

32-5616: Recombinant Epstein-Barr Virus (HHV-4) EBNA1, His Tag

Description

Source : The E.Coli derived recombinant protein contains the HHV-4 EBNA1 regions 408-641 amino acids having a molecular mass of 24.8kDa. The EBNA1 protein is fused to His Tag at N-terminus. The Epstein-Barr virus (EBV), also called Human herpes virus 4 (HHV-4), is a virus of the herpes family (which includes Herpes simplex virus and Cytomegalovirus. On infecting the B-lymphocyte, the linear virus genome circularizes and the virus subsequently persists within the cell as an episome. The virus can execute several distinct programs of gene expression which can be broadly categorized as being lytic cycle or latent cycle. The lytic cycle or productive infection results in staged expression of a host of viral proteins with the ultimate objective of producing infectious virions. Formally, this phase of infection does not inevitably lead to lysis of the host cell as EBV virions are produced by budding from the infected cell. The latent cycle (lysogenic) programs are those that do not result in production of virions. A very limited, distinct set of viral proteins are produced during latent cycle infection. These include Epstein-Barr nuclear antigen (EBNA)-1, EBNA-2, EBNA-3A, EBNA-3B, EBNA-3C, EBNA-leader protein (EBNA-LP) and latent membrane proteins (LMP)-1, LMP-2A and LMP-2B and the Epstein-Barr encoded RNAs (EBERs).

Product Info

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| Amount : | 0.1 mg |
| Purification : | Protein is >95% pure as determined by 10% PAGE (coomassie staining). |
| Content : | PBS lyophilized. It is recommended to add deionized water to prepare a working stock solution of approximately 0.5mg/ml and let the lyophilized pellet dissolve completely. |
| Storage condition : | EBV EBNA1 although stable at 4°C for 1 week, should be stored below -18°C. Please prevent freeze thaw cycles. |
| Amino Acid : | MKHHHHHPV GEADYFEYHQ EGGPDGEPDV PPGAIEQGPA DDPGEGPSTG PRGQGDGRR KKGWFGKHR GQGSNPKFE NIAEGLRALL ARSHVERTTD EGTWVAGVFV YGGSKTSLYN LRRGTALAIP QCRLTPLSRL PFGMAPGPGP QGPLRESIV CYFMVFLQTH IFAEVLKDAI KDLVMTKPAP TCNIRVTVCS FDDGVDLPPW FPPMVEGAAA EGDDGDDGDE GGDGDEGEEG QE |