

# Agilent AdvanceBio N-Glycanase-PLUS Peptide-N-Glycosidase F (PNGase F), ≥10 U/mL

(Peptide-N-Glycosidase F) Specifications	
Product Code	GKE-5010
Specific Activity	≥10 U/mg One unit of N-Glycanase-PLUS is defined as the amount of enzyme required to catalyze the release of N-linked oligosaccharides from 1 μmole of denatured ribonuclease β per minute at pH 7.5 and 37 °C.
Activity	≥10 U/mL
Shipping	Shipped on ice pack for next day delivery.
Storage	2 to 8 °C or –20 °C, but avoid repeated freeze-thawing.
Formulation	A sterile-filtered solution in 20 mM Tris-HCl, 50 mM NaCl, 1 mM EDTA (pH 7.5)

## Introduction

N-Glycanase-PLUS [PNGase F: Peptide-N<sup>4</sup>-(acetyl-β-glucosaminy)-asparagine amidase, EC 3.5.1.52] is isolated from a strain of *E. coli*, expressing a cloned gene from *Chryseobacterium [Flavobacterium] meningosepticum* (formerly ProZyme).

N-Glycanase is widely used for the removal of N-glycans from glycoproteins and glycopeptides. It is well known that denaturation of glycoprotein substrates before enzyme digestion dramatically increases the efficiency of their deglycosylation, allowing complete removal of most classes of N-glycans from glycoproteins. In contrast, deglycosylation proceeds rather slowly, and in some cases incompletely, with native glycoprotein substrates presumably due to steric constraints.<sup>11</sup> However, it is often desirable to obtain deglycosylation of glycoprotein in the absence of denaturants and detergents to allow structural or functional studies of the folded protein. Studies have shown that to obtain efficient deglycosylation of the native glycoprotein substrates, it is important to use a high starting concentration of the enzyme. Recombinant N-Glycanase-PLUS is recommended for all applications requiring deglycosylation of glycoproteins in the absence of denaturants. The high activity also allows microscale reaction volumes and shorter reaction times to be explored. The highly concentrated enzyme preparation is the reagent of choice for efficient deglycosylation in the absence of detergents, which facilitates subsequent analysis by electrospray or MALDI-TOF mass spectrometry.<sup>19</sup>



Sulfhydryl reagents such as  $\beta$ -mercaptoethanol used for glycoprotein denaturation do not interfere with enzyme activity. N-Glycanase-PLUS tolerates most chaotropic agents, and is at least 80% active in the presence of <5 M urea, <2 M guanidine HCl and 0.25 M NaSCN; however, the enzyme is inactivated by the presence of guanidine thiocyanate.<sup>4</sup>

N-Glycanase-PLUS is compatible with a wide range of buffers.<sup>5</sup> The purified enzyme is free from detectable protease activity. Additional protease inhibitors (for example, PMSF, pepstatin A, benzamidine, aprotinin, leupeptin, and 1,10-phenanthroline) can be included in enzyme digestions to inhibit any other types of proteases present in samples. This is particularly important when deglycosylation under native conditions is performed, and retention of protein conformation is desirable. Deglycosylation efficiency against metalloprotein substrates has been suggested to be enhanced by inclusion of EDTA at between 0.1 and 1 mM final concentration. Deglycosylation can be conveniently analyzed using SDS-PAGE if the loss of glycans results in a significant lowering of the protein's molecular weight.

#### **Procedure for deglycosylation (denaturing conditions)**

1. Prepare 50 to 500  $\mu$ g glycoprotein solution in 45  $\mu$ L of 1x Incubation Buffer. Add 2.5  $\mu$ L of SDS/ $\beta$ -mercaptoethanol (final reaction concentration 0.1% SDS, 50 mM  $\beta$ -mercaptoethanol).
2. Denature glycoprotein by heating at 100 °C for five minutes. Allow the mixture to cool.
3. Add 2.5  $\mu$ L of NP-40 (final reaction concentration 0.75% detergent).
4. Add 1  $\mu$ L N-Glycanase-PLUS to reaction mixture and incubate for two hours to overnight at 37 °C.

## References

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