

## **Product** Data Sheet

# HLA-A\*0201 HPV16 E7 complex Tetramer Protein, Human (YMLDLQPET, HEK293, His-Avi)

Cat. No.: HY-P700918

Synonyms: rHuHLA-A\*0201 HPV16 E7 complex Protein

Species: Human
Source: HEK293

Accession: P04439-1 (G25-T305)&P61769 (I21-M119)&YMLDLQPET

Gene ID: /&567

Molecular Weight: 260-265 kDa under Non reducing (N) condi

# Appearance Lyophilized powder. Formulation Lyophilized from a 0.22 μm filtered solution of PBS, pH 7.4. Normally 8% trehalose is added as protectant before lyophilization. Endotoxin Level <1 EU/μg, determined by LAL method. Reconsititution It is not recommended to reconstitute to a concentration less than 100 μg/mL in ddH<sub>2</sub>O. Storage & Stability Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage. Shipping Room temperature in continental US; may vary elsewhere.

## **DESCRIPTION**

### Background

The HLA-A\*0201 HPV16 E7 complex, a member of the antigen-presenting major histocompatibility complex class I (MHCI) family, plays a pivotal role in immune recognition and response. When associated with β2 microglobulin (B2M), it presents predominantly viral and tumor-derived peptides on the surface of antigen-presenting cells, facilitating recognition by CD8-positive T cells through their alpha-beta T cell receptors (TCR). This interaction guides an antigen-specific T cell immune response, crucial for eliminating infected or transformed cells. Beyond viral peptides, it may also present self-peptides originating from the signal sequence of secreted or membrane proteins. The complex displays a diverse repertoire of intracellular peptide antigens, typically 8 to 13 amino acids in length, processed through immunoproteasomes or insulindegrading enzyme. The binding of different peptides, defined by specific anchor residues, contributes to the fine specificity of antigen recognition. Notably, HLA-A\*0201, a representative allele, presents a restricted peptide repertoire, including viral epitopes from influenza and SARS-CoV-2, as well as tumor-associated antigens such as MAGE1, MAGEA3, and WT1. The complexity of peptide presentation highlights the crucial role of this MHCI complex in orchestrating immune responses against various pathogens and cancerous cells.

 $\label{lem:caution:Product} \textbf{Caution: Product has not been fully validated for medical applications. For research use only.}$ 

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