

E-ISSN: 2320-7078 P-ISSN: 2349-6800 www.entomoljournal.com

JEZS 2020; 8(6): 1326-1330 © 2020 JEZS Received: 15-08-2020 Accepted: 28-09-2020

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Journal of Entomology and Zoology Studies

Available online at www.entomoljournal.com



Detection of virulence Hemolysin gene (HlyA) in Photobacterium damselae subspecies damselae isolates

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DOI: https://doi.org/10.22271/j.ento.2020.v8.i6r.8015

Abstract

Polymerase chain reaction (PCR) technique was used to assay for the detection of virulence gene in the genomes of the *Photobacterium damselae* subsp. *damselae* isolated from cage culture and wild source fishes, particularly *hlyA* gene responsible of hemolysin toxins production in this genus. Sixty one strains of *P. damselae* subsp. *damselae* were isolated from cage and wild caught fishes. A pair of primers was designed to detect the hemolysin gene (*HlyA*) of *P. damselae* subsp. *damselae* by PCR method. The expected product size (417bp) was obtained from *P. damselae* subsp. *damselae* isolates. *Photobacterium damselae* grew on TCBS agar plate producing green colonies whereas *P. damselae* subsp. *piscicida* did not grow. The PCR methods used are cost and labour effective when compared with the other molecular methods and commercially available kits.

Keywords: HlyA gene, PCR, Photobacterium damselae subsp. damselae, cage culture, wild fish, primer

Introduction

Aquaculture production of aquatic organisms including fish, crustaceans, and mollusks is the fastest growing aquaculture industry globally; while wild capture fisheries have become relatively stable in recent years, aquaculture production has increased by about 9% per year since 1985 ^[1]. Water pollution is one of the most important global problems in aquaculture field. Aquaculture and brackish water fishes can be infected when urban wastes mix with farm water resulting in contamination with bacterial and other microorganism's infection ^[2]. Photobacterium damselae subsp. damselae which was formely classified as Vibrio damselae is a halophilic bacterium causing skin ulcers in warm and cold water fishes [3, 4, 5]. Photobacterium damselae subsp. damselae has been reported to cause wound infections and fatal necrotizing fasciitis in humans ^[6, 7, 8]. Outbreaks of photobacteriosis may occur in the temperature range of 14-29 °C and at salinities of 3-21ppt however the optimum range for acute disease is 18-25 °C and 5-15ppt. Photobacterium damselae subsp. piscicida is considered an obligate pathogen and its survival is short lived outside the host even in saltwater conditions ^[9]. Hemolysins contribute to the virulence of *P. damselae* subsp. damselae for mice and fish ^[10]. Strains of particular virulence produce plasmid-encoded damselysin (Dly) and phobalysin P (PhlyP), a phospholipase D, and a small pore-forming toxin (PFT), respectively. All hemolytic strains express chromosomally encoded phobalysin C (PhlyC), which is closely related to PhlyP^[11, 12]. Extracellular products (ECPs) from P. damselae subsp. damselae have been shown to display cytotoxic activity for different fish and mammalian cell lines. Only virulent strains produce toxic ECPs and the cytotoxic components are thermo labile ^[13]. Stephens *et al.* ^[14] investigated the fish infected with *P. damselae* subsp. piscicida suffer from an acute septicemia, lethargic, swim slowly near the surface and ultimately sink and rest on the bottom prior to death. Infected fish show haemorrhagic septicaemia and the development of whitish areas or granulomas in the spleen, kidney and liver [15].

Molecular diagnosis of disease outbreaks is important for proper management practice and effective control. Recent microbiological methods of culture and biochemical characterization are time consuming and labour intensive. Diagnosis methods using immunological techniques such as agglutination or the ELISA are comparatively quicker ^[9, 16, 17]. PCR is not only more sensitive, but also faster ^[18], cost and labour effective when compared with biochemical analysis. The aim of the present study was to develop a convinced PCR-based test for *P. damselae* subsp. *damselae*.

The two subspecies have an identical *16S rRNA* gene sequences ^[19] but they differ in their biochemical and physiological characteristics ^[20, 21]. A pair of primers was designed to detect *P. damselae* subsp. *damselae* by PCR. Also, thiosulphate citrate bile salts–sucrose agar (TCBS) was used to differentiate *P. damselae* subsp. *damselae* from *P. damselae* subsp. *piscicida*.

Materials and Methods Bacterial strains

The sixty one strains were obtained from cage culture and wild fishes in south east coast of India. All strains of *P. damselae* subsp. *damselae* were cultured in TCBS agar with 2% NaCl.

DNA primers

A pair of primers was selected to amplify a 417bp fragment of hemolysin gene (*HlyA*) derived from *P. damselae* subsp. *damselae*. The forward primer, HlyA30FW, was 20 nucleotides long (5'- CGCTTTCGGACCATCTTTAC-3') corresponding to positions 13321-13340 of the *P. damselae* subsp. *damselae* gene for hemolysin (Fig. 1) and the reverse primer, HlyA30RV, was 20 nucleotides long (5'-TGGGATAGAGGAAGTCGTTC-3') corresponding to positions 13718-13730). Primers were designed using Primer3plus software (www.bioinformatics.nl/primer3plus) and then the analysis of the designed primers was done using DNA MAN and manually check was the target gene.

HlyA30FW						
	13321	13331	13341	13351	13301	13371
13321	CGCTTTCGGA GCGAAAGCCT	GGTAGAAATG	TAACTTCAGC ATTGAAGTCG	ACCACCAGTA TGGTGGTCAT	ACACCAATAG TGTGGTTATC	TGAATCCAGA ACTTAGGTCT
	13381	13301	13401	13411	15421	13431
13381	AGATTCATTT TCTAAGTAAA	ATATTATAAT TATAATATTA	TAGAATTAAT ATCTTAATTA	ATTCTCCCGA TAAGAGGGCT	GGGAACGTTT CCCTTGCAAA	TTAATATCTG AATTATAGAC
	13441	13451	13401	13471	13481	13401
13441	AGCTTTTTTA TCGAAAAAAT	TTTGAAGTAG AAACTTCATC	ATATATCAAA TATATAGTIT	TAAATAATCT ATTTATTAGA	TGAGCAATAG ACTCGTTATC	CACTGGCTGA GTGACCGACT
	13501	13877	13521	13531	12541	13551
13501	CCACTCTCTC GGTGAGAGAG	GCCCAACCAG CGGGTTGGTC	ATACCACCTG TATGGTGGAC	ATAATTAGCC TATTAATCGG	CAATAATATT GTTATTATAA	TTTGCGTCAG AAACGCAGTC
	13501	13871	13581	13501	13601	13011
13561	TGTATCATTT ACATAGTAAA	AGATGTATAC TCTACATATG	CGGCACCAGT GCCGTGGTCA	CGTATTATCA GCATAATAGT	TCTAAGCTAA AGATTCGATT	TTCGAACTAT AAGCTTGATA
	13621	13631	13541	13481	13001	13671
13621	TTTTGCATCA AAAACGTAGT	GGCGTTGCAA CCGCAACGTT	CGCCGTTAGT GCGGCAATCA	ACCAAATGAT TGGTTTACTA	AAAGAACGCT TTTCTTGCGA	CTAATATAAC GATTATATTG
	13581	13601	13701	13711	13721	13731
13681	TTGATAAATA AACTATTTAT	AGTGAAATAT TCACTTTATA	TACCATTTTT ATGGTAAAAA	ACAGAAAGAA TGTCTTT <mark>CTT</mark>	CGACTTCCTC GCTGAAGGAG	TATCCCAAAT ATAGGGTTTA
					HlyA30RV	

Fig 1: Nucleotide sequence of Photobacterium damselae subsp. damselae hemolysin gene (*HlyA*). Underlined nucleotides (HlyA30FW, HlyA30RV) correspond to the pair of primers used to amplify DNA in PCR Assays

Preparation of samples DNA extraction

Pure cultures were inoculated in trypticase soya broth and incubated for 24 hours. After 24hrs the cultures was centrifuged at 6000rpm and the supernatant is discarded. The bacterial pellet was collected aseptically and 1ml of DNA extraction solution (GeNeiTM HIMEDIA, Mumbai) was added into the cells. The mixture was homogenized well and incubated at 60°C for 60 min. The mixture was centrifuged at 10,000 x g for 10 min and the supernatant was collected in fresh sterile tube. To the supernatant, equal volume of absolute alcohol was added and centrifuged at 10,000 x g for 5 min. The supernatant was discarded and the DNA pellet was washed twice with 1 ml of 95% alcohol and centrifuged at 6000 x g for 5 min. The supernatant was removed and the DNA pellet was allowed to air dry for 5 min and 100 µl of deionized water was added and stored at 4°C for further use.

PCR reaction

The PCR condition includes 25 μ L reaction included 2 μ L forward and reverse primers, respectively, 6.5 μ L ultrapure water, 12.5 μ L master mix and 2 μ L template DNA. The thermal cycle (Biorad) program comprised 35 cycles of 5 min

at 95°C, 1min at 95°C, 1 min at 56°C, and 1min at 72°C, and a final 10 min extension at 72°C. The cycling conditions and sequence of the primer listed in the Table 1. The 10 μ l of amplified products were separated by electrophoresis on a 1% (w/v) agarose at 100V gel run for 50-90min in Trisborate – EDTA (TBE) buffer (0.89 M tris, 0.89 M boric acid, 0.02 M EDTA, pH 8.0) and a 100bp DNA ladder was used as a size reference. This gel was then stained with 0.5µg/ml ethidium bromide solution for 20 min and visualized by UV – induced fluorescence. The presence of a clear fragment with the correct amplification size was assessed as a positive signal indicating the presence of the gene.

 Table 1: Primer designed for target gene of Photobacterium

 damselae
 subsp. Damselae

Primer Name	Sequence	Product Size	
HlyA30FW	CGCTTTCGGACCATCTTTAC	417hm	
HIYA30 RV	TGGGATAGAGGAAGTCGTTC	417bp	

Results and discussion

The specific-PCR was performed using the primers HlyA30FW and HlyA30RV to determine whether a 417bp of *hlyA* gene fragment, respectively, could be detected among

the 61 P. damselae subsp. damselae isolates. Out of the 61 strains, 36/ 61 of P. damselae subsp. damselae (Fig. 2) isolates were PCR positive for the hlyA gene. This could be easily explained as the two subspecies differed only by a single nucleotide [22]. Rajan et al. [23] evaluated the pair of primers, CPSF and CPSR, designed from the capsular polysaccharide gene successfully amplified an expected 410bp DNA fragment of all P. damselae ssp. piscicida and P. damselae strains tested. Osorio et al. [19] reported that the two subspecies have the same 16S rRNA gene sequence and have used the nested PCR to differentiate P. damselae subsp. piscicida from P. damselae subsp. damselae. Toranzo et al. ^[24] and Zorrilla *et al.* ^[25] reported that *P. damselae* subsp. piscicida did not grow on TCBS agar plate. Conversely, it was reported that P. damselae subsp. damselae could grow on TCBS agar plate ^[3, 26]. The *Photobacterium damselae* subsp. damselae isolated from different cage culture and wild caught fishes (Table 2) in south east coast of India. Out of the 36 positive hlyA genes, 22.2% of the cage culture system isolates, 75% of wild caught isolates and 2.8% of marine ornamental fish isolates present the hlyA gene. Cage culture fish's incident of P. damselae subsp. damselae isolates during the summer season. The pH, salinity and temperature of the cage water were measured during the sampling period. Temperature of the water during the sampling period was 31 ± 0.5 °C, salinity (36%) and pH (8.0-8.1).

In order to identify P. damselae subsp. damselae, bacterial colonies of the bacteria that gave a PCR product were placed on TCBS plate. In previous reports, bacteria grown on BHIA were scraped off from the plates, resuspended in saline to extract DNA of the mixed cultures and were then used for PCR ^[12]. Photobacterium damselae subsp. piscicida could not grow on TCBS whereas P. damselae subsp. damselae. The sensitivity of the PCR was evaluated by serial dilution of P. damselae subsp. Piscicida cell suspension as detailed elsewhere ^[27]. Amplification resulted in detectable levels of PCR product when a minimum of 13CFU of P. damselae subsp. damselae was used (Fig. 3 and 4). In conclusion, Screening of specific hemolysin gene (HlyA) appeared to be the most effective way of detecting and characterizing P. damselae virulence factors. The high throughput and costeffective specific-PCR system used in this study could provide a powerful supplement to the conventional methods for a more accurate risk assessment and monitoring of P. damselae species in the wild and cage culture fishes.



Fig 2: Agarose gel electrophoretic analysis of the PCR amplified product using HlyA30FW and HlyA30RV primers (417bp) for detection of virulence hemolysin gene. A: Lane 1-MP66 (HlyA), Lane 2-MP76 (HlyA), Lane 3-MP140, Lane 4-MP147, Lane 5-MP148, Lane 6-MP157, Lane 7-MP160, Lane 8- MP162 (HlyA), Lane 9-MP189 (HlyA), Lane 10-MP190 (HlyA), Lane 11-MP191 (HlyA), Lane 12-MP192, Lane 13-MP193, Lane 14-MP194 (HlyA), Lane 15-MP195, Lane 16-Negative control, Lane 17-100bp DNA marker.



Fig 3: Agarose gel electrophoretic analysis of the PCR amplified product using HlyA30FW and HlyA30RV primers (417bp) for detection of virulence hemolysin gene. B: Lane 16-MP196 (HlyA), Lane 17-MP197 (HlyA), Lane 18-MP198, Lane 19-MP199, Lane 20-MP224, Lane 21-MP233, Lane 22-MP234, Lane 23- MP235, Lane 24-MP236 (HlyA), Lane 25-MP237, Lane 26-MP238 (HlyA), Lane 27-MP239 (HlyA), Lane 28-MP240 (HlyA), Lane 29-MP241, Lane 30-MP242, Lane 31-MP244, Lane 32-MP245 (HlyA), Lane 33-MP246 (HlyA), Lane 34-MP247, Lane 35-MP248 (HlyA), Lane 36-MP249 (HlyA), Lane 37-MP250, Lane 38-MP251, Lane 39-MP252 (HlyA), Lane 40-MP253 (HlyA), Lane 41-MP254 (HlyA), Lane 42-MP255 (HlyA), Lane 43-MP270, Lane 44-MP271 (HlyA), Lane 45-MP272 (HlyA), Lane 46-Negative control, Lane 47-100bp DNA marker.



Fig 4: Agarose gel electrophoretic analysis of the PCR amplified product using HlyA30FW and HlyA30RV primers (417bp) for detection of virulence hemolysin gene. C: Lane 46-MP273 (HlyA), Lane 47-MP274 (HlyA), Lane 48-MP276, Lane 49-MP277 (HlyA), Lane 50-MP278 (HlyA), Lane 51-MP279 (HlyA), Lane 52-MP280 (HlyA), Lane 53- MP281 (HlyA), Lane 54-MP282 (HlyA), Lane 55-MP283 (HlyA), Lane 56-MP284 (HlyA), Lane 57-MP285 (HlyA), Lane 58-MP286 (HlyA), Lane 59-MP287, Lane 60-MP288 (HlyA), Lane 61-289MP, Lane 62-Negative control, Lane 63-100bp DNA marker.

Table 2. Details of internotysin gene(11) detection in 1 notobacter turn autisetate subsp. autisetate isola	t haemolysin gene(A) detection in <i>Photobacterium damselae</i> subsp. <i>damselae</i> isolates
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Sl. No	Isolate No	Source	Fish Name	Hly A gene
1.	MP66	Marine ornamental fish farm	Chaetodon octofasciatus	Р
2.	MP76	Landing center (Theraspuram)	Scarus globiceps	Р
3.	MP162	Cage culture system (Thangachimadam)	Lates calcarifer	Р
4.	MP189	Cage culture system (Thangachimadam)	Lates calcarifer	Р
5.	MP190	Cage culture system (Thangachimadam)	Lates calcarifer	Р
6.	MP191	Cage culture system (Thangachimadam)	Lates calcarifer	Р
7.	MP194	Cage culture system (Thangachimadam)	Lates calcarifer	Р
8.	MP196	Cage culture system (Thangachimadam)	Lates calcarifer	Р
9.	MP197	Cage culture system (Thangachimadam)	Lates calcarifer	Р
10.	MP236	Cage culture system (Thangachimadam)	Lates calcarifer	Р
11.	MP238	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
12.	MP239	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
13.	MP240	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
14.	MP245	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
15.	MP246	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
16.	MP248	Landing centers (Tharuvaikulam)	Scarus globiceps	Р
17.	MP249	Landing centers (Theraspuram)	Brama brama	Р
18.	MP252	Landing centers (Theraspuram)	Brama brama	Р
19.	MP253	Landing centers (Theraspuram)	Brama brama	Р
20.	MP254	Landing centers (Theraspuram)	Brama brama	Р
21.	MP255	Landing centers (Theraspuram)	Brama brama	Р
22.	MP271	Landing centers (Theraspuram)	Brama brama	Р
23.	MP272	Landing centers (Theraspuram)	Brama brama	Р
24.	MP273	Landing centers (Theraspuram)	Brama brama	Р
25.	MP274	Landing centers (Pamban)	Lutjanus sebae	Р
26.	MP277	Landing centers (Pamban)	Lutjanus sebae	Р
27.	MP278	Landing centers (Pamban)	Lutjanus sebae	Р
28.	MP279	Landing centers (Pamban)	Lutjanus sebae	Р
29.	MP280	Landing centers (Pamban)	Lutjanus sebae	Р
30.	MP281	Landing centers (Pamban)	Lutjanus sebae	Р
31.	MP282	Landing centers (Pamban)	Lutjanus sebae	Р
32.	MP283	Landing centers (Pamban)	Lutjanus sebae	Р
33.	MP284	Landing centers (Pamban)	Lutjanus sebae	Р
34.	MP285	Landing centers (Pamban)	Lutjanus sebae	Р
35.	MP286	Landing centers (Pamban)	Lutjanus sebae	Р
36.	MP288	Landing centers (Pamban)	Lutjanus sebae	Р

Acknowledgements

The study was part of the PhD thesis of the first author in Aquatic Animal Health of TNJFU. Financial support from National Surveillance Programme for Aquatic animal disease - sub project no: 21 is gratefully acknowledged.

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