

DESCRIPTION

Source Chinese Hamster Ovary cell line, CHO-derived human MAN1A1 protein
Pro63-Glu653, with a C-terminal 6-His tag
Accession # P33908.3

N-terminal Sequence Analysis Pro63

Predicted Molecular Mass 67.7 kDa

SPECIFICATIONS

SDS-PAGE 63-65 kDa, under reducing conditions.

Activity Measured by its ability to remove α -mannose from the high mannose glycan Man-9.
A distinct band is observed in the rhMAN1A1 digested sample on SDS-PAGE gel, as measured under the described conditions.

Endotoxin Level <1.0 EU per 1 μ g of the protein by the LAL method.

Purity >95%, by SDS-PAGE visualized with Silver Staining and quantitative densitometry by Coomassie® Blue Staining.

Formulation Supplied as a 0.2 μ m filtered solution in Tris, NaCl and CaCl₂. See Certificate of Analysis for details.

Activity Assay Protocol

- Materials**
- Digestion Buffer: 50 mM MES, 15 mM CaCl₂, 1 mg/mL BSA, pH 6.0
 - Labeling Buffer: 25 mM Tris, 150 mM NaCl, 10 mM MnCl₂, pH 7.5
 - Recombinant Human MAN1A1 (rhMAN-1A1) (Catalog # 10665-GH)
 - Oligomannose-9 (Man-9) (Dextra Laboratories, Catalog # MC1131), 0.1 mg/mL stock in deionized water
 - Recombinant Human MGAT1 (rhMGAT1) (Catalog # 8334-GT)
 - UDP-GlcNAc (Sigma, Catalog # U4375), 50 mM stock in 50% ethanol, 50% deionized water
 - Recombinant Human FUT8 (rhFUT8) (Catalog # 5768-GT)
 - GDP-Cy5-Fucose (Catalog # ES301)
 - 15% SDS-PAGE gel
 - Reducing SDS-PAGE gel loading buffer
 - Fluorescent imager

- Assay**
- Digestion:**
1. Dilute rhMAN1A1 to 20 μ g/mL in Digestion buffer.
 2. Dilute Man-9 to 20 μ g/mL in Digestion Buffer.
 3. Combine 5 μ L of 20 μ g/mL Man-9, 5 μ L of 20 μ g/mL rhMAN1A1 and 10 μ L of Digestion Buffer. Include a Control containing 5 μ L of 20 μ g/mL Man-9 and 15 μ L of Digestion Buffer.
 4. Incubate at 37 °C for 2 hours.

- Labeling:**
1. Dilute rhMGAT1 to 100 μ g/mL in Labeling Buffer.
 2. Dilute UDP-GlcNAc to 1 mM in Labeling Buffer.
 3. Dilute rhFUT8 to 100 μ g/mL in Labeling Buffer.
 4. Dilute GDP-Cy5-Fucose to 0.05 mM in Labeling Buffer.
 5. Transfer 10 μ L of each digestion to a new tube and add 5 μ L of 100 μ g/mL rhMGAT1, 5 μ L of 1 mM UDP-GlcNAc, 5 μ L of 100 μ g/mL rhFUT8 and 5 μ L of 0.05 mM GDP-Cy5-Fucose.
 6. Incubate at 37 °C for 60 minutes.
 7. Add 6 μ L of Reducing SDS-PAGE gel loading buffer to each reaction.
 8. Load 12 μ L of each reaction onto a 15% SDS-PAGE gel and perform electrophoresis.
 9. Analyze gel on a fluorescent imager.

- Final Assay Conditions**
- Per Reaction:
- rhMAN1A1: 0.05 μ g
 - Man-9: 0.05 μ g
 - rhMGAT1: 0.5 μ g
 - UDP-GlcNAc: 5 nmol
 - rhFUT8: 0.5 μ g
 - GDP-Cy5-Fucose: 0.25 nmol

PREPARATION AND STORAGE

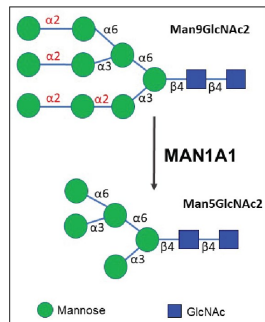
Shipping The product is shipped with polar packs. Upon receipt, store it immediately at the temperature recommended below.

- Stability & Storage** Use a manual defrost freezer and avoid repeated freeze-thaw cycles.
- 6 months from date of receipt, -20 to -70 °C as supplied.
 - 3 months, -20 to -70 °C under sterile conditions after opening.

DATA

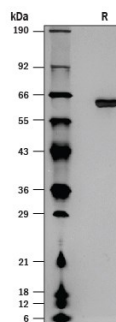
Rev. 11/2/2022 Page 1 of 2

Enzyme Activity



Recombinant Human MAN1A1 His-tag Protein Enzyme Activity Diagram. MAN1A1 is involved in the removal of 4 distinct α -1,2-linked mannose residues from Man9GlcNAc2 to produce Man5GlcNAc2, an essential step of N-glycan maturation.

SDS-PAGE



Recombinant Human MAN1A1 His-tag Protein SDS-PAGE. 1 μ g/lane of Recombinant Human MAN1A1 His-tag (Catalog # 10665-GT) was resolved with SDS-PAGE under reducing (R) conditions and visualized by silver staining, showing a band at 63-65 kDa.

BACKGROUND

N-glycan maturation in Golgi apparatus starts with high-mannose glycan Man-9 that is capped with four 1,2- α -linked mannose residues at its non-reducing ends. During the process, these mannose residues are removed to generate Man-5 oligomannose glycan, a precursor for complex and hybrid N-glycans (1). Failure of removing these mannose residues will result in the display of high-mannose glycans on cell surface and extracellular matrix. Increased levels of high-mannose glycans on cell surface are usually associated with disease progress such as tumorigenesis and viral infection (2). The removal of 1,2- α -linked mannose residues are catalyzed by 4 α -mannosidases, including MAN1A1, MAN1B1, MAN1A2 and MAN1C1, that have overlapping substrate specificity and slight differences in enzyme activity (3). MAN1A1 is also a tumor-suppressor (4) and low levels of expression of MAN1A1 correlate with poor prognosis in breast cancer patients (5, 6).

References:

1. Oliveira-Ferrer, L. *et al.* (2014) Br. J. Cancer **110**:753.
2. Moremen K.W. *et al.* (2012) Nat. Rev. Mol. Cell Biol. **13**:448.
3. Moremen, K.W. and Nairn, A.V. (2014) Handbook of Glycosyltransferases and Related Genes p1297.
4. Liu, T. *et al.* (2014) PLoS One **9**:e107941.
5. Milde-Langosch, K. *et al.* (2014) Breast Cancer Res. Treat. **145**:295.
6. Karen Legler, *et al.* (2018) Br. J. Cancer **118**:847.