Clinical significance of immunocytochemistry for PIK3CA as a carcinogenesis-related marker on liquid-based cytology in cervical intraepithelial neoplasia

TOMOKO GOTO1, MASASHI TAKANO2, HIDENORI SASA2, HITOSHI TSUDA3, KAZUHIRO YAMAUCHI4 and YOSHIHIRO KIKUCHI2

1Department of Obstetrics and Gynecology, Saitama Medical College, 38 Morohongo, Moroyama-cho, Iruma-gun, Saitama 350-0495; Departments of 2Obstetrics and Gynecology, 3Pathology II, National Defense Medical College, 3-2 Namiki, Tokorozawa, Saitama 359-8513; 4Pathology and Cytology Center, PCL Japan, 1361-1 Matoba, Kawagoe, Saitama 350-1101, Japan

Received July 11, 2005; Accepted September 13, 2005

Abstract. The catalytic subunit α of phosphatidylinositol 3-kinase (PIK3CA) has been expected to play a role as an important oncogene in uterine cervical carcinoma, whose expression in non-invasive lesions has received considerable attention. We investigated the potential of PIK3CA as a carcinogenesis-related marker for early intraepithelial lesion of the uterine cervix in cytology samples. Seventy-four cases with abnormal cytology suggesting the existence of cervical intraepithelial neoplasia (CIN) lesions, whose findings were histologically confirmed, were selected; they consisted of 20 CIN1, 21 CIN2, and 33 CIN3, respectively. In addition, 17 normal cases, whose cervical cytology revealed no abnormality over three occasions, were selected. Liquid-based cytology specimens were applied for human papillomavirus (HPV) DNA typing and immunocytochemistry using three different antibodies for p16INK4a, Ki-67 and PIK3CA, respectively. The fraction of immunopositive cells on the slides was calculated and expressed as mean numbers. Receiver operating characteristic (ROC) plots were generated to determine the diagnostic accuracy of each immunocytochemistry test. The mean number of immunopositive cells in the CIN3 group was significantly higher than other groups for all three antibodies. Among all groups, PIK3CA showed a superior specificity to distinguish CIN3 from other groups. Comparison of three markers by ROC curves also revealed that PIK3CA provided the best method for distinguishing CIN3. Thus, expression of PIK3CA was observed in liquid-based cytology in CIN lesions, which suggested its diagnostic significance in addition to the use of routine cervical cancer smear and the HPV screening program.

Introduction

Uterine cervical cancer is one of the most common female cancers worldwide. Although a beneficial impact from cytological screening of the uterine cervix in reducing the incidence and mortality of uterine cervical cancer has been well recognized, cytology has a limited sensitivity and reproducibility as well as relatively frequent equivocal results. Also, Papanicolaou smear is affected by a certain rate of false-negative results (1). Considerable efforts have been made to improve the quality of primary screening of cervical cancer and there has been an important evolution with the introduction of new technology such as liquid-based cytology (LBC). Combination with human papillomavirus (HPV) testing has been encouraged because laboratory and epidemiological research suggests a strong association between HPV infection and cervical cancer (2,3). So far, the specificity of HPV testing is not high enough to be performed well in a primary screening setting (4). There is a strong demand for additional, more sensitive and specific markers to improve screening programs. Objective and decisive biomarkers would improve standardization and quality control for the diagnosis of cervical cancer.

Immunocytochemistry provides information that is closely related to the morphological information provided by cytology and histology. This could be a useful diagnostic asset because it is relatively easy and fast. Several studies have shown the feasibility of applying p16INK4a immunocytochemistry to cervical cytology (5-7). Sahebali et al showed that p16INK4a and Ki-67 immunocytochemistry were applied to liquid-based cytology, respectively (8,9). p16INK4a could have the same significant clinical utility as measuring HPV gene expression.
Ki-67 positivity has also been found to increase with the number of positive cells in dysplasia and carcinoma and can be applied as a surrogate marker for HPV-related neoplasia (10). Although infection with HPV is known to play a crucial role in the immortalization of cervical epithelial cells, HPV infection in women does not always develop dysplasia or carcinoma (11) and the long latency of the transition from CIN3 to carcinoma strongly suggests that other factors in addition to HPV infection are required for the malignant transformation of epithelial cells (12). Mutations affecting tumor-suppressor genes or cellular oncogenes may be promoting factors.

PIK3CA encodes the p110α catalytic subunit of phosphatidylinositol 3-kinase (PI3K), which is an important component of the lipid signaling pathway. The subsequent activation of the downstream serine/threonine protein kinase is involved in multiple cellular processes, including cell survival, proliferation and apoptosis, and has been related to carcinogenesis (13,14). Amplification and overexpression of PIK3CA in ovarian cancer and in squamous cell carcinoma of the neck and uterine cervix have been reported (15-17). Zhang et al (18) showed that frequent amplification of PIK3CA in primary cervical tumors might be functionally involved in cervical carcinogenesis. Although screening for cervical cancer precursors has evolved due to the introduction of new technology, there is no cytological study to test the PIK3CA function immunocytochemically in a clinical setting. In the present study, the significance of immunocytochemistry of PIK3CA using liquid-based samples was studied.

Materials and methods

Cases. Of the patients who visited the Department of Obstetrics and Gynecology at the National Defense Medical College Hospital between July 2002 and December 2003, 74 cases with abnormal cytology suggesting the existence of a CIN lesion were selected for the current study. All cytological findings were histologically confirmed by punch biopsy or conization within two months of cytological evaluation. The cases with CIN consisted of 20 CIN1, 21 CIN2, and 33 CIN3, respectively. In addition, 17 normal cases were selected from patients whose cervical cytology was annually analyzed and HPV-DNA testing was also performed with residual cell suspension. For each specimen, 6 ml was used to make 3 slides of thin layer liquid-based cytology (LBC) with the fully robotic AutoCyte PREP System (Tripath Imaging) and each slide was used for immunocytochemical staining and HPV typing.

Immunocytochemical staining and antibodies. Immunostaining for p16INK4a primary antibody, clone E6H4 (MTM Laboratories, Heidelberg, Germany), was performed using p16INK4a Research Kit (DakoCytomation, Glostrup, Denmark). Cells were regarded as positive if both the nucleus and the cytoplasm were stained.

Immunocytochemistry was performed using the prediluted primary anti-human Ki-67 mouse monoclonal antibody (DakoCytomation, 1:150 dilution). The slides were incubated for 60 min. Detection was accomplished using the EnVision+ system (DakoCytomation) for 60 min and incubation with DAB (DakoCytomation) for 5 min. All steps were performed at room temperature and slides were washed in phosphate-buffered saline. Slides were counterstained with hematoxylin. Cells were considered immunopositive if the nuclei showed homogeneous or punctuate staining. Cytoplasmic staining without nuclear staining was not considered to be positive.

Immunocytochemistry was performed using the prediluted primary goat polyclonal antibody against PI 3-kinase 110α (Santa Cruz Biotechnology, Inc., CA, USA, 1:200 dilution). The slides were incubated for 60 min, followed by labeling of LSAB+ System-HRP (DakoCytomation) and incubation with DAB (DakoCytomation) for 2 min. Slides were counterstained with hematoxylin. Cells were considered immunopositive if the cytoplasm showed homogeneous staining.

Evaluation of immunocytochemistry. The evaluation of immunocytochemistry was made by light microscopy. The staining intensity was not graded to avoid subjective interpretation. All cells in three high power fields (x400), which were considered to be representative of the whole cell area, were counted and the total amount of cells on the slide was calculated. This referred to the Bethesda system 2001 concerning the assessment of LBC cellularity (19). All positively stained cells in the whole cell area were counted and the fraction of positive cells on the slide was calculated by two independent pathologists with no knowledge of the patient's clinical information. This fraction was expressed as the number of immunopositive cells/1000 cells to compare the results of all samples.

HPV-DNA detection and typing. The remaining cell suspension was transferred to an Eppendorf tube and the cells were pelleted by centrifugation. DNA extraction was performed using QiaAmp DNA Blood Mini Kit (Qiagen, Tokyo, Japan) and its protocol. The DNA extracts were stored at -20°C until PCR was carried out. All samples were tested with L1-PCR consensus primers (L1C1, 5′-CGTAAACGTTTTTCCCTATT TTTT-3′; L1C2, 3′-GTTATGTCATATAAAATCCCAT-5′) (20). Typing of HPV-DNA was performed by the restriction fragment length polymorphism method and was carried out with type-specific PCRs for high-risk HPV (HR-HPV) types (HPV6, 11, 16, 18, 30, 31, 33, 34, 39, 42, 43, 44, 45, 51, 52, 53, 54, 55a, 55b, 56, 58, 59, 61, 66). Samples that were positive for the consensus PCR but negative for the type specific primers were classified as unidentified HPV genotypes of unknown malignant potential.

Statistical analysis. Data were analyzed using the software package, STATMATE III (Atoms, Tokyo, Japan). A p-value <0.05 was considered to be statistically significant. All
analyzed data are expressed as the mean number of immunopositive cells per 1000 cells and standard error of the mean (SEM). Receiver operating characteristic (ROC) plots were generated to determine the diagnostic accuracy of each immunocytochemistry test for CIN3 compared to CIN1, CIN2 and normal. The curve which comes close to the ideal values of 100% sensitivity and 100% specificity, which is the top left corner of the ROC graph, provided the best combination of sensitivity and specificity, assuming an equal importance of each. ROC curves were used to further establish optimal thresholds.

Results

Immunostaining. Immunopositive cells for p16\textsuperscript{INK4a} exhibited a medium-brown cytoplasm with a darker brown nucleus. Cytoplasmic staining was diffuse. Immunoreactive cells for Ki-67 showed dark brown, homogeneous or punctuated staining, limited exclusively to the nucleus. Immunopositivity for PIK3CA exhibited medium-brown homogeneous staining localized to the cytoplasm (Fig. 1).

Cytological diagnosis. Table I shows the distribution of the study group into CIN categories together with their age distribution and the number of infectious cases with HR-HPV types. The mean number of cells on a slide was 16,970 (95% CI = 15,005-18,936).

Table I. The study population.

<table>
<thead>
<tr>
<th></th>
<th>Number (%)</th>
<th>Mean age (range)</th>
<th>HR-HPV+ HPV positive/HPV negative (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal</td>
<td>17 (18.7)</td>
<td>38 (27-55)</td>
<td>17/33/67</td>
</tr>
<tr>
<td>CIN1</td>
<td>20 (22.0)</td>
<td>38 (21-54)</td>
<td>42/67/33</td>
</tr>
<tr>
<td>CIN2</td>
<td>21 (23.1)</td>
<td>36 (22-52)</td>
<td>50/58/42</td>
</tr>
<tr>
<td>CIN3</td>
<td>33 (36.3)</td>
<td>38 (26-57)</td>
<td>58/67/33</td>
</tr>
<tr>
<td>Total</td>
<td>91</td>
<td>38</td>
<td>45/60/40</td>
</tr>
</tbody>
</table>

*p16\textsuperscript{INK4a} immunostaining. When comparing the number of p16\textsuperscript{INK4a} immunopositive cells/1000 cells between the different CIN groups, the CIN3 group showed a significantly higher count (mean, 7.27; SEM, 1.30) than the normal group (mean, 0.35; SEM, 0.19; p<0.001), the CIN1 group (mean, 0.87; SEM, 0.31; p<0.001) and the CIN2 group (mean, 2.03; SEM, 0.44; p<0.01) (Fig. 2). In addition, the CIN2 group showed a significantly higher count of immunopositive cells than the normal group (p<0.01).

Ki-67 immunostaining. When comparing the number of Ki-67 immunopositive cells/1000 cells between the different CIN groups, the CIN3 group showed a significantly higher count (mean, 6.31; SEM, 0.98) than the normal group (mean, 0.58; SEM, 0.20; p<0.001), CIN1 group (mean, 2.34; SEM, 0.58; p<0.01) and CIN2 group (mean, 2.23; SEM, 0.72; p<0.01) (Fig. 2). The CIN1 and CIN2 group also showed higher mean counts of immunopositive cells than the normal group (p<0.05).

Figure 1. Representative immunopositive dysplastic cells for PIK3CA in a case of severe dysplasia. x40.

Figure 2. Mean number and standard error of the mean (SEM) of immunopositive cells/1000 cells were shown in normal and CIN lesions. Immunopositive cell counts in the CIN3 group were significantly higher than all other groups in each method. PIK3CA especially showed superior specificity to distinguish CIN3 from other groups. *p<0.05, **p<0.01, ***p<0.001.
However, only 10% of infected women develop cervical infection and cervical dysplasia/carcinoma is well established. The infection of cervical cells by HPV manifests itself in changes in the function or expression of host genes, and the detection of these alterations can play a role in screening and diagnosis. Actually, mild cytological abnormalities are extremely common in young women, but most of these lesions regress spontaneously, even when caused by oncogenic types of HPV (24,25) which play a crucial role in the pathogenesis of cervical cancer. This has prompted efforts to discover other biomarkers and other screening techniques with the potential to supplement cytological screening. Accordingly, there is a need to explore the screening system detecting women with lesions that need medical intervention, such as CIN3, but p16INK4a does not seem to be suitable for this because of its lower specificity. The development of assays with high specificity for detecting cancer precursors as well as high sensitivity would represent an attractive alternative as primary screening tests or as tests to complement cytology and HPV typing.

Redon et al (17) identified the PIK3CA gene as the most likely oncogene to be overexpressed as a consequence of 3q gain. PIK3CA, encoding the catalytic subunit p110α of PI3-kinase 1A, has recently been suggested to play a critical role as an oncogene in ovarian and cervical cancer (15,16). Information about abnormal PIK3CA gene status in squamous cell carcinoma (SCC) has predominantly been obtained from experiments on cell lines. Employing ovarian carcinoma cell lines and uterine cervix SCC cell lines, they both pointed out that increased PI3K activity is solely caused by gene-dependent overexpression of the catalytic subunit p110α. Woenckhaus et al (26) showed that not only invasive SCCs in the head and neck but also high-grade dysplasias showed PIK3CA copy number gain, providing strong evidence that genomic gain of 3q26 is involved in the progression from high-grade dysplasia (21,22). HPV-DNA testing seems to be useful in the triage of difficult cases but, as a screening device, it lacks specificity (4,23). p16^{INK4a} immunocytochemistry could be used as a surrogate marker for HR-HPV infection (5-7,9). In our results, p16^{INK4a} immunopositive cells in LBC for CIN3 were higher in the HR-HPV-positive group than in the negative group. The infection of cervical cells by HPV manifests itself in changes in the function or expression of host genes, and the detection of these alterations can play a role in screening and diagnosis. Actually, mild cytological abnormalities are extremely common in young women, but most of these lesions regress spontaneously, even when caused by oncogenic types of HPV (24,25) which play a crucial role in the pathogenesis of cervical cancer. This has prompted efforts to discover other biomarkers and other screening techniques with the potential to supplement cytological screening. Accordingly, there is a need to explore the screening system detecting women with lesions that need medical intervention, such as CIN3, but p16^{INK4a} does not seem to be suitable for this because of its lower specificity. The development of assays with high specificity for detecting cancer precursors as well as high sensitivity would represent an attractive alternative as primary screening tests or as tests to complement cytology and HPV typing.

Redon et al (17) identified the PIK3CA gene as the most likely oncogene to be overexpressed as a consequence of 3q gain. PIK3CA, encoding the catalytic subunit p110α of PI3-kinase 1A, has recently been suggested to play a critical role as an oncogene in ovarian and cervical cancer (15,16). Information about abnormal PIK3CA gene status in squamous cell carcinoma (SCC) has predominantly been obtained from experiments on cell lines. Employing ovarian carcinoma cell lines and uterine cervix SCC cell lines, they both pointed out that increased PI3K activity is solely caused by gene-dependent overexpression of the catalytic subunit p110α. Woenckhaus et al (26) showed that not only invasive SCCs in the head and neck but also high-grade dysplasias showed PIK3CA copy number gain, providing strong evidence that genomic gain of 3q26 is involved in the progression from high-grade dysplasia (21,22). HPV-DNA testing seems to be useful in the triage of difficult cases but, as a screening device, it lacks specificity (4,23). p16^{INK4a} immunocytochemistry could be used as a surrogate marker for HR-HPV infection (5-7,9). In our results, p16^{INK4a} immunopositive cells in LBC for CIN3 were higher in the HR-HPV-positive group than in the negative group. The infection of cervical cells by HPV manifests itself in changes in the function or expression of host genes, and the detection of these alterations can play a role in screening and diagnosis. Actually, mild cytological abnormalities are extremely common in young women, but most of these lesions regress spontaneously, even when caused by oncogenic types of HPV (24,25) which play a crucial role in the pathogenesis of cervical cancer. This has prompted efforts to discover other biomarkers and other screening techniques with the potential to supplement cytological screening. Accordingly, there is a need to explore the screening system detecting women with lesions that need medical intervention, such as CIN3, but p16^{INK4a} does not seem to be suitable for this because of its lower specificity. The development of assays with high specificity for detecting cancer precursors as well as high sensitivity would represent an attractive alternative as primary screening tests or as tests to complement cytology and HPV typing.

Redon et al (17) identified the PIK3CA gene as the most likely oncogene to be overexpressed as a consequence of 3q gain. PIK3CA, encoding the catalytic subunit p110α of PI3-kinase 1A, has recently been suggested to play a critical role as an oncogene in ovarian and cervical cancer (15,16). Information about abnormal PIK3CA gene status in squamous cell carcinoma (SCC) has predominantly been obtained from experiments on cell lines. Employing ovarian carcinoma cell lines and uterine cervix SCC cell lines, they both pointed out that increased PI3K activity is solely caused by gene-dependent overexpression of the catalytic subunit p110α. Woenckhaus et al (26) showed that not only invasive SCCs in the head and neck but also high-grade dysplasias showed PIK3CA copy number gain, providing strong evidence that genomic gain of 3q26 is involved in the progression from high-grade dysplasia (21,22). HPV-DNA testing seems to be useful in the triage of difficult cases but, as a screening device, it lacks specificity (4,23). p16^{INK4a} immunocytochemistry could be used as a surrogate marker for HR-HPV infection (5-7,9). In our results, p16^{INK4a} immunopositive cells in LBC for CIN3 were higher in the HR-HPV-positive group than in the negative group. The infection of cervical cells by HPV manifests itself in changes in the function or expression of host genes, and the detection of these alterations can play a role in screening and diagnosis. Actually, mild cytological abnormalities are extremely common in young women, but most of these lesions regress spontaneously, even when caused by oncogenic types of HPV (24,25) which play a crucial role in the pathogenesis of cervical cancer. This has prompted efforts to discover other biomarkers and other screening techniques with the potential to supplement cytological screening. Accordingly, there is a need to explore the screening system detecting women with lesions that need medical intervention, such as CIN3, but p16^{INK4a} does not seem to be suitable for this because of its lower specificity. The development of assays with high specificity for detecting cancer precursors as well as high sensitivity would represent an attractive alternative as primary screening tests or as tests to complement cytology and HPV typing.

Redon et al (17) identified the PIK3CA gene as the most likely oncogene to be overexpressed as a consequence of 3q gain. PIK3CA, encoding the catalytic subunit p110α of PI3-kinase 1A, has recently been suggested to play a critical role as an oncogene in ovarian and cervical cancer (15,16). Information about abnormal PIK3CA gene status in squamous cell carcinoma (SCC) has predominantly been obtained from experiments on cell lines. Employing ovarian carcinoma cell lines and uterine cervix SCC cell lines, they both pointed out that increased PI3K activity is solely caused by gene-dependent overexpression of the catalytic subunit p110α. Woenckhaus et al (26) showed that not only invasive SCCs in the head and neck but also high-grade dysplasias showed PIK3CA copy number gain, providing strong evidence that genomic gain of 3q26 is involved in the progression from high-grade dysplasia (21,22). HPV-DNA testing seems to be useful in the triage of difficult cases but, as a screening device, it lacks specificity (4,23). p16^{INK4a} immunocytochemistry could be used as a surrogate marker for HR-HPV infection (5-7,9). In our results, p16^{INK4a} immunopositive cells in LBC for CIN3 were higher in the HR-HPV-positive group than in the negative group. The infection of cervical cells by HPV manifests itself in changes in the function or expression of host genes, and the detection of these alterations can play a role in screening and diagnosis. Actually, mild cytological abnormalities are extremely common in young women, but most of these lesions regress spontaneously, even when caused by oncogenic types of HPV (24,25) which play a crucial role in the pathogenesis of cervical cancer. This has prompted efforts to discover other biomarkers and other screening techniques with the potential to supplement cytological screening. Accordingly, there is a need to explore the screening system detecting women with lesions that need medical intervention, such as CIN3, but p16^{INK4a} does not seem to be suitable for this because of its lower specificity. The development of assays with high specificity for detecting cancer precursors as well as high sensitivity would represent an attractive alternative as primary screening tests or as tests to complement cytology and HPV typing.
dysplasia to invasive SCC. In the study of Heselmeyer et al (27), which performed CGH and DNA cytometry on dysplasia and carcinomas of the uterine cervix, chromosome 3q gain was suggested to represent a key lesion in the transition from pre-invasive neoplasia to invasive SCC of the uterine cervix. The increase of PIK3CA DNA in dysplasia may provide an initial growth advantage to the cell. However, to date, there is no information about the genetic and post-transcriptional status of PIK3CA within premalignant lesions of uterine cervical cancer.

To our knowledge, the present study is the first report to examine PIK3CA status in LBC samples. PIK3CA was significantly expressed in the protein level by immunocytochemistry in CIN3 lesions. The possibility of using PIK3CA to identify cytological abnormalities and function as a surrogate marker for discriminating precancerous lesions was investigated. By applying ROC, we established function as a surrogate marker for discriminating precancerous lesions. In our study, p16 INK4a and Ki-67 epithelial lesions do not need to be followed because of their lower malignancy potential. In our study, p16 INK4a and Ki-67 also showed high cell counts of immunopositivity in CIN3 lesions but not as high as PIK3CA. From the results of HPV typing, Ki-67 and PIK3CA immunopositive cell counts were not related to HR-HPV infection, which suggests the utility of adjunct tools to HPV-DNA testing for Ki-67 immunocytochemistry in reflecting their potential of cell growth and PIK3CA as a carcinogenesis-related marker in early precancerous lesions. In conclusion, the diagnostic accuracy of PIK3CA to distinguish CIN3 was shown to be the best method among three markers to use as an adjunct method to LBC and HPV testing. Although these results need to be studied in a prospective situation, tests can be performed on the same sample concurrently and might be able to automate the staining procedure including counting of immunopositive cells.

References