



Full length article

Antimicrobial properties and phenoloxidase activation of the lectin isolated from kadal shrimp (*Metapenaeus doboni*)

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ABSTRACT

The present study reveals purification and characterization of the lectin from the haemolymph of *Metapenaeus doboni*. The *Md*-Lec was purified by affinity chromatography with mannose coupled sepharose CL-4B column and it exhibits single band with a molecular weight of 68 kDa in SDS-PAGE. Furthermore, the molecular mass was confirmed by MALDI-TOF and functional groups present were analysed by FTIR. The surface morphology of purified *Md*-Lec displays the homogeneous nature of protein. The X-ray diffraction (XRD) analysis expresses three peaks at 10.7716°, 21.6258° and 31.7523° which indicate the crystalline nature of the protein and the retention time of 3.068 min evident from HPLC reveals the purity of the sample. Functional analysis of purified *Md*-Lec exhibits yeast agglutination activity against *Saccharomyces cerevisiae* and has the ability to agglutinate the human erythrocytes, which was observed by light microscopy. It also exhibited phenoloxidase activation, encapsulation and phagocytic activities. In addition, purified *Md*-Lec showed the broad spectrum of bacterial agglutination activity against Gram negative *Vibrio parahaemolyticus* and *Aeromonas hydrophila*, important fish pathogens.

1. Introduction

Lectins are glycoproteins which can bind to carbohydrates like mannose, galactose, lactose, N-acetyl glucosamine, N-acetyl galactosamine, fucose, and rhamnose with significant specificity [1]. They possess at least one carbohydrate recognition domain (CRD) that specifically and reversibly binds to different sugar moieties present on the surfaces of pathogens, and lectins are present in almost all organisms including plants, animals, viruses, bacteria, cyanobacteria and yeasts [2–5].

Lectins are involved in many biological functions including cell adhesion, phagocytosis, complement activation and innate immunity [6]. Based on their structure, binding specificities and calcium dependency, lectins are classified into different families like C-type lectins, F-type lectins, galectins, intelectins, rhamnose binding lectins, I-type lectins, Lily-type lectins etc. [7]. C-type lectin (CTL), one of major marine lectins, is characterized by Ca²⁺-dependent binding to mono and oligosaccharides, and they are classified in several groups; collectins, proteoglycan core proteins, selectins directly or indirectly involved in immune function [8]. Role of different types of lectins, their therapeutic applications and the tissue distribution of lectins in the fishes is de-

scribed by our group [9].

In invertebrates like shrimp, based on structural differences, expression patterns and functions, there are seven groups of lectins: C-type, L-type, P-type, M-type, fibrinogen-like domain lectins, galectins, and calnexin/calreticulin [10]. The shrimp lectins take part in innate immune mechanisms by activating prophenoloxidase, encapsulation, melanization and promotion of phagocytosis [11–15].

Metapenaeus doboni is distributed along the Indian coast, off Sri Lanka, Malaysia, Indonesia to Philippines and New Guinea. In India it is common along the South west coast- Kerala, Goa, Karnataka and the south east coast-Orissa, Visakhapatnam, Tamil Nadu and Union territory of Puducherry [16]. Presently little information is available about the immune system and immune response in kadal shrimp. Therefore, the present study addressed to characterize the immune response in kadal shrimp, with focus on isolation and characterization of a lectin from haemolymph and substantiate its functional roles in agglutination, antimicrobial and anticancer properties.

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2. Materials and methods

2.1. Collection and preparation of samples

Live and healthy kadal shrimps (*Metapenaeus dobsoni*) were obtained from a hatchery at Varapuzha, Kerala, India. Haemolymph was collected and extracted in aseptic conditions using gauge 23 hypodermic needles and diluted with 1:1 anticoagulant solution (0.45 M NaCl, 0.1 M glucose, 30 mM sodium citrate, 26 mM citric acid, 10 mM EDTA, pH 7.5). A total of 30 ml haemolymph was pooled for purification of lectin. Then it was centrifuged at 1000 rpm for 10 min at 4 °C and the supernatant was collected into fresh tubes and stored at –20 °C until use.

2.2. Purification of lectin from kadal shrimp haemolymph

Lectin was purified according to the method described by Jayanthi et al. [17] with slight modifications. Briefly, mannose coupled sepharose CL-4B column was used for the purification of *Md-Lec*. The column was extensively washed with TBS/CaCl₂ buffer (10 mM Tris–HCl, 150 mM NaCl, 10 mM CaCl₂, pH 7.4). The haemolymph was equilibrated with an equal volume of TBS/CaCl₂ buffer. Followed by this equilibration, the equilibrated sample was gently applied into a mannose coupled sepharose CL-4B column. Purified fractions were eluted with elution buffer (10 mM TrisHCl, 140 mM NaCl, 3 mM EDTA, pH 8.0) containing EDTA.

2.3. Characterization of purified *Md-Lec*

2.3.1. Molecular weight determination

Polyacrylamide gel electrophoresis (12%) was performed on eluted fractions under reduced conditions as described by Laemmli [18]. After electrophoresis, the gel was stained with Coomassie brilliant blue (GE Healthcare Bio-Sciences, India) and image captured through ChemiDoc XRS + system (Bio-rad, USA). The molecular mass of the purified *Md-Lec* was determined by comparison of its electrophoretic mobility with those of molecular mass marker proteins (Takara BIO INC, Japan). Total protein concentration was determined by Lowry's method [19] using bovine serum albumin (BSA) as standard.

2.3.2. High performance liquid chromatography and X-ray diffraction analysis

HPLC separation was carried out using a reversed phase C18 column (7.8 mm × 30 cm) previously equilibrated with TBS-I at a flow rate of 0.8 ml min⁻¹. HPLC system (Zorbax Bio-series GF-250, Du Pont, Willington, DE, USA), was used for the homogeneity analysis. In order to determine the spatial distribution of atomic coordinates and the arrangement of atoms, an XRD analysis (XRD, Scintag-SDS 2000) was performed on purified *Md-Lec* at 40 kV/20 mA, using continuous scanning 2θ mode. The average grain size and shape of the purified *Md-Lec* was determined using Scherrer's formula [$d = (0.9\lambda/\beta\cos\theta)$], where d is the mean diameter of purified *Md-Lec*, λ is the wavelength of the X-ray radiation source, and β is the angular FWHM of the XRD peak at the diffraction angle (θ).

2.3.3. Fourier transform infrared spectroscopy (FTIR) and MALDI-TOF analysis

For FTIR spectra studies, purified *Md-Lec* (1–50 μl) was placed in a thermostated cell fitted with CaF₂ windows (with 6 μm Teflon spacer for measurements in water). The spectra of biological molecules were recorded at a resolution of 4 cm⁻¹.

For MALDI-TOF analysis, the purified 68 kDa protein was excised from SDS-PAGE gel and subjected to trypsin digestion. In brief, the gel

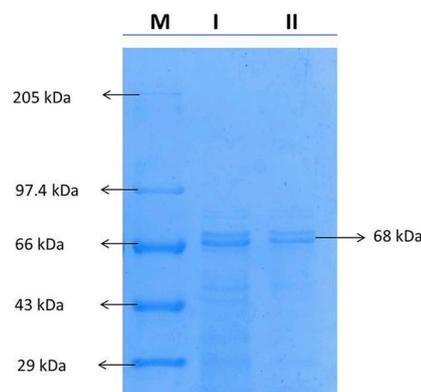


Fig. 1. Polyacrylamide gel electrophoresis (12%) in the presence of sodium dodecyl sulfate (SDS) of *Md-Lec* purified by Mannose-Sepharose CL 4B affinity chromatography. Lane M contains High molecular weight marker. Lane I contains reducing pattern and Lane II contains non-reducing pattern of purified *Md-Lec* from *Metapenaeus dobsoni* haemolymph. The protein bands were stained with Coomassie brilliant blue – R 250. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

bands corresponding to *Md-Lec* were finely cut and added to 1.5 ml eppendorf tubes containing stain removal solution (100 mM ammonium bicarbonate: 50% acetonitrile (1:1)) vortexed for 30 min. After complete removal of stain, the gel pieces were dehydrated with 100% acetonitrile (ACN). Then, the gelpieces were subjected to reduction and alkylation. Trypsin solution was formulated in 40 mM NH₄HCO₃ and added to each eppendorf tubes containing gel pieces and stored at ice cold condition for absorption of trypsin. After absorption, NH₄HCO₃ was added and incubated at 37 °C for 12–16 h. The digestion was stopped by adding 4% formic acid. The peptides were extracted by washing gel pieces 3 times with 4% FA in 50% ACN. The resultant peptides were analysed by MALDI-TOF/TOF MS (Shimadzu, Biotek Axima Performance) [30].

2.4. Functional analysis

2.4.1. Haemagglutination assay

The haemagglutination assay (HA) was performed in microtiter plates according to Correia and Coelho [20]. Lectin preparations (50 μl) were serially two fold diluted in PBS before addition of 50 μl 2% (v/v) suspension of human erythrocytes. In control, purified *Md-Lec* was replaced by BSA. The titer was expressed as the highest dilution exhibiting haemagglutination. The HA was defined as the inverse of the last dilution at which the sample still showed haemagglutination. Specific HA (SHA) was defined as the ratio between the titer and protein concentration (milligrams per milliliter). These haemagglutinated titres of *Md-Lec* were visualized by light microscopy at the magnification of 40X (Leica DMIL).

2.5. Phenoloxidase enhancing activity

The ability of *Md-Lec* in activating prophenoloxidase was studied by measuring the formation of dopachromes from L-DOPA. As per the method described by Iswarya et al. [21], 50 μg/ml of purified *Md-Lec* was mixed with equal volume of laminarin (1 mg ml⁻¹) and incubated at 25 °C for 15 min. Then 50 μl haemocyte lysate supernatant was introduced and re-incubated for 45 min in the presence of 5 mM CaCl₂. Subsequently, 50 μl L-DOPA (3 mg ml⁻¹) as enzyme substrate was added and incubated at 25 °C for different time intervals (30, 60, 90, 120 min). In control *Md-Lec* is replaced by TBS I buffer. After

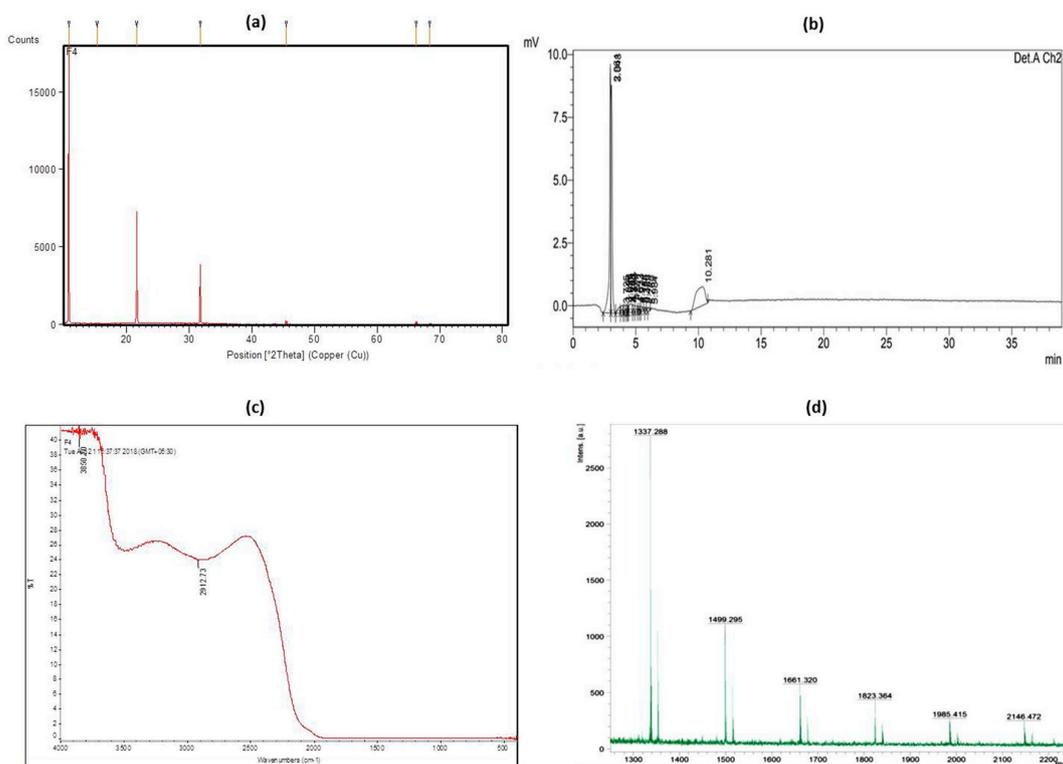


Fig. 2. Biochemical characterization of *Md-Lec*. (a) XRD analysis of purified *Md-Lec* showed diffraction peaks at 10.7716° , 21.6258° and 31.7523° which indicate the crystalline nature of the protein (b) The HPLC analysis of purified *Md-Lec* showing a single peak with a retention time at 3.068 min which indicates the purity of the sample (c) FTIR analysis for functional group identification. The attributions of the main absorption characteristics of glycosidic structures are related to O–H stretching ($3000\text{--}3500\text{ cm}^{-1}$) (d) MALDI–TOF/TOF MS peptide analysis of 68 kDa *Md-Lec*.

incubation, the formation of dopachrome was spectrophotometrically measured in the wavelength of 490 nm and expressed as unit/min/mg/protein.

2.5.1. Encapsulation assay

To evaluate the encapsulation activity of *Md-Lec* was carried out by the method reported earlier [21], with slight modifications. The haemocyte suspension was mixed with sepharose beads suspension and *Md-Lec* (25, 50 and $100\text{ }\mu\text{g/ml}$) in V- bottomed microtitre plate and allowed to incubate at 25°C for 45 min with intermediate mixing at every 15 min. In control wells, *Md-Lec* was replaced by TBS I buffer. The entire volume from each suspension was spread on sterilized glass slide and kept undisturbed for 10 min. Then the slides were observed under light microscope (40X) (Leica DMIL).

2.5.2. Yeast agglutination assay

To determine the ability of agglutination of *Md-Lec* to yeast cells (*Saccharomyces cerevisiae*), $50\text{ }\mu\text{l}$ of different concentrations of the purified *Md-Lec* was added to a U-shaped 96-well microtitre plate containing equal volume of Tris buffer. The same volume of a suspension of yeast ($10^6\text{ cells ml}^{-1}$) was then added to the wells and incubated for 4 h at 25°C . In control, purified *Md-Lec* was replaced by BSA. The pattern of agglutination was monitored with an inverted light microscope (40X) (Leica DMIL).

2.5.3. Bacterial agglutination assay

The bacterial agglutination by purified *Md-Lec* against *V. parahaemolyticus* and *A. hydrophila* was evaluated as described by Rubeena et al. [22]. Different concentrations of the purified *Md-Lec* were added to a 96-well microtitre plate and an equal volume of Tris buffer was

added. In control, purified *Md-Lec* was replaced by BSA. The same volume of a suspension of bacterial cells ($10^6\text{ cells ml}^{-1}$) was then added to the wells and incubated for 30 min and then the pattern of agglutination was monitored with an inverted light microscope (10X) (Leica DMIL).

2.5.4. Antibiofilm assay

The effect of purified *Es-Lec* on biofilm-forming Gram-negative bacteria (*Aeromonas hydrophila* and *Vibrio parahaemolyticus*) was tested using 24-well polystyrene plates containing glass pieces immersed in Luria Bertani broth inoculated with the bacterial suspension of 1% inoculum from overnight cultures (10^7 CFU ml^{-1}). Different concentrations of *Md-Lec* were introduced into the wells and incubated at 37°C for 48 h. For examining the biofilm, the media were discarded and weakly adherent cells were removed by thorough washing with deionized water and allowed to air dry before staining. The biofilm were stained with 1 ml of 0.4% acridine orange (w/v) for 10 min. The biofilm inhibition in 3D view was observed by confocal laser scanning microscopy (Carl Zeiss LSM 710 Germany). In control, *Md-Lec* was replaced by BSA.

2.6. Statistical analysis

All experiments were performed in triplicates ($n = 3$). Data are presented as mean \pm standard error mean (S.E.M.) of control and treated samples. The data were subjected to one way analysis of variance (ANOVA) and the significance of differences between means were calculated by Tukey's HSD test and the significance accepted at $P < 0.05$.

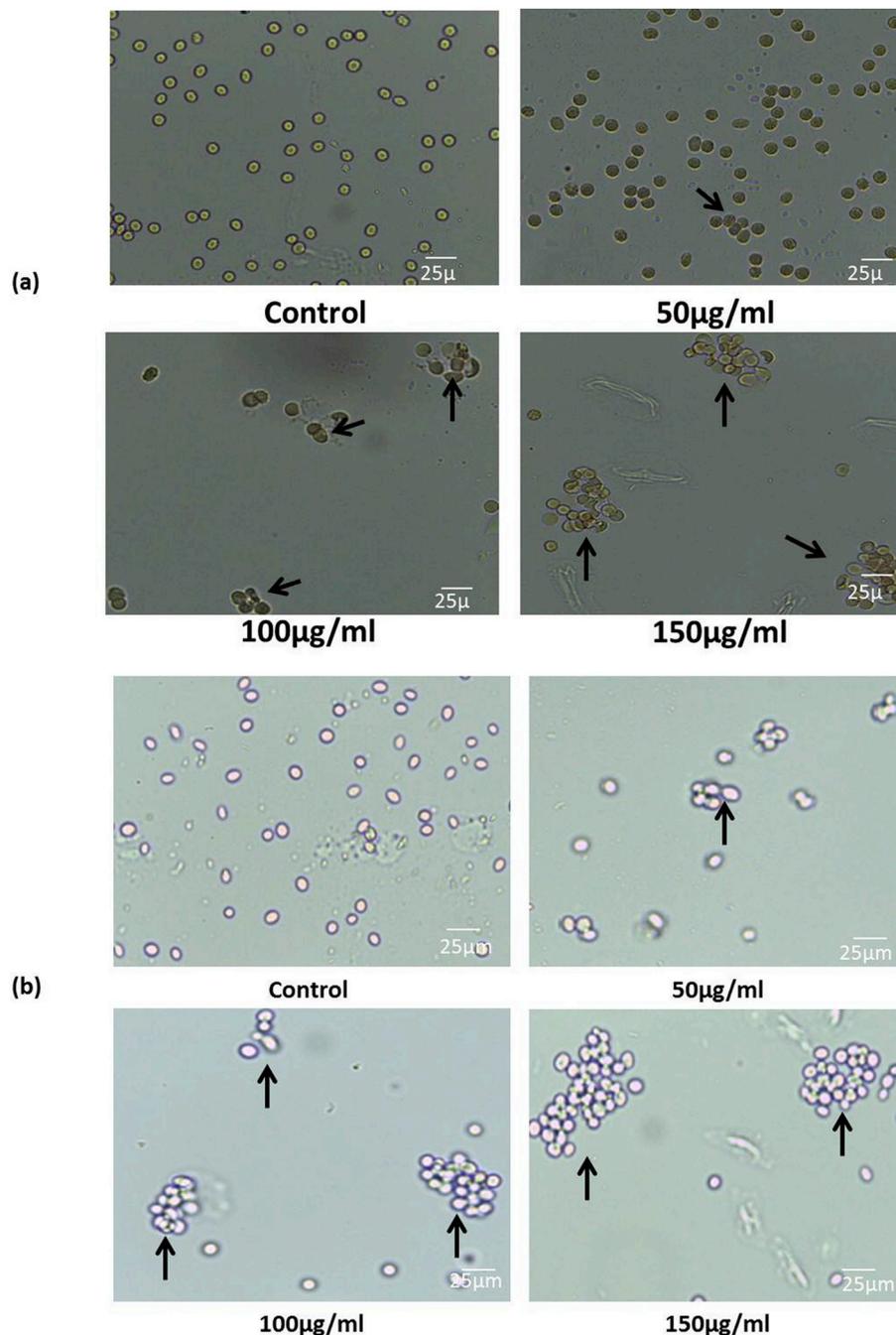


Fig. 3. (a) Haemagglutination assay with against 2% human RBC suspension and (b) yeast agglutination assay of *Md-Lec* at different concentrations. Highest concentration (150 µg/ml) showed maximum agglutination. Black arrows indicate the agglutination.

3. Results

3.1. Purification of lectin from kadal shrimp

Lectin from haemolymph of kadal shrimp was purified by affinity chromatography and the SDS-PAGE analysis revealed a doublet band of approximately 68 kDa in 12% SDS-PAGE under reducing conditions (Fig. 1).

3.2. HPLC and X-ray diffraction analysis

The XRD analysis of purified *Md-Lec* showed one major diffraction peak at 10.7716° and other peaks at 21.6258° and 31.7523° (Fig. 2a) which shows the lattice arrangement and crystalline nature of protein.

The HPLC analysis revealed a single peak of retention time of 3.068 min (Fig. 2b).

3.3. FTIR and MALDI-TOF analysis

FTIR analysis was done for functional group identification. The attributions of the main absorption characteristics of glycosidic structures are related to O–H stretching ($3000\text{--}3500\text{ cm}^{-1}$) (Fig. 2c). The intensity of the peaks obtained in MALDI-TOF analysis refers to the 68 kDa protein (Fig. 2d) and the mass measurement was done following Henzel et al. [36].

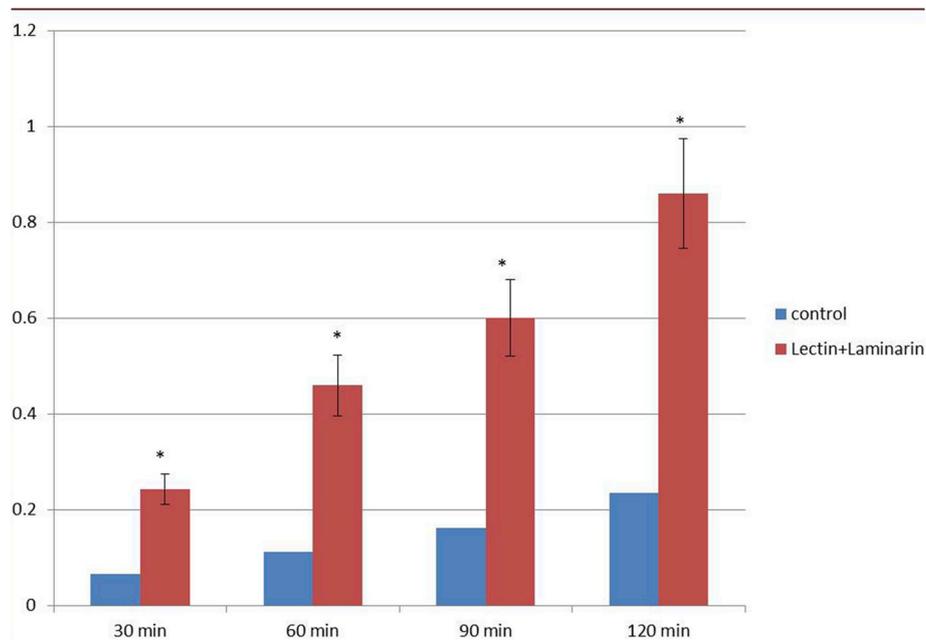


Fig. 4. Specific PO activity of *Md-Lec* at different time intervals of incubation. PO activity increased with time of incubation.

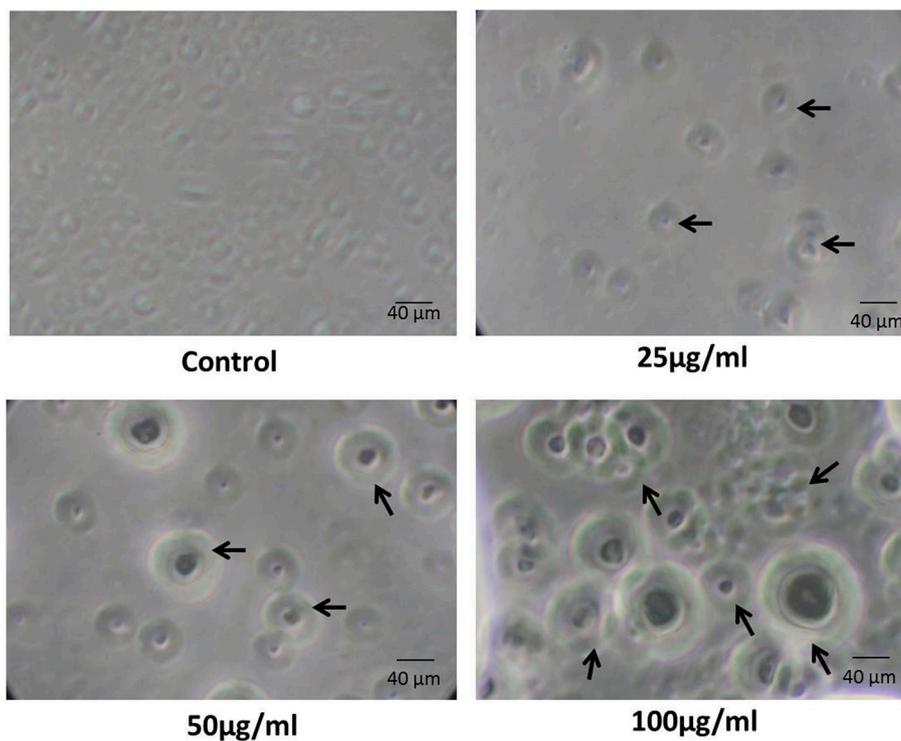


Fig. 5. Light microscopic images showed encapsulation of Sepharose CL4B beads with *Md-Lec*. The arrow indicates the encapsulation of beads.

3.4. Haemagglutination assay

Purified *Md-Lec* had the ability to agglutinate human erythrocytes, and agglutination was highest in 150 µg/ml of purified *Md-Lec* vs. the lower concentration (50 µg/ml). No agglutination occurred when purified *Md-Lec* was replaced by BSA (Fig. 3a).

3.5. Phenoloxidase enhancing activity

PO assay indicates that purified *Md-Lec* could enhance phenoloxidase activity together with laminarin and it strongly depends on time

as it was proved that longer the incubation time higher the PO activity (Fig. 4).

3.6. Encapsulation assay

Encapsulation ability of the haemocytes from *M. dobsoni* was evaluated using sepharose beads *in vitro*. The haemocytes extensively encapsulated sepharose beads up on the addition of *Md-Lec*. High encapsulation activity was observed with 100 µg/ml of purified *Md-Lec*. In control, encapsulation activity was not observed due to the absence of lectin (Fig. 5).

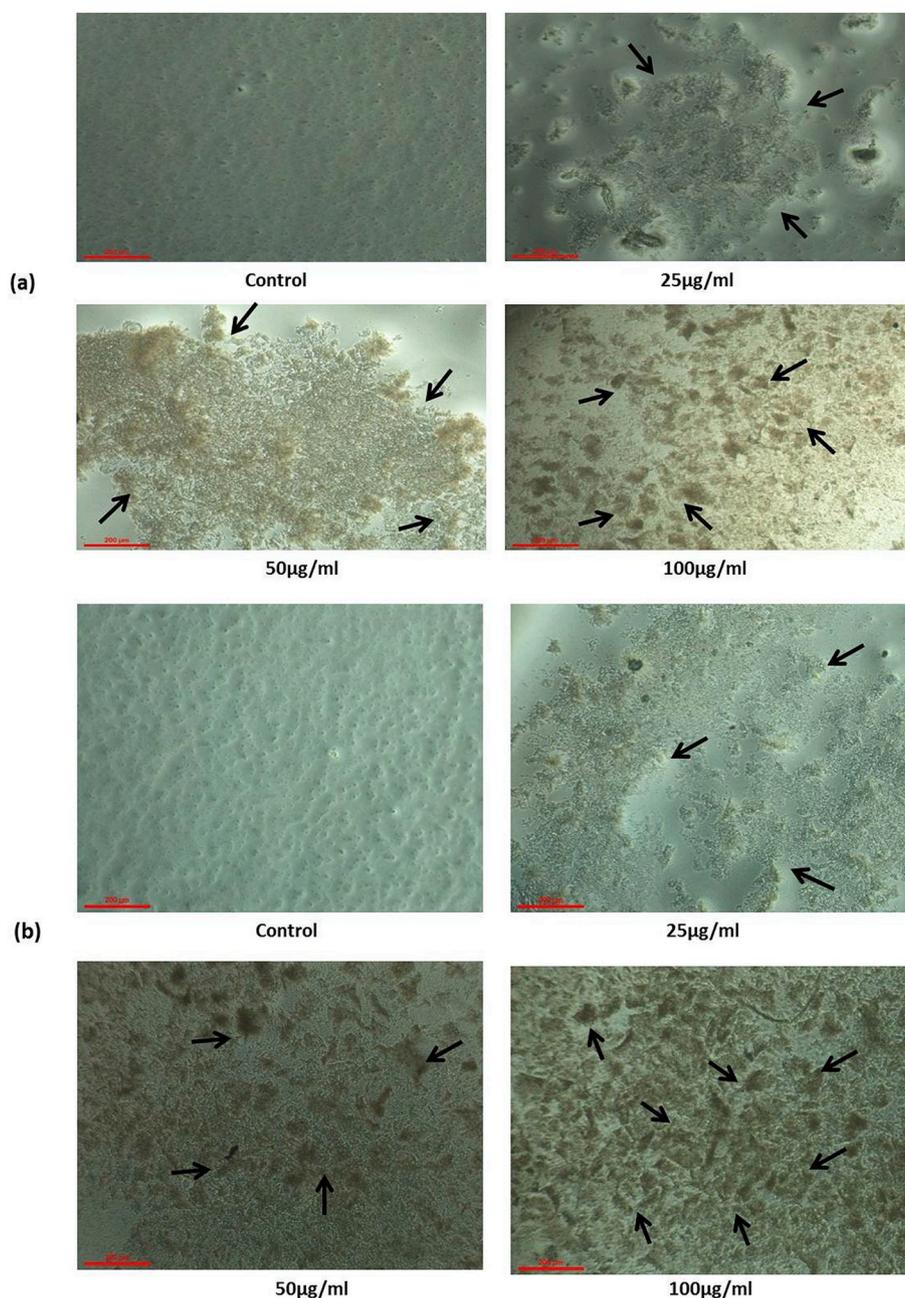


Fig. 6. Bacterial agglutination effect of *Md-Lec* against clinically important (a) Gram negative *A. hydrophila* and (b) Gram negative *V. parahaemolyticus* demonstrated by light microscopy. Higher concentration of lectin showed maximum agglutination activity. Black arrows indicate the agglutination.

3.7. Yeast agglutination assay

The purified *Md-Lec* showed ability to agglutinate *S. cerevisiae* and highest agglutination was revealed in 150 µg/ml of purified *Md-Lec* compared to the lowest concentration (50 µg/ml). No agglutination occurred when purified *Md-Lec* was replaced by BSA (Fig. 3b).

3.8. Bacterial agglutination assay

The purified *Md-Lec* could bind and agglutinate *A. hydrophila*, and *V. parahaemolyticus* and agglutination was evident as clumps of bacteria as revealed in 100 µg/ml of purified *Md-Lec* vs. the lowest concentration (25 µg/ml). No agglutination occurred when purified *Md-Lec* was replaced by BSA (Fig. 6a and b).

3.9. Antibiofilm assay

The biofilm architecture was interrupted the increased concentrations of *Md-Lec* which was evident by decreased bacterial count of *A. hydrophila* and *V. parahaemolyticus* observed by confocal laser scanning microscopy (Carl Zeiss LSM 710 Germany) (Fig. 7).

4. Discussion

Till date several lectins are purified from crustaceans with a higher range of molecular weights of 452 kDa from *Penaeus japonicus* (Kuruma shrimp) [23] and 220 kDa *Litopenaeus schmitti* (White shrimp) [24] to a lower range of molecular weights of 14 kDa from *Tachypleus tridentatus* (Tri-spine horse-shoe crab) [25] and 20 kDa from *Scylla serrata* (Mud crab) [26]. In the present study, we isolated and purified a mannose binding lectin of 68 kDa from the haemolymph of Kadal shrimp which

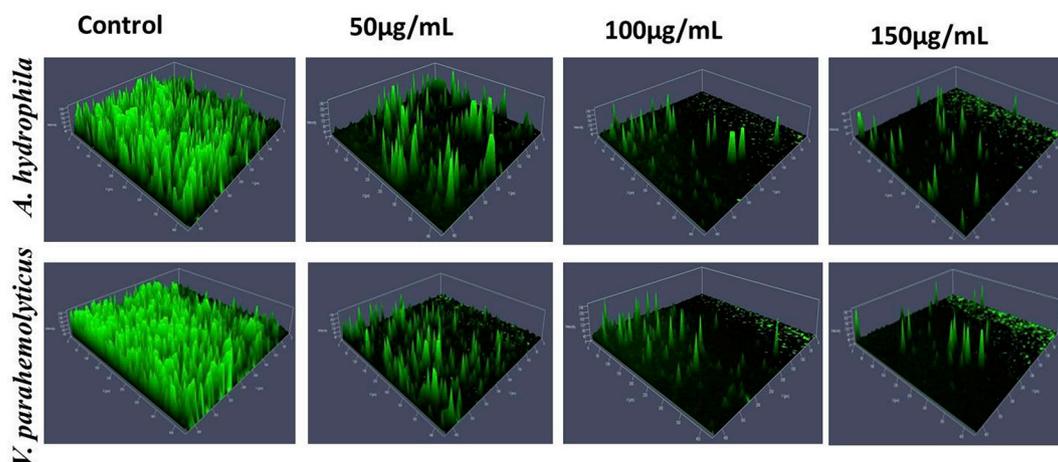


Fig. 7. Antibiofilm effect of *Md-Lec* on biofilm inhibition against clinically important Gram negative *A. hydrophila* and Gram negative *V. parahaemolyticus* with the stain of acridine orange demonstrated by confocal microscopy. Higher concentration of lectin showed maximum activity against biofilm formation. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

was confirmed by SDS-PAGE. The doublet band indicates the post translational modifications (phosphorylation) [35]. Previously, a lectin with a molecular weight of 66 kDa was reported in *Ictalurus punctatus* (Ictalurid catfish) [27]. The mannose coupled sepharose CL-4B affinity column was used for the purification of lectin from Kadal shrimp haemolymph. The calcium ions in the buffer facilitated the binding of lectin on to the mannose ligands in the column, while during elution of lectin from the column, the EDTA present in the elution buffer chelated the calcium ions and hence the lectin was released. Furthermore, the purity and crystalline nature was confirmed by HPLC and XRD analysis respectively, as reported by Jayanthi et al. [28] previously. The functional group analysis and confirmation of molecular mass was done by FTIR analysis and MALDI-TOF analysis which goes in parallel with those reported by Divya et al. [29].

The functional properties of *Md-Lec* were evaluated by haemagglutination and yeast agglutination assay in which agglutination of RBC and yeast cells occurred in a dose-dependent manner. This suggests that when more lectin was available the more they could bind to the sugar moieties present on the cell surfaces. The haemagglutination activity reported by Silva et al. [30] agrees with the findings of the present study. In addition to the role of *Md-Lec* in agglutination, it also plays vital role in encapsulation and phagocytosis. *Md-Lec* was able to facilitate encapsulation against the sepharose beads and this property is previously reported in some other invertebrates also [28,31,32]. Prophenoloxidase (ProPO) mechanism is actively involved in the extermination of pathogens. Sivakamavalli et al. [31] reported that the complex formed between PAMPs and lectin could activate the ProPO (inactive form) to PO (active form) as a result of the degranulation of haemocytes [33]. In the present study, the PO system was triggered by *Md-Lec* along with laminarin and the PO activity was more efficient over time.

Furthermore, the antimicrobial property of *Md-Lec* was studied. *Md-Lec* could agglutinate the Gram negative aquatic pathogens tested in a dose-dependent manner and inhibited the biofilm formation which in turn prevented the pathogens from colonizing and proliferating on host surface as reported by Jayanthi et al. [28]. The intensity of biofilm disruption was studied by confocal microscopy [29]. To conclude, in this present study, we report the purification of immune related lectin *Md-Lec* with a molecular mass of 68 kDa from the haemolymph of *Metapenaeus dobsoni* by affinity column chromatography technique which was confirmed by SDS-PAGE and strongly establish the agglutination, encapsulation, PO mechanism and antibiofilm properties which were not extensively studied as evident by the available literature. Furthermore, the biochemical properties of *Md-Lec* were studied by MALDI-TOF, HPLC, XRD and FTIR. A significant antibacterial and

antibiofilm activity against the most important aquatic pathogens-Gram-negative bacteria *V. parahaemolyticus* and *A. hydrophila* was exhibited by *Md-Lec* indicating its therapeutic potential which demands further molecular studies to develop this molecule into a reliable biological tool in the field of aquaculture.

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