

Recombinant Human SIAE His-tag

Catalog Number: 11497-SI

Diotecnne
RDSYSTEMS

DESCRIPTION		
Source	Chinese Hamster Ovary cell line, CHO-derived human SIAE protein lle24-Lys523, with a C-terminal 6-His tag Accession # Q9HAT2	
N-terminal Sequence Analysis	lle24	
Predicted Molecular Mass	57 kD	

SPECIFICATIONS	
SDS-PAGE	63-69 & 88-97 kD, under reducing conditions.
Activity	Measured by its ability to hydrolyze p-nitrophenylacetate. The specific activity is >25000 pmol/min/µg, as measured under the described conditions.
Endotoxin Level	<0.10 EU per 1 µg 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 µm filtered solution in Tris and NaCl. See Certificate of Analysis for details.

Activity Assay Protocol Materials Assay Buffer: 50 mM Tris, pH 8.0 Recombinant Human SIAE His tag (rhSIAE) (Catalog # 11497-SI) Substrate: 4-Nitrophenyl Acetate, 100 mM stock in acetone Clear 96 Well Plate Plate Reader with Absorbance Read Capability Assay 1. Dilute rhSIAE to 1.2 $\mu g/mL$ in Assay Buffer. 2. Dilute Substrate to 8 mM in Assay Buffer. 3. Load into a plate 50 µL of 1.2 µg/mL rhSIAE, and start the reaction by adding 50 µL of 8 mM Substrate. For Substrate Blank, load 50 µL of Assay Buffer and 50 μL of 8 mM Substrate. 4. Read plate at 410 nm in kinetic mode for 5 minutes. 5. Calculate specific activity: Adjusted V_{max}^* (OD/min) x well volume (L) x 10^{12} pmol/mol Specific Activity (pmol/min/µg) = ext. coeff** (M⁻¹cm⁻¹) x path corr.*** (cm) x amount of enzyme (µg) *Adjusted for Substrate Blank **Using the extinction coefficient 18500 M⁻¹cm⁻¹ ***Using the path correction 0.32 cm Note: the output of many spectrophotometers is in mOD Final Assay Per Well: Conditions rhSIAE: 0.06 µg

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.	
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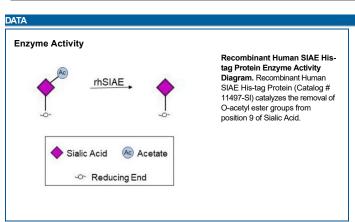
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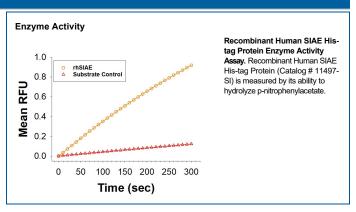
Substrate: 4 mM

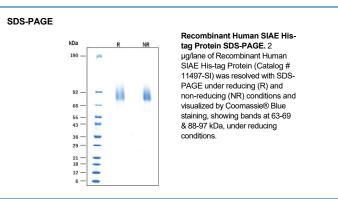


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BACKGROUND

Recombinant human Sialate O-acetylesterase (SIAE) is a glycosylated lysosomal protein that has two transcript isoforms: the cytosolic sialic acid esterase (Cse) that lacks 35 amino acids at the N-terminus and the lysosomal/membrane-associated sialic acid esterase (Lse) (1, 2). The Lse form is the predominantly expressed form responsible for the acetyl esterase activity on NeuAc in most tissues (2) and represents the recombinant product form. SIAE contains a signal sequence, a variable N-terminus, and a core protein structure that is unique within the SGNH hydrolase family (2) and contains a shallow active site pocket that accommodates binding of various complex substrates with acetylated sialic acid. SIAE Lse form catalyzes the removal of O-acetyl ester groups from position 9 of Sialic acid and is highly expressed in the testis, colon, and prostrate (3). SIAE deficiency has been shown to impact B cell development, signaling, and immunological tolerance through modulation of Siglec binding (4). Many SIAE variants in humans that result in defective activity or improper secretion have been identified in patients with autoimmune diseases including Crohn's disease, Sjogren's syndrome, lupus, rheumatoid arthritis, and type I diabetes (5). In addition, SIAE modulation of Siglec binding and sialic acid levels may play a key role in the immune evasion pathway associated with cancer (6). SIAE has also been implicated to play an important role in immune modulation during pregnancy (7).

References:

- 1. Higa, H.H. et. al. (1987) Biochem. Biophys. Res. Commun. 144:1099
- 2. Orizio, F. et. al. (2015) Glycobiology 9:992.
- 3. Zhu, H. et. al. (2004) J. Biomed. Biotechnol. 3:130.
- 4. Cariappa, A. et. al. (2009) J. Exp. Med. 206:125.
- 5. Surolia, I. et. al. (2010) Nature 466:243.
- 6. Grabenstein, S. et. al. (2021) Glycobiology **31**:1279.
- 7. Erickson, J. et. al. (2022) Nature 606:769.