Original Article Altered expression of fibroblast growth factor receptor 2 isoform IIIc: relevance to endometrioid adenocarcinoma carcinogenesis and histological differentiation

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Abstract: Fibroblast growth factor receptor 2 (FGFR2) is activated in many cancers and considered as a potential therapeutic molecular target including for endometrial endometrioid carcinoma (EEC). Overexpression of FGFR2 isoform IIIc (FGFR2IIIc) has been shown to be associated with carcinogenesis in various cancers, but its expression in EEC has not been reported yet to the best of our knowledge. In this study, we identified roles for FGFR2IIIc in EEC carcinogenesis and demonstrated its diagnostic and prognostic values in EEC. FGFR2IIIc expression was compared between 10 normal endometrium, 10 atypical endometrial hyperplasias, and 47 EEC specimens using immunohistochemistry and quantitative real-time PCR. Atypical hyperplasia, Grade 1 (G1), and Grade 2 (G2) differentiated EEC tissues showed significantly higher FGFR2IIIc expression than normal endometrium tissue. However, as compared to G1 and G2 EECs, Grade 3 (G3) differentiated EEC tissue showed lower FGFR2IIIc expression (P<0.05). There was no significant association between FGFR2IIIc expression and patient age, lymph node metastasis, and EEC stage. These results suggest that altered FGFR2IIIc expression plays an important role in EEC carcinogenesis and may occur in precancerous tissues. However, FGFR2IIIc appears to be not related to EEC progression. Some G3 EECs may develop through different carcinogenic processes than G1 and G2 EECs.

Keywords: Endometrial carcinoma (EC), endometrial endometrioid carcinoma (EEC), fibroblast growth factor receptor 2 (FGFR2), IIIc, carcinogenesis, differentiation

Introduction

Endometrial carcinoma (EC) is one of the most common cancers among women worldwide. About 80% of EC are endometrial endometrioid carcinomas (EECs). EEC may be preceded by precancerous lesions such as hyperplasia and atypical hyperplasia [1, 2]. Atypical hyperplasia always shows similar findings to G1 EEC in not only histological appearance but also molecular features [3]. EECs are divided into 3 grades according to histological differentiation: grade 1 (G1), well differentiated; grade 2 (G2), moderately differentiated; and grade 3 (G3), poorly differentiated. Histological grade determines prognosis. As compared to patients with G1 and G2 EECs, patients with G3 EEC present with significantly worse prognosis [4, 5].

Despite being a serious public health problem, EC has not drawn enough attention. Moreover,

only a few potential molecular therapeutic targets have been identified, the most recent being fibroblast growth factor receptor 2 (FGFR2). FGFR2 and its signalling pathway are reported to be activated in many cancers due to gene amplification and point mutations. In addition, FGFR2 is a potential therapeutic molecular target for patients with FGFR2 activation-associated cancer such as EC [6-12].

Alternative splicing of FGFR2 produces 2 isoforms, FGFR2IIIb and FGFR2IIIc. In normal tissue, FGFR2IIIb is mainly expressed in epithelial cells, while FGFR2IIIc is mainly expressed in mesenchymal or stromal cells [13]. FGFR2IIIc expression has been reported in various cancers including bladder cancer and ovarian cancers [14-19]. However, to our knowledge, the role of FGFR2IIIc in EEC has not yet been clarified. The purpose of the present study was to identify the role of FGFR2IIIc expression in EEC

	Normal endometrium	Hyperplasia	ECC	
	n=10	n=10	n=47	
High TS	1 (10%)	8 (80%)	23 (49%)	
Low TS	9 (90%)	2 (20%)	24 (51%)	
P-value		p<0.01	p<0.05	

Table 1. Association of FGFR2IIIc expression asevaluated by total score (TS) with different histo-logical types

carcinogenesis and to investigate association of FGFR2IIIc expression with clinicopathological features in EEC.

Methods and materials

Case selection

We reviewed the pathology data for 47 EEC patients who underwent surgery at Nippon Medical School hospital. Normal and atypical hyperplasia tissue samples were also obtained from 10 patients respectively. Two certified pathologists (WX Peng and Z Naito) reviewed all cases in order to verify the original histopathological diagnosis, grading, and EEC stage according to the World Health Organization (WHO) classification system and 2008 International Federation of Gynecology and Obstetrics (FIGO) grading of EC. This study was conducted in accordance with the ethical principles of the Declaration of Helsinki, Informed consent for the usage of tissues was obtained from all patients.

Immunohistochemical analysis

Paraffin-embedded tumors were cut into 3.5um thick sections and placed on silane-coated glass slides. The sections were de-waxed in xylene and rehydrated in a series of graded ethanol solutions, and the endogenous peroxidase activity was blocked with a 0.3% H₂O₂ methanol solution. Before application of the primary antibody, the slides were subjected to antigen retrieval by heating in 10 mM citrate buffer (pH 6.0) for 15 min at 121°C in an autoclave oven. Then, anti-FGFR2IIIc antibody (in house; 1:200 dilution) [20] was applied to the slides, which were incubated overnight at 4°C. The slides were rinsed in 0.01 mol/L phosphate- buffered saline, and bound antibodies were detected with the Simple Stain MAX PO (R) (Nichirei Corp. Tokyo, Japan) using 3,3'-diaminobenzidine tetrahydrochloride as the substrate. The peroxidase reaction was visualized with 0.02% 3,3'-diaminobenzidine tetrahydrochloride containing 0.005% H_2O_2 in 0.01 M Trisphosphate buffer (pH 7.4). Finally, the sections were lightly counterstained with hematoxylin.

Blinded immunohistochemical evaluation of each case was carried out independently by the above pathologists.

The immunohistochemical expression of FGF-R2IIIc was graded using the immunostain intensity score (IS) (0, completely negative; 1, weekly stained; and 2, moderately to strongly stained), and the graded percentage score (PS) of positive cells (0, less than 5%; 1, 5-50%; and 2, more than 50%). The total score (TS) was then calculated using the following formula: TS=IS×PS. The cases with TS of 4 were classified into high and the others into low expression groups. Subsequently, we analyzed the level of FGFR2IIIc expression according to patient age, tumor differentiation grade, lymph node metastasis, and EEC stage.

Quantitative real-time polymerase chain reaction (qRT-PCR)

To confirm FGFR2IIIc expression in the collected tissues, gRT-PCR was conducted using tissue isolated from formalin-fixed paraffinembedded (FFPE) specimens. RNA samples were then extracted using RNeasy FFPE kit (Qiagen, Crawley, UK), according to the manufacturer's instructions. 18S was used as internal control. Then, cDNA was synthesized from total RNA using SuperScript[®]VILO[™] cDNA Synthesis Kit following the manufacturer's protocol (Invitrogen by Life Technologies, Carlsbad, CA). The corresponding cDNA was amplified using specific primers for FGFR2IIIc, 18S rRNA, and a TaqMan probe (all from Applied Biosystems), with denaturation at 95°C for 10 minutes, followed by 40 cycles of denaturation at 95°C for 15, and annealing at 60°C for 60 seconds.

Statistical analysis

Fisher's exact test was used for statistical analyses of the results of the immunohistochemical studies. The qRT-PCR findings were analyzed using GraphPad Prism 5 (GraphPad Software, Inc., La Jolla, CA). Differences among means were evaluated by a 2×2 contingency table using Fisher's exact test or ANOVA followed by

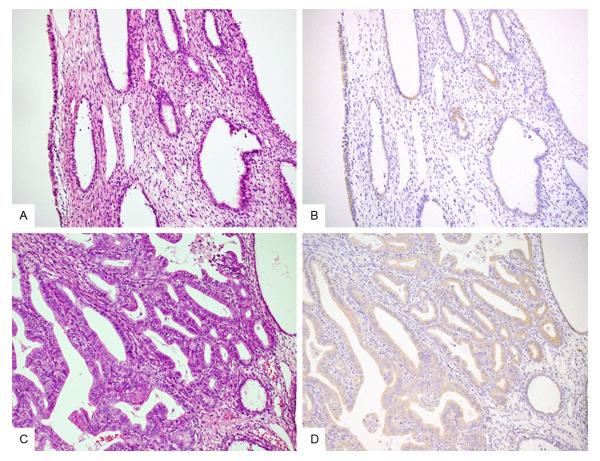


Figure 1. Immunohistochemical results of FGFR2IIIc in normal tissue and hyperplasia. As compared to normal endometrium (A: HE stain; B: immunostain; ×200), atypical hyperplasia cases (C: HE stain; D: immunostain; ×200) show significantly higher expression of FGFR2IIIc.

Table 2. Association of FGFR2IIIc expression asevaluated by intensity score (IS) with differenthistological types

	Normal endometrium	Hyperplasia	ECC
	n=10	n=10	n=47
High IS	1 (10%)	10 (100%)	32 (68%)
Low IS	9 (90%)	0	15 (32%)
P-value		p<0.001	p<0.01

Dunnett's post-hoc test. P<0.05 was considered significant. All results shown represent mean \pm SEM.

Results

In normal endometrium tissue, although all 10 cases showed FGFR2IIIc expression, the level in 9 (90%) of them was low (**Table 1**). In contrast, 80% of atypical hyperplasia and 49% of EEC cases showed high expression of FGFR2IIIc. As compared to normal tissue, FGFR2IIIc

expression in both atypical hyperplasia and ECC showed significantly higher expression (P<0.01 and P<0.05, respectively) (**Table 1**) (**Figure 1**). However, there was no significant difference between hyperplasia and ECC cases. Similar results were also observed using IS assessment (**Table 2**).

Clinicopathological features of each EEC case are listed in **Table 3**. The patients ranged in age from 32-84 years, with a median age of 57 years. Therefore, we divided the patients into 2 age groups: below 57 years (n=22) and 57 years or older (n=25). With regard to histological differentiation, 21 cases were G1, 15 cases were G2, and 11 cases were G3. Among them, 24 patients underwent lymphadenectomy, of whom 3 had lymph node involvement. Moreover, 31 patients had stage I tumors, 3 had stage II tumors, and 13 had stage III or higher stage tumors. In order to investigate the role of FGFR2IIIc in EEC, we analyzed the association between FGFR2IIIc expression level and clinico-

patients	
Clinicopathological features	Cases
Age (yrs)	
Under 57	22
57 or older	25
Grade	
1	21
2	15
3	11
LN metastasis	
Positive	3
Negative	21
No data	23
Stage	
I	31
II	3
III or higher	13

Table 3. Clinicopathological features of	
patients	

pathological features. In total, 23 patients (49%) showed high FGFR2IIIc expression (Table 4). Among them, (1) 10 patients were younger than 57 years and 13 were 57 years or older; (2) 14 patients had G1, 7 patients had G2, and 2 patients had G3 EEC; (3) 1 patient had lymph node metastasis while 8 were negative; and (4) 16 patients had stage I tumors and 7 had stage II or higher stage tumors. Although FGFR2IIIc expression was not statistically different between G1 and G2 EECs, G3 EEC showed significantly decreased FGFR2IIIc level as compared to G1 and G2 EECs (P<0.05) (Table 4) (Figure 2). Similar results were obtained using IS assessment (P<0.005) (Table 5). No association was detected between FGFR2IIIc expression and patient age, lymph node metastasis, or stage.

To determine FGFR2IIIc mRNA level, we performed qRT-PCR using FFPE tissue, and found that FGFR2IIIc mRNA levels in atypical hyperplasia and G1 EEC were significantly higher than that in normal endometrium (**Figure 3**). However, the FGFR2IIIc mRNA level in G3 EEC was significantly lower than that in G1 EEC and atypical hyperplasia, results which validate the immunohistochemical analysis.

Discussion

In EC, different histological types may have different precancerous status and may be associated with distinct molecular and genetic altera-

Table 4. Association between clinicopatho-	
logical features and TS	

logical leatures and 15			
Clinicopathological features	High TS	Low TS	P-value
Age (yrs)			NS*
Under 57 (n=22)	10 (45%)	12 (55%)	
57 or older (n=25)	13 (52%)	12 (48%)	
Grade			p<0.05
1 and 2 (n=36)	21 (78%)	15 (22%)	
3 (n=11)	2 (18%)	9 (82%)	
LN metastasis			NS*
Positive (n=3)	1 (33%)	2 (67%)	
Negative (n=21)	8 (38%)	13 (62%)	
No data (n=23)	14 (61%)	9 (39%)	
Stage			NS*
I (n=31)	16 (52%)	15 (48%)	
II or higher (n=16)	7 (44%)	9 (56%)	
*: not significant			

*: not significant.

tions. EEC occurs most frequently in pre- and peri-menopausal women and is strongly associated with excessive estrogen exposure. EEC is usually found to coexist or succeed atypical endometrial hyperplasia, also known as precancerous lesion. The development of EEC is a multistep process, and each step involves accumulation of genetic aberrations. Genomic instability such as microsatellite instability and chromosomal aneuploidy, inactivation of tumor suppressor genes such as PTEN and p53, and activation of oncogenes such as K-Ras and β -catenin are important events in this process [21, 22].

Recently, the FGF/FGFR signalling pathway has attracted attention as an important mechanism of carcinogenesis and tumor development. According to the study of Soufla et al, FGF2 up-regulation was found to be strongly related to endometrial carcinogenesis [23]. FGFR2 is one of the most important receptors of the FGF/FGFR pathway. FGFR2 single-nucleotide polymorphisms are strongly related to the oncogenesis of breast cancer and EC [10, 24-27]. According to the recent report by Byron et al, activating mutations in FGFR2 were found in 16% of EC cases, and up-regulated FGFR2 mRNA expression was observed in these EC specimens [7]. In EC cells with activated FGFR2, knockdown of FGFR2 induces cell death, suggesting that FGFR2 is important to EC cell proliferation [7]. FGFR2IIIc activation also drives

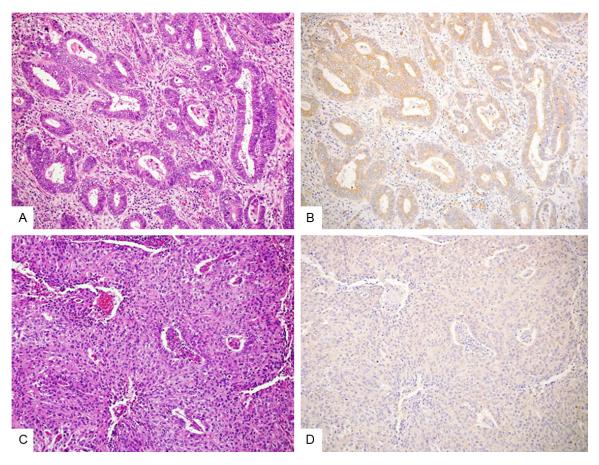


Figure 2. Immunohistochemical results of FGFR2IIIc in G1 and G3 EECs. G1 EEC (A: HE stain; B: immunostain; ×200) shows high FGFR2IIIc expression, but G3 EEC (C: HE stain; D: immunostain; ×200) shows significantly decreased expression of FGFR2IIIc.

Table 5. Association betweetures and IS	en clinico	pathologic	cal fea-
Clinicopathological features	High IS	Low IS	P-value
			NC*

Clinicopathological features	High IS	Low IS	P-value
Age (yrs)			NS*
Under 57 (n=22)	15 (65%)	7 (35%)	
57 or older (n=25)	17 (68%)	8 (32%)	
Grade			p<0.005
1 and 2 (n=36)	29 (81%)	7 (19%)	
3 (n=11)	3 (27%)	8 (73%)	
LN metastasis			NS*
Positive (n=3)	2 (67%)	1 (33%)	
Negative (n=21)	12 (57%)	9 (43%)	
No data (n=23)	18 (78%)	5 (22%)	
Stage			NS*
I (n=31)	20 (65%)	11 (35%)	
II or higher (n=16)	12 (75%)	4 (25%)	

*: not significant.

cell proliferation. In prostate cancer, FGFR2IIIc expression correlates with tumor progression

[28, 29]. In rat bladder cancer cells, FGFR2IIIc expression is correlated with epithelial-to-mesenchymal transition, a phenomenon that is associated with tumor progression and invasion [30]. Furthermore, immunohistochemical analysis of uterine cervical tissue revealed a correlation between FGFR2IIIc expression and the progression of cervical dysplasia. The same result was obtained in an in-situ hybridization validation study [20], suggesting that abnormal FGFR2IIIc expression is an early event in uterine cervix carcinogenesis, and that immunohistochemical staining is a convenient and reliable method to evaluate FGFR2IIIc expression in human tumor tissue.

In this study, we examined the expression and localization of FGFR2IIIc in normal endometrial tissue, atypical hyperplasia, and EEC. Significantly higher FGFR2IIIc expression

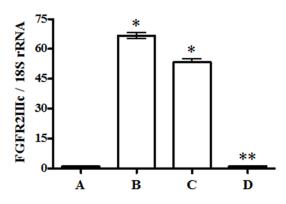


Figure 3. mRNA expression of FGFR2IIIc in tissues. Tissues from atypical hyperplasia (B) and G1 EEC (C) show higher FGFR2IIIc mRNA expression than normal tissue (A). However, G3 EEC (D) shows significantly decreased expression compared with G1 EEC. The data represent mean \pm SEM (n=3). *P<0.05 versus A and D. **P<0.05 versus B and C.

was observed in atypical hyperplasia and G1 EEC. This provides evidence that FGFR2IIIc expression increases in endometrial tissue as it progresses from normalcy to atypical hyperplasia, and maintains a high level of expression in G1 EEC. This indicates that FGFR2IIIc plays an important role in the carcinogenesis of EEC, and that alteration in its expression may occur during the atypical hyperplasia stage. In the early stage of EEC, high FGFR2IIIc expression is important for tumor growth and maintenance. However, G3 EEC showed significantly lower FGFR2IIIc expression than G1 and G2 EECs. suggesting that FGFR2IIIc may not be related to tumor progression. This result is different from that in prostate and bladder cancers [28-30], suggesting that FGFR2IIIc may play distinct roles in different tissues. Our data are supported by investigations of Voss et al. as well as Kuwabara et al. who found that G3 EEC have similar clinical, immunohistochemical, and prognostic characteristics with other endometrial cancers such as papillary serous carcinoma and clear cell carcinoma rather than G1 and G2 ECCs [4, 5]. More genetic aberration accumulations are needed to drive EEC progression and some G3 EEC cases may develop through different carcinogenic processes than G1 and G2 EECs.

G3 differentiation is considered one of the strongest predictors of recurrence and progression in EEC [31, 32]. Patients with G3 EEC are recommended for more aggressive therapy. Therefore, correct assessment of grade differ-

entiation is important. It is very common that grade differentiation is often misinterpreted using curettage biopsy specimens because of the limited amount of the sample. Our results show that high FGFR2IIIc expression correlates with G1 and G2 EECs, especially using the immunostain intensity evaluation, but not with G3 ECC. Using limited biopsy specimens, immunohistochemical results for FGFR2IIIc may not only aid in predicting the risk for recurrence and prognosis of patients, but also in deciding the treatment plan.

Recent reports revealed that endometrial cancer cell lines with activating FGFR2 mutations are selectively sensitive to the pan-FGFR inhibitor, PD173074 [32]. Targeted therapy for ECs with overexpression of FGFR2 is under consideration. FGFR2IIIc is an important isoform of FGFR2 and may serve as another good candidate for targeted therapy with pan-FGFR inhibitors in EEC. Additional studies are needed to determine the association between high FGF-R2IIIc expression in EEC and susceptibility to FGFR2 inhibitors.

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Disclosure of conflict of interest

There are no conflicts of interest to declare.

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References

- Mutter GL. Histopathology of genetically defined endometrial precancers. Int J Gynecol Pathol 2000; 19: 301-309.
- [2] Boruban MC, Altundag K, Kilic GS, Blankstein J. From endometrial hyperplasia to endometrial cancer: insight into the biology and possible

medical preventive measures. Eur J Cancer Prev 2008; 17: 133-138.

- [3] Zakharov V, Lin HK, Azzarello J, McMeekin S, Moore KN, Penning TM, Fung KM. Suppressed expression of type 2 3alpha/type 5 17beta-hydroxysteroid dehydrogenase (AKR1C3) in endometrial hyperplasia and carcinoma. Int J Clin Exp Pathol 2010; 3: 608-617.
- [4] Voss MA, Ganesan R, Ludeman L, McCarthy K, Gornall R, Schaller G, Wei W, Sundar S. Should grade 3 endometrioid endometrial carcinoma be considered a type 2 cancer-a clinical and pathological evaluation. Gynecol Oncol 2012; 124: 15-20.
- [5] Kuwabara Y, Susumu N, Banno K, Hirao T, Kawaguchi M, Yamagami W, Suzuki N, Aoki D, Nozawa S. Clinical characteristics of prognostic factors in poorly differentiated (G3) endometrioid adenocarcinoma in Japan. Jpn J Clin Oncol 2005; 35: 23-27.
- [6] Langheinrich MC, Schellerer V, Perrakis A, Lohmüller C, Schildberg C, Naschberger E, Stürzl M, Hohenberger W, Croner RS. Molecular mechanisms of lymphatic metastasis in solid tumors of the gastrointestinal tract. Int J Clin Exp Pathol 2012; 5: 614-623.
- [7] Byron SA, Gartside MG, Wellens CL, Mallon MA, Keenan JB, Powell MA, Goodfellow PJ, Pollock PM. Inhibition of activated fibroblast growth factor receptor 2 in endometrial cancer cells induces cell death despite PTEN abrogation. Cancer Res 2008; 68: 6902-6907.
- [8] Heiskanen M, Kononen J, Bärlund M, Torhorst J, Sauter G, Kallioniemi A, Kallioniemi O. CGH, cDNA and tissue microarray analyses implicate FGFR2 amplification in a small subset of breast tumors. Anal Cell Pathol 2001; 22: 229-234.
- [9] Shin EY, Lee BH, Yang JH, Shin KS, Lee GK, Yun HY, Song YJ, Park SC, Kim EG. Up-regulation and coexpression of fibroblast growth factor receptors in human gastric cancer. J Cancer Res Clin Oncol 2000; 126: 519-528.
- [10] Dutt A, Salvesen HB, Chen TH, Ramos AH, Onofrio RC, Hatton C, Nicoletti R, Winckler W, Grewal R, Hanna M, Wyhs N, Ziaugra L, Richter DJ, Trovik J, Engelsen IB, Stefansson IM, Fennell T, Cibulskis K, Zody MC, Akslen LA, Gabriel S, Wong KK, Sellers WR, Meyerson M, Greulich H. Drug-sensitive FGFR2 mutations in endometrial carcinoma. Proc Natl Acad Sci U S A 2008; 105: 8713-8717.
- [11] Davies H, Hunter C, Smith R, Stephens P, Greenman C, Bignell G, Teague J, Butler A, Edkins S, Stevens C, Parker A, O'Meara S, Avis T, Barthorpe S, Brackenbury L, Buck G, Clements J, Cole J, Dicks E, Edwards K, Forbes S, Gorton M, Gray K, Halliday K, Harrison R, Hills K, Hinton J, Jones D, Kosmidou V, Laman R, Lugg R,

Menzies A, Perry J, Petty R, Raine K, Shepherd R, Small A, Solomon H, Stephens Y, Tofts C, Varian J, Webb A, West S, Widaa S, Yates A, Brasseur F, Cooper CS, Flanagan AM, Green A, Knowles M, Leung SY, Looijenga LH, Malkowicz B, Pierotti MA, Teh BT, Yuen ST, Lakhani SR, Easton DF, Weber BL, Goldstraw P, Nicholson AG, Wooster R, Stratton MR, Futreal PA. Somaticmutations of the protein kinase gene family in human lung cancer. Cancer Res 2005; 65: 7591-7595.

- [12] Jang JH, Shin KH, Park JG. Mutations in fibroblast growth factor receptor 2 and fibroblast growth factor receptor 3 genes associated with human gastric and colorectal cancers. Cancer Res 2001; 61: 3541-3543.
- [13] Miki T, Bottaro DP, Fleming TP, Smith CL, Burgess WH, Chan AM, Aaronson SA. Determination of ligand-binding specificity by alternative splicing: two distinct growth factor receptors encoded by a single gene. Proc Natl Acad Sci U S A 1992; 89: 246-250.
- [14] Chaffer CL, Brennan JP, Slavin JL, Blick T, Thompson EW, Williams ED. Mesenchymal-toepithelial transition facilitates bladder cancer metastasis: role of fibroblast growth factor receptor-2. Cancer Res 2006; 66: 11271-11278.
- [15] Valve E, Martikainen P, Seppänen J, Oksjoki S, Hinkka S, Anttila L, Grenman S, Klemi P, Härkönen P. Expression of fibroblast growth factor (FGF)-8 isoforms and FGF receptors in human ovarian tumors. Int J Cancer 2000; 88: 718-725.
- [16] Drugan CS, Paterson IC, Prime SS. Fibroblast growth factor receptor expression reflects cellular differentiation in human oral squamous carcinoma cell lines. Carcinogenesis 1998; 19: 1153-1156.
- [17] Cha JY, Lambert QT, Reuther GW, Der CJ. Involvement of fibroblast growth factor receptor 2 isoform switching in mammary oncogenesis. Mol Cancer Res 2008; 6: 435-445.
- [18] Kwabi-Addo B, Ropiquet F, Giri D, Ittmann M. Alternative splicing of fibroblast growth factor receptors in human prostate cancer. Prostate 2001; 46: 163-172.
- [19] Marek L, Ware KE, Fritzsche A, Hercule P, Helton WR, Smith JE, McDermott LA, Coldren CD, Nemenoff RA, Merrick DT, Helfrich BA, Bunn PA Jr, Heasley LE. Fibroblast growth factor (FGF) and FGF receptor-mediated autocrine signalling in non-small-cell lung cancer cells. Mol Pharmacol 2009; 75: 196-207.
- [20] Kawase R, Ishiwata T, Matsuda Y, Onda M, Kudo M, Takeshita T, Naito Z. Expression of fibroblast growth factor receptor 2IIIc in human uterine cervical intraepithelial neoplasia and cervical cancer. Int J Oncol 2010; 36: 331-340.

- [21] Kapucuoglu N, Aktepe F, Kaya H, Bircan S, Karahan N, Ciris M. Immunohistochemical expression of PTEN in normal, hyperplastic and malignant endometrium and its correlation with hormone receptors, bcl-2, bax, and apoptotic index. Pathol Res Pract 2007; 203: 153-162.
- [22] Liu F. Molecular carcinogensis of endometrial cancer. J Obstet Gynecol 2007; 64: 26-32.
- [23] Soufla G, Sifakis S, Spandidos DA. FGF2 transcript levels are positively correlated with EGF and IGF-1 in the malignant endometrium. Cancer Lett 2008; 259: 146-155.
- [24] Eswarakumar VP, Lax I, Schlessinger J. Cellular signaling by fibroblast growth factor receptors. Cytokine Growth Factor Rev 2005; 16: 139-149.
- [25] Katoh Y, Katoh M. FGFR2-related pathogenesis and FGFR2-targeted therapeutics. Int J Mol Med 2009; 23: 307-311.
- [26] Katoh M. Cancer genomics and genetics of FGFR2. Int J Oncol 2008; 33: 233-237.
- [27] Mutch DG. The new FIGO staging system for cancers of the vulva, cervix, endometrium and sarcomas. Gynecol oncol 2009; 115: 325-328.
- [28] Oltean S, Sorg BS, Albrecht T, Bonano VI, Brazas RM, Dewhirst MW, Garcia-Blanco MA. Alternative inclusion of fibroblast growth factor receptor 2 exon Illc in Dunning prostate tumors reveals unexpected epithelial mesenchymal plasticity. Proc Natl Acad Sci U S A 2006; 103: 14116-14121.

- [29] Yan G, Fukabori Y, McBride G, Nikolaropolous S, McKeehan WL. Exon switching and activation of stroma and embryonic fibroblast growth factor (FGF)-FGF receptor genes in prostate epithelial cells accompany stromal independence and malignancy. Mol Cell Biol 1993; 13: 4513-4522.
- [30] Baum B, Settleman J, Quinlan MP. Transitions between epithelial and mesenchymal states in development and disease. Semin Cell Dev Biol 2008; 19: 294-308.
- [31] Lutman CV, Havrilesky LJ, Cragun JM, Secord AA, Calingaert B, Berchuck A, Clarke-Pearson DL, Soper JT. Pelvic lymph node count is an important prognostic variable for FIGO stage I and II endometrial carcinoma with high-risk histology. Gynecol oncol 2006; 102: 92-97.
- [32] Byron SA, Pollock PM. FGFR2 as a molecular target in endometrial cancer. Future Oncology 2009; 5: 27-32.