# **Research Paper**



# Investigating Ki67 and p16 Positivity Pattern in Human Papillomavirus 16 and 18 Versus Other High-risk Human Papillomaviruses

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# ABSTRACT

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**Objectives:** The expression of Ki67 and p16 has been used for the diagnosis of human papillomavirus (HPV) 16 and 18; however, limited research has been conducted on the comparison with other high-risk HPVs (HR-HPVs) in this respect. Accordingly, this study compares the positivity pattern of Ki67 and p16 in HPV16/18 with other HR-HPVs.

**Methods:** This descriptive study included women with positive screening test (Pap smear test) and positive HR-HPV referred to Shahid Mohammadi Hospital, Bandar Abbas, Iran, from 2018-2019 for colposcopy and cervical biopsy. Biopsy specimens were stained for Ki67 and p16. Data on age, education, and menopause status were also recorded.

**Results:** Of the 80 women included in this study with a mean age of  $33.16\pm8.55$  years, 48.8% had positive HPV16/18, and 51.2% were positive for other HR-HPVs. The positivity pattern of Ki67 and the type of HR-HPV (HPV16/18 vs other HR-HPVs) were significantly correlated (P=0.006). Full-thickness positive Ki67 was only observed in HPV16/18 positive specimens. Positive Ki67 in the upper and middle thirds were also significantly more frequent with HPV16/18. The correlation between positive Ki67 and HR-HPVs remained significant in women aged >30 years, with university education, and of childbearing age.

**Discussion:** The positivity pattern of Ki67 is significantly correlated with HPV16/18, and age, education, and menopause status are the influential factors. Meanwhile, the strongly positive p16 pattern was more frequent in HPV16/18 compared to other HR-HPVs; however, the relationship between the positivity pattern of p16 and the type of HR-HPV was not significant.

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# Introduction

n its recent consensus statements, the World Health Organization (WHO) has emphasized the evaluation of cervical cancer. To date, no single effective method has been recognized for the prediction of cervical intraepithelial neoplasia (CIN) and disease progression in cervical cancer [1, 2]. The current screening program includes periodical Pap smear (cervical cytology) and high-risk human papillomavirus (HR-HPV) testing based on the WHO and the American Society for Colposcopy and Cervical Pathology (ASCCP) protocols [3, 4]. Evidence shows that HR-HPV testing has been more effective than cytology for the reduction of CIN 3 and cervical cancer [5].

P16 is a down-regulator of proliferation in normal cells by reducing the activity of cyclin-dependent kinase 4 (CDK4) and cyclin-dependent kinase 6 when the retinoblastoma protein is deactivated. HPV E7 can accelerate the progression of cancer cells by deactivating retinoblastoma protein leading to the overexpression of p16 in the injured cells. On the other hand, Ki67 is a nuclear protein and an indicator of cell proliferation. Accordingly, in normal cells, p16 and Ki67 are not concurrently expressed. The co-expression of p16 and Ki67 shows the reduction of cell cycle due to HPV oncoproteins which mainly occurs during viral persistency and indicates the potential correlation of p16 and Ki67 co-expression (p16/Ki67) with HPV infection [6].

Multiple studies have used the expression of p16 and Ki67 for the diagnosis of HPV16 and HPV18. For instance, Nam et al. evaluated the relationship between p16 and Ki67 expression with CIN 3 and HR-HPV and found a positive correlation between the CIN grade with p16 and Ki67 expression [7]. In another study, Keating et al. reported a 91% positive predictive value with p16 expression and 82% with Ki67 for the diagnosis of HPV [8]. Moreover, Ordi et al. have recommended the assessment of p16 expression along with histologic examination of the cervical biopsy specimens from HR-HPV-positive women [9].

Although many studies have evaluated the expression of p16 and Ki67 for the diagnosis of HPV16 and 18, limited research has been done to compare the expression of these two markers between HPV16/18 and other HR-HPVs. Accordingly, this study compares the positivity pattern of Ki67 and p16 in HPV16/18 with other HR-HPVs.

# **Materials and Methods**

### Study participants

This descriptive study evaluated all women with positive screening test (Pap smear) and positive HR-HPV who had been referred to Shahid Mohammadi Hospital, Bandar Abbas, Iran, in 2018-2019 for colposcopy and cervical biopsy. The participants were selected through census sampling and all eligible patients were included in the study. Inappropriate cervical specimens for immunohistochemistry were excluded from the study.

### Study design

The age, education, and menopause status of the patients were recorded. Cervical biopsy specimens underwent HPV genotyping using a polymerase chain reaction. The specimens were marked as HPV16/18 or other HR-HPVs (31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 68, 69). Ki67 and p16 staining were performed using Sanatashkhis products (manufactured in Iran). The staining steps were as follows. At first, 100 µL peroxidase was added to each slide. Then, the specimens were incubated at room temperature and darkness for 10 min. Afterward, each slide was rinsed with distilled water three times for 5 min. In the next step, a 100  $\mu L$  primary antibody amplifier was added to the specimens and remained for 15 min. Subsequently, each slide was rinsed three times for 5 min. Meanwhile, 100 µL master polymer plus horseradish peroxidase was added to each slide with specimens incubated at room temperature for 30 min followed by rinsing each slide 3 times for 5 min. The chromogen solution was added to each slide and was left for 5 min at room temperature followed by rinsing the slides 3 times for 5 min. Then, the 3,3'-diaminobenzidine enhancer was added to the slides and remained for 1 to 2 min at room temperature and then rinsed. For counterstaining, the specimens were covered with hematoxylin for 1 min and then rinsed with distilled water.

Ki67 positivity patterns included the lower third, the middle third, the upper third, and full-thickness, while p16 staining patterns were either weakly or strongly positive. Also, the CIN grades were determined.

### Data analysis

The SPSS software, version 25 (Armonk, NY: IBM Corp., USA) was used for data analysis. Mean $\pm$ SD, frequencies, and percentages were used to describe the variables. The Fisher exact tests were used to compare frequencies between groups. Meanwhile, P $\leq$ 0.05 were considered statistically significant.

# Results

Of the 80 women evaluated in this study (mean age of  $33.16\pm8.55$  years), 39(48.8%) had positive HPV16/18, and 41(51.2%) were positive for other HR-HPVs. Ki67 was positive in the lower third of the specimen in 60(75%), up to the middle third in 13(16.3%), up to the upper third in 5(6.3%), and full-thickness in 2(2.5%). Three specimens (3.8%) were negative for p16, while 52(65%) were weakly and 25(31.3%) strongly positive for p16. The general characteristics of the study population are demonstrated in Table 1.

Positive Ki67 up to the middle and upper third, and full-thickness positivity were significantly higher in positive HPV16/18 specimens compared to other HR-

Table 1. Study population characteristics

HPVs (P=0.006). Nevertheless, no significant correlation was detected between the p16 positivity pattern and HR-HPV type (P=0.163) (Table 2).

When the study participants were divided into different groups based on age, education, menopause status, and CIN grade, a significant relationship was found between Ki67 positivity pattern and HR-HPV type in women aged >30 years, those with university education, and women of childbearing age (Table 3). However, no significant correlation was observed between the positivity pattern of p16 and the type of HR-HPV in women of different age groups, education, or menopause status (Table 4).

	No. (%)	
Age (y)	≤30	37(46.3)
	>30	43(53.8)
Education	High school diploma or less	46(57.5)
Education	University	34(42.5)
Menonause status	Childbearing age	74(92.5)
Menopause status	Menopause	6(7.5)
	HPV16/18	39(48.8)
TIN-THE V	Other HR-HPVs	41(51.2)
	Negative	6(7.5)
	CIN 1	53(66.3)
CIN	CIN 2	14(17.5)
	CIN 3	5(6.3)
	Carcinoma in situ	2(2.5)
	Lower 3 <sup>rd</sup>	60(75.0)
Ki67 Staining	Middle 3 <sup>rd</sup>	13(16.3)
Kitor Stalling	Upper 3 <sup>rd</sup>	5(6.3)
	Full-thickness	2(2.5)
	Negative	3(3.8)
p16 staining	Weakly positive	52(65.0)
	Strongly positive	25(31.3)

Abbreviations: HPV: Human papillomavirus; HR-HPV: High-risk HPV; CIN: Cervical intraepithelial neoplasia.

Discussion

This study found a significant correlation between the positivity pattern of Ki67 with HR-HPV type (HPV16/18 and other HR-HPVs). Full-thickness positive Ki67 was only observed in HPV16/18 positive specimens. Also, positive Ki67 in the upper and middle third of the specimens were significantly higher with HPV16/18 compared to other HR-HPVs. However, although the strongly positive pattern of p16 was more frequent with HPV16/18 compared to other HR-HPVs, the difference was not statistically significant.

Cervical cancer is the fourth most common malignant tumor in women worldwide and HPV infection is considered its major etiology [10]. Specific HPV types, especially HPV16 and HPV18 can lead to cervical dysplasia which is a reversible precancerous lesion. Consistent cervical infection with HPV can cause irreversible alterations that can result in carcinoma in situ and consequently invasive cervical cancer [11]. Early diagnosis using different screening methods is the primary measure for the prevention and treatment of cervical cancer. Currently, three methods, including cervical cytology, HPV genotyping, and cytology together with HPV genotyping are used for cervical cancer screening [12]. Due to the low sensitivity of cytology, many HPV-positive women require multiple cytology follow-ups. Thus, more effective markers are required to triage HPV-positive women with normal cytology or without HPV16/18, or to differentiate women with high-grade CIN from low-grade squamous intraepithelial lesions and atypical squamous cells of undetermined significance. Evidence shows that

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cytology with p16/Ki67 dual staining can be a substitute biomarker with high sensitivity and specificity for high-grade CIN [13-17].

In line with our findings, Lewitowicz et al. also showed a significant correlation between HPV16 and positive p16/Ki67 [18]. Similarly, Liao et al. demonstrated that p16/Ki67 expression in HPV16/18 positive and other 12 HR-HPV positive specimens was significantly higher compared to specimens with negative HPV [19]. The results of this study were also consistent with ours; however, we only evaluated HR-HPV-positive specimens and showed significantly higher expression of Ki67 with HPV16/18 compared to other HR-HPVs. On the other hand, Jiang et al. reported an increase in p16/Ki67 positivity with increased HPV risk (HR-HPVs) from 65% in women with positive HPV51/39/68/35 to 88% in women with positive HPV16/18 [20]. Moreover, Yu et al. observed that the odds of positive p16/Ki67 in women with persistent HPV16/18 infection was approximately 4-fold compared to women with other HR-HPVs [6]. In addition, Dona et al. reported that the correlation between positive p16/Ki67 with HPV16/18 was twice stronger than its correlation with other HR-HPVs [21]. In previous studies, the concurrent expression of p16 and Ki67 has been evaluated, while this study assessed each marker separately and showed no significant correlation between positive p16 and HR-HPV type. Furthermore, in the previous studies, specimens were assessed regarding the positivity of these markers, i.e. each specimen was either positive or negative, while we also evaluated the pattern of positivity for each marker, which can be regarded as a major strength for the current study. By

Variables		No. (%)		-
	Distribution	HPV16/18 (n=39)	Other HR-HPVs (n=41)	Ρ
Ki67 staining	Lower 3 <sup>rd</sup>	23(59.0)	37(90.2)	
	Middle 3 <sup>rd</sup>	10(25.6)	3(7.3)	0.000
	Upper 3 <sup>rd</sup>	4(10.3)	1(2.4)	0.006
	Full-thickness	2(5.1)	0(0.0)	
P16 staining	Negative	1(2.6)	2(4.9)	
	Weakly positive	22(56.4)	30(73.2)	0.163
	Strongly positive	16(41.0)	9(22.0)	

HPV: Human papillomavirus; HR-HPV: High-risk HPV.

\*Fisher exact test.

Table 2. Ki67 & p16 patterns: HPV16/18 vs other HR-HPVs

Ki67 Staining	Distribution	No. (%)		- D*
		HPV16/18	Other HR-HPVs	٢
Age(y) ≤30	Lower 3 <sup>rd</sup>	11(55.0)	14(82.4)	
	Middle 3 <sup>rd</sup>	8(40.0)	3(17.6)	0.157
	Upper 3 <sup>rd</sup>	1(5.0)	0(0.0)	
	Lower 3 <sup>rd</sup>	12(63.2)	23(95.8)	
Aco(u) > 20	Middle 3 <sup>rd</sup>	2(10.5)	0(0.0)	0.021
Age(y) >50	Upper 3 <sup>rd</sup>	3(15.8)	1(4.2)	0.021
	Full-thickness	2(10.5)	0(0.0)	
	Lower 3 <sup>rd</sup>	13(68.4)	24(88.9)	
High school	Middle 3 <sup>rd</sup>	3(15.8)	3(11.1)	0.110
diploma or less	Upper 3 <sup>rd</sup>	1(5.3)	0(0.0)	0.118
	Full-thickness	2(10.5)	0(0.0)	
	Lower 3 <sup>rd</sup>	10(50.0)	13(92.9)	
University	Middle 3 <sup>rd</sup>	7(35.0)	0(0.0)	0.015
	Upper 3 <sup>rd</sup>	3(15.0)	1(7.1)	
Childbearing age	Lower 3 <sup>rd</sup>	21(60.0)	35(89.7)	
	Middle 3 <sup>rd</sup>	10(28.6)	3(7.7)	0.010
	Upper 3 <sup>rd</sup>	4(11.4)	1(2.6)	
Menopause	Lower 3 <sup>rd</sup>	2(50.0)	2(100.0)	0.467
	Upper 3 <sup>rd</sup>	2(50.0)	0(0.0)	0.467
CIN 1	Lower 3 <sup>rd</sup>	20(100.0)	33(100.0)	-
CIN 2+	Lower 3 <sup>rd</sup>	0(0.0)	2(40.0)	
	Middle 3 <sup>rd</sup>	10(62.5)	3(60.0)	0.070
	Upper 3 <sup>rd</sup>	4(25.0)	0(0.0)	0.076
	Full-thickness	2(12.5)	0(0.0)	

Table 3. Ki67 pattern: HPV16/18 vs other HR-HPVs by age, education, menopause, CIN grade

Abbreviations: HPV: Human papillomavirus; HR-HPV: High-risk HPV; CIN: Cervical intraepithelial neoplasia.

\*Fisher exact test.

solely taking positive and negative results into account, most probably we could have achieved similar results concerning these markers.

Another strength of our study was that we evaluated the correlation between the positivity pattern of Ki67 and p16 with HR-HPV types in women of different age groups, education levels, and menopause status and found a significant correlation between Ki67 positivity pattern with HR-HPV types in women aged >30 years, individuals with university education, and women of childbearing age. Previous studies lacked such analyses.

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Ki67 Staining	Expression Levels	No. (%)		D*
		HPV16/18	Other HR-HPVs	- r
Age (y) ≤30	Negative	0(0.0)	2(11.8)	
	Weakly positive	11(55.0)	11(64.7)	0.150
	Strongly positive	9(45.0)	4(23.5)	
Age (y) >30	Negative	1(5.3)	0(0.0)	
	Weakly positive	11(57.9)	19(79.2)	0.228
	Strongly positive	7(36.8)	5(20.8)	
High school	Weakly positive	13(68.4)	21(77.8)	0.542
diploma or less	Strongly positive	6(31.6)	6(22.2)	0.513
	Negative	1(5.0)	2(14.3)	
University	Weakly positive	9(45.0)	9(64.3)	0.223
	Strongly positive	10(50.0)	3(21.4)	
Childbearing age	Negative	1(2.9)	2(5.1)	
	Weakly positive	20(57.1)	28(71.8)	0.315
	Strongly positive	14(40.0)	9(23.1)	
Menopause	Weakly positive	2(50.0)	2(100.0)	0.467
	Strongly positive	2(50.0)	0(0.0)	0.467
CIN 1	Negative	0(0.0)	2(6.1)	
	Weakly positive	18(90.0)	28(84.8)	0.827
	Strongly positive	2(10.0)	3(9.1)	
CIN 2+	Weakly positive	2(12.5)	0(0.0)	1 000
	Strongly positive	14(87.5)	5(100.0)	1.000

Table 4. P16 pattern: HPV16/18 vs other HR-HPVs by age, education, menopause, CIN grade

Abbreviations: HPV: Human papillomavirus; HR-HPV: High-risk HPV; CIN: Cervical intraepithelial neoplasia.

\*Fisher exact test.

Since the CIN grade of each specimen was also available, this study assessed the correlation between Ki67 and p16 positivity patterns with HR-HPV types. According to the findings, regardless of the HR-HPV type, all CIN 1 specimens were positive for Ki67 in the lower third. Moreover, there was no significant correlation between the positivity pattern of Ki67 and p16 with HR-HPV types in CIN 2+ specimens; however, 4 and 2 HPV16/18 positive specimens with CIN 2+ were positive for Ki67 in the upper third and full-thickness, respectively, while Ki67 was positive in none of the CIN 2+ specimens that were also positive for other HR-HPVs. On the other hand, in CIN 1 and CIN 2+ specimens, the correlation between the p16 positivity pattern and HR-HPV types was statistically significant. Nonetheless, 18 and 2 HPV16/18 positive specimens with CIN 1 had weakly and strongly positive patterns of p16, which was higher compared to other HR-HPVs with CIN 1. Moreover, 2 HPV16/18 positive specimens with CIN 2+ had a weakly positive pattern of p16, while none of the specimens with other HR-HPVs and CIN 2+ had such a pattern. In this regard, Wentzensen et al. showed that for triaging HPV-positive women, p16/Ki67 staining can better classify the risk of CIN 3+ compared to Pap smear cytology. They reported that women with positive p16/Ki67 staining had

a higher risk of CIN 3+ compared to women with positive Pap smear [22]. Besides, in the study by Liao et al., the sensitivity of p16/Ki67 for the diagnosis of CIN 2+ and CIN 3+ was 94.1% and 92.9%, respectively [19]. Based on the results of the study by Stanczuk et al., the absolute sensitivity and specificity of p16/Ki67 for the diagnosis of CIN 2+ in women with positive HR-HPV were 85% and 76.7%, respectively [23]. Therefore, further studies are required to evaluate the relationship between the positivity pattern of Ki67 and p16 with HR-HPV types and its role in determining the CIN grade. The difference between previous studies and this research in this regard may rely on different demographic characteristics of the study populations, the concurrent evaluation of the two markers in other studies, and different study designs.

## Conclusions

The present study showed a significant correlation between the positivity pattern of Ki67 with HPV16/18, with age, education, and menopause status as influential factors. Meanwhile, the strongly positive p16 pattern was more frequent in HPV16/18 compared to other HR-HPVs; however, the relationship between the positivity pattern of p16 and the type of HR-HPV was not significant.

### **Study limitations**

The primary limitation of the current study was its relatively small sample size which limits the generalizability of our findings. Future studies with a larger sample size are required to confirm our findings.

## **Ethical Considerations**

### Compliance with ethical guidelines

This study received ethical approval from the Ethics Committee of Hormozgan University of Medical Sciences (Code: IR.HUMS.REC.1398.421). It complies with the statements of the Declaration of Helsinki. Written informed consent was obtained from the participants.

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### **Authors' contributions**

Conceptualization, study validation and supervision: Ali AtashAbParvar; Data analysis, data interpretation, writing and review: Masoumeh Baniasadi; Final approval: All authors.

#### Conflict of interest

The authors declared no conflict of interest.

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