

Purification, partial characterization and antimicrobial activity of Lectin from *Chenopodium Quinoa* seeds

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Abstract

A novel lectin was isolated from the seeds of *Chenopodium quinoa*. To achieve this end, the crude extract from the quinoa was submitted to two purification steps, Sephadex G50 and Mono Q. The hemagglutinating activity showed that this lectin agglutinates human erythrocytes. Its activity is inhibited by glucose and mannose, and remained stable under a wide range of pH levels and temperatures. The quinoa lectin was found to be a heterodimeric lectin of approximately 60 kDa, consisting of two subunits of approximately 25 kDa and 35 kDa. This lectin had its antimicrobial activity tested against several bacteria strains and effectively inhibited three strains. These strains were all Gram-negative, making this lectin a promising antimicrobial tool.

Keywords: antimicrobial; *Chenopodium quinoa*; glucose/mannose-specific; lectin; seeds.

Practical Application: The purification and characterization of this novel lectin allow its study for a diversity of applications such as antibacterial, anticancer, anti-inflammatory and biotechnology and its further use as drugs for human diseases.

1 Introduction

Lectins are oligomeric carbohydrate-binding proteins that are involved in various biological recognition processes (Ghosh & Mandal, 2012). They are a large and diverse group of proteins that have the ability to bind reversibly to monosaccharides and oligosaccharides, which can be defined as a class of structurally diverse proteins or glycoproteins (Sharon, 2008). These proteins are widely distributed in nature, being found in microorganisms (Visini et al., 2015), animals (Lundbo et al., 2015) and plants (Silva et al., 2007).

More than a hundred lectins have been isolated and characterized to varying degrees, primarily in seeds. However, the structural diversity of these proteins requires further studies to improve the understanding of their different molecular properties, biochemical and functional. This would allow for a wider application of lectins as tools in various fields (immunological, biochemical and biotechnological), especially those that exploit the protein-carbohydrate interaction. The ability of certain lectins to distinguish between different cell types has encouraged and justified the purification and characterization of lectins originating from various species (Silva et al., 2007).

It is unlikely that all of the functions and applications of lectins have been identified. Their easy occurrence, their high thermal stability and mainly, their well-defined carbohydrate binding properties are important tools in the field of glyco-science, immunology and biotechnology, healthcare and pharmaceutical industries as biochemical tools for different studies (Nasi et al., 2009).

Their ability to reversibly bind to a specific saccharide has attracted the attention of scientists. Each lectin has a characteristic sugar-binding specificity profile indicating that they are able to recognize different glycotopes (Kamiya et al., 2012). Based on the specificity of the sugar bound, plant lectins are classified into groups, such as Gal/GalNAc-specific, glucose/mannose-specific, fucose specific, GlcNAc specific, and sialic acid specific lectins (Goldstein & Poretz, 1986). Since mannose is widespread in animals, insects and microorganisms, mannose-binding lectins are considered biologically important proteins (Wong et al., 2008).

Despite their differences in specificity for carbohydrates, lectins are similar relative to their physicochemical properties. They are usually composed of two or four subunits with molecular weights of 25-45 kDa, each one with a carbohydrate-binding site (Nasi et al., 2009). C-type lectins present only homodimers linked by disulfide bonds (Arlinghaus & Eble, 2012).

The physiological function of plant lectins has not been openly proved. It is understood that the lectins are responsible for the plant defense and specific protein-carbohydrate interaction mediators within the plant cells (Ahmad et al., 2011).

Many plants produce seeds that are rich in protein and carbohydrates and therefore are commonly infested by pests (Coelho et al., 2010). Lectins are a potential type of molecule candidate for controlling pests and diseases (Araújo-Filho et al., 2010). The mechanisms of recognition involve specific interactions between the protein receptors and saccharide residues (Lee &

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Peng, 2012). The lectins that bind to accessible carbohydrate residues from the cell wall or cell membrane trigger a cascade of biological responses.

The capacity for identifying and binding glycoconjugates from the microorganism's surface is exclusive to lectins. Consequently, they are capable of inhibiting the motility and multiplication of microorganisms (Oliveira et al., 2007). The interaction between carbohydrates and lectins act in many biological processes, and intermediate a wide range of activities, including bacterial and fungal growth inhibition (Klafke et al., 2013).

The pseudo cereal *Chenopodium quinoa* was originally cultivated in Peru, Bolivia, Equator, Chile and Argentina at various latitudes and altitudes (Bhargava et al., 2006; Pesoti et al., 2015). Due to its potential as a food source and the limited production in South America, quinoa is now being introduced in Europe, North America, Africa and Asia (Bhargava et al., 2006). This has contributed to its popularization, especially in developing countries, as an alternative food source. It is an important food source considering its high nutritional content, in addition to its high levels of essential amino acids (Silva et al., 2015).

There is abundant literature on legume lectins, however, there have been a lack of studies regarding cereal lectins. There was a corn grain study (Molina et al., 2004) where the lectin was characterized, but this is one of only a few available examples using cereal seeds.

The purpose of this article is to describe the purification process and partial characterization of a novel cereal lectin from *Chenopodium quinoa* seeds, in addition to showing its antimicrobial activity.

2 Materials and methods

2.1 Extraction and protein purification

Chenopodium quinoa seeds, donated by Embrapa, were pulverized to obtain a finely powdered material. For the extraction, the protein flour was diluted in 0.1M saline solution with a pH of 7.6 (rate of 1:5 (w/v)) before being placed under stirring for one hour at 25 °C. The supernatant was recovered using centrifugation (3500 rpm for 30 minutes) (Fanem, Excelsa II 206 BL, Brazil), and was then dialyzed against ultrapure water before being lyophilized (Liotop, L101, Brazil).

First, the supernatant (herein referred to as CqCE [*Chenopodium quinoa* crude extract]) was diluted (0.3 mg/3 mL) in 0.2 M Ambic buffer with a pH of 7.6. The volume of 3 mL of this solution was then chromatographed on a Sephadex G-50 equilibrated with 0.2 M Ambic buffer with a pH of 7.6, under a flow of 0.5 mL/min (3 ml/tube), using a Fast Protein Liquid Chromatography system (FPLC) (BioRad BioLogic LP, USA). The fractions collected had an absorbance of 280 nm. After chromatography, two different fractions were recovered. These two fractions were clearly separated by two peaks, called Peak 1 and Peak 2. Both peaks were lyophilized (Liotop, L101, Brazil) and stored in a freezer at -20 °C.

Peak 1, which had shown hemagglutinating activity, was submitted to an ion exchange chromatography. It was dissolved in Tris-HCl 20 mM pH 8.0 buffer and applied (1 mL) to an ion exchange column (MonoQ-GE) previously equilibrated with a Tris-HCl 20 mM pH 8.0 buffer. Fractions of 3.0 mL were collected and monitored at 280 nm in a FPLC system (BioRad BioLogic LP, USA). The binding protein was eluted after the application of Tris-HCl 20 mM + NaCl 500 mM pH 8.0. Both peaks were tested and, the one that had shown HA was then dialyzed against the Tris-HCl buffer 10 mM and then lyophilized (Liotop, L101, Brazil) and stored in a freezer at -20 °C.

The protein content was verified using the Bradford Method (Bradford, 1976) and hemagglutinating activity, according to Sharon & Lis (1972). The results from the hemagglutinating activity were expressed as Hemagglutinating Units (HU), which were calculated as the inverse of the highest dilution, presenting hemagglutination per 50 µL (HU. 50 µL⁻¹).

Determination of hemagglutinating activity

The hemagglutinating activity was tested on intact and trypsinized human erythrocytes of the ABO-human blood system. The collected blood was kept in a saline solution containing EDTA, following Sharon & Lis's (1972) method. The blood was centrifuged at room temperature in a table centrifuge (Eppendorf, Eppendorf centrifuge 5417, USA) at 450 rpm × g for 10 min. Then, the supernatant was rejected and the cell pellet (erythrocytes) was washed 4-5 times with 0.15 M NaCl at a rate of 5 mL of NaCl for each mL of sedimented erythrocytes. For one mL of washed erythrocytes, 25 mL of PBS was added. For each 10 parts of this suspension, a part of the trypsin solution at 1% w/v was added before being incubated at 37 °C for 1 hour. The trypsinized erythrocytes were washed 4-5 times with 0.15 M NaCl to remove the enzyme. The erythrocytes, intact and trypsinized, were finally resuspended in a TBS-Ca²⁺ buffer to reach the final concentration of 2% of the erythrocytes.

To determine the hemagglutinating activity of the fractions extracted from the quinoa, microtiter plates of 96 wells were used. In this step, 50 µL of the sample was added to the wells of the first column, and the following wells were serially diluted. After the dilutions, 50 µL intact or trypsinized erythrocytes solution at 2% were added to all wells. The wells of column 12 containing only the TBS-Ca²⁺ solution and erythrocytes were considered controls. Over the course of two hours, the plates were incubated at 37 °C, and at the end of this period, the minimum concentration of the sample showing hemagglutinating activity was observed.

Affinity for carbohydrate test

The test for inhibition of hemagglutinating activity of the carbohydrates was performed using agar agar, agarose, corn starch, cellulose, fructose, fucose galactose, glucose, glucosamine, gum arabic, lactose, maltose, mannitol, mannose, methyl-α-piranoside, pectin, raffinose, ribose and sucrose. In the first row of wells of the microtiter plates, 50 µL of carbohydrate were added in an initial concentration of 100 mM. The carbohydrates were serially diluted with 50 µL of TBS-Ca²⁺ already present in the plate

wells. Thereupon, 50 μL of sample was added to each well for a final concentration of 9.0 mg/mL. The plates were kept at room temperature for 30 minutes before 50 μL of a 2% suspension of trypsinized type A human erythrocyte was added. After two hours, the lowest concentration of carbohydrate that was able to inhibit agglutination of erythrocytes at 37 °C was determined. Columns 11 and 12 were considered controls, where the negative controls (no HA) consisted of a 50 μL carbohydrate solution and 50 μL of cell suspension, and the positive (HA) consisted of 50 μL of lectin and 50 μL of suspension of erythrocytes.

Chelating agents action on the hemagglutination activity

Next, 50 μL EDTA (25 mM) was added and subsequently serially diluted in TBS- Ca^{2+} 25 μL , which was already present. Then, 25 μL of a solution containing lectin was added, followed by a 50 μL of cell suspension trypsinized human type A erythrocytes 2% in TBS- Ca^{2+} . The control, in column 12, contained 25 μL of the lectin solution, 25 μL of TBS- Ca^{2+} and 50 μL of the cell suspension.

Denaturing the agent's action on the hemagglutination activity: pH

Aliquots of lectin were separated in microtubes, concentrated in a Speed Vac, and resuspended in buffers with pH varying from 2.0 to 10.0. They were then incubated in a water bath at 37 °C for one hour. After this period, they were removed and balanced with NaOH or HCl until the pH stabilized between 7.0 and 8.0. Then, the hemagglutination assay was used to test the samples. The activity was compared to the control that had been prepared with the lectin diluted in TBS- Ca^{2+} .

Denaturing agent's action on the hemagglutination activity: thermal stability

The stability test of the lectin at different temperatures was performed in a water bath at a range of 37 °C to 100 °C for a period of 30 minutes. After stopping the trial with an ice bath, the samples were analyzed for their hemagglutinating activity using human type A erythrocytes. The control was done at a rate that remained at room temperature.

2.2 Characterization of the lectin

Electrophoresis under denaturing conditions

The electrophoresis for the polyacrylamide-sodium dodecyl sulfate (SDS-PAGE) was prepared using 12.5% running gel and a gel separation of 5%. The samples (10-50 mg protein) were dissolved in a sample buffer (1.0 M Tris-HCl pH 6.8, Bromophenol Blue, glycerol, 20% SDS, ultrapure water) with and without DDT. The electrophoretic run was carried out at room temperature in a running buffer of Tris-HCl/Glycine 0.025 M, which lasts approximately three hours in a current of 25mA (Biorad, Mini Protean, USA).

After the run, the gel was removed from the plates and placed in a solution containing coomassie blue R-250 0.25% in

acetic acid, methanol and water (10:40:50) for protein staining. A process of washing in acetic acid, methanol, and water was used to destain the gel and remove excess dye, helping with the visualization of the protein bands (10:40:50). Finally, the molecular weight was determined using molecular mass references.

Native acidic electrophoresis

The native electrophoresis under acidic conditions was performed using 12.5% running gel and stacking gel 5%. The sample was resuspended in the sample buffer ((KOH – 1M; glacial acetic acid; ultrapure water; pH 6.8), glycerol, basic fuchsin, ultrapure water).

The electrophoretic run (Biorad, Mini Protean, USA) was performed at room temperature in 1X running buffer glacial acetic acid/alanine at a pH of 4.3. Initially, a pre-race was performed under a constant current of 1.0 mA, lasting approximately one hour. Subsequently, the electrodes were reversed under constant voltage. The final race was indicated by the arrival of a basic fuchsin marker in the middle of the plate.

After the run, the gel was stained using Coomassie blue R-250 0.25% methanol, acetic acid and ultrapure water (40:10:50). The bleaching gel to remove excess dye and visualization of the protein bands was effected by washing it in methanol, acetic acid and ultrapure water (40:10:50).

2.3 Antimicrobial activity

The antimicrobial activity was conducted using 11 strains of bacteria, seven of which were gram negative (*Escherichia coli* ATCC 11229, *Pseudomonas aeruginosa* ATCC 25619, *Proteus vulgaris* ATCC 13315, *Salmonella enterica* ATCC 10708, *Klebsiella pneumoniae* ATCC 4352, *Serratia marcescens* ATCC 14756 and *Enterobacter cloacae* ATCC 39978) and, four of which were gram positive (*Staphylococcus aureus* ATCC 29213, *Enterococcus faecalis* ATCC 19433, *Micrococcus luteus* ATCC 53598, *Staphylococcus epidermidis* ATCC 12228). The microorganisms were activated in Mueller-Hinton broth after 24 hours of incubation and a groove was made for isolating the colonies. The cell number standardization was done using the tube number 0.5 from the McFarland scale, which corresponds to 1.5×10^8 cells, followed by two serial dilutions (1 mL broth with cells in 9 mL of saline) in order to reach the concentration of 10^6 cells. Then, the microorganisms were inoculated in petri dishes containing Mueller-Hinton agar. Three concentrations of the lectin were tested (100 $\mu\text{g}/\text{mL}$, 250 $\mu\text{g}/\text{mL}$ and 500 $\mu\text{g}/\text{mL}$). The control was the buffer used to dilute the proteins (20 mM Tris-HCl pH8.0). Approximately 10 μL of each concentration of *C. quinoa* lectin was added to the agar's surface. The plates were incubated for 24 hours at 37 °C.

2.4 Statistical analysis

Data were analyzed using a one-way analysis of variance (ANOVA) (General Linear Models). A Tukey test was used to identify the means that differed when the ANOVA indicated statistical significance. For this test, p values <0.05 were considered significant. All the experiments were done triplicated.

3 Results

3.1 Purification of *C. quinoa* lectin

The sample was applied to a column of size exclusion Sephadex G-50, which separated the proteins by different molecular weights. The presence of three protein peaks was observed (Figure 1a). Hemagglutinating tests for both peaks revealed that only the first was able to cause erythrocyte hemagglutination.

The sample containing the first peak from Sephadex G-50 chromatography was applied to an ion exchange column (MonoQ), in which two protein fractions were detected: One that did not bind to the resin and the other that was eluted after a saline solution (Tris-HCl 20 mM + NaCl 500 mM) (Figure 1b). The first peak, which demonstrated hemagglutinating activity, was named CqLec (*Chenopodium quinoa* Lectin). After its purification, the CqLec yield was 9.3% (Table 1).

Hemagglutinating activity determination

The hemagglutinating activity was tested on intact and trypsinized ABO system human erythrocytes, observing, at the end of the incubation period, the minimum concentration of lectin isolated from *C. quinoa* seeds with hemagglutinating activity. The tests showed that the isolated CqLec agglutinates human erythrocytes, having a higher specificity for B Type (Table 2), followed by A and O.

Test for carbohydrate affinity

The specificity of carbohydrate binding for the *C. quinoa* lectin was investigated using the hemagglutinating inhibition method. Though this method is semi-quantitative, it provides information regarding various saccharides capabilities that inhibit lectin activity (Nasi et al., 2009). CqLec demonstrated specificity to glucose and mannose (Table 3).

Chelating agents action on the hemagglutinating activity

The CqLec hemagglutinating activity was shown to be independent of the divalent cations, since EDTA did not inhibit it (not shown).

Denaturing agents action on the hemagglutinating activity

After being diluted and remaining in buffers at a pH ranging from between 2 and 10, aliquots of CqLec underwent a pH adjustment to a range of between 7 and 8, before being submitted to the hemagglutinating assay. The CqLec activity remained stable even under the wide range of pH levels (Figure 2a), suggesting great resistance to extreme pH levels.

The CqLec thermostability was performed by placing the samples at temperatures ranging from between 37 °C and 100 °C

Table 1. Purification table.

Step	Hemagglutinating Activity (HU.100 μL^{-1})*	Total Protein (mg)	Yield (%)
Crude Extract	64	28.8	100.0
Sephadex G50	32	8.99	31.2
MonoQ (GE)	16	0.836	9.3

*HU: Hemagglutinating Unit: the inverse of the hemagglutinating title, calculated as $\frac{1}{\text{number of well presenting the highest dilution with visible hemagglutination}}$.

Table 2. Minimum hemagglutinating concentration of CqLec on human erythrocytes.

Human Erythrocytes (ABO system)	Lower hemagglutinating concentration ($\mu\text{g}/\text{mL}$)
A	7.73
B	3.86
O	7.73
AB	15.46

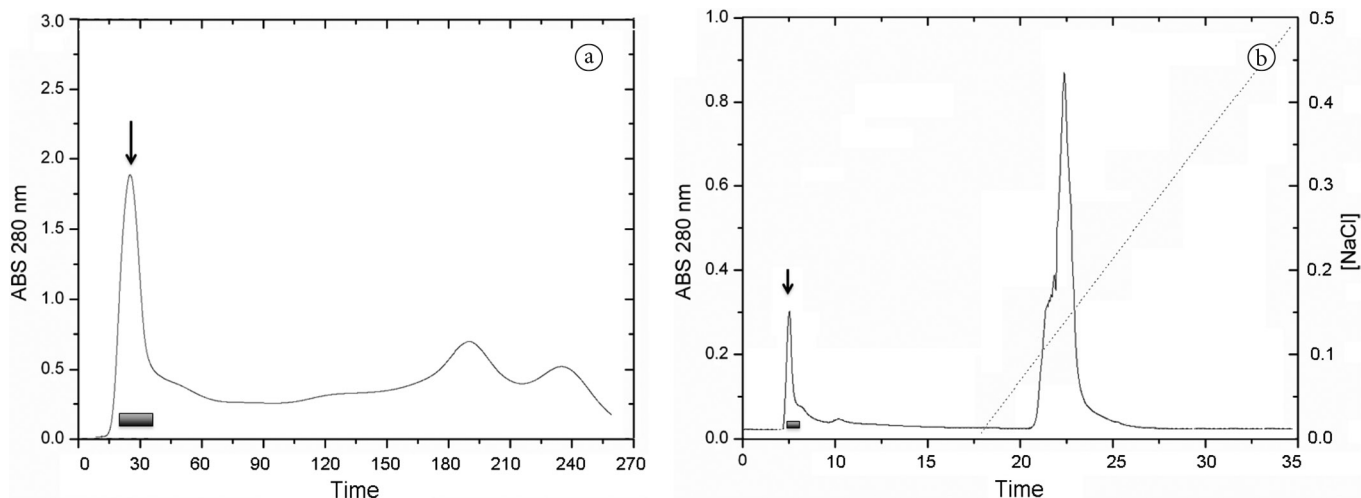


Figure 1. (a) CqCE gel-filtration chromatography column on Sephadex G-50 equilibrated with 0.2 M Ambic buffer at a flow of 0.5 mL/min; (b) Ion exchange chromatography. The peak originating from the Sephadex G-50 column was applied on a MonoQ resin equilibrated with Tris-HCl 20 mM pH 8.0. The first peak was eluted before the saline solution addition. The second peak has been released after the addition of saline solution. The absorbance was realized at 280 nm.

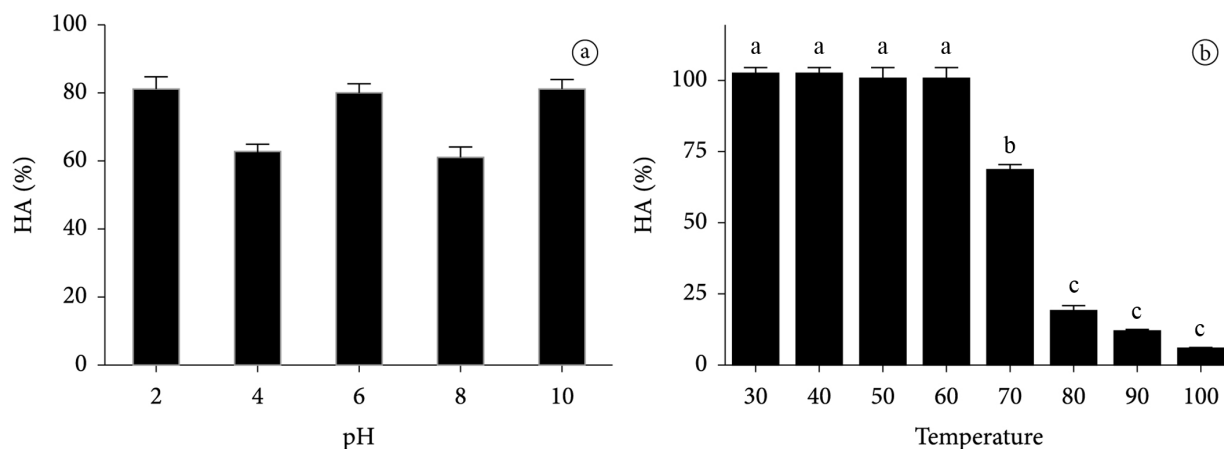


Figure 2. (a) Effect of pH on hemagglutinating activity of CqLec. Between the treatments no significant difference was found (ANOVA, $p < 0.05$); (b) Effect of temperature on hemagglutinating activity of CqLec. Different letters represent significant difference between the treatments (ANOVA, $p < 0.05$).

Table 3. Effects of carbohydrates on the hemagglutinating activity induced CqLec.

Carbohydrates	MIC*
Agar-agar	NI**
Agarose	NI**
Corn starch	NI**
Celullose	NI**
Fructose	NI**
Fucose	NI**
Galactose	NI**
Glucose	3.25 mM
Glucosamine	NI**
Gum arabic	NI**
Lactose	NI**
Maltose	NI**
Mannitol	NI**
Mannose	3.25 mM
Methyl α -piranoside	NI**
Pectin	NI**
Raffinose	NI**
Ribose	NI**
Saccharose	NI**

*MIC: Minimum inhibitory activity. **No Inhibition.

for 30 minutes. After the sample's temperature returned to ambient temperature, hemagglutinating activity was observed in each. CqLec samples exposed to temperatures of between 37 °C and 80 °C demonstrated activity similar to that of the control, while those subjected to 80 °C lost part of this activity. At temperatures of 100 °C, the sample almost completely lost the activity (Figure 2b).

3.2 Structural and biochemical characterization of the lectin

Electrophoresis under denaturing conditions

The SDS-PAGE showed two bands (Figure 3). The molecular weight of one of the bands was estimated to be 35 kDa and the other band was estimated to be approximately 25 kDa.

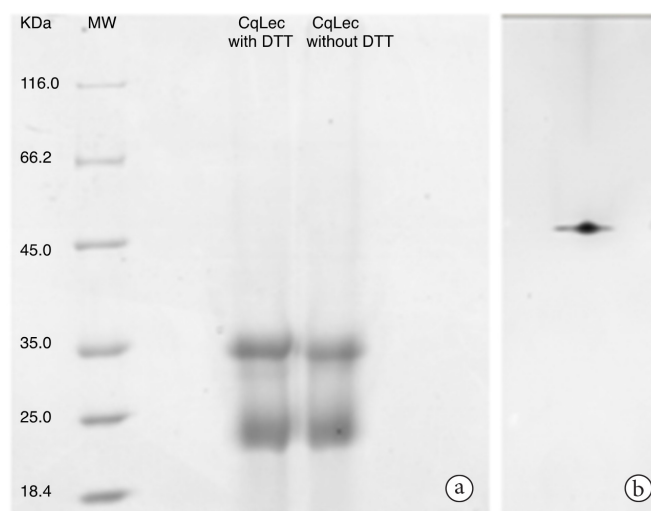


Figure 3. (a) SDS-Page Electrophoresis; (b) Acidic native electrophoresis.

The SDS-PAGES were made using a sample buffer with DTT with and without the presence of this reagent. No differences could be seen between the two samples, indicating that bisulfide bonds do not link the two distinct subunits of CqLec.

Acidic native electrophoresis

The purified CqLec showed a single band on the acidic native electrophoresis. Even though the SDS-PAGE demonstrated two bands, the native electrophoresis showed that the CqLec was completely purified and was apparently composed of two subunits with different molecular weights, making it likely that CqLec is probably a heterodimeric lectin of approximately 60 kDa.

3.3 Antimicrobial activity

The inhibition halos were obtained only on the gram-negative bacteria (Figure 4), specifically, *E. coli* ATCC 11229, using a CqLec concentration of 500 $\mu\text{g}/\text{mL}$ and 250 $\mu\text{g}/\text{mL}$, *Pseudomonas*

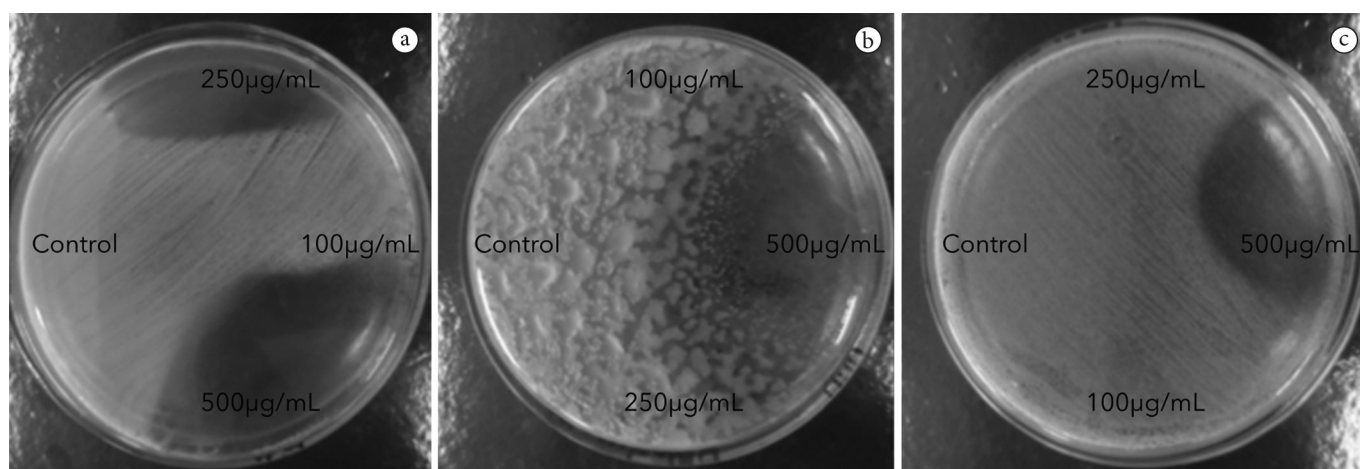


Figure 4. Antimicrobial activity of CqLec. (a) *Escherichia coli*; (b) *Pseudomonas aeruginosa*; (c) *Salmonella enterica*.

aeruginosa ATCC 25619 at a concentration of 500 µg/mL and *Salmonella enterica* ATCC 10708 at a concentration of 500 µg/mL.

4 Discussion

Lectins are defined as proteins/glycoproteins that have at least one domain with no catalytic properties that reversibly bind to some mono or oligosaccharides (Van Damme et al., 2003). The specific amino acid residues are essential for maintaining the carbohydrate binding and hemagglutinating activity of the lectins (Baiimiev et al., 2007). The best-known property of lectin activity is erythrocytes hemagglutination. The lectin detection was carried out mainly through the hemagglutination assay. Thus, most researchers have focused their research on human erythrocytes ABO system antigens.

Some lectins do not agglutinate erythrocytes that are not pretreated with proteases, such as papain and trypsin, which remove obstructions from the membrane surface receptor. This makes them inaccessible to the lectin-binding site. In any case, the interaction of the lectin with the erythrocytes occurs because these receptors are similar to those for which the lectin has specificity. Certainly, the occurrence of lectins points to additional roles and it is appropriate to think of lectins as multifunctional molecules that have more than one function in a particular species or different functions in different species.

The bonding process between lectins and carbohydrates is generally dependent on the presence of metallic ions. The metal's binding site has been described in detail for Con-A, demonstrating that it is very conserved in other plant lectins (Loris et al., 1998). Studies using *Castanea crenata* lectin (Nomura et al., 1998) demonstrated that this protein is independent of the presence of metallic ions, as is CqLec.

Stability, even under highly acidic and basic conditions, is a common property for lectins. The lectin from quinoa seeds showed stability and kept its activity at a high level when submitted to pH levels from 2 to 10. Sharma et al. (2009) obtained similar results when studying lectin, finding considerable stability in the pH range of 1-14.

Thermal stability is also an interesting property for lectins. The results obtained for CqLec are similar to the lectin that Martínez-Cruz et al. (2001) and Wong et al. (2006) purified and characterized. CqLec is fairly thermostable due to the stability of the hemagglutinating activity between 0 °C and 80 °C, although it demonstrated a considerable loss in activity level at temperatures of above 90 °C.

Structurally, CqLec was apparently shown to be a heterodimeric lectin, and bisulfide bonds do not link its subunits once the presence and absence of DTT did not show any differences. Similar results were found for JFL and GCL, in which the SDS PAGE, both in the presence and absence of 6-mercaptoethanol JFL, showed two bands with molecular weight 11 kDa and 17 kDa, and the GCL showed two bands with molecular weight 17 kDa and 26 kDa (Laija et al., 2010). The lectin from *Dioclea lasiophylla* seeds is another example of lectins presenting bands of different molecular weights. DlyL is composed of three different chains with different molecular weights and no bisulfide bonds present (Pinto-Júnior et al., 2013).

Several lectins present antibacterial activity. Some studies demonstrated that components of the bacterial cell wall, such as lipopolysaccharides (LPS), peptidoglycans and teichuronic and teichoic acids, are responsible for the antibacterial effects of lectin (Klafke et al., 2013). In the other hand, *A. jiringa* seed lectin presented higher levels of antimicrobial activity against gram-positive bacteria than against gram-negative bacteria (Charungchittrak et al., 2010), demonstrating that lectin acts against all bacteria groups. These abilities are important tools for biotechnology.

In gram-negative bacteria, the interaction between lectins and LPS is especially interesting, and according to these results, we can infer that bacterial species that have suffered growth interference from CqLec have, in their LPS composition, one or more of the sugars for which the lectin showed specificity. The (sugar) cores of *E. coli* and *S. enterica*, for example, are composed of the common hexoses, glucose, galactose, and *N*-acetylglucosamine (Klafke et al., 2013), two of which are CqLec-specific. The concentration needed for bacterial growth inhibition by CqLec is similar to those presented by *Indigofera*

heterantha lectin (Qadir et al., 2013) and by *Canavalia ensiformis* lectin (Kulkarni & Tayade, 2013).

5 Conclusion

The evaluation of the results demonstrates that the lectin present in *Chenopodium quinoa* seeds is probably a heterodimeric protein with a total molecular mass of approximately 60 kDa (one subunit of approximate 25 kDa and another one of 35 kDa). In addition, bisulfide bonds are not important for the linkage of these subunits. The evaluation can also verify that CqLec agglutinates type B human erythrocytes at a greater intensity compared to the other blood types. CqLec is resistant to different temperature and pH ranges, demonstrating its independence to the presence of divalent cations for the activity. The sugar specificity pattern shows that this lectin belongs to the glucose/mannose-binding group. It demonstrated antimicrobial activity against three gram-negative bacteria (*Pseudomonas aeruginosa*, *Escherichia coli* and *Salmonella enterica*), probably due to their specific bond to sugars that are present at the LPS of those bacteria. The present study demonstrated the potential of *C. quinoa* lectin as an antimicrobial and biotechnological tool.

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