



Designing of Polyepitopic Vaccine by Targeting E7 Protein Sequences: An Immuno-Informatics Approach in Human Papillomavirus 16

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Abstract: Cervical cancer is the forth common type of cancer among women worldwide. Infection with high-risk human papillomavirus (HPVs) types is necessary for the development of cervical cancer and its precursors. HPV-16 is the most frequently known HPV type in cervical lesions. HPV-16 express E6 and E7 oncoproteins that require for the uncontrolled cellular proliferation. An efficient approach to treat lesions related to HPV infection is based on therapeutic vaccines against tumors induced by HPV. Unlike prophylactic vaccines based on the induction of antibodies, therapeutic anti-tumor vaccines need to induce cell mediated immune responses capable of identifying and eliminating abnormal cells. For this purpose, therapeutic vaccine should be efficiently stimulate innate and adaptive immune system, in particular CD 4⁺ and CD 8⁺ T cells. In the present study we report the design of a recombinant multiepitope protein containing immunogenic epitopes of HPV-16 E7. To design the vaccine, we assembled a database containing 1070 different sequences of HPV-16 E7 identified worldwide. We then predicted 20 poteintial HPV-16 E7 epitope using bioinformatics approaches. On the other, we exracted experimental epitopes from IEDB database. Using combination of predicted (theoretical) and experimental epitopes, we achived potent epitopic construct that can induce both CD 4⁺ and CD 8⁺ T cells efficiently. To increase the magnitude and quality of E7-specific immune response, linkers were inserted between epitopes (AAY used between CD 8+ epitopes and GPGPG used between CD 4+ epitope). Finally, we developed a therapeutic vaccine candidate employing a recombinant protein consisting of a string of multi-immunogenic T cell epitopes of E7. Although the study requires further in vitro and in vivo screening, it seems this epitope-focused peptide vaccine designing create a potent therapeutic vaccine that can develop alternative approach for the treatment of patients with this cancer.

Keywords: Human Papilloma virus; Therapeutic vaccine E7

References

- [1] L.Oliviera, M. Morale, "Design, Immune Responses and Anti-Tumor Potential of an HPV16 E6E7 Multi-Epitope Vaccine" Plos one, 21 (2015):e0138686.
- [2] S. Sabah, M. Gazi, "Designing of Epitope-Focused Vaccine by Targeting E6 and E7 Conserved Protein Sequences: An Immuno-Informatics Approach in Human Papillomavirus 58 Isolates" Interdiscip Sci. 2016, DOI 10.1007/s12539-016-0184-5
- [3] JT. Schiller, P. Davies, "Delivering on the promise: HPV vaccines and cervical cancer".2004, Nat Rev Microbiol. 2(2004):343-7.