

Structure-activity relationship of Citrus segment membrane RG-I pectin against Galectin-3: The galactan is not the only important factor

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ABSTRACT

Rhamnogalacturonan I (RG-I) pectin are regarded as strong galectin-3 (Gal-3) antagonist because of galactan sidechains. The present study focused on discussing the effects of more structural regions in pectin on the anti-Gal-3 activity. The water-soluble pectin (WSP) recovered from citrus canning processing water was categorized as RG-I pectin. The controlled enzymatic hydrolysis was employed to sequentially remove the α -1,5-arabinan, homogalacturonan and β -1,4-galactan in WSP. The Gal-3-binding affinity K_D (kd/ka) of WSP and debranched pectins were calculated to be 0.32 μ M, 0.48 μ M, 0.56 μ M and 1.93 μ M. Moreover, based on the more sensitive cell line (MCF-7) model, the IC_{30} value of WSP was lower than these of modified pectins, indicating decreased anti-Gal-3 activity. Our results suggested that the total amount of neutral sugar sidechains, the length of arabinan and cooperation between HG and RG-I played important roles in the anti-Gal-3 activity of WSP, not the Gal/Ara ratio or RG-I/HG ratio. These results provided a new insight into structure-activity relationship of citrus segment membrane RG-I as a galectin-3 antagonist and a new functional food.

1. Introduction

Cancer is one of the biggest health problems with global deaths reaching 9.6 million (WHO 2019). Galectin-3, a β -galactoside binding protein, has been regarded as a “culprit molecule” promoting the development of cancers by accelerating the proliferation, adhesion, metastasis and inhibiting the cell apoptosis (Nguyen et al., 2018; Rajput et al., 2016). The expression of Gal-3 differs in various cell types, but the immune-cells and collagen-producing cells seem to be the primary sources of Gal-3 (Suthahar et al., 2018). Gal-3 is up-regulated in many cancers including colorectal, breast and lung cancer (Jiang, Dang, Gong, & Guo, 2019; Newlaczyl & Yu, 2011). The biological properties of Gal-3 are associated with its binding activity, which is due to the interaction of several glycoconjugates with its conserved carbohydrate-recognition domain (CRD) (Laaf, Bojarova, Elling, & Kren, 2019).

Pectin, a plant cell wall polysaccharide, consist predominantly of

homogalacturonans (HG), a few rhamnogalacturonan I (RG-I) and a minor rhamnogalacturonan II (RG-II) fraction (Yapo, Wathelet, & Paquot, 2007; Voragen, Coenen, Verhoef, & Schols, 2009; Yapo, Lerouge, Thibault, & Ralet, 2007, 2011). RG-I is viewed as one of the structural components containing the highest rhamnose content ramified in pectin, consisting of repeating $[-2)\text{-}\alpha\text{-L-Rhap-(1}\rightarrow\text{4)-}\alpha\text{-D-GalpA-(1}\rightarrow\text{)]}$ units, where a portion of the rhamnose residues are substituted with varying neutral and branched sugar chains, including linear or branched α -(1 \rightarrow 5)-arabinan, linear β -(1 \rightarrow 4)-galactan, branched β -(1 \rightarrow 3,6)-galactan and arabinogalactan (Liu et al., 2018; Maxwell, Belshaw, Waldron, & Morris, 2012; Mohnen, 2008; Ridley, O'Neil, & Mohnen, 2001). Due to the high-temperature and acid extraction method for commercial pectin, RG-I usually exists in agro-industrial by-products after the extraction process (Jiang & Du, 2017; Mao et al., 2020; Pi, Liu, Guo, Guo, & Meng, 2019; Yan et al., 2018), with much more amount than HG regions (Khodaei & Karboune, 2018). Recently, many RG-I-

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from fruits and vegetables have been reported to possess excellent anti-Gal-3 activity, such as pumpkin RG-I, citrus peel RG-I, and ginseng RG-I (Wu et al., 2020; Xue et al., 2019; Zhao et al., 2017; Zhang, Lan et al., 2016; Gao et al., 2013). Thus, the novel RG-I pectin is a promising galectin-3 antagonist. Published studies have also revealed that ginseng pectins and chemically modified citrus pectin with different structure exhibited various effects on galectin-3 action, and the short β -1,4 galactan side chains in RG-I play important roles (Zhang, Lan et al., 2016; Gao et al., 2013). However, most of these studies focus on the binding affinity of natural pectic polymers or degraded oligomers. No systematic evaluation of the specific structural region required for anti-Gal-3 activity has been carried out. Such an evaluation requires a modification method that can target the removing specific regions of pectin instead of its random and violent chemical degradation.

Enzyme digestion is an environment-friendly and green approach to achieve mild modification. Different carbohydrate active enzymes can remove targeted sugar chains without any changes in the other domains of pectin (Oosterveld, Pol, Beldman, & Voragen, 2001; Zheng & Mort, 2008). Hence, enzyme catalyzed modification is a preferred method for studying the structure-function relationship of RG-I pectin.

In this study we obtained citrus segment membrane RG-I pectin, the water-soluble pectin (WSP) from citrus canning processing water. The hypothesis was that WSP can antagonize Gal-3 and all of the side chain structures contribute to its anti-Gal-3 activity. The aim of this study is to determine the anti-Gal-3 bioactivity of WSP and discuss the structure-functionality relationship. Three enzymes were used to partly remove the arabinan, homogalacturonan and galactan. We then analyzed the structural changes between WSP and debranched pectin samples to assess their ability to antagonize Gal-3 and determine the important bioactive structural factors. This study is significant, as it should lead to the development of novel Gal-3 antagonists and safe food additives from this natural product and also provide for the potential valorization of a citrus derived agriculture industry waste.

2. Material and methods

2.1. Material

The basic pectin source was collected from citrus fruit canning factories (Ningbo, China). Endo-1,5- α -arabinanase (EC 3.2.1.99, 400 U), endo-polygalacturonase (EC 3.2.1.15, 5000 U), endo-1,4- β -galactanase (EC 3.2.1.89, 1000 U) were purchased from Megazyme International Ireland Ltd. (Wicklow, Ireland). Modified citrus pectin (MCP) was purchased from EcoNugenics (Santa Rosa, CA, USA). Monosaccharide standards, 1-phenyl-3-methyl-5-pyrazolone (PMP) and D₂O were all purchased from Sigma-Aldrich (Shanghai, China). All other chemicals were analytical grade and purchased from Aladdin Co. (Shanghai, China).

2.2. Preparation of WSP and enzymatic modification

The crude pectin was recovered from alkaline water during the citrus segment membrane removal process from fruit canning factories (Chen et al., 2017). The basic processing water underwent a two-step filtration using 200 and 400 meshes filters, neutralization in a pH adjusted reactor, vacuum concentration at 70 °C, and precipitation using ethanol. The precipitates were recovered by alcohol washing and oven-dried at 55 °C for 24 h. The dried pectin powder was dissolved in distilled water (0.5 g/100 mL) followed by gentle heating (60 °C, stirring at 600 rpm for 40 min), centrifugation (8000 rpm, 10 min) to obtain the original sample, water-soluble pectin (WSP), and finally freeze-dried.

The method for enzyme-modification of WSP is shown in Fig. 1. WSP (200 mg) was re-dissolved in 0.1 M citric acid-sodium acetate buffer with a final concentration of 2 mg mL⁻¹ and subjected to sequential enzymatic hydrolysis by endo-1,5- α -arabinanase (0.2 U mL⁻¹ at 40 °C for 48 h, pH 4.0), endo-polygalacturonase (0.05 U mL⁻¹ at

40 °C for 72 h, pH 5.5) and endo-1,4- β -galactanase (0.05 U mL⁻¹ at 40 °C for 24 h, pH 4.0) under gentle stirring (150 rpm). The resultant hydrolysates were heated at 90 °C for 5 min for terminating enzymatic activity and then, dialyzed (MWCO 10 kDa) against deionized water for three days, and freeze-dried. EP-1 referred to WSP debranched by endo-1,5- α -arabinanase; EP-2 was modified EP-1 treated by endo-polygalacturonase; EP-3 was endo-1,4- β -galactanase-modified EP-2.

2.3. Molecular weight determination of SEC-MALLS

The homogeneity and relative molecular weight of WSP and enzyme-treated pectins were determined using high-performance (HP) size exclusion chromatography (SEC) (on a OHPak SB-G guard column, SB-806 HQ and SB-804 HQ column, 7.8 × 300 mm, Shodex, Japan), and detected with a multi-angle laser light scattering (DAWN HELEOS II, Wyatt Technology, USA), by viscometry (ViscoStar™ III, Wyatt Technology, USA) and by refractive index (SEC-MALLS-VISC-RI) at 25 °C, according to Wei et al., 2019. Samples were dispersed directly in purified water (5 mg mL⁻¹) and filtered through a 0.22 μ m filter membrane twice. Solution (50 μ L) was injected through a sample loop and the running time was 50 min. The mobile phase was 0.15 M NaCl solution containing 0.02 % NaN₃ (pH 7.0) with a flow rate of 0.5 mL min⁻¹. Data analysis was performed using ASTRA software, version 7.1.2 (Wyatt Technology).

2.4. Monosaccharide analysis

The monosaccharide composition of every pectic sample was analyzed after derivatization with 1-phenyl-3-methyl-5-pyrazolone (PMP) using high performance liquid chromatography (HPLC) on a Waters e2695 (Waters, US) equipped with a Zorbax Eclipse XDB-C18 column (250 mm × 4.6 mm, 5 μ m, Agilent, USA) and detected using a 2489 UV/Vis Detector (Waters, US) at 250 nm as described by Zhang et al., 2018. Briefly, 2 M trifluoroacetic acid (TFA) was first employed to hydrolyze the pectin samples (2 mg) at 110 °C for 8 h in a sealed ampoule and then evaporated with methanol under nitrogen blower to completely remove TFA. Pre-column derivatization was performed using PMP solution (0.5 M, in methanol) at 70 °C for 30 min. Finally, chloroform was used to extract excess reagent three-times. The supernatant was filtered through a 0.22 μ m nylon membrane and 10 μ L of the resulting solution was injected for analysis. The mobile phases were: solvent A, potassium phosphate buffer (0.05 M, pH 6.9) with 15 % acetonitrile, and solvent B, the same buffer with 40 % acetonitrile.

2.5. ¹H NMR analysis and FT-IR spectroscopy

NMR spectroscopy was performed on pectins (10–15 mg) dissolved in 0.5 mL of D₂O (99.96 %) that had been lyophilized twice to replace exchangeable protons (Li et al., 2019) before final dissolution in 0.5 mL D₂O. ¹H NMR spectra was recorded on Agilent DD2–600 MHz spectrometer (Agilent, USA) at room temperature.

The FT-IR spectra of four pectin samples were recorded by using a Nicolet iN10 instrument (Thermo Fisher Scientific, USA) (Zhang et al., 2018). Samples (2 mg) were mixed with KBr powder and then pressed into pellets for FT-IR measurements in the range of 4000–400 cm⁻¹.

2.6. Atomic force microscopy (AFM)

Microstructures of pectic samples were acquired by AFM according to Zhang et al., 2018. Briefly, the pectic samples were dissolved in ultrapure water with a final concentration of 1 mg mL⁻¹ under continuous stirring for 2 h and then, incubating at 80 °C for 2 h. Subsequently, the diluted pectin solutions (10 μ g mL⁻¹) containing equal mass of sodium dodecyl sulfate (SDS) were stirred for 24 h to depolymerize the macromolecules. A 10 μ L of sample solution was filtered through a 0.22 μ m filter and dropped onto freshly cleaved mica

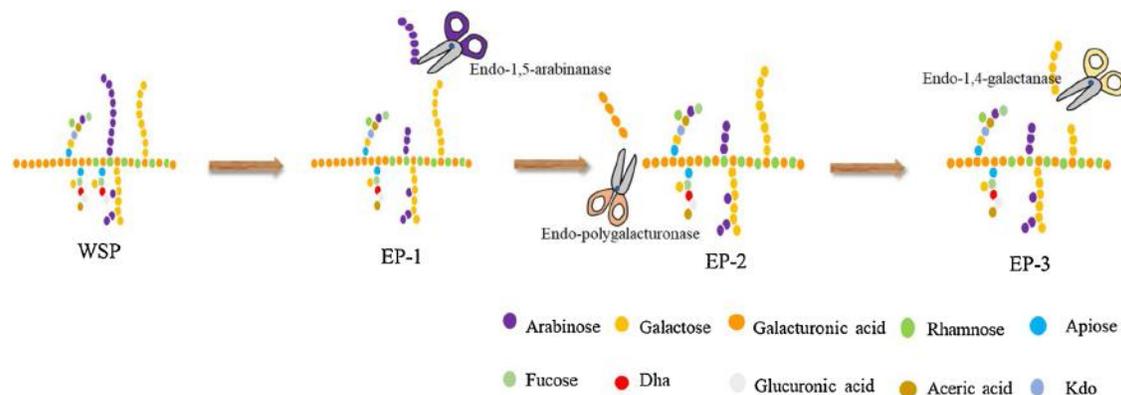


Fig. 1. The sequence for enzyme-modification.

substrate. Afterwards, the mica substrates were air-dried and then observed by AFM (XE-70, Park Scientific Instruments, Suwon, Korea), operated in a tapping mode at room temperature. The probe was a classical silicon cantilever (Si_3N_4) with a spring constant of 0.2 N/m and a resonance frequency of approximately 13 kHz. XEI data processing software was used for image manipulation.

2.7. Binding assay by surface plasmon resonance (SPR) assay

The SPR assay was performed following our previous method (Wei et al., 2019). Binding affinities of WSP and treated pectins to Gal-3 were performed on a BIAcore 3000 system (GE Healthcare, USA). Gal-3 protein was immobilized on an activated CM5 sensor chip (GE Healthcare, USA). Pectic samples were dispersed in HBS-EP buffer (0.01 M HEPES, 0.15 M NaCl, 3 mM EDTA, and 0.005 % surfactant P20, pH 7.4) and gradiently diluted to different concentrations. Each concentration of these sample was injected individually at a flow rate of 30 $\mu\text{L}/\text{min}$ for 3 min, followed by dissociation and regeneration. Data analysis was performed through BIAcore 3000 evaluation software.

2.8. Anti-proliferation activity assay

Anti-proliferation activity was evaluated in MCF-7 (human breast cancer, cultured in DMEM with 10 % FBS) and A549 (human lung carcinoma, cultured in RPMI 1640 with 10 % FBS) cell lines. Both of the cells were incubated at 37 °C and 5% CO_2 . Briefly, 100 μL of the cells were seeded in a 96-well plate at a concentration of $5 \times 10^3/\text{well}$. After 24 h of incubation, 100 μL of pectin sample solution at various concentration (2000 $\mu\text{g mL}^{-1}$, 1000 $\mu\text{g mL}^{-1}$, 500 $\mu\text{g mL}^{-1}$, 250 $\mu\text{g mL}^{-1}$, 125 $\mu\text{g mL}^{-1}$, 63 $\mu\text{g mL}^{-1}$) were slowly added to each well. MCP served as the positive control. After incubating for 48 h, 20 μL of methyl thiazolyl tetrazolium (MTT, 5 mg mL^{-1}) was added and the cells were

further incubated for 4 h at 37 °C. After removing the medium, a volume of 150 μL dimethyl sulfoxide (DMSO) was added to each well to dissolve the resulting formazan. The absorbance was measured at 490 nm using a microplate reader (Bio-Rad, USA) and the anti-proliferation effect was calculated as

$$\text{Anti-proliferation effect (\% control)} = [1 - (A_2 - A_0) / (A_1 - A_0)] \times 100.$$

(A_0 , A_1 , A_2 respectively represent the absorbance of the system containing only medium, the absorbance of the system with cells, the absorbance of the system with pectin sample or MCP.)

2.9. Statistical analysis

All experiments were carried out in triplicate and the average of three independent experiments were used as the statistical result. The one-way analysis of variance (ANOVA) was conducted, followed by Duncan's multiple range test at significance level of 0.05 using SPSS software (version 19; IBM Corporation, New York, NY, USA).

3. Results and discussion

3.1. Molecular properties

The relative molecular information of WSP, EP-1, EP-2, EP-3 were revealed in Table 1, which were calculated using Astra 6.1. The weight average of molar mass (Mw) of these four samples were $5.93 \times 10^5 \text{ g mol}^{-1}$, $5.26 \times 10^5 \text{ g mol}^{-1}$, $4.72 \times 10^5 \text{ g mol}^{-1}$, $3.70 \times 10^5 \text{ g mol}^{-1}$, respectively. The molecular weight of WSP and these enzyme-modified pectins gradually decreased, but all of these samples had a molecular weight significantly larger than the Gal-3 protein. Enzymatic modification removed specific side chains and no remarkable changes in the average molecular weight was observed. Polydispersity coefficients (Mw/Mn) were 3.15, 5.79, 2.36, 2.10, respectively, which indicated

Table 1
Molecular and conformation parameters of WPS and enzyme-modified fractions.

Parameters	Samples			
	WSP	EP-1	EP-2	EP-3
Mn (g/mol)	$(1.88 \pm 0.03) \times 10^5$	$(0.91 \pm 0.01) \times 10^5$	$(2.00 \pm 0.03) \times 10^5$	$(1.76 \pm 0.01) \times 10^5$
Mp (g/mol)	$(1.62 \pm 0.02) \times 10^5$	$(0.41 \pm 0.01) \times 10^5$	$(1.46 \pm 0.03) \times 10^5$	$(1.05 \pm 0.03) \times 10^5$
Mw (g/mol)	$(5.93 \pm 0.02) \times 10^5$ ^a	$(5.26 \pm 0.08) \times 10^5$ ^a	$(4.72 \pm 0.01) \times 10^5$ ^b	$(3.70 \pm 0.06) \times 10^5$ ^c
Mw/Mn	3.15 ± 0.03 ^b	5.79 ± 0.01 ^c	2.36 ± 0.03 ^a	2.10 ± 0.02 ^a
Rh (nm)	0.449 ± 0.02 ^a	0.576 ± 0.02 ^a	0.635 ± 0.05 ^a	0.654 ± 0.03 ^a
[η] (mL/g)	10.788 ± 0.01	8.22 ± 0.01	12.612 ± 0.04	10.12 ± 0.03
K (mL/g)	$(4.22 \pm 0.04) \times 10^{-3}$	$(1.77 \pm 0.01) \times 10^{-3}$	$(1.18 \pm 0.04) \times 10^{-4}$	$(5.54 \pm 0.04) \times 10^{-4}$
α	0.72 ± 0.01 ^a	0.77 ± 0.01 ^a	0.98 ± 0.04 ^a	0.83 ± 0.03 ^a

Mn: number-average of molar mass; Mp: peak-average of molar mass; Mw: weight-average of molar mass; Rh: hydrodynamic radius. Means with the different letter are significantly different between columns ($p < 0.05$). ^aSignificant against the Mw of Gal-3 (31 kDa) at $p < 0.05$.

Table 3
1H NMR chemical shifts assignments of WSP, EP-1, EP-2 and EP-3.

Residues	Samples			
	WSP	EP-1	EP-2	EP-3
→3)-α-Ara-(1→	5.18(H1);4.16(H2); 4.25(H4)	5.18(H1);4.15(H4)	5.18(H1)	5.21(H1)
→5)-α-Ara-(1→	5.11(H1);4.32(H2); 4.13(H3);4.20(H4); 4.92(H5)	5.12(H1);4.30(H2); 4.13(H3)	5.14(H1);4.30(H2); 4.19(H4)	5.16(H1); 4.30(H2)
→3,5)-α-Ara-(1→	5.14(H1);4.07(H3); 4.44(H4)	5.14(H1);4.41(H2); 4.15(H3);4.46(H4)	5.14(H1);4.43(H2); 4.16(H3);4.47(H4)	4.43(H2); 4.15(H3)
→4)-α-GalA-(1→	5.09(H1);3.99(H3); 4.44(H4);4.69(H5)	5.09(H1);4.44(H4)	5.11(H1);4.68(H5); 4.46(H4)	5.11(H1); 4.46(H4); 4.68(H5)
→3)-β-Gal-(1→	4.16(H3)	4.61(H1)	4.64(H1)	4.64(H1)
→4)-β-Gal-(1→	4.66(H1);4.44(H4)	4.63(H1);3.54(H2)	4.66(H1); 3.55(H2)	4.67(H1); 3.55(H2)

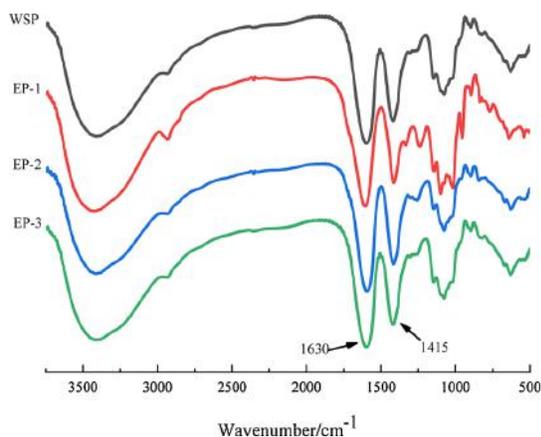


Fig. 3. IR spectra (% transmittance as a function wavenumber) of WSP, EP-1, EP-2, EP-3.

treatment, these treated pectin still exhibited low level of esterification, which indicated that enzymatic modification didn't change the esterification pattern. The wavenumbers between 1010 and 1150 cm^{-1} (e.g. 1024 cm^{-1} , 1076 cm^{-1} , 1145 cm^{-1}) corresponded to C–O and C–C vibration of glycosidic bonds and the pyranose (Li et al., 2019; Liu et al., 2016; Sucheta, Misra, & Yadav, 2020) and no changes caused by enzyme modification were found in these pyranose configuration.

3.4. Surface morphology

AFM technique was next employed to image the surface morphology of WSP, EP-1, EP-2 and EP-3 (Fig. 4). The long branching sugar chains could be observed in WSP, which might be the α -1,5-arabinan side-chains since WSP contained high amount of arabinose. After hydrolysis of endo-1,5- α -arabinanase, many short straight chains occurred, suggesting the notably decreasing in the amount of neutral sugar in EP-1. And then endo-polygalacturonase degraded the arborization-like structure, which made EP-2 appear to be short chains with many short branches, and the chain-chain interaction was more intensive in EP-2. It seemed that some HG regions were embedded in the trunk of RG-I and some existed as sidechain of RG-I. At last, the endo-1,4- β -galactanase resulted in significant sidechains depolymerization and EP-3 contains short backbone with very few branches. In addition, some irregular spherical conformation with the heights between 1–5 nm were observed within these molecule shapes, which implicated the existence of polymer. The loss of branching and homogalacturonan can reduce aggregated pectic complexes (Round, Rigby, MacDougall, & Morris, 2010), which could be observed in EP-2 image. Taken together, these AFM results were consistent with the semi-flexible chain conformation in SEC-MALLs analysis.

3.5. Binding affinity on Gal-3

The result of SPR analysis presented smooth binding curves between the samples and Gal-3 (Fig. 5). Gal-3 is a β -galactose-binding and tumor-related protein with key roles in accelerating cancer, fibrosis, carcinogenesis, and several other diseases. Therefore, the excellent anti-Gal-3 activity of RG-I pectin make it a non-toxic and effective inhibitory agent in disease therapy. Gal-3 inhibitors, such as MCP and ginseng pectin, have been reported to occupy the carbohydrate recognition domain (CRD) inside Gal-3 protein molecule via the galactan side chain (Gao et al., 2013; Zhang, Lan et al., 2016), which can explain the decrease of KD value between EP-2 and EP-3 (Table 4). Due to the effect of endo-1,4- β -galactanase, half of the galactan sidechains were removed in EP-2, which led up to approximately 3-fold decrease in the binding affinities. In addition, binding kinetics of WPS, EP-1 and EP-2 to Gal-3 were 0.321 μM , 0.48 μM , 0.56 μM , respectively, which suggested that partly removal of α -1,5-arabinan and HG region could impair the WSP's activity against Gal-3. Regarding to the results of SEC-MALLS analysis, all of the four molecular weights were between 3×10^5 and 6×10^5 g mol^{-1} , over 10-times larger than the molecular weight of Gal-3 (3.1×10^4 g mol^{-1}) (Fukumori et al., 2007). Compared to chemical modification, which decreases the Mw of pectin to 7.9×10^3 g mol^{-1} , enzyme digestion did not cause remarkable differences in molecular weight of the four samples, and no evident changes in the structural pattern were observed by FT-IR. Thus, the effects of different sidechains on the anti-Gal-3 activity could be assessed. Different domains inside Gal-3 can bind to different types of polysaccharides (Miller et al., 2016). For instance, the N-terminal (NT) tail of Gal-3 can bind to rhamnoga-lacturonan and F-face of CRD can interact with larger different polysaccharides (Miller et al., 2017). Therefore, the WSP might exert its arabinan sidechains to interact with the F-face in β -strands, which might account for the loss of activity caused by α -1,5-arabinanase (Gao et al., 2012). According to previous report, arabinose residues have positive or negative influence on the anti-Gal-3 activity due to different location (Gao et al., 2013) and α -1,5-arabinan exhibited negative effects, but the results in this study suggested that α -1,5-arabinan in the citrus RG-I also contributed to the activity, which may due to different length of sugar chains. In addition, the structure mode of RG-I backbone also make sense to the activity against Gal-3 (Zhang, Lan et al., 2016; Gao et al., 2013), so some HG segment is essential for WSP's activity, which is consistent with previous result (Maxwell et al., 2015). It has been reported that Gal/Ara ratio plays important role in binding Gal-3 (Gao et al., 2013). However, the ratio of Gal/Ara in WSP could not remarkably affect the anti-Gal-3 property. Compared to that of WSP (0.24), the Gal/Ara ratio of EP-1 increased to 1.56, but the binding affinity of WSP still decreased. Also, the importance of RG-I/HG ratio was not observed, though the RG-I and HG can work cooperatively. Conversely, more amount of neutral sugar sidechains might promote the anti-Gal-3 activity, since the decrease of binding affinity between pectin and Gal-3 was accompanied by a reduction in the degree of branching. In general, the SPR assay data suggested that all of the

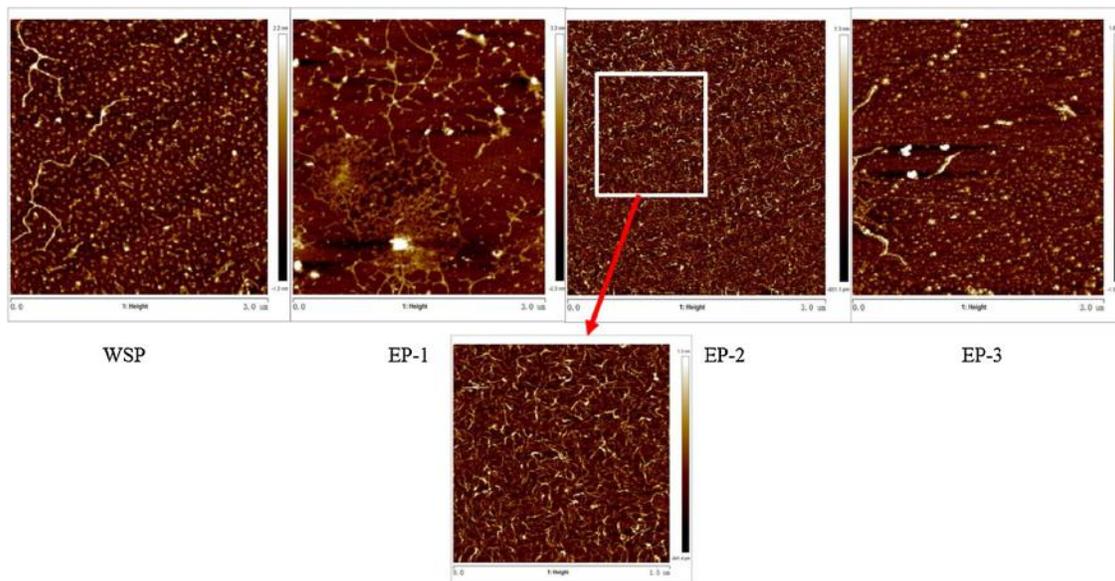


Fig. 4. Representative topographical AFM images of WSP, EP-1, EP-2, EP-3 (Each sample were at the concentration of $10 \mu\text{g mL}^{-1}$).

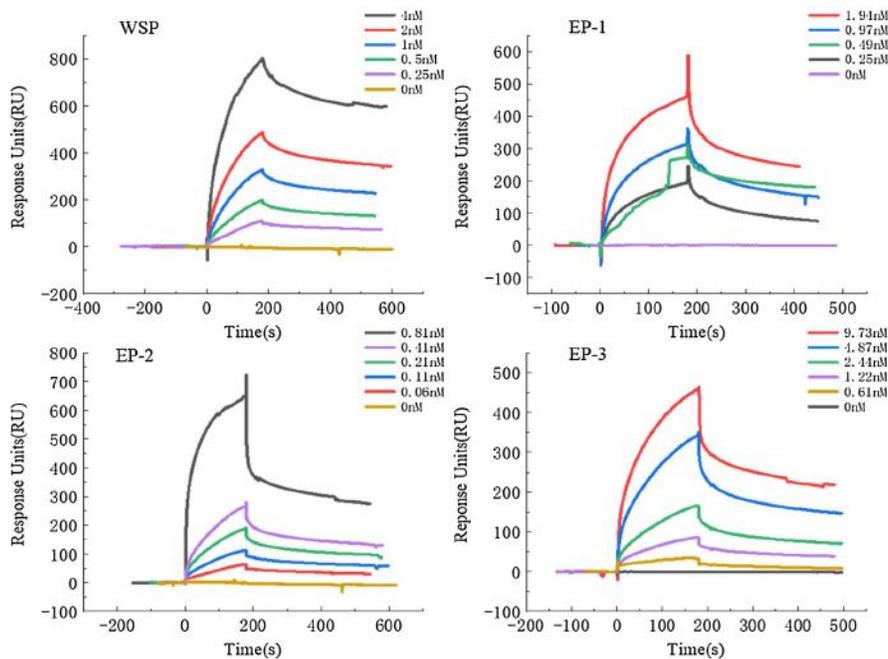


Fig. 5. SPR analysis of the binding affinities of WSP, EP-1, EP-2, EP-3 to galectin-3.

Table 4

The binding affinities of the four samples to galectin-3.

Samples	Mw (KDa)	k_a (1/Ms)	k_d (1/s)	KD(k_d/k_a) (μM)
WSP	593	1.86×10^6	5.98×10^{-4}	0.321
EP-1	526	5.34×10^6	2.56×10^{-3}	0.48
EP-2	472	2.20×10^6	1.23×10^{-3}	0.56
EP-3	370	7.85×10^5	1.52×10^{-3}	1.93

structural regions of WSP played an essential role in anti-Gal-3 activity, and the galactan sidechains possessed the largest effects. Moreover, WSP appeared to be a stronger Gal-3 inhibitor than most other

polysaccharides (Gao et al., 2013; Wei et al., 2019; Zhao et al., 2017).

3.6. Anti-proliferation activities

One of the biological properties of Gal-3 is promoting the proliferation of cancer cells, therefore we employed MTT assay to determine the effects of WSP on Gal-3 function. The anti-proliferation activities of the four samples were tested on A549 (Fig. 4A) and MCF-7 (Fig. 4B) cell lines *in vitro* and MCP was employed as the positive control. As shown in Fig. 6, the proliferation of A549 and MCF-7 cell lines were evidently inhibited by all the samples in a dose-dependent manner. The IC_{20} values of the four samples against A549 were

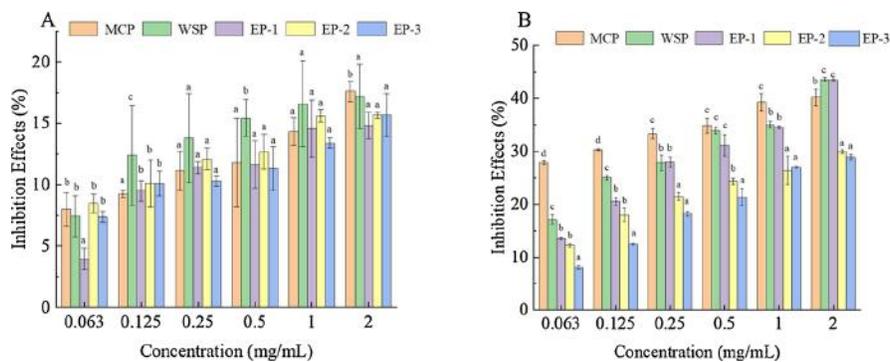


Fig. 6. Inhibition effects on A549 (A) and MCF-7 (B) cell lines viability. Means with the different letter are significantly different ($p < 0.05$), and error bars represent standard deviation.

0.12 mg mL⁻¹, 3.83 mg mL⁻¹, 11.19 mg mL⁻¹, 19.36 mg mL⁻¹, respectively. But in Fig. 6A, it seemed that no significant difference appeared among the groups treated with different samples, indicating that the anti-proliferation effects on A549 cells of these pectic fractions was not consistent with the anti-Gal-3 activity. The the IC₃₀ values against MCF-7 were 0.33 mg mL⁻¹, 0.43 mg mL⁻¹, 1.84 mg mL⁻¹, 1.97 mg mL⁻¹, respectively. In addition, the four samples exhibited the remarkably inhibitory effects of WSP > EP-1 > EP-2 > EP-3 against MCF-7 cells, especially at the concentration of 0.125 mg mL⁻¹, which showed the same decreasing trend as the anti-Gal-3 activity. So MCF-7 was more sensitive to Gal-3 inhibitors, which may be owing to the higher expression level of Gal-3 compared to A549 (Wei et al., 2019). These data suggested that WSP and these debranched samples might exert the anti-tumor activity as a galactin-3 antagonist against cancer cells with high expression of Gal-3. Moreover, WSP even exhibited significantly stronger inhibitory effect than that of MCP at the concentration of 2 mg mL⁻¹ in Fig. 6B, implicating the potential of WSP for curing cancers. Compared to other plant polysaccharides, the anti-proliferation activity of WSP against A549 cells was higher than the polysaccharides from *Flammulina velutipes* (Kom.) (Yang et al., 2012) and *Allium macrostemon Bunge* (Zhang, Wang, & Zeng, 2015), the inhibition effect against MCF-7 was stronger than an acidic polysaccharide from *Angelica sinensis* (Oliv.) Diels (Zhang, Zhou et al., 2016), indicating the potential antitumor activity.

4. Conclusions

The WSP recovered from canning processing water was determined to be RG-I type pectin with high amount of neutral sidechains, especially arabinan. Functionally, WSP exhibited very strong binding affinity with Gal-3, compared to most other polysaccharides. After partly removing the α -1,5-arabinan and β -1,4-galactan, the anti-Gal-3 activity of WSP decreased, indicating that besides the galactan sidechains, arabinan also contributed to the binding affinity between RG-I pectin and Gal-3 and long α -1,5-arabinan sidechains showed high activity. The total amount of neutral sidechains played an important role, not the ratio of Gal/Ara. In addition, our results suggested that RG-I and HG regions in WSP acted cooperatively, but the bioactivity of WSP was not significantly related to the ratio of RG-I/HG. This study provides some important information of the structure-function relationship of the citrus segment membrane pectin. These findings are also helpful in developing novel Gal-3 antagonists with low toxicity as well as the improved food additives in the industry.

CRedit authorship contribution statement

Dongmei Wu: Data curation, Formal analysis, Investigation, Project administration, Writing - original draft. Jiaqi Zheng: Software. Weiwei

Hu: Supervision. Xiaoliang Zheng: Methodology. Qiaojun He: Resources. Robert J. Linhardt: Writing - review & editing. Xingqian Ye: Funding acquisition, Writing - review & editing. Shiguo Chen: Methodology, Funding acquisition, Writing - review & editing.

Declaration of Competing Interest

The authors declare that they have no conflict of interest.

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