

**DESCRIPTION**

**Source** *E. coli*-derived *f. meningosepticum* PNGase F protein  
Ala41-Asn354 with N-terminal Met and 6-His tag  
Accession # P21163

**N-terminal Sequence Analysis** Met

**Predicted Molecular Mass** 36 kDa

**SPECIFICATIONS**

**SDS-PAGE** 34 kDa, reducing conditions

**Activity** Measured by its ability to deglycosylate ribonuclease B under denatured conditions.  
>50% ribonuclease B (10 µg) is deglycosylated by 2.5 ng rFmPNGase F within 30 minutes, as measured under the described conditions.

**Endotoxin Level** <1.0 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: 0.1 M Tris, pH 7.5
  - Denaturing Buffer (10X): 5% SDS, 0.8 M β-Mercaptoethanol
  - Recombinant *F. meningosepticum* PNGase F (rFmPNGase) (Catalog # 9109-GH)
  - Ribonuclease B, from bovine pancreas (RNase B) (Sigma, Catalog # R7884), 2.5 mg/mL stock in 25 mM Tris, pH 7.5
  - 10% Triton® X-100 (Amresco, Catalog # M236)
  - Reducing SDS-PAGE Sample Buffer
  - SDS-PAGE or Western Blot

- Assay**
1. Dilute Denaturing Buffer to 5X in deionized water.
  2. Create a Substrate Mixture containing 0.8 mg/mL RNase B and 1X Denaturing Buffer in deionized water.
  3. Heat Substrate Mixture at 100 °C for 10 minutes. Cool to room temperature and microcentrifuge briefly.
  4. Add 10% Triton® X-100 to a final concentration of 1.67%.
  5. Dilute rFmPNGase F to 0.167 ng/µL in Assay Buffer.
  6. Combine 15 µL of Substrate Mixture and 15 µL 0.167 ng/µL rFmPNGase F. Include a control containing 15 µL of Substrate Mixture and 15 µL of Assay Buffer.
  7. Incubate reaction mixture at 37 °C for 30 minutes.
  8. Combine equal volumes of incubated reaction mixture and reducing SDS-PAGE sample buffer and boil samples at 100 °C for 3-5 minutes.
  9. Load 15 µL (2.5 µg RNase B) per lane on a 4-20% SDS-PAGE gel.
  10. Stain gel and analyze for percent deglycosylation using densitometry.

- Final Assay Conditions** Per Reaction:
- rFmPNGase F: 2.5 ng
  - RNase B: 10 µg

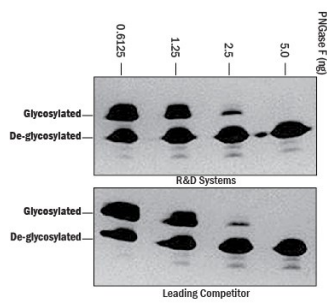
**PREPARATION AND STORAGE**

**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.

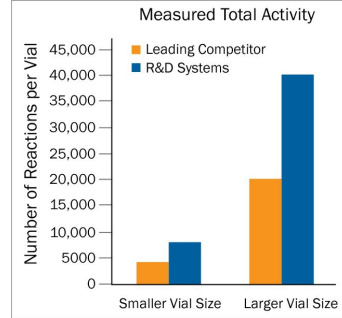
**DATA**

**Enzyme Activity**



Recombinant *F. meningosepticum* PNGase F (Catalog # 9109-GH) from R&D Systems and a leading competitor are able to deglycosylate 10 µg of RNase B at 37 °C in one hour. The *E. coli*-produced enzyme from R&D Systems offers a better value than the competition

**Bioactivity**



Total activity per vial of Recombinant *F. meningosepticum* PNGase F (Catalog # 9109-GH) compared to the leading competitor. R&D Systems® PNGase F gives you 2x more enzyme at a comparable price.

**BACKGROUND**

PNGase F, peptide N-glycosidase F from *Flavobacterium meningosepticum*, catalyzes the hydrolysis of asparagine-linked high mannose, as well as hybrid and complex oligosaccharides from glycoproteins (1). Unlike glycosidases that hydrolyze glycosidic bonds, PNGase F is an amidase that cleaves the beta-aspartylglucosamine bond between the innermost GlcNAc of N-glycans and asparagine residues of glycoproteins (2). The enzyme is highly active on various N-glycans except those with the innermost GlcNAc modified with alpha 1-3-linked core fucose, which is commonly found on plant glycoproteins (3). Cleavage with PNGase F will convert the asparagine residue to an aspartic residue, allowing identification of the glycosylation site by mass spectrometry (4). This purified enzyme is compatible with glycan analysis using mass spectrometry.

**References:**

1. Elder, J.H. and Alexander, S. (1982) Proc. Natl. Acad. Sci. USA **79**:4540.
2. Maley, F. *et al.* (1989) Anal. Biochem. **180**:195.
3. Tarentino, A.L. and Plummer, T.H. (1994) Methods Enzymol **230**:44.
4. Zhang, H. *et al.* (2003) Nat. Biotechnol. **21**:660.