

N2 Neuraminidase (NA) Protein with N-Terminal Histidine Tag from Influenza Virus, A/turkey/Wisconsin/1/1966 (H9N2), Recombinant from Baculovirus

Catalog No. NR-43783

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Contributor and Manufacturer:

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Product Description:

A recombinant form of the N2 neuraminidase (NA) protein from influenza A virus, A/turkey/Wisconsin/1/1966 (H9N2) containing an N-terminal histidine tag was produced in Sf9 insect cells using a baculovirus expression vector system and purified by nickel affinity chromatography. The predicted ectodomain coding region of the NA gene was fused to a synthetic gene segment encoding an N-terminal octa-histidine tag followed by a 43 amino acid tetramerization domain from vasodilator-stimulated phosphoprotein (VASP) and a thrombin cleavage site, as described for the 1918 pandemic virus.^{1,2} The predicted protein sequence is shown in Figure 1. NR-43783 has a theoretical molecular weight of approximately 51.22 kilodaltons. The crystal structure of the 1918 human N2 NA precursor has been solved at 2.40 Å resolution (PDB: [2HT8](#)). The full-length NA precursor protein is 469 residues (GenPept: [CAE00569](#)).

Material Provided:

Each vial contains approximately 150 to 200 µg of purified recombinant NA protein in phosphate buffered saline PBS (pH 7.4). The protein content in micrograms and the concentration, expressed as micrograms per milliliter, are shown on the Certificate of Analysis.

Packaging/Storage:

Purified recombinant NA protein was packaged aseptically in screw-capped plastic cryovials. This product shipped on dry ice and should be stored at -20°C immediately upon arrival.

Functional Activity:

NR-43783 was demonstrated to be functionally active based on its ability to cleave the fluorogenic substrate 2'-(4-methylumbelliferyl)-α-D-N-acetylneuraminic acid (4-MUNANA).³

Citation:

Acknowledgment for publications should read "The following reagent was obtained through BEI Resources, NIAID, NIH: N2 Neuraminidase (NA) Protein with N-Terminal Histidine Tag from Influenza Virus, A/turkey/Wisconsin/1/1966 (H9N2), Recombinant from Baculovirus, NR-43783."

Biosafety Level: 1

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the following publication: U.S. Department of Health and Human Services, Public Health Service, Centers for Disease Control and Prevention, and National Institutes of Health. Biosafety in Microbiological and Biomedical Laboratories. 6th ed. Washington, DC: U.S. Government Printing Office, 2020; see www.cdc.gov/biosafety/publications/bmbl5/index.htm.

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References:

1. Kühnel, K., et al. "The VASP Tetramerization Domain is a Right-Handed Coiled Coil Based on a 15-Residue Repeat." *Proc. Natl. Acad. Sci. USA* 101 (2004): 17027-17032. PubMed: 15569942.
2. Xu, X., et al. "Structural Characterization of the 1918 Influenza Virus H1N1 Neuraminidase." *J. Virol.* 82 (2008): 10493-10501. PubMed: 18715929.
3. Wetherall, N. T., et al. "Evaluation of Neuraminidase Enzyme Assays Using Different Substrates to Measure Susceptibility of Influenza Virus Clinical Isolates to Neuraminidase Inhibitors: Report of the Neuraminidase

Inhibitor Susceptibility Network." J. Clin. Microbiol. 41 (2003): 742-750. PubMed: 12574276.

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Figure 1: Predicted Protein Sequence

1	ADPHHHHHH	HSSSDYSDLQ	RVKQELLEEV	KKELQKVKEE	IIEAFVQELR
51	KRGSLVPRGS	PSRSEFEVCP	EVAEYRNWSK	PQCQITGFAP	FSKDNSVRLS
101	AGGDIWVTRE	PYVSCDPGKC	YQFALGQGT	LDNKHSNGTI	HDRIPHRTLL
151	MNELGVPFHL	GTKQVCIAWS	SSSCHDGKAW	LHVCVTGDDR	NATASFIYDG
201	MLVDSIGSWS	QNILRTQESE	CVCINGTCTV	VMTDGSASGN	ADTRVLFIRE
251	GKIIHISPLS	GSAQHIEECS	CYPRYPDVRC	VCRDNWKGSN	RPVIDIKMAD
301	YSINSGYVCS	GLVGDTPRSD	DSSSNSNCRD	PNNERGNPGV	KGWAFDNGDD
351	VWMGRTISKD	SRSGYETFRV	IGGWTTPN SK	SQVNRQVIVD	SNNWSGYSGI
401	FSVEGKSCIN	RCFYVELIRG	RPQETRVWWT	SNSIVVFCGT	SGTYGTGSWP
451	DGANINFMPI				

Plasmid-derived amino acids – Residues 1 to 3 and 61 to 66

His Tag – Residues 4 to 11

Tetramerization domain – Residues 12 to 54

Thrombin cleavage sequence – Residues 55 to 60

NA protein – Residues 67 to 460 [represents amino acid residues 76 to 469 of the native NA protein (GenPept: [CAE00569](#))]