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Identification of sugar moieties in chief cells of the rat fundic gastric

glands

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Running title: Sugar moieties of gastric chief cells

#### **Abstract**

Many studies have been conducted to determine the composition of the glyconjugates of the mucus-secreting cells of the fundic glands of the stomach. However, the chief cells of these glands have been largely ignored because they secrete mainly zymogens with a lower glycosylation. The aim of this work was to analyze the glycoconjugates of the gastric chief cells by a battery of 17 different lectins, recognizing Fucose, N-AcetylGalactosamine, Galactose, N-Acetylneuramiinic acid, N-AcetylGlucosamine and Mannose containing oligosaccharides.

Histochemical techniques were performed with several lectins and also combined with two pre-treatments;  $\beta$ -elimination, which removes O-linked oligosaccharides, and incubation with Peptide-N-Gycosidase F, which removes N-linked oligosaccharides. In addition, acid hydrolysis was performed before WGA histochemistry, and incubation with glucose oxidase before Con A labeling.

Many lectins did not stain the chief cells. In addition, the presence of O-glycans in the apical cell membrane was demonstrated with the lectins AAL, HPA, MPA/MPL, PNA, RCA-I, and WGA. Some of these O-glycans were resistant to short-term β-elimination pre-treatments. Mannose-binding lectins stained the basal cytoplasm of the chief cells. The level of glycosylation of the chief cells was lower than that of the mucous cells. The presence of O-glycans in the apical cell membrane is consistent with the presence of mucins such as MUC1 in the apical membrane of chief cells. Moreover, Mannose-binding lectins revealed N-glycosylation in the basal cytoplasm. The knowledge of gastric chief cell glycoconjugates is relevant because of their potential involvement not only in in physiological but also in pathological processes, such as cancer.

# Miniabstract

Lectin histochemistry of gastric chief cells showed a low level of glycosylation, identified O-glycans in the apical membrane (attributed to MUC1), and revealed N-glycosylation processes in the basal cytoplasm.

# **Keywords:**

Gastric glands, glycoconjugates, lectin histochemistry, mucins, zymogens.

#### Introduction

The study of glycoconjugates has been relatively uncommon for a long time, probably due to their great variety and structural complexity. However, in recent years, the analysis of glycans present in glycoproteins and glycolipids –called glycobiology– has become a rapidly growing field in the life sciences, with a great relevance in many areas of basic research, biomedicine and biotechnology (Varki and Kornfeld, 2017). In addition, glycans have been shown to play a large variety of biological roles, from the very subtle to those that are essential for development, growth, or organism survival. Moreover, glyconjugates have important roles in pathological processes, including cancer and infection by microorganisms (Varki and Gagneux, 2017; Varki and Kornfeld, 2017; Varki et al., 2017). Glycans often have a complex structure that greatly hinders their study. Combinations of various techniques (physical, chemical and enzymatic) are frequently used to analyze them, but very interesting and basic information on the composition of glycans can be obtained *in situ* by means of lectins or other glycan-binding proteins and antibodies (Cummings et al., 2017; Mulloy et al., 2017).

The cells of the gastric mucosa produce a glycoconjugate-rich secretion that has attracted the attention of researchers (Holmen Larsson et al., 2013; Rodríguez-Piñeiro et al., 2013; Jin et al., 2017). The rat gastric mucosa is divided into several regions. The *corpus* is the medial region and contains the fundic tubular glands with several types of secretory cells that arise from stem cells. Some of them are mucus-secreting cells. Others, the zymogenic or chief cells, located at the bottom of the gland, are serous cells that secrete digestive enzymes (Hoffmann, 2013). Several lectin histochemical studies have been performed to analyze the glycoprotein composition of these cells and their secretion, mainly for the mucous cells (Madrid et al., 1990, 1998; Gómez-Santos et al.,

treatments allows not only to find out the carbohydrate residues present in the glycoproteins, but also the type of oligosaccharide they belong to, i.e., N- or O-glycan (Brooks et al. 2012). Mucous cells have been studied more extensively as they secrete highly glycosylated mucins, a very diverse family of O-glycosylated proteins. However, the chief cells produce mainly a zymogenic secretion, formed by proteins with a very low or absent glycosylation level, although they also express some mucins. The characterization of the fundic gland glycoconjugates is interesting because of their important role in normal physiological processes, in development and in pathological processes, such as gastric cancer (Su et al., 2004; Duarte et al. 2016). Notably, mucins, O-glycosylated proteins produced in the gastric glands, are involved in the development of gastric cancer. The MUC1 protein, located in the apical membrane of chief cells, is known to be in the binding of *Helycobacter pilori* to the gastric mucosa, which can produce chronic gastritis and result in gastric cancer. The most common types of gastric cancer are also known to arise from chief cells (Duarte et al., 2016; Mills and Goldenring, 2017). MUC1 is also known to be an oncogene with an anti-apoptotic function in cancer cells, but its MUC1 protein product has also been suggested to play a protective role against inflammation and carcinogenesis in the normal gastric mucosa (Saeki et al., 2014). However, there is no report analyzing the glycan composition of chief cells using lectins combined with deglycosylation pre-treatments. This would allow a better characterization of the presence of glycoconjugates even at a subcellular level, showing precisely where they are located.

2018). The combination of lectin histochemistry with various deglycosylation pre-

The aim of this work was to analyze the presence of glyconjugates in the chief cells of rat fundic glands by means of histochemistry with a battery of 17 lectins combined with several deglycosylation pre-treatments.

#### **Material and methods**

#### Reagents

We purchased several reagents from EY (San Mateo, CA): biotinylated *Aleuria* aurantia lectin (AAL), *Galanthus nivalis* agglutinin (GNA), *Glycine max* agglutinin (SBA), *Ulex europaeus* agglutinin-I (UEA-I), *Canavalia ensiformis* agglutinin (Concanavalin A, Con A), and peroxidase-labeled agglutinin from *Limax flavus* (LFA). We used some other reagents from Sigma Aldrich (Spain): type III glucose oxidase from *Aspergillus niger*, Bovine serum albumin (BSA), 3,3'-diaminobenzidine (DAB), peroxidase-labeled agglutinins from *Helix pomatia* (HPA) and *Maclura pomifera* (MPA/MPL), and biotinylated *Arachis hypogaea* (PNA) and *Lotus tetragonolobus* (LTA) agglutinins.

Roche (Spain) supplied the following reagents: the enzyme Peptide-*N*-glycosidase F (PNGase F) from *Flavobacterium meningosepticum* and expressed in *Escherichia coli*, peroxidase-labeled anti-digoxigenin antibody (HRP-anti-DIG), digoxigenine-labeled *Datura stramonium* (DSA) and *Sambucus nigra* (SNA) agglutinins.

In addition, we purchased other reagents from Atom (Barcelona, Spain): avidin-biotin-peroxidase complex (Vectastain ABC kit peroxidase standard), Avidin-biotin blocking kit, biotinylated *Maackia amurensis* haemagglutinin (MAH), *Bandeiraea simplicifolia* lectin I-B<sub>4</sub> (BSI-B<sub>4</sub>), *Ricinus communis* agglutinin-I (RCA-I), *Dolichos biflorus* (DBA) and *Triticum vulgaris* (WGA) agglutinins.

#### Samples

We used some paraffin-embedded samples obtained during the years 2002-2003 from six adult male Sprague-Dawley rats weighing between 250-300 g, that were previously

employed in other studies (Gómez-Santos et al., 2007, 2018). We also used samples of gall bladder, testis and intestine from our archives for the controls of deglycosylation techniques (Madrid et al., 1994). At the time we obtained the samples, we followed all the applicable international, national and institutional guidelines for the care and use of experimental animals. Samples were stored at room temperature in the dark until use.

#### Histochemical techniques

The lectins used are listed in table 1. Their specificities were indicated in one of our previous works (Gómez-Santos et al., 2018) and are summarized in table 1. We obtained 4 µm-thick sections from the paraffin embedded samples. After removal of paraffin and hydration, we blocked the endogenous peroxidase and carried out the histochemical techniques. In addition, when digoxigenin- or biotin-labeled lectins were used, the nonspecific binding of antibodies was blocked by incubating the samples with BSA. All the procedures were performed as previously described (Gómez-Santos et al., 2018). The lectin dilutions were as follows: 6µg/ml HPA, 20µg/ml MPA/MPL, 25µg/ml LFA, 10µg/ml DSA, 30µg/ml SNA, 5µg/ml RCA-I, WGA and MAH, 3µg/ml BSI-B4, 50µg/ml DBA, PNA, LTA and SBA, 10µg/ml AAL, UEA-I and Con A, and 60µg/ml GNA.

Peroxidase-labeled lectins were developed with DAB. When Digoxigenin-labeled lectins were used, samples were incubated with 0.6U/ml HRP-anti-DIG antibody and then the peroxidase was developed. The samples treated with biotin-labeled lectins were incubated with ABC kit before peroxidase development. Finally, we counterstained the sections with hematoxylin.

#### Deglycosylation pretreatments

In addition to performing histochemical techniques with each lectin, in other tissue sections the same techniques were performed after different deglycosylation procedures. The deglycosylation procedures used were chemical deglycosylation by  $\beta$ -elimination, which removes O-linked oligosaccharides (Ono et al., 1983), and enzymatic deglycosylation with PNGase F, which removes N-linked oligosaccharides. We performed the  $\beta$ -elimination procedure for every lectin in different sections for 1 and 5 days, to discriminate both labile and resistant O-glycans (Gómez-Santos et al., 2007). In these procedures, unpublished observations have shown an increase of endogenous biotin labeling by ABC kit in gastric gland cells, so in these cases it is also necessary to block endogenous biotin. The detailed procedure for  $\beta$ -elimination and PNGase F incubation, including blocking of endogenous biotin, has been previously described (Martínez-Menárguez et al., 1993; Gómez-Santos et al., 2018).

To determine specifically which carbohydrate was labeled by WGA, a lectin that recognizes both N-Acetylglucosamine (GlcNAc) and sialic acid (NeuAc), we also tested the lectin after removal of terminal NeuAc by acid hydrolysis. This was carried out by immersing the sections in 0.1N HCl at 82-84°C for 3h before the lectin histochemical procedure (Madrid et al., 1994).

ConA can label Glucose (Glc) and Mannose (Man), thus in some sections we made a pre-incubation with 50U/ml type III Glucose oxidase at 37°C overnight to convert Glc into gluconic acid, which is not recognized by Con A (Alonso et al., 2006).

#### Controls

We used controls for all the procedures as previously reported (Gómez-Santos et al., 2018). Briefly, they were the following: 1) substitution of the lectins or the enzyme by buffer alone, 2) preincubation of the lectins with the corresponding hapten sugar

inhibitor at a concentration of 0.2M, and 3) staining of sections of other tissues of known altered binding pattern for each of the chemical and enzymatic pretreatments. As specific control of the β-elimination procedure, we combined the deglycosylation treatment with HPA staining of rat testis sections, because this tissue loses the HPA-labeling when β-elimination works correctly (Martínez-Menárguez et al., 1993). For deglycosylation with PNGase F, the control consisted of verifying the absence of labeling in sections of rat testicle with AAL, since incubation with PNGase F prevents the binding of this lectin (Martínez-Menárguez et al., 1993).

Semi-quantitative evaluation of the staining

For each lectin and each deglycosylation procedure, three samples were evaluated by three independent observers. They classified the staining intensity in the chief gastric cells into six categories: no labeling (0), very weak (1), weak (2), moderate (3), strong (4), and very strong (5). Sometimes a different staining intensity was observed in different sections or in different fields of the same section. When this happened, the staining intensity was classified in a range indicated with the minimum and maximum values.

#### Results

We analyzed the expression of glyconjugates in the chief cells of gastric glands in the rat stomach by means of lectin histochemistry. For each lectin, in addition to the lectin histochemical staining in some histological sections, the same staining was performed in other sections after different deglycosylation methods: the  $\beta$ -elimination technique, which removes O-linked oligosaccharides, and incubation with the enzyme PNGase F, which removes N-linked oligosaccharides. For the lectin WGA, removal of NeuAc was performed by acid hydrolysis with HCl. Moreover, for the lectin Con A some sections were preincubated with the enzyme glucose oxidase, which converts Glc into gluconic acid.

We used 17 lectins that recognize the most abundant sugar moieties in glycoconjugates, i.e., Fucose (Fuc), N-Acetylgalactosamine (GalNAc), Galactose (Gal), NeuAc, GlcNAc and Mannose (Man). The results are shown in table 2. Below, we highlight the most important by grouping the lectins according to the sugar that they recognize. In contrast to other neighboring cell types, such as parietal cells, none of the Fucrecognizing lectins labeled the cytoplasm of the chief cells. However, AAL clearly labeled the apical surface of these cells. This labeling remained after incubation with PNGase F, but pretreatment of the sections with a  $\beta$ -elimination protocol completely removed it (Fig. 1).

The only GalNAc-labeling lectin which labeled the cytoplasm of the chief cells was MPA/MPL. This lectin strongly labeled the supranuclear cytoplasm, corresponding to the area occupied by the secretory vesicles, in a lattice-like pattern. In addition, the apical surface of the cells was strongly labeled. The labeling was unaltered by pretreatment with PNGAse F and disappeared after \(\beta\)-elimination for 5 days. Chief cells were negative for the remaining GalNAc-binding lectins, except HPA, which labeled

the apical membrane of the cells. This labeling remained after PNGase F pretreatment but was not longer observed after  $\beta$ -elimination for 1 day procedure. (Fig. 2).

The cytoplasm of the chief cells was negative for most Gal-labeling lectins (PNA, BSI-B4 and RCA-I). A weak cytoplasmic labeling was seen with PNA, but this was not detectable when  $\beta$ -elimination was previously carried out. Furthermore, both PNA and RCA-I labeled the apical surface of the chief cells, but this staining disappeared after  $\beta$ -elimination in both cases. Moreover, the chief cells showed a very weak lattice-like cytoplasmic labeling when RCA-I was performed after  $\beta$ -elimination for 1 day, but not after 5 days (Fig. 3).

LFA was the only NeuAc-binding lectin showing positive labeling in the chief cells. Labeling was detected at the supranuclear cytoplasm and it was resistant to incubation with PNGase F or  $\beta$ -elimination. The chief cells were negative for MAH and SNA, except when MAH staining was performed after  $\beta$ -elimination for 5 days, when a very weak lattice-like labeling could be seen (Fig. 4).

WGA labeled the cytoplasm and the apical surface of the chief cells. This labeling was also resistant to most of the deglycosylation pre-treatments employed (including acid hydrolysis), except for  $\beta$ -elimination for 5 days, which reduced cytoplasmic staining and abolished apical surface staining. However, DSA, the other lectin that recognizes GlcNAc, very weakly labeled the chief cells, and this labeling was also resistant to  $\beta$ -elimination. After PNGase-F procedure the apical surface became very slightly positive (Fig. 5).

Finally, both Man-binding lectins, GNA and Con A, moderately labeled the cytoplasm of the chief cells. The staining was more intense in the basal than in the supranuclear cytoplasm. When lectin histochemistry was performed after incubation with PNGase F, no labeling was seen in either case. When samples were pre-treated with Glucose-

oxidase, the basal cytoplasm Con A-labeling was absent or very weak, but the supranuclear cytoplasm and the apical surface of the chief cells appeared labeled instead (Fig. 6).

#### **Discussion**

In this work we examined the expression of glycoconjugates in the rat gastric chief cells by means of lectin histochemistry alone and in combination with deglycosylation pretreatments: β-elimination to remove O-linked oligosaccharides (Ono et al., 1983) and incubation with PNGase F to remove N-linked glycans (Martínez-Menarguez et al., 1993). Beta-elimination was carried out for each lectin in different sections for 1 and 5 days, to discriminate between labile and resistant O-linked oligosaccharides (Gómez-Santos et al., 2007). In the case of WGA histochemistry, it was also carried out after acid hydrolysis with HCl, which removes NeuAc, to distinguish whether this lectin recognized GlcNAc or NeuAc (Madrid et al., 1994). Finally, Con A, which can bind to Man or Glc, was also tested after incubation with glucose oxidase to convert Glc into gluconic acid, which is not labeled by the lectin (Alonso et al., 2006).

We can simplify the results obtained with lectins into three labeling patterns: 1) lectin did not label any cell structure, 2) lectin stained relatively large areas of the internal cytoplasm, and 3) the apical cell surface was positive for lectin. The last two staining patterns could be observed with some of the 17 tested lectins.

The first observed pattern, i.e. the absence of labeling, was shown by several lectins that recognize Fuc, GalNAc, Gal or NeuAc. Other lectins that also recognize these sugars showed one or both of the other two labeling patterns before mentioned. This discrepancy between lectins recognizing the same carbohydrate can be explained by the different affinities displayed by each of the lectins towards different carbohydrate sequences summarized in Table 1.

The second staining pattern, the labeling of the internal cytoplasm, with stronger or weaker intensity, was observed with only a few of the lectins that bind to Fuc, GalNac,

Gal or NeuAC and with all those that recognize GlcNAc and Man. Among the lectins that recognize Fuc, GalNAc, Gal or NeuAc, only MPA/MPL and LFA stained the internal cellular cytoplasm with some intensity. Labeling with MPA/MPL disappeared after β-elimination for 5 days, indicating the presence of GalNAc in resistant O-linked oligosaccharides (Gómez-Santos et al., 2007). The LFA staining did not disappear with any of the deglycosylation pre-treatments, which indicates that NeuAc may be present in both N- and O-linked oligosaccharides. Both MPA/MPL and LFA labeling showed a lattice-like pattern in the supranuclear cytoplasm, which corresponds to the cell region typically occupied by the secretory granules.

WGA can label to GlcNAc or NeuAC. Removal of NeuAc by acid hydrolysis did not alter the WGA-labeling suggesting that WGA is recognizing GlcNAc. This labeling decreased but did not completely disappear after  $\beta$ -elimination. On the contrary, the weak labeling seen after histochemistry with DSA slightly increased after  $\beta$ -elimination. All these results suggest that lectins are recognizing different GlcNAc moieties, some in N- and other in O-linked oligosaccharides.

We used two lectins, GNA and Con A, that recognize Man. Con A can also bind to Glc. Both lectins stained the basal and supranuclear cytoplasm of the chief cells. When histochemistry was performed with Con A after incubation with glucose oxidase, the cells remained positive, with minimal differences in the pattern of labeling, suggesting that Con A may be identifying mostly Man residues. Although most Man residues are usually located in N-linked oligosaccharides, Man has also been described in O-linked oligosaccharides (Varki and Kornfeld, 2017). Our results showed that removal of the N-linked oligosaccharides by PNGase F incubation turned the gastric chief cells negative for the two Man-labeling lectins, indicating that most of the Man in chief cells was located in N-linked oligosaccharides.

Most lectins that labeled the cytoplasm stained the supranuclear region, occupied by the Golgi apparatus and secretory granules. Usually, this labeling showed a lattice-like staining pattern, which was apparently due to the fact that the cytoplasm surrounding the secretion granules was positive but the granules themselves were negative. Comparing the results obtained in the chief cells with those obtained in a previous work on the mucus-producing cells of the gastric glands, it can be pointed out that the chief cells show a lower content of oligosaccharides (Gómez-Santos et al., 2018). In addition, lectins usually label the secretion granules in mucous cells, whereas in the chief cells the granules were rarely stained. Surely, this is related to the type of secretion produced by each cell type. Whereas mucous cells have a mucin-rich secretion, which are highly glycosylated proteins, the chief cells secrete mostly zymogens, which are proteins with few or no glycosylation and once secreted they are converted into digestive enzymes (Baudyš and Kostka, 1983; Dekker et al., 1989; Kageyama 2002; Holmén Larsson et al., 2013). The degree of glycosylation of zymogenic proteins from the gastric chief cells is diverse. Some of them are somewhat glycosylated, whereas others lack glycans. Even the same protein can be glycosylated in some species and not glycosylated in others (Kageyama and Takahashi, 1978; Green and Roberts, 2004; Hassam et al., 2010). One of the highly glycosylated zymogen proteins secreted by gastric chief cells is gastric lipase (Barrowman et al., 1970; Moreau et al., 1989). In particular, it has been suggested that 10-15% of the human gastric lipase mass is due to oligosaccharides, probably N-glycans (Wicker-Planquart et al., 1999). Since lectins recognizing mannose stained the cytoplasm surrounding the granules, but not the granules themselves, we cannot conclude that the staining is due to recognition of gastric lipase. Zymogenic protein expression by chief cells is known to be clinically relevant. For example, the expression of progastriscin, one of the main proteolytic enzymes present in the gastric

fluid, has been shown to gradually decrease from the healthy stomach to benign lesions to precancerous lesions, and finally to gastric cancer (Hassan et al., 2010).

Only the Man-recognizing lectins labeled with some intensity the basal cytoplasm, where the rough endoplasmic reticulum is typically located in chief cells. These results are consistent with the fact that mannosylation occurs mostly in the rough endoplasmic reticulum, whereas other sugar moieties are added in the Golgi apparatus, (Stanley et al., 2017).

Finally, some lectins (AAL, HPA, MPA/MPL, PNA, RCA-I and WGA) labeled the apical surface of the chief cells, probably because they were bound to glycoconjugates of the apical cell membrane. This labeling was absent when O-linked oligosaccharides were removed before lectin staining. This is consistent with what was previously observed in immunohistochemical studies to identify mucin 1 (MUC1). These studies showed a diffuse staining in the cytoplasm, which was more distinguishable in the apical membrane of the cells (Saeki et al. 2014). MUC1 is a transmembrane protein that has a VNTR (variable number of tandem repeat) region rich in serine, threonine and proline that is highly O-glycosylated. O-linked oligosaccharides of mucins can be diverse. The simplest mucin O-glycan is a single N-acetylgalactosamine linked to serine or threonine, which can be recognized by MPA/MPL. The most common O-linked oligosaccharide in mucins is Galβ(1-3)GalNAc, which can be labeled by HPA, MPA/MPL and PNA. Other common O-glycans in mucins can be recognized by lectins that labeled the apical cell membrane of chief cells (Brockhausen and Stanley 2017). MUC1 has a prominent role as a barrier against external insults to the gastric cells, but its expression changes during pathological events: the MUC1 gene is silenced in intestinal metaplasia, a pre-neoplastic lesion, but is reactivated in gastric cancer. Furthermore, MUC1 is regarded as an oncoprotein because it suppresses the

transcription of the p53 gene, a major tumor suppressor gene (Wei et al., 2007; Saeki et al., 2014). Our lectin-labeling results are consistent with the presence of MUC1 in the apical membrane of chief gastric cells. Moreover, this lectin histochemistry method could be clinically used as a prognostic tool to monitor early changes in the expression of mucins in the gastric epithelial surface, and correlate these changes with the development of cellular events contributing to gastric cancer. Experiments with gastric cancer patient samples have already pointed out an aberrant expression of glycoconjugates and the altered binding to some lectins as potential prognostic factors for this disease (Li et al. 2016).

In conclusion, we have demonstrated the presence of different glycoconjugates in the chief cells of the fundic glands of the rat stomach by means of a battery of 17 lectins, coupled to different deglycosylation treatments. Many of the lectins did not stain chief cells, and only a few of them stained some regions of the cytoplasm, suggesting a low level of glycosylation, especially when comparing to gastric mucous cells. Most of the labeling obtained with the lectins was observed in the apical membrane of the chief cells and could be unequivocally attributed to O-glycans. This labeling can be explained by the presence of MUC1 and possibly other mucins expressed by chief cells. Labeling of Man-binding lectins at the cytoplasmic region where rough endoplasmic reticulum is located would reflect the N-glycosylation of some secretory proteins. It could be very interesting to compare these lectin binding patterns of the chief cells in the healthy gastric mucosa with those of precancerous and cancerous lesions, to test this methodology as a possible tool for gastric cancer prognosis.

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### **Author contributions**

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Curation, Writing - Review & Editing, Funding acquisition. Edurne Alonso: Formal
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José Sáez: Conceptualization, Methodology, Writing - Original Draft, Visualization,
Supervision, Project administration, Funding acquisition.

#### **Conflict of interest**

The authors declare that they have no conflict of interest.

The funding sources had no role in the study design; in the collection, analysis and interpretation of data; in the writing of the report; or in the decision to submit the article for publication.

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## Figure legends

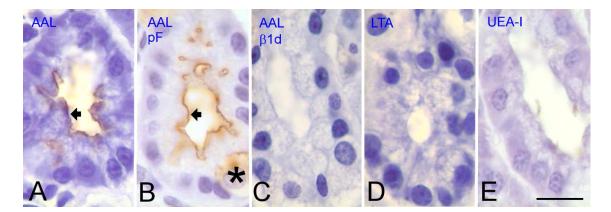


Figure 1. Histochemical staining of the rat fundic glands for Fuc-labeling lectins. AAL labeled the apical surface of the chief cells (a, arrow); the staining remained with the pretreatment with PNGAse F (b, arrow) and was absent after β-elimination for 1 day pretreatment (c). The other two Fuc- binding lectins, LTA and UEA-I were negative. Abbreviations: AAL, *Aleuria aurantia* lectin; LTA, *Asparagus pea* agglutinin; UEA-I, *Ulex europaeus* agglutinin-I; pF, pre-incubation with PNGase F; β1d, β-elimination for 1 day. Asterisk: parietal cell. Scale bar: 20 μm.

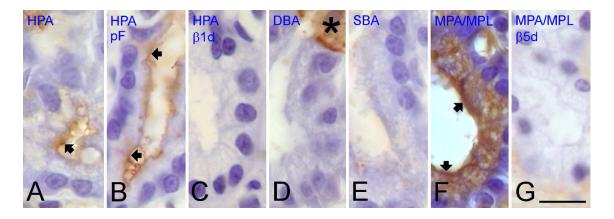


Figure 2. GalNAc-binding lectins. HPA stained the apical surface of the chief cells when it was used alone (a, arrow) or in combination with PNGase F pretreatment (b, arrows); however this staining was not visible when β-elimination for 1 day was used (c). DBA and SBA were negative in all cases (d, e), although DBA stained the parietal cells (d, asterisk). MPA/MPL (f) was positive in the cytoplasm and also at the apical

surface (arrows) and negative after β-elimination for 5 days (g). Abbreviations: HPA, *Helix pomatia* agglutinin; DBA, *Dolichos biflorus* agglutinin; SBA, soybean agglutinin; MPA/MPL, *Maclura pomifera* lectin; pF, incubation with PNGase F; β1d, β-elimination for 1 day; β5d, β-elimination for 5 day. Scale bar: 20 μm.

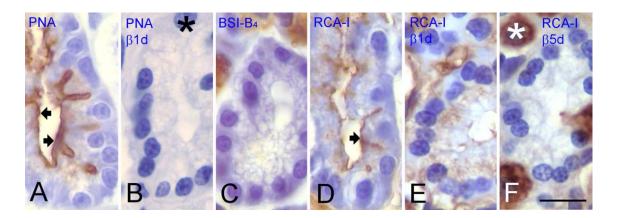


Figure 3. Gal-binding lectins. PNA showed a very weak labeling in the cytoplasm (a) and a stronger staining at the apical surface (a, arrows), neither labeling was observed when the sections were pre-treated with β-elimination procedure for 1 day (b). BSI-B4 was negative (c). The apical surface of the chief cells was moderately positive to RCA-I (d, arrow) and this staining was absent after β-elimination (e, f). The cytoplasm turned from negative to RCA-I (d) to slightly positive in a lattice-like manner after β-elimination for 1 day (e) and again negative if the β-elimination pretreatment was extended to 5 days (f). Abbreviations: PNA, peanut agglutinin; BSI-B<sub>4</sub>, *Bandeiraea simplicifolia*-I-B<sub>4</sub> lectin; RCA-I, *Ricinus communis* agglutinin; β1d, β-elimination for 1 day; β5d, β-elimination for 5 day. Asterisks: parietal cells. Scale bar: 20 μm.

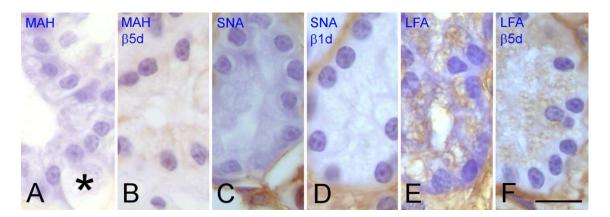


Figure 4. NeuAc-binding lectins. MAH was initially negative (a), but the cytoplasm was very slightly stained after β-elimination for 5 days pre-treatment (b). SNA was negative (c, d), while LFA was positive in the cytoplasm in a lattice-like way (e, f). Abbreviations: MAH, *Maackia amurensis* hemagglutinin; SNA, *Sambucus nigra* agglutinin; RCA-I, *Limax flavus* agglutinin; β1d, β-elimination for 1 day; β5d, β-elimination for 5 day. Asterisk: parietal cell. Scale bar: 20 μm.

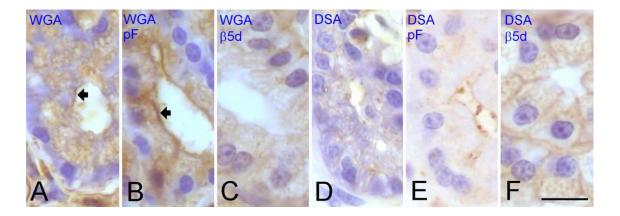


Figure 5. GlcNAc-binding lectins. WGA was positive in the cytoplasm (a) and in the apical surface (a, arrow). After PNGase-F the WGA-staining increased in the apical surface (b, arrow), and the labeling was absent in the apical surface after  $\beta$ -elimination for 5 days (c). DSA showed a lattice-like pattern in the cytoplasm (d) and after PNGase-F (e) and this staining increased after  $\beta$ -elimination for 5 days (f). Abbreviations: WGA, wheat germ agglutinin; DSA, *Datura stramonium* agglutinin; pF, incubation with PNGase F;  $\beta$ 5d,  $\beta$ -elimination for 5 day. Scale bar: 20  $\mu$ m.

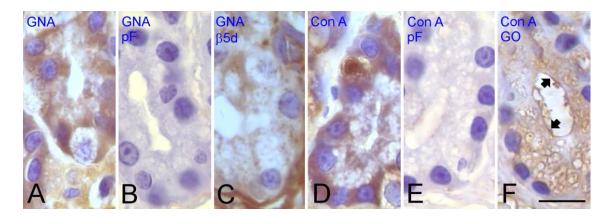


Figure 6. Man-binding lectins. GNA and Con A were positive, mainly in the basal cytoplasm (a, d) and this staining was absent after PNGase-F (b, e); however β-elimination for 5 days did not modify it (c). Pretreatment with Glucose oxidase removed the staining to Con A in the basal cytoplasm (f) while the apical surface was slightly positive (f, arrows). Abbreviations: GNA, *Galanthus nivalis* agglutinin; Con A, Concanavalin A agglutinin; pF, incubation with PNGase F; β5d, β-elimination for 5 day, GO, incubation with Glucose oxidase. Scale bar: 20 μm.

# **Tables**

Table 1. Lectins and their binding specificity.

Lectin	Abreviation	Oligosaccharide specificity		
Orange peel (Aleuria aurantia) lectin	AAL <sup>(1)</sup>	$Fuc\alpha(1,6) > Fuc\alpha(1,2)Gal\beta(1,4)GlcNAc > Fuc\alpha(1,3)$		
Asparagus pea ( <i>Lotus</i> tetragonolobus) agglutinin	LTA	$Fuca(1,3) > Fuca(1,2)Gal\beta(1,3-4)GlcNAc$		
Gorse seed ( <i>Ulex</i> europaeus) agglutinin-I	UEA-I	$Fuca(1,2)Gal\beta(1,4)GlcNAc\beta$		
Snail ( <i>Helix pomatia</i> ) agglutinin	HPA	$GalNAca(1,3)GalNAc > Gal\beta(1,3)GalNAc$		
Horse gram ( <i>Dolichos</i> biflorus) agglutinin	DBA	GalNAca(1,3)GalNAc >> GalNAca(1,3)Gal		
Soybean ( <i>Glycine max</i> ) agglutinin	SBA	$\alpha GalNAc > \alpha Gal > \beta GalNAc$		
Osage orange tree (Maclura pomifera) lectin	MPA/MPL	GalNAcα(1 $\rightarrow$ )Ser/Thr; Galβ(1,3)GalNAcα(1 $\rightarrow$ )Ser/Thr		
Peanut (Arachis hypogaea) agglutinin	PNA	$Gal\beta(1,3)GalNAc > Gal\beta(1,4)GlcNAc > Gal$		
Griffonia (Bandeiraea simplicifolia)I-B4 lectin	BSI-B4	$Gal\alpha(1,3)Gal>Gal\alpha(1,2)Gal;Gal\alpha(1,4)Gal$		
Castor bean ( <i>Ricinus</i> communis) agglutinin I	RCA-I	$Gal\beta(1,4)GlcNAc > Gal\beta(1,3)GlcNAc > Gal\alpha(1,3)Gal$		
Amur maackia (Maackia amurensis) hemagglutinin	MAH <sup>(2)</sup>	NeuAc $\alpha(2,3)$ Gal $\beta(1,3)$ ; SO <sub>4</sub> -3-Gal $\beta(1,3)$		
Elderberry bark (Sambucus nigra) agglutinin	SNA	NeuAcα(2,6)Gal/GalNAc		
Yellow slug ( <i>Limax</i> flavus) agglutinin	LFA	NeuAc		

Lectin	Abreviation	Oligosaccharide specificity		
Wheat germ ( <i>Triticum</i> vulgaris) agglutinin	WGA	(GlcNAc) <sub>2-3</sub> >>> GlcNAc		
		NeuAc		
Thorn apple ( <i>Datura</i> stramonium) agglutinin	DSA	$(GlcNAc)_{2-3} > GlcNAc > Gal\beta(1,4)GlcNAc$		
Snowdrop (Galanthus nivalis) agglutinin	GNA	$Man\alpha(1,3)Man > Man$		
Concanavalin A or Jack bean ( <i>Canavalia</i> ensiformis) agglutinin	Con A	oligosaccharide( $\rightarrow$ 2) Man $\alpha$ (1,6) Man $\beta$ (1,4)GlcNAc $\beta$ (1,4)GlcNAc $\beta$ (1 $\rightarrow$ )Asn oligosaccharide ( $\rightarrow$ 2) Man $\alpha$ (1,3)		

<sup>&</sup>lt;sup>1</sup>AAL is sometimes abbreviated as AAA, but then could be confused with the lectin from fresh water eel (*Anguilla anguilla*), whis is usually abbreviated as AAA.

<sup>&</sup>lt;sup>2</sup>MAH is also abbreviated as MAA-II, MAA-2 and MAL-II. MAH should not be confused with MAL (*Maackia amurensis* leukoagglutinin, also abbreviated as MAM, MAL-I and MAA-I) or MAA (an undefined mixture of MAL and MAH).

Table 2. Semiquantitative evaluation of lectin-binding labeling of the chief cells in the rat fundic gastric glands. The table shows the staining intensity of the cytoplasm of the cells by a numerical scale from 0 to 5. When there was some variability in staining the range is indicated with the minimum and maximum values separated by a dash. When there is no value, it means that the procedure was not applicable. In addition, it is noted when there was a lattice-like cytoplasmic labeling (\*) and also when the apical surface of the cell was stained (#) whatever the intensity of this staining.

Lectin	Alone	PNGase F	β-elimination	β-elimination	Acid	Glucose
			(1d)	( <b>5d</b> )	hydrolysis <sup>(a)</sup>	oxidase <sup>(b)</sup>
Fuc-labeling le	ectins					
AAL	0#	0#	0	0		
LTA	0	0	0	0		
UEA-I	0	0	0	0		
GalNAc-labeli	ng lectins	;				
HPA	0#	O#	0	0		
DBA	0	0	0	0		
SBA	0	0	0	0		
MPA/MPL	4-5*#	4-5*#	4*#	0		
Gal-labeling le	ectins					
PNA	0-1#	0-1#	0	0		
BSI-B <sub>4</sub>	0	0	0	0		
RCA-I	0#	0#	1*	0		
NeuAc-labeling lectins						
MAH	0	0	0	0-1*		

Lectin	Alone	PNGase F	β-elimination	β-elimination	Acid	Glucose
			(1d)	(5d)	hydrolysis <sup>(a)</sup>	oxidase <sup>(b)</sup>
SNA	0	0	0	0		
LFA	2-3*	2-3*	2-3*	2-3*		
GlcNAc-labe	ling lectins	,				
WGA	2-3#	2-3#	2-3#	0-1	3#	
DSA	0-1*	0-1*#	2*	2*		
Man-labeling	g lectins					
GNA <sup>(c)</sup>	3	0	2	2		
Con A	3	0	3	2-3		2-3

<sup>\*</sup>The labeling of the cytoplasm was lattice-like.

<sup>#</sup>The apical surface of the cells was labeled.

<sup>&</sup>lt;sup>(a)</sup>Acid hydrolysis was only performed with WGA to elucidate if the staining was due to GlcNAc or to NeuAc.

<sup>(</sup>b) Incubation with Glucose oxidase was only carried out with Con A to elucidate if the staining was due to Glc or Man.

<sup>&</sup>lt;sup>(c)</sup>Some data of GNA histochemistry have been previously published (Gómez-Santos et al., 2017).