

## HA (aa 17-529)(A/Aichi/2/1968)(H3N2)

CATALOG NUMBER: IA-H3-0043t, 50 µg

### Introduction

Influenza hemagglutinin (HA) is a type of hemagglutinin found on the surface of the influenza viruses. HA is an antigenic glycoprotein, like all other hemagglutinins, it causes red blood cells to agglutinate. HA is responsible for binding the virus to the cell that is being infected. HA proteins bind to cells with sialic acid on the membranes, such as cells in the upper respiratory tract or erythrocytes.

HA is a homotrimeric integral membrane glycoprotein. HA monomer is synthesized as a single polypeptide that is subsequently cleaved into two smaller polypeptides, the HA1 and HA2 subunits. Each HA monomer consists of a long, helical chain anchored in the membrane by HA2 and topped by a large HA1 globule.

### Description

Viral protein purified from 293 cell culture.

### Viral Protein

C-terminal 6xHis tagged HA (amino acid 17-529)(H3N2)(A/Aichi/2/1968)(GenBank No. AAA43178). A trimerization domain (Isoleucine Zipper) sequence has been introduced into the C-terminal of HA to stabilize the formation of trimer HA.

### Applications

Western blot standard, antibody ELISA, antigen, etc.

### Storage

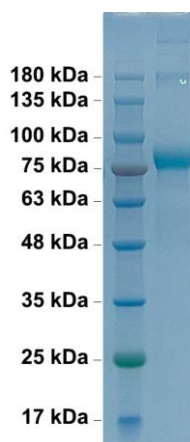
Store at -20 °C; Stable for 3 months from the date of shipment when kept at 4 °C. Non-hazardous.

### Concentration

1 µg/µl in PBS.

### Purity

>95% pure by 10% SDS PAGE gel.



**SDS-PAGE:** purified HA (H3N2) (A/Aichi/2/1968) protein with a trimerization domain

### HA (aa 17-529)(A/Aichi/2/1968/H3N2) SEQ:

QDLPGNNDSTATLCLGHHA VPNGTLVKTITDDQIEVTNATELVQSSSTGKICNNPHRILDGIDCTLIDALLGDPHCDVFNQNETWDLFVERSKAFSNCYP  
YDVPDYASLRSLVASSGTLEFITEGFTWTGVTQNGGSNACKRGP GSGFFSRLNWLTKSGSTYPVLNVTMPNNDNFDKLYIWGIHHPSTNQEQTSLYVQA  
SGRVTVSTRSQQTIIIPNIGSRPWVRLGSSRIS IYWTIVKPGDVLVINSNGNLIAPRGYFKMRTGKSSIMRSDAPIDTCISECITPNGSIPNDKPFQNV  
NKITYGACPKYVKQNTLKLATGMRNVPEKQTRGLFGA IAGFIENGWEGMIDGWYGFRHQNSEGTGQAADLKSTQA AIDQINGKLN RVIEKTNEKPFHQIE  
KEFSEVEGRIQDLEKYVEDTKIDLWSYNAELLVALENQHTIDLTDSEMKNLFEKTRRQLRENAEEMGNCGCFKIYHKCDNACIESIRNGTYDHDVYRDEA  
LNNRFQIKGVELKSGYKD