E6/E7 proteins are potential markers for the screening and diagnosis of cervical pre-cancerous lesions and cervical cancer in a Chinese population

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Received October 6, 2015; Accepted March 14, 2017

DOI: 10.3892/ol.2017.6932

Abstract. The present prospective study aimed to evaluate the effects of E6/E7 protein detection by western blotting on cervical cancer (CC) early screening compared with detection by Hybrid Capture 2 (HC2) test and ThinPrep cytological test (TCT) in a Chinese population. A total of 450 cases of cervical intraepithelial neoplasia (CIN) suspected samples (positive in ≥ 1 indicator of TCT and HC2 test) were recruited from women who were treated at the International Peace Maternity and Child Health Hospital (Shanghai, China) from March 2014 to February 2015. Each sample was analyzed by cytological test. In addition, human papillomavirus (HPV) DNA examination by Hybrid Capture Tube test and E6/E7 protein expression detection by western blotting were performed in all samples, as well as histologic diagnosis to determine the stage of CIN. The results revealed that, for the diagnosis of CIN2+, although the sensitivity of E6/E7 protein detection was lower than that of HC2 test (71.3 vs. 96.6%, respectively), the specificity was markedly improved (67.6 vs. 5.9%, respectively). Compared with that of TCT, the sensitivity of E6/E7 protein detection was much higher (36.2 vs. 71.3%, respectively), but the specificity was lower (88.2 vs. 67.6%, respectively). In the present study, HPV E6/E7 protein expression was evaluated as a potential new biomarker for CC, with satisfactory diagnostic values for HPV types 16 and 18. The relative diagnostic value may be further improved by combination of E6/E7 messenger RNA detection.

Introduction

Cervical cancer (CC), which is the third most common cause of cancer in women in the world, is mainly associated with infection by human papillomavirus (HPV), which is a small, circular, double-stranded DNA virus that is also associated with cervical neoplasia, anogenital warts and other anogenital cancers (1,2). According to previous reports from the International Agency for Research on Cancer, HPV could be classified into high-risk (HR) types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 68, 73 and 82), probable HR types (26, 53 and 66) and low-risk types (6, 11, 40, 42, 43, 44, 54, 61, 70, 72, 81 and CP6108) (3).

HR HPV, which is considered to be a necessary biological factor for CC, can express specific viral genes such as E6 and E7, which have been observed in CC cell lines and biopsies (4,5). These specific viral genes express oncoproteins (such as E6 and E7 proteins) that can be inserted into the host's cell genome (6). Therefore, E6 and E7 proteins may be critical for identifying HR HPV, and consequently could be a potential new biomarker for CC diagnosis.

Currently, the established methods for the screening of CC in clinical trials include Pap test, ThinPrep cytological test (TCT), HPV DNA determination by Hybrid Capture 2 (HC2) test, HPV messenger RNA (mRNA) determination by PreTect HPV-Proofer and colposcopy (7). However, due to the low sensitivity of the aforementioned methods, females with high risk of developing CC require regular re-tests to confirm the accuracy of the results (7). Among patients with negative cytology results but positive HPV DNA test results, a small percentage will develop CC (8). Therefore, an efficient diagnostic method is required for these particular patients (TCT/HPV⁺).

The present study aimed to evaluate E6/E7 proteins as the main biomarkers of pre-CC lesions, and to determine the suitability of E6/E7 protein detection as a potential screening method for CC diagnosis.

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Key words: E6/E7 proteins, HPV DNA, HPV mRNA, prognostic detection, diagnosis

Materials and methods

Subjects. A total of 450 samples from female patients with suspected cervical intraepithelial neoplasia (CIN) (positive in ≥ 1 indicator of TCT and HC2 test, excluding carcinoma patients) according to routine physical examination were collected from the Cervical Department (International Peace Maternity and Child Health Hospital, Shanghai, China) from March 2014 to February 2015. Of these patients, 348 were diagnosed as CIN2⁺ and the rest were diagnosed as CIN2⁻. The patients were aged from 18 to 45 years old, and had not been subjected to any particular treatment, including hysterectomy, radiotherapy or chemotherapy. Each sample was analyzed by cytological test, HPV DNA detection by polymerase chain reaction and E6/E7 protein expression detection by western blotting (9,10). In addition, histologic diagnosis was applied to determine the stage of CIN (11). Since histopathology is recognized as the gold standard for tumor diagnosis, the results of E6/E7 protein detection, HPV DNA evaluation and TCT were compared with the histopathologic results in terms of their sensitivity, specificity, positive predictive value (PPV) and negative predictive value (NPV) (11).

Preparation and preservation of samples. Cervical TCT samples obtained from biopsy were prepared in two formats, either using PreservCyt® Solution (Hologic Inc., Bedford, MA, USA) or SurePath[™] Preservative Solution (TriPath Imaging Inc., Burlington, NC, USA). A total of 2 ml cervical specimen preparation (containing $\sim 75 \,\mu$ l cell pellets per 1 ml solution) were collected in 20 ml PreservCyt® Solution as specimens for TCT. All samples were maintained at -20°C prior to use, and were tested within 1-2 weeks after collection. Specimens with abnormal TCT findings were rinsed into Specimen Transport Medium (Digene Corporation, Gaithersburg, MD, USA) to detect HPV DNA by Hybrid Capture Tube test (HCT; Digene Corporation) following the manufacturer's protocol for PreservCyt® Solution-containing specimens. The specimens were stored at 4°C. The threshold of positive HPV detection was a relative light unit/cut-off ratio (RLU/CO) of \geq 1.0. Samples containing <5 µl cell precipitate were discarded.

TCT. TCT was used to identify high-grade squamous intraepithelial lesions (HSILs) (7). A liquid based cell plastic brush was inserted into the cervix at the squamo-columnar junction, and was rotated 5-8 circles. The exfoliated cells were stored in liquid-based cell preservation solution purchased from Beijing TCT Medical Technology Co., Ltd. (Beijing, China). The ultra-thin coating was made by ThinPrep 2000 system (Cytyc Corporation, Marlborough, MA, USA). The cells were fixed with 95% alcohol aqueous solution for 15 min at room temperature and pap stained for 3 min at room temperature, and then observed under the 15JF microscope (magnification, x40) from Shanghai CSOIF Co., Ltd. (Shanghai, China). Liquid-based cell plastic brushes and liquid-based cell preservation solution were purchased from Beijing TCT Medical Technology Co., Ltd. (Beijing, China). All steps (sample collection, preparation and final staining) were followed according to the manufacturer's protocol (Beijing TCT Medical Technology Co., Ltd.). The samples were classified into five grades according to The Bethesda System: Within normal limitation (normal); atypical squamous cells with undetermined significance; low grade squamous intraepithelial lesion (LSIL); HSIL; and ASCs where HSIL cannot be excluded (12).

HC2 test. HC2 test (Qiagen Sciences, Inc., Gaithersburg, MD, USA), a signal amplification test based on the hybridization of a RNA probe cocktail for 13 high-risk oncogenic types with the target DNA, and capture and detection of the DNA-RNA hybrid by chemiluminescence, was applied to quantitate the level of HPV DNA for its standardization approved by the USA Food and Drug Administration (13). This test was performed with 4 ml of PreservCyt samples, and then mixed with 400 µl transfer buffer (digeneHC2HPVDNA detection kit; Qiagen Sciences, Inc.). Following centrifugation (135 x g for 15 min at 25°C), 75 μ l of the supernatant was added to the microwell plate in a water bath at 65°C for 15 min. The solution was mixed for 30 sec, and then kept in a water bath at 65°Cfor 15 min. Denaturant (150 µl; digeneHC2HPVDNA detection kit; Qiagen Sciences, Inc.) was then added to the mixture. Following TCT, the residual samples were denatured by asymmetric PCR (Piko[®] Thermal Cycler 96-well system; Thermo Fisher Scientific, Inc., Waltham, MA, USA) to obtain single-stranded DNA (13), which was then mixed and reacted with an RNA probe cocktail (BD Pharmingen; BD Biosciences, San Jose, CA, USA) for 13 HR-HPV oncogenic types (16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59 and 68) to capture oncogenic HPV subtypes using DML-2000 gene hybridization amplifiers (Digene Corporation). The type of positive point HPV was determined according to the distribution pattern of HPV. RLU/CO values of ≥1.0 were determined as positive. All samples were evaluated in triplicate.

Western blot assay. The human cervical cancer cells obtained from biopsy were frozen for 2 days. Protein extraction was performed from frozen cells by adding RIPA Rapid Cell Lysates (catalog no. BYL40825; Shanghai Jierdun Biotech Co., Ltd., Shanghai, China), which contained Halt Protease and Phosphatase Inhibitor Cocktail (Thermo Fisher Scientific, Inc.) as previously described (14). The protein concentration in the cell lysates was determined with Pierce BCA Protein Assay kit (catalog no. PICPI23223; Thermo Fisher Scientific, Inc.). A total of 25 μ g protein was loaded on 15% polyacrylamide gels, according to the molecular weight of E6/E7 proteins (HPV16 E6/E7 and HPV18 E6, 17 kDa; and HPV18 E7, 12 kDa), for electrophoresis as described previously (15). GAPDH was applied as an internal control to confirm equal loading of cell lysates. Upon electrophoresis, samples were transferred to polyvinylidene fluoride membranes and blocked with 5% skimmed milk powder diluted in PBS-Tween-20 buffer. Next, primary antibodies (C1P5; catalog no., sc-460) against the E6/E7 proteins of HPV16 (HPV16 E6 dilution, 1:200; and HPV16 E7 dilution, 1:1,500), antibodies (BF7; catalog no., ab20192) against the E6 proteins of HPV18 or antibodies (8E2; catalog no., ab100953) against the E7 proteins of HPV18 (HPV18 E6; dilution, 1:600; and HPV18 E7 dilution, 1:1,000) for 1 h at room temperature were added, followed by addition of horseradish peroxidase-labeled secondary antibodies (dilution 1:1,000) incubated for 1 h at room temperature. All

Characteristic	CIN2+ (n=348)	CIN2 ⁻ (n=102)	P-value
Mean age ± SD, years	39.7±8.9	42.2±9.7	0.166
HPV E6/E7 protein detection, n (%)			< 0.001
Positive	248 (71.3)	33 (32.4)	
Negative	100 (28.7)	69 (67.6)	
HR-HPVHC2 test, n (%)			0.270
Positive	336 (96.6)	96 (94.1)	
Negative	12 (3.4)	6 (5.9)	
TCT, n (%)			< 0.001
HSIL	126 (36.2)	12 (11.8)	
LSIL	108 (31.0)	39 (38.2)	
ASC-H	48 (13.8)	6 (5.9)	
ASC-US	45 (12.9)	21 (20.6)	
Normal	21 (6.0)	24 (23.5)	

Table I. Characteristics of the patients according to histological diagnosis.

CIN, cervical intraepithelial neoplasia; SD, standard deviation; HPV, human papillomavirus; HC2, Hybrid Capture 2; TCT, ThinPrep cytological test; HSIL, high-grade squamous intraepithelial lesion; LSIL, low-grade squamous intraepithelial lesion; ASC, atypical squamous cell; US, undetermined significance; ASC-H, ASCs where HSIL cannot be excluded.

antibodies, together with anti-GAPDH antibody (6C5; catalog no., ab8245) were purchased from Abcam (Cambridge, UK), with the exception of anti-HPV16 E6 antibody, which was purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Subsequently, samples were subjected to enhanced chemiluminescence detection (GE Healthcare, Chicago, IL, USA). Finally, the representative gray value of protein expression was scanned (Fig. 1) and analyzed by Quantity One 1-D analysis software version 4.6.9 (Bio-Rad Laboratories, Inc., Hercules, CA, USA). Based on the positive/negative value of 1,393 volunteers from the Pap test between April 2012 and February 2015 in south China, the E6/E7 for the western blot analysis with an optical density under 450 nm value of $\geq 0.4 \pm 0.005$ were considered (16). The test was used to observe the exfoliated cells from the cervix fixed with physiological saline under the microscope.

Histological diagnosis. Histologic diagnosis, which is deemed as the gold standard for the diagnosis of CC, was used in the present study to clarify the stage of CIN with CIN2⁺ serving as the disease endpoint (17). The main steps were performed according to the procedure described by Ratnam *et al* (17). Cervical biopsy results read by one or more pathologists were obtained from participating centers and accepted as the disease endpoint for the study purposes. Pathologists were blinded to HPV results.

Statistical analysis. Categorical variables were assessed in the form of percentages. Comparison among groups based on percentage values was assessed for statistical significance. Differences in the quantitative values (such as age) between groups were analyzed by the Student's t-test. Statistical analysis was performed by SPSS13.0 for Windows (SPSS, Inc., Chicago, IL, USA). P<0.05 was considered to indicate a statistically significant difference.

Results

Results of histological diagnosis. The results from all four methods are shown in Table I. The proportion of E6/E7 protein-positive patients with CIN2⁺ was >2-fold higher than that of CIN2⁻ patients, which was statistically significant (P=0.000). However, this high specificity was not observed in HC2-positive patients with high risk of developing CC, although the sensitivity of the HC2 test was much higher than E6/E7 protein-positive patients (96.6 vs. 71.3%). According to the results of TCT, only HSIL⁺ cases were statistically significant (P<0.0001).

Comparison of diagnostic value of HPV E6/E7 proteins with other tests about sensitivity and specificity. For the diagnosis of CIN2⁺ cases in Table II, the specificity of E6/E7 protein detection was markedly improved compared with that of the HC2 test (67.6 vs. 5.9%), although the sensitivity was lower (71.3 vs. 96.6). Compared with that of TCT, the sensitivity of E6/E7 protein detection was much higher (71.3 vs. 36.2%), while the specificity was lower (67.6 vs. 88.2%). According to previous studies, the sensitivity of E6 protein detection by OncoE6 Cervical Test was only 42.4%, which is lower than the 71.3% of HPV E6/E7 protein detection, but it exhibited 99.1% specificity for E6 protein detection (18). The sensitivity of E6/E7 mRNA detection gradually increases when the number of HR-HPV types increases (17,19,20). When the number of HR-HPV types was >9, the sensitivity was >90%, with a satisfactory specificity (19,20). In addition, the sensitivity of HPV E6/E7 protein detection was higher than that of E6/E7 mRNA detection (71.3 vs. 56.3%, respectively) with regard to HPV types 16 and 18 (19).

As shown in Table III, the specificity of E6/E7 protein detection for patients with CIN2⁺ plus ASC-US or LSIL was higher than that of HPV DNA testing, suggesting that

Table II. Diagnostic v	alue of HPV E6/E7 protei	in detection, HC2 tes	t and TCT or	n high-grade histological dia	gnosis (positive cervical intra	epithelial neoplasia2).	
Indicator	HPV type	Triage	Cases, n	Sensitivity ^a , % (95% CI)	Specificity ^b , % (95% CI)	PPV°, % (95% CI)	NPV ^d , % (95% CI)
HPV E6/E7 proteins HPV DNA Morphocytology	16 and 18 13 types of HR-HPV -	Western blotting HC2 test TCT	450 450 450	71.3 (66.5-76.0) 96.6 (94.6-98.5) 36.2 (31.1-41.3)	67.6 (58.4-76.9) 5.9 (1.2-10.5) 88.2 (81.2-94.6)	88.3 (84.5-92.0) 77.8 (73.8-81.7) 91.3 (86.5-96.1)	40.8 (33.3-48.3) 33.3 (9.2-57.5) 28.8 (23.8-33.9)
"Sensitivity=true positiv tive + false negative). Hl value; 13 types of HR-H	e/(true positive + false nega PV, human papillomavirus; F PV: 16, 18, 31, 33, 35, 39, 4	ttive). ^b Specificity=true IR, high risk; HC2, Hyl 5, 51, 52, 56, 58, 59 an	negative/(true orid Capture 2; d 68 type HVI	: negative + false positive); ^c PF (TCT, ThinPrep cytological test P.	V=true positive/(true positive + ; CI, confidence interval; PPV, pc	false positive). ^d NPV=tr ositive predictive value; N	ue negative/(true nega- PV, negative predictive
Table III. Diagnostic v	/alue of HPV E6/E7 prote	sin detection and HC	2 test in CIN	2+ plus LSIL or ASC-US pa	tients.		
A, CIN2 ⁺ plus LSIL							
Indicator	HPV type	Triage	Cases, n	Sensitivity ^a , $\%$ (95% CI)	Specificity ^b , % (95% CI)	PPV°, % (95% CI)	NPV ^d , % (95% CI)
HPV E6/E7 proteins HPV DNA	16 and 18 13 types of HR-HPV	Western blotting HC2 test	147 147	77.8 (69.8-85.8) 97.2 (94.1-100.0)	56.4 (40.1-72.7) 7.7 (-1.1 to 16.4)	83.2 (75.8-90.6) 74.5 (67.2-81.8)	47.8 (32.8-62.8) 7.7 (-1.1 to 16.4)
B, CIN2 ⁺ plus ASC-U	S						
Indicator	HPV type	Triage	Cases, n	Sensitivity ^a , $\%$ (95% CI)	Specificity ^b , % (95% CI)	PPV°, % (95% CI)	NPV ^d , % (95% CI)
HPV E6/E7 proteins HPV DNA	16 and 18 13 types of HR-HPV	Western blotting HC2 test	66 66	78.4 (68.0-88.7) 93.5 (88.5-98.6)	66.7 (47.7-85.7) 11.1 (-1.6 to 23.8)	83.3 (73.1-93.6) 78.4 (70.6-86.2)	27.3 (16.2-38.3) 33.3 (-5.1 to 71.8)
^a Sensitivity=true positiv tive + false negative). H predictive value; LSIL, l type HVP.	e/(true positive + false nega PV, human papillomavirus; []] ow-grade squamous intraepi	tive). ^b Specificity=true HR, high risk; HC2, H. tithelial lesion; ASC, aty	negative/(true ybrid Capture 'pical squamou	 negative + false positive); ^cPF CIN, cervical intraepithelial us cell; US, undetermined signi 	V=true positive/(true positive + neoplasia; CI, confidence interve icance; 13 types of HR-HPV: 16	false positive). ^d NPV=tr ul: PPV, positive predictiv , 18, 31, 33, 35, 39, 45, 5	ue negative/(true nega- e value; NPV, negative 1, 52, 56, 58, 59 and 68

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Patients	Indicator	HPV type	Triage	Cases, n	Sensitivity ^a , % (95% CI)	Specificity ^b , % (95% CI)	PPV°, % (95% CI)	NPV ^d , % (95% CI)
CIN2 ⁺ plus HSIL ⁻ and HPV DNA ⁺ CIN2 ⁺ plus LSIL and HPV 16 DNA CIN2 ⁺ plus ASC-US and HPV 16 DNA	HPV E6/E7 proteins E6/E7 mRNA E6/E7 mRNA	16 and 18 16, 18, 31, 33 and 45 16, 18, 31, 33 and 45	Western blotting Proofer Proofer	295 254 103	71.8 (65.7-77.9) 67 (52-80) 72 (51-88)	58.3 (47.6-69.1) 72 (65-78) 74 (63-84)	81.4 (75.8-87.0) 36 (26-46) 47 (31-64)	45.0 (35.5-54.4) 90 (85-94) 89 (79-96)
"Sensitivity=true positive/(true positive + fa tive + false negative). HPV, human papillo atypical squamous cell; US, undetermined si	lse negative). ^b Specificit mavirus; CIN, cervical i ignificance; mRNA, mes	y=true negative/(true ne; ntraepithelial neoplasia; senger RNA; CI, confide	gative + false positi HSIL, high-grade s nce interval; PPV, po	ve); °PPV= quamous in ositive pred	true positive/(true p atraepithelial lesion; lictive value; NPV, n	ositive + false posi LSIL, low-grade a legative predictive v	tive). ^d NPV=true n squamous intraepitl value.	egative/(true nega nelial lesion; ASC

Table IV. Diagnostic value of HPV E6/E7 protein detection in CIN2+ patients with HSIL- and HPV DNA+.

Previous evidence indicated that E6/E7 protein detection exhibited better sensitivity than TCT or E6/E7 mRNA detection for the same HPV types, and better specificity than HPV DNA testing by HC2. For CIN2+ plus ASC-US or LSIL cases, the sensitivity and specificity of E6/E7 protein detection were similar to those of E6/E7 mRNA detection for the same HPV types. However, markedly different results were obtained in terms of PPV and NPV (83.2 vs. 37.1% for PPV; and 47.8 vs. 88.9% for NPV, respectively) (23). Therefore, E6/E7 protein detection exhibited a higher PPV than that of E6/E7 mRNA detection. The sensitivity and specificity of E6/E7 protein detection for CIN2+ patients with HSIL and positive HPV DNA testing were 71.8 and 58.3%, respectively (Table IV), while E6/E7 mRNA testing exhibited higher specificity and similar sensitivity compared with those of E6/E7 protein detection. According to previous studies on HPV mRNA detection, there may be no significant differences in terms of sensitivity or specificity between the screening indicators of E6/E7 mRNA and E6/E7 protein detection (21). However, the PPV in the E6/E7 protein test was higher than the NPV, while the NPV in the E6/E7 mRNA test was higher than the PPV.

Discussion

Approximately 565,000 new cases of CC occur in the world each year, and the incidence rate in developing countries is 3-fold higher than that in developed countries (24,25). Approximately 50% of all CC cases in the world were recorded in China and India (26). Nearly 181,500 people are diagnosed with CC each year in China, while >30,000 women succumbed to CC (27). In Europe, ~38,000 CC cases are diagnosed each year, and >2/3 of those would be expected to be cured and survive (28).

Usually, women with abnormal results of TCT or/and HPV DNA detection should undergo subsequent pathological biopsy in order to diagnose the existence and staging of CINs. Histology is recognized as the gold standard for diagnosing the pathological process of CC. CIN2⁺ is considered a precancerous lesion and, if left untreated, this may progress to cervical cancer; therefore, if a patient exhibits a CIN2⁺ lesion, this should be treated (17,29). However, it is unknown which kind of lesions will finally lead to infiltrative cancers. In developed countries, primary screening based on TCT could prevent >80% of CC. Abnormal diseases are often missed or misdiagnosed due to the limitations of testing sensitivity and sampling techniques (28,30).

As a result, numerous studies have focused on improving the techniques of CC screening. Chen *et al* (31) evaluated the efficacy of the Pap test combined with TCT in CC screening, and observed that the combination had high sensitivity and specificity. Subsequently, other reports revealed that the combined detection of HR-HPV by HC2 test and TCT may



Figure 1. Representative image of HPV16/18 E6/E7 protein detection by western blotting with specific antibodies against E6/E7 proteins and GAPDH. HPV, human papillomavirus. Patients in the Pap test with positive results were detected for western blot analysis, and numbers represent patients.

improve the sensitivity and specificity of CIN diagnosis and the prediction of its postoperative recurrence (32). In 2010, Ratnam *et al* (17) assessed Proofer and HC2 tests in a cross-sectional study, and noticed that Proofer is more specific than HC2 in identifying women with CIN2⁺, but has a lower sensitivity. The introduction of the above methods may improve the accuracy of CC screening, and may avoid a large number of unnecessary colposcopy and biopsy procedures, while effectively predicting the development of tumor lesions.

Although the aforementioned methods are popular and are recommended worldwide by guidelines for primary CC screening, a few potential patients are still missed and could not be dug out as detection of traditional activity showed lower sensitivity or specificity. Previous studies on the diagnosis of HPV and the occurrence of CC were usually conducted at the cellular, RNA or DNA level, with few studies performed at the protein level. Therefore, further evaluations are necessary for improved accurate screening.

In 2011, Ratnam *et al* (33) demonstrated that the sensitivity and specificity of the E6/E7 transcription-mediated amplification method for the detection of CIN2⁺ were 100.0 and 88.4%, respectively, in a study comprising 1,373 women. The Aptima HPV test, which detected E6/E7 mRNA of 14 oncogenic types, was more specific for detecting CIN2⁺ than the HC2 test. However, in that study, Aptima and HC2 testing were performed upon routine CC screening, subjects of which were confirmed CIN2⁺ already (33). Following infection by HPV, HPV E6/E7 proteins would be overexpressed upon HPV invasion into the host's cervical cells in the form of episomal HPV DNA or upon viral integration into the host's genome (34). These events were closely and directly associated with the development of cancer; thus, the major cause of pre-CC lesions appears to be the functional expression of HR-HPV E6/E7 proteins (34,35). Therefore, E6/E7 protein expression may be directly associated with CC risk.

A pilot clinical study on the application of OncoE6 Cervical Test indicated that HPV 16/18/45 E6 protein detection had higher specificity than HPV DNA testing for CIN3⁺ detection (36). In another clinical trial, this new technology was demonstrated to have higher specificity compared with that of HC2 testing [98.9%; 95% confidence interval (CI)=98.6-99.2% vs. 86.8%; 95% CI=85.9-87.7%, respectively], but lower sensitivity (67.3%; 95% CI=52.5-80.1% vs. 98.0%; 95% CI=89.1-99.9%, respectively) for CIN3⁺ detection only (37). Based on those findings, the present study introduced a new type of biomarker, E6/E7 protein detection, and evaluated this new methodology for the diagnosis and progression determination of cervical pre-cancerous lesions and CC.

In the present study, the combination of E6 and E7 protein testing could enhance the accuracy of the test; thus, the differences between E6/E7 protein detection, HC2 test and TCT were assessed in a Chinese population. According to the present results, the application of E6/E7 protein detection in CC screening could reduce the limitations of other recent technologies, such as the Pap test and HC2 test, and the indicators (E6/E7 proteins) for both cervical pre-cancer and progression to CC could be quantified with the standard detection method established in the present study (17,21). However, not all types of anti-E6/E7 monoclonal antibodies exhibit high specificity currently (8,38). Therefore, further studies about the combination of E6/E7 mRNA and protein testing may be introduced to improve the specificity and sensitivity of CC screening. Further clinical studies with larger samples and multi-center studies are required to evaluate the efficiency of E6/E7 protein detection as a new indicator for the screening and diagnosis of CC.

In conclusion, the present study provided evidence that E6/E7 proteins may be potential new biomarkers with satisfactory diagnostic values for HPV types 16 and 18. The relative diagnostic value may be further improved by combination with E6/E7 mRNA detection. Furthermore, the number of HPV types being tested could be increased in the future.

Acknowledgements

The authors would like to thank the patients, volunteers, clinicians and registry staff involved in the project, as well as the Natural Science Foundation of China (grant no. 09ZR1435000) for providing financial support.

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