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Molecular Characterization of Enterotoxigenic *Bacillus cereus* in Tropical Fisheries Environment of Cochin, India

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INTRODUCTION

Bacillus cereus is a Gram positive spore-forming organism, which causes two different types food poisoning viz. diarrhoeal type and emetic type. Apart from these, *B. cereus* has also been reported to be associated with ophthalmitis, respiratory tract infection and central nervous system disorder (Bekemeyer and Zimmerman, 1985; Beer *et al.*, 1990; Barrie *et al.*, 1992). In *B. cereus*, *hbla*, *bceT* and *entFM* are the important genes, which are responsible for diarrhoeal enterotoxin production (Heinrichs *et al.*, 1993; Agata *et al.*, 1995; Asano *et al.*, 1997). Being a spore-former organism, it can survive heat-treated foods. Some strains of *B. cereus* are psychrotrophic in nature and thus can grow even in refrigerated food items (TeGiffel *et al.*, 1997). This organism is also capable of producing biofilm on different food-contact surfaces (Oosthuizen *et al.*, 2002) and the possibility of transmission from those surfaces to different food products can't be ruled out. According to Centers for Disease Control, United States of America, approximately 2% of the total food-borne illnesses in USA are due to enterotoxigenic *B. cereus* (Centers for Disease Control and Prevention, 1994). In European countries, food poisoning caused by enterotoxigenic *B. cereus* is very common and attributes to 47, 33 and 22% of the total food poisoning cases reported in Iceland, Norway and Finland, respectively (Granum and Lund, 1997). Although in India, no major outbreak due to this

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pathogen has been reported so far, enterotoxigenic *B. cereus* has been frequently isolated from different types of food items (Kamat *et al.*, 1989; Agarwal *et al.*, 1997; Das *et al.*, 2009). In one study carried out among diarrhoea patients in and around Kolkata, India, *B. cereus* was isolated from 3.5% of the stool samples of the patients (Banerjee *et al.*, 2011). As per the tropical seafood is concerned, it was observed that 24.91% of the samples collected from Cochin and surrounding area were positive to enterotoxigenic *B. cereus* (Das *et al.*, 2009). So, this is very much required to study the occurrence of this pathogen in the tropical fisheries environment. Keeping this in view, the present study was designed to assess the occurrence and molecular characterization of *B. cereus* from tropical fisheries environmental samples of Cochin and surrounding areas.

MATERIALS AND METHODS

Samples and Isolation of the Organism

A total of 51 samples collected during 2009–10 from fish markets and landing centres of Cochin and surrounding areas comprising of 30 ice, 18 mud and sand and 3 sea water were screened for the presence of *B. cereus*. Selective plating technique on Polymixin-pyruvate-egg yolk-mannitol-bromocresol purple agar (PEMPA) was done for isolation of *B. cereus* (Szabo *et al.*, 1984). Twenty five grams of sample was blended in 225 ml of sterile normal saline in a stomacher blender (Seward Ltd, U.K.) and serial 10 fold dilutions were made on normal saline. From each dilution, 0.5 ml was spread-plated on PEMP plates. After incubating at 37°C for 24 h, suspected *B. cereus* colonies showing lecithinase reaction were subcultured and subjected to identification using Gram's staining and standard biochemical reactions as mentioned in Bacteriological analytical manual of United States Food and Drug Administration (Rhodehamel and Harmon, 1998). The starch hydrolytic ability of the organisms was checked on nutrient agar containing 1% starch.

Screening of *B. cereus* Isolates for Haemolytic Activity and Production of Enterotoxin

The haemolytic activity of the suspected isolates was checked on spot inoculation on blood agar viz. Columbia agar base with 5% human blood. The blood agar plates were incubated at 37°C for 24 h. The ability to produce diarrhoeal enterotoxin was detected by Reverse passive latex agglutination (RPLA) test using BCET-RPLA kit (Oxoid) as per manufacturer's instruction.

Preparation of Crude PCR Template

Crude PCR template from the isolates was prepared by boiling method. Overnight grown culture of the isolate was centrifuged at $7000 \times g$ for 10 minutes at 4°C . The pellet was washed once with normal saline and suspended in autoclaved MilliQ distilled water. The suspended bacterial cells were lysed on boiling water bath for 10 min. Then it was immediately frozen at -20°C . After freezing, the lysate was thawed and centrifuged at $4000 \times g$ to pellet the debris. $5 \mu\text{l}$ of the supernatant was used as template in the PCR reaction.

Polymerase Chain Reaction

Total 4 sets of primers were used for this study. For *B. cereus* group specific PCR, BalF and BalR primers were used. All the isolates were screened for the presence of three virulent genes viz. *hbla*, *entFM* and *bceT* genes, which are generally responsible for enterotoxin production. The primer pairs HblA1/HblA2, BceT1/BceT2 and ENT A/ENT B were used to detect the presence of *hbla*, *bceT* and *entFM* genes, respectively. The details of primers including sequences have been depicted in Table 1. PCRs were carried out in $25 \mu\text{l}$ of reaction mixture, which consisted of $5 \mu\text{l}$ of template, 10 mM Tris-HCl (pH 9), 50 mM KCl, 1.5 mM MgCl_2 , 200 μM of each dNTP (Fermentas), 1 U *Taq* DNA polymerase (Fermentas) and 10 p mol of respective primers. For each PCR assay at the beginning of the reaction, the initial

Table 1: PCR Primers and Cycling Condition Details

Primer	Sequence	Target and Reference	PCR Cycling Condition
BalF BalR	5'- TGCAACTGTATTAGCACAAAGC T -3' 5'- TACCACGAAGTTTGTTCACTACT -3'	<i>Bacillus cereus</i> Group (Chang <i>et al.</i> , 2003)	94°C for 45 seconds (denaturation), 55°C for 45 seconds (annealing) and 72°C for 45 seconds (extension).
HblA1 HblA2	5'- CTAATGTAGTTTCACCTGTAGCAAC- 3' 5'- AATCATGCCACTGCGTGGACATATAA- 3'	<i>hbla</i> gene (Mantynen and Lindstrom, 1998)	94°C for 30 seconds, 58°C for 45 seconds and 72°C for 1 minute.
ENT A ENT B	5'- ATGAAAAAAGTAATTTCAGG- 3' 5'- TTAGTATGCTTTTGTGTAACC- 3'	<i>entFM</i> gene (Asano <i>et al.</i> , 1997)	94°C for 45 seconds, 52°C for 45 seconds and 72°C for 1 minute.
BceT1 BceT2	5'- GAATTCCTAACTTGCAACCATCTC G- 3' 5'- CTGCGTAATCGTGAATGTAGTCAAT- 3'	<i>bceT</i> gene (Mantynen and Lindstrom, 1998)	94°C for 45 seconds, $55/65^{\circ}\text{C}$ for 45 seconds and 72°C for 45 second

denaturation was done at 95°C for 5 min and at the end of the cycle, the final extension was carried out at 72°C for 5 min. The other details of PCR cycling condition for each PCR have been mentioned in Table 1. *bceT* gene specific PCR was run at two different annealing temperatures viz. 55 and 65°C. Following PCR, the amplified products were resolved in 1.5% agarose gel prepared in 0.5 × TBE buffer (45 mM Tris, 45 mM boric acid and 2 mM EDTA, pH 8.0) containing ethidium bromide (0.3 µg/ml). Electrophoresis was done at 7-volts/cm for 1 hour and the gel was photographed by an UV gel documentation system (Alpha Innotech Corporation, USA).

RESULTS AND DISCUSSION

Occurrence of *B. cereus* and Enterotoxin Production

Among the 51 samples screened, the presence of *B. cereus* was detected in 16 samples (4 ice and 12 mud and sand). Total 41 isolates have been recovered from the positive samples. All the isolates were haemolytic on blood agar and 38 isolates showed starch hydrolytic property. By Reverse passive latex agglutination (RPLA) test, 36 isolates were found to produce diarrhoeal enterotoxin. Those 36 isolates were sourced from 4 ice and 11 mud and sand samples. All the enterotoxin producing isolates were found to hydrolyze starch (Table 2).

Detection by *B. cereus* Group Specific PCR

In PCR using *B. cereus* group specific primers BalF/BalR, all the 41 isolates yielded 533 bp amplified product. Standard strains of *Bacillus subtilis* NCIM 2545 and *Staphylococcus aureus* NCIM 2079 were also employed in this PCR as negative controls, but none of them yielded any amplified product in this PCR.

Screening for Virulence Genes

The isolates were screened for the presence of three enterotoxin producing genes viz. *hbla*, *entFM* and *bceT* genes. *hbla* gene specific PCR using primers HbIA1/HbIA2 yielded 834 bp amplified product in case of 36 enterotoxigenic isolates. The isolates which were negative to enterotoxin production by RPLA method, didn't yield any amplified product in this PCR. In case of *entFM* gene specific PCR using ENT A/ENT B primers, 1300 bp amplified product was observed in case of 35 isolates. On the other hand, *bceT* gene specific PCR didn't yield any amplified product in any of the isolates tested. The results of each PCR, enterotoxin production by RPLA test and starch hydrolysis test for each isolate has been depicted in Table 2.

Table 2: Details of *B. cereus* Isolates

Isolates	Starch Hydrolysis	RPLA*	BCG**	Hemolysis	hbla PCR	entFM PCR	bceT PCR
BC-env-1	+	+	+	+	+	+	-
BC-env-2	+	+	+	+	+	+	-
BC-env-3	+	+	+	+	+	+	-
BC-env-4	+	+	+	+	+	-	-
BC-env-5	+	+	+	+	+	+	-
BC-env-6	+	+	+	+	+	+	-
BC-env-7	+	+	+	+	+	+	-
BC-env-8	+	+	+	+	+	+	-
BC-env-9	+	+	+	+	+	+	-
BC-env-10	+	-	+	+	-	+	-
BC-env-11	+	+	+	+	+	+	-
BC-env-12	+	+	+	+	+	+	-
BC-env-13	+	+	+	+	+	+	-
BC-env-14	+	+	+	+	+	+	-
BC-env-15	+	+	+	+	+	+	-
BC-env-16	+	+	+	+	+	+	-
BC-env-17	-	-	+	+	-	-	-
BC-env-18	+	+	+	+	+	+	-
BC-env-19	+	+	+	+	+	+	-
BC-env-20	+	+	+	+	+	+	-
BC-env-21	+	+	+	+	+	+	-
BC-env-22	+	+	+	+	+	+	-
BC-env-23	+	+	+	+	+	+	-
BC-env-24	+	+	+	+	+	+	-
BC-env-25	+	+	+	+	+	+	-
BC-env-26	+	+	+	+	+	-	-
BC-env-27	+	+	+	+	+	-	-
BC-env-28	+	+	+	+	+	+	-
BC-env-29	+	+	+	+	+	+	-
BC-env-30	-	-	+	+	-	+	-
BC-env-31	-	-	+	+	-	+	-
BC-env-32	+	+	+	+	+	+	-
BC-env-33	+	+	+	+	+	+	-
BC-env-34	+	+	+	+	+	+	-
BC-env-35	+	+	+	+	+	+	-
BC-env-36	+	+	+	+	+	+	-
BC-env-37	+	+	+	+	+	+	-
BC-env-38	+	+	+	+	+	+	-
BC-env-39	+	+	+	+	+	+	-
BC-env-40	+	-	+	+	-	-	-
BC-env-41	+	+	+	+	+	-	-

* BCG = *Bacillus cereus* group specific PCR

** RPLA = Enterotoxin production by RPLA test

The aim of the this study was to study the occurrence of enterotoxigenic *B. cereus* in fishery environmental samples collected from fish markets and landing centers and also to study the presence of different virulence genes in the isolates of *B. cereus*. In this study, 31.37% (16 out of 51) of the samples were found to possess *B. cereus* and enterotoxigenic *B. cereus* was isolated from 15 samples (Occurrence rate 29.41%). The occurrence of enterotoxigenic *B. cereus* was found 13.33% (4 out of 30) and 61.11% (11 out of 18) in case of ice and mud/sand, respectively. On other hand, all the three sea water samples were found negative for this organism. The comparatively high occurrence was found in case of mud and sand collected from the fishery environment. In previous study, it was observed that 29.41% of fishes of tropical area of Cochin possessed enterotoxigenic *B. cereus* (Das *et al.*, 2009). As per this study, it can be stated that occurrence of enterotoxigenic *B. cereus* is moderately high in tropical fisheries environment and the fisheries environment may be possible source of transmission of *B. cereus* to fish.

On screening for virulence genes, it was observed that all the enterotoxigenic isolates possessed *hbla* gene while this gene was absent in all non-enterotoxigenic isolates (Table 2). Similar type of finding was reported by Das *et al.* (2009) in case of *B. cereus* isolates from fish. *entFM* gene was detected in three non-enterotoxigenic isolates, while this gene was not detected in four enterotoxigenic isolates (Table 2). So, it can be stated that no relationship can be established between presence of *entFM* genes and enterotoxin production. *bceT* gene was not detected in any of the isolates. So, this gene can be considered rare in case of tropical fishery environmental isolates of Cochin. So based on this study, it can be concluded that *hbla* gene can be considered as marker for diarrhoeal enterotoxin production in case of *B. cereus* isolates from tropical fishery environment.

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