

Endoglycosidase H (Endo H)

Cat no. LDG0002RG

Product Overview

Package component

Specification	Item	Amount
10,000 U	Endoglycosidase H (Endo H)	1 vial (500 U/ μ L)
	10X Glycoprotein Denaturing Buffer	1 vial (1 mL)
	10X Reaction buffer	1 vial (1 mL)
50,000 U	Endoglycosidase H (Endo H)	1 vial (500 U/ μ L)
	10X Glycoprotein Denaturing Buffer	1 vial (1 mL)
	10X Reaction buffer	1 vial (1 mL)

Description

Protein glycosylation is a complex posttranslational modification that manipulates biological activity such as protein folding, intracellular trafficking, stability, and half-life, affecting protein function. Endoglycosidase H is a glycosidase that cleaves asparagine-linked oligomannose and hybrid, but not glycan complex, from N-linked glycoproteins. It hydrolyses the bond connecting the two N-acetylglucosamine residues comprising the chitobiose core, leaving an N-acetylglucosamine residue on the asparagine.

Source

Escherichia coli

Activity

One unit of Endoglycosidase H cleaves > 95% of the carbohydrate from 10 μ g of denatured RNase B in a total reaction volume of 10 μ L at 37°C for 1 h.

Storage buffer

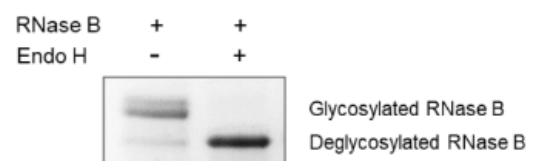
The enzyme is supplied in 20 mM Tris-HCl, 50 mM NaCl, 5 mM EDTA, pH 7.5

Storage and Stability

This product is stable after storage at -20°C for long-term storage under sterile conditions. Avoid repeated free-thaw cycles.

Procedure

1. Add 1–20 μ g of the target glycoprotein, 1 μ L of 10X Glycoprotein Denaturing Buffer, and an appropriate volume of H₂O to a final 10 μ L total reaction volume.
2. Heat the sample for protein denature at 100°C for 10 minutes.
3. Add 2 μ L of 10X Reaction Buffer, 1–5 μ L of Endo H (500–2500 units), and an appropriate volume of H₂O to a final 20 μ L total reaction volume.
4. Incubate reaction mixture at 37°C for 1 hour.
5. Determine the glycosylated level of the samples by SDS-PAGE analysis.



The standard assay was performed by incubating 1 unit of Endo H and 10 μ g of RNase B under the above conditions. SDS-PAGE analysis of RNase B digested with Endoglycosidase H.

Important notes

1. Please fine-tune the input sample volume to find the optimal condition for your assay.
2. Once optimize for the cleavage condition, the cleavage reactions can be scaled up to cleave a large amount of the target fusion protein.

Disclaimer

This product is for research use only and is not intended for diagnostic use.

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