Enhancement of humoral and cell mediated immune response to HPV16 L1-derived peptides subsequent to vaccination with prophylactic bivalent HPV L1 virus-like particle vaccine in healthy females

MASATO YOKOMINE1, SATOKO MATSUEDA2,5, KOUICHIRO KAWANO1, TETSURO SASADA2,6, AKIMASA FUKUI1, TAKUTO YAMASHITA3, NOBUKAZU KOMATSU4, SHIGEKI SHICHIJO2, KAZUTO TASHI1, KEN MATSUKUMA1, KYOHO ITOH2, TOSHIHARU KAMURA1,7 and KIMIO USHIJIMA1

1Department of Obstetrics and Gynecology, Kurume University School of Medicine, Kurume, Fukuoka 830-0011; 2Cancer Vaccine Center, Kurume University Medical Center, Kurume, Fukuoka 839-0863; 3Biostatistics Center and 4Department of Immunology, Kurume University School of Medicine, Kurume, Fukuoka 830-0011, Japan

Received December 1, 2015; Accepted November 25, 2016

DOI: 10.3892/etm.2017.4150

Abstract. Currently prophylactic HPV16/18 L1 virus-like particle (VLP) vaccines are employed with great success for the prevention of HPV infection. However, limited information is available regarding the immune responses against human papillomavirus (HPV) 16/18 L1 subsequent to HPV16/18 L1 VLP vaccination, primarily due to the lack of widely used assays for immune monitoring. The aim of the present study was to identify HPV16 L1-derived B and T cell epitopes for monitoring the immune responses after HPV16/18 L1 VLP vaccination in healthy females. The levels of immunoglobulin G (IgG), IgE, IgA and IgM reactive to HPV16 L1-derived peptides were measured by multiplex bead suspension assay. Following detailed B cell epitope mapping, T cell responses specific to HPV16 L1-derived peptides were evaluated by an IFN-γ ELISPOT assay. The levels of IgG, IgM and IgA reactive to 20-mer peptides (PTPSGSMVTSDAQIFNKPYW) at positions 293-312 and 300-319 of HPV16 L1 were significantly increased in the plasma after 2, 7, and 12 months after first vaccination. Detailed epitope mapping identified the amino acid sequence (TSDAQIFNKP) at position 301-310 of HPV16 L1 as an immunogenic B cell epitope. In addition, T cell responses to an HLA-A2- and HLA-A24-restricted epitope (QIFNKPYWL) at position 305-313 of HPV16 L1 were increased following immunization, suggesting that the HPV16/18 L1-VLP vaccination as able to induce specific immune responses in T and B cells simultaneously. The identified B and T cell epitopes may be useful as a biomarker for monitoring the immune responses subsequent to HPV16/18 L1 VLP vaccination. Thus, the present study may provide novel information to improve the understanding of the immune responses to HPV16 L1.

Introduction

Cervical cancer is the fourth most prevalent cancer in women worldwide (1). The main cause of this disease has been known to be an infection of specific types of human papillomavirus (HPV), including HPV types 16 and 18 (2,3). Global epidemic studies of cervical screening with cervical cytology have demonstrated its efficacy. The greatest decline in cervical cancer-associated mortality was 3% per year between 1950 and 1970 (4). Currently, prophylactic HPV16/18 L1-virus-like particle (VLP) vaccines, which induce neutralizing antibody responses (5,6), have also been used with great success and show extremely high preventive effect against HPV16/18 infection (7,8). For instance, the preventive effect of the HPV16/18 L1-VLP vaccines has been reported to last up to 8.4 years (9,10). Nevertheless, only limited information is currently available regarding the immune responses against HPV16/18 L1, primarily due to the lack of widely used assays for immune monitoring (11,12). Several assays, such as pseudovirion-based neutralization assay, enzyme-linked immunosorbent assay, competitive luminex immunoassay, and the in situ-purified
glutathione-S-transferase L1 fusion protein-based multiplex immunoassay (GST-L1 MIA), have been developed to monitor antibody responses to HPV (13-16). However, no standardized serological assay is currently available for the assessment of HPV-specific antibody responses, particularly for large scale examination. Therefore, the biomarkers to precisely monitor the effects of HPV16/18 L1-VLP vaccines remain to be identified.

In the present study, B and T cell responses to HPV16/18 L1 were investigated in healthy females in order to identify a biomarker for monitoring immune responses subsequent to the HPV16/18 L1-VLP vaccination.

**Materials and methods**

**Immunization with the bivalent HPV16/18 L1-VLP vaccine and sample collection.** The present study was approved by the Kurume University Ethical Committee (Kurume, Japan). After a full explanation of the protocol, a written informed consent was obtained from 10 healthy females prior to enrollment. The enrolled females, who were aged between 23 and 33 years with a mean age of 25.5±2.9 years, had a human leukocyte antigen (HLA) -A2 and/or HLA-A24. They were not subjected to cervical cytology examination or an HPV DNA test prior to participation to the present study as titration of neutralizing antibody against HPV is undetectable in 50% of infected women and is at a low level if detected. In addition, obtaining a cervical sample would cause an unnecessary burden to these healthy young women. The participants received intra-muscular injection (0.5 ml) of the bivalent HPV16/18 L1-VLP vaccine (Cervarix®; GlaxoSmithKline, London, UK), which contained HPV16 L1-VLP (20 µg) and HPV18 L1-VLP (20 µg) mixed with AS04 adjuvant [aluminum hydroxide (500 µg) and 3-deacylated monophosphoryl lipid A (50 µg)]. The vaccine was administered in three doses at 0, 1 and 6 months. In all cases, 20 ml of peripheral blood was collected prior to immunization and at 1, 2, 7, 12 and 18 months after the first immunization. Plasma was separated from whole blood and frozen at -80˚C until further use. Peripheral blood mononuclear cells (PBMCs) were obtained by Ficoll-Paque Plus (GE Healthcare, Uppsala, Sweden) density gradient centrifugation, and cryopreserved until further use.

**Peptides.** In total, 10 different HPV16 L1-derived peptides (20-mer) with binding motifs to both HLA-class I (A2 or A24) and HLA-class II (DR) were selected using MULTIPRED web software (antigen.12.a-star.edu.sg/multipred/; accessed on 12 February 2012), as reported previously (17), and are listed in Table I. For epitope mapping, 8 different 10-mer overlapping peptides were selected from the amino acid sequence from positions 295 to 325 of HPV16 L1. The synthetic peptides were purchased from Greiner Bio-One (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Each peptide was dissolved in dimethyl sulfoxide, and stored at -80˚C until use in further experiments.

**Measurement of immunoglobulin (Ig) levels reactive to HPV16 L1-derived peptides.** The Ig levels reactive to HPV16L1-derived peptides were measured by multiplex bead suspension array using the Luminex system (Luminex Corp., Austin, TX, USA) as previously described (18). In order to detect peptide-specific IgG, IgM and IgE levels, the beads were washed with wash buffer (0.05% Tween-20 in PBS) and incubated with 100 µl of biotinylated goat anti-human IgG (1:200; BA-3080), biotinylated goat anti-human IgM (1:200; BA-3020; both Vector Laboratories Inc., Burlingame, CA, USA) or biotinylated goat anti-human IgE (1:200; AHI0509; BioSource; Thermo Fisher Scientific, Inc.) antibodies for 1 h at 30˚C. To detect peptide-specific IgA, the beads were incubated with 100 µl goat anti-human IgA antibody (1:200; A80-102A; Bethyl Laboratories, Inc., Montgomery, TX, USA) for 1 h at 30˚C, followed by washing and subsequent incubation with 100 µl of biotinylated rabbit anti-goat IgG antibody (1:200; 55366; Cappel, MP Biomedicals, LLC, Solon, OH, USA) for 30 min at 30˚C. Following washing, the beads were incubated with 100 µl of streptavidin-PE (1:200; S-866; Molecular Probes; Thermo Fisher Scientific, Inc.) for 30 min at 30˚C, followed by washing and detection of fluorescence intensity units on the beads using the Luminex system (18).

**Measurement of T cell responses to an HPV-16 L1-derived peptide.** T cell responses specific to HPV16 L1-derived peptides were evaluated by IFN-γ ELISPOT assay (8223; Medical & Biological Laboratories Co., Ltd., Nagoya, Japan). Briefly, PBMCs (2x10^5 cells/well) isolated from whole peripheral blood samples of the participants were cultured with 10 µg/ml of synthetic peptides in 96 round well plates (Nunc; Thermo Fisher Scientific, Inc.) in 200 µl culture medium containing 45% AIM-V (Thermo Fisher Scientific, Inc.), 45% RPMI-1640 (Thermo Fisher Scientific, Inc.), 10% FBS (MP Biomedicals, Solon, OH, USA), 20 IU/ml interleukin-2 (Bio-Rad Laboratories, Inc., Hercules, CA, USA) and 0.1 mM MEM non-essential amino acid solution (Thermo Fisher Scientific, Inc.). Half of the medium was replaced with new medium containing the peptide (20 µM) at every 3 or 4 days. After 5 stimulations, T cells were stimulated with peptide-pulsed T2 cells (ATCC, Manassas, VA, USA) or CIR-A24 cells (kindly provided by Dr Masaumi Takiguchi, Kumamoto University, Kumamoto, Japan), and IFN-γ production in response to the specific peptide was determined in comparison with the response to irrelevant peptide. Cells were also tested for IFN-γ production in response to a negative control peptide, HLA-A2 restricted peptide (SLYNTVATL, synthesized; Greiner Bio-One, Kremsmünster, Austria) from human immunodeficiency virus (HIV) or HLA-A24 restricted HIV peptide (RYLRQQLLGI, synthesized; Greiner Bio-One). HIV peptides were used as negative control peptides since these peptides were irrelevant to HPV. Spot numbers of IFN-γ-secreting cells after 18-h incubation were determined by ELISPOT assay with an ELISPOT reader (ImmunoSpot S5 Versa Analyzer; Cellular Technology Ltd., Shaker Heights, OH, USA). All assays were performed in duplicate. Peptide-specific T cell responses were shown as the differences between the spot numbers per 1x10^5 PBMCs in response to the specific peptide and those in response to the control peptide. When the PBMCs from females with both HLA-A2 and HLA-A24 were used, mean values of spot numbers from the culture with peptide-pulsed T2 cells and those with peptide-pulsed CIR-A24 cells were used for calculation. PBMCs were available for this analysis in only 9 of 10 individuals at 2 months after vaccination, but were
Comparison of IgM, IgE and IgA levels prior to and following vaccination. The IgM, IgE, and IgA levels were then measured against HPV16 L1-derived peptides in plasma obtained from healthy females (n=10) before immunization and at 1, 2, 7, 12 and 18 months after the first immunization. As shown in Fig. 1, the IgG levels specific to Peptide 6 at 2 months after the first vaccination also demonstrated an increasing trend (P=0.080). IgE and IgA levels against Peptide 6 did not change significantly after any time of vaccination. However, there were no significant changes in the IgM levels against any peptides. Furthermore, the increase of IgG and IgA levels against Peptide 8 at 2 and 7 months after the first immunization showed a positive correlation (Fig. 3).

Results

The peptide-specific IgG responses (levels in FIU) to HPV16 L1-derived peptides following the HPV16/18 L1-VLP vaccination. The levels of IgG against HPV L1-derived 20-mer peptides were investigated in the plasma (100-fold dilution) of 10 healthy females at 1, 2, 7, 12 and 18 months after the first immunization and compared with the levels before immunization. Median levels of IgG from the 10 healthy females are shown.

Table I. HPV16 L1-derived peptides used in the present study and their binding motifs to HLA-A2 and HLA-A24.

<table>
<thead>
<tr>
<th>Peptide</th>
<th>Position</th>
<th>Sequence</th>
<th>Position</th>
<th>Sequence</th>
<th>Score</th>
<th>Position</th>
<th>Sequence</th>
<th>Score</th>
</tr>
</thead>
<tbody>
<tr>
<td>Pep 1</td>
<td>54-73</td>
<td>KPNNKILVPKVSGLQYRFV</td>
<td>60-68</td>
<td>ILVPKVSGL</td>
<td>30</td>
<td>59-68 KILVPKVSGL</td>
<td>14</td>
<td></td>
</tr>
<tr>
<td>Pep 2</td>
<td>392-422</td>
<td>HSMNSTILEDWNGFLQPPGG</td>
<td>398-406</td>
<td>ILEDWNFGGL</td>
<td>23</td>
<td>397-406 TILEDWNFGGL</td>
<td>16</td>
<td></td>
</tr>
<tr>
<td>Pep 3</td>
<td>62-81</td>
<td>VPKVSGLQYRFVRIHLDPDN</td>
<td>67-75</td>
<td>GLQYRFVRI</td>
<td>22</td>
<td>66-75 SGLQYRFVRI</td>
<td>24</td>
<td></td>
</tr>
<tr>
<td>Pep 4</td>
<td>112-131</td>
<td>PLGVGSIHGPLNKLQDelen</td>
<td>118-126</td>
<td>SGHPPLNKLQ</td>
<td>22</td>
<td>117-126 ISGHPPLNKLQ</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>Pep 5</td>
<td>243-262</td>
<td>GDSLLLFRQVFHRFLLFN</td>
<td>249-257</td>
<td>YLREQMFV</td>
<td>22</td>
<td>248-257 FYLRERQMFV</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>Pep 6</td>
<td>300-319</td>
<td>VTSQIFQIFNPYYQWLQRAQGH</td>
<td>305-313</td>
<td>QIFQIFNPYYQW</td>
<td>21</td>
<td>305-313 QIFQIFNPYYQW</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>Pep 7</td>
<td>144-162</td>
<td>RECISSMDYKQTLQCLIGCK</td>
<td>148-156</td>
<td>SMDYKQTLQCLIGCK</td>
<td>20</td>
<td>148-156 SMDYKQTLQCLIGCK</td>
<td>11</td>
<td></td>
</tr>
<tr>
<td>Pep 8</td>
<td>293-312</td>
<td>PTPSGSMVTSDAQNFKNKPYW</td>
<td>298-306</td>
<td>QNFKNKPYW</td>
<td>20</td>
<td>298-306 QNFKNKPYW</td>
<td>10</td>
<td></td>
</tr>
<tr>
<td>Pep 9</td>
<td>384-403</td>
<td>TADVMTYIHSMNSTILEDWN</td>
<td>390-399</td>
<td>YIHSMNSTILEDWN</td>
<td>20</td>
<td>389-398 YIHSMNSTILEDWN</td>
<td>10</td>
<td></td>
</tr>
<tr>
<td>Pep 10</td>
<td>152-171</td>
<td>KQTLQCLIGCKPPIGEHWG</td>
<td>157-165</td>
<td>CLIGCKPPIGEHWG</td>
<td>23</td>
<td>156-165 CLIGCKPPIGEHWG</td>
<td>12</td>
<td></td>
</tr>
</tbody>
</table>

Scores indicate the prediction of peptide binding with HLA on MULTIPRED. Scores are not presented for the peptide binding in HLA-DR as each peptide listed binds with several types of HLA-DR and the scores are different among each type. Anchor residues for HLA class I are shown in bold. HPV, human papillomavirus; HLA, human leukocyte antigen; Pep, peptide.

Statistical analysis. The levels of IgG, IgM, IgE and IgA specific to HPV-16 L1-derived peptides were compared at the indicated time-points prior to and following the vaccination using Wilcoxon signed-rank test. All reported P-values were two-sided and statistically significant differences were indicated by P<0.05. Comparison of IgG and IgA response was compared by analysis of variance. Analyses were performed with the use of JMP version 11 (SAS Institute Inc., Cary, NC, USA).

available in all 10 individuals at other time points prior to and following vaccination.
to these findings, it is hypothesized that the overlapped position of Peptides 6 and 8 may be a B cell epitope.

**Detailed B cell epitope mapping.** To obtain further detailed information on immunogenic epitopes, the levels of IgG reactive to each of the 8 different 10-mer peptides derived from the amino acid sequence between positions 295 and 325 of HPV16 L1 were determined, which contained the overlapped position of Peptide 6 and Peptide 8 sequences (position 300-312: VTSDAQIFNKPYW). As shown in Fig. 4, the IgG levels specific to the peptide at position 301-310 (VTSDAQIFNKP) of HPV16 L1, but not specific to other 10-mer peptides, were significantly increased at 7 months after first vaccination compared with before vaccination (P=0.004). Notably, this amino acid sequence was shared by the immunogenic 20-mer Peptides 6 and 8.

The present study also analyzed the T cell responses to an HPV16 L1-derived peptide in PBMCs subsequent to vaccination. T cell responses to an HPV16 L1-derived peptide at position 305-313 (QIFNKPYWL) of HPV16 L1 in the PBMC cultures before and after vaccinations were further assessed. The peptide at position 305-313 (QIFNKPYWL) of HPV16 L1 (was used in the current experiment since it was previously demonstrated that it binds well to both HLA-A*0201 and HLA-A*2402 (17). PBMCs were stimulated in vitro with specific or control peptides, and then their IFN-γ production in response to the peptide-pulsed T2 (HLA-A2+) or CIR-A24 (HLA-A24+) cells was examined by ELISPOT assay. As shown in Fig. 5, the peptide-specific T cell responses at 1 month after the first immunization demonstrated significant increase compared with the response prior to vaccination (P=0.026). These results suggest that the HPV16/18 L1-VLP vaccination was able to induce specific immune responses in T and B cells simultaneously.

**Discussion**

In the present study, the humoral and cellular immune responses to HPV16 L1-derived peptides were analyzed in healthy females subsequent to immunization with the HPV16/18 L1-VLP vaccine. When B cell responses to 10 different 20-mer peptides were screened, the levels of IgG of Peptides 6 and 8 were significantly elevated in the plasma at 2, 7 and 12 months after first vaccination compared with before vaccination, and the levels of IgG of Peptide 4 were significantly elevated in the plasma at 2 and 7 months after the first vaccination, suggesting an isotype switch to IgE and IgA. Similarly, the levels of IgA against Peptide 6 showed an increasing trend at 2 months after the first immunization. Although it is reported that the dominant antibody isotype in the cervical mucosa after immunization with HPV 16/18 L1-VLP is IgG, IgA is known to be the dominant antibody isotype in the mucosal immune system. Hence, IgA against the Peptide 6 and 8 may transudate and/or exudate from the systemic circulation to the cervical mucosa, and thus can be detectable in the mucosa after HPV16/18 L1-VLP vaccination.

Previously, we identified the B cell epitopes derived from HPV16 L1 in BALB/c and C57BL/6 N mice (17). Similar to the results obtained in humans in the present study, Peptide 6 was the major B cell epitope shared by female Balb/c (H-2d) and C57BL/6 N (H-2b) mice in our previous study (17). Nevertheless, detailed mapping analysis of B cell epitopes in humans indicated that a 10-mer peptide (TSDAQIFNKP) at the position 301-310 of HPV16 L1 was immunogenic, whereas another 10-mer peptide (QIFNKPYWL) at position 304-313, which was identified as the B cell epitope in mice (17), was not immunogenic. This finding suggested that immunogenic B cell epitopes may be slightly different between mice and humans, although the reason for this discrepancy remains to be determined.
suggested that the HPV16/18 L1-VLP vaccination was able to induce specific immune responses in T and B cells simultaneously.

In the present study, the epitope recognized by B cells (TSDAQIFNKP) was not completely identical to that recognized by T cells (QIFNKPYWL); however, 6 out of 9 or 10 amino acids were shared by these two epitopes. Since it has been well recognized that cellular and humoral immune responses are crucial in the induction of effective immunity in animal models (19-21), simultaneous induction of both B and T cell responses against HPV16 L1 may be important for efficient prevention of HPV infection and subsequent HPV-associated neoplasia.

In conclusion, a 10-mer amino acid sequence (TSDAQIFNKP) derived from HPV16 L1 was identified in the present study as an immunogenic B cell epitope in the females after HPV16/18 L1-VLP immunization. In addition, T cell responses to another HLA-A2- and HLA-A24-restricted epitope (QIFNKPYWL) derived from HPV16 L1 in the vaccinated individuals were detected. The identified B and T cell epitopes may be useful for monitoring the immune responses to HPV16/18 L1-VLP subsequent to vaccination. Whether the
vaccine-induced IgG and/or IgA response to the identified B cell epitope possesses the biological activity to neutralize the HPV 16/18 L1 vaccine remains to be clarified in the near future. However, the present study may provide novel information for better understanding the immune responses to HPV 16 L1-VLP vaccines.

Acknowledgements

The present research was supported in part by The Fukuoka Obgyn Researcher’s Charity Foundation Fund, Japan.

References


