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Product Information

PNGase Fast Kit

PNGase F from *Elizabethkingia meningoseptica* recombinant, expressed in *E. coli*

Catalog Number **EMS0001** Storage Temperature 2–8 °C

TECHNICAL BULLETIN

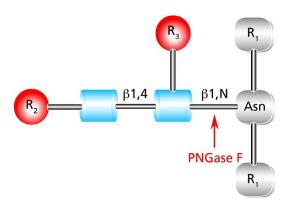
CAS RN 83534-39-8

EC 3.5.1.52

Synonyms: N-Glycosidase F, Peptide N-glycosidase

Product Description

PNGase F (Peptide N-Glycosidase F) cleaves asparagine-linked high mannose as well as hybrid and complex oligosaccharides from glycoproteins. It deaminates the asparagine to aspartic acid, but leaves the oligosaccharide intact. PNGase F will not remove oligosaccharides containing $\alpha(1\rightarrow 3)$ -linked core fucose, commonly found in plant glycoproteins. A tripeptide with the oligosaccharide-linked asparagine as the central residue is the minimal substrate for PNGase F.



Molecular mass: 36 kDa

pH range: 6–10 Optimal pH: 8.6.

Components

Sufficient reagents for $50 \times 100 \mu g$ reactions

- Denaturing Buffer, 5× Reaction Buffer 500 μL Proprietary detergent solution in buffer Catalog Number EMS0002
- PNGase F Enzyme Solution 50 μL Supplied in a solution of 20 mM Tris-HCl, pH 7.5, containing 1 mM EDTA and 50 mM NaCl Catalog Number EMS0003

Reagents and Equipment Required but Not Provided.

- 2-Mercaptoethanol (βME, Catalog Number M6250 or equivalent)
- 30 kDa MWCO Centrifugal Filtration Units, 0.5 mL (Catalog Number MRCF0R030, UFC5030, or equivalent). 10 kDa filters will also work, but spin time must be increased.
- 50 mM ammonium bicarbonate solution

Precautions and Disclaimer

This product is for R&D use only, not for drug, household, or other uses. Please consult the Safety Data Sheet for information regarding hazards and safe handling practices.

Preparation Instructions

The enzyme and reaction buffer are ready to use.

Storage/Stability

The kit should be stored at 2–8 °C. Components have been shown to be stable for at least 2 years if stored properly. The detergent solution may crystallize at 2–8 °C, but can easily be resolubilized by heating at 70 °C for 5 minutes and vortexing. Repeat if necessary.

Procedures

Sample preparation and handling

For best results, it is recommended to start with protein concentrations of 5–10 mg/mL in a low salt buffer. NaCl concentrations of greater than 250 mM can cause proteins to aggregate and precipitate during denaturation.

The following procedure is scaled for 10 μ L of a protein solution, which should correspond to 50–100 μ g of protein mass. Volumes of reagents can be scaled appropriately if larger or smaller sample inputs are desired. If concentration of the sample is necessary, it is recommended to desalt, dry down, and reconstitute the protein in water to achieve the desired concentration.

The detergents in the Denaturing Buffer can interfere with subsequent glycan labeling and analysis. These can be efficiently removed using centrifugal filtration units noted in the optional reagents section. For analysis of deglycosylated proteins, detergent removal may not be necessary. Procedure, Step 2 can be omitted.

Note: Some proteins are not amenable to fast deglycosylation techniques. It is always best to compare the results of a fast digest to a traditional overnight digest or the established protocol in use.

Deglycosylation Procedure

- 1. Denature glycoprotein
 - a. Transfer 10 μL of sample solution (50–100 μg of protein) to a small tube.
 - b. Add 2.5 μ L of Denaturing Buffer (Catalog Number EMS0002).
 - c. Dilute βME 100-fold in water and add 0.5 μL to the reaction mix to make ~5 mM in βME .
 - d. Mix well.
 - e. Incubate at 70 °C for 10 to 15 minutes to denature. For proteins which have shown a tendency to aggregate or precipitate when heated, a lower denaturing temperature may be used. For example, 50 °C has been shown to be effective for some proteins, but in general deglycosylation is most efficient after denaturing at 70 °C.
- 2. Detergent removal
 - a. Transfer the denatured protein solution to the filter cup of a 30 kDa centrifugal filtration unit.
 - b. Centrifuge at $10,000-14,000 \times g$ for 5 minutes.
 - c. Add 100 μ L of 50 mM ammonium bicarbonate.
 - d. Centrifuge at $10,000-14,000 \times g$ for 5 minutes.
 - e. Transfer the filter cup to a clean tube.
 - f. Add 50 μ L of 50 mM ammonium bicarbonate to the filter cup and transfer the filter unit to 37 °C.
- 3. Deglycosylation
 - a. Allow to equilibrate at 37 °C for 5 minutes.
 - b. Add 1 μL of PNGase F Enzyme Solution (Catalog Number EMS0003).
 - c. Mix well.
 - d. Incubate at 37 °C for 10-20 minutes.
 - e. The majority of glycans will be released in 10 minutes.
 - f. More difficult glycans may require longer incubations.
- 4. Recover the glycans by centrifugation.
 - a. Centrifuge at $10,000-14,000 \times q$ for 5 minutes.
 - b. Add 50 μ L of 50 mM ammonium bicarbonate to the filter cup and centrifuge again.
 - c. The glycans will be in the flow-through in the lower part of the centrifugal unit.
 - d. Label as desired for analysis.
- 5. Deglycosylated protein can be recovered from the top of the filtration unit.
 - a. Invert the filter insert into a clean tube.
 - b. Centrifuge at $10,000-14,000 \times q$ for 5 minutes.
 - c. Rinsing the filtration unit with ammonium bicarbonate can improve recovery.

JK,JAB,MAM 03/18-1