important in preventing the propagation of cells with damaged DNA. Mutations in *p53* are present in nearly 90 % of serous carcinomas. After DNA damage, nuclear *p53* accumulates and causes cell cycle arrest by inhibiting cyclin-D1 phosphorylation of the *Rb* gene and thereby promoting apoptosis. Mutations in *p53* are present in almost 80 % of endometrial intraepithelial carcinoma lesions, the precursor lesion to serous carcinomas. It is postulated that mutation in one allele occurs early during the development of serous carcinoma, and loss of the second normal allele occurs late in the progression to carcinoma [46].

Inactivation of *p16* and overexpression of HER2/neu are other alterations that occur in type 2 EC. *p16* inactivation was found in approximately 45 % of serous carcinomas and some clear cell cancers. The *p16* tumor suppressor gene is located on chromosome 9p21 and encodes a regulatory protein in the cell cycle. Furthermore, inactivation of *p16* leads to uncontrolled cell growth and tumor progression [47]. HER2/neu overexpression and gene amplification were found in about 45 and 70 % of clear cell and serous carcinomas, respectively. HER2/neu is an oncogene that codes a transmembrane receptor tyrosine kinase involved in cell signaling [48, 49]. Negative and reduced E-cadherin expression occurred in 62 and 87 % of serous and clear cell cancers, respectively. E-cadherin-negative tumors are more likely to be poorly differentiated or non-endometrioid and are associated with poorer prognosis [50, 51].

The hereditary endometrial carcinomas

Comprising the Lynch syndrome, the most common extracolonic malignancy is the EC, developed at a significant earlier age than in the general population. Currently, the

most important pathway of HNPCC is the MSI [48, 49, 52, 53]. The EC arising in HNPCC is related to type 1 tumors, since they occur at young age and are histologically of mucinous or endometrioid type, but their pathway is driven by germ line mutations and is, thus, distinctive [54–57].

New approaches for endometrial cancer

In case of recurrent advanced disease, when surgery is not effective in curing and chemotherapy cannot be used to control the disease progression, the median survival among these patients is <1 year. However, the use of the new targeted agents alone or in combination with cytotoxic therapy is emerging as a promise to prolong survival and improve the quality of life. The results and characteristics of the completed trials evaluating the molecular-targeted drugs are summarized in Table 1.

Angiogenesis inhibitors

Bevacizumab, aflibercept and thalidomide are the antiangiogenic agents that have been recently evaluated in the treatment of EC, as well as sunitinib, a multi-targeted agent. Bevacizumab is a humanized monoclonal antibody against vascular endothelial growth factor (VEGF)-A. VEGF is a cytokine associated with the promotion of cell division and permeability of vascular endothelial cells, essential for angiogenesis, and is enhanced in cancer cells. In a phase II GOG trial of single-agent bevacizumab in recurrent EC, seven of 52 patients (13.5 %) exhibited a clinical response [one complete response (CR) and six partial responses (PR)] and 21 patients (40.4 %) had a progression-free survival (PFS) of at least 6 months. The median PFS was 4.2 months, and the overall survival (OS) was 10.5 months. The adverse reactions were the same as those seen with conventional bevacizumab therapy, although no perforation of the digestive tract or fistula formation was reported [58]. Results of an ongoing three-arm phase II trial that compares the combination of standard chemotherapy or ixabepilone-containing regime with bevacizumab versus the combination of the standard cytotoxic scheme with temsirolimus are anxiously expected to disclose how well they work in patients with stage III, stage IV and recurrent EC [59].

Aflibercept exhibits high-affinity binding to VEGF-A, VEGF-B and placental growth factor. In a GOG phase II trial, three out of 44 patients (7 %) had PR, and 10 patients (23 %) met the endpoint of 6 months PFS. The median PFS and OS were 2.9 and 14.6 months, respectively. Significant toxicities were uncommon, but there were two treatment-related deaths recorded—one GI perforation and one arterial rupture [60].

Table 1 Clinical trials with target therapy in ovarian cancer

Drug	Class	Phase	Number of patients	Response
Sunitinib monotherapy [62]	TKI multi-targeted inhibitor: VEGFR and PDGFR	II	Chemotherapy treated: 33	PR: 18.1 % SD: 18.1 % PFS > 6 m: 30.3 % PFS: 3 m
Erlotinib monotherapy [63]	TKI EGFR inhibitor	II	Chemotherapy naive: 34	PR: 12.5 % SD: 46.9 %
Gefitinib monotherapy [64]	TKI EGFR inhibitor	II	Chemotherapy treated: 26	CR: 3.8 % SD: 27.2 % PFS ≥ 6 m: 15.4 %
Lapatinib monotherapy [70]	TKI HER2 inhibitor	II	Chemotherapy treated: 30	PR: 3.3 % SD: 23.3 % PFS > 6 m: 10 %
Aflibercept monotherapy [60]	Fusion protein anti-VEGF	II	Chemotherapy treated: 27	PR: 7 % SD: 32 % PFS ≥ 6 m: 41 %
Thalidomide monotherapy [61]	Immunomodulatory agent	II	Chemotherapy treated: 24	PR: 12.5 % SD: 8.3 % PFS: 1.7 m PFS ≥ 6 m: 8.3 %
TEM monotherapy [72]	mTOR inhibitor	II	Chemotherapy naive: 29 Chemotherapy treated: 25	Chemotherapy naive: PR: 14 %; SD: 69 %; PFS: 7.33 m. Chemotherapy treated: PR: 4 %; SD: 48 %; PFS 3.25 m.
Ridaforolimus monotherapy [78]	mTOR inhibitor	II	Chemotherapy treated: 24 Chemotherapy naive: 7	PR: 8.8 % SD: 52.9 %
Bevacizumab monotherapy [58]	Anti-VEGF Mab	II	Chemotherapy treated: 50	CR: 1.9 % PR: 11.5 % PFS ≥ 6 m: 40.4 %
Trastuzumab monotherapy [68]	Anti-HER2 receptor Mab	II	Chemotherapy naive: 8 Chemotherapy treated: 26	PR: 0 % SD: 58.8 % PFS: 1.84 m
Combination of TEM and bevacizumab [74]	mTOR inhibitor and anti-VEGF Mab	II	Chemotherapy treated: 53	CR: 2 % PR: 22.5 % SD: 55.1 % PFS ≥ 6 m: 46.9 % PFS: 5.6 m
Combination of temsirolimus and PLD [75]	mTOR inhibitor and anthracycline	Ib	Chemotherapy treated: 5	PR: 20 % SD: 60 % PFS ≥ 6 m: 80 %

Table 1 continued

Drug	Class	Phase	Number of patients	Response
Dovitinib [89]	TKI FGFR2 inhibitor	II	Mutated: 22 Non-mutated: 31	Mutated: PR: 5 %; SD: 59 %; PFS: 4.1 m; OS: 20.2 m. Non-mutated: PR: 16 %; SD: 35 %; PFS: 2.7 m; OS: 9.3 m
Brivanib [90]	TKI multi-targeted inhibitor: anti-FGFR and anti-VEGF	II	Chemotherapy treated: 43	OS: 10.7 m CR: 2.3 % PR: 16.3 % PFS \geq 6 m: 30.2 % PFS: 3.3 m
Nintedanib [91]	TKI multi-targeted inhibitor: PDGFR, FGFR and VEGFR	II	Chemotherapy treated: 32	OS: 10.1 m PFS > 6 m: 21.9 % CR: 0 % PR: 9.4 %
Pilaralisib [85]	TKI PI3K inhibitor	II	Chemotherapy treated: 67	CR: 3 % PR: 3 % PFS > 6 m: 11.9 %

TKI tyrosine kinase inhibitor, TEM temsirolimus, VEGF vascular endothelial growth factor, PDL pegylated liposomal doxorubicin, PR partial response, SD stable disease, PFS progression-free survival, PFS > 6 m, progression-free survival >6 months

The exact mechanism of angiogenesis inhibition by thalidomide is unknown. The GOG performed a phase II trial evaluating the anti-tumor activity and adverse effects of thalidomide in persistent or recurrent EC. Only three of 24 patients (12.5 %) had PR, two (8.3 %) had stable disease (SD), and 15 (62.5 %) had progressive disease (PD); two patients (8.3 %) remained progression free after 6 months, and the median PFS and OS were 1.7 and 6.3 months, respectively. Thalidomide demonstrated limited ability to reduce angiogenic markers levels and to delay progression [60].

Sunitinib, an oral multi-targeted receptor tyrosine kinase inhibitor with antiangiogenic activity, was evaluated in a Canadian phase II trial, where six (18.1 %) out of 33 women had PR and six (18.1 %) had SD; in total, 10 patients (30.3 %) had disease control for at least 6 months [62].

EGFR inhibitors

The recent attempt to inhibit the EGFR pathway in EC was performed mainly with low molecular weight tyrosine kinase inhibitors (gefitinib, erlotinib) and cetuximab, a monoclonal antibody against EGFR. Amit et al. [63] assessed 32 patients for response to erlotinib in a phase II trial: 12.5 % had PR and 46.8 % had SD. Of the 26 patients evaluable for gefitinib efficacy in other similar phase II trial, four patients experienced PFS more than 6 months, one had CR, and the clinical benefit was 31 % [64].

HER2 inhibitors

Trastuzumab and pertuzumab are monoclonal antibodies that act in different domains of HER2 and are used in the treatment for HER2-enriched breast cancer. Many case reports suggested responses to trastuzumab in patients with recurrent and metastatic EC who had HER2 overexpressed [65–67]. The GOG group performed a phase II trial to evaluate the efficacy of single-agent trastuzumab in advanced or recurrent HER2-positive EC. Even with the amendment to require HER2 amplification by FISH, of the 34 patients enrolled, no major tumor response was observed and 20 patients experienced stable disease [68]. A preclinical study evaluated pertuzumab activity separately or in combination with trastuzumab against primary uterine serous papillary adenocarcinoma cell lines expressing different levels of HER2/neu. The results revealed that pertuzumab significantly increases trastuzumab-induced antibody-dependent cell-mediated cytotoxicity and may represent a new therapeutic option [69]. Lapatinib and neratinib are HER2 and EGFR inhibitors. The GOG 229D was a phase II trial assessing the efficacy of single-agent lapatinib in persistent and recurrent EC. Three out of 30 evaluable patients

had a PFS more than 6 months. Only one had a PR, seven had SD, 21 had PD, and just 8 % was HER2 positive. The study concluded that lapatinib did not have enough activity to be used as a single agent [70]. Data demonstrated that cell lines with HER2 amplification were strongly more sensitive to neratinib compared with non-amplified cell lines; thus, neratinib may be a potential treatment for HER2-amplified EC [71].

mTOR inhibitors

Of particular interest is the PI3K/AKT/mTOR inhibition in the treatment of recurrent or persistent EC, since this pathway is frequently activated. Temsirolimus (TEM), an ester derivative of rapamycin, is the most studied mTOR inhibitor in this context. A phase II study looking at 29 chemotherapy-naïve patients showed that 14 % had PR and 69 % had SD as best response. On the other hand, 25 patients were in the chemotherapy-treated group: one (4 %) had PR and 12 (48 %) had SD [72]. A Canadian study concluded that women previously treated with chemotherapy were at 7.3 times greater risk of progression and experienced 20.9 % increased tumor growth when compared to chemotherapy-naïve women [73]. The combination of TEM and bevacizumab assessed in previously treated patients showed 24.5 % of clinical responses (CR or PR) and 46.9 % progression free for at least 6 months [74]. A phase Ib trial showed that the combination of TEM and pegylated liposomal doxorubicin (PLD) is safe and manageable [75]. The phase II trial performed by Fleming et al. [76] assessing TEM with megestrol alternating with tamoxifen in women with metastatic and recurrent EC was closed early because of an excess of venous thrombosis and the absence of activity. Ridaforolimus was assessed in two phase II trials. In the first one, Colombo et al. [77] demonstrated that 13 of 45 patients (28.9 %) achieved a clinically beneficial response (CR, PR or SD) for more than 16 weeks. In a most recent trial, the response rate was even higher; Tsoref et al. [78] showed that 21 of 34 patients (61.7 %) had clinical benefit. Recently, the anti-diabetic medication metformin has been demonstrated to exert anti-neoplastic effects in several cancer cell lines, including EC cell lines. Metformin also reduces the AKT activity through inhibition of insulin receptor substrate 1 [79]. A retrospective multi-institutional cohort analysis compared EC patients with diabetes mellitus who used metformin to those who did not use from 2005 to 2010 and concluded that non-metformin users had 1.8 times worse recurrence free survival (95 % CI 1.1–2.9, $p = 0.02$) and 2.3 times worse overall survival (95 % CI 1.3–4.2, $p = 0.005$) [80]. Prospective ongoing trials are evaluating the association of metformin with other therapies. MD Anderson is recruiting patients for a phase II single-arm study to evaluate whether the combination of

everolimus, letrozole and metformin can help control recurrent or progressive EC [81]. Also GOG is now conducting a randomized phase II/III two-arm placebo-controlled trial to evaluate the addition of metformin to the cytotoxic regimen paclitaxel and carboplatin in treating patients with EC stage III, IV or recurrent disease [82].

New agents

The phosphatidylinositol 3-kinase (PI3K) pathway, considered an attractive target for therapeutic intervention, is one of the most altered oncogenic pathways in EC. Two phase II trials evaluating the safety and efficacy of the PI3K inhibitors BKM-120 [83] and XL147 [84] for recurrent EC were completed not long ago and are under analysis. A recent phase II multicenter single-arm study evaluated the use of pilaralisib (XL147) in previously treated advanced or recurrent EC. Of the 67 assessed patients, only 6 % presented overall response rate and the rate of PFS >6 months was 11.9 %. There was no association between molecular alterations and clinical activity, and the most commonly reported adverse events grade >3 were rash (9.0 %), diarrhea (4.5 %) and increased alanine aminotransferase (4.5 %) [85].

The inhibition of the AKT pathway could play an important therapeutic role in the treatment of advanced EC. A preclinical analysis has demonstrated specificity of MK-2206, an orally active allosteric inhibitor for AKT. Used in the context of progesterin resistance, the MK-2206 stabilized and increased the level of the progesterone receptor B and improved progesterone response in EC cells that have hyperactivated AKT. A phase II investigating the efficacy of the MK-2206 in recurrent EC is ongoing [86]. Others AKT inhibitors (ARQ 092, AZD5363, GDC-0068 and GSK2141795) are being tested in early-stage clinical trials for solid tumors.

The FGFR2 has emerged as a novel target-based therapy for the treatment of EC. A preclinical study explored the anti-tumor activity of the FGFR2 inhibitor dovitinib in EC xenograft models and concluded that its action is beyond the *FGFR2* mutational status [87]. Another preclinical assay demonstrated synergistic antitumor activity combining ponatinib, an oral multi-targeted kinase inhibitor that potently inhibits all four FGFR family members, with ridaforolimus in EC cells [88].

The results of a phase II trial evaluating the efficacy and safety of dovitinib as second-line therapy in mutated and non-mutated patients with advanced EC were recently presented at the ESMO Congress 2014. The median PFS (4.1 versus 2.7 months) and OS (20.2 versus 9.3 months) trended to be higher in the *FGFR2* mutated group; the disease control rate was 64 % (59 % SD, 5 % PR) in the *FGFR2* mutated group and 51 % (35 % SD, 16 % PR) in

the *FGFR2* non-mutated group, respectively. The most common grade 3/4 adverse events were hypertension (17 %) and diarrhea (9 %) [89].

The phase II trial published by Pollock et al. [90] evaluated the efficacy and tolerability of brivanib, an oral multi-targeted inhibitor with anti-VEGF and anti-FGFR activity, as single agent in recurrent or persistent EC. Of the 43 evaluable patients, 18.6 % had response (1 CR and 7 PR) and the median PFS and OS were 3.3 and 10.7 months, respectively. Brivanib was reasonably well tolerated; nine patients had grade 3 hypertension and one experienced grade 4 confusion.

Recently, a GOG single-arm phase II study evaluated nintedanib, a potent small molecule triple receptor tyrosine kinase inhibitor of PDGFR α and β , FGFR 1/3 and VEGFR 1–3, as monotherapy in the treatment of recurrent or persistent EC. Of the 32 eligible patients, there were no CR and only three partial responses with an overall response rate of 9.4 %. The PFS, OS and the PFS at 6 months were, respectively, 3.3, 10.1 months and 21.9 %; gastrointestinal toxicity was the most common serious adverse event (5). Based on the insufficient activity, the enrollment to second stage of the study was not carried out [91].

Theoretically, dual mTOR/PI3K competitive inhibitors attached to the ATP-binding cleft of both class I PI3Ks and mTORC1/2 should suppress more efficiently the PI3K/AKT/mTOR pathway than agents that act at single points of inhibition. Preclinical trials using GDC-0980 and BEZ235 have demonstrated cell growth inhibition in several cancer cell lines and tumor xenograft models [92, 93]. Apparently, the BEZ235 had more anti-tumor activity in tumor cells with PI3K and/or *PTEN* mutations [94]. Recently, GDC-0980 phase II study has been completed, but so far results have not been published [95].

Conclusion

The recent advances in the comprehension of the molecular mechanism of endometrial genetic alterations and pathways that explain the carcinogenesis have led to the development of clinical trials addressing the use of new targeted therapies for EC. Unfortunately, in spite of the efforts of the pharmaceutical industry and academic institutions, only few molecular-targeted agents have shown a significant impact on survival and response in clinical trials. Many agents that demonstrated significant antitumor activity in preclinical studies have failed to reproduce these results when tested in the clinical studies, and most of the patients with advanced disease remain incurable and refractory to conventional therapy.

Among the targeted therapy drugs, as shown in Table 1, the mTOR inhibitors had the highest response rates,

reaching considerable PR (14 %), SD (69 %) and median PFS (7.3 months) for the chemotherapy-naïve group [72]. The FGFR inhibitors remain a promise for personalized treatment of advanced endometrial disease, but we must wait for unpublished results of recently closed clinical trials. Even more, it has become clear that chemotherapy-naïve patients respond better than previously treated patients. Perhaps this fact may be explained by changes in the molecular profile over the course of the disease. However, these outcomes still remains lower in comparison with the results presented by the classical trials that evaluated cytotoxic chemotherapy regime adriamycin and cisplatin where objective responses (CR and PR) reached 60 % in the first trial [96] and 43 % in the EORTC study [97], but with greater toxicity profile.

As exposed above, EC has a heterogeneous complex carcinogenesis process. Several genetic abnormalities accumulate in the cellular machinery affecting components of the same or different pathways. Consequently, there is considerable difficulty in identifying a specific target that, when pharmacologically inhibited, would make the cancer cells susceptible, shifting them toward the mechanism of apoptosis.

In conclusion, even with limited results obtained so far, the personalized approach should be the mainstay for future alternative treatment in metastatic disease with even more specific and less toxic agents.

Conflict of interest None.

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