

## **PRODUCT INFORMATION**

Tag C-Flag Tag

Target ACHB3

Synonyms N/A

**Description**Human ACHB3 full length protein-synthetic

nanodisc

Delivery 6~8weeks

Uniprot ID Q05901

Expression Host HEK293

**Protein Families** Ion Channels: Cys-loop Receptors

Protein Pathways N/A

Formulation &

Reconstitution

Storage & Shipping

Background

Molecular Weight

The human full length ACHB3 protein has a MW of

52.7kDa

Lyophilized from nanodisc solubilization buffer (20 mM Tris-HCl, 150 mM NaCl, pH 8.0). Normally 5% – 8% trehalose is added as protectants before lyophilization. Please see Certificate of Analysis for specific instructions. Do not use solvents with

for specific instructions. Do not use solvents with a pH below 6.5 or those containing high concentrations of divalent metal ions (greater than 5 mM) in subsequent experiments. Store at -20°C to -80°C for 12 months in lyophilized form. After reconstitution, if not

intended form. After reconstitution, if not intended for use within a month, aliquot and store at -80°C (Avoid repeated freezing and thawing). Lyophilized proteins are shipped at ambient

temperature.

The nicotinic acetylcholine receptors (nAChRs) are members of a superfamily of ligand-gated ion channels that mediate fast signal transmission at synapses. The nAChRs are (hetero)pentamers composed of homologous subunits. The subunits that make up the muscle and neuronal forms of nAChRs are encoded by separate genes and have different primary structure. There are several subtypes of neuronal nAChRs that vary based on which homologous subunits are arranged around the central channel. They are classified as alphasubunits if, like muscle alpha-1 (MIM 100690), they have a pair of adjacent cysteines as part of

the presumed acetylcholine binding site. Subunits lacking these cysteine residues are classified as beta-subunits (Groot Kormelink and Luyten, 1997 [PubMed 9009220]). Elliott et al. (1996) [PubMed 8906617] stated that the proposed structure for each subunit is a conserved N-terminal extracellular domain followed by 3 conserved

transmembrane domains, a variable cytoplasmic loop, a fourth conserved transmembrane domain, and a short C-terminal extracellular

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region.[supplied by OMIM, Apr 2010]

Usage Research use only
Conjugate Unconjugated

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