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Naive and radiolabeled antibodies to E6 and E7 HPV-16 oncoproteins show pronounced antitumor activity in experimental cervical cancer

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Abstract

Background—In spite of profound reduction in incidence, cervical cancer claims >275,000 lives annually. Previously we demonstrated efficacy and safety of radioimmunotherapy directed at HPV16 E6 oncoprotein in experimental cervical cancer.

Materials & methods—We undertook a direct comparison of targeting E7 and E6 oncoproteins with specific ¹⁸⁸Rhenium-labeled monoclonal antibodies in CasKi subcutaneous xenografts of cervical cancer cells in mice.

Results—The most significant tumor inhibition was seen in radioimmunotherapy-treated mice, followed by the unlabeled monoclonal antibodies to E6 and E7. No hematological toxicity was observed. Immunohistochemistry suggests that the effect of unlabeled antibodies is C3 complement mediated.

Conclusion—We have demonstrated for the first time that radioimmunotherapy directed toward E7 oncoprotein inhibits experimental tumors growth, decreases E7 expression and may offer a novel approach to cervical cancer therapy.

Keywords

¹⁸⁸Rhenium; apoptosis; complement cytotoxicity; E6 and E7 oncoproteins; HPV16 positive cervical cancer; p53 expression; radioimmunotherapy; retinoblastoma

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It has been well established and is universally accepted that cervical cancer is caused by Human Papilloma Virus, HPV [1,2]. Molecular epidemiologists have been able to trace strains of HPV back to the prehistoric era noting conservation of the viral epitopes for over 200,000 years [3]. Throughout the course of time the preservation of the active oncogenic portions of the virus, the reliance of malignant transformation on early oncogenes expression has remained the same; namely of the *E6* and *E7* viral oncogenes [4]. Persistence of HPV infection leads to the malignant change of normal cervical epithelium to cancer. In concert with other factors such as loss of heterozygosity, *E6* oncogene deregulates the *p53* pathway, while *E7* de regulates retinoblastoma gene (*Rb*) [5]. In spite of the profound reduction in incidence of cervical cancer due to Papanicolaou screening and initiation of HPV vaccination, cervical cancer remains a pandemic with more than 85% of the global burden of disease occurring in developing countries and claims more than 275,000 lives annually (WHO, 2011). In early stages of cervical cancer, surgery can be curative, however, when cancer is metastatic at the time of a patient presentation or is recurrent, treatment options are limited.

Our approach to address this limitation is the development of radioimmuno-therapy (RIT) to target the viral specific oncoproteins of HPV-induced cervical cancer by labeling monoclonal antibodies (mAbs) to *E6* or *E7* oncoproteins, the products of *E6* and *E7* expression, with a radionuclide which emits cytotoxic radiation, such as ¹⁸⁸Rhenium (¹⁸⁸Re) [6]. To access the intranuclearly located target *E6* and *E7* oncoproteins, this approach initially relies on tumor necrosis that occurs as the tumor outgrows its blood supply causing the release of intranuclear contents into the interstitial spaces where it becomes accessible to mAbs while further damage is being mediated via beta-emission of a radionuclide. Our previous studies focused on RIT targeting *E6* oncoprotein. We have demonstrated the ability of a mAb C1P5 which is specific for HPV-16 *E6* to target *E6* antigen in experimental cervical cancer models with both high and low HPV copy numbers [7,8]. We observed abrogation of xenografted cervical tumors in nude mice by ¹⁸⁸Re-labeled C1P5 mAb to *E6* and a surprising effect of tumor growth inhibition in the unlabeled C1P5 mAb treatment group.

The molecular arrangement of *E7* oncogene is intricately linked to another early gene *E2* in the HPV genome and although the relative expression in actual human tumors (and commercially available cell lines) is blunted compared with *E6* as shown in [9], the impact of the linkage might render targeting it with RIT equally efficacious. In our previous studies we detected *E7* expression by western blot analysis in CasKi, SiHa and HeLa human cervical cancer cell lines with the *E7* specific mAb TVG701Y as in [10], however, the potential of *E7* as an RIT target *in vivo* remains unexplored.

This report focuses on the direct comparison of efficacy of targeting *E7* and *E6* oncoproteins with specific mAbs labeled with ¹⁸⁸Re in CasKi subcutaneous xenografts of cervical cancer cells in nude mice. We hypothesized that the effects of RIT directed against *E7* oncoprotein will be comparable to those of RIT directed against *E6*. We also compared the effect of unlabeled mAbs to *E6* and *E7* on the tumors. To our knowledge, this is the first report on comparative targeting of *E6* and *E7* oncoproteins with specific mAbs for developing novel immunotherapy for cervical cancer.

Materials & methods

Cell lines, antibodies & reagents

The commercially available CasKi human cervical cancer cell line, expressing both E6 and E7 oncoproteins, was purchased from the American Type Culture Collection (VA, USA). Cells were grown in RPMI-1640 medium containing 10% FBS (Sigma) and 1% Penicillin-streptomycin solution (Sigma, penicillin 10,000 U and streptomycin 10 mg/ml) at 37°C in a 5% CO₂ incubator. Matrigel, used in development of tumors, was purchased from BD Biosciences (MD, USA). Murine mAbs C1P5 (IgG1) to HPV-16 E6 + HPV-18 E6 and TVG-701Y (IgG2a to HPV-16 E7) were procured from Abcam.

Radiolabeling of antibodies

The beta-emitter ¹⁸⁸Re (half-life, 16.9 h) was produced from beta decay of a parent radionuclide ¹⁸⁸W (half-life 69 days) using a ¹⁸⁸W/¹⁸⁸Re generator (ITG Isotope Technologies Garching GmbH, Germany). After ¹⁸⁸Re was eluted in the form of sodium perrhenate, the antibodies were labeled with ¹⁸⁸Re ‘directly’ through binding of reduced ¹⁸⁸Re to the generated sulfhydryl groups on the antibodies, as described previously [11]. The radiolabeling yields were measured by instant thin layer chromatography by developing silica gel (SG) 10 cm strips in saline. In this system the ¹⁸⁸Re-labeled antibodies stay at the point of application while free ¹⁸⁸Re moves with the solvent front. The typical radiolabeling yields for both C1P5 and TVG-701Y mAbs were 85%. The radiolabeled mAbs were purified by HPLC using TosoHaas size exclusion column with PBS at 1 ml/min as an eluent using Waters HPLC system equipped with UV and radiation (Bioscan) flow-through detectors. The stability of the ¹⁸⁸Re-radiolabel on the mAbs was determined by incubating the radiolabeled mAbs in mouse serum for 48 h (~3 physical half-lives for ¹⁸⁸Re) at 37°C and analyzing the aliquots on the HPLC size exclusion column described above at 0, 1, 2, 4, 8, 24 and 48 h during the incubation in serum. No loss of ¹⁸⁸Re radiolabel from the mAbs was noted.

Tumor model

All animal studies were carried out in accordance with the guidelines of the Institute for Animal Studies at the Albert Einstein College of Medicine. Twenty-five six-week-old female athymic balb/c nude mice were purchased from Charles River Laboratories. 8×10^6 CasKi cells mixed with Matrigel were implanted into the flanks of mice in a subcutaneous fashion and allowed to grow to tumor size of 3–5 mm.

Biodistribution of ¹⁸⁸Re-C1P5 mAb in tumor-bearing mice & dosimetry calculations

For biodistribution the CasKi tumor-bearing mice were randomized into the groups of four mice per group and injected intraperitoneally (IP) with 50 µCi ¹⁸⁸Re-C1P5. The mice were sacrificed at the preset time points of 24, 48 and 72 h, their tumors and major organs removed, blotted from blood, weighed and counted for radio activity in the gamma counter, and percentage of injected dose per gram organ/tissue was calculated. These radioactive decay-corrected data were used to perform the radiation dosimetry calculations. Each mouse organ or tissue has a uniquely calculated absorbed fraction depending on geometry (and

density for tissue, bone and lungs) and complete energy spectrum of the applicable radionuclide. The absorbed fractions for ^{188}Re vary from 0.172 to 0.709 [12]. The effective clearance values for each organ and time point were obtained from the bio-distribution data, then fit each to a single exponential. The correlation coefficients are given in the last column of Table 1 under the heading 'r²'. The areas under curve were then integrated, multiplied by the equilibrium dose constant and absorbed fraction, and corrected for units. The equilibrium dose constant for ^{188}Re was obtained from the updated radionuclide emission data provided in [13]. The equilibrium dose constant was determined only for electrons (because only negligible photon radiation is absorbed by the small mouse organs). The results were expressed in cGy per unit μCi administered to a typical mouse.

RIT of CasKi tumor-bearing mice with ^{188}Re -labeled mAbs to E6 & E4 oncoproteins

CasKi tumor-bearing mice were randomized to five treatment groups of five mice per group: untreated controls, 300 μCi ^{188}Re -C1P5, 300 μCi ^{188}Re -TVG-701Y, unlabeled TVG-701Y and unlabeled-C1P5 groups. The total amount of an antibody per mouse in all groups was 10 μg . As patients with the metastatic cervical cancer have significant peritoneal disease, we administered the radiolabeled mAbs as IP injections to approximate the clinical situation. After IP injection administered as 200 μl volume, the mice were observed for 33 days and the tumors were measured in three dimensions over the observation period every three days. The volume of the tumors was calculated in the following fashion: Volume (initial) = $V_0 = (x * y * z)/2$. The subsequent measurements (V_2) were calculated as % change from initial volume $(V_2 - V_0)/V_0$. The experiment was performed two times.

Analyses of white blood cells & platelets

Baseline blood samples were taken from the tail vein of five non-tumor bearing mice and compared with the published data for balb/c athymic mice from Charles River Laboratories. Weekly blood samples were obtained in a random fashion from three mice per treatment group by ocular aspiration of blood. An aliquot of 1.5 μl of blood was added to 200 μl of ammonium oxalate solution (1 ml ammonium oxalate mixed with 3 ml of sterile water) and platelets were counted according to the published protocol [14]. White blood cells were counted according to standard protocol after diluting 10 μl of blood to 190 μl with 2% acetic acid solution.

Immunohistochemical analyses

Complement deposition was assessed using the murine mAb 6F64 to C3 (Santa Cruz Biotechnology) with the dilution of 1:1000. In addition, the tumors were stained for apoptosis (TUNEL), p53, Rb, p16 (BD Pharmingen) and CD3 (Dako). The immunohistochemistry (IHC) procedures were performed according to published protocols. The evaluation/scoring of the immunoreactivity was based on published criteria, and by two independent observers. TUNEL assay was performed according to the *in situ* cell death detection protocol (Roche). Retinoblastoma (Leica Biosystems) and p53 (Dako) assays were performed at a dilution of 1:20. For CD3 staining, the positive control tissue was tonsil using the CD3 antibody and the negative control tissue was tonsil with nonspecific rabbit IgG. For Rb the positive control tissue was tonsil using the Rb antibody and the negative control was

tonsil using nonspecific rabbit IgG. Positive control for p16 was metastatic squamous cell carcinoma; and positive control for p53 was endometrial intraepithelial serous carcinoma. The tumors were also stained and evaluated for the expression of E7 and E6 oncoproteins. C1P5 mAb was used for staining E6 oncoprotein at a concentration of 1:200 and a mAb to E7 oncoprotein (Cervimax, Germany) was used for E7 detection according to the Cervimax HPV E7 kit protocol.

Statistical analysis

The differences between the tumor sizes measured in the control and treatment groups were analyzed using Graphpad Prism Software two-way ANOVA. The differences were considered statistically significant when p-values were <0.05. The standard error of the repeated measurements of white blood cells and platelets was derived using Graphpad Prism Software.

Results

Treatment of CasKi derived tumors with TVG701Y mAb to E7 demonstrates tumor growth inhibition comparable to that achieved with C1P5 mAb

Mice bearing CasKi xenografts were treated with equal activities of the ¹⁸⁸Re-labeled mAbs TVG701Y and C1P5 to E7 and E6 oncoproteins, respectively, as well as the matching amounts of the unlabeled mAbs. Statistically significant tumor growth inhibition was observed in all treatment groups by 12 days status post-intraperitoneal injection compared with untreated controls (Figure 1A). In the untreated group, tumors grew aggressively and mice had to be sacrificed when tumors exceed 1 cm³ as per the Institute for Animal Studies protocol. Therefore the absolute number of mice of the control did not remain the same during the observation period. All mice otherwise survived the 33 day observation period. The experiment was performed two times. The most pronounced effect on the tumor growth was seen in mice treated with RIT dose of 300 µCi ¹⁸⁸Re-C1P5 or ¹⁸⁸Re-TVG701Y. There was not a significant difference with respect to tumor growth abrogation by the radiolabeled mAbs based on the target oncoprotein, E6 or E7. Unlabeled C1P5 and TVG701Y mAbs also inhibited tumor throughout the observation period. The effect of the unlabeled mAbs on the tumor, though less pronounced than that of the radiolabeled mAbs, was nevertheless statistically significant in comparison with the untreated controls (Figure 1A). Similarly to the radiolabeled mAbs, there was no difference between unlabeled C1P5 and TVG701Y mAbs effect on the tumors. This experiment not only demonstrated the potential of E7 oncoprotein as a target for RIT but also, exactly as observed previously with unlabeled C1P5 mAb in [7,8,10], revealed the inhibition of tumor growth by unlabeled TVG701Y mAb to E7. It should be noted that we did not use an irrelevant radiolabeled mAb as a control in this study because to be a proper control such mAb also has to bind to the intracellular antigen and as it was pointed out in [15] describing using a mAb to a different intracellular antigen – a single strand DNA in nonviable tumor cells – such radiolabeled mAb will also have some therapeutic effect on the tumor.

Dosimetry calculations predicted relatively low radiation doses to all major organs

The dosimetry calculations (Table 1) allowed to estimate that the dose to the tumor delivered by 300 μCi ^{188}Re -C1P5 mAb used in this study was approximately 300 cGy. The doses to all major organs were well below the maximum tolerated limits for these organs.

Antibody treatment results in limited toxicity to bone marrow

Weekly assessment of white blood cells (WBCs) and platelet did not demonstrate any depression of counts to the diagnosis of neutropenia or thrombocytopenia throughout the observation period nor were there significant differences in between treatment groups (Figure 1B & C). The evaluation of the WBCs showed that the main effect on WBCs was some decrease in counts in the untreated group while the counts in the treatment groups increased (Figure 1B), however, no infections were noted in any groups. There were no significant differences in platelet counts between the groups (Figure 1C), pointing to the lack of toxicity related to the bone marrow suppression with the 300 μCi doses of RIT.

Radiolabeled & unlabeled TVG701Y & C1P5 mAbs decreased the expression of E7 oncoprotein in the tumors

We evaluated the impact of RIT and unlabeled mAbs on the expression of E7 oncoprotein by IHC in the tumors procured at the completion of the *in vivo* experiments described above. Positive staining for E7 HPV oncoprotein is characterized by brown intranuclear uptake that should not be confused with nonspecific uptake in the cytoplasm (Figure 2). It can be seen that untreated CasKi cells show intranuclear staining for E7 (Figure 2A) which decreases in the tumors treated with ^{188}Re -TVG701Y (Figure 2B) or unlabeled TVG701Y (Figure 2D). Interestingly, treatment of CasKi tumors with ^{188}Re -C1P5 mAb also resulted in somewhat decreased staining for E7 oncoprotein (Figure 2C). This could be explained by the fact that RIT kills the tumor cells which express both E6 and E7 oncoproteins thus resulting in the decreased expression of either oncoprotein.

Radiolabeled TVG701Y & C1P5 mAbs increased the levels of apoptosis in the CasKi tumors

TUNEL analysis was performed on the untreated tumors (Figure 3A) and the results compared with those obtained from tumors treated with ^{188}Re -C1P5 or ^{188}Re -TVG701Y mAbs (Figure 3B & C). Increased number of apoptotic cells was detected in tumors treated with ^{188}Re -C1P5 or ^{188}Re -TVG701Y mAbs, however, the most pronounced response was noted in the ^{188}Re -C1P5 treatment group (Figure 3B). RIT is known to cause apoptotic cell death which is precipitated by the impact of charged particles on the nucleus and cellular machinery [16].

Active host immune response is seen in the tumors post-treatment with C1P5 & TVG701Y mAbs

We first sought to determine if there was an increase in C3 complement deposition in the tumors of mice treated with TVG701Y mAb. The significance of C3 complement deposition in the unlabeled C1P5 (Figure 4B) and TVG701Y (Figure 4C) mAbs treated tumors can be seen as compared with the untreated control (Figure 4A). To further investigate the host

immune response post-treatment with mAbs to E6 and E7, we stained the tumors from all groups with CD3-specific mAb to evaluate for the presence and proportion of tumor infiltrating lymphocytes. There was a distinct presence of tumor infiltrating lymphocytes in the tumors of treated mice with either C1P5 or TVG-701Y mAbs. Figure 5B shows the presence of CD3 positive cells in the tumors from the mice treated with unlabeled TVG701Y mAb. All tumors across the treatment groups showed diffuse and strong immunoreactivity for p16, supporting their high-risk HPV associated oncogenesis (Figure 5D). None of the tumors stained positive for p53 indicating the absence of protein expression at the IHC level (Figure 5F). There was no consistent pattern of Rb pathway expression in the treatment groups. Figure 5G displays the staining for Rb in the unlabeled TVG701Y mAb treatment group.

Discussion

In the era of personalized medicine, it is imperative that we look closely at the malignant portions of the well-preserved HPV oncoproteins and seek to develop less toxic, more specific treatment options for women world-wide who still succumb to this disease at an alarming rate. Cervical cancer is the only known curable gynecologic malignancy that remains pandemic in many areas in the world, in spite of the availability of a preventive vaccine and early treatment. RIT is currently clinically approved in the USA and Europe for treatment of B-cell lymphomas with ⁹⁰Y- and ¹³¹I-labeled CD20-binding mAbs and in China – for treatment of advanced lung cancer with ¹³¹I-labeled mAb to single strand DNA. However, all previous clinically approved and still experimental RIT applications target human ‘self’ antigens which are also expressed to certain extent on healthy tissues. Our approach of targeting virally associated antigens on tumors is unique as viral antigens have no homology to human antigens, thus allowing for much higher specificity of targeting and less toxicity of the treatment.

We have published previously on the ability of RIT against E6 oncoprotein to inhibit the growth of experimental cervical tumors [7,8,10]. However, in spite of the significance of these results, tumor growth after one intraperitoneal dose administration was maximally inhibited for about 18 days in all models with the subsequent regrowth of tumors while less rapid than that of control groups, still being observed. It was therefore logical to determine if targeting E7 oncoprotein would be comparable to these data or perhaps even be superior to the results obtained with ¹⁸⁸Re-C1P5 mAb to E6. Secondly, due to the rapid clearance from non-target organs as demonstrated in ¹⁸⁸Re-C1P5 biodistribution studies in [8], we anticipated that this treatment modality would also have limited toxicity to the bone marrow – non-significant neutropenia or thrombocytopenia as this often is the dose limiting toxicity in radiation-associated therapies. We also hypothesized that the unlabeled mAb TVG701Y to E7 would exert tumor growth inhibition as noted in previous studies with C1P5 mAb to E6. Finally, we studied the potential mechanisms of the unlabeled mAbs to determine if their action is in part related to the restoration of p53 and Rb, stimulation of complement mediated toxicity (CDC) and/or induction of apoptosis based on the actions of US FDA-approved unlabeled mAbs currently in clinical use [17].

During the 33-day observation period and serial measurements of tumor growth, we observed that the abundance of the target oncoproteins did not influence the effect on the tumor growth inhibition as the tumor suppressive effects of both ^{188}Re C1P5 and ^{188}Re -TVG701Y were not statistically significant from each other and no depression of WBCs or platelets was recorded during the observation period. Though 300 cGy dose delivered to the tumors by the radiolabeled mAbs was clearly much lower than the doses routinely delivered to the tumors by external beam radiation therapy, it resulted in significant retardation of the tumor growth as RIT works through different mechanisms than external beam radiation therapy [16,18].

We also observed the profound effects for unlabeled C1P5 and TVG701Y mAbs on the tumor growth. It is important to note that in any preparation of a radiolabeled mAb only one in 1000 to one in 50 mAb molecules are actually combined with a radionuclide, thus, the fact that unlabeled mAbs also mediate a profound effect on the tumor constitutes an important addition to the therapeutic profile of RIT.

To investigate the findings of unlabeled mAbs efficacy, we sought to gain insights into the antitumor activities of these mAb via IHC analyses of complement deposition, patterns of T-cell infiltration, intensity of E6 and E7 oncoprotein staining and presence of p53 and Rb. The classical pathway, triggered by antigen-bound antibody, alternative pathway and the mannose-binding lectin pathway all converge at C3 level which then stimulates subsequent steps that causes the assembly of the membrane attack complex as described in [19] which conceivably can attack the membranes of the nearby cells. We observed C3 complement deposition in the tumors treated with either C1P5 mAb to E6 or TVG701Y mAb to E7 as well as in our previous work when treating experimental melanoma tumors with unlabeled mAb to melanin which is also an intracellular antigen available for antibody binding only in nonviable tumor cells [20]. In this regard, an extensive profile was established via IHC analyses of complement regulatory proteins decay-accelerating factor CD55, membrane cofactor protein, CD46 and CD59 to investigate the comparative expression profiles in the spectrum of clinically relevant samples of normal human uterine cervix, premalignant and malignant disease [21]. The authors noted differential expression patterns in response to adjacent nests of tumor cells, stromal factors and hypothesized that these regulators may have functions to protect against complement. There is a complex interplay in the tumor between the necrosis and the complement as inflammation is a critical component of tumor progression. It has been demonstrated that although complement and TNF- α potentially exert significant antitumor effects, both mediators may also promote tumor progression [22]. For the purposes of this study aimed at demonstrating that E7 oncoprotein could also be an RIT target in cervical cancer we showed that there was much more complement deposition in both C1P5 and TVG-701Y mAbs treated tumors than in untreated tumor (Figure 4) which strongly suggests that the complement was induced by the mAbs rather than necrosis. It is much too early in our investigation to determine if mAbs to E6 and E7 reverse or inhibit some of the local tumor microenvironment effects, however, along with other investigators who have observed sporadic deposition of C3 on tumor cells and stroma prior to treatment with mAbs which subsequently increased twofold after treatment as shown in [23], we are planning further investigations along this avenue.

The finding of peripheral infiltration of T cells in the tumor is consistent with the published literature that demonstrates that T-cell infiltration in cervical cancer is oftentimes non-homogenous, with rare invasion in tumor maintaining either periphery or within stroma [24,25]. Therefore we did not expect to see a differential pattern of staining from the periphery of the tumor versus the center. These CD3 positive cells are most likely intestinal intraepithelial lymphocytes which have been shown to be present in significant amounts in athymic nude mice [26]. Our next steps are to perform double staining analysis to look at the ratios of CD4⁺ and CD8⁺ regulatory cells as well as procure tumors at various timepoints to determine if the profiles of regulatory cells are affected by treatment with RIT or unlabeled antibody [27]. Our current findings support the observations that the microenvironments of cervical cancer cells are somehow resisting the infiltration of the cells.

IHC for p53 showed that none of the tumors stained positively for this protein regardless of treatment group. We postulate that the differences in levels of p53 perhaps may not be noted via IHC and would be best analyzed in a quantitative fashion with real-time PCR as the increased deposition of apoptotic bodies noted in the TUNEL assay may correlate directly with radiation-induced damage versus that of antibody-dependent cytotoxicity as a result of the function inhibition of E6 and E7 oncoproteins. On the other hand, the absence of immunoreactivity for p53 may possibly suggest that there is a downstream pathway and perhaps incomplete restoration of the *p53* pathway for a sufficient specific protein expression that can be detected at the level of IHC. The absence of a consistent pattern in Rb expression can potentially be related to the actions of *E7* in destabilizing *Rb* by making it a target for proteasome mediated degradation, however, when the host cell recognizes the ongoing proliferation, apoptosis or senescence by *p53* activation is triggered.

The RIT approach will have the utility in cervical cancer as all patients are pretreated with chemotherapy such as cisplatin, and chemotherapy kills some cells in the tumors and thus creates the opportunity for the mAbs to E6 and E7 to reach their targets. In fact, in our previous studies we have shown that pretreatment of mice bearing HPV-16 positive cervical and head and neck tumors with cisplatin greatly enhances the efficacy of chemotherapy [28]. Furthermore, we anticipate that this treatment approach will be able to overcome some of the limitations of treatment with chemotherapy as a radiosensitizer. As tumors grow in size and become necrotic, the interstitial pressures noted at a cellular level oftentimes are such that the chemotherapeutic agent is not able to enter the cell, but rather bypass the tumor and is excreted without effect. Factors of better oxygenation and improved drug access from a lower interstitial pressure have best been studied in anal cancers (similar origin of HPV) using chemotherapy prior to definitive treatment or other small molecules to interrupt the natural history of the tumor prior to definitive management [29].

In summary, this report unequivocally demonstrates that the relative low expression of HPV E7 oncoprotein compared with E6 does not negatively affect its therapeutic potential seen with administration of RIT and of unlabeled mAb to E7. In fact, E7 therapeutic potential may indeed be synergistic or additive as we develop multimodal therapy approaches for future study. The limited toxicity to the bone marrow portends a favorable toxicity profile.

Conclusion

We performed direct comparison of targeting E7 and E6 oncoproteins with specific ^{188}Re -labeled mAbs in CasKi subcutaneous xenografts of cervical cancer cells in mice. The most significant tumor inhibition was seen in RIT-treated mice, followed by the unlabeled mAbs to E6 and E7. No hematological toxicity was observed. Immunohistochemistry suggests that the effect of unlabeled antibodies is C3 complement mediated. We have demonstrated for the first time that RIT directed toward E7 oncoprotein inhibits experimental tumors growth, decreases E7 expression and may offer a novel approach to cervical cancer therapy.

Future perspective

In future studies, we will evaluate for potential toxicity to the kidney and liver. Furthermore, combination therapy using both C1P5 and TVG701Y or serial injections in a once a week fashion may result in complete eradication of the tumors in future studies. By taking a closer look at the mechanisms of action of the unlabeled antibodies we are further intrigued by the findings and seek to develop further depths of understanding in order to magnify the effects in a clinically relevant manner. We anticipate that this treatment will be initially offered to patients who have recurrent or metastatic disease, a group who would have more commonly undergone chemotherapy with or without radiation therapy.

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Executive summary

- In spite of profound reduction in incidence, cervical cancer claims >275,000 lives annually.
- Previously we demonstrated efficacy and safety of radioimmunotherapy (RIT) directed at HPV16 E6 oncoprotein in experimental cervical cancer.
- Here we compare for the first time targeting E7 and E6 oncoproteins with specific ¹⁸⁸Rhenium-labeled monoclonal antibodies in CasKi subcutaneous xenografts of cervical cancer cells in mice.
- The most significant tumor inhibition was seen in RIT-treated mice, followed by the unlabeled monoclonal antibodies to E6 and E7. RIT reduced E7 expression in the tumors. No hematological toxicity was observed.
- Immunohistochemical evaluation of the tumors suggested active involvement of the immune system in the process of tumor growth abrogation with C3 complement deposition and lymphocytes infiltration. The levels of p53 and Rb were not affected by the treatment.
- We demonstrated for the first time that RIT directed toward E7 oncoprotein inhibits experimental tumor growth, decreases E7 expression and may offer a novel approach to cervical cancer therapy.

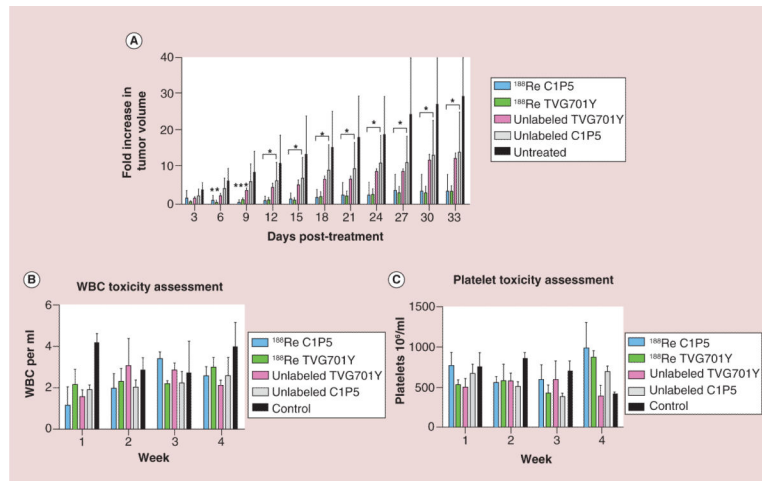


Figure 1. Radioimmunotherapy of CasKi tumors with $^{188}\text{Rhenium-C1P5}$ and $^{188}\text{Rhenium-TVG-701Y}$ monoclonal antibodies
(A) Average change in tumor volume ($V_2 - V_0/V_0$) over time among treatment groups was compared with untreated CasKi control tumors. **(B)** WBC counts during the observation period. X-axis shows 10^6 WBC per ml; **(C)** platelet counts during the observation period. * $p < 0.05$; ** $p < 0.005$; *** $p < 0.0005$.
 ^{188}Re : $^{188}\text{Rhenium}$; WBC: White blood cell.

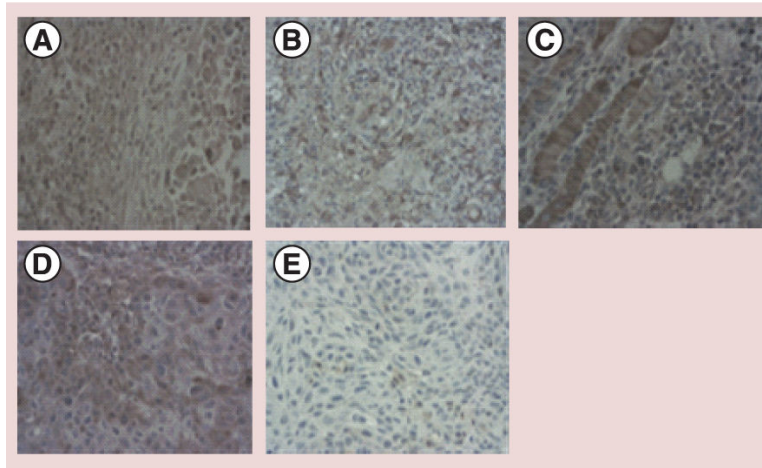


Figure 2. Immunohistochemical staining for E7 protein in CasKi tumors
(A) Untreated tumor; (B) treated with $^{188}\text{Rhenium-TVG-701Y}$ monoclonal antibodies (mAb); (C) treated with $^{188}\text{Rhenium-C1P5}$ mAb; (D) treated with unlabeled TVG701Y mAb; (E) non-HPV expressing, untreated control tumor. Focused brown staining indicates the presence of E7 protein.

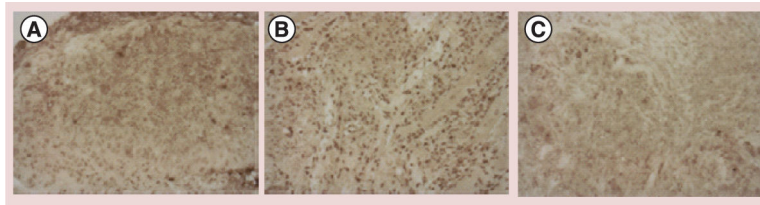


Figure 3. TUNEL assay for DNA fragmentation

(A) Untreated CasKi tumors; (B) treated with 300 μCi $^{188}\text{Rhenium-C1P5}$ monoclonal antibodies; (C) treated with 300 μCi $^{188}\text{Rhenium-TVG-701Y}$. Apoptotic cells stain brown.

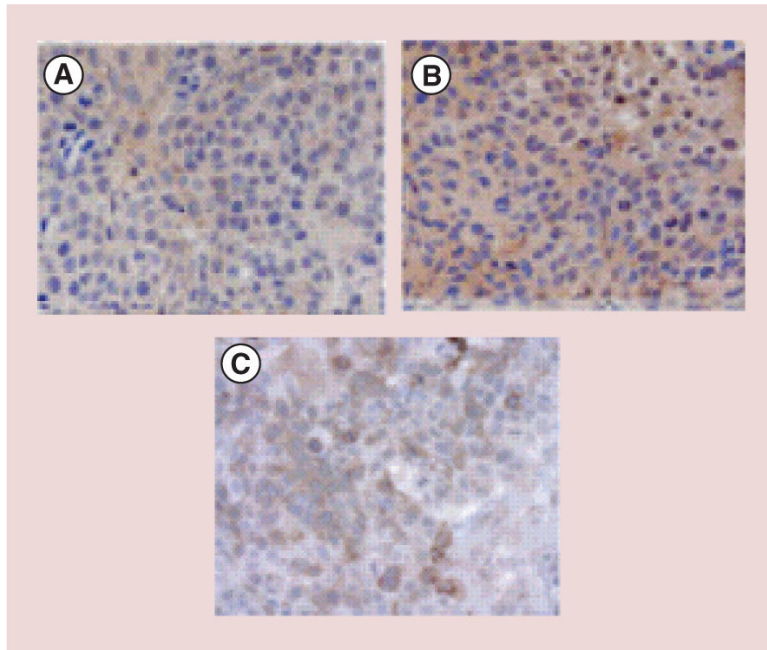


Figure 4. Immunohistochemical staining for C3 complement deposition
(A) Untreated CasKi tumors; (B) CasKi tumors treated with unlabeled C1P5 monoclonal antibodies; (C) CasKi tumors treated with unlabeled TVG701Y monoclonal antibodies. Brown staining indicates C3 complement presence.

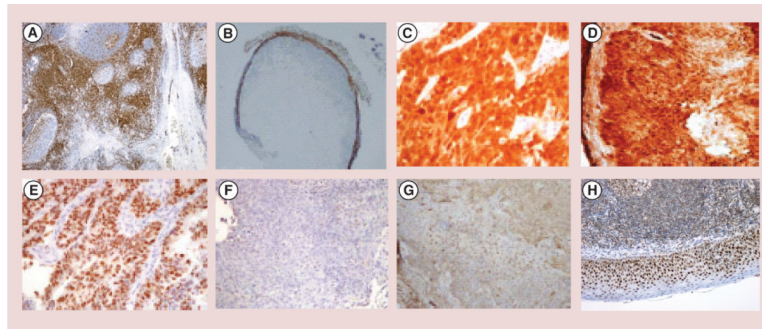


Figure 5. Immunohistochemical analyses of immune system interaction with the antibodies treated CasKi tumors

(A) Positive control for CD3 positive cells (tonsil); (B) lymphocytic invasion with CD3 positive cells of the tumors treated with unlabeled TVG701Y monoclonal antibodies (mAb) to E7. CD3 positive cells stain brown; (C) positive control stained for p16 (metastatic squamous cell carcinoma); (D) tumor treated with unlabeled C1P5 mAb stained positive for p16; (E) positive control for p53 (endometrial intraepithelial serous carcinoma); (F) tumor treated with unlabeled C1P5 mAb stained negative for p53; (G) tumor treated with unlabeled TVG701Y mAb stained positive for retinoblastoma; (H) positive control for retinoblastoma (tonsil).

Table 1

Dosimetry results for ^{188}Re Rhenium-C1P5 monoclonal antibodies administered to CasKi tumor-bearing mice.

Organ	cGy/ μCi	Label	r^2
Blood	5.292	Re-188	0.98
Lung	1.066	Re-188	0.99
Heart	1.633	Re-188	0.98
Spleen	0.903	Re-188	0.99
Liver	1.400	Re-188	0.98
Kidney	2.138	Re-188	0.99
Stomach	1.071	Re-188	0.99
Tumor	0.907	Re-188	0.99
Muscle	0.712	Re-188	0.96
Bone	0.698	Re-188	0.98

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