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Expression of a recombinant *Human papillomavirus* 16 E6GT oncoprotein fused to N- and C-termini of *Potato virus X* coat protein in *Nicotiana benthamiana*

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Abstract Infection with Human papillomaviruses (HPVs) is linked to cervical cancer, which is one of the most common cancers found among women worldwide. Despite preventive immunization, a therapeutic vaccine, targeting infected individuals, is required. Vaccine strategies for treatment of HPV-induced cervical cancer are based on E7 and E6 oncoproteins. In this work we report the transient expression of chimeric particles containing the E6 oncoprotein from Human papillomavirus type 16 (HPV-16) in plants. We fused a mutagenized coding sequence of the E6 oncoprotein (E6GT) with the coding sequence of Potato virus X coat protein (PVX CP) both with the 5'- and 3'-terminus using linkers of different length (0, 4 or 15 amino acids). The expression in E. coli was performed to assess the characteristics of the recombinant protein prior to plant expression. The yield and immunological reactivity of the expressed proteins were screened with anti-PVX CP and E6 antibodies. The highest yields of chimeric particles were observed in the transgenic N. benthamiana expressing Potato virus A HC-Pro protein and the Tobacco

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I. Polakova · M. Duskova · M. Smahel Department of Experimental Virology, Institute of Hematology and Blood Transfusion, U Nemocnice 1, 12820 Prague 2, Czech Republic *mosaic virus* movement protein. When inoculated on host plants, these recombinant viruses were not able to spread systemically. The obtained results revealed the new relations for design of expression cassettes for plant-based vaccine production.

Keywords Transient expression \cdot Molecular farming \cdot *Potato virus X* \cdot *Human papillomavirus* \cdot Oncoprotein E6 \cdot Bacterial expression

Introduction

Human papillomaviruses (HPVs) cause cervical carcinoma and are associated with several other human anogenital cancers (Ganguly and Parihar 2009). They are small viruses with a double-stranded circular DNA, which replicates in the nucleus of infected cells. Among the HPV proteins, the early proteins E6 and E7 attract most attention as they transform and immortalize cells (Ganguly and Parihar 2009; McLaughlin-Drubin and Munger 2009; Hsu et al. 2011).

The availability of prophylactic vaccines against HPVs (based on HPV-16, -18, -6, and -11 L1 structural proteins assembled in virus-like particles—VLPs) represents a milestone in the prevention of this infection (Koutsky et al. 2002; Villa et al. 2005; Harper et al. 2006), establishing the base for a significant reduction of the rate of cervical cancer in future. Despite this preventive immunization, a therapeutic vaccine, targeting already infected individuals, is also required. Many candidate vaccines with therapeutic potential are currently tested in ongoing trials; however, there is low expectation that any of the current therapeutic vaccines will have a substantial public health impact in the near future (Ma et al. 2010). Several

sed on E7 oncopro- Materials and methods herapeutic vaccines,

Cloning of fusion constructs

To avoid safety concerns associated with the administration of an oncogene in humans, we used an attenuated HPV-16 E6 gene, termed E6GT, that was demonstrated to prevent E6-mediated p53 degradation: it carries two amino acid substitutions, C70G and I135T (Polakova et al. 2010).

E6GT was amplified by PCR from vector pBSC-E6GT using either G9-R and G9-U or G9-P and G9-O primers, respectively (Table 1; Fig. 1). The first PCR product was inserted into the PVX CP-BH/pMPM-A4 Ω plasmid (Plchova et al. 2011) using the *Xba*I and *BgI*II restriction sites to create the final expression plasmid E6GT-PVX CP/ pMPM-A4 Ω carrying the E6GT fused to the 5'-terminus of PVX CP. The second PCR product was cloned into the PVX CP-NB/pMPM-A4 Ω plasmid (Plchova et al. 2011) using the *BgI*II and *SaI*I restriction sites to produce the final PVX CP-E6GT/pMPM-A4 Ω expression plasmid carrying the E6GT fused to the 3'-terminus of PVX CP.

The fusions of E6GT with the 4 amino acids linker L4 (amino acids GPGP) (Massa et al. 2008) either at the 3'-terminus of E6GT or at the 5'-terminus of E6GT were prepared by PCR from vector pBSC-E6GT using the primer pairs G9-R and G9-V or G9-Q and G9-O, respectively (Table 1). The first PCR product was cloned into the PVX CP-BH/pMPM-A4 Ω plasmid using the *XbaI* and *BglII* restriction sites to create the final expression plasmid, which was termed E6GT-L4-PVX CP/pMPM-A4 Ω . The product of the second PCR reaction was ligated into the PVX CP-NB/pMPM-A4 Ω plasmid using the *BglII* and *SaII* restriction sites to produce the final PVX CP-L4-E6GT/pMPM-A4 Ω expression plasmid.

The 'splicing by overlap extension by PCR' method (Warens et al. 1997) was used to fuse the E6GT coding sequence with the 15 amino acids beta-sheet linker L15 [amino acids (SGGGG)₃] (Alfthan et al. 1995) either at the 3'- or at the 5'-terminus. The E6GT sequences (named E6GT-RS and E6GT-NO) were PCR amplified from vector pBSC-E6GT using the primer pairs G9-R and G9-S or G9-N and G9-O, respectively. The L15 linkers (named L15-TJ and L15-AM, respectively) were PCR amplified from vector pTZ-CP-Linker-E7 K + rc (unpublished results) using the primer pairs G9-T and G9-J or G9-A and G9-M, respectively. Two PCR products (either E6GT-RS and L15-TJ or E6GT-NO and L15-AM) were mixed in equimolar amounts. The second-stage PCR was performed using external primers (either G9-R and G9-J or G9-A and G9-M) to produce E6GT-L15 or L15-E6GT products. The E6GT-L15 product was inserted into the PVX CP-BH/ pMPM-A4 Ω plasmid using the XbaI and BglII restriction sites to create the final expression plasmid, which was

therapeutic HPV-specific vaccines based on E7 oncoprotein, which is the primary target of therapeutic vaccines, designed to treat later stages of the disease, have been tested in animal models and some have advanced into phase II and III clinical trials (Münger et al. 1989; Billich 2003; Frazer 2004). There is an increasing demand for gene expression systems that allow for high levels of foreign protein production in plants, whether transgenically, transiently or by means of a virus vector (Scholthof et al. 2002; Canizares et al. 2005; Faye and Gomord 2010; Anwar et al. 2011; Kim et al. 2011; Martínez-González et al. 2011).

Vectors derived from *Potato virus X* (PVX) are widely used to rapidly express high levels of recombinant proteins e.g. vaccines in plants. PVX coat protein (PVX CP) fusion with its ability to form virus–like particles is a suitable carrier for display of antigens accessible to the immune system.

There were numerous attempts to produce vaccines using N-terminal fusions of antigens with PVX CP which were able to form virus-like particles (Marusic et al. 2001; Lico et al. 2006, 2009; Uhde-Holzem et al. 2007; Marconi et al. 2009; Morgenfeld et al. 2009). There is also number of works regarding expression of HPV antigens in plants (Franconi et al. 2002; Biemelt et al. 2003; Varsani et al. 2003; Liu et al. 2005; Maclean et al. 2007; Fernández-San Millán et al. 2008; De la Rosa et al. 2009). However, successful expression of the E6 oncoprotein in plants has not yet been reported. Therefore we designed a transient expression system for this antigen as N- or C-terminal PVX CP fusions first in *E. coli* to determine their properties, and then for expression in plants.

The mutagenized HPV16 E6 protein (E6GT) has been proposed as a safer candidate than the wild type E6 for vaccine development. The introduction of double mutation into the E6 protein led to the loss of the ability to induce the p53 degradation (Polakova et al. 2010).

We previously focused on the development of transient expression systems based on plant viral vectors for production of different epitopes from *Human papillomavirus* type 16 (HPV-16) minor coat protein L2 and oncoprotein E7 or on the expression of mutagenized HPV-16 E7 oncoprotein in plants (Cerovska et al. 2008, 2012; Hoffmeisterova et al. 2012; Plchova et al. 2011).

To display E6GT on the surface of PVX CP VLPs, we used a transient expression system previously designed for expression of the E7ggg protein (Plchova et al. 2011). This system allows to express different full-length antigens fused either to the N- or C-terminus of PVX CP using linkers of different lengths to find the optimal expression. We studied the impact of the topology of the fusion and the effect of linker length on expression levels both in *Escherichia coli* and *Nicotiana benthamiana*.

Table 1 Oligonucleotides used for preparation of chimeric constructs in pMPM-A4 Ω

Primers	5'-3' Sequence	Restriction sites	Name of construct
G9-A	CATT <u>AGATCT</u> GGTGGAGGTGGCTCTG	BglII	PVX CP-L15-E6GT/pMPM-A4Ω
G9-M	CAGTTCTCTTTTGGTGTGATCCGCCGCCACC	-	
G9-N	TGGCGGCGGATCACACCAAAAGAGAACTGCAA	-	
G9-O	CATT <u>GTCGAC</u> TTACAGCTGGGTTTCTCTA	SalI	
G9-P	CATT <u>AGATCT</u> CACCAAAAGAGAACTGCAA	BglII	PVX CP-E6GT/pMPM-A4Ω
G9-Q	CATT <u>AGATCT</u> GGTCCGGGTCCTCACCAAAAGAGAACTGCAA	BglII	PVX CP-L4-E6GT/pMPM-A4Ω
G9-R	CATT <u>TCTAGA</u> GATGCACCAAAAGAGAACTG	XbaI	E6GT-L15-PVX CP/pMPM-A4Ω
G9-S	AGCCACCTCCACCCAGCTGGGTTTCTCTACG	-	
G9-T	GAGAAACCCAGCTGGGTGGAGGTGGCTCTG	-	
G9-J	CATT <u>AGATCT</u> TGATCCGCCGCCACCA	BglII	
G9-U	CATTAGATCTCAGCTGGGTTTCTCTACG	BglII	E6GT-PVX CP/pMPM-A4 Ω
G9-V	CATT <u>AGATCT</u> AGGACCCGGACCCAGCTGGGTTTCTCTACG	BglII	E6GT-L4-PVX CP/pMPM-A4Ω

Underlined sequences are restriction endonuclease sites

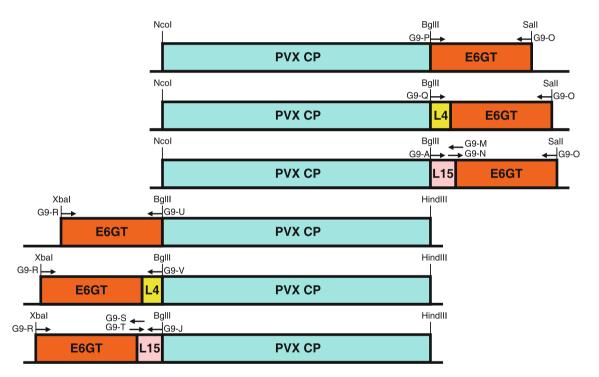


Fig. 1 Schematic illustration of the genetic constructs in bacterial expression vector pMPM-A4 Ω . Restriction endonuclease sites and positions of oligonucleotides (*black arrows*) used for preparation of

termed E6GT-L15-PVX CP/pMPM-A4 Ω . The L15-E6GT product was cloned into the PVX CP-NB/pMPM-A4 Ω plasmid using the *BgI*II and *SaI*I restriction sites to produce the final PVX CP-L15-E6GT/pMPM-A4 Ω expression plasmid (Fig. 1). All PCR reactions were performed using Phusion High-Fidelity DNA polymerase (Finnzymes, Espoo, Finland).

The 3'-terminal part of constructs having the E6GT fused to the 3'-terminus of PVX CP was then inserted into the infectious PVX full length clone pGR106 (kindly

chimeric constructs are indicated. *E6GT*—mutated form of E6 from HPV16; *L4*—4 amino acids linker; *L15*—15 amino acids linker; *PVX CP*—*Potato virus X* coat protein

provided by DC Baulcombe, The Sainsbury Laboratory, Norwich, UK; Chapman et al. 1992), cleaved by *XhoI* to produce the PVX CP-E6GT/pGR106, PVX CP-L4-E6GT/ pGR106 and PVX CP-L15-E6GT/pGR106 expression plasmids. Thus, these constructs contained the E6GT sequence followed by the repetition of the 3'-terminus of PVX CP and usual pGR106 sequence with its *cis* acting elements in the 3'-untranslated region. All constructs with E6GT fused to the 5'-terminus of PVX CP were then inserted into pGR106 cleaved by *XbaI* and *XhoI*, so the PVX CP present in pGR106 was replaced by chimeric constructs to form E6GT-PVX CP/pGR106, E6GT-L4-PVX CP/pGR106 and E6GT-L15-PVX CP/pGR106, respectively. The constructs were checked for their accuracy by DNA sequencing.

Protein expression in E. coli

The chimeric proteins were expressed from the pMPM-A4 Ω plasmid in *E. coli* MC1061 cells. The optimal conditions for protein expression required incubation for 12 h at 25 °C following induction with 0.2 % arabinose.

Partial purification of PVX CP fused to E6GT – enriched fractions

Fusion proteins between PVX CP and E6GT were partially purified from 500 mL of the bacterial culture. The cells were centrifuged at $15,000 \times g$ for 15 min at 4 °C. The pellet was resuspended in 50 mL 20 mM Tris-HCl pH 7.5 with 0.2 mg/mL lysozyme (Sigma) and 0.1 mg/mL DNAse I (Roche), and incubated at 37 °C for 10 min. After homogenisation in glass homogenisator on ice the bacterial lysate was centrifuged. The supernatant containing most of the soluble protein was subjected to further purification on sucrose cushion (Cerovska et al. 2004). The presence of fusion proteins was tested by ELISA (dilution 1 and 10 µg/mL in 15 mM Na₂CO₃, 35 mM NaHCO₃, 3 mM NaN₃, pH 9.6). The insoluble fusion proteins were solubilized using 8 M urea. Protein concentration in each fraction was determined by the Bradford method using BSA as a protein standard (Bradford 1976). The fusion proteins from plant material were purified according to Cerovska et al. (2004).

Plant propagation

Four genotypes of Nicotiana benthamiana were evaluated as hosts for expression of modified PVX CPs: N. benthamiana wild type (wt), N. benthamiana carrying the coding sequence of Potato virus A HC-Pro protein (HC-Pro; kindly provided by E. Savenkov, Department of Plant Biology, Sveriges Lantbruksuniversitet, Uppsala, Sweden), N. benthamiana 3H carrying the coding sequence of *Tobacco mosaic virus* movement protein (3H; kindly provided by R. Beachy, Donald Danforth Plant Science Center, St. Louis, USA) and N. benthamiana K7 plants, which were obtained in our laboratory by crossing N. benthamiana HC-Pro and N. benthamiana 3H. All plants were grown in a growth chamber under controlled conditions (20-25 °C over day, 15-20 °C over night, 16 h-light/8 h-dark). Eighteen plants per group were inoculated in their four-leaf stage. All experiments were performed in triplicate.

Agrobacterium transformation and plant inoculation

The binary vectors with full-length infectious cDNA carrying the PVX CP fusion genes were isolated from E. coli and introduced into chemically competent Agrobacterium tumefaciens (GV3101) using freeze-thaw method (An 1987). Bacteria containing the plasmids were selected on LB-agar plates with 30 µg/mL kanamycin. For transient expression, the resulting Agrobacterium strains induced by acetosyringone were used to infiltrate N. benthamiana leaves. Bacterial suspension for agroinfection of plants was prepared in LB-broth containing 30 µg/mL kanamycin at 28 °C. The primary infection of the host plants was performed by infiltration using 1 mL hypodermic syringe without needle (Hoffmeisterova et al. 2008). As a negative control, leaves inoculated with LB medium were used. As a positive control, N. benthamiana K7 agroinfiltrated with A. tumefaciens containing the pGR106 vector was used.

Polyacrylamide gel electrophoresis (SDS-PAGE)

1 mL aliquots of *E. coli* MC1061 cells were pelleted by 1 min centrifugation, resuspended in 100 μ L of the Laemmli buffer, boiled for 2 min and aliquots were loaded on 12 % polyacrylamide gel containing SDS (Laemmli 1970). The plant samples were homogenized using the FastPrep (MP Biomedicals, Solon, OH USA) system and crude transgenic plant extracts were diluted to a concentration of 1 mg/mL total soluble proteins in the Laemmli buffer containing 6 M urea, placed for 10 min into boiling water, cooled to room temperature and then subjected to SDS-PAGE.

Western blot analysis

The proteins separated by SDS-PAGE were electroblotted to a nitrocellulose membrane (0.2 µm, Sleicher&Schuell Protran) in semidry system (OMNI-TRANS apparatus, Omnibio Brno, Czech Republic) (Hirano and Watanabe 1990). The membrane was then incubated for 1 h in 4 %BSA in PBS + 0.5 % Tween 20 (PBST) and then washed 4×5 min in PBST. The primary rabbit polyclonal anti-PVX CP antibodies (dilution 1:1,000 in PBST; PrimeDiagnostics, Wageningen, Netherland) and secondary goat anti-rabbit antibodies conjugated to alkaline phosphatase (AP) (1:30,000 in PBST; Sigma, St. Louis, MO, USA) were used to detect fusion proteins expressed both in E. coli and in N. benthamiana. The primary goat polyclonal anti-HPV16 E6 (N-17) (1:1,000; Santa Cruz Biotechnology, Santa Cruz, CA, USA) and the secondary rabbit antigoat-AP antibodies (1:30,000 in PBST; Sigma, St. Louis, MO, USA) were applied to detect fusion proteins expressed in N. benthamiana. The bands of interest were visualized

by reaction with a substrate of AP (BCIP/NBT; Sigma, St. Louis, MO, USA) (Sambrook et al. 1989).

Plate-trapped antigen ELISA (PTA-ELISA)

The PTA-ELISA test was performed as described by Mowat (1985). As primary antibodies the rabbit polyclonal anti-PVX CP (1:1,000; Sigma, St. Louis, MO, USA) or goat anti-HPV16 E6 (N-17) (1:1,000; Santa Cruz Biotechnology, Santa Cruz, CA, USA) were used. As secondary antibodies the AP-conjugated goat anti-rabbit or rabbit anti-goat IgG (Sigma, St. Louis, MO, USA) in dilutions recommended by manufacturers were used. All ELISA measurements were performed in duplicates and repeated three times.

Electron microscopy

Copper grids with carbon support layer were floated on droplets of purified chimeric VLPs (obtained from bacterial or plant expression; 0.8 mg/mL) at room temperature for 40 min. The grids were washed with PBS for 2 min, then with H₂O for 2 min (repeated three times). The non-labeled samples were stained with drops of 2 % uranyl acetate at room temperature for 15 min. For the purposes of immunogold-labeling, polyclonal antibodies recognizing PVX CP or E6 were bound to the appropriate parts of the samples by incubating the grid with the rabbit anti-PVX CP IgG (dilution 1:1,000 in PBS; PrimeDiagnostics, Wageningen, Netherland) or goat-anti HPV16 E6 (N-17) (dilution 1:1,000 in PBS; Santa Cruz Biotechnology, Santa Cruz, CA, USA) at room temperature for 40 min. After washing with PBS, samples were decorated with gold-labeled goat anti-rabbit IgG or rabbit-anti goat (10 nm or 15 nm gold particles, Sigma, St. Louis, MO, USA) by floating the grids on a 1:20 PBS dilution of the IgG at room temperature for 40 min. After washing with PBS, the samples were stained with drops of 2 % uranyl acetate (15 min at room temperature). Electron microscopy was carried out using a JEM 1010 transmission electron microscope (Jeol, Japan, facility of Biology Centre of the Academy of Sciences of the Czech Republic, Ceske Budejovice, Czech Republic).

Results

Cloning of fusion constructs

Five different PVX CP constructs (E6GT-PVX CP, E6GT-L4-PVX CP, PVX CP-E6GT, PVX CP-L4-E6GT and PVX CP-L15-E6GT) with the HPV-16 E6GT gene fused either to 5'- or 3'-terminus of PVX CP were prepared (Fig. 1) and then in pGR106 (Fig. 2a, b). We did not obtain the E6GT-

L15-PVX CP construct, since in all repeated cloning experiments the L15 linker was about from 21 to 30 nucleotides shorter than expected.

Expression and purification of fusion proteins in E. coli

All fused proteins were expressed at high levels in *E. coli* except for the E6GT-L15-PVX CP. The theoretical molecular weight of the fusion proteins was approximately 44 kDa, with small differences caused by the presence of linkers, in the case of L4 about 0.3 kDa and in the case of L15 about 1 kDa. All fusion proteins expressed in *E. coli* showed bands of the appropriate size on the Western blots (Fig. 3a, b).

The yields of purified C-fusion proteins expressed in *E. coli* were 1.85 ± 0.4 mg/mL. In comparison with the C-fusions significantly larger quantities of N-fusion proteins were produced in *E. coli* inclusion bodies. The yields of the N-terminal fusion proteins (both the soluble part and pellets) were 2.3 ± 0.4 mg/mL (P < 0.05).

Although most of the N-terminal fusion proteins were soluble, a small part of them remained in the pellet; purification of C-terminal fusion proteins showed higher amounts of recombinant proteins in the soluble fraction.

Production of virus-like particles in E. coli

The partially purified fusion proteins expressed in *E. coli* were subjected to electron microscopy. The chimeric VLPs were visualized with 2 % uranyl acetate. The putative VLPs showed similar shape and length as the other particles formed by recombinant PVX CP expressed in *E. coli* (Maclean et al. 2007; Fernández-San Millán et al. 2008). Immunogold-labeling using polyclonal antibodies against PVX CP proved that all our fusion proteins form chimeric particles. For illustration, chimeric VLPs of expressed fusion protein PVX CP-L4-E6GT are shown in Fig. 4a.

Selection of the most suitable plant genotype

The *N. benthamiana* K7 plants were shown to provide the highest expression of target proteins, about 50-times more than wt and HC-Pro plants and about twice as much as 3H plants, as evidenced by SDS-PAGE/Western blot analysis. For illustration, expression of fusion protein PVX CP-L15-E6GT is shown in Fig. 5.

Expression and purification of fusion proteins in *N*. *benthamiana* K7 plants

All constructs were subsequently expressed in *N. benth-amiana* K7 plants. The infection was initiated by agroin-filtration using a syringe without needle (Hoffmeisterova

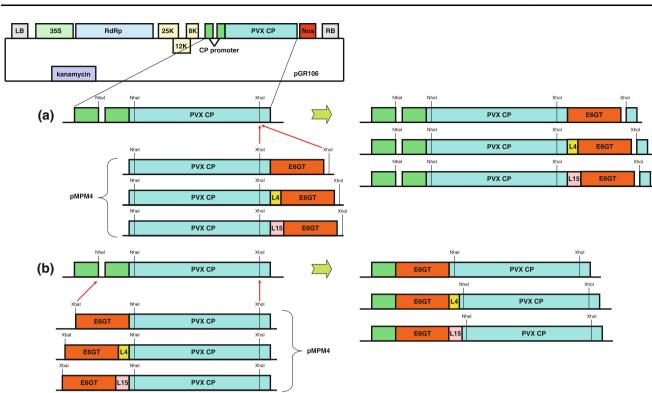


Fig. 2 Cloning strategy. **a** Cloning of 3'-terminally fused constructs into pGR106. To fuse PVX CP with E6GT and linkers L4 or L15 the restriction sites *Xho*I were used. The final constructs contain the duplicated 3'-terminus of PVX CP. **b** Cloning of 5'-terminally fused constructs into pGR106. The constructs were cut out from pMPM-A4 Ω by *Xba*I and *Xho*I and inserted into pGR106 cleaved by *Nhe*I

and *XhoI*. *LB* left border, *RB* right border, *35S* Cauliflower mosaic virus 35S promoter, *RdRp* RNA-dependent RNA polymerase, *25 K*, *12 K and 8 K* triple gene block proteins, *PVX CP Potato virus X* coat protein, *CP promoter* promoter of PVX CP, *Nos* nopaline synthase terminator, *E6GT* mutated form of E6 from HPV16, *L4* s4 amino acids linker; *L15* 15 amino acids linker

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et al. 2008). In the following mechanical inoculations the extracts from agroinfiltrated leaves were not infectious.

The presence of fusion genes and their products in harvested *N. benthamiana* K7 leaves was determined by immunocapture (IC) RT-PCR (Fig. 6), plate-trapped antigen (PTA) ELISA and Western blot, using IgG antibodies specific to PVX CP and E6. The ELISA absorbances of samples prepared from leaves inoculated with LB medium (negative controls) were comparable to the absorbances of healthy leaves without LB treatment.

We could not confirm the expression of N-terminal fusion proteins in plants by immunoblotting, therefore we isolated total RNA from the appropriate infected leaves and performed RT-PCR. After sequencing the RT-PCR products, we found out that nucleotide sequences of the E6GT fused to the 5'-terminus of PVX CP were mutated. The mutations occured in the region of nucleotides 170–324 in the case of the E6GT-PVX CP construct and in the region of nucleotides 74–324 in the case of the E6GT-L4-PVX CP construct. Both regions accumulated different nucleotide substitutions that resulted in amino acid changes and premature stop codons. It seems that after the infection of plants the PVX virus does not tolerate the 5'-terminal E6GT fusion. On the contrary, the presence of all C-terminal fusion proteins in *N. benthamiana* K7 plants was confirmed both by PTA ELISA and Western blot analysis with antibodies against PVX CP and E6 (Fig. 3c) and by IC RT-PCR (Fig. 6). The amounts of chimeric proteins were determined by SDS-PAGE/Western blot analysis using anti-E6 antibody (Fig. 7). The levels of all expressed C-terminal fusion proteins were determined using serial dilutions of PVX CP-L4-E6GT expressed in bacteria as a standard. The average yields of all C-terminal PVX CP fusion proteins after purification were approximately 5 ± 1.2 mg/kg (P < 0.05) of fresh leaf tissue.

Western blot analysis of different plant parts showed that the fusion proteins were present only in inoculated leaves and that the infection did not spread systemically as did the wild type virus (Fig. 5).

Production of virus-like particles in plants

Immunogold-labeling with polyclonal antibodies against PVX CP and E6 confirmed the presence and accessibility of the desired epitopes on the surface of the purified

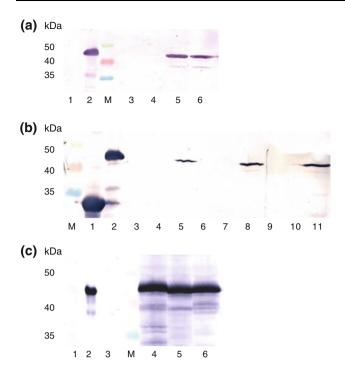


Fig. 3 Western blot analysis of fusion proteins PVX CP E6GT expressed in bacteria and plants. a Western blot analysis of N-terminally fused proteins PVX CP E6GT expressed in E. coli MC1061. 1 pMPM-A4Ω in MC1061; 2 positive control (E7GGG-PVX CP); M M SpectraTM Multicolor Broad Range Protein Ladder (10-260 kDa), Fermentas; 3 E6GT-PVX CP before induction; 4 E6GT-L4-PVX CP before induction; 5 E6GT-PVX CP after induction; 6 E6GT-L4-PVX CP after induction. b Western blot analysis of C-terminally fused proteins PVX CP E6GT expressed in E. coli MC1061. M M SpectraTM Multicolor Broad Range Protein Ladder (10-260 kDa), Fermentas; 1 PVX CP in MC1061; 2 positive control (E7GGG-PVX CP); 3 PVX CP-L15-E6GT before induction; 4 PVX CP-L15-E6GT not induced; 5 PVX CP-L15-E6GT after induction; 6 PVX CP-E6GT before induction; 7 PVX CP-E6GT without induction; 8 PVX CP-E6GT after induction; 9 PVX CP-L4-E6GT before induction; 10 PVX CP-L4-E6GT without induction; 11 PVX CP-L4-E6GT after induction. c Western blot analysis of C-terminally fused proteins PVX CP E6GT expressed in N. benthamiana K7 plants. 1 250 ng of purified PVX; 2 250 ng of PVX CP-L4-E6GT purified from *E. coli*, $\overline{3}$ not infected plant; \widetilde{M} M SpectraTM Multicolor Broad Range Protein Ladder (10-260 kDa), Fermentas; 4 PVX CP-L15-E6GT in inoculated leaves 12 days after inoculation (dpi); 5 PVX CP-E6GT in inoculated leaves 12 dpi; 6 PVX CP-L4-E6GT in inoculated leaves 12 dpi

particles. Chimeric VLPs of PVX CP-L4-E6GT fusion protein are shown for illustration (Fig. 4b).

Discussion

Cervical carcinomas are almost universally associated with high-risk human papillomavirus infections and are a leading cause of cancer death in women worldwide. HPV oncoproteins contribute to cancer initiation and progression and their expression is necessary for the maintenance of the

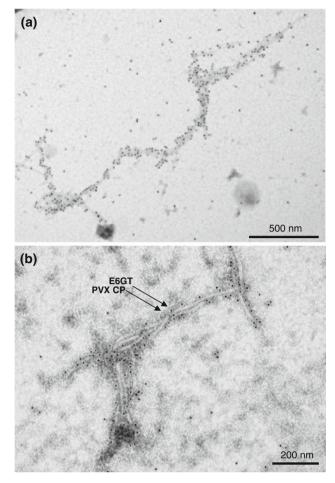


Fig. 4 Immunoelectron microscopy of chimeric virus-like particles of expressed fusion protein PVX CP-L4-E6GT using negative staining by 2 % uranyl acetate. a Immunochemical labeling of PVX CP-L4-E6GT isolated from E. coli MC1061. The particles were decorated with rabbit anti-PVX CP (1:1,000 in PBS; PrimeDiagnostics, Wageningen, Netherland); secondary goat anti-rabbit 10 nm gold conjugate (1:20 in PBS; Sigma, St. Louis, MO, USA). Bar represents 500 nm; b Immunochemical labeling of PVX CP-L4-E6GT partially purified from N. benthamiana K7 plants. The particles were decorated with polyclonal goat anti-HPV16 E6 (N-17) antibodies (1:1,000 in PBS; Santa Cruz Biotechnology, Santa Cruz, CA, USA), secondary rabbit anti-goat antibodies conjugated with 10 nm Au (1:20; Sigma, St. Louis, MO, USA), polyclonal rabbit anti-PVX CP antibodies (1:1,000 in PBS; PrimeDiagnostics, Wageningen, Netherland), secondary goat anti-rabbit antibodies conjugated with 5 nm Au (1:20 in PBS; Sigma, St. Louis, MO, USA). Bar represents 200 nm

transformed state. Disease prevention has been realized by introduction of prophylactic vaccines that prevent transmission of specific high-risk HPVs. However, these vaccines have no therapeutic efficacy and since HPVassociated cervical cancers may arise years or even decades after the initial infection, it has been estimated that there will be no measurable decline of HPV-associated tumors before 2040. Hence, therapeutic efforts to combat high-risk HPV-associated disease remain of critical importance (Hellner and Munger 2011).

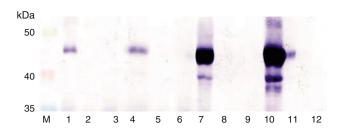


Fig. 5 Western blot analysis of fusion protein PVX CP-L15-E6GT expressed in *N. benthamiana* plants. *M* M SpectraTM Multicolor Broad Range Protein Ladder (10–260 kDa), Fermentas; *1 N. b.* inoculated leaves; *2 N. b.* systemic leaves; *3 N. b.* roots; *4 N. b.* HC-Pro inoculated leaves; *5 N. b.* HC-Pro systemic leaves; *6 N. b.* HC-Pro roots; *7 N. b.* 3H inoculated leaves; *8 N. b.* 3H systemic leaves; *9 N. b.* 3H roots; *10 N. b.* K7 inoculated leaves; *11 N. b.* K7 systemic leaves; *12 N. b.* K7 roots. To detect the fusion protein the primary rabbit polyclonal anti-PVX CP antibodies were used

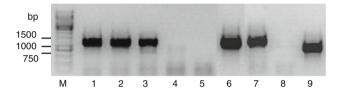


Fig. 6 Immunocapture RT-PCR analysis of fusion genes in inoculated *N. benthamiana* K7 leaves. *M* GeneRuler 1 kb DNA Ladder, Fermentas; *I* PVX CP-L15-E6GT; *2* PVX CP-E6GT; *3* PVX CP-L4-E6GT; *4* PVX CP; *5* healthy leaves; *6* E6GT-PVX CP; *7* E6GT-L4-PVX CP; *8* healthy leaves; *9* PVX CP. The E6GT (G9-O) and pGR106-specific primers were used to amplify 3'-terminal fusion genes (1–5). The pGR106-specific primers were used to detect 5'-terminal fusion genes (6–9)

High-risk human papillomavirus infections are generally transient and believed to be controlled by cell-mediated immune response because the majority of HPV-induced lesions spontaneously regress (Eiben et al. 2003). Nevertheless, a major feature of HPV that might impair HPV therapeutic vaccination is the poor presentation of viral antigens, which are expressed at a low level (Tindle 2002). As the virus cannot be propagated in culture, vaccines have been based on recombinant antigens with inherent highcost production. Therefore, in a search for alternative formulations able to improve the immunogenicity of HPV-16 E6 protein, we took advantage of our extensive experience with PVX based expression platform (Cerovska et al. 2004, 2008, 2012; Pokorná et al. 2005; Hoffmeisterova et al. 2008; Plchova et al. 2011) and used this system for the transient expression of chimeric PVX particles bearing mutagenized HPV-16 E6 in plants. Other studies have also demonstrated the advantage of PVX as an epitope presentation system in vaccination (Franconi et al. 2002; Marusic et al. 2001).

We successfully expressed all full-length C- and two N-terminal (no linker, L4) fusion proteins in *E. coli* MC1061 using the pMPM-A4 Ω plasmid.

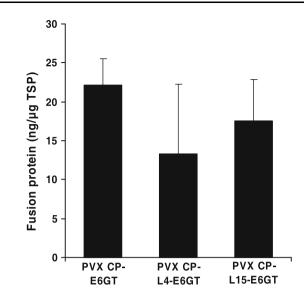


Fig. 7 Transient expression of three PVX CP C-terminus fusion proteins in *Nicotiana benthamiana* K7 plants. *N. benthamiana* K7 leaves were agroinfiltrated and the accumulation of each fusion protein was quantified by Western blot analysis using anti-E6 (N-17) antibodies (1:1,000 in PBST; Santa Cruz Biotechnology, Santa Cruz, CA, USA) and by rabbit anti-goat-AP secondary antibodies (1:30,000 in PBST; Sigma, St. Louis, MO, USA), where the bands corresponding to the respective constructs were compared with the dilution series of the PVX CP-L4-E6GT expressed in bacteria. Each *column* represents the mean value of 3 independent experiments. In each experiment 2 agroinfiltrated leaves per plant (18 plants in total) were collected. The standard deviation is represented by error bars. *TSP* total soluble protein

After optimization of expression conditions in *E. coli* high yield and proper serological reactivity of all constructs were observed. Both N-terminal and C-terminal fusion proteins accumulated to approximately the same levels. Also the length of the linker did not have substantial impact on yield of fusion proteins. Based on immunoelectron microscopy analysis with anti-PVX CP antibodies we can conclude that all fusion proteins were able to form VLPs.

On the contrary, only the C-terminal fusion proteins accumulated to detectable levels in plants. The highest yields of chimeric protein were observed in the transgenic *N. benthamiana* K7. When inoculated on host plants, these recombinant viruses were not able to spread systemically, the viral infection was restricted only to the inoculated leaves. All three constructs were expressed in plant tissue at approximately similar levels and there were no substantial differences when linkers of different lengths were used. All three fusion proteins expressed in plants formed VLPs with serologically accessible parts of PVX CP and E6GT protein.

The main advantage of transient-expression system in plants is the relative rapidity of the procedure, while the main constraint of this approach is generally a low stability and infectivity of the chimeric structures carrying exogenous sequences fused to the N-terminus of PVX CP (Uhde-Holzem et al. 2007). We have previously expressed E7GGG fused to both N- and C-terminus of PVX CP in *N. benthamiana* HC-Pro transgenic plants (Plchova et al. 2011). The chimeric coat proteins carrying E7GGG fusions were expressed in plant tissues at approximately the same concentrations and, similarly to E6 fusions, were not able to cause infection of upper systemic leaves. This observation correlated with inability of E7GGG to form viral particles probably due to the steric obstructions or change in the charge.

On the contrary, in this work, the attenuated E6GT HPV oncoprotein accumulated to detectable levels only when fused to the C-terminus of PVX CP. Again, the fusion proteins were not able to spread systemically in transgenic K7 *N. benthamiana*, even though all of them were able to form viral particles as evidenced by electron micrographs and IC-RT-PCR. The K7 plants carry both HC-Pro from PVA and MP from TMV. HC-Pro, which is known as a viral suppressor of RNA silencing is known to elevate the level of potexvirus expression while the TMV MP is known to complement the lack of cell-to-cell transport activity of wild type PVX CP (Fedorkin et al. 2001). Both these traits have likely contributed to obtain the highest expression in this host genotype.

We could only speculate why the chimeric proteins, that formed the particles, were unable to spread systemically. It is possible that the characteristics of the fusion with E6GT (e.g. fused sequence length and amino acid composition), and/or its location in the vector prevented the virus from interacting with some host factors, which are indispensable for longdistance movement or entry from phloem to cells.

In our future research we would like to address the instability of N- and C-terminal PVX-CP fusions. One of the possible strategies we would like to follow is to identify novel positions in the PVX-CP sequence allowing stable surface presentation of heterologous antigens. Also the comparison of immunogenicity of VLPs with that of peptide vaccines containing corresponding E6-derived epitopes is of great interest. Last but not least combined immunization with DNA vaccines and VLPs will also be tested, as this heterologous prime/boost strategy proved to be very efficient in other systems (Woodland 2004).

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