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Antimicrobial activities of lactic acid bacteria isolated from Malaysian prawn, *Macrobrachium rosenbergii*



Authors Info

N.Z. Azahar^{1*}, S. Iehata¹, F. Fadhil¹, M. Bulbul² and Md. A. Kader^{1,2}

¹School of Fisheries and Aquaculture Sciences, Universiti Malaysia Terengganu, 21030 Kuala Terengganu, Terengganu, Malaysia

²Institute of Tropical Aquaculture, Universiti Malaysia Terengganu, 21030 Kuala Terengganu, Terengganu, Malaysia

*Corresponding Author Email : syafiqahzahidahazahar@gmail.com

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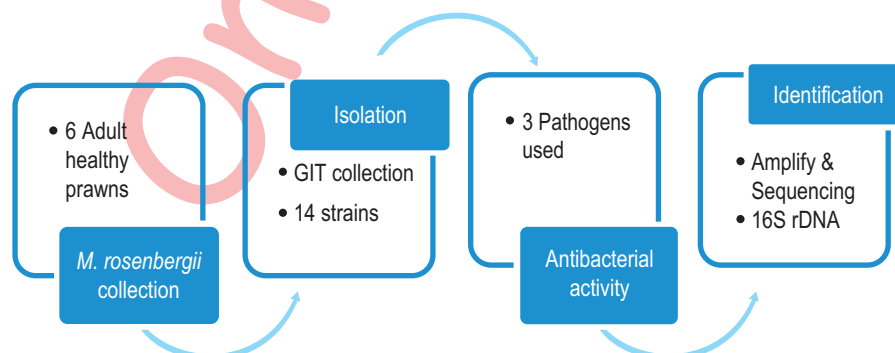
Abstract

Aim: This study aimed to identify lactic acid bacteria (LAB) isolated from *Macrobrachium rosenbergii* with an emphasis on their antimicrobial activities against pathogens.

Methodology: Six *M. rosenbergii* broodstocks were collected from local fishermen and brought to the laboratory where all of them were sacrificed and their digestive tracts were collected. Fourteen isolates grew in MRS agar medium and subjected to pH between 6.3 - 6.6 at 37°C. The strains were tested for their ability to inhibit the growth of pathogens; *Vibrio parahaemolyticus*, *V. alginolyticus* and *Aeromonas hydrophila*. Molecular identification of the strains were performed by amplifying and sequencing the 16S rDNA by which the results were then compared to the database of known 16S rDNA sequences.

Results: The results showed that 57% of the strains inhibited growth against both *V. alginolyticus* and *V. parahaemolyticus*, whereas only 21% of the strains inhibited growth against all the pathogens tested. The identification result showed that 6 strains had high similarity with *Enterococcus faecalis*, whereas 4 isolates had high similarity with *Lactococcus garviae* and 4 isolates were affiliated with *Lactococcus lactis*.

Interpretation: All strains were identified as lactic acid bacteria, and strains with the best ability to inhibit pathogens will be selected for further studies on their potential as dietary additive.



Introduction

Crustaceans such as prawns, shrimps, lobsters and crabs play an important role in aquaculture and are good protein sources, essential amino acids and polyunsaturated fatty acids (Muralisankar *et al.*, 2014). While the giant freshwater prawn, *Macrobrachium rosenbergii* is a commercially important species that is currently being farmed throughout the world, but this industry has experienced little growth over the past decade (FAO, 2016). Part of this limited growth include diseases such as *Macrobrachium* Muscle Virus (MMV), White Spot Syndrome Baculovirus (WSBV) and Idiopathic Muscle Necrosis (IMN) (FAO, 2004-2016).

In the past, farmers had to use large quantities of drugs such as antibiotics in order to improve the prawn survival rates and production, which has led to the emergence of antibiotic resistant bacteria and imbalance of microbiota in the gastrointestinal tract of aquatic species (Nakano, 2007). Therefore, probiotics have been used as an alternative to antibiotics for inhibiting pathogens and diseases in aquaculture species (Cruz *et al.*, 2012). Probiotics comes from the Greek words; 'Pro' and 'bios' which means "for life" (Schrezenmeir and De Vrese, 2001). Probiotics are defined as live microorganisms and can promote beneficial health to the host when they are administered in an appropriate amount (FAO/WHO, 2001). Their mechanisms of actions include competitive exclusion against pathogens, host digestion improvement through production of enzymes, supplying essential nutrients, enhancement of immune response, as well as improving water quality (Balcazar *et al.*, 2006).

There are many criteria to be met while selecting probiotic microorganisms that could confer beneficial health effects to the host. firstly, they should cause no harm to the host, should be accepted by the host via ingestion, should have ability to proliferate and colonize within the host, and should contain no virulent resistance or antibacterial resistance genes (Verschuere *et al.*, 2000; Kesarcodi-Watson *et al.*, 2008). The potential probiotics must be tested with inhibitory activity against pathogens, first in order to access whether they can provide protection to the hosts when exposed to the pathogens (Vijayan *et al.*, 2006; Hai *et al.*, 2007; Irianto and Austin, 2002; Vaseeharan *et al.*, 2004). Therefore, potential probiotics that possess these characteristics are most suitable to be applied to the cultured species in aquaculture.

The objective of the study was to isolate lactic acid bacteria (LAB) strains from gastrointestinal tracts of healthy adult prawn, *M. rosenbergii*, based on their antimicrobial activities against three pathogens.

Materials and Methods

Source of prawns and bacteria: Adult freshwater prawn *M.*

rosenbergii, with an average weight of 10 g were obtained from Sungai Manir, Kuala Terengganu, Terengganu, Malaysia. MRS agar, Marine agar, Tryptic Soy Agar (TSA), MRS broth and Marine broth were prepared according to manufacturer's instruction (Merck, USA). Both the species of *Vibrio* (*V. alginolyticus* and *V. parahaemolyticus*) were cultivated and maintained in Marine broth. *Aeromonas hydrophilla* was cultured and maintained in TSA. The incubation temperature was maintained at 37°C.

Isolation of putative probiotics and determination of antibacterial activity against pathogens :

Prawns were washed with 70% ethanol before dissecting out the intestines using sterile tweezers and scissors. The intestines with digestive contents were weighed and suspended in phosphate buffered saline (0.01 M phosphate buffer, 0.0027 M potassium chloride, and 0.137 M sodium chloride, pH 7.4) to prepare the homogenate by using homogenizer. The suspensions were serially diluted between 10^{-2} to 10^{-5} for each dilution, and 100 μ l of the suspension was sampled and spread onto the surface of MRS agar with 1% CaCO₃. The plates were then incubated under aerobic conditions at 37°C for 24 - 48 hrs. A total of 14 isolates were obtained and all of them were tested for antibacterial activity against the following pathogens: *A. hydrophilla*, *V. parahaemolyticus* and *V. alginolyticus* by well diffusion assay. Briefly, after 24 hrs of culturing *V. parahaemolyticus* and *V. alginolyticus* in a marine broth, a sterile cotton swab was dipped into the broth and spread onto marine agar, which was allowed to dry for 20 min. Wells of 5.0 mm in diameter were punched into the agar with a sterile pipette tip. Each agar plate was punched with 4 wells. All bacterial isolates were cultured in a MRS broth and incubated at 37°C for 24 hrs, and then 5ml of broth culture was sampled and centrifuged for 5 min at 10,000 rpm. A 60 μ l of supernatant was taken from the tube and added into the well of agar plates loaded with corresponding pathogens. All the plates were incubated for 24 hrs at 37°C. The plate was observed for any zones of inhibition (ZOI) around the well. The results were considered positive if the diameter (mm) of the ZOI was greater than 1 mm.

Identification of probiotic strains : Probiotic strains with positive antagonistic activity against the pathogens were cultured in a MRS broth for 24 hrs and centrifuged at 15,000 x g for 2 min to obtain a pellet. The pellet obtained was used for DNA extraction using a DNA extraction kit (Promega, Madison, WI, U.S.A.) following the manufacturer's protocol. Following DNA extraction, bacterial 16S rDNA gene was amplified using 8F (5'AGAGTTTGATCCTGGCTCAG 3') and 1492R (5'CGGGAACGATTCACCG 3') primers for amplification using Polymerase Chain Reaction (PCR). Product quality was verified through electrophoresis and analyzed using 1.5% agarose gel stain with sybr safe, then sequencing was sent to 1st Base Company. The obtained 16S rDNA gene sequences were added to publically available bacterial 16S rDNA sequences and were integrated to the database with the automatic alignment tool in Chromas software. A phylogenetic tree was generated by

performing a distance matrix analysis using neighbor joining method in Mega 6 Software (Tamura *et al.*, 2013). A database search and comparisons were done with BLAST search.

Results and Discussion

The use of strong antimicrobial chemicals, including antibiotics, is becoming increasingly regulated in disease treatments and thus, more environmentally friendly options are required. Alternatives to such probiotics may be a better choice over antibiotics (Cruz *et al.*, 2012). The demand for the probiotics is growing larger due to the increasing demand for hygienic and environmentally friendly products (Wang *et al.*, 2008). In this study, 14 probiotics strains from the gastrointestinal tract of healthy freshwater prawns, *M. rosenbergii*, were identified.

Table 1 shows the antimicrobial activities against three pathogens. As shown in Table 1, strain no. 2, 9 and 12 had inhibitory effects against all the three pathogens tested. On the other hand, 8 strains (no. 3, 13, 16, 20, 27, 30, 31 and 33) showed inhibitory effects against both *Vibrio* spp. Strain no. 18 and 39 showed inhibitory effects on *V. alginolyticus* only, whereas strain no. 17 had inhibitory effects against *V. parahaemolyticus* only.

Molecular identification of the strains was performed by amplifying and sequencing the 16S rRNA gene sequences and

comparing the results to the database of known 16S rRNA gene sequences. As shown in Fig. 1, the results of the identification showed that the 14 isolated strains belonged to *Enterococcus faecalis*, *Lactococcus lactis* and *Lactococcus garvieae*. *Enterococcus faecalis* have antagonistic activity against many pathogenic bacteria. According to Allameh *et al.* (2017), isolation of *E. faecalis* from intestine of snakehead fish, *Channa striatus* showed high inhibitory effect against pathogens; *A. hydrophila*, *Pseudomonas aeruginosa* and *Shewanella putrefaciens* when tested by well diffusion method. On the other hand, the colonization of *E. faecalis* with *C. butyrium*, *B. mesenteries* and *L. sporogenes* in prawn gut improved the survival and growth of *M. rosenbergii* postlarvae (Jayanthi *et al.*, 2015). This proved that *E. faecalis* is a potential probiotic and portrays an important role in improving the host defense mechanism against bacterial infection (Allameh *et al.*, 2017). Besides, *L. lactis* are used in aquaculture industry in order to prevent cultured species from pathogenic bacteria (Balcazar *et al.*, 2009).

The growth of several pathogens such as *A. hydrophila* and *V. alginolyticus* have been inactivated by candidate probiotics, isolated from the gastrointestinal tract of the host (Vine *et al.*, 2004). Probiotics *in vivo* can produce antimicrobial metabolites, and therefore can serve as prophylactic and therapeutic agents (Vine *et al.*, 2004). Probiotics can be used to

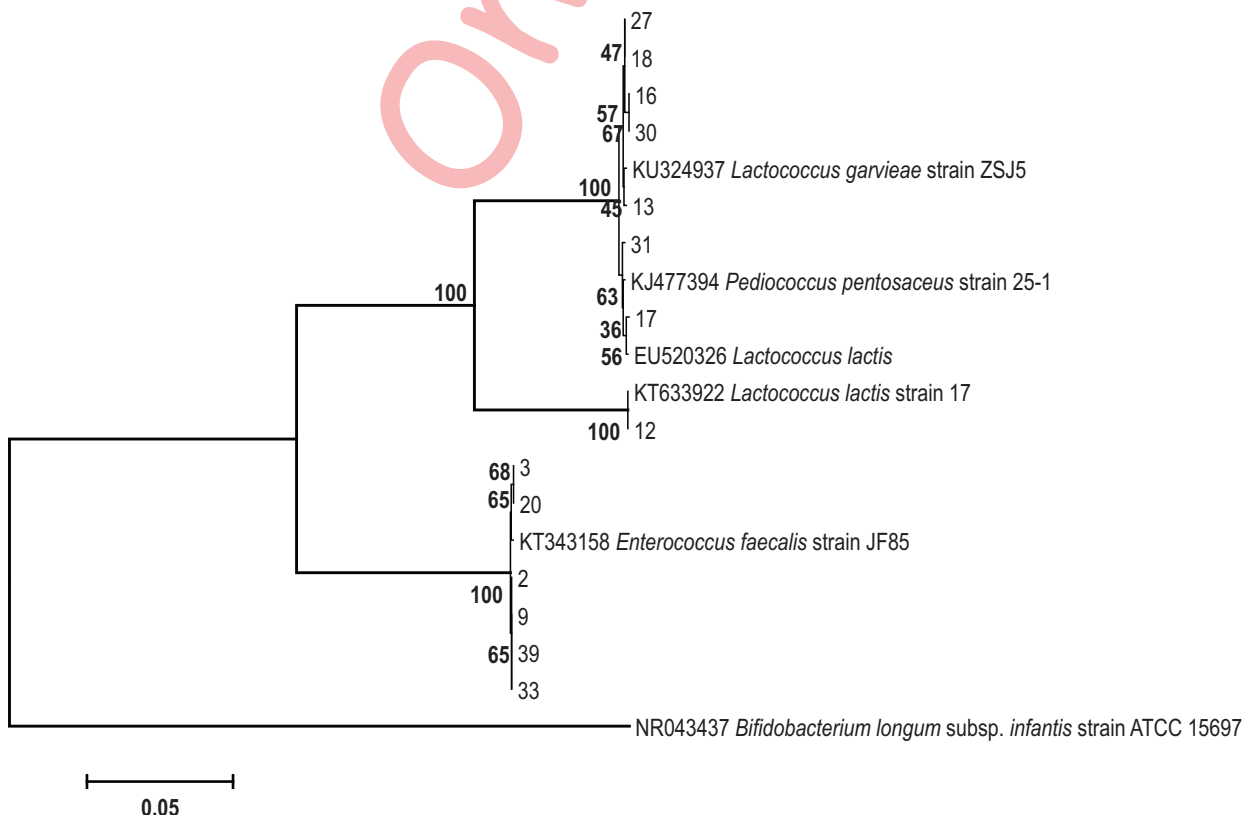


Fig. 1 : Phylogenetic neighbour-joining tree deduced from analysis nucleotide sequences of isolated bacteria

Table 1 : Antimicrobial activities of 14 strains against various pathogens. Clear zones: +: inhibition formed; -: no inhibition by well diffusion method

Strains	Growth of inhibition of pathogens		
	<i>V. parahaemolyticus</i>	<i>V. alginolyticus</i>	<i>A. hydrophila</i>
2	+	+	+
3	+	+	-
9	+	+	+
12	+	+	+
13	+	+	-
16	+	+	-
17	+	-	-
18	-	+	-
20	+	+	-
27	+	+	-
30	+	+	-
31	+	+	-
33	+	+	-
39	-	+	-

restrain the bacterial pathogens in fish and shrimp hatchery (Panigrahi et al., 2005; Balcazar et al., 2009; Ng et al., 2014). Based on positive results of this study, *Lactococcus lactis*, *Enterococcus faecalis* and *Lactococcus garvieae* should be further studied in feeding trials to investigate the effects of these probiotic bacterial on growth performance and immune response in *M. rosenbergii*.

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