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Author Comments:	This work claims to expand the use of PGA technology to investigate repertoire (specificities) and levels of circulating anti-carbohydrates antibodies, both in health and during any pathological condition. This kind of study can be extrapolated to different species (rat, primates, chickens, humans). We anticipate the potentiality that this approach may bring in the early diagnosis and derived treatment in some of the pathological conditions where antibodies seem to play an important role. As JoVE is a leading video journal, we consider it the perfect platform to disseminate the mentioned methodology to the scientific community.
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TITLE:

Printed Glycan Array: A Sensitive Technique for the Analysis of the Repertoire of Circulating Anti-Carbohydrate Antibodies in Small Animals

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SHORT ABSTRACT:

This work shows the potential of printed glycan array (PGA) technology for the analysis of circulating anti-carbohydrate antibodies in small animals.

LONG ABSTRACT:

The repertoire of circulating anti-carbohydrate antibodies of a given individual is often

associated with its immunological status. Not only the individual immune condition determines the success in combating internal and external potential threat signals, but also the existence of a particular pattern of circulating anti-glycan antibodies (and their serological level variation) could be a significant marker of the onset and progression of certain pathological conditions. Here, we describe a PGA-based methodology that offers the opportunity to measure hundreds of glycan targets with very high sensitivity; using a minimal amount of sample, which is a common restriction present when small animals (rats, mice, hamster, etc.) are used as models to address aspects of human diseases. As a representative example of this approach, we show the results obtained from the analysis of the repertoire of natural anti-glycan antibodies in BALB/c mice. We demonstrate that each BALB/c mouse involved in the study, despite being genetically identical and maintained under the same conditions, develops a particular pattern of natural anti-carbohydrate antibodies. This work claims to expand the use of PGA technology to investigate repertoire (specificities) and the levels of circulating anti-carbohydrates antibodies, both in health and during any pathological condition.

INTRODUCTION:

Antibodies play a central role in our defense against invading pathogens by directly neutralizing viruses^{1,2} and bacteria^{2,3}, by activating the complement system^{4,5} and the enhancement of phagocytosis⁶. Additionally, they are essential elements in cancer targeting and elimination of malignant cells⁷, and homeostasis maintenance^{8,9}.

Disorders of the immune system can result in autoimmune and inflammatory diseases¹⁰ and cancer¹¹. All these pathological conditions ideally demand a prompt diagnosis for an efficient treatment. In the case of autoimmune disorders, the serological presence of autoantibodies in most of the cases is a predictor for diagnostic of autoimmunity^{10,12}. These antibodies react with the cell surface and extracellular autoantigens and, they are often present for many years before the presentation of autoimmune disease^{10,12}. Immune deficiencies and cancer are also diagnosed with blood tests that either measure the level of immune elements such as antibodies, or their functional activity¹¹.

The identification of the repertoire of circulating antibodies and their serological levels are paramount to set a prognosis and evaluate the progression of all of the mentioned pathological conditions. We have previously demonstrated the potential of PGA technique for the analysis of circulating antibodies in different animal species¹³⁻¹⁶, minimizing the use of large volumes of serological samples, avoiding the problem associated with antibodies cross-reactivity¹⁷ and allowing high-throughput profiling of an extensive repertoire of antibodies¹⁵.

Glycan-based immunoassays are mainly conditioned, among other factors, by the origin and production of carbohydrates, which determine the affinity and binding of ligands^{15,18-21}. Glycan-based immunoassays can be developed in suspension (microspheres)^{15,21,22} or in flat-activated surfaces^{15,21,23,24}. The last include ELISA (the most conventional of these methods) and PGA. There is not much data comparing these methodologies in the same experimental setting^{15,25-27}. We have previously compared the efficacy and selectivity of these immunoassays to profile anti-glycan antibodies in individual human plasma samples¹⁵. For some antibodies such as those

targeting anti-A/B blood group, all the immunoassays could detect them with statistical significance and they positively correlated with each other^{15,18,21}. Meanwhile, anti-P1 antibodies were primarily detected by PGA with the highest discriminative power, and there was no correlation in the determinations made by the different glycan-based immunoassays^{15,18,21}. These differences between methods were mainly related to the antibody/antigen ratio and glycan orientation¹⁵. ELISA and suspension arrays are more susceptible to unspecific binding than PGA because there is an excess of antigen over antibodies in these methods¹⁵. Additionally, the orientation of glycans in the PGA is more restricted than in ELISA and suspension arrays¹⁵. ELISA is convenient when the study includes a limited panel of glycans. Along with suspension arrays, ELISA offers broader flexibility regarding assay reconfiguration. PGA is exceptionally convenient for discovery approaches^{15,18,21,28}. Despite these clear advantages and disadvantages, the three mentioned immunoassays could be used to study different aspects of glycan-antibody interactions. The final goal of the study is the one will guide the selection of the more suitable methodology.

The present work aims to extend the use of PGA technology for the analysis of the repertoire of circulating anti-glycan antibodies in small animals. As a representative result, we present here a detailed protocol to assess the repertoire of natural anti-carbohydrate antibodies in adult BALB/c mice by PGA.

PROTOCOL:

1. Glyochips Production

1.1. Microarray preparation

1.1.1. Print the glycans (50 mM) and polysaccharides (10 µg/mL) in 300 mM phosphate buffered saline (PBS, pH 8.5) at 6 replicates onto N-hydroxysuccinimide-derivatized glass slides, using non-contact robotic arrayer (drop volume ~ 900 pL). Each slide contains 4 different blocks of sub-arrays (**Figure 1A**, in colors) repeated 6 times. Every single sub-array is formed by 112 different glycan spots, including controls (8 rows × 14 columns) (**Figure 1B**).

Note: Glycan-related information is provided in the **Supplementary Table 1**. The glycan library used for printing microchips is the result of a long-term synthetic effort of the IBCh team; examples of synthesis are described in related publications²⁹⁻³⁸. The glycan library included blood group antigens and some of the most frequently occurring terminal oligosaccharides, as well as core motifs of mammalian N- and O-linked glycoproteins and glycolipids, tumor-associated carbohydrate antigens, and polysaccharides from pathogenic bacteria.

1.1.2. Incubate the slides in moisture box (relative humidity ~70%) at room temperature (25 °C) for 1 h.

1.1.3. Blocking microarrays: incubate the slides for 1.5 h with blocking buffer at room temperature (100 mM boric acid, 25 mM ethanolamine, 0.2% (v/v) Tween-20 in ultrapure water).

1.1.4. Wash the glycochip with ultrapure water and dry it by air.

1.2. Glycochip quality control

1.2.1. Analyze two microarrays from each batch using 1mg/mL solution of complex immunoglobulin preparation (CIP, containing IgG, IgM and IgA), 10 µg/mL solution of biotinylated goat anti-human immunoglobulins as a secondary antibody (IgM + IgG + IgA), followed by 1 µg/mL solution of the corresponding fluorescent streptavidin conjugate (via protocol described below, see step 2).

1.2.2. Scan and analyze the glycochips (see step 3, analysis of glycan array).

1.2.3. Use microarray batches with intra- and inter-chip correlation higher than 0.9.

2. Glycan array technique

2.1. Prepare the following aqueous solutions (in ultrapure water) and store them at room temperature (25 °C):

- Buffer-1: 1% (w/v) bovine serum albumin (BSA) in PBS, 1% (v/v) Tween-20 and 0.01% (w/v) NaN₃
- Buffer-2: 1% (w/v) BSA in PBS, 0.1% (v/v) Tween-20 and 0.01% (w/v) NaN₃.
- Buffer-3: 0.1% (v/v) Tween-20 in PBS.
- Buffer-4: 0.001% (v/v) Tween-20 in PBS.

2.2. Glycochip and sample preparation

2.2.1. Put the storage box with the slides on the table until they reach room temperature (25 °C).

Note: Use powder-free latex gloves. The glycochip must be manipulated by the bottom part of the glass slide, where the barcode is located. The barcode will help you to identify the right side, avoiding the contact with the surface where the glycans are printed.

2.2.2. Open the box, take the glycochip and place it in the incubation chamber (25 °C), already conditioned with wet filter paper to keep humidity constant inside the chamber.

2.2.3. Meanwhile, dilute the mice serum with Buffer-1 (1:20) in 1.5 mL tubes. Homogenize the serum solution (5 s) with a vortex mixer.

Note: The volume needed to totally cover a single glycochip surface is approximately 1 mL.

2.2.4. After the homogenization, incubate the diluted serum at 37 °C for 10 min in a water bath to avoid immunoglobulin aggregation. Centrifuge the tubes for 3 min at 10,000 x g and 25 °C, collect the supernatant and discard any precipitated material.

2.2.5. Place the glycochip carefully in the incubation chamber. Incubate it for 15 min at 25 °C with 1 mL of Buffer-3 to eliminate any residual material on the surface of the glycochip.

2.2.6. Hold the glycochip in a vertical position and rewash it with some drops of Buffer-3 using a plastic Pasteur pipette. Carefully remove the buffer from the glycochip surface using filter paper.

2.3. Reaction: antibodies binding

2.3.1. Place the glycochip in the incubation chamber. Spread the diluted serum sample over the glycochip surface using a micropipette. Incubate with orbital agitation (30 rpm) at 37 °C for 1.5 h. Ensure that all dry area of the glycochip surface is covered by the diluted serum sample using the tip of the pipette.

2.3.2. Remove any excess sample and immerse the glycochip for 5 min in Buffer-3 at 25 °C. Then, pass the glycochip to a container with Buffer-4 (5 min) and finally wash (5 min) the glycochip with ultrapure water. Centrifuge the glycochip for 1 min at 175 x g and 25 °C to remove the liquid.

2.4. Detection: secondary antibody

2.4.1. Place the glycochip in the incubation chamber. Spread over the glycochip surface a solution (5 µg/mL) of goat anti-mouse (IgG + IgM) conjugated to biotin in Buffer-2. Incubate with orbital agitation (30 rpm) at 37 °C for 1 h.

2.4.2. Remove the unbound fraction and repeat the washing steps.

2.4.3. After the centrifugation, incubate the glycochip in darkness at 25 °C for 45 min (30 rpm) with 2 µg/mL of the corresponding fluorochrome-labeled streptavidin solution (in Buffer-2).

2.4.4. In darkness, remove the unbound fraction and repeat the washing steps.

2.4.5. Dry the glycochip by air.

Note: Glycochip should be scanned as soon as possible. But if it's impossible to do scan immediately after staining, glycochips can be stored in a cool and dry place in darkness.

3. Analysis of Glycan Array

3.1. Scan the array

3.1.1. Leave the glycochip on the table until it reaches room temperature in dark. At the same time, turn on the slide scanner and the laser (excitation wavelength of 633 nm).

3.1.2. Holding the microarray, slide the glycochip into the slot until it touches the back.

3.1.3. Scan the glycochip (**run easy scan**), and save the scan as a “.TIFF” file.

3.2. Array quantification

3.2.1. Quantify the array using a ScanArray Analysis System. Open previously scanned images, by clicking **File** in the **Configure & File group** on the Main Window (**Figure 1B-D**)

3.2.2. Load the corresponding array file template in GAL format (disposition of printed glycans on the glass slide) (**Figure 1C**).

3.2.3. Adjust GAL template by carefully aligning the array (grids) with the spots in the image and initiate quantification (**Figure 1D**).

3.2.4. Select the quantification parameters:

- Quantification type: Run Easy Quant.
- Quantification method: Fixed Circle
- Auto find Spots: unclick all options
- Normalization Method: LOWESS (Locally Weighted Scatter Plot Smoothing).

3.2.5. Save the quantified data as “.CSV” file (**Figure 1D**). Transfer this data into a common spreadsheet file using Microsoft Excel or another appropriate application.

3.2.6. Use the interquartile range (IQR) as the main statistical method: calculation of the median (Quartile 2) of all signals for each ligand and the interquartile deviation (75th and 25th percentiles, or upper and lower quartiles Q3 and Q1, respectively).

3.2.7. Perform interactive exploration of data by using a Hierarchical Clustering Explorer application.

3.2.8. Use clustering parameters: Average Linkage (UPGMA) and Euclidean distance as similarity distance measure. Perform hierarchical clustering by rows without normalization.

REPRESENTATIVE RESULTS:

Here, we present a summary of representative results obtained from the quantification of the repertoire of natural anti-glycan antibodies in a population of 20 BALB/c mice. The glycochips used in this study contained 419 different glycan structures. Most glycans were synthesized as –

CH₂CH₂CH₂NH₂ spacer-armed O-glycosides, in several cases as –CH₂CH₂NH₂ or –NHCOCH₂NH₂ glycosides. All glycan structures were characterized by high resolution (700- or 800 MHz) NMR spectroscopy, purified and tested by HPLC, indicating their >95% purity. We have simultaneously determined IgM + IgG anti-glycan antibodies due to a restriction in the amount of mouse serum. In the PGA, we considered values above 4,000 RFU as a positive signal of antibody binding (this value is ~10% of the top glycans RFU). The results presented in this work follow most of the guidelines for reporting glycan microarray-based data³⁹. Only 17% of carbohydrate structures demonstrated ≥4,000 RFU in the PGA (**Figure 2**, in red). Most of the glycan structures exposed in the glycochips were not recognized by the repertoire of circulating anti-glycan antibodies of BALB/c mice (**Figure 2**, in blue and white)²⁸. The conserved pattern of natural anti-carbohydrate antibodies of BALB/c included 12 different glycan specificities, with very high median signal intensities of antibodies binding (≥10,000 RFU **Table 1**)²⁸.

Figure 1. Schematic representation (not at scale) of the glycan array configuration, printing, and analysis. (A) Printed microchips are developed with a library of 419 different glycan structures, followed by the detection with an appropriate fluorescently labeled secondary antibody. Each slide contains 4 different blocks of sub-arrays (in colors), repeated 6 times. Every single sub-array is formed by 112 different glycan spots (8 rows × 14 columns), including controls. (B) A representative example of the images obtained from microchip scanning using a fluorescence scanner (third part of the image). (C) The process of aligning the "grid" to spots in every single sub-array (template adjustment during quantification). (D) The fluorescence is detected for each spot and results are transferred into a common spreadsheet file.

Figure 2. Repertoire of natural circulating anti-carbohydrate antibodies of BALB/c mice. Mouse serum (1:20) was incubated with the glycochips and scanned using a ScanArray reader. Data were analyzed with a microarray analysis system and results were expressed in relative fluorescence units (RFU) as the median ± median absolute deviation (MAD). Blue and white colors represent binding signals lower than 4,000 RFU (background); red color represents signals ≥4,000 RFU (positive binding). F, female; M, male (n = 20). This figure has been reproduced from Bello-Gil, D. *et al.*²⁸.

Table 1. Top rank glycan structures recognized by natural antibodies of BALB/c mice. Glycans with binding signals above 4,000 RFU in at least 80% of examined mice (n = 20). ^bsp means aminoethyl, aminopropyl or glycyI spacer. ^cfuranose; all other monosaccharides are in a pyranose form; Fuc residue has L-configuration, all other monosaccharides - D-configuration. This table has been modified from Bello-Gil, D. *et al.*²⁸.

Supplementary Table 1. List of glycans, their binding to natural circulating antibodies (IgM + IgG) of BALB/c mice (n=20), expressed in relative fluorescence units (RFU) as median ± MAD, and the number of animals exceeding cut off (≥ 4000 RFU). This table has been reproduced from Bello-Gil, D. *et al.*²⁸.

DISCUSSION:

Glycan microarrays have become indispensable tools for studying protein-glycan interactions⁴⁰.

The present work describes a protocol based on PGA technology to study the repertoire of circulating of anti-carbohydrate antibodies in BALB/c mice. Since PGA offers the possibility to screen large numbers of biologically unknown glycans, it is an exceptionally convenient discovery tool^{13,15,28}. The proposed method offers the possibility to measure, in the same experimental setting, hundreds of glycan structures using a reduced amount of serological sample (50 µL). This is especially critical in the case of small animals (little circulating blood volume), or when it's necessary to extract blood several times from the same experimental animal.

We demonstrated, as representative results, that genetically identical mice should not be considered as immunological equivalents; because they develop different patterns of natural anti-carbohydrate antibodies (only 12 glycan specificities were conserved). Serological levels for the rest of the repertoire of natural anti-carbohydrate antibodies varied considerably among the examined animals. Analysis of the gut microbiota of inbred animals⁴¹ could explain this heterogeneity⁴²⁻⁴⁶. If the production of natural anti-glycan antibodies is mediated by the antigenic stimulation of microbiota, and this is different among inbred mice⁴¹, fine specificity of these antibodies will not be identical.

The main drawback for PGA development is the access to well-defined glycan structures^{40,47}. Glycans produced in biological systems are heterogeneous^{40,47,48}, and their biosynthesis relies on the differential expression of carbohydrate enzymes, resulting in heterogeneous mixtures of glycoforms, each with a distinct physiological activity⁴⁷. The complex composition and configuration of the glycans present in the biological systems make their productions challenging^{40,47,48}. Along with chemo-enzymatic synthesis, glycans isolated from natural sources will continue to be the major source of glycans for arrays development⁴⁰. Low synthetic yields and the complex purification process from glycoproteins and glycosphingolipids make the efficient production of glycans at large scale difficult^{40,47,48}. Hence, the availability and the prices of glycans continue being a very limiting condition to expand the use of PGA as a discovery tool.

Additionally, within the protocol, critical steps mostly relating to the correct distribution of solutions (serum, secondary antibodies) over the glycochip surface must be executed with caution. The methodology requires, at least, 1 mL of these solutions, to homogenously soak all dry areas of the glycochip surface. This is crucial to obtain minimal differences between glycan replicates and also to avoid excessive background during quantification.

Despite the mentioned limitations, PGA is a very sensitive tool for approaches related to study protein-glycan interactions⁴⁰, or to study the repertoire of anti-glycan antibodies in a particular experimental setting or condition^{13,15,28}. This study can be extrapolated to different species (including human samples)^{13,15,23,28}, providing a versatile methodology for identifying the repertoire of circulating anti-carbohydrate antibodies.

We also anticipate the potentiality that this approach may bring in the early diagnosis and derived treatment in some of the pathological conditions where antibodies directed to glycan structures seem to play an important role.

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DISCLOSURES:

Nailya Khasbiullina and Alexey Nokel are employees of Semiotik LLC, who is the supplier of the glycochips used in this study.

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Figure 1

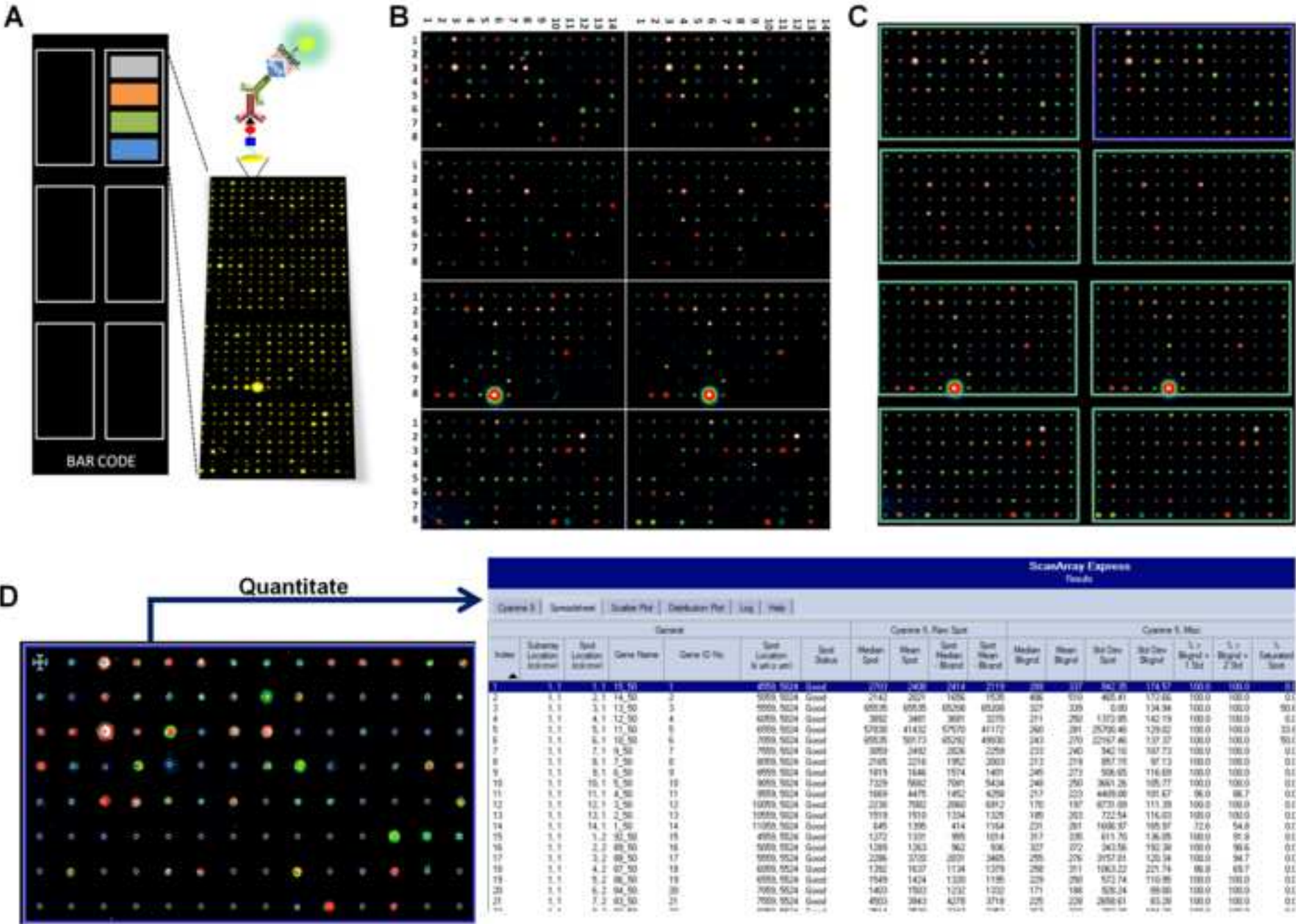
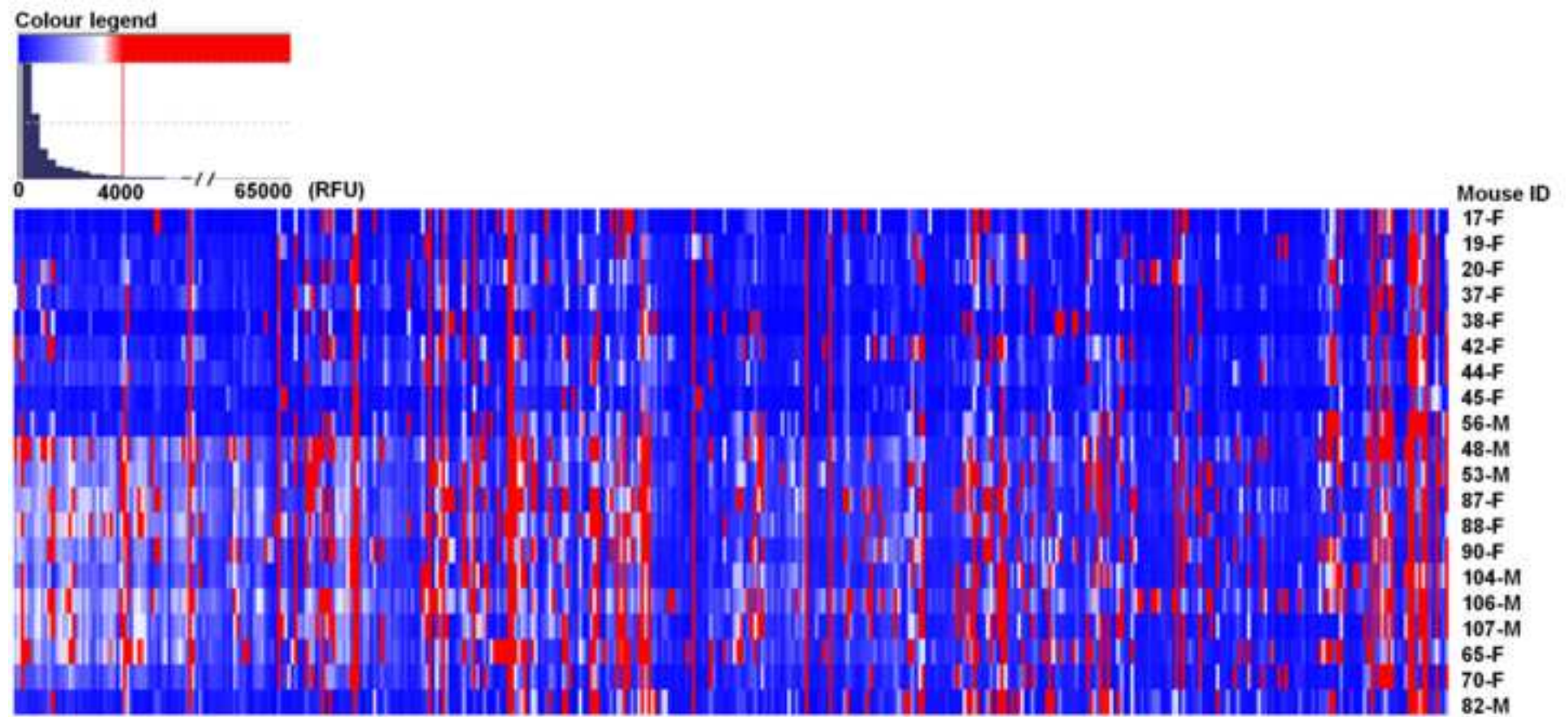


Figure 2



Glycan ID (#)	Structure
60	6-O-Su-Galβ-sp ^b
271	Galβ1-6Galβ1-4Glcβ-sp
802	Galβ1-3GalNAc(fur ^c)β-sp
176	3-O-Su-Galβ1-4(6-O-Su)Glcβ-sp
166	GlcAβ1-6Galβ-sp
150	3-O-Su-Galβ1-3GalNAcα-sp
437	GalNAcα1-3(Fucα1-2)Galβ1-3GalNAcβ-sp
125	6-Bn-Galβ1-4GlcNAcβ-sp
154	3-O-Su-Galβ1-3GlcNAcβ-sp
177	3-O-Su-Galβ1-4(6-O-Su)GlcNAcβ-sp
287	3-O-Su-Galβ1-3(Fucα1-4)GlcNAcβ-sp
234	Galβ1-4(Fucα1-3)GlcNAcβ-sp

Common name	Median and MAD as RFU		Number of mice showing RFU \geq 4000 (%)
	61113	1156	100
	53622	1934	100
	51348	2324	100
	43008	9342	100
	39105	2993	85
	37943	3232	100
A(type 4)	33886	3193	90
	32674	5389	95
	32651	3954	100
	32496	7215	100
SuLe ^a	20063	4962	95
Le ^x	13573	2635	80

MATERIAL/ EQUIPMENT	
Antibodies	
	biotinylated goat anti-human Igs
	biotinylated goat anti-mouse IgM + IgG
Equipment	
	Robotic Arrayer sciFLEXARRAYER S5
	Stain Tray (slide incubation chamber)
	Centrifuge
	Pipettes
	Slide Scanner
	Shaking incubator
Biological samples	
	BALB/c mice sera
	Complex Immunoglobulin Preparation (CIP)
Chemicals, Reagents and Glycans	
	Glycan library
	Bovine serum albumin (BSA)
	Ethanolamine
	Tween-20
	Phospahte buffered saline (PBS)
	Sodium azide
	Streptavidin Alexa Fluor 555 conjugate
	Streptavidin Cy5 conjugate
Materials	
	N-hydroxysuccinimide-derivatized glass slides H
	Whatman filter paper
	1.5 mL tubes
Software and algorithms	
	ScanArray Express Microarray Analysis System
	Hierarchical Clustering Explorer application

N/ A: not applicable

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Thermo Fisher Scientific, Waltham, MA, USA
Thermo Fisher Scientific
Scienion AG, Berlin, Germany
Simport, Beloeil, QC, Canada
Eppendorf, Hamburg, Germany
Gilson, Middleton, WI, USA
PerkinElmer, Waltham, MA, USA
Cole-Parmer, Staffordshire, UK
This paper
Immuno-Gem, Moscow, Russia
Institute of Bioorganic Chemistry (IBCh), Moscow, Russia
Sigma-Aldrich, St. Louis, MO,
Sigma-Aldrich
Merck Chemicals & Life Science S.A., Madrid, Spain
VWR International Eurolab S.L, Barcelona, Spain
Sigma-Aldrich
Thermo Fisher Scientific
GE Healthcare, Little Chalfont, Buckinghamshire, UK
Schott-Nexterion, Jena, Germany
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Eppendorf
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University of Maryland, MD, USA

IDENTIFIER
Ref. #: 31782
Ref. #: 31807
http://www.sciencion.com/products/sciflexarrayer/
Ref. #: M920-2
Ref. #: 5810 R
http://www.gilson.com/en/Pipette/
ScanArray G _x Plus
Ref. #: SI50
N/ A
http://www.biomedservice.ru/price/goods/1/17531
N/ A
Ref. #: A9418
Ref. #: 411000
Ref. #: 655204
Ref. #: E404
Ref. #: S2002
Ref. #: S21381
Ref. #: PA45001
Ref. #: 1070936
Ref. #: WHA10347509
Ref. #: 0000120000
http://www.per
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Analysis of the repertoire of circulating anti-carbohydrate antibodies in small animals by printed glycan array technology

Author(s):

Sara Olivera-Ardid, Nailya Khasbiullina, Alexey Nokel, Andrey Formanovsky, Inna Popova, Tatiana Tyrtysh, Roman Kunetskiy, Nadezhda Shilova, Nicolai Bovin, Daniel Bello-Gil, Rafael Mañez

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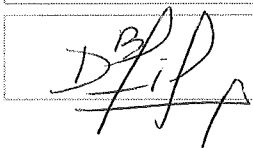
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Bing Wu, Ph.D.
Review Editor
Journal of Visualized Experiments (JoVE)

Barcelona, January 23th, 2018

Dear Dr. Wu,

Please find enclosed the reviewed version of the manuscript entitled "Printed Glycan Array: a sensitive technique for the analysis of the repertoire of circulating anti-carbohydrate antibodies in small animals" by Olivera-Ardid *et al.*, which we are submitting to consider for publication in JoVE.

First of all, thank you so much for your time and useful contribution. We appreciated the inputs the editorial team has given, which helped to improve our manuscript significantly.

Specific comments from the editorial team included:

1. Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammar issues.

Many thanks.

2. Please provide a title for Supplementary Table in Figure and Table Legends.

Thanks. (see line 299).

3. Milli-Q, Streptavidin Alexa Fluor 555 conjugate, commercial language.

Thanks. It was removed and substituted by:

- Ultrapure water.
- Fluorescent streptavidin conjugate.

4. Protocol step 2.1.1: Please write this step in imperative tense.

Thanks. It was corrected (line 150).

5. 2.2.2: What's the temperature of the incubation chamber?

25 °C (line 166).

6. 2.2.4: How is homogenization performed?

Homogenize the serum solution (5 s) with a vortex mixer (line 169).

Daniel Bello-Gil, PhD

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Infectious Pathology and Transplantation Division,
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7. Please revise the Discussion to explicitly cover the following in detail in 3-6 paragraphs with citations:

- a) Critical steps within the protocol

[Discussion line 335.](#)

- b) Any modifications and troubleshooting of the technique

[Please see the notes in the protocol.](#)

- c) Any limitations of the technique

[Discussion line 324.](#)

- d) The significance with respect to existing methods

[Introduction line 82-102.](#)

[Discussion line 305, 341.](#)

- e) Any future applications of the technique

[Discussion line 347](#)

We look forward to hearing from you regarding our resubmission. We would be glad to respond to any further questions and comments that you may have.

Sincerely,

Daniel.

Supplementary Material

Printed Glycan Array: a sensitive technique for the analysis of the repertoire of circulating anti-carbohydrate antibodies in small animals.

Sara Olivera-Ardid^{1,*}, Nailya Khasbiullina^{2,3,*}, Alexey Nokel³, Andrey Formanovsky², Inna Popova², Tatiana Tyrtysheva², Roman Kunetskiy², Nadezhda Shilova², Nicolai Bovin^{2,4}, Daniel Bello-Gil¹, Rafael Mañez^{1,5}

* **Correspondence:** dbello@idibell.cat

Table S1. List of glycans, their binding to natural circulating antibodies (IgM+IgG) of BALB/c mice (n=20), expressed in relative fluorescence units (RFU) as median \pm median absolute deviation (MAD), and the number of animals exceeding *cut off* (≥ 4000 RFU). This table has been reproduced from [28].

Glycan ID (#)	Structure	Common Name	Median and MAD as RFU		Animals showing RFU ≥ 4000
001	Fuc α -sp3	T _n Ser T _n	1590	342	1
002	Gal α -sp3		1522	316	0
003	Gal β -sp3		11858	1420	12
004	GalNAc α 1-OSer		2145	395	2
005	GalNAc α -sp3		1642	192	2
006	GalNAc β -sp3		1001	250	0
007	Glc α -sp3		1242	222	0
009	Glc β -sp3		1576	325	0
010	GlcNAc β -sp3		4124	471	1
011	GlcNAc β -sp2		4634	607	2
012	GlcNAc β -sp7	aminoglucitol	2068	523	2
013	GlcNAc β -sp8		7267	1117	9
014	GlcN(Gc) β -sp4		914	203	0
015	HOCH ₂ (HOCH) ₄ CH ₂ NH ₂		1374	316	0
016	Man α -sp3		1164	263	0
017	Man α -sp4		1567	369	0
018	Man β -sp4		1842	319	2
019	ManNAc β -sp4		4752	390	4
020	Rha α -sp3		970	308	0
021	Gal β -sp4		1338	269	0
022	GlcNAc β -sp4		932	230	0
023	GalNAc β -sp4		1246	271	0
024	GlcNAc α -sp3		1463	260	2
025	GalNAc β -sp10		848	209	0
026	Rha β -sp4		1098	227	0

027	3,6-Me2Glc	DMG	673	123	0
028	Xyl β -sp4		1411	302	0
029	Fuc β -sp4		1139	301	0
030	Glc β -sp4		1449	429	1
031	L-Ara α -sp4		1099	369	0
032	GalNGc β -sp3		483	119	0
037	3-O-Su-Gal β -sp3		2317	503	3
038	3-O-Su-GalNAc β -sp3		3657	689	7
041	6-O-Su-GalNAc α -sp3		5504	1443	7
043	6-O-Su-GlcNAc β -sp3		1040	253	1
044	GlcA α -sp3	α -glucuronic acid	1275	327	0
045	GlcA β -sp3	β -glucuronic acid	1671	335	1
046	6-H ₂ PO ₃ Glc β -sp4	β -Glc6P	1369	316	0
047	6-H ₂ PO ₃ Man α -sp3	α -Man6P	1300	322	0
048	Neu5Ac α -sp3		583	131	0
049	Neu5Ac α -sp9		1349	334	3
050	Neu5Ac β -sp3		2184	526	3
051	Neu5Ac β -sp9		1698	609	3
052	Neu5Gc α -sp3		847	286	0
053	Neu5Gc β -sp3		1202	307	0
054	9-NAc-Neu5Ac α -sp3		256	77	0
055	3-O-Su-GlcNAc β -sp3		840	214	0
056	Gal α -sp7		1016	263	0
057	Rib -sp4		1153	248	0
058	Fuc β -sp3		2266	422	1
059	Crypted	Crypted	1360	389	0
060	6-O-Su-Gal β -sp3		61113	1156	20
061	3-OSu-GalNAc α -sp3		2692	622	3
071	Fuc α 1-2Gal β -sp3	H _{di}	1077	245	1
072	Fuc α 1-3GlcNAc β -sp3		1424	300	2
073	Fuc α 1-4GlcNAc β -sp3	Le	755	217	0
074	Fuc β 1-3GlcNAc β -sp3		533	130	0
075	Gal α 1-2Gal β -sp3		790	184	0
076	Gal α 1-3Gal β -sp3	B _{di}	775	192	0
077	Gal α 1-3GalNAc β -sp3	T _{$\alpha\beta$}	668	155	0
078	Gal α 1-3GalNAc α -sp3	T _{$\alpha\alpha$}	581	113	0
080	Gal α 1-3GlcNAc β -sp3		384	76	0
081	Gal α 1-4GlcNAc β -sp3	α -LN	815	183	1
082	Gal α 1-4GlcNAc β -sp8	α -LN	713	144	1
083	Gal α 1-6Glc β -sp4	melibiose	2285	348	2
084	Gal β 1-2Gal β -sp3		1332	301	1

086	Galβ1-3GlcNAcβ-sp2	Le ^C	609	130	0
087	Galβ1-3Galβ-sp3		1488	282	1
088	Galβ1-3GalNAcβ-sp3	T _{ββ}	1468	174	2
089	Galβ1-3GalNAcα-sp3	TF	297	59	0
092	Galβ1-4Glcβ-sp2	Lactose, Lac	889	182	0
093	Galβ1-4Glcβ-sp4	Lactose, Lac	1089	290	0
094	Galβ1-4Galβ-sp4		1124	278	0
096	Galβ1-4GlcNAcβ-sp2	N-acetyllactosamine, LN	2604	893	1
097	Galβ1-4GlcNAcβ-sp3	N-acetyllactosamine, LN	491	103	0
098	Galβ1-4GlcNAcβ-sp5	N-acetyllactosamine, LN	341	65	0
099	Galβ1-4GlcNAcβ-sp8	N-acetyllactosamine, LN	357	65	1
100	Galβ1-6Galβ-sp4		18406	2396	15
101	GalNAcα1-3GalNAcβ-sp3	Fs-2	1158	307	1
102	GalNAcα1-3Galβ-sp3	A _{di}	1199	295	1
103	GalNAcα1-3GalNAcα-sp3	core 5	451	87	0
104	GalNAcβ1-3Galβ-sp3		603	144	0
105	GalNAcβ1-3GalNAcβ-sp3	para-Fs	5975	1065	11
106	GalNAcβ1-4GlcNAcβ-sp3	LacdiNAc	332	79	0
107	GalNAcβ1-4GlcNAcβ-sp2	LacdiNAc	390	95	0
110	Glcα1-4Glcβ-sp3	maltose	10870	2172	10
111	Glcβ1-4Glcβ-sp4	cellobiose	3363	407	3
112	Glcβ1-6Glcβ-sp4	gentiobiose	6968	1229	5
113	GlcNAcβ1-3GalNAcα-sp3	core 3	4852	620	3
114	GlcNAcβ1-3Manβ-sp4		1474	423	2
115	GlcNAcβ1-4GlcNAcβ-Asn	chitobiose-Asn	12575	2669	14
116	GlcNAcβ1-4GlcNAcβ-sp3	chitobiose	1040	186	1
117	GlcNAcβ1-4GlcNAcβ-sp4	chitobiose	8585	1418	11
118	GlcNAcβ1-6GalNAcα-sp3	core 6	7860	714	4
119	Manα1-2Manβ-sp4		1200	259	0
120	Manα1-3Manβ-sp4		1224	338	0
121	Manα1-4Manβ-sp4		1064	328	0
122	Manα1-6Manβ-sp4		1484	380	0
123	Manβ1-4GlcNAcβ-sp4		2103	428	2
125	6-Bn-Galβ1-4GlcNAcβ-sp2		32674	5389	19
126	6-Bn-Galα1-4(6-Bn)GlcNAcβ-sp3		15410	3185	15
127	Galβ1-4Glcβ-dipeptide		559	105	0
128	Galβ1-4Glcβ-dipeptide		1181	263	0
129	Galβ1-3(6-O-Bn)GlcNAcβ-sp3		683	184	0
130	(6-O-Bn-Galβ1)-3GlcNAcβ-sp3		378	104	0
131	(6-O-Bn-Galβ1)-3(6-O-Bn)GlcNAcβ-sp3		1908	494	2
132	Galβ1-3GalNAcα-sp5	TF	376	72	0

133	Galβ1-4Glcβ-dipeptide		6816	465	4
134	Galβ1-4Glcβ-dipeptide		1197	271	0
135	Galβ1-4Glcβ-dipeptide		941	330	0
136	Galβ1-4Glcβ-dipeptide		476	98	0
137	Galβ1-4Glcβ-dipeptide		589	88	0
138	Galβ1-4Glcβ-dipeptide		571	130	0
139	Galβ1-4Glcα-sp3		395	65	0
140	Galα1-3GalNAc(fur)β-sp3		608	117	0
142	GlcNAcα1-3GalNAcβ-sp3		4523	1339	4
143	Fucα1-2(3-O-Su)Galβ-sp3		468	138	0
144	Galβ1-3(6-O-Su)GlcNAcβ-sp2		297	73	0
145	Galβ1-3(6-O-Su)GlcNAcβ-sp3		388	76	0
146	Galβ1-4(6-O-Su)Glcβ-sp2		2044	469	2
149	GlcNAcβ1-4(6-O-Su)GlcNAcβ-sp2		2569	475	5
150	3-O-Su-Galβ1-3GalNAcα-sp3		37943	3232	20
151	6-O-Su-Galβ1-3GalNAcα-sp3		1840	250	2
152	3-O-Su-Galβ1-4Glcβ-sp2	SM3	2564	413	2
153	6-O-Su-Galβ1-4Glcβ-sp2		1796	504	2
154	3-O-Su-Galβ1-3GlcNAcβ-sp3		32651	3954	20
156	3-O-Su-Galβ1-4GlcNAcβ-sp2		2305	458	1
158	4-O-Su-Galβ1-4GlcNAcβ-sp2		2927	860	2
159	4-O-Su-Galβ1-4GlcNAcβ-sp3		1656	736	2
160	6-O-Su-Galβ1-3GlcNAcβ-sp2		346	142	0
161	6-O-Su-Galβ1-3GlcNAcβ-sp3		536	141	0
162	6-O-Su-Galβ1-4GlcNAcβ-sp2		5615	1540	7
164	GlcAβ1-3GlcNAcβ-sp3		979	218	1
165	GlcAβ1-3Galβ-sp3		1035	308	1
166	GlcAβ1-6Galβ-sp3		39105	2993	17
167	GlcNAcβ1-4-[HOOC(CH ₃)CH]-3-O-GlcNAcβ-sp4	GlcNAc-Mur	1012	269	1
168	GlcNAcβ1-4Mur-L-Ala-D-i-Gln-Lys	GMDP-Lys	3668	152	1
169	Neu5Acα2-3Galβ-sp3	GM4	2305	596	3
170	Neu5Acα2-6Galβ-sp3		4157	633	3
171	Neu5Acα2-3GalNAcα-sp3	3-SiaT _n	844	168	1
172	Neu5Acα2-6GalNAcα-sp3	SiaT _n	3250	124	1
173	Neu5Acβ2-6GalNAcα-sp3	b-SiaT _n	3221	855	5
174	Neu5Gcα2-6GalNAcα-sp3	Neu5Gc-T _n	1596	178	1
175	Neu5Gcβ2-6GalNAcα-sp3		1453	451	2
176	3-O-Su-Galβ1-4(6-O-Su)Glcβ-sp2		43008	9342	20
177	3-O-Su-Galβ1-4(6-O-Su)GlcNAcβ-sp3		32496	7215	20
178	6-O-Su-Galβ1-4(6-O-Su)Glcβ-sp2		3506	763	5
179	6-O-Su-Galβ1-3(6-O-Su)GlcNAcβ-sp2		1770	580	0

180	6-O-Su-Galβ1-4(6-O-Su)GlcNAcβ-sp2		2991	832	4
181	3,4-O-Su ₂ -Galβ1-4GlcNAcβ-sp3		2993	820	5
182	3,6-O-Su ₂ -Galβ1-4GlcNAcβ-sp2		487	151	0
183	4,6-O-Su ₂ -Galβ1-4GlcNAcβ-sp2		4401	944	4
184	4,6-O-Su ₂ -Galβ1-4GlcNAcβ-sp3		1945	649	2
186	Neu5Acα2-8Neu5Acα2-sp3	(Sia) ₂	190	64	0
187	3-O-Su-Galβ1-4(6-O-Su)Glcβ-sp2		290	91	0
188	Neu5Acα2-8Neu5Acβ-sp9		1162	382	2
189	3,6-O-Su ₂ -Galβ1-4(6-O-Su)GlcNAcβ-sp2		2054	543	3
190	Galβ1-4-(6-P)GlcNAcβ-sp2		1217	347	1
191	6-P-Galβ1-4GlcNAcβ-sp2		926	194	1
192	GalNAcβ1-4(6-O-Su)GlcNAcβ-sp3		483	106	0
193	3-O-Su-GalNAcβ1-4GlcNAcβ-sp3		3103	626	4
194	6-O-Su-GalNAcβ1-4GlcNAcβ-sp3		2884	455	4
195	6-O-Su-GalNAcβ1-4-(3-O-Ac)GlcNAcβ-sp3		589	122	0
196	3-O-Su-GalNAcβ1-4(3-O-Su)-GlcNAcβ-sp3		605	123	1
197	3,6-O-Su ₂ -GalNAcβ1-4-GlcNAcβ-sp3		287	54	0
198	4,6-O-Su ₂ -GalNAcβ1-4GlcNAcβ-sp3		2251	407	1
199	4,6-O-Su ₂ -GalNAcβ1-4-(3-O-Ac)GlcNAcβ-sp3		690	178	0
200	4-O-Su-GalNAcβ1-4GlcNAcβ-sp3		466	134	0
201	3,4-O-Su ₂ -GalNAcβ1-4-GlcNAcβ-sp3		3790	859	6
202	6-O-Su-GalNAcβ1-4(6-O-Su)GlcNAcβ-sp3		4471	933	8
203	Galβ1-4(6-O-Su)GlcNAcβ-sp2		5424	1203	8
204	4-O-Su-GalNAcβ1-4GlcNAcβ-sp2		753	120	1
205	Neu5Acα2-6GalNAcβ-sp3		606	111	1
206	Neu5Gcα2-3Gal-sp3		1295	371	0
207	Neu5Acβ2-6GalNAcβ-sp3		2913	646	3
208	Galβ1-3GlcNAcβ-sp4	Le ^C	658	172	0
209	Crypted	Crypted	6503	1331	9
210	Crypted	Crypted	1743	558	1
211	Crypted	Crypted	8440	1533	13
212	Crypted	Crypted	2129	530	3
213	Neu5Acβ2-6Galβ-sp3		2672	827	3
214	ΔGlcAβ1-3Galβ-sp3	deltaGlcAβ3Gal	589	161	0
215	Fucα1-2Galβ1-3GlcNAcβ-sp3	Le ^d , H (type 1)	2727	444	3
216	Fucα1-2Galβ1-4GlcNAcβ-sp3	H (type 2)	10845	2895	15
217	Fucα1-2Galβ1-3GalNAcα-sp3	H (type 3)	3108	468	6
219	Fucα1-2Galβ1-4Glcβ-sp4	H (type 6)-Gly	8251	1687	10
220	Galα1-3Galβ1-4Glcβ-sp2		901	275	0
221	Galα1-3Galβ1-4Glcβ-sp4		1230	393	2
222	Galα1-3Galβ1-4GlcNAcβ-sp3	Galili (tri)	254	106	0
223	Galα1-4Galβ1-4Glcβ-sp2	P ^k , Gb3, GbOse ₃	756	247	0

225	Gal α 1-4Gal β 1-4GlcNAc β -sp2	P ₁	573	165	0
226	Gal α 1-3(Fuc α 1-2)Gal β -sp3	B _{tri}	1227	312	1
227	Gal α 1-3(Fuc α 1-2)Gal β -sp5	B _{tri} -C8	1044	248	1
228	Gal β 1-2Gal α 1-3GlcNAc β -sp3		321	87	0
229	Gal β 1-3Gal β 1-4GlcNAc β -sp4		323	81	0
230	Gal β 1-4Gal β 1-4Glc β -sp2	Mc3	248	84	0
232	Gal β 1-4GlcNAc β 1-6GalNAc α -sp3		624	81	1
233	Gal β 1-3(Fuc α 1-4)GlcNAc β -sp3	Le ^A	210	58	0
234	Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	Le ^X	13573	2635	16
235	GalNAc α 1-3(Fuc α 1-2)Gal β -sp3	A _{tri}	586	306	1
236	GalNAc α 1-3(Fuc α 1-2)Gal β -sp5	A _{tri}	498	262	1
237	GalNH α 1-3 Gal β -OCH ₂ CH ₂ CH ₂ NHAc Fuc α 1-2		357	135	0
239	GalNAc β 1-3(Fuc α 1-2)Gal β -sp3		849	654	1
240	(Glc α 1-4) ₃ β -sp4	maltotriose	11233	1567	6
241	(Glc α 1-6) ₃ β -sp4	isomaltotriose	520	163	0
242	GlcNAc α 1-3Gal β 1-4GlcNAc β -sp2		278	90	0
243	GlcNAc α 1-3Gal β 1-4GlcNAc β -sp3		215	61	0
245	GlcNAc α 1-6Gal β 1-4GlcNAc β -sp2		3394	1053	3
246	GlcNAc β 1-2Gal β 1-3GalNAc α -sp3		434	94	0
247	GlcNAc β 1-3Gal β 1-3GalNAc α -sp3		321	63	0
248	GlcNAc β 1-3Gal β 1-4Glc β -sp2		822	179	0
249	GlcNAc β 1-3Gal β 1-4GlcNAc β -sp2		1652	322	2
250	GlcNAc β 1-3Gal β 1-4GlcNAc β -sp3		1942	488	2
251	GlcNAc β 1-4Gal β 1-4GlcNAc β -sp2		1067	216	1
252	GlcNAc β 1-4GlcNAc β 1-4GlcNAc β -sp4	chitotriose	1111	206	2
253	GlcNAc β 1-6Gal β 1-4GlcNAc β -sp2		2877	653	3
254	GlcNAc β 1-6(Gal β 1-3)GalNAc α -sp3	core 2	2499	525	4
255	GlcNAc β 1-6(GlcNAc β 1-3)GalNAc α -sp3	core 4	5853	425	4
256	GlcNAc β 1-6(GlcNAc β 1-4)GalNAc α -sp3		5020	613	8
258	Man α 1-6(Man α 1-3)Man β -sp4	Man ₃	743	160	0
259	Gal β 1-4(Gal β 1-3)GlcNAc β -sp3		372	58	0
260	Fuc β 1-4 GlcNAc β -sp3		663	140	0
	Gal β 1-3				
261	Fuc β 1-3 GlcNAc β -sp3		299	62	0
	Gal β 1-4				
262	Gal β 1-3GalNAc β 1-3Gal-sp4		437	120	0
263	(GalNAc β -PEG2)3- β -DD		657	114	0
264	Gal β 1-4Gal β 1-4GlcNAc β -sp3		461	87	0

266	Gal α 1-4Gal β 1-4GlcNAc β -sp3	P ₁	1017	227	0
267	GlcNAc β 1-3Gal β 1-3GlcNAc β -sp3		718	138	0
268	GlcNAc β 1-4(Fuc α 1-6)GlcNAc β -sp3		1291	311	1
269	Gal β 1-3Gal β 1-4Glc β -sp4		518	137	0
270	Gal β 1-4Gal β 1-4Glc β -sp4		939	202	0
271	Gal β 1-6Gal β 1-4Glc β -sp4		53622	1934	20
272	Neu5Ac α 2-3Gal β 1-4Glc β -sp4-Cit	3'-SL -citrullin	236	60	0
273	Fuc β 1-2Gal β 1-4GlcNAc β -sp3		242	60	0
274	GalNAc α 1-3Gal β 1-4GlcNAc-sp3		426	90	0
275	GalNAc β 1-3Gal β 1-4GlcNAc-sp3		403	103	0
276	GlcNAc β 1-4Gal β 1-4GlcNAc β -sp3		1357	456	2
277	GalNGc α 1-3(Fuc α 1-2)Gal β -sp3	N-Gc-A _{tri}	375	99	0
287	3-O-Su-Gal β 1-3(Fuc α 1-4)GlcNAc β -sp3	3'-OSu-Le ^A	20063	4962	19
288	3-O-Su-Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	3'-OSu-Le ^X	593	146	0
289	Neu5Ac α 2-6(Gal β 1-3)GalNAc α -sp3		296	88	0
290	Neu5Ac α 2-6(Gal α 1-3)GalNAc α -sp3		262	68	0
291	Neu5Ac β 2-6(Gal β 1-3)GalNAc α -sp3		1887	727	3
292	Neu5Ac α 2-3Gal β 1-3GalNAc α -sp3		4477	1186	4
293	Neu5Ac α 2-3Gal β 1-4Glc β -sp3	3'SL	1684	349	1
294	Neu5Ac α 2-3Gal β 1-4Glc β -sp4	3'SL	363	99	0
295	Neu5Ac α 2-6Gal β 1-4Glc β -sp2	6'SL	290	66	0
296	Neu5Ac α 2-6Gal β 1-4Glc β -sp4	6'SL	360	96	0
297	Neu5Ac β 2-6Gal β 1-4Glc β -sp2		1645	409	3
298	Neu5Ac α 2-3Gal β 1-4GlcNAc β -sp3	3'SLN	5486	1855	3
299	Neu5Ac α 2-3Gal β 1-3GlcNAc β -sp3	3'SiaLe ^C	3968	1266	6
300	Neu5Ac α 2-6Gal β 1-4GlcNAc β -sp3	6'SLN	936	527	1
302	Neu5Ac β 2-6Gal β 1-4GlcNAc β -sp3		1100	368	2
303	Neu5Gc α 2-3Gal β 1-4GlcNAc β -sp3		2147	451	1
304	Neu5Gc α 2-6Gal β 1-4GlcNAc β -sp3		792	191	0
305	Neu5Gc β 2-6Gal β 1-4GlcNAc β -sp3		601	112	0
306	9-Nac-Neu5Ac α 2-6Gal β 1-4GlcNAc β -sp3		1431	437	1
307	KDN α 2-3Gal β 1-3GlcNAc β -sp2		1441	304	2
308	KDN α 2-3Gal β 1-4GlcNAc β -sp2		903	204	0
309	Neu5Ac α 2-6(Neu5Ac α 2-3)GalNAc α -sp3		487	107	1
310	3'SiaLacNAc β -OCH ₂ CH ₂ CH ₂ NH-(3'SiaLacNAc-amide-sp3)		1932	665	1
315	Neu5Ac α 2-3Gal β 1-4-(6-O-Su)GlcNAc β -sp3		1930	477	3
316	Neu5Ac α 2-3Gal β 1-3-(6-O-Su)GlcNAc β -sp3		2752	1014	5
318	Neu5Ac α 2-6Gal β 1-4-(6-O-Su)GlcNAc β -sp3		1004	234	1
319	Neu5Ac α 2-3-(6-O-Su)Gal β 1-4GlcNAc β -sp3		7963	705	7
320	4-O-Su-Neu5Ac α 2-3-(6-O-Su)Gal β 1-4GlcNAc β -sp3		9621	1580	10
321	(Neu5Ac α 2-8) ₃ -sp3	(Sia) ₃	226	57	0

322	(Neu5Ac α 2-8)3 β -sp3		553	112	0
323	Neu5Ac α 2-6Gal β 1-3GlcNAc-sp3	6'-SiaLe ^c	317	100	0
324	Neu5Ac α 2-6Gal β 1-3(6-O-Su)GlcNAc-sp3		361	80	0
325	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		275	57	0
326	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		311	65	0
327	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		495	125	0
328	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		298	55	0
329	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		586	133	0
331	Neu5Gc α 2-3Gal β 1-3GlcNAc β -sp3		2552	467	5
332	Neu5Ac α 2-3Gal β 1-4Glc β -dipeptide		381	91	0
333	Neu5Gc α 2-3Gal β 1-3-(6-O-Su)GlcNAc β -sp3		4811	1109	5
334	Neu5Gc α 2-3Gal β 1-4-(6-O-Su)GlcNAc β -sp3		1639	274	2
335	Neu5Ac α 2-3Gal β 1-3-(6-O-Su)GlcNAc β -sp3		6401	1102	6
336	α Kdo-(2 \rightarrow 8)- α Kdo-(2 \rightarrow 4)- α Kdo-sp11		5921	1649	10
337	GalNAc α 1-4Gal β 1-4GlcNAc β -sp3		4219	657	7
338	Neu5Ac α 2-6Gal β 1-3GalNAc α -sp3		306	59	0
339	Neu5Ac β 2-6Gal β 1-3GalNAc α -sp3		1931	365	3
340	Gal α 1-3(Neu5Ac β 2-6)GalNAc β -sp3		2907	760	4
341	Neu5Ac α 2-3-(6-Su)Gal β 1-4GlcNAc β -sp2		5964	1255	9
359	Gal α 1-3(Fuc α 1-2)Gal β 1-3GlcNAc β -sp3	B (type 1)	400	112	0
360	Gal α 1-3(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3	B (type 2)	5071	644	6
362	Gal α 1-3(Fuc α 1-2)Gal β 1-3GalNAc α -sp3	B (type 3)	8283	2150	14
363	Gal α 1-3(Fuc α 1-2)Gal β 1-3GalNAc β -sp3	B (type 4)	2728	589	6
364	Gal α 1-3Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	α GalLe ^x	541	116	0
365	Gal α 1-4(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3		962	361	1
366	GalNAc α 1-3(Fuc α 1-2)Gal β 1-3GlcNAc β -sp3	A (type 1)	351	71	0
368	GalNAc α 1-3(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3	A (type 2)	2621	625	2
369	GalNAc α 1-4(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3		3246	374	3
370	Crypted	Crypted	2629	835	3
371	Fuc α 1-2Gal β 1-3(Fuc α 1-4)GlcNAc β -sp3	Le ^B	307	107	0
372	Fuc α 1-2Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	Le ^Y	3936	1040	8
373	Gal α 1-3Gal β 1-4GlcNAc β 1-3Gal β -sp3	Galili (tetra)	777	128	0
374	Gal α 1-3(Gal α 1-4)Gal β 1-4GlcNAc β -sp3	Gal α 2-3',4'LN	280	72	0
375	Gal α 1-4GlcNAc β 1-3Gal β 1-4GlcNAc β -sp3		1032	190	1
377	Gal β 1-3GlcNAc β 1-3Gal β 1-3GlcNAc β -sp2		827	166	1
378	Gal β 1-3GlcNAc α 1-3Gal β 1-4GlcNAc β -sp3		10106	2009	9
379	Gal β 1-3GlcNAc β 1-3Gal β 1-4GlcNAc β -sp3		1304	387	1
380	Gal β 1-3GlcNAc α 1-6Gal β 1-4GlcNAc β -sp2		2306	344	1
381	Gal β 1-3GlcNAc β 1-6Gal β 1-4GlcNAc β -sp2		2581	827	3
382	Gal β 1-3GalNAc β 1-4Gal β 1-4Glc β -sp3	GA1, asialo-GM1	642	125	0
383	Gal β 1-4GlcNAc β 1-3Gal β 1-4Glc β -sp4	LNnT	280	74	0

384	Galβ1-4GlcNAcβ1-3Galβ1-4GlcNAcβ-sp2	i	290	76	0
386	Galβ1-4GlcNAcα1-6Galβ1-4GlcNAcβ-sp2		3822	638	2
387	Galβ1-4GlcNAcβ1-6Galβ1-4GlcNAcβ-sp2		2719	521	2
388	Galβ1-4GlcNAcβ1-6(Galβ1-3)GalNAcα-sp3		229	77	0
389	GalNAcβ1-3Galα1-4Galβ1-4Glcβ-sp3	Gb4, P	399	90	0
390	(Glcα1-4) ₄ β-sp4	maltotetraose	13504	2947	10
394	GlcNAcβ1-4				
	Galβ1-4GlcNAcβ-sp2		2286	487	3
	GlcNAcβ1-3				
395	GlcNAcβ1-6(GlcNAcβ1-3)Galβ1-4GlcNAcβ-sp2	T _k	1249	316	0
396	(GlcNAcβ1) ₃ -3,4,6-GalNAcα-sp3		4966	727	4
397	(GlcNAcβ1) ₃ -3,4,6-GalNAcα-sp3		7793	1480	9
398	Galβ1-3GlcN(Fm)β1-3Galβ1-4GlcNAcβ-sp3		557	106	0
399	Galβ1-3GlcNAcα1-3Galβ1-3GlcNAcβ-sp2		6788	1178	10
401	Galβ1-3GlcNAcβ1-3Galβ1-3GlcNAcβ-sp3		717	181	0
402	GalNAcα1-3				
	Galα1-4GlcNAcβ-sp3		175	45	0
	Fucα1-2				
403	Galβ1-3GlcNAcβ1-3Galβ1-4GlcNAcβ-sp2		2047	376	4
404	GalNAcα1-3Galβ1-4(Fucα1-3)GlcNAcβ-sp3	αGalNAcLe ^x	3611	1529	5
405	Galα1-3(Fucα1-2)Galα1-4GlcNAcβ-sp3		2929	227	1
406	GalNAcα1-3(Fucα1-2)Galα1-3GalNAcβ-sp3		304	60	0
	3-O-SuGalβ1-4GlcNAcβ1-3Galβ1-4GlcNAcβ-sp3		2425	448	3
419					
420	4-O-SuGalβ1-4GlcNAcβ1-3Galβ1-4GlcNAcβ-sp3		894	183	1
423	Neu5Acα2-3Galβ1-4(Fucα1-3)GlcNAcβ-sp3	SiaLe ^x	1673	483	1
425	Neu5Acα2-3Galβ1-4				
	GlcNAcβ-sp3		2164	673	2
	Fucβ1-3				
426	Neu5Acα2-3Galβ1-3(Fucα1-4)GlcNAcβ-sp3	SiaLe ^A	216	60	0
428	Neu5Acα2-3Galβ1-4(Fucα1-3)(6-O-Su-)GlcNAcβ-sp3	6-OSu-SLe ^x	429	149	0
429	Neu5Acα2-3(6-O-Su)Galβ1-4(Fucα1-3)GlcNAcβ-sp3	6'-OSu-SiaLe ^x	762	222	2
431	Neu5Acα2-3Galβ1-4(2-O-Su-Fucα1-3)GlcNAcβ-sp3		918	242	2
432	Neu5Acα2-3Galβ1-4				
	GlcNAcβ-sp3		546	207	0
	3-O-Su-Fucα1-3				
433	Neu5Acα2-6(Neu5Acα2-3Galβ1-3)GalNAcα-sp3		203	66	0
434	Neu5Acα2-8Neu5Acα2-3Galβ1-4Glcβ-sp4	GD3	287	84	0
435	Neu5Acα2-3Galβ1-4(2-O-Su-Fucα1-3)(6-O-Su-)GlcNAcβ-sp3		425	133	0

436	4-O-Su-Neu5Ac α 2-3Gal β 1-4(Fuc α 1-3)(6-O-Su-)GlcNAc β -sp3		963	415	2
437	GalNAc α 1-3(Fuc α 1-2)Gal β 1-3GalNAc β -sp3	A(type 4)	33886	3193	18
438	Fuc β 1-2Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3		460	131	0
439	α Kdo-(2 \rightarrow 4)- α Kdo-(2 \rightarrow 4)- α Kdo-(2 \rightarrow 6)- β GlcNAc-sp11		5179	1212	9
440	Neu5Ac β 2-6(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3		2150	416	3
441	Neu5Ac α 2-6(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3		719	160	1
442	Neu5Ac α 2-3 (GalNAc β 1-4)Gal β 1-4Glc β -sp4	GM2	499	111	0
479	Fuc α 1-2Gal β 1-3GlcNAc β 1-3Gal β 1-4Glc β -sp4	LNFP I	512	286	0
480	Fuc α 1-2Gal β 1-3GlcNAc β 1-3Gal β 1-4GlcNAc β -sp2	H(type 1) penta	3826	911	8
481	Gal α 1-3Gal β 1-4GlcNAc β 1-3Gal β 1-4Glc β -sp4	Galili (penta)	241	60	0
482	Gal α 1-3(Fuc α 1-2)Gal β 1-3(Fuc α 1-4)GlcNAc β -sp3	BLe ^B	1726	229	3
483	Gal α 1-3(Fuc α 1-2)Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	BLe ^Y	2689	293	3
484	GalNAc α 1-3(Fuc α 1-2)Gal β 1-3(Fuc α 1-4)GlcNAc β -sp3	ALe ^B	175	49	0
485	Gal β 1-4GalNAc α 1-3(Fuc α 1-2)Gal β 1-4GlcNAc β -sp3		2643	642	4
488	Gal β 1-4GlcNAc β 1-6(Gal β 1-4GlcNAc β 1-3)GalNAc α -sp3		284	69	0
489	Gal β 1-4GlcNAc β 1-3(GlcNAc β 1-6)Gal β 1-4GlcNAc β -sp2		541	172	0
490	Gal β 1-4GlcNAc β 1-6(GlcNAc β 1-3)Gal β 1-4GlcNAc β -sp2		527	119	0
491	GalNAc α 1-3(Fuc α 1-2)Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3	ALe ^Y	213	65	0
492	(Glc α 1-6) ₅ β -sp4	isomaltopentaose	1151	263	2
493	(GlcNAc β 1-4) ₅ β -sp4	chitopentaose	1798	509	1
495	Man α 1-6(Man α 1-3)Man α 1-6(Man α 1-3)Man β -sp4	Man ₅	892	319	1
496	Fuc α 1-2Gal β 1-3(Fuc α 1-4)GlcNAc β 1-3Gal β 1-4Glc β -sp4	Le ^b (hexa)	223	72	0
497	Fuc α 1-2Gal β 1-4(Fuc α 1-3)GlcNAc β 1-3Gal β 1-4Glc β -sp4	Le ^Y (hexa)	2057	212	4
498	Gal β 1-4GlcNAc β 1-3Gal β 1-4GlcNAc β 1-3Gal β 1-4GlcNAc β -sp3	(LN) ₃	201	62	0
499	Gal β 1-4GlcNAc β 1-6(Gal β 1-4GlcNAc β 1-3)Gal β 1-4GlcNAc β -sp2	I	200	45	0
501	Gal β 1-3GalNAc β 1-3Gal α 1-4Gal β 1-4Glc β -sp4	Gb5	1360	254	1
502	(Glc α 1-6) ₆ β -sp4	maltohexaose	3033	653	5
503	(GlcNAc β 1-4) ₆ -sp4	chitohexaose	2396	314	1
504	(A-GN-M) ₂ -3,6-M-GN-GN β -sp4	9-OS	200	57	0
505	(GN-M) ₂ -3,6-M-GN-GN β -sp4	7-OS	728	195	0
506	Araf6	Ara6	158	37	0
507	GalNAc α 1-3GalNAc β 1-3Gal α 1-4Gal β 1-4Glc β -sp3	Fs-5	2021	988	3

508	GalNAc β 1-3(Fuc α 1-2)Gal β 1-4(Fuc α 1-3)GlcNAc β -sp3		402	93	0
509	Gal β 1-3GalNAc β 1-4(Neu5Ac α 2-3)Gal β 1-4Glc β -sp4	GM1	1553	448	3
527	Neu5Ac α 2-3Gal β 1-4GlcNAc β 1-3Gal β 1-4GlcNAc β -sp2		179	48	0
528	Neu5Ac α 2-3Gal β 1-4(Fuc α 1-3)GlcNAc β 1-3Gal β -sp3	SiaLe ^x -3Gal	203	49	0
529	Neu5Ac α 2-6(Gal β 1-3)GlcNAc β 1-3Gal β 1-4Glc β -sp4	LSTb	297	70	0
530	(Neu5Ac α 2-3Gal β 1)2-3,4-GlcNAc β -sp3		1529	352	1
531	Neu5Ac α 2-8Neu5Ac α 2-3(GalNAc β 1-4)Gal β 1-4Glc β -sp2	GD2	434	116	0
534	Neu5Ac α 2-3Gal β 1-4GlcNAc β 1-3Gal β 1-4GlcNAc β -sp3		193	58	0
535	Neu5Ac α 2-8Neu5Ac α 2-3(GalNAc β 1-4)Gal β 1-4Glc-sp4	GD2	234	61	0
536	Neu5Ac α 2-3Gal β 1-3GlcNAc β 1-3Gal β 1-4Glc β -sp4	LSTa	342	92	0
537	Neu5Ac α 2-3Gal β 1-4GlcNAc β 1-3Gal β 1-4Glc β -sp4	LSTd	202	61	0
538	Gal β 1-4(Fuc α 1-3)GlcNAc β 1-6(Gal β 1-3GlcNAc β 1-3)Gal β 1-4Glc β -sp4	MFLNH III	2473	712	3
539	Gal β 1-4GlcNAc β 1-6(Fuc α 1-2Gal β 1-3GlcNAc β 1-3)Gal β 1-4Glc β -sp4	MFLNH I	780	239	0
540	Gal β 1-4(Fuc α 1-3)GlcNAc β 1-6(Neu5Ac2-6Gal β 1-4GlcNAc β 1-3)Gal β 1-4Glc β -sp4	MSMFLNnH	2262	337	3
541	Gal β 1-4(Fuc α 1-3)GlcNAc β 1-6(Fuc α 1-2Gal β 1-3GlcNAc β 1-3)Gal β 1-4Glc β -sp4	DFLNH (a)	6347	878	7
542	Gal β 1-3GlcNAc β 1-3Gal β 1-4(Fuc α 1-3)GlcNAc β 1-6(Gal β 1-3GlcNAc β 1-3)Gal β 1-4Glc β -sp4	MF(1-3)iLNO	2867	344	6
543	Fuc α 1-2Gal β 1-3(Fuc α 1-4)GlcNAc β 1-3[Gal β 1-4(Fuc α 1-3)GlcNAc β 1-6]Gal β 1-4Glc β -sp4	TFLNH	3573	657	6
545	α Kdo-(2 \rightarrow 8)- α Kdo-(2 \rightarrow 4)- α Kdo-(2 \rightarrow 6)- β GlcNAc-(1 \rightarrow 6) α GlcNAc-sp11		3329	757	6
625	(GlcA β 1-4GlcNAc β 1-3) ₁₁₋₁₂ -NH ₂ -ol	hyaluronic acid	776	257	1
627	(Sia2-6A-GN-M) ₂ -3,6-M-GN-GN β -sp4	11-OS, YDS	184	54	0
629	Trehalose-ethanolamine	negative control	323	71	0
630	(GlcA β 1-3GlcNAc β 1-4) ₂₀ -NH(<i>p</i> -C ₆ H ₄)CH ₂ CH ₂ NH ₂	hyaluronic acid , 8kDa	921	236	1
631	(GlcA β 1-3GlcNAc β 1-4) ₃₈ -NH(<i>p</i> -C ₆ H ₄)CH ₂ CH ₂ NH ₂	hyaluronic acid, 15-30kDa	456	174	1
632	(GlcA β 1-3GlcNAc β 1-4) ₁₃ -NH(<i>p</i> -C ₆ H ₄)CH ₂ CH ₂ NH ₂	hyaluronic acid, 4-8kDa	481	109	0
800	GlcNAc α 1-4GlcNAc β -sp3		1697	257	1
801	GalNAc α 1-3GalNAc(fur) β -sp3		3988	289	2
802	Gal β 1-3GalNAc(fur) β -sp3		51348	2324	20
804	[Gal β 1-4GlcNAc β -OCH ₂ CH ₂] ₂ NH	LN dimer	908	252	1
805	GalNAc β 1-4(6-O-Bn)GlcNAc β -sp3		6754	1034	11
806	Gal α 1-6Glc α -sp3	α -melibiose	5469	949	4

807	GlcNAc β 1-4GlcNAc α -sp4		10158	984	8
808	Gal α 1-6Glc β -sp3	melibiose	9724	1131	7
809	GalNAc β 1-3GalNAc α -sp3		8521	1509	13
810	GalNGc α 1-3GalNAc α -sp3	core 5(Gc)	328	76	0
850	Gal β 1-3(6-O-Su)GalNAc α -sp3	6-SuTF	341	102	0
851	Gal α 1-3(6-O-Su)GalNAc α -sp3	6-SuT $_{\alpha\alpha}$	267	86	0
852	GlcNAc β 1-4-[HOOC(CH ₃)CH]-3-O-GlcNAc α -sp4	GlcNAc-Mur α	1056	322	1
900	H-(Gly) ₆ -NH ₂ Gly6-amide, linear	Gly ₆ -amide, linear	25036	2262	16
901	biot-CMG ₂ -NH ₂	biot-CMG₂	52213	3728	20
902	Peptide, crypted		5630	1228	6
903	Peptide, crypted		2617	897	4
904	Peptide, crypted		6111	1154	12
905	Suc-LPWYRAPK-NH ₂	RhD	16175	3630	20
906	Peptide, crypted		475	163	0
907	Peptide, crypted		457	93	0
908	Peptide, crypted		6687	2162	13
909	Peptide, crypted		812	166	1
910	Npentyl-Indol-linked-3-NH ₂ (AcOH salt)		686	168	0
911	N-C5-indazole-3-CONH-PEG-NH ₂ (AcOH salt)		6037	1212	11

sp2 = -O(CH₂)₂NH₂

sp3 = -O(CH₂)₃NH₂

sp4 = -NHCOCH₂NH₂

sp5 = -O(CH₂)₃NH-CO(CH₂)₅NH₂

sp8 = -(OCH₂CH₂)₆NH₂