Abstract. Greece is one of the first countries of the European Union to introduce a human papilloma virus (HPV) vaccine in its national vaccination programme. Geographical variations in the prevalence of the different HPV types have been demonstrated. The aim of this preliminary case control study was to investigate HPV infection in women with low- and high-grade squamous intraepithelial lesions (SILs) from two different geographical areas of Greece; Central Greece and Crete. Seventy-five cervical specimens were collected from women with SILs from Crete and Central Greece. HPV detection and typing were performed by using polymerase chain reaction (PCR) techniques. HPV-18 was detected more frequently in Crete than in the Central Greece samples (29.7 vs 13.1%). HPV-16 was predominant in Central Greece (34.2%), while in Crete it was detected in 23% of the studied women. Non-16/-18 HPV types were detected in 45.9% of the women from Crete and 52.6% of the women from Central Greece. No relationship was observed between the geographical distribution of HPV and the presence of K-ras or B-raf point mutations in either group. Our pre-vaccination data indicate a high prevalence of HPV-18 in Crete. A trend for difference was observed in the rates of non-HPV-16/-18 women between the two areas. A large epidemiological study is required to investigate the prevalence of the different HPV types to further investigate the effectiveness of HPV vaccination in the Greek population.

Introduction

Human papilloma viruses (HPV) are small double-stranded DNA viruses that comprise a remarkably heterogeneous family of more than 130 types (1,2). Different HPV types can cause a wide range of infections in adults and children, including common warts, genital warts, recurrent respiratory papillomatosis, low-grade and high-grade squamous intraepithelial lesions and cervical cancer. 'High-risk' HPV types have been implicated in the development of intraepithelial lesions (SILs) and its progression to cervical cancer (2,3). Fifteen HPV types have been classified as ‘high risk’, HPV-16,-18,-31,-33,-35,-39,-45,-51,-52,-56,-58,-59,-68,-73 and -82 (4,5).

It has been demonstrated that the distribution of different HPV types in women with cervical neoplasia varies according to region (6-8). These geographic variations can play an important role in the effectiveness of an HPV vaccination and must be taken into consideration when designing policies against HPV. Current and proposed HPV vaccine candidates are HPV type-specific conferring only type-specific immunity against ‘high-risk’ types HPV-16 and -18 (9-11). HPV-16 and -18 are the most common HPV types and are responsible for ~70% of all cervical cancer cases (5,6). Vaccination against HPV-16 and -18 potentially prevents more than two thirds of cervical cancer cases worldwide. However, the impact of an HPV vaccination in different geographical regions will be related to the prevalence of HPV-16 and -18 in the different populations (6).

Interestingly, among women with low- and high-grade SILs, the prevalence of HPV-16/18 is lower compared to cervical cancer; however, it follows similar patterns in different regions (6,7). The aim of this preliminary case control study was to investigate HPV infection in women with low-grade and high-grade SILs from two different geographical areas of Greece, the island of Crete and Central Greece. We also examined the presence of K-ras codon 12 and B-raf exon 15 point mutations and their possible relationship to the geographical distribution of HPV. Greece is one of the first European Union countries to introduce an HPV vaccine in its national vaccine schedule after the approval of the quadrivalent vaccine by the European Union in 2006. Pre-vaccination data concerning the prevalence of HPV genital infection in women will enable the investigation of the effectiveness of an HPV vaccination and to prevent HPV-induced carcinogenesis.

Correspondence to: Professor D.A. Spandidos, Laboratory of Virology, Faculty of Medicine, University of Crete, Heraklion, Crete, Greece
E-mail: spandidos@spandidos.gr

Key words: human papilloma virus, vaccination programme, geographic variations, Greece
Materials and methods

Our study included cervical specimens that were obtained from women with low-grade or high-grade cervical intraepithelial lesions (SILs), who were treated at the University Hospital of Heraklion on the island of Crete and the University Hospital of Larissa in Central Greece during the period 2003 to 2005. The histological analysis of the material was performed at the Departments of Pathology of the University Hospitals of Heraklion and Larissa, Greece. Clinicopathological data (age, origin, residence and medical history) were available for all patients included in the study. Tissue specimens obtained during colposcopy were divided into two parts. One portion was fixed in buffered formalin (pH 7.4) for routine histopathological assessment, while the other portion was paraffin-embedded for DNA extraction. DNA was extracted as previously described (12) and stored at -20°C.

For the detection of the HPV the general primers GP5+ and GP6+ were used. The extracted DNA (1 µl) of each sample was amplified in a total volume of 30 µl containing 5 µM of 10X PCR reaction buffer (200 mM Tris-HCl, pH 8.4 and 500 mM KCl), 1.5 mM MgCl2, 200 µM of each dNTP, 0.5 µM of each primer and 0.6 U Platinum Taq DNA polymerase (Invitrogen Ltd., UK). The amplification conditions, the general primers for HPV detection, the specific primers used for HPV detection and typing have been previously described (13,14). PCR products for HPV were analysed on 2% agarose gel and photographed on a UV light transilluminator. PCR assay was carried out in a PTC-200 programmable thermal controller (MJ Research Inc., USA). All PCR reactions included appropriate negative controls. DNA extracted from HeLa cells and plasmid DNA of HPV-16 and HPV-18 were used as positive controls. The detection of exon 15 B-raf and codon 12 K-ras point mutations was performed using polymerase chain reaction (PCR) and restriction fragment length polymorphism (RFLP) analysis as previously described (15).

Statistical analyses were performed using SPSS software (SPSS, version 11.5). The Pearson χ² test was performed in order to compare the two different group populations according to age, origin, residence, histology, point K-ras and B-raf mutations and HPV status. The limit of statistical significance was set at p<0.05.

Results

Our study included 75 Greek women with squamous epithelial neoplasia (SIL). The clinicopathological characteristics of our sample are presented in Table I. In the population group from Crete, 23 women had low-grade SILs and 14 had high-grade SILs. Of the women from Central Greece, 23 had low-grade SILs and 15 had high-grade SILs. The mean age of the population groups from Crete and Central Greece was 42 (range 28-58) years and 44 (range 31-61) years, respectively. No differences were observed between samples from Crete and Central Greece according to age, ethnicity or residence (Pearson χ² test, p>0.05). Codon 12 point mutations in K-ras were detected in 4 out of 37 women from Crete and 5 out of 38 from Central Greece. All K-ras mutations were detected in women with high-grade SILs. No B-raf exon 15 point mutation was detected in either population. There was no statistically significant difference in the presence of K-ras or B-raf mutation between samples from Crete and Central Greece (Table I).

The results of the HPV detection and typing of our study samples are presented in Table II. All women included in our study were HPV-positive (75/75, 100%). HPV-18 was detected in 11 women (29.7%) from Crete and in 5 women (13.1%) from Central Greece. HPV-16 was predominant in Central Greece (34.2%), while in Crete it was detected in 9 women (27%). Non-16/-18 HPV types were detected in 17 (45.9%) women from Crete and 20 (52.6%) women from Central Greece; however, this difference was not statistically significant (Pearson χ² test, p>0.05).

In the total sample, HPV-16 was detected more frequently than HPV-18 (30.7 vs 21.3%). HPV-16 was detected in 10 (21.7%) women with low-grade SILs and in 44.8% of women
with high-grade SILs. The respective rates for HPV-18 among patients with low- and high-grade SILs were 15.2 and 31.3%. Co-infection with HPV-16 and -18 was detected in one woman with high-grade SIL from Crete. In the total sample, the ratio HPV-16/-18 to non-HPV-16/-18 was 1.03 (38/37), the ratio among women with low-grade SILs was 0.55 (16/29), and the ratio among women with high-grade SILs was 2.63 (21/8). The respective ratios among women from Crete were 1.17 (20/17), 0.77 (10/13) with low-grade SILs, and 2.5 (10/4) with high-grade SILs, and among women from Central Greece 0.9 (18/20), 0.44 (7/16) with low-grade SILs, and 2.75 (11/4) with high-grade SILs.

Discussion

The human papilloma virus (HPV) has a global distribution (6). The prevalence of the different types of HPV worldwide has implications for the effectiveness of HPV vaccinations against HPV-induced carcinogenesis. This is the first report in the literature to document differences in the geographical distribution of HPVs in Greece. It has been proposed that a worldwide vaccine against HPV-16/-18 may prevent a larger proportion of cervical carcinoma cases in Europe, North America and Australia than in Africa, South America and Asia (6). Notably, in our sample from Crete, HPV-16 was detected in 21.7% in low-grade SILs and 35.7% in high-grade SILs, while among women from Central Greece the respective prevalence was 21.7% in low-grade SILs and 53.3% in high-grade SILs. It seems that HPV-16, which dominates in mainland Europe and worldwide, is detected less frequently in women from Crete.

Among low-grade SILs, the prevalence of HPV-18 has previously been estimated as 5.3% in Africa, 7.1% in Asia, 9.2% in North America, 3.6% in South/Central America and 5.2% in Europe (7). Among high-grade SILs, the respective prevalence ranges from 6.5% in Europe to 10% in North America, while the pattern is consistent among SCC (6,8). In our sample, HPV-18 was detected in 21.7% of low-grade SILs and 42.9% of high-grade SILs. The frequency of HPV-18 in the sample from Crete was higher compared to the frequencies that have been reported by others in Europe. Notably, the frequency of HPV-18 found in women with SIL from Crete is one of the highest that has been reported in the literature (6,8). Further research is required to study the possible causes of HPV-18 dominance in Crete.

The distribution of HPV types in different geographical regions may be related to host immunologic and genetic factors (16). Crete, an island at the southeast edge of Europe, close to Africa, represents a remote population target with unique geographical characteristics. Knowledge on the distribution of different HPV types in a region is essential for designing and implementing an effective HPV vaccination programme. Currently, the only ‘high-risk’ HPV types that are covered by the different current virus-like particle vaccines are HPV-16 and -18 (9,10). The geographical distribution of

Table II. HPV detection and typing in women with squamous intraepithelial neoplasia in two geographical areas of Greece: Central Greece vs Crete.

<table>
<thead>
<tr>
<th>Sample group</th>
<th>HPV (+)</th>
<th>HPV-16 (+)</th>
<th>HPV-18 (+)</th>
<th>Non-HPV-16/-18 (+)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Crete</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LSIL</td>
<td>23</td>
<td>5</td>
<td>5</td>
<td>13</td>
</tr>
<tr>
<td>HSIL</td>
<td>14</td>
<td>5</td>
<td>6</td>
<td>4</td>
</tr>
<tr>
<td>Total</td>
<td>37 (100%)</td>
<td>10 (27.0%)</td>
<td>11 (29.7%)</td>
<td>17 (45.9%)</td>
</tr>
<tr>
<td>Central Greece</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LSIL</td>
<td>23</td>
<td>5</td>
<td>2</td>
<td>16</td>
</tr>
<tr>
<td>HSIL</td>
<td>15</td>
<td>8</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>Total</td>
<td>38 (100%)</td>
<td>13 (34.2%)</td>
<td>5 (13.1%)</td>
<td>20 (52.6%)</td>
</tr>
<tr>
<td>Total (%)</td>
<td>75 (100%)</td>
<td>23 (30.7%)</td>
<td>16 (21.3%)</td>
<td>37 (49.3%)</td>
</tr>
</tbody>
</table>

LSIL, low-grade squamous intraepithelial lesions; HSIL, high-grade squamous intraepithelial lesions.
Table III. HPV infection in Greece. Review of the literature.

<table>
<thead>
<tr>
<th>Author</th>
<th>Region</th>
<th>Methodology</th>
<th>n</th>
<th>HPV-16 (%)</th>
<th>HPV-18 (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Koffa, et al (24)</td>
<td>Attica</td>
<td>Multiplex PCR</td>
<td>31</td>
<td>11/31 (35.5%)</td>
<td>3/31 (9.7%)</td>
</tr>
<tr>
<td>Agorastos, et al (12)</td>
<td>Attica</td>
<td>Separate DNA blot</td>
<td>99</td>
<td>4/99 (4.0%)</td>
<td>8/99 (8.0%)</td>
</tr>
<tr>
<td>Noutsou, et al (14)</td>
<td>Attica</td>
<td>Multiplex PCR</td>
<td>35</td>
<td>3/35 (8.6%)</td>
<td>14/35 (40.0%)</td>
</tr>
<tr>
<td>Rousakis-Schulze, et al (28)</td>
<td>Attica</td>
<td>Multiplex PCR</td>
<td>88</td>
<td>4/88 (4.5%)</td>
<td>36/88 (40.9%)</td>
</tr>
<tr>
<td>Mammas, et al (29)</td>
<td>Attica</td>
<td>Multiplex PCR</td>
<td>102</td>
<td>11/102 (10.8%)</td>
<td>22/102 (21.6%)</td>
</tr>
<tr>
<td>Lyronis, et al (30)</td>
<td>Crete</td>
<td>Multiplex PCR</td>
<td>47</td>
<td>5/47 (10.6%)</td>
<td>12/47 (25.5%)</td>
</tr>
<tr>
<td>Prokopakis, et al (31)</td>
<td>Crete</td>
<td>Specific PCR</td>
<td>31</td>
<td>15/31 (48.4%)</td>
<td>12/31 (38.7%)</td>
</tr>
<tr>
<td>Efstathiou, et al (32)</td>
<td>Crete</td>
<td>Multiplex PCR</td>
<td>57</td>
<td>25/57 (45.6%)</td>
<td>20/57 (35.1%)</td>
</tr>
<tr>
<td>Lyronis, et al (33)</td>
<td>Crete</td>
<td>Multiplex PCR</td>
<td>102</td>
<td>6/102 (5.9%)</td>
<td>0/102 (0.0%)</td>
</tr>
</tbody>
</table>

SIL, squamous intraepithelial lesions; CC, cervical cancer; PCR, polymerase chain reaction; RFLP, restriction fragment length polymorphism.

**Different HPV types suggests that the effect of vaccination against HPV will vary depending on the region.**

No correlation was found in HPV distribution and K-Ras/ B-Raf mutational activation. The Ras-Raf signaling pathway is a key membrane-to-nucleus signaling system that regulates fundamental cellular functions in response to extracellular signals by activating a number of effector proteins (17,18). Mutational activation of the Ras-Raf signaling pathway at different levels has been implicated in the initiation and progression of several types of human cancer (19,20). K-ras mutations occur in the early stages of cervical neoplasia and represent the most frequent ras point mutations in women with SIL (21). B-raf mutations have recently been found in several types of cancer and have been proposed as a common mechanism of Ras-Raf signaling pathway activation which leads to cellular transformation (22). However, the absence of B-Raf mutations in our population indicates that B-Raf mutational activation does not seem to be involved in the pathogenesis of SIL.

Although HPV-16 and -18 are the dominant HPV types detected in women with cervical cancer and its precursors worldwide, in several geographical areas, other HPV types have been detected more frequently than HPV-16 and -18 (6,7). Among low-grade SILs, other 'high-risk' HPV types, such as -31, -51, -52, -56 and -58 were detected more frequently than HPV-18 (7). In Asia, a high prevalence of HPV-58 and in Europe HPV-31 has been demonstrated in low-grade and high-grade SILs (6-8). Among women with SILs, the frequency of non-16/-18 HPV types ranges from 34 to 68%. High prevalence of non-16/-18 HPV types are of great importance since these types are not covered by the existing vaccines. In our study, the ratio HPV-16/18 to non-HPV-16/-18 in Central Greece [0.9 (18/20)] was lower compared to the ratio of 1.17 (20/17) in Crete.

Both current and future HPV vaccine candidates are HPV type-specific conferring only type-specific immunity (9-11). Data on cross-protection of the current HPV L1 virus-like particle vaccines against other HPV types are still limited. It has been shown that prophylactic quadrivalent human papilloma-virus (types -6, -11, -16, and -18)-like particle vaccine results in a neutralizing antibody response which is HPV type specific (9). Recently, Harper et al reported that the protection conferred by the bivalent virus-like particle vaccine against HPV-16 and -18 was extended to 'high-risk' types HPV-45 and -31 (10). Taxonomically, the DNA genome of different types of HPV differs by at least 10% of the nucleotide sequence of the three open reading frames E6, E7 and L1 from that of any other known type (23). Long-term follow-up studies will clarify the role of HPV vaccination against non-16/-18 HPV types.

The differences in the geographical distribution of HPV types in Greece remain to be clarified. In the future, it is crucial to examine a larger number of clinical samples and to detect the prevalence of all fifteen 'high-risk' HPV types in the Greek population. These findings will provide pre-vaccination data of the population enabling the evaluation of the efficacy of the vaccination programme and to promote policies against HPV-induced carcinogenesis in Greece. Moreover, possible geographical variations in specific areas demonstrate the necessity for the development of polyvalent
heterogeneous vaccines in the future which will include types that are more prevalent in specific regions.

References