

Production of Human Papilloma Virus Type 16 E6 Oncoprotein as a Recombinant Protein in Eukaryotic Cells

Mirshahabi H¹, Soleimanjahi H¹, Pourpak Z², Meshkat Z¹, Hassan ZM¹

Abstract

Background: Cervical cancer is one of the most important and widespread cancer which affects women. There are several causes of cervical cancer; among them HPV types 16 and 18 are the most prominent ones which are recurrent and persistent infections. These genotypes are currently about 70% of cervical cancer causes in developing countries. Due to the importance of these viruses in cervical cancer, we pioneered the production of Human Papilloma Virus type 16 E6 oncoprotein as a recombinant protein in order to develop a vaccine. Two HPV oncoproteins, E6 and E7, are consistently expressed in HPV-associated cancer cells and are responsible for malignant transformation. These oncogenic proteins represent ideal target antigens for developing vaccine and immunotherapeutic strategies against HPV-associated neoplasm.

Methods: In the present study, the cloned E6-oncoprotein of HPV16 in pTZ57R/T-E6 vector was used to produce professional expression vector. The target gene was subcloned in a eukaryotic expression vector. The pcDNA3-E6 vector was propagated in E.coli strain DH5 α and transfected into CHO cells 72 hours post-transfection.

Results: The transfected cells were harvested; mRNA detection and the interest protein production were confirmed by western blot analysis using specific anti E6 monoclonal antibody.

Conclusion: HPV16-E6 target protein recognized by specific antibody could be an appropriate form of protein, which can be used for further studies. Due to potential effect of this protein, its DNA construction can be used for DNA vaccine in future studies.

Keywords: E6 protein; Human Papilloma Virus type 16; Uterine Cervical Neoplasms; Gene expression

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Introduction

There are approximately more than 100 different types of Human Papilloma Virus (HPV) [1,2], which can be further classified as low risk types, found primarily in benign lesions, and high risk types, found in malignant or potentially malignant tumors [3,4]. HPVs are intimately associated with development of cervical cancer [4], which is the second leading cause of cancer-related death among women worldwide [5, 6]. It was shown that in some region of Iran such as Gorgan, it is considered very important cancer as the genital cancer is the third malignancy [7]. More than 99% of cervical cancers contain DNA of HPV

[8], particularly the high-risk HPV type 16 [9]. About 80% of 500,000 new cases of cervical cancer occur each year in developing countries [5, 10].

The oncogenic activity of high-risk HPV E6 gene has been shown in tissue culture and transgenic mouse model systems extensively [1, 8]. The E6 oncoprotein plays an important role in HPV-associated cervical carcinoma and also in the Human Papilloma Virus life cycle [11]. The E6 is expressed through all stages of cervical neoplasia [12]. A novel and powerful strategy for anti-tumor vaccine is direct inoculation of plasmid DNA encoding tumor-associated antigens. This technique called DNA immunization, which is

known to induce both antigen-specific cellular and humoral immune response [13, 14]. DNA vaccine has become an appealing approach to generate antigen-specific immunotherapy as a result of simplicity, stability, safety, and capacity for repeated administration [9]. The major concern on HPV DNA vaccine is the possibility of DNA integration into the host genome [15]. Several studies have demonstrated in different animal models that DNA immunization with plasmid encoding HPV oncogenic proteins can prevent or delay development of invasive carcinoma [16].

The HPV E6 protein is a small protein containing of approximately 150 amino acids [1, 11], which degrades p53 tumor suppressor through an ubiquitin-dependent pathway, interfering with both cell cycle control and apoptotic pathway [17, 18]. In this study we showed construction, expression, and characterization of E6 protein isolated from HPV 16 which can be used for DNA vaccine.

Materials and Methods

Construction of expression vector

The E6 gene of HPV16 was isolated and cloned in pTZ57R/T-E6 to use for initial cloning of HPV16-E6 gene and sequencing [19]. Preparative digestion was done by Hind III (all enzymes were obtained from Roch, Germany) and BamH I sites of cloning

vector and then subcloned into eukaryotic expression vector pcDNA3 (Invitrogen, USA). Competent Ecoli cells were transformed using constructed pcDNA3-E6. The accuracy of constructed plasmid was confirmed by colony-PCR and restriction enzyme analysis. Using E6 forward

5'-TAATCGAAGCTTAAACTAAGGGCCT-3' and reverse

5'-GCAAAGGATCCATATATTCATGCAATGTAG-3' primers. The colonies were used directly as template in colony-PCR, using PCR mix (50 pmol each primer, 1.5 mM MgCl₂, 0.2 mM each dNTP, 1U Taq DNA polymerase (Cinagen, Iran) in a total reaction volume of 15 µl). Amplification was carried out for 35 cycles (94°C for 30s, 61°C for 30s, 72°C for 45s) after an initial denaturation at 94°C for 5 min, on a Techne Thermal Cycler. The cycles were followed by a 5 min extension at 72°C and PCR product was visualized by UV on a 1.5% agarose gel electrophoresis. Colonies containing expected plasmids were selected and purified after large scale preparation. Purified pcDNA3-E6 was confirmed by restriction enzyme analysis using Hind III and BamH I.

Cell culture

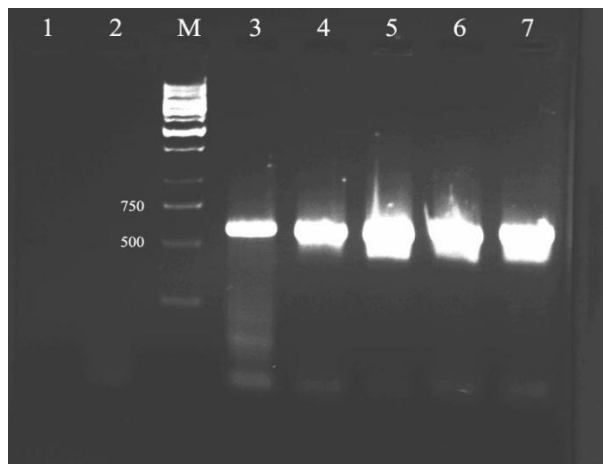


Figure 1. Shows agarose gel electrophoresis of colony PCR amplification products which stained with Ethidium Bromide and visualized under UV illumination. Lane 1: PCR negative control, lane 2: negative control plasmid (pcDNA3), lane M: DNA marker, lane 3: PCR positive control and lanes 4, 5, 6, 7: the colonies containing desired product of 600 bp belong to HPV16-E6 were confirmed.

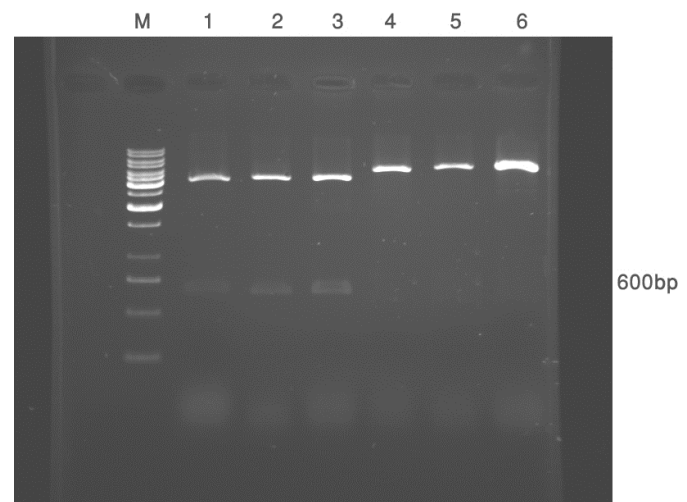


Figure 2. Shows digestion pattern of pcDNA3-E6 vector using BamH I and Hind III (lanes 1, 2 and 3) or HindIII (lanes 4, 5 and 6).

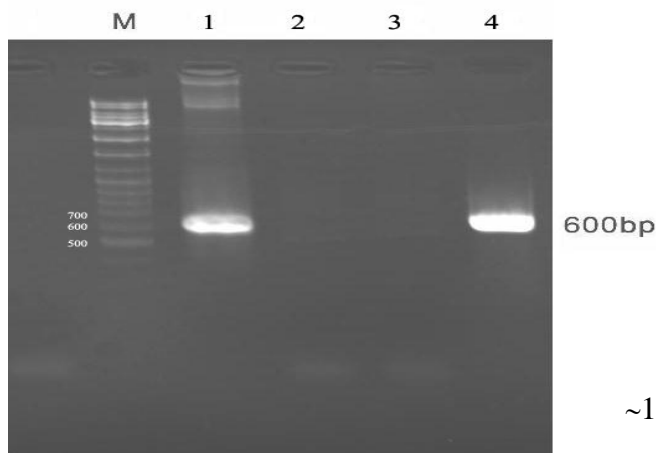


Figure 3. Shows confirmation of E6 mRNA production in transfected CHO cells. Lane M: DNA size marker, lane 1: PCR positive control, lane 2: PCR negative control, lane 3: untransfected cell, lane 4: transfected cell containing E6 mRNA which can be analyzed by RT-PCR.

CHO cells were cultured in RPMI 1640 supplemented with 10% Fetal Calf Serum (FCS), 2mM L-glutamine, 100 U/ml of penicillin, 100 µg/ml of streptomycin, and 1 mM sodium pyrovate. The cells incubated at 37°C in a humidified 5% CO₂ atmosphere.

Transfection by calcium phosphate precipitation

Expressing E6 protein, from the respective plasmid construct was tested in transient transfection assay. In a transient transfection experiment, two well of microplate untransfected cells were used as negative controls. To monitor expression of E6 protein, CHO cells were seeded into 6-well microtiter plates and incubated overnight in complete medium without any antibiotic. The confluent CHO cells were transfected with pcDNA3-E6 or pcDNA3 (as negative control) using calcium-phosphate protocol [20, 21]. Fourty eight hours after transfection, the cells were washed with PBS and then were analyzed by RT-PCR.

RNA extraction and Reverse Transcription (RT)-PCR

Total RNA were isolated from the cells using RNX-plus solution (Cinagen, Iran) according to manufacturer's instructions. Contaminated DNA was

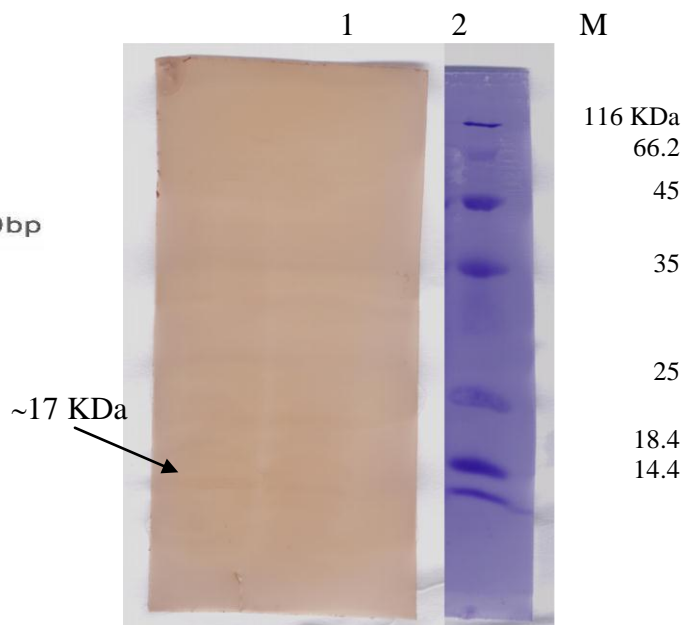


Figure 4. Shows western blot analysis of HPV-E6 gene using mouse monoclonal antibody. Lane 1: protein size marker, lane 2: transfected cell, lane 3: untransfected cells.

removed by DNase I (Fermentase, Germany) and extracted RNA was used in RT-PCR.

The first strand of cDNA was synthesized at 37°C for 60 min in presence of 1µg RNA and 200 U/MMLV reverse transcriptase (Fermentase, Germany), 20U RNase inhibitor, dNTP mix (final concentration of 1 mM) with oligo (dT) priming in a 20µl reaction. The PCR amplification of cDNA was carried out using E6 primers. The condition of PCR mix and programs were the same as previously described.

Western Blot analysis

For characterization of expressed HPV16-E6 protein, total cell lysates were prepared by sonication (60HZ, 0.5 amplitude) after 5 times freezing and thawing. The samples were lysed by addition of 2X sample buffer at 100°C for 5 min on 12% gel and stained with coomassie blue. The proteins were also transferred into nitrocellulose paper by electrophoresis for 1 hour at 90 V, blocked with 3% BSA in PBS for more than 1 h. The proper antigenicity of E6 protein was assessed using mouse monoclonal anti-E6 antibody (abcam, UK) and specific bands were visualized with Horse-Radish-Peroxidase (HRP)-conjugated anti-mouse immunoglobulin G antibody (TEBSUN Company, Iran)

and development was carried out using Diamino Benzidine (DAB) substrate (Biogene, Iran).

Results

Construction and expression of HPV 16-E6

The expression vector containing HPV16-E6 gene was constructed as described in Material and Methods. The accuracy of resulted plasmid was confirmed by colony-PCR using designed specific E6 primers (Figure 1).

The colonies containing desired plasmid were shown in lane 4, 5, 6 and 7 behind positive control in lane 3 on agarose gel and existence 600 bp bands belong to HPV16-E6.

The colonies which were positive in colony-PCR were subjected to restriction enzyme analysis using BamH I and Hind III (Figure 2).

In vitro evaluation

Plasmid pcDNA3-E6 was transfected in CHO cells using calcium phosphate. The expression of HPV16-E6 was analyzed by RT-PCR 48 hr post-transfection.

As shown in Figure 3, using E6 specific primers the 600 bp fragments was amplified on resulted cDNA of target produced mRNA on transfected cells.

Characterization of HPV16-E6 protein by western blotting

The transfected cells by pcDNA3-E6 plasmid were subjected to SDS-PAGE electrophoresis and transferred into a Polyvinylidene Fluoride (PVDF) membrane and confirmed by monoclonal antibody and visualized with 3,3'-diaminobenzidine tetrahydrochloride. The calculated molecular mass of HPV16-E6 gene was approximately 17 KDa which were appeared in transfected cell extract. The band is absent in untransfected or mock transfected cells (Figure 4).

Discussion

Considering the important role of Human Papilloma Virus in cervical cancer and high frequency genital infection of HPV, detecting high-risk genotype of HPV infection in high risk population is of paramount importance to prevent malignancy. As the incidence of cervical cancer continues to increase, the need for a safe and effective vaccine is more felt. Since the E6 and E7 proteins represent attractive target to develop anti-cancer vaccine, availability of HPV16-E6 constructions can greatly help for developing potent vaccines such as subunit or DNA vaccine [22].

In the present study, isolation of HPV 16 was subjected to isolation of E6 gene by PCR. The gene

of interest was subcloned in pcDNA3 expression vector. After confirmation, the CHO cells were transfected with pcDNA3-E6 and protein production was evaluated by RT-PCR, SDS-PAGE and western blotting. It was shown that there is always a minor expression of protein in transfected cells.

In optimal condition only 2.5% of total proteins in eukaryotic cells belong to recombinant protein. As shown by Rapp L, the E6 protein was presented at very low level in cells. In vitro analysis of E6 was very complicated [11]. In transfected CHO cell suspension E6 protein was detected as a faint band migrating approximately 17 KDa, according to its calculated molecular mass. The expression of E6 protein was done by using efficient expression vector and optimization of conditions. Previous attempts to express E6 protein have demonstrated that the level of expression was very low and immunoprecipitation was needed to detect the desired protein. These primary findings warrant further studies for DNA vaccine against HPV16.

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Conflict of Interest

The authors state that there is no conflict of interest.

Authors' Contribution

MH and MZ carried out the tests and helped in writing the paper. SH designed the study, supervised the project, interpreted the results, draft and revised the manuscript.

PZ and HZM contributed to the literature review. All authors read and approved the final manuscript.

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