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## Probiotic Potential of Gut Associated Bacteria from Indigenous Fresh Water Ornamental Fishes of Kerala, South India

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**Abstract** Probiotic potential of gut associated aerobic and facultative anaerobic bacterial flora of indigenous freshwater fishes such as *Puntius filamentosus* and *Barilius bakeri*, were analysed in this study. Total viable count (TVC) of heterotrophic bacteria ranged between  $0.64 \times 10^7$  to  $1.31 \times 10^7$  and  $0.59 \times 10^7$  to  $1.92 \times 10^7$  per gram in gut of *Barilius bakeri* and *P. filamentosus* respectively. While bacteria belonging to the genus *Corynebacterium* dominated the gut of *P. filamentosus*, *Bacillus* was found to be dominant genus in the gut of *B. bakeri*. More than 50% of bacterial isolates from both these fishes were capable of producing various exoenzymes such as amylase, gelatinase and lipase, with 15% of them showing excellent amylolytic and gelatinolytic activity. Selected bacterial isolates were tested for antagonistic activity against fish, shrimp and human pathogens, which revealed 15% of isolates having antagonistic activity against at least one pathogenic *Vibrio* species tested. These isolates were further tested for their ability to grow under different temperature, pH and salinity conditions in order to evaluate their suitability for application under different field conditions. The result of the present study offer scope for further research to evaluate probiotic potential of these gut associated bacteria in the larval rearing system and hatchery operations.

**Keywords** Probiotics; Ornamental fish culture; Gut microflora; *Puntius filamentosus*; *Barilius bakeri*

### Introduction

Ornamental fish keeping is one of the most popular hobbies of the world today. With the increase in demand for ornamental fishes especially in developed countries, many countries in Asia have started farming and nearly 60% of the international trade in ornamental fishes originates from developing countries. The export of ornamental fishes from India is at present mainly confined to freshwater varieties and the export is limited to fishes collected from wild (Madhu et al., 2009). More than 230 fish species have been reported from various river systems of Kerala and almost half of them are ornamental species (Radhakrishnan and Kurup, 2006). Extensive collection of these fishes from wild stock resulted in the decline of fish diversity in riverine system.

Hatchery rearing is an alternative to overcome this situation. However, disease outbreaks during the early developmental stages results in economic loss and is

the main constraint in preventing the success (Verschuere et al., 2000). Controlling diseases through antibiotic treatment is also leading to the emergence of drug resistant pathogens, which are difficult to treat in the long run. In recent years, control of diseases by environment friendly methods such as probiotic bacteria and immunostimulants has gained attention of researchers (Castex et al., 2008). The use of probiotic bacteria to control potential pathogens is showing good results (Gomez- Gil et al., 2000).

Kerala is blessed with 42 rivers and most of them support unique fish fauna, which include potential ornamental fishes. Considering this potential, government of Kerala has also initiated steps to promote ornamental fish culture in small scale house hold ponds and tanks. As a precondition to commercial exploitation, techniques need to be developed for successful breeding and rearing of potential indigenous ornamental fishes. In the present study

fishes such as *Puntius filamentosus* and *Barilius bakeri*, which have ornamental value, were chosen to study their gut microbiota and to evaluate their probiotic potential. The study has been taken on the assumption that an understanding of the gut microbiota of these indigenous ornamental fishes, might lead to valuable insight for the development of possible strains that could be used as probiotics in the breeding, larval rearing, and growth enhancement.

## 1 Materials and Methods

### 1.1 Description of the Collection site and Fishes selected for the Study

Fresh water fishes namely *Puntius filamentosus* (Filament Barb) and *Barilius bakeri* (Malabar Baril) (Plate 1b & c) were collected from its natural habitat Chalakudi River, Latitude: 10 09' 44' N; Longitude: 76 15' 56'' E (Plate 1 a) and brought alive to the laboratory. Fishes were collected with the help of professional fisherman, who caught them using cast net. After taking the morphometric measurements such as total

length (TL) and standard length (ST) the fishes were dissected out aseptically using sterile surgical blade. The entire gut region was aseptically removed, weighed and homogenized using sterile glass homogenizer, and serially diluted up to  $10^{-6}$  using 10% phosphate buffer solution of pH 7.2.

Aliquots of 0.2 ml samples from each dilution were spread plated in duplicate on nutrient agar with media composition of peptic digest of animal tissue and NaCl - 5g/L each, Beef extract and Yeast extract - 1.5 g/L each, Agar - 15g/L (Himedia- Mumbai) for the enumeration of cultivable bacterial flora aerobic heterotrophic bacteria. The plates were then incubated at 30°C for 24 hours. Colonies developed on the plate were counted and expressed as colony forming units (cfu) /mL of fish gut. Well separated and morphologically different colonies were picked up using a sterile inoculation needle and transferred to sterile nutrient agar slants. The isolates were purified by quadrant streaking and were stored in nutrient agar

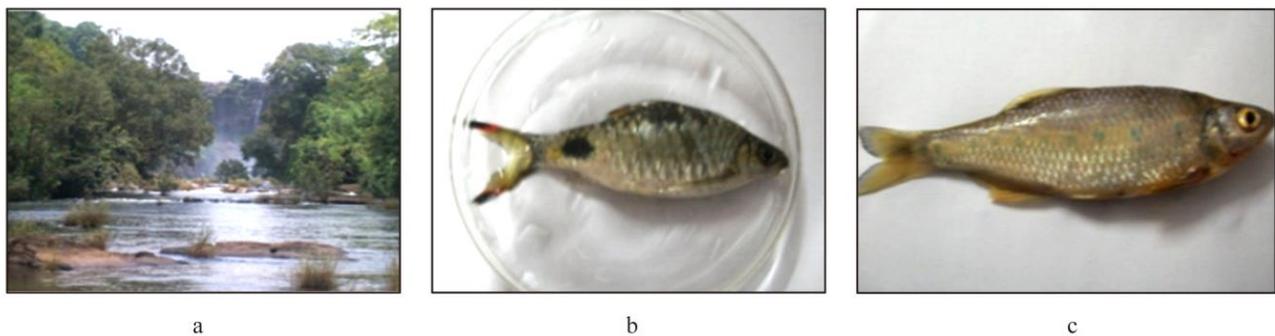


Plate 1 a) Chalakudi river system b) *Puntius filamentosus* c) *Barilius bakeri*

slants for the further study. Estimation of microbiota of the gut was done using standard methods (Ringø et al., 1995). The isolated bacterial strains were identified up to generic level using the taxonomic key by Buchanan & Gibbons (1984). For generic level classification of the isolates they were subjected to various tests such as Gram stain, spore stain, motility, Kovacs oxidase test, catalase activity, and oxidation / fermentation test along with morphological characteristics of the colony.

### 1.2 Hydrolytic Enzyme production potential of the isolates

Ability of the gut associated bacteria from *P. filamentosus* and *B. bakeri* to produce various hydrolytic enzymes

such as amylase; lipase and gelatinase were checked by plate assay. For amylase and gelatinase activity, nutrient agar plates with 1% soluble starch and 1% gelatin respectively were prepared and the cultures were spot inoculated in the plates. After incubation of the plates at room temperature ( $28 \pm 2^\circ\text{C}$ ) for 48 h, nutrient agar plates with 1% soluble starch were flooded with Lugol's iodine and nutrient agar plates with 1% gelatin flooded with mercuric chloride solution (Frazier, 1926). The presence of a clear zone around the colonies was noted as a positive result in both tests. For lipase activity, nutrient agar plates with 1% Tween-80 were used. Test organisms were spot inoculated and plates were incubated at room temperature for 3-4 days. A positive result was

indicated by zone of clearing around the colonies of lipolytic organisms (Rhodes, 1959).

### 1.3 Determination of Antibacterial activity of the Isolates

Antibacterial activity of heterotrophic bacteria isolated from the gut of *P. filamentosus* and *B. bakeri* were determined following the method described by (Mujeeb et al., 2010). The antibacterial activity was assayed by disc diffusion method against different fish, shrimp and human pathogens such as *Vibrio harveyi*, *V. alginolyticus*, *V. splendidus*, *V. fluvialis*, *V. cholera*, *Pseudomonas aeruginosa*, *Aeromonas hydrophila*, *A. caviae* and *Salmonella* strain 14 and 44. Filter paper discs of 6 mm diameter were cut out from Whatmans No. 1 filter paper and sterilized at 15 lbs for 15 min. The bacterial isolates were enriched in nutrient broth by overnight incubation at 37°C for 16 hours. The filter paper discs were then impregnated with 20 µl of broth cultures of the test organisms. These discs were then placed over sterile tryptic soy agar (TSA) seeded with test pathogen. The seeded plates along with test culture were then incubated at 37°C for 24 hours. Formation of clearing zone around the discs was considered positive indication of inhibitory activity.

### 1.4 Determination of Growth abilities of the Selected Isolates

Growth of any organism depends on the certain physical and chemical parameters, and each organism has a minimum and maximum range of tolerance beyond which, their growth is inhibited. Metabolic activity of bacteria is directly linked to the environmental parameters like temperature, salinity, and pH (Monod, 1949). Isolates showing good enzyme activity were selected for the growth studies. Isolate numbers T3, T6, T8, T14 and T23 were selected from *B. bakeri* and P3, P7, P8, P14 and P20 from *P. filamentosus*.

In order to check the effect of temperature on growth, these isolates were then inoculated into 5ml sterile nutrient broth (pH 7) and incubated at various temperatures such as 4, 20, 27, 37 and 50°C. Similarly effect of pH on growth of these isolates was monitored by inoculating the cultures into sterile nutrient broth prepared with various pH such as 2, 4, 6 and 8. Ability to grow in various salinity was tested similarly by monitoring the growth in nutrient broth maintained at various salinities such as 0, 10, 15 and 25 ppt. The

inoculated nutrient broth tubes for checking the effect of pH and salinity were incubated at (28 ± 2°C). Growth is measured in terms of optical density at 620 nm, after 10, 24 and 48 hours.

## 2 Results and Discussion

### 2.1 Enumeration and Characterisation of Gut bacteria

Total viable count (TVC) of heterotrophic bacteria ranged between 0.64 x 10<sup>7</sup>-1.31 x 10<sup>7</sup> cfu/mL in the gut of *Barilius bakeri*, 0.59 x 10<sup>7</sup> -1.92 x 10<sup>7</sup> cfu/mL guts of the fish *P. filamentosus*. A total of fifty eight isolates were selected for characterisation and further analysis from the fish gut samples, (32 from the *B. bakeri* and 26 from the *P. filamentosus*). Characterisation of the isolates revealed that *Corynebacterium* is the predominant genus in the gut of *P. filamentosus* followed by genera such as *Kurthia*, *Micrococcus* and *Staphylococcus*. *Bacillus* sp. was found to be the dominant genera in the gut of *B. bakeri* followed by genera *Corynebacterium*, *Micrococcus*, *Planococcus*, *Kurthia* and *Staphylococcus*. (Figure 1). In both these fishes the gut bacterial flora was found to be dominated by Gram positive forms. These results are in agreement with the microbiota that has been reported previously from the guts of different species of fishes (Ringø et al., 1998; Ray et al., 2012). Both marine and freshwater water fish have been shown to have a specific indigenous gut microbiota and it may change with age, nutritional status, and environmental conditions (Olafsen, 2001; Vine et al., 2006). In general, indigenous microbiota of freshwater fish species tends to be dominated by members of the genera *Aeromonas*,

Table 1 Details of pathogenic bacteria used in the study

SI No	Name of the Pathogen	Source of the isolate
1	<i>Vibrio harveyi</i>	Shrimp farm at Cochin
2	<i>V. alginolyticus</i>	Shrimp farm at Cochin
3	<i>V. splendidus</i>	Shrimp farm at Cochin
4	<i>V. fluvialis</i>	Shrimp farm at Cochin
5	<i>Pseudomonas aeruginosa</i>	Shrimp farm at Cochin
6	<i>V. cholerae</i>	Cochin estuary
7	<i>Aeromonas hydrophila</i>	Freshwater ornamental fish farm at Cochin
8	<i>A. caviae</i>	Freshwater ornamental fish farm at Cochin
9	<i>Salmonella</i> 14	World Health Organization
10	<i>Salmonella</i> 44	World Health Organization

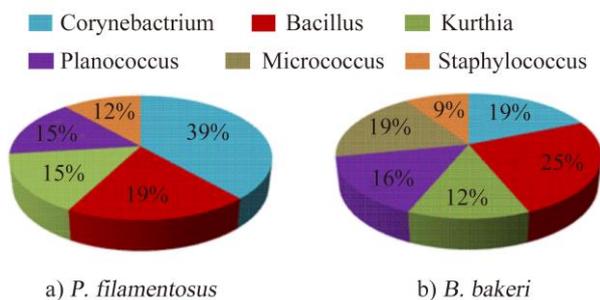


Figure 1 Distribution of various genera of heterotrophic bacteria in the gut of a) *P. filamentosus* and b) *B. bakeri*

*Plesiomonas*, representatives of the family Enterobacteriaceae, and obligate anaerobic bacteria of the genera *Bacteroides*, *Fusobacterium*, and *Eubacterium* (Sakata, 1990). However, the aerobic heterotrophic bacteria of the gut of *P. filamentosus* and *B. bakeri* were found to be dominated by Gram positive forms. These variations are accounted due to factors like bacterial host specificity, food type, and water resource (Verner et al., 2003).

## 2.2 Hydrolytic enzyme production potential of the Bacterial isolates from Fish Gut

These diverse bacterial flora of the gastrointestinal tract represent a diversified enzymatic potential. Resident intestinal bacteria in fish are known to accelerate the digestion process by producing extracellular enzymes (Stickney and Shumway 1974; Cahill, 1990; Subha, 2013). Understanding of the enzyme producing potential of gut microbiota may help in formulating feeds and probiotics for larval rearing. All isolates were screened for their ability to produce hydrolytic enzymes such as amylase, gelatinase, and lipase. Results revealed widespread hydrolytic enzyme production potential among the isolates from *B. bakeri* (Figure 2). The isolates P3, P6, P7, P8, P14, P 21, and P 25 from *P. filamentosus* showed maximum activity against the all the screened enzymes. In the case of *B. bakeri*, maximum activity was showed by T3, T5, T8, T20, and T23. Sixty five percentage of the isolates are capable of producing two of the three enzymes in *B. bakeri*, in case of *P. filamentosus* 84% of the isolates were able to do so.

The composition of enzyme producing bacterial flora in the fish digestive tracts are correlated to their feeding habits. Kar and Ghosh (2008) reported higher densities of proteolytic bacterial strains in carnivore

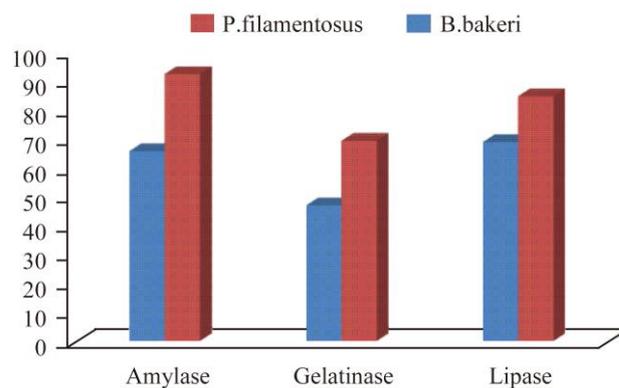


Figure 2 Percentage of isolates capable of producing hydrolytic enzymes

bottom feeder *Channa punctatus* and cellulolytic strains in herbivore column feeder, *Labeo rohita*. Lipase producing bacteria were abundant in the guts of fishes which prey on a variety of live organisms (copepods, mysids, amphipods) rich in highly unsaturated fatty acids (Murugan et al., 2009). The occurrence of proteolytic, cellulolytic, and amylolytic bacteria in the gut has been suggested as an omnivorous feeding habit of the fish (Ghosh et al., 2010). Diverse bacterial flora with different enzyme producing activity reflects the omnivorous feeding nature of these two fishes analyzed. This result indicates that there is also a distinct microbial source of enzymes, apart from the endogenous sources in fish gastrointestinal tracts. In fish, it has been reported that *Bacteroides* and *Clostridium sp.* have contributed to the host's nutrition, especially by supplying fatty acids and vitamins (Sakata, 1990). Depending on the diameter of clearing zone the isolates were classified as those with very good, good and poor activity (Table 2) on enzyme assay plates. Results revealed that around 15% of the isolates (Table 2) have very good amylolytic and gelatinolytic activity.

## 2.3 Antibacterial activity of Fish Gut bacteria

Bacterial strains with inhibitory activity have been shown to inhibit pathogenic bacteria both *in vitro* and *in vivo* through several different mechanisms. Selected bacterial isolates that showed good hydrolytic enzyme potential were further screened for their antagonistic activity against shrimp, fish and human pathogens belonging to the genera *Vibrio*, *Aeromonas*, *Pseudomonas* and *Salmonella* that generally causes diseases in fish, shrimp and human. Maximum inhibitory activity was

Table 2 Range of enzyme activity of the bacterial isolates from gut of *B. bakeri* and *P. filamentosus*

Hydrolytic Ezymes	Very Good*		Good**		Poor***	
	Bb	Pf	Bb	Pf	Bb	Pf
Amyase	15.62	15.38	50.00	76.92	42.30	7.69
Gelatinase	15.62	15.38	31.25	53.84	65.38	30.76
Lipase	3.12	11.53	65.62	73.07	38.46	15.38

Note: \* $\geq 20$ mm; \*\*10-19mm; \*\*\* $\leq 9$ mm; Bb – *Barilius bakeri*, Pf: *Puntius filamentosus*

shown by isolates T14 and T3 against *Vibrio alginolyticus*. None of the isolates were antagonistic against *Aeromonas spp.*, *Salmonella spp.*, and *Pseudomonas aeruginosa*. Out of 58 isolates, 25% showed antagonism against at least one pathogenic *Vibrio* (Table 3). These results suggest that intestinal bacteria with antibacterial abilities may inhibit the growth of invading bacteria in intestine of fish, to some extent. The antibacterial effect of bacteria is generally due to production of antibiotics, bacteriocins, siderophores, lysozymes or proteases, and alteration of pH values by the organic acids either singly or in combination: (Gatesoupe, 1999; Balcazar et al., 2006). Though the antibacterial activity of the isolates in general was found to be less some of the isolates showed very good activity shrimp pathogens such as *V. alginolyticus*, *V. splendidus* and *V. fluvialis*.

Table 3 Antibacterial activity of selected isolates from the gut of *Barilius bakeri* and *Puntius filamentosus*

Name of Pathogen	Isolate Number									
	T	T	T	T	T	P	P	P	P	P
	3	6	8	1	2	3	7	8	1	2
				4	3				4	0
<i>Vibrio harveyi</i>	-	-	-	-	-	-	-	-	-	-
<i>V. alginolyticus</i>	+	+	-	+	+	+	+	-	-	+
<i>V. splendidus</i>	+	-	+	-	-	-	-	+	-	-
<i>V. fluvialis</i>	+	-	-	+	+	+	-	-	+	+
<i>Pseudomonas aeruginosa</i>	-	-	-	-	-	-	-	-	-	-
<i>V. cholerae</i>	-	-	-	-	-	-	-	-	-	-
<i>Aeromonas hydrophila</i>	-	-	-	-	-	-	-	-	-	-
<i>A. caviae</i>	-	-	-	-	-	-	-	-	-	-
<i>Salmonella 14</i>	-	-	-	-	-	-	-	-	-	-
<i>Salmonella 44</i>	-	-	-	-	-	-	-	-	-	-

++ > 20 mm; + = 10-19 mm; - No activity

## 2.4. Effect of environmental parameters on growth of selected isolates

Although different bacterial genera can invade in to the gut of the fish, colonisation and survival depends on their adaptability to the microenvironment of the gut. The process of colonisation in early developing fish larvae or fry is complex and seems to be affected by bacterial load in the water (Ringø et al., 1995), which in turn a proxy of physico chemical parameter of the system. Understanding the optimum condition required for the colonisation of the isolates may help in the development of culture system and as probiotics in later stage. Optimum growth is obtained at 37°C, while temperature range preferred by majority of the isolates was in between 27°C-37°C; none of the isolates were able to grow at 4°C. Selected isolates showed growth in all range of salinity (0-25ppt) used in the experiment, with maximum growth rate at 25 ppt, showing the requirement of salt content for better growth. No growth is noted below pH 2, and optimum range favoured by most of the isolates was between 6 and 8. (Table 4).

## 3 Conclusion

Bacteria present in the aquatic environment may influence the composition of the gut microbiota in fish and impossible to avoid them being a component of the diet (Cahill, 1990). The present study shows gut of *B. bakeri* and *P. filamentosus* were colonized by diverse genera of bacteria and are capable of producing hydrolytic enzyme at varying levels. It seems logical to think that the enzymatic mass lodged in the digestive tract might interfere in a considerable way with a major part of the metabolism of the host animal (Rasiah et al., 2009). Population levels of amylolytic strains were highest in the gut of both fishes. This can be correlated with the omnivorous feeding habit of the fishes. This also indicates that there is a distinct microbial source of digestive enzymes, such as amylase, gelatinase and lipase apart from the endogenous sources in fish gastrointestinal tracts. This information along with the antibacterial activity of the selected isolates justifies their ability as a potential probiotic in ornamental fish culture.

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Table 4 Effect of environmental parameters on growth of selected isolates from the gut of *B. bakeri* and *P. filamentosus*

Isolate No	Growth parameter												
	Temperature					Salinity (ppt)				pH			
	4 °C	20 °C	27 °C	37 °C	50 °C	0	10	15	25	2	4	6	8
T 3	--	+	+	+	+	+	+	+	+	--	+	+	+
T 6	--	+	+	+	+	+	+	+	+	--	+	+	+
T 8	--	+	+	+	+	+	+	+	+	--	+	+	+
T 14	--	+	+	+	+	+	+	+	+	--	+	+	+
T 23	--	+	+	+	+	+	+	+	+	--	+	+	+
P 3	--	+	+	+	+	+	+	+	+	--	+	+	+
P 7	--	+	+	+	+	+	+	+	+	--	+	+	+
P 8	--	+	+	+	+	+	+	+	+	--	+	+	+
P 14	--	+	+	+	+	+	+	+	+	--	+	+	+
P 20	--	+	+	+	+	+	+	+	+	--	+	+	+

-- No Growth                      + Growth

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