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Produktinformation



Forschungsprodukte & Biochemikalien



Zellkultur & Verbrauchsmaterial



Diagnostik & molekulare Diagnostik



Laborgeräte & Service

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Lieferung & Zahlungsart

siehe unsere [Liefer- und Versandbedingungen](#)

Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

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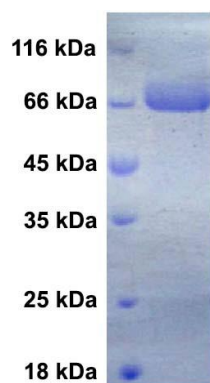
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HA (aa 16-548)(B/Massachusetts/02/2012)

CATALOG NUMBER: IA-BM7-005WP, 50 µg

Introduction	<p>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.</p> <p>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.</p>
Description	Viral protein purified from 293 cell culture
Viral Protein	C-terminal 6xHis tagged HA (B/Massachusetts/02/2012) protein (amino acid 16-548) (GenBank Accession No. AGL06036)
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. No MSDS required.
Endotoxin Level	<0.01 EU per 1 µg of the protein by LAL test
Concentration	1 µg/µl in PBS
Purity	>95% pure by 10% SDS PAGE gel



SDS-PAGE: purified HA (B/Massachusetts/02/2012) protein

Reference:

1. COMMITTEE ON INFECTIOUS DISEASES. Recommendations for prevention and control of influenza in children, 2014-2015. Pediatrics. 134:e1503-1519, 2014.

HA (aa 16-548)(B/Massachusetts/02/2012) SEQ:

DRICTGITSSNSPHVVKTATQGEVNVTVGVIPLTTTPTKSYFANLKGTKTRGKLCPDCLNCTDLVALGRPMC VGTTPSAKASILHEVRP
VTSGCFPIIMHDRTKIRQLANLLRGYENIRLSTQNVIDAEKAPGGPYRLGTSGSCP NATSKSGFFATMAWAVPKDNNKNATNPLTVEVPY
ICAEGEDQITVWGFHSDDKTQMKNLYGDSNPQKFTSSANGVTTHYVSQIGGFDPQTEDGGLPQSGRIVVDYMMQKPGKTGTIVYQRGVL
LPQKVCASGRSKVIKGSPLIGIADCLHEKYGGLNKSHPYYTGEHAKAIGNCPIWVKTPLKLANGTKYRPPAKLLKERGFFGAIAGFL
EGGWEGMIAGWHGYTSHGAHGVAADLKSTQEAINKITKNLNSLSELEVKNLQRLSGAMDELHNEIILELDEKVDLDRADTISSQIELA
VLLSNEGIINSEDEHLLALERKLLKMLGPSAVDIGNGCFETKHKCNQTCLDRIAAGTFNAGEFSLPTFDSL NITAASLNDGGLDNHTI**H**
HHHHH