

# **Certificate of Analysis**

# Human IgG Glycoprotein Standard

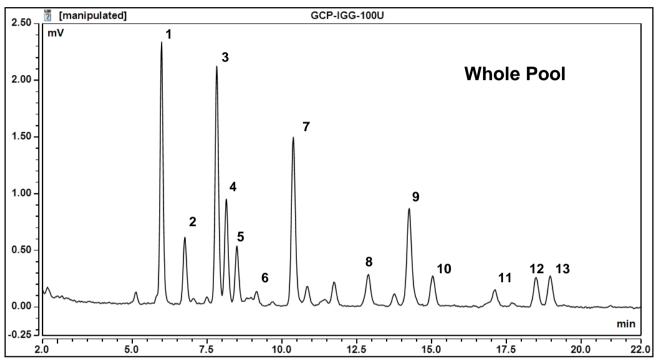
Cat. #: GCP-IGG-10	00U Batch: B87J-02	Nominal size: 100µg	
Description:	A glycoprotein standard for use during glycan release and labeling.		
Source:	This product is purified from human blood. Human IgG glycoprotein standard contains a mixture of gamma-type antibodies.		
Glycans:	(Table 1). These glycans exist in both fucosylated and non-fu without bisecting GlcNAcs, and terminate either with 0, 1 or 2	najority of glycans present on human antibodies are N-link bi-antennary types e 1). These glycans exist in both fucosylated and non-fucosylated forms, with and ut bisecting GlcNAcs, and terminate either with 0, 1 or 2 galactoses (the G0, G1 G2 glycan forms). Sialic acid terminating glycans are also present on native human	
Form:	Dry. Lyophilised powder.		
Molecular Weight:	~150kDa (protein weight only)		
Amount:	109 $\mu$ g protein (In comparison to BSA standard, determined b rounded to nearest $\mu$ g)	by BCA assay. Value	
Storage:	Refrigerate (-20°C) both before and after dissolving. This product is stable for at least 5 years as supplied.		
Shipping:	The product is shipped at ambient temperature.		
Handling:	Once dissolved avoid repeated thawing and refreezing, storage temperature or above, exposure to light and long term expose cause glycan desialylation.	-	
Safety:	This product is non-hazardous and has been purified from na be free of all hazardous material including pathogenic biologic <b>For research use only. Not for human or drug use</b>		



#### Analysis:

IgG glycans were released from human IgG Glycoprotein (Cat# GCP-IGG-100U) using PNGase F.

Following release the glycans were labeled using 2-Aminobenzamide (2-AB) using the LudgerTag<sup>™</sup> 2-AB Glycan Labeling Kit (Cat# LT-KAB-A2).



*Figure 1: HILIC HPLC profile of 2-AB labelled IgG N-glycans, released by PNGase F from GCP-IGG-100U batch B87J-02.* See Table 1 for peak assignment.

Figure 1 shows a LudgerSepN2 HPLC profile of IgG N-glycans. To thoroughly investigate the N-glycans we first separate them based on charge on a LudgerSepC3 column (Figure 2) and then run each fraction on a LudgerSepN2 column. From these studies, combined with exoglycosidase investigation (Figure 3) ,we identified the glycans shown in Table 1. For further information on Glycoprofiling please contact us at info@ludger.com



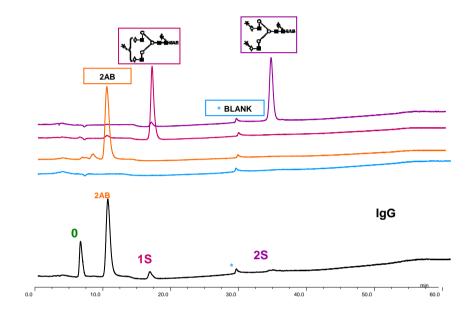


Figure 2: LudgerSep-C3 profile analysis of human IgG PNGaseF released N-glycans from a similar batch of GCP-IGG. 2-AB labelled glycans were separated on the LudgerSepC3 column. It is evident from this analysis that the proportion of sialylated glycans is relatively low (see 1S and 2S peaks above). A combination of LudgerSepC3/N2 and exoglycosidase digestion is required to identify the glycans and their relative abundance, see Table 1.



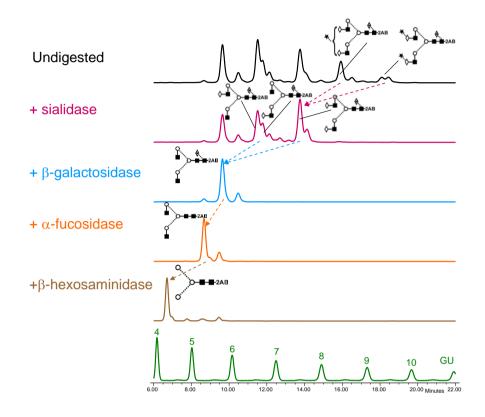


Figure 3: Example exoglycosidase analysis of Human IgG N-glycans from a similar batch of GCP-IGG. LudgerSepN2 chromatograms are shown.



Peak No.	<b>2</b> 4		Whole Pool %
	Structure	GU	Area
1	F(6)A2	5.76	16.5
2	F(6)A2B	6.14	
	A2[6]G(4)1	6.14	4.9
	A2[6]BG(4)1	6.14	
3	A2[3]G(4)1	6.59	
	A2[3]BG(4)1	6.59	18.2
	F(6)A2[6]G(4)1	6.59	
4	F(6)A2[3]G(4)1	6.70	7.2
5	F(6)A2[6]BG(4)1	6.85	4.4
	F(6)A2[3]BG(4)1	6.85	4.4
6	A2G(4)2	7.08	2.1
7	F(6)A2G(4)2	7.53	19.4
	F(6)A2BG(4)2	7.53	19.4
8	A2G(4)2S1	7.98	3.0
9	F(6)A2G(4)2S1	8.42	11.6
10	F(6)A2BG(4)2S1	8.66	3.3
11	A2G(4)2S2	8.90	1.6
	A2BG(4)2S2	8.90	
12	F(6)A2G(4)2S2	9.32	3.6
13	F(6)A2BG(4)2S2	9.47	3.3

**Table 1: Summary of Human IgG N-glycans found in a similar batch of GCP-IGG.** See the end of this document for details of the glycan nomenclature used. GU – glucose units – a system of comparing glycans to a glucose homopolymer standard. Many common N-glycans have reported GU values. A combination of GU value, mass spectrometry and exoglycosidase digestion (Figure 3), can be used to unambiguously identify most N-glycans.



### **Structure Abbreviations**

All N-glycans have two core GlcNAcs; F at the start of the abbreviation indicates a core fucose, (6) after the F indicates that the fucose is  $\alpha$ 1-6 linked to the inner GlcNAc; Mx, number (x) of mannose on core GlcNAcs; Ax, number of antenna (GlcNAc) on trimannosyl core; A2, biantennary with both GlcNAcs as  $\beta$ 1-2 linked; A3, triantennary with a GlcNAc linked  $\beta$ 1-2 to both mannose and the third GlcNAc linked  $\beta$ 1-4 to the  $\alpha$ 1-3 linked mannose; A3', triantennary with a GlcNAc linked  $\beta$ 1-2 to both mannose and the third GlcNAc linked  $\beta$ 1-6 to the  $\alpha$ 1-6 linked mannose; A4, GlcNAcs linked as A3 with additional GlcNAc  $\beta$ 1-6 linked to  $\alpha$ 1-6 mannose; B, bisecting GlcNAc linked  $\beta$ 1-4 to  $\beta$ 1-3 mannose; Gx, number (x) of linked galactose on antenna, (4) or (3) after the G indicates that the Gal is  $\beta$ 1-4 or  $\beta$ 1-3 linked; [3]G1 and [6]G1 indicates that the galactose is on the antenna of the  $\alpha$ 1-3 or  $\alpha$ 1-6 mannose; Sx, number (x) of sialic acids linked to galactose; the numbers 3 or 6 in parentheses after S indicate whether the sialic acid is in an  $\alpha$ 2-3 or  $\alpha$ 2-6 linkage.

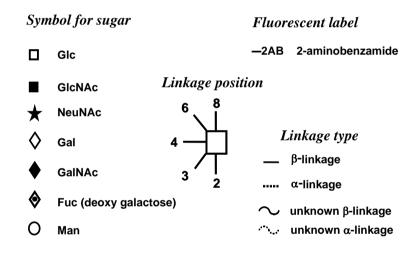


Figure 4. Symbols used to depict glycan structures.



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#### This product is intended for *in vitro* research only.

### Address

Ludger Ltd, Culham Science Centre, Oxford OX14 3EB United Kingdom

Tel: +44 1865 408 554 Fax: +44 870 163 4620 Email: info@ludger.com www.ludger.com