Obín Science

Glycosidase Selection Strategies

Glycosylation is a common post-translational modification in eukaryotic cells, crucial for protein folding, stability, activity, and intercellular signaling. It can be categorized into N-linked and O-linked glycosylation. N-linked glycans are further classified into high-mannose, hybrid, and complex types.



Fig. 1 Three main types of glycans

However, glycosylation poses challenges: (1) it reduces protein homogeneity, hindering crystallization, and (2) abnormal glycosylation can cause diseases. Deglycosylation enzymes are valuable tools for studying glycoproteins and related diseases.





Fig. 5 The types of glycans recognized by Endo S2 and its cleavage site

It is important to note that the deglycosylation activity of Endo S, CU43, and Endo S2 is highly dependent on the intact conformation of human IgG. If human IgG is denatured, they cannot cleave the glycans. Additionally, Endo S, CU43, and Endo S2 are unable to remove glycans from human IgA, IgM, IgE, or IgGs from other species.

The following is a summary of the four enzymes:

Enzyme	Catalog No.	Target Glycan Types	Recognition Site	Dependence on Protein Conformation
PNGase F	JN842012	High-mannose, hybrid, complex	None	Works on both denatured and native proteins
Endo S	JN936012	Complex glycans	Human lgG Asn297	Specific to native human IgG
CU43	JN946012/JN946011	Hybrid, complex glycans	Human lgG Asn297	Specific to native human IgG
Endo S2	JN865012	High-mannose, hybrid, complex	Human lgG Asn297	Specific to native human lgG

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