Abstract 1842

HPV POSITIVE CERVICAL SQUAMOUS CELL CARCINOMAS: ASSOCIATION WITH EXPRESSION OF P16, P53 AND KI67.

Type: Abstract Submission

AS03. Public Health, Epidemiology and Implementation Science / AS03c. Screening for HPV-related Topic: Disease: Implementation, Evaluation and Impact

Authors: Liba Sokolovska<sup>1</sup>, Karina Biserova<sup>1,2</sup>, Daira Krisane<sup>2,3</sup>, Alesja Dudorova<sup>1,2,3</sup>, Viola Daniela Kiselova<sup>1</sup>, Simons Svirskis<sup>1</sup>, Dmitrijs Perminovs<sup>4</sup>, Jurijs Nazarovs<sup>2,3,5</sup>, Maria Isaguliants<sup>1</sup>; <sup>1</sup>Riga Stradins University, Institute of Microbiology and Virology, Riga, Latvia, <sup>2</sup>Pauls Stradins Clinical University Hospital, Pathology Institute, Riga, Latvia, <sup>3</sup>Riga East Clinical University Hospital, Pathology Centre, Riga, Latvia, <sup>4</sup>E. Gulbis Laboratory Ltd, Riga, Latvia, <sup>5</sup>Riga Stradins University, Department of Pathology, Riga, Latvia

### Introduction

Cervical cancer is the fourth most common malignancy in females. The majority of cervical carcinomas (CC) are associated with high-risk HPVs (hrHPV). Here, we investigated the association between hrHPV genotypes, virus load, p16, p53, and Ki-67 expression in hrHPV-positive CCs.

# **Methods**

FFPE cervical tissues of 76 patients (median age 60 years) with primary cervical squamous cell carcinomas were assessed. Tumor grades were evaluated histologically, and expression of p16, p53, and Ki-67 immunohistochemically using Flex kits, Autostainer Link-instrument (Dako), and Eclipse 55i microscope (Nikon). DNA isolated from FFPE samples (Qiagen) was used for hrHPV DNA detection and semiguantification (Anyplex Seegene).

## Results

All 76 cancer cases were positive for at least one hrHPV, mainly HPV16, followed by HPV33. Other genotypes were less prevalent (Table 1)

### table 1.png

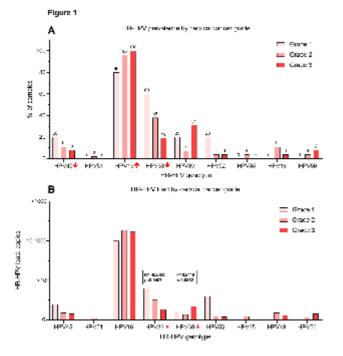
	Fotal n=76	Grade 1 n=5	Garde 2 n=45	Grade 3 n=26
HPV66	0	41	19 19	0
HPV45	<b>8</b> 10.3%	1 20%	5 11.1%	<b>2</b> 7.7%
HPV58	0	41	19	0
HPV51	1 1.3	0	1 2.2%	0
HPV69	0	41	19	D
HPV16	73 96.1%	4 80%	43 95.6%	26 1012%
HPV35	<b>25</b> 32.9%	3 60%	17 37.8%	5 19.2%
HPV39	12 15.8%	1 20%	3 6 7%	<b>8</b> 30.8%
HPV52	4 5 7%	1 20%	2 4 4%	1 7 8%
HPV35	2 2.6%	- e	2 4 4%	0
HPV18	6 7.9%	સ	5 11,1%	3.8%
HPV56	4 5.3%	0	2 4,4%	2 7.7%
HPV68	0	ei -	6	D

Table 1. HR-HPV genotype prevalence in cervical cancer tissues by cancer y

### 🕀 enlarge

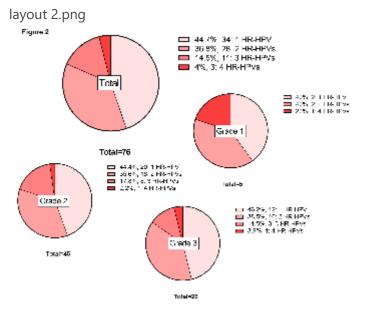
Prevalence of HPV16 increased, while HPV33 and HPV45 decreased with tumor grade (Figure 1A(F1A)). HPV33 load correlated negatively, while HPV39 - positively with tumor grade (F1B).

### layout 1.png



#### <u>⊕ enlarge</u>

Many CC samples were positive for multiple hrHPVs (F2), without correlating with tumor grade.



#### **<u>enlarge</u>**

74 CCs (97.4%) were p16(+). Median % of ki67-expressing cells was 43.5% (IQR: 25.3 – 70.0%). ki67-expressing cells positively correlated with HPV39 (R=0,2406, p=0,0362) and negatively with HPV18 load (R=-0,3079, p=0,0067), and tended to increase with increasing cancer grade (median: 52 vs 56 vs 65). Aberrant p53 expression was detected in six CC cases (7.9%), most often in Grade 3 (5/6) in patients older than 60 (4/6) and positively correlated with cancer grade (R=0,2936, p=0,0100), and HPV18 virus load (R=0,2783, p=0,0149).

### Conclusions

p16 positivity acted as a surrogate marker of hrHPV(+) cancer, with 97% sensitivity. Percent of Ki67-positive cells was not informative in distinguishing tumor grades. Least informative was p53-staining. Of note, HPV33 and 39 correlated with tumor grade. Thus, immunostaining for p16 is sufficient to confirm an hrHPV(+) CC.

Acknowledgments. Latvian Science Council grant LZP-2021/1-0484.

Print