

Enzymatically-derived oligo-carrageenans interact with α -Gal antibodies and Galectin-3

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Abstract

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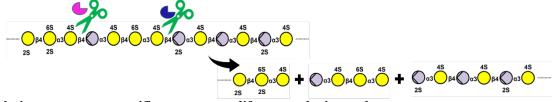
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Carrageenans are linear sulfated galactans synthesized in the Gigartinales, Rhodophyceae species with a varied range of biological properties that are of value to the pharmaceutical and cosmetic sectors. It is unknown how the fine structure of carrageenans dictates their capacity to affect molecular and cellular responses important to wound healing, or the ability to mitigate oxidative, hemostatic and inflammatory processes. Here we use specific endo-carrageenases from the marine bacterium Zobellia galactanivorans, to produce enzymatically defined neoseries oligosaccharides from carrageenans with 3,6-anhydro-D-galactose on the non-reducing end. Further enzymatic modification of the oligosaccharides was done by treating with the 3,6anhydro-D-galactosidases from the same bacterium which hydrolyze non-reducing end 3,6anhydro-D-galactose moieties from neo-carrageenan oligosaccharides. Using the enzymatically produced oligosaccharides, we demonstrate binding to natural human serum antibodies and a monoclonal anti-αGal Ab (m86). The significant interactions with the Galα(1,3)Gal reactive antibodies produced by humans makes them potential potent inducers of complementdependent reactions and attractive for therapeutic applications. We also demonstrate modulation of the galectin selectivity for the Gal-3 Carbohydrate Recognition Domain (CRD) relative to Gal-1 which has implications to targeting specific biological pathways regulated by the galectins.

1. Endolytic cleavage of carrageenans by specific enzymes

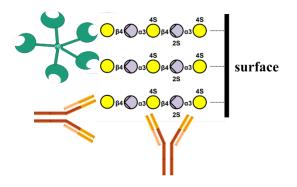


2. Exolytic carrageenan specific enzymes modify non-reducing ends



3. Galectin-3 binds non-reducing ends of carrageenan oligosaccharides

Galactose specific antibodies bind along the length and non-reducing ends of oligo-carrageenans



Key words. carrageenans, carrageenases, 3,6-anhydro-D-galactosidase, glycan-binding antibody, alpha-Gal antigen, galectin, thrombin, anti-tumor

Introduction

The discovery that cellular communication occurs via the 'sugar code' which, unlike nucleic acid and protein code, largely takes place on the cell surface, has led to significant interest in these conjugate molecules (Gabius, 2018). The major players in sugar communication are surface bound glycans responsible for the generation of biochemical messages and carbohydrate-binding proteins transferring those signals into bioactivity. This communication is informed by protein recognition of sugar epitope structures and topology (Romero & Gabius, 2019). Among glycan ligands, galactans stand out for their medical potential, mostly because humans produce staggering amounts of galactosyl-binding (glyco)proteins as anti-Gal antibodies and galectins (Galili, 2013; Laaf, Bojarova, Elling, & Kren, 2019).

A common galactose-based sequence found in the majority of mammals is the α Gal epitope, α -Galp-(1 \rightarrow 3)- β -Galp-(1 \rightarrow 4)-GlcpNAc-R. It is an antigen that has been eliminated from human cells during primate evolution. Synchronically with the elimination, the evolution of molecules capable of reacting with the excluded glycan occurred, *i.e.* natural anti- α Gal antibodies (Macher & Galili, 2008). Maintenance of high levels of natural anti- α Gal antibodies is believed to be due to continual environmental exposure, for instance, to members of the adult human gut microbiome producing an α Gal epitope (e.g. *Escherichia coli, Haemophilus influenza* and *Lactobacillus* species) (Montassier et al., 2019). It is estimated that the amount of anti- α Gal antibodies reaches up to 1-3% of all circulating Ig produced by about 1% of all B cells, ensuring a strong innate immunity reaction. Upon precipitation of surfaces marked with the epitope, the constant Fc parts of Ig triggers the natural immunity into action to battle virus and tumors or recruit and activate tissue macrophages for skin regeneration (Galili, 2013), thus making the antibodies interesting targets for improved wound healing.

Galectins are an ancient family of endogenous soluble β -galactoside-binding proteins abundantly produced in humans (13 out of the 19 known are human) (Laaf et al., 2019). Once outside the cells, galectins bind cell surface glycoproteins through their carbohydrate recognition domains (CRDs). CRDs bind β -galactoside containing epitopes of di- to tetra-saccharides abundant in N- and O-glycans found on glycoproteins and glycolipids of vertebrate tissues. CRDs can operate in a traditional or nontraditional mode. The traditional mode is mediated through direct connection with a receptor glycan to restrict mobility of transmembrane

receptors such as nutrient transporters, cytokines and adhesion receptors. The nontraditional mode assembles multiple glycosylated partners and forms a dynamic and gel-like layer of membrane organization — often termed a lattice — that regulates diffusion, complex formation and domain interactions in the plasma membrane (Nabi, Shankar, & Dennis, 2015). Both modes translate glycan-encoded information into biological cues with modulating effects on intracellular glycoprotein trafficking, cell-matrix adhesion, cell signaling, cancer growth, angiogenesis and effector functions of immune cells (Rabinovich & Toscano, 2009). The galectins have complex biological functions and sometimes opposing effects. Tuning their impacts is tissue, cell type and context-dependent, but yet also attractive as pharmacological targets. Inhibition of extracellular Gal-1 by glycomimetic substances holds potential for pharmacotherapy of several disorders such as skin cancer and hypertrophic scar formation (Astorgues-Xerri et al., 2014; Blanchard, Bum-Erdene, Bohari, & Yu, 2016; Kirkpatrick et al., 2021). Moreover, Gal-3 is considered a preserver of the tumor microenvironment, thus playing an important role in cancer growth (Filipova et al., 2020).

Epitopes for galectins and anti-αGal antibodies contain β-galactose in their motifs and the αGal antigen has proven to be a ligand for galectins as well (Gal-1 and Gal-3) (Hirabayashi et al., 2002). Seaweed matrix polysaccharides are potential epitope mimics as they have regiospecific sulfations (Ficko-Blean, Hervé, & Michel, 2015) but differ structurally from animal sulfated mucosal glycans and glycosaminoglycans. Biologically active substances derived from seaweeds are often sulfated carbohydrates covering multiple biological properties like anticoagulant, antitumor, immunomodulatory and many others (Deniaud-Bouet, Hardouin, Potin, Kloareg, & Herve, 2017; Huang, Tan, & Nie, 2022). Among these algal sulfated polysaccharides, carrageenans are a natural source of galactose-containing glycans with an appealing array of biological properties (Bhattacharyya et al., 2010; Carlucci et al., 1997; Cheong, Qiu, Du, Liu, & Khan, 2018; Cosenza, Navarro, Pujol, Damonte, & Stortz, 2015). Notably, they exhibit strong antiviral properties when they are administered topically to the mucosa of the nasal cavity and the upper respiratory tract (Graf et al., 2018; Pereira & Critchley, 2020). These linear polymers are composed of alternating 3-linked β-D-galactopyranosyl units (G-unit) and 4-linked α-D-galactopyranosyl units (D-unit). In gelling carrageenans the latter is mainly 4-linked 3,6-anhydro-α-D-galactopyranosyl (DA-unit) such as found in β- (G4-DA), κ-(G4S-DA) or 1- (G4S-DA2S) carrageenan motifs. The structures of carrageenans are well known and have been reviewed (Ciancia, Matulewicz, & Tuvikene, 2020; Ficko-Blean et al., 2015). The designation of G-, D-, DA- is made in accordance to the nomenclature proposed by Knutsen and co-authors (Knutsen, Myslabodski, Larsen, & Usov, 1994).

The repeating disaccharide units of G, D and/or DA interchanging along the carrageenan polymer resemble the sequence of the α Gal disaccharide [α -Galp-(1 \rightarrow 3)- β -Galp-(1 \rightarrow 4)]. A recent investigation determined that carrageenan polymers bind to total human serum antibodies with a preference for IgM (Sokolova et al., 2021). This finding led to the conclusion that carrageenans present a highly multifaceted ligand platform for human serum antibody binding; however, the molecular basis of the interaction of carrageenans with anti- α Gal antibodies has never been investigated.

Understanding the fine structural characteristics of carrageenans influencing in vitro biological properties could allow the controlled and reproducible creation of carrageenan-based structures with pharmaceutical potential. This can be approached by using carrageenan-specific enzymes with their regio-, stereo-, and enantio-selectivity, mild reaction conditions, and sequential or one-pot reactions. The model marine bacterium Zobellia galactanivorans (Barbeyron et al., 2016) is capable of catabolizing carrageenans by means of a carrageenaninduced regulon containing a non-canonical polysaccharide utilization locus (PUL) (Ficko-Blean et al., 2017). Z. galactanivorans produces endolytic enzymes capable of releasing oligocarrageenans. The family GH16 κ-carrageenases (Matard-Mann et al., 2017; Michel, Chantalat, Duee, et al., 2001) and family GH82 1-carrageenases (Barbeyron, Michel, Potin, Henrissat, & Kloareg, 2000; Michel, Chantalat, Fanchon, et al., 2001) are glycoside hydrolases that specifically cleave β-1,4 linkages within carrageenan chains, producing varying lengths of oligosaccharides with a DA moiety at the non-reducing end and a G moiety at the reducing end. The sulfation motifs on the released oligosaccharides will vary according to the carrageenases used. To this end, both the carrageenan oligosaccharides and the αGal epitope, have nonreducing end galactose-based structures, though the non-reducing end terminal 3,6-anhydro-α-D-galactose structure has undergone a ring flip due to the steric restraints of the 3,6-anhydrobridge.

The exolytic 3,6-anhydro-D-galactosidase enzymes (ZgGH127-1, ZgGH127-2, ZgGH127-3 and ZgGH129), were recently discovered in the first described carrageenan PUL (Ficko-Blean et al., 2017). They are specific for the removal of the α -1,3-DA monosaccharide residue from the non-reducing end of neo-carrageenan oligosaccharides, this results in a non-reducing end β -1,4-G residue after exohydrolase treatment. One 3,6-anhydro-D-galactosidase, ZgGH129, has undergone a thorough kinetic characterization (Wallace, Guee, et al., 2020) and

its fine specificity has also been examined, revealing a preference for unsulfated neo-β-carrabiose motifs on the non-reducing end (Wallace, Ficko-Blean, & Stubbs, 2020; Wallace, Guee, et al., 2020). While the catalytic activity of the other 3,6-anhydro-D-galactosidases from the GH127 family has been confirmed, qualitative analyses indicate they may not all share the same fine specificities; what is known is that they are able to remove the 3,6-anhydro-D-galactose residue from natural neo-β-κ-κ DP6 oligosaccharides obtained from *Furcellaria lumbricalis* (Ficko-Blean et al., 2017).

Since carrageenans have modified $Gal\alpha(1,3)Gal$ motifs and internal β -galactose in their motifs, they are probably able to intervene in processes involving sugar codes. Thus, the aim of the study is to explore the possibilities of carrageenan-based galactans to interact with the human galactoside binding proteins Galectins 1 and 3 (Gal-1 and Gal-3) and human serum glycan binding antibodies. Using carrageenan-specific marine bacterial enzymatic treatments to change the carrageenan structures should enable us to postulate which galactan components, either terminal or along the glycan chain itself, are responsible for binding to the galactoside-specific proteins.

Materials and methods

κ-Carrageenan was a kind gift from Sanofi Bio-Industries.

The reagents purchased from Sigma, St. Louis, USA. 1-Carrageenan from Euchema denticulatum (cat no. C1138, lot no. SLCC4631), Ampicillin sodium salt (cat no. A9518), R-2A Agar (cat no. 17209, lot no. BCCB8737), sodium deoxycholate (cat no. D6750, lot no. MKBS0373V), sucrose (cat no. S0389, lot no. SLCF6079), lysozyme (cat no. L6876), Ammonium persulfate, for electrophoresis (cat no. A3678, lot no. 113K0656), silver nitrate (cat no. 31630, lot no. STBJ5526), Human Serum (cat no. P2918, lot no. SLCC8367), Anti-Human IgM (μ-chain specific)-Peroxidase antibody produced in goat (cat no. A6907, lot no. SLCH6187), 2,2'-Azino-bis(3-ethylbenzothiazoline-6-sulfonic acid) (cat no. A 3219, lot no. MKCM9914), Streptavidin-HRP Solution (ORO3L, lot no. 3792178), Thiodigalactoside (cat. no. SML2310, lot no. 0000093539), Thrombin Inhibitor Screening Kit (Fluorometric) (cat. no. MAK243), Factor Xa Inhibitor Screening Kit (Fluorometric) (cat. no. MAK239)

Reagents from other companies. Celite 545 (Diatomeceous silica for filtration) was from PROLABO (cat no. 22552.290, lot no. 93103, Paris, France), tryptone (cat no. BP9727, lot no. 137-676, Fisher BioReagents, USA), yeast extract (cat no. 1349-9539, lot no. A1985, Fisher

Scientific, USA), DNase I (cat no. 18405J, Interchim, France), imidazole (cat no. G3H4N2, lot no. 211276, ThermoScientific, Germany), N,N,N',N'-tetramethylethylenediamine (TEMED) (cat no. 87689, Fluka, USA), acrylamide/bis-acrylamide (37.5:1) solution 30% (cat no. 1610158, Bio-Rad Inc., France), Alcian Blue (cat no. J60122, lot no. U19F032, AlfaAesar, Germany), Carbo-bind plate (Costar, Corning Inc., NY, USA), 96-well tissue culture treated sterilized polystyrene Imagelock microplate (cat no. 4379, Sartorius, Essen BioScience, Ann Arbor, MI, USA). Human VEGF DuoSet ELISA (cat no. DY293B, R&D Systems, USA). Galα1-3Galβ1-4Glc carbohydrate sequence (cat no. OG44766, Biosynth Carbosynth)

Reagents from Abcam. Biotinylation Kit / Biotin Conjugation Kit (Fast, Type B) -Lightning-Link® (ab201796) composed of Quencher reagent, ab273995, lot no. GR3401609-1, Modifier reagent, ab273994, lot no. GR3404190-4; Biotin (Typ B) Conjugation Mix, ab274078, lot no. GR3395269-1. Human Galectin 1 Matched Antibody Pair Kit (ab218170) or Human Galectin 1 Detector antibody (Abcam, SNT-12499, lot no. 2101008040). Human Galectin 3 Detector antibody (Abcam, cat no. ab 270341, lot no. GR3405992-1). Human Galectin 3 Capture antibody (ab270340, lot no. GR3404762-1). human monoclonal antibodies to Alpha Gal [m86] (ab281899, lot no. GR3422108-1). TMB ELISA Substrate (High Sensitivity) (cat. no. ab171523, lot no. 2101010452). Recombinant human Galectin 1 protein (ab50237, lot no. GR322443-22 and GR322443-23). Recombinant human Galectin 3 protein

ÄKTA Avant Chromatographic System equipped with Unicorn™ software, SP Histrap™ FF column (5ml), Sephadex 200 gel packing were purchased from GE Healthcare Co., Ltd. (Chicago, IL, USA).

Polysaccharide extraction

(active) (ab89487, lot no. GR339273-2).

C. crispus gametophytes and *F. lumbricalis* were harvested during low tide along the shore of Roscoff (Brittany, France) on December 15^{th} , 2020 and September 3^{rd} , 2015, respectively. After the harvest the algae were washed with water to remove impurities, dried in an oven at 40° C for 24 h, ground in powder by Waring blender and weighed. The polysaccharides from both algae were isolated by aqueous extraction in MilliQ water at 80° C for 4 h (w:v = 30 g: 2 L). To remove the algal biomass residue, Celite 545 (20 g) was added to the resultant solution followed by filtration by a paper filter or nylon membrane. The carrageenan polymer was recovered by precipitation with 4 volumes of ethanol, washed with ethanol and then dried overnight in a $45 ^{\circ}$ C oven. In order to remove excess salt and non-polymeric materials, the carrageenans were dialyzed (12-14 kDa MWCO Spectra/Por ®)

against distilled water for 3 days at room temperature followed by precipitation with ethanol (v = v = 1 : 4) and drying at 45°C overnight. The carrageenan yield was calculated from the percentage of dried algae.

Nuclear magnetic resonance

One dimensional 1H NMR spectra were recorded at 353K on a BRUKER AVANCE III HD 500 spectrometer equipped with an indirect 5 mm probehead BBFO 1H/ [BB]. NMR analyses were performed on samples dissolved in D2O (10 mg/mL). The spectra were collected according BRUKER's pulse programs with standard pulse sequences of 2 s of delay, a 30° pulse and 64 scans. Chemical shifts were expressed in ppm relative to TMS as external reference.

Media

LB-media: Luria-Bertani medium (tryptone 10 g, yeast extract 5 g, NaCl 5 g, 1 L H₂O, autoclaved for 20 min at 115°C), LB/Amp-media: LB-media supplemented with ampicillin sodium salt (0.01%, w : v = 0.1 g : 1 L), LB/ Amp/agar-media: LB-media supplemented with agar (1.5 %) autoclaved for 15 min at 121°C after cooling down to 45-50°C ampicillin sodium salt was added (0.01 % w/v). ZYP 5052 media: After autoclavation of ZY solution (10 g tryptone, 5 g yeast extract, 925 mL water) the media was supplemented with the following solutions 1 mL 1M MgSO₄, 1 mL 1000×trace metal mixture (1×conc = 50 μ M Fe, 20 μ M Ca, 10 μ M Mn, 10 μ M Zn, 2 μ M Co, 2 μ M Cu, 2 μ M Ni, 2 μ M Mo, 2 μ M Se, 2 μ M H₃BO₃), 20 mL 50×5052 (1×5052 = 0.5 % glycerol, 0.05% glucose, 0.2% α -lactose), 50 mL 20×NPS (1×NPS = 100 mM PO₄, 25 mM SO₄, 50 mM NH₄, 100 mM Na, 50 mM K) + 1 mL ampicillin stock (100 mg/mL).

Cloning, enzyme production and purification

All genes coding for endolytic and exolytic carrageenan-related enzymes were cloned from genomic DNA extracted from the marine bacterium *Zobellia galactanivorans* (Barbeyron et al., 2016) into the ampicillin resistant plasmid pFO4 as previously described (Groisillier et al., 2010). The pFO4 plasmid incorporates a 6His tag on the N-terminus of the proteins for immobilized metal affinity chromatography (IMAC). The plasmids were transformed into the *E. coli* BL21 (DE3) strain for protein expression. The list of recombinant enzymes is the following: κ-Carrageenase (Label: ZgGH16, locus ID: Zgal_236, MW: 33.5 kDa) (Matard-Mann et al., 2017), ι-carrageenase (ZgGH82, Zgal_4265, MW: 51.0 kDa (Barbeyron et al., 2000; Groisillier et al., 2010) and four 3,6-anhydro-D-galactosidases (3 GH127 enzymes: ZGAL 3147, MW: 77.0 kDa; ZGAL 3148, MW: 75.0 kDa; ZGAL 3150, MW: 78.8 kDa; and

one GH129 enzyme: ZGAL_3152, MW: 77.8 kDa) (Ficko-Blean et al., 2017; Wallace, Ficko-Blean, et al., 2020; Wallace, Guee, et al., 2020).

Prior to experimental use, the *E. coli* BL21(DE3) strains containing the target plasmids were maintained as frozen glycerol stocks at -80°C. The culture cells were streaked out on LB/Ampicillin agar plates and incubated overnight at 37°C. The next day single cell colonies from the plates were used to initiate the starter cultures in 5 mL LB/Ampicillin media, incubated overnight at 37°C and shaking at 175 rpm. The starter culture (1mL) was used to inoculate 1 L of ZYP 5052 media (Studier, 2005) and which was cultivated 180 rpm and 20°C. After 3 days, the cells were harvested by centrifugation (3000 g, 30 min, 4°C).

A chemical lysis procedure was used to lyse the cells. First, the harvested cells were suspended in 10 mL of resuspension buffer (25 % sucrose, 50mM Tris pH 8.0). After 10 min on ice with lysozyme (150 μ L, 100 mg/mL), 20 mL of lysis buffer containing 100 mM NaCl, 1% Na deoxycholate, 1% Triton X-100, 20 mM Tris pH 8.0 was added and incubated 5 min on ice. The final step of the chemical lysis consisted in the addition of MgCl₂ (to 5mM) and DNase I (150 μ L, 500 000 Units/mL in CaCl₂ 10 mM, MgCl₂ 10 mM, Tris 10 mM pH8) at room temperature for 5 minutes. The lysed cells were centrifuged at 20 000 g for 45 min at 4°C and the supernatants containing the soluble protein were passed through 0.2 μ m filter.

Enzyme purification was performed by means of an ÄKTA Avant system with a 5 mL HisTrap FF column immobilized with Ni²⁺. After loading the column with the filtered supernatant, the column was washed with 20 mM Tris pH 8.0 and 100 mM NaCl until there was no longer a change in the UV. The proteins were eluted in a linear imidazole gradient up to 0.5M imidazole, 25 mM Tris pH 8, 50mM NaCl. A second purification step was performed on a Sephadex-200 gel filtration column by isocratic elution with 50 mM HEPES pH 7.5 and 100 mM NaCl. Purity of the protein samples was analyzed by SDS-PAGE.

Digestion of polymers by endolytic enzymes

Endolytic carrageenases (κ- and ι-carrageenases) were incubated with the carrageenan polymer samples to release oligosaccharides. The carrageenans were treated three different ways: 1) digested with ι-carrageenase, 2) digested with κ-carrageenase or 3) digested with κ-and ι-carrageenase enzymes sequentially. All samples were kept in identical conditions (100 rpm, 30°C, 0.5% carrageenan solutions (3 g/600 mL) in 50 mM NH₄HCO₃ at pH 7). The sequential reactions were started by treatment with ι-carrageenase (1.2 μg/mL) over 5 days, heat denaturing for 25 mins at 70°C, followed by treatment with κ-carrageenase (1.2 μg/mL)

over 4 days. The single enzymatic reactions were done by adding either 1-carrageenase (1.2 μ g/mL) or κ -carrageenase (1.2 μ g/mL). All the reactions were stopped by heating for 25 min at 70°C in a water bath.

Carbohydrate polyacrylamide gel electrophoresis

Carbohydrate polyacrylamide gel electrophoresis (C-PAGE) (Zablackis & Perez, 1990) was used to qualitatively visualize the polydispersity of charged oligo-carrageenans after digestion with endolytic enzymes (ι- and/or κ-carrageenases). The concentration of acrylamide in the stacking and running gels was 6% and 27%, respectively. The upper and lower buffer chambers contained 50 mM Tris, pH 8.7 with 1 mM EDTA. The samples (20 μL, 0.5 %) were mixed with 60 μL of sucrose-phenol red solution (10 % sucrose, 0.1 % phenol red), after that 2-3 μL of the mixture were loaded, and the samples were electrophoresed at ambient temperature and 200 V until 75% of the migration front of the gel length (visualized with the phenol red). The gels were placed into a big petri dish. The carbohydrates were fixed and stained in the gel with 0.5% alcian blue in MilliQ H₂O (10 minutes, with agitation) and destained in MilliQ H₂O with agitation until the gel showed clear again. The silver staining of the glycans was performed by silver nitrate (0.4%, 10 min, without agitation). After washing 3 times with H₂O under agitation, a solution of sodium carbonate/formaldehyde (7 g sodium carbonate and 80 μL formaldehyde in 100 mL MilliQ H₂O) was added, manually agitated and the reaction was stopped with 1 mL acetic acid added directly to the petri dish.

Reducing sugar assay

A reducing sugar assay was performed to measure the level of hydrolysis of the samples by the enzymes following the method developed by Lever (Lever, 1972). It is based on the reaction of reducing ends of sugars with p-hydroxybenzoic acid hydrazide (PAHBAH) which is stored at 5% in 0.5M HCl and prior to the assay diluted 10-fold in 0.5 M NaOH. The test was done in triplicate by heating the mixture of the sample (20 μ L) and the PAHBAH reagent (180 μ L) at 100°C for 5 min. The absorbance of the reaction mixtures was detected at 410 nm by a Spark 10 M microplate reader (Tecan, France).

Modification of carrageenans by 3,6-anhydro-D-galactosidases

The oligosaccharides from C. crispus and F. lumbricalis (0.7 % solutions in H_2O) produced after endolytic digestion by κ - and ι -carrageenases were treated with the 3,6-anhydro-D-galactosidases or with the enzyme buffer as a control, otherwise all samples were treated the same. The enzymes or enzyme buffer were added to their respective solutions two times at 24

h intervals and incubated a total of 72 h at 25 °C. After the first enzyme addition the concentration of GH127-2, GH127-3, GH129 was 0.05 mg/mL and of GH127-1 was 0.005 mg/mL. After the second enzyme addition the final enzyme concentration was 0.1 mg/mL for GH127-2, GH127-3, GH129 and 0.01 mg/mL for GH127-1. Carrageenan polymers were also treated with a mixture of the 3,6-anhydro-D-galactosidases with final concentrations of 0.025 mg/mL for GH127-2, GH127-3, GH129 and 0.0025 mg/mL for GH127-1. Controls contained only enzyme buffer solution in a volume equal to the volume of enzymes added.

Exolytic 3,6-anhydro-D-galactosidase activity on artificial substrate

The activity of the 3,6-anhydro-D-galactosidases was tested on an artificial substrate 4-nitrophenyl 3,6-anhydro-α-D-galactopyranoside (3,6-ADG-pNP) (Wallace, Guee, et al., 2020). The enzymes (ZgGH127-1, ZgGH127-2, ZgGH127-3 and ZgGH129, 5 μg/mL, final) were incubated with 2 mM 3,6-ADG-pNP in water in an endpoint digestion over 2 days at ambient temperature, the development of the color reaction with the substrate was registered at 405 nm. The blank was with water instead of the 3,6-anhydro-D-galactosidases and control was with boiled mixture of enzymes.

Purification of the carrageenan oligosaccharides

Carrageenan oligosaccharides, treated with the 3,6-anhydro-D-galactosidases and the buffer controls, were ultrafiltrated via a 10 kDa CutOff membrane on Amicon® Ultra-15 10K centrifugal filter devices for volumes up to 15 mL (cat no. UFC901008D, UFC901024D, UFC901096D, Millipore, Sigma, USA) at 2500 g. Monosaccharides under 100 – 500 Da were removed by dialysis with Spectra/Por dialysis tubes (MWCO 100 - 500 Da) against ultrapure water.

High performance size exclusion chromatography (HPSEC)

Filtered samples (200 μ L, 0.22 μ m) were injected on analytical Superdex 200 and Superdex peptide columns (Cytiva) connected in series. The elution was conducted in 0.1 M LiNO₃ using an isocratic Thermo Scientific Ultimate 3000 pump working at a flow rate of 0.5 mL/min. The elution was monitored using a Wyatt Optilab Rex refractive index detector. Chromeleon software (Thermo Scientific) was used for data acquisitions. For comparison, inhouse purified standards oligosaccharides were also injected (Guibet et al., 2008; Jouanneau, Boulenguer, Mazoyer, & Helbert, 2010).

Thrombin and Factor Xa inhibitor screening capacity of carrageenans

The ability of carrageenans to inhibit the coagulation factors Thrombin and Factor Xa was tested by fluorometric assay using kits: Thrombin Inhibitor Screening Kit (Fluorometric, cat. no. MAK243), Factor Xa Inhibitor Screening Kit (Fluorometric, cat. no. MAK239). The protocol follows Thrombin and Factor Xa enzymes which cleave a synthetic 7-Amino-4-methylcoumarin (AMC)-based peptide substrate to release fluorophore AMC. Fluorescence (ex = 350 nm/em = 450 nm) was measured in kinetic mode for 30–60 minutes at 37°C. In the presence of irreversible inhibitors (GGACK Dihydrochloride for Factor Xa, 10 mM, and PPACK Dihydrochloride, 2 mM, for Thrombin) the reaction was abolished. The concentration for carrageenans for anticoagulant activity testing was 1 mg/ml in water. Both Thrombin and Factors Xa screening assays were performed and the results calculated according to the manufacturer's instructions.

Binding human serum antibodies and monoclonal human antibodies to aGal [m86]

To determine the binding ability of normal human serum (NHS) antibodies and monoclonal antibodies (mAb) to the alpha-Gal [m86] epitope, an indirect ELISA-based technique using sugars covalently linked to Carbo-bind® plate wells was applied. Briefly, Carbo-bind® plates were coated with 50 µL of carrageenans [0.02, 0.2, and 2 mg/mL in phosphate-buffered saline (PBS), pH 7.5 were initially tested and 2 mg/mL was chosen as the optimal concentration], sealed with plate seal tape and incubated overnight at 37°C. The plates were then heated at 60°C for 25 min, rinsed 2-4 times with washing buffer (1X PBS, 0.05% Tween 20) and blocked with 1 mg/mL of bovine serum albumin (BSA) for 90 min while shaking then washed. Human serum (2000-fold dilution) in 1X PBS containing 0.1 mg/mL BSA was added to the wells and incubated, shaking, for 2 h at room temperature. The mAb to αGal (0.1 µg/mL) in 1X PBS containing 0.1 mg/mL BSA was added to the wells and incubated, shaking, overnight at 4°C with mAb to αGal. After washing, horseradish peroxidase-conjugated goat anti-human IgM (μ-chain specific) was diluted 2000-fold in PBS with 0.1 mg/mL BSA buffer and 50 µL was added per well for 1 h at room temperature, 400 rpm shaking. The bound antibodies were detected by the reaction with the TMB substrate and the absorbance at 450 nm was measured. The mean absorbance of negative control wells (coating with buffer alone) was used to define the endpoint titer. Commercial Galα1-3Galβ1-4Glc (Biosynth Carbosynth) was used as positive control.

Binding human Gal-1 and Gal-3 inhibition assay with thiodigalactoside

The galectin binding assay was done on CARBO-bind® 96-well microwell plates. All reagents were added at $50 \mu L$ /well unless otherwise stated, after each step of the protocol the

wells were rinsed with washing buffer (1X PBS, 0.05% Tween 20), and incubation, with the exception of the coating of the plates, was at ambient temperature. First, plates were coated with carrageenans (varying concentrations in 1X PBS, pH 7.5), covered with 96-well plate seal tape and incubated overnight at 37°C. This was followed by a brief shaking down of the overnight condensate and heating at 60°C for 25 min. Blocking was done with BSA (1% in 1X PBS, pH 7.5), 90 min and shaking (400 rpm). Galectin 1 (12.5 ng/well or 50 μL of 18.66 μM) and Galectin 3 (3.125 ng/well or 50 µL of 19.23 µM) were added in 1X PBS containing 2 mM EDTA, pH 7.5 either with or without the inhibitor thiodigalactoside (TDG; 1.4 mM and incubated shaking for 2 h. BSA served as a negative control. The plate-bound Galectins 1 or 3 were conjugated with biotin conjugated monoclonal Detector Galectin 1 (0.25 µg/mL) or Detector Galectin 3 (1 µg/mL) antibodies in blocking buffer [prior to this stage, unconjugated Detector antibody to Galectin 3 was biotinylated by the Biotinylation Kit / Biotin Conjugation Kit (Fast, Type B) - Lightning-Link® according to the manufacturer's instructions]. A streptavidin-peroxidase conjugate (0.5 µg/mL) was added with a TMB chromogenic substrate as an indicator at 450 nm to detect the plate-bound galectins bound to the Detector Galectins. The results are expressed as percentage mean value relative to the negative control (100 %).

Statistical analysis

All data are expressed as the arithmetic mean \pm standard deviation. Activity data were tested for differences using one-way repeated measures analysis of variance (ANOVA) with Tukey post-hoc test. A probability value (P) less than 0.05 was considered significant.

Results and discussion

With growing understanding of the potential of biological responses originating from specific glycan interactions, tuning of the sugar code is important to achieving therapeutic benefit. Following the examples of polymers like heparin, which provides the basis for therapeutic oligosaccharides and glycomimetics (Courtois, 2009; Varki & Gagneux, 2015), we produced a library of oligo-galactans from the red algal carrageenans with varying degrees of sulfation. During the current investigation the focus was narrowed down to gelling type carrageenans (which contain high amounts of the 3,6-anhydro-D-galactose monosaccharide), to examine the effect of the exolytic 3,6-anhydro-D-galactosidases after the action of the endolytic carrageenases.

Four gelling carrageenan samples have been used in the current study. Commercial κ-and ι-carrageenans were used as controls as well as two natural, hybrid carrageenans. Hybrid carrageenans have different sulfation motifs along the carrageenan chains which can affect their functional and rheological properties (Van de Velde, 2008) and are thus also of interest for bioactivity studies. The native carrageenans were gently extracted from the red algae *C. crispus* gametophyte and *F. lumbricalis* giving a yield after extraction, precipitation and dialysis at 12-14 kDa of 9.52% and 16.44%, respectively. The structures of all carrageenan samples, including commercial, were determined by 1H-NMR (Fig. 1a, Supplementary Table 1) and the MWs were also determined by MALS (Supplementary Table 2). The 1H-NMR data confirmed the attribution of commercial samples to ι- and κ-types (Supplementary Table 1). Not unexpectedly, the native carrageenan structures found in the *C. crispus* gametophyte are mainly κ/β-motifs (Guibet et al., 2008) and the structures of the *F. lumbricalis* are mainly κ/β-motifs with a small portion of ι-carrageenan (Correc et al., 2012; Rodriguez Sanchez, Matulewicz, & Ciancia, 2022).

The oligo-carrageenans produced after hydrolysis of the β -1,4-linkage by κ - and t-carrageenases (Fig. 1b) were characterized by carbohydrate polyacrylamide gel electrophoresis (C-PAGE) and the PAHBAH reducing sugar assay (Fig. 1c-d). The predominant electrophoretic pattern of oligo-carrageenan fragments in C-PAGE reflects separation by charge-to-mass ratio, with the less charged species migrating more slowly and the smaller species migrating more rapidly. Undigested (controls) or digested but still high-molecular-weight and/or less-charged carrageenans exhibit limited penetration into the polyacrylamide gel matrix and mostly remain in the stacking area. Our results demonstrate that after treatment of the commercial (τ and τ) and hybrid *C. cripsus* and *F. lumbricalis* (τ) and τ) carrageenans with either τ - or τ -carrageenase alone, or both of the carrageenases sequentially (τ -carrageenase first, followed by τ -carrageenase) it results in oligosaccharides with different migration patterns by C-PAGE. This is particularly evident with the hybrid carrageenans (Fig. 1c).

In some cases, the separation of the carrageenans is clearly visible in the C-PAGE gel, with the stacking area appearing emptier with the samples digested by κ - and ι -/ κ -carrageenases relative to the polymer control (Fig 1c, samples k3, k4, c3, c4). The κ / β -type oligo-carrageenans from *F. lumbricalis* poorly penetrate the gel matrix - probably due to their limited charge density. Since the C-PAGE technique depends largely on the charge of the fragments to be qualitatively analyzed, we also evaluated the depth of the endolytic degradation by assaying the increase in free reducing ends by the PAHBAH reducing sugar method (Lever, 1972). This

provided a quantitative method to monitor the substantial hydrolysis of low sulfated κ/β -carrageenan from F. lumbricalis. It was also confirmed that degradation of commercial κ - and ι -carrageenans was highly specific to their appropriate enzymes, κ -carrageenase and ι -carrageenase, respectively, though we did see some significant crossover activity of the κ -carrageenase on commercial ι -carrageenan likely due to the presence of some κ -carrabiose motifs in the mainly ι -carrageenan chain (Fig 1. acd), and vice versa for the ι -carrageenase on commercial κ -carrageenan (Fig 1. ad). None of the carrageenans were treated chemically, they were only treated enzymatically to modify their length.

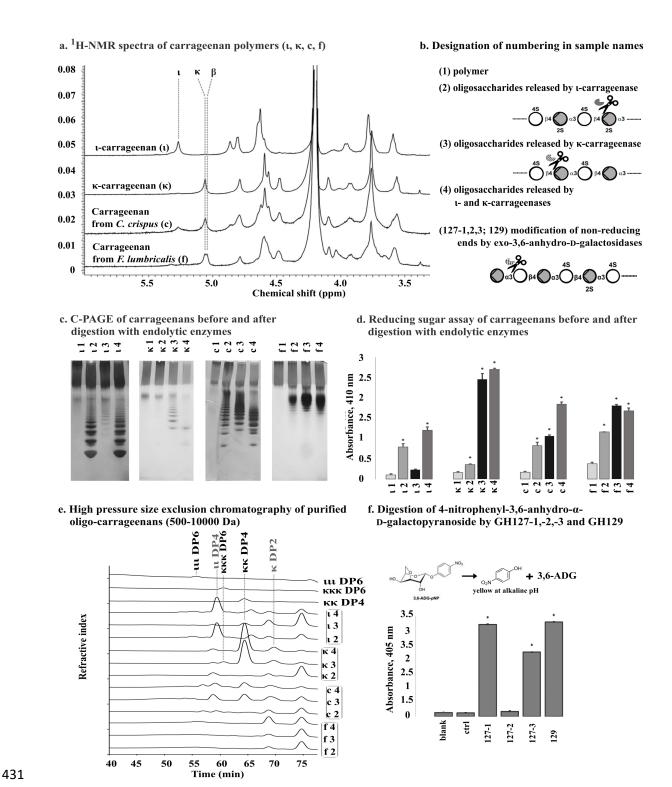


Figure 1. Characterization of carrageenans and enzymatically-produced oligocarrageenans. (a).

¹H-NMR spectra of the carrageenans used in this study. The designation of samples is assembled by numbers (visualized in section b) and letters: "t" stands for commercial t-carrageenan "κ" - κ-carrageenan, "c" - carrageenan from *C. crispus*, and "f" - carrageenan from *F. lumbricalis*. (b). A scheme to visualize the nomenclature used to describe the polymers and oligosaccharides released by endolytic enzymes and modified by exolytic ones: 1 - polymer, 2 - t-carrageenase, 3 - κ-carrageenase, 4 - sequential double digestion with both endolytic enzymes; 127-1, -2, -3 or 129 -modification by 3,6-anhydro-D-galactosidases GH127-1, GH127-2, GH127-3 or GH129. (c-d). C-PAGE (c) and reducing sugar assay (PAHBAH) (d) of carrageenans before and after digestion with endolytic enzymes. (e). High performance size exclusion chromatography (HPSEC) of purified oligo-carrageenans (500 -

10 000 Da). (**f**). Digestion of artificial substrate 4-nitrophenyl-3,6-anhydro-α-D-galactopyranoside (3,6-ADG-pNP) by 3,6-anhydro-D-galactosidases GH127-1, GH127-2, GH127-3 and GH129 where blank was with buffer instead of enzyme and control (ctrl) was with a boiled mixture of the enzymes. Statistical significance difference (P<0.05) of degradation in comparison to each initial polymer is marked with an asterisk (*). Bars represent mean values of 2 independent experiments with each of them done in triplicates ±SD.

HPSEC analysis (Fig. 1e) confirms the production of oligosaccharides mixtures containing at least κ – di, tetra- and hexa-saccharides, and ι – tetra and hexa-saccharides. κ -carrageenan hydrolyzed by κ -carrageenase or by both enzymes contained κ -tetrasaccharide as a major product, while ι -carrageenan hydrolyzed by ι -carrageenase or by both enzymes contained ι -tetrasaccharide as a main product. Hydrolysates of carrageenans from C. C crispus and E lumbricalis gave more complex mixtures, containing hybrid oligosaccharides for which standards were not available.

After treatment with the specific endohydrolases, the oligosaccharides produced are from the neo-series, with 3,6-anhydro-D-galactose (DA) on the non-reducing end, though the sulfation pattern may vary. These oligosaccharides were then further modified by the exolytic 3,6-anhydro-D-galactosidases. We first verified the activity of the 3,6-anhydro-Dgalactosidases (ZgGH127-1, ZgGH127-2, ZgGH127-3, ZgGH129) on the artificial colorimetric substrate 3,6-ADG-pNP by endpoint reaction (Fig. 1f) (Wallace, Guee, et al., 2020). The data show that 3,6-ADG-pNP substrate was hydrolyzed by all the 3,6-anhydro-Dgalactosidases except GH127-2 under assay conditions. In order to select an upper size limit of oligosaccharides, the samples were ultrafiltrated at 10 kDa. To ensure removal of the DA monosaccharide, which could alone have bioactive properties (Lee et al., 2019; Yun et al., 2013), a dialysis into water with a 100-500 Da membrane was also performed and the yields were calculated after lyophilization (Supplementary Table 3). All oligosaccharide samples were analyzed using HPAEC (Supplementary Figure 1) to help inform on exolytic enzymatic activity in the highly complex carrageenan oligosaccharide mixtures. The polymers also underwent a treatment with the mix of 3,6-anhydro-D-galactosidases, followed by dialysis (100-500 MWCO) into water for use as controls in the bioactivity experiments and their yields were also calculated (Supplementary Table 4).

Glycan-protein interactions were assayed in a system where the reducing ends of sugars were covalently attached to the surface via a reaction of the open chain aldehydes with the hydrazide coated surface. The methods used to detect interactions between covalent surface-adhered glycans and proteins were ELISA-based and thus efficiently able to profile a large number of samples (Wolfenden, Cousin, Nangia-Makker, Raz, & Cloninger, 2015). The

orientation in space and elimination of access to reducing ends gave us an opportunity to delve into the biochemistry of binding favored by eukaryotic galactose-binding proteins. The tested proteins were polymorphic antibodies of total human serum, a monoclonal IgM anti- α Gal antibody (m86 mAb) and Galectins 1 and 3 (Gal-1 and Gal-3). The usefulness of using the CARBO-bind® plates is that they provide a multivalent surface for the antibody and lectin binding, which likely contributes to the avid binding of the protein partners.

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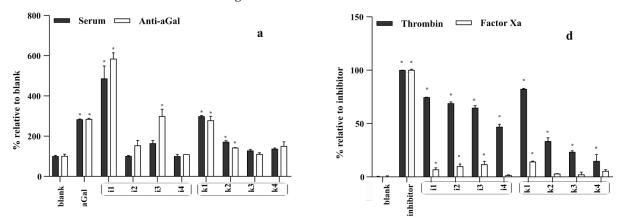
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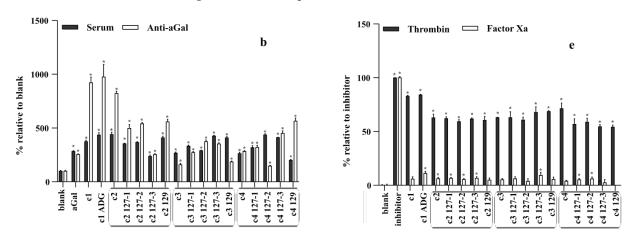
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Total human serum antibodies and monoclonal antibodies against Galα(1,3)Gal, a minimal αGal epitope (Cunningham et al., 2013; Milland et al., 2007), (m86 mAb (Galili, LaTemple, & Radic, 1998)) were analyzed for capacity to bind carrageenans (Fig. 2a-c). As a positive control we used the trisaccharide Gala1-3Gal\beta1-4Glc covalently linked to the plate surface and leaving free the minimal αGal epitope on the non-reducing end. M86 mAb were IgM type and, out of human serum antibodies, only IgM variety was selected for determination because IgM human serum was shown to be far more reactive to the aGal epitope than IgG (Galili, Macher, Buehler, & Shohet, 1985; Sandrin, Vaughan, Dabkowski, & McKenzie, 1993). Our results demonstrated that IgM of both natural human serum and mAb were able to bind carrageenans (Fig. 2a-c). Similar binding intensities to natural antibodies and mAb were detected only for ι- and κ-commercial carrageenans and their derivatives. Less signal was detected on the oligosaccharides, suggesting that the antibodies have enhanced binding to the carrageenan polymer 3D structure rather than the non-reducing ends of the carrageenan oligosaccharides (Fig. 2a). This endo-binding could be intra- or intermolecular. The signal intensity for binding is similar between the trisaccharide of the positive control and kcarrageenan polymer, but the signal with 1-carrageenan polymer is twofold higher, thus sulfation degree may also affect antibody binding (Fig. 2a). The primary theory of binding by antibodies to the aGal epitope is end-on insertion (Milland et al., 2007), but the groove type binding of Galα(1,3)Gal for some lectins has already been reported and the possibility of similar binding type in antibodies is implied (Yuriev et al., 2009). Our results support the possibility of groove-type antibody binding, at least for the carrageenan polymer based substrates.

ι- and κ-carrageenans and their derivatives



и/к-carrageenan from C. crispus and its derivatives



κ/β-carrageenan from F. lumbricalis and its derivatives

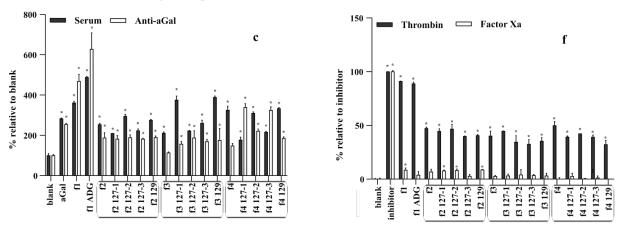


Figure 2. Activity of carrageenans towards human antibodies and anticoagulant factors. (a, b, c). Carrageenans ability to bind antibodies of human serum and m86 (monoclonal anti-αGal Ab) with a trisaccharide α-Gal sequence (Galα1-3Galβ1-4Glc) as positive control. (d, e, f). Inhibition of anticoagulant Factor Xa and Thrombin by carrageenans. Irreversible inhibitors GGACK Dihydrochloride (10 mM) and PPACK Dihydrochloride (2 mM) were used for Factor Xa and Thrombin, respectively. (a) and (d) corresponds to commercial k and i-carrageenans and their enzymatic derivatives; (b) and (e) corresponds to *C. crispus* ι/κ-carrageenans and their enzymatic derivatives. The designation of the samples is assembled by letters and figures: "t" stands for commercial ι-carrageenan, "κ" - commercial κ-carrageenan, "c" - carrageenan from *C. crispus*, "f" - carrageenan

from *F. lumbricalis*. The numbering attributes to the initial polymer and after treatment with endolytic enzymes: **1** - polymer, **2** - ι-Carrageenase, **3** - κ-carrageenase, **4**- digestion with both endolytic enzymes, 127-1, 127-2, 127-3, 129 correspond to the 3,6-anhydro-D-galactosidases ZgGH127-1, ZgGH127-2, ZgGH127-3 and ZgGH129, respectively; ADG – mixture of the four 3,6-anhydro-D-galactosidases. The asterisk (*) indicates probability value *P* <0.05 relative to blank (100%) where no sugars were bound to the plate. The data are representative of triplicates from 2-4 independent experiments.

As with the commercial carrageenan polymers, the anti-αGal mAb interactions showed more signal intensity with the sulfated C. crispus κ/ι -carrageenans relative to the F. lumbricalis κ/β -carrageenans; for the serum antibodies the preference is less clear. The nuances of antibody exo-binding to the hybrid oligo-carrageenans were more profoundly studied by using 3,6anhydro-D-galactosidases to remove non-reducing 3,6-anhydro-D-galactose. C. crispus derivatives released by 1-carrageenase (c2, c2 127-1, c2 127-2, and c2 129) and the double digest (c4 129) showed substantially more signal intensity for the anti-αGal mAb relative to the natural serum antibodies (Fig. 2b); however, the C. crispus oligosaccharide with the strongest signal is the c2 oligosaccharide, only treated with 1-carrageenase, not with the 3,6-anhydro-Dgalactosidases. This leads us to suggest that the increased signal for the c2 oligosaccharide with mAb appears when there is a DAnr, possibly sulfated in 2S since it was prepared with the 1carrageenase. On the other hand, treatment of the double digest c4 129 sample is particularly interesting because there seems to be a much stronger selectivity for the mAb relative to the serum antibodies. Furthermore, the signal intensity of c4 129 is higher relative to the c4 for the mAb interaction. This leads us to suggest that the double digest followed by the ZgGH129 3,6anhydro-D-galactosidase treatment exposes a motif which has higher specificity for the mAb relative to the serum antibodies.

The *F. lumbricalis* carrageenan and its derivatives released by 1-carrageenase (f2, f2 127-1, f2 127-2, f2 127-3 and f2 129) and κ -carrageenase (f3, f3 127-1, f3 127-3, and f3 129) showed more signal intensity with serum antibodies relative to anti- α Gal mAb (Fig. 2c). Interestingly, this trend was not evident for the double digested f4 samples which gave a mixed preference for either serum antibodies (f4, f4 127-2, f4 129) or the anti- α Gal mAb (f4 127-1, f4 127-3) (Fig. 2c). The highest signal for the *F. lumbricalis* serum antibodies with oligosaccharide was seen on f3 127-1 and f3 129 (treated with κ -carrageenase followed by the 3,6-anhydro-D-galactosidases), suggesting a preference for Gnr or G4Snr. This contradictory result between mAb and the serum Abs is possibly related to the different specificities found in human serum antibodies. The monoclonal antibodies are specific to a minimal α Gal epitope and this pivotal specificity is shared by some of the natural antibodies (Milland et al., 2007). However, the fine specificity to the full epitope is conferred to the natural antibodies by sugar moieties

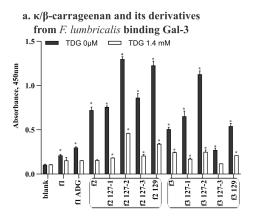
downstream to the minimal recognition unit, such as Glc or GlcNAc residues (Milland et al., 2007). Carrageenans do not contain either Glc or GlcNAc and the galactose units themselves are often modified: firstly, by negatively charged sulfate groups and, secondly, by a hydrophobic 3,6-anhydro-bridge which sterically forces an inversion of the galactopyranose conformation (from 4C_1 to 1C_4) (Ciancia, Noseda, Matulewicz, & Cerezo, 1993; Ficko-Blean et al., 2015). These structural differences in the carrageenan derivatives may contribute to the differences in preference between the monoclonal and natural antibodies.

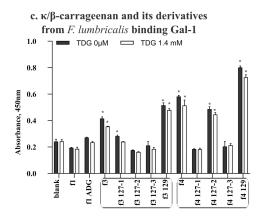
Strikingly, carrageenans can modulate blood protease cascades, with their sulfation patterns helping to determine their pro- or anti-coagulant properties (Pomin, 2012). Regiosulfation on the C2 of the anhydro-α-D-galactose and the C6 of the β-D-galactose has been shown to increase the carrageenan's ability to inhibit thrombin activity (de Araujo et al., 2013), however the importance of the 3,6-anhydro-D-galactoside has never been evaluated for anticoagulant activity. To this end, we evaluated the effect of the carrageenans and their potential for anticoagulation properties on thrombin and Factor Xa through an inhibition assay. For commercial ι - and κ -carrageenan polymer we see a similar inhibition of thrombin and poor Factor Xa inhibition (Fig 2d). Decreasing the MW reduced the thrombin inhibition more significantly for the κ-oligosaccharides than the ι-oligosaccharides (Fig 2d). For C. crispus carrageenans the decrease in MW by the endohydrolases (c2, c3, c4) and modification of the non-reducing ends by the 3,6-anhydro-D-galactosidases (GH127-1, -2, -3, and GH129) did not drastically affect the thrombin inhibitory activity relative to the polymer (c1) (Fig. 2e). For κcommercial and F. lumbricalis carrageenans and their oligosaccharide derivatives, decreasing the MW reduced the thrombin inhibition relative to the polymer (Fig. 2d,f). This may also be related to sulfation degree as both the κ -carrageenan and F. lumbricalis carrageenans (mainly hybrid κ/β-motifs (Correc et al., 2012)) are significantly less sulfated than ι-carrageenan. In all cases, the inhibitory potential of carrageenans towards Factor Xa was not substantial in comparison to the positive irreversible inhibitor control (Fig. 2d-f).

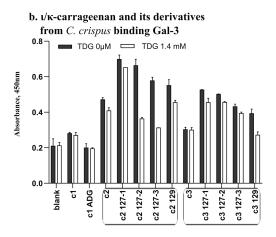
The carrageenans were further tested with regard to their interaction with galectins. In general, galectins are a large family of glycan-binding proteins with preference for β -galactosides. Their subfamilies are classified by fine-tuned specificity for longer glycans of saccharides linked on either side of the β -galactosides (Nielsen et al., 2018). Galectins bind through two major types of interactions: a large hydrogen-bond network and hydrophobic CH- π interactions between the glycan and aromatic amino acid sidechains in the CRD (Modenutti, Capurro, Di Lella, & Marti, 2019). Gal-1 and Gal-3 have been chosen as target galectins

because inhibitors selectively favoring one of them would be promising since both galectins have an assortment of functions (Nabi et al., 2015). Gal-3 has a binding affinity to oligosaccharide structures bearing the lacNAc motif internally while Gal-1 prefers terminal disaccharide lacNAc sequences (Laaf et al., 2019). Carrageenans contain endo- β -D-galactose but no N-acetyl groups or nitrogen atoms. The α Gal epitope has been listed among galectin inhibitors (Hirabayashi et al., 2002) and since carrageenans bind the mAb anti- α Gal, they may also have potential for binding the galectins' CRD region. We thus chose to evaluate the inhibition potential of the carrageenans and their derivatives with Gal-1 and Gal-3. The design of the experiment included a competitive inhibition assay with thiodigalactoside (TDG), a CRD inhibitor of Gal-1 (K_d = 24 μ M) and Gal-3 (K_d = 49 μ M) since binding may be within the CRD or elsewhere on the galectins, such as observed with modified citrus pectin and galactomannan polysaccharides (Stegmayr et al., 2016).

Preliminary tests were done on all the carrageenans and their derivatives to determine the fractions that showed binding signal with Gal-1 and Gal-3. The carrageenans that showed strong signal intensity with the galectins were chosen for inhibition studies (Fig 3). The derivatives of commercial κ- and ι-carrageenans did not exhibit binding (data not shown). Our results indicate that carrageenans can be naturally-sourced glycan partners of Gal-1 and Gal-3 and they function as such when some structural requirements are met (Fig. 3a-d). Molecular mass, the structure of the non-reducing ends of oligo-carrageenans, and sulfation degree are features affecting galectin-binding properties of carrageenans. Thus, a decrease in molecular mass relative to the polymer improved binding signal intensity significantly with Gal-3 for all of the carrageenan derivatives tested and, to a lesser degree, with Gal-1 with only some of the carrageenan derivatives (Fig. 3a-d). This might be due to increased exo-binding as there are less endo-motifs in the shorter chain oligo-carrageenans. Unlike the oligo-carrageenans, with varied and significant capacities to bind Gal-1 and Gal-3 depending on sulfation degree and the treatment with the 3,6-anhydro-D-galactosidases, the polymers demonstrated less interactions with the galectins. Of the hybrid oligosaccharides tested, the oligo-carrageenans from F. lumbricalis were particularly interesting as the concentration used to show active binding was 0.5 mg/mL, whereas for the C. crispus oligo-carrageenans it was necessary to increase the concentration to 4 mg/mL. This is a striking difference between active concentrations of the f2 and c2 oligosaccharides for Gal-3 which is tempting to relate to the degree of sulfation on the oligosaccharides (Fig. 3 a,b).







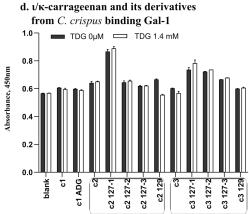


Figure 3. Binding of carrageenans and their derivatives from C. crispus (4 mg/mL) and F. lumbricalis (0.5 mg/mL) to Galectin 1 and 3 in the presence or absence of TDG (1396.3 μ M). Data are representative of three replicates in three independent experiments (Mean \pm Standard deviation). The designation of the samples is assembled by letters and figures: " ι " stands for commercial ι -

carrageenan, "κ" - commercial κ-carrageenan. "c" - carrageenan from *C. crispus*, "f" - carrageenan from *F. lumbricalis*. The numbering attributes to the initial polymer and after treatment with endolytic enzymes: 1 - polymer, 2 - ι-carrageenase, 3 - κ-carrageenase, 4 - digestion with both endolytic enzymes, 127-1, 127-2, 127-3, 129 correspond to the 3,6-anhydro-D-galactosidases ZgGH127-1, ZgGH127-2, ZgGH127-3 and ZgGH129, respectively; ADG – mixture of the four 3,6-anhydro-D-galactosidases. The asterisk (*) indicates significant differences <0.05 relative to blank, where no sugars were bound to the plate. The data are representative of triplicates from 2-4 independent experiments.

There appears to be two modes of binding to Gal-3. The first mode of binding is via the CRD region as seen with the inhibition by TDG and the κ/β -oligo-carrageenans from F. *lumbricalis* (Fig. 3a). Further hydrolysis with the 3,6-anhydro-D-galactosidases improves the binding signal significantly in some cases (f2 127-2, f2 129, f3 127-2). In general, the interaction with Gal-3 CRD was improved after 3,6-anhydro-D-galactosidase treatment of the F. *lumbricalis* oligosaccharides (Fig 1a). This suggests that a non-reducing β -galactose moiety is preferable for binding the Gal-3 CRD. ZgGH129 requires neo- β -carrabiose on the non-

reducing end, but can tolerate a subsequent ι - or κ -motif during hydrolysis (Wallace, Ficko-Blean, et al., 2020; Wallace, Guee, et al., 2020). Hence, the products after reaction would contain a neutral G on the nr end but the subsequent disaccharide motif may be sulfated. The tangible increase in binding might also be due to improved avidity to Gal-3 after hydrolysis of DAnr. The second mode of binding seems to be non-CRD binding-site mediated as in many cases the TDG shows poor to no inhibition with Gal-3 when incubated with the *C. crispus* oligocarrageenans (Fig 3b). Notably, their active concentration was also 8-fold higher than for the *F. lumbricalis* κ/β -carrageenan derivatives. In any case, these κ/ι -derivatives demonstrated decreased interaction with Gal-3 relative to the κ/β -carrageenan derivatives using this binding methodology. Overall, the *F. lumbricalis* κ/β -oligo-carrageenans have strong potential as saccharide-based ligands selective and specific for the Gal-3 CRD.

Gal-1 binding to the carrageenans and oligosaccharide derivatives was not inhibited by TDG, suggesting that all the interactions with the carrageenans are distal to the CRD carbohydrate-binding region (Fig 3cd). Still some of the hybrid κ/β -oligo-carrageenans from *F. lumbricalis* showed improved signal intensity (Fig 3c, f3 129, f4, f4 127-2, f4 129). The *C. crispus* κ/ι -oligo-carrageenans showed poor binding in general to Gal-1 (Fig 3d) with a signal similar to that of the blank, with the exception of the c2 and c3 samples treated with ZgGH127-1.

Conclusion

The binding of carrageenan polymers relative to the enzymatically produced oligosaccharides suggests anti-αGal antibody binding through both groove-type and end-on insertion. Gal-3 CRD revealed exo-binding as the CRD demonstrated a preference for the oligocarrageenans. Furthermore, treatment with one of the four 3,6-anhydro-D-galactosidases, ZgGH127-1, ZgGH127-2, ZgGH127-3, or ZgGH129, inflicted a substantial impact on Gal-3 CRD binding to the oligosaccharides and, thus, the structure of the non-reducing ends of oligocarrageenans are a molecular determinant to binding the Gal-3 CRD. Our findings make carrageenans alluring glycomimetics for future exploration as multivalent ligands for human serum anti-αGal IgM antibodies and as specific ligands selective for Gal-3 over Gal-1. These revelations could suggest a molecular basis for widely studied oligo-carrageenans' antitumor properties and provide more prospective potential of these molecules to anti-tumor research (Calvo et al., 2019; Torres, Florez-Fernandez, & Dominguez, 2021). One of the mechanisms to turn cold tumors with low responsiveness to immunotherapy into so-called hot or inflamed

669	tumors is by increasing local inflammation (Ochoa de Olza, Navarro Rodrigo, Zimmermann,
670	& Coukos, 2020). Local inflammation is a natural outcome of such potent inducers of
671	complement as the Gala(1,3)Gal reactive antibodies. Carrageenans and their derivatives are
672	also prospective therapeutic agents in wound healing since they have the potential to affect both
673	immune response and the activity of coagulation cascade components.
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