

Recombinant HAV/Hepatitis A Virus VP1-P2A (aa 722-830)

Catalog No.	CSI15704A	Quantity:	100 µg
	CSI15704B		0.5 mg
	CSI15704C		1.0 mg

Description: Forty-two antigenic domains were identified across the hepatitis A virus (HAV) polyprotein by using a set of 237 overlapping 20-mer synthetic peptides spanning the entire HAV polyprotein. Nineteen antigenic domains were found within the structural proteins, and 22 were found within the nonstructural proteins, with 1 domain spanning the junction of VP1 and P2A proteins. Five of these domains were considered immunodominant, as judged by both the breadth and the strength of their immunoreactivity. One domain is located within the VP2 protein at position 57-90 aa. A second domain, located at position 767-842 aa, contains the C-terminal part of the VP1 protein and the entire P2A protein. A third domain, located at position 1403-1456 a.a, comprises the C-terminal part of the P2C protein and the N-terminal half of the P3A protein. The fourth domain, located at position 1500-1519 a.a, includes almost the entire P3B and the last domain, located at position 1719-1764 aa, contains the C-terminal region of the P3C protein and the N-terminal region of the P3D protein. Four of the five most immunoreactive domains are derived from small HAV proteins and/or encompass protein cleavage sites separating different HAV proteins. The *E. Coli* derived 51.2 kDa recombinant protein contains the VP1-P2A immunodominant regions, amino acids 722-830

Source: *E. coli*

Molecular Weight: 51.2 kDa

Formulation: 10 mM CBB, pH9.6 + 0.1% SDS +50% glycerol.

Purity: HAV VP1-P2A protein is >90% pure as determined by 10% PAGE (coomassie staining).

Purification Method: Inclusion bodies.

Specificity: Immunoreactive with sera HAV-infected individuals.

Storage & Stability: HAV VP1-P2A protein although stable at 4°C for 1 week, should be stored below -18°C. **Please prevent freeze thaw cycles.**

Applications: HAV VP1-P2A antigen is suitable for ELISA and Western blots, excellent antigen for detection of HAV with minimal specificity problems.

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