

Occurrence of *vibrio* spp. in marked shellfish

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Abstract

Seventy- five samples of the common marketed shellfish Shrimp, Crab, Oyster, lobster and squilla mantis (15 of each) were collected from the fish market in Sharkia Governorate, Egypt. All shellfish samples were subjected to bacteriological examination for isolation and identification. The overall prevalence of vibrio in examined shellfish was 72%. The identified species *Vibrio. fluvialis* were detected in 1(7%), 4(33%) and 3(25%) of examined crab, squilla mantis and oyster, respectively. *Vibrio mimicus* was detected in 2(13%), 2 (17%) and 3(50%) of examined crab, mantis and lobster, respectively. *Vibrio alginolyticus* was detected in 3(20%), 3 (33%), 4(33%) and 3(25%) in examined crab, shrimp, Oyster and squilla mantis, respectively. *Vibrio vulnificus* was detected in 3(20%), 1(11%) and 3 (25%) of examined crab, shrimp and oyster, respectively. *Vibrio cholera* was detected in 3(20%), 2(22%), 1(8%) and 1 (17%) of examined crab, shrimp, Oyster and Lobster, respectively. *Vibrio parahaemolyticus* was detected in 3(20%), 3 (33%), 2(17%), 2(17%) and 2 (33%) of examined crab, shrimp, Oyster, squilla mantis and Lobster, respectively.

1- Introduction

shellfish considered as important food element for broad section of world population and considered the main protein of animal source. Recently, more people exchange red meat to fish (Adebayo-Tayo *et al.*, 2012).

shellfish is a main dietary Supply of proteins of rich biological value, vitamins, minerals and omega-3 which are essential for optimum neural development, and is low in saturated fatty acid. They are very healthy foods.(National Public Health Institute of Finland, 2004)

Although Seafood is a nutritious food that constitutes one of the desirable components of a healthy diet. One of the main risks involves the consumption of uncooked or improperly cooked seafood that may be naturally contaminated by foodborne pathogens present in the marine environment. Such risk is further increased if the food is

mishandled during processing where pathogens could multiply exponentially under favorable conditions.(Food and Environment Hygiene Department, 2005).

The main microorganism identified in foodborne diseases of marine shellfishes are *Vibrio anguillarum*, *V. alginolyticus*, *V. fluvialis*, *V. furnissii*, *V. harveyi*, *V. parahaemolyticus* and *V. Vulnificus* as reported by (Buck 1990 and Liao *et al.*, 2004).

The public health hazard of consumption of raw, inadequately cooked or insufficiently processed fish, and fish products include nausea, vomiting, diarrhea, abdominal cramps, headache as well as muscle and joint pain. Good manufacturing practices should be in place to minimize the risk of food poisoning associated with the consumption of fish and seafood products.(Pal *et al.*, 2016)

This study was aimed to investigate the prevalence and distribution vibrio spp. in different marked shellfish.

2- Materials and Methods

2.1. Collection of shellfish samples

A total of 75 random samples of shellfish (Shrimp, Crab, Oyster, lobster, squilla mantis) 15 of each were collected from Sharkia governorate fish markets and from different retail seafood stores during winter of 2018. The collected samples were packaged separately and transferred with a minimum of delay to the laboratory in a sterile ice box. All samples were subjected to bacteriological examination for enumeration, isolation and identification.

2.2. Preparation of samples:

The samples were rapidly washed with tap water. Then the surfaces were swabbed with 70% ethyl alcohol for surface disinfection. The shells were removed. Then the muscles were removed with a sterile forceps and a scalpel and placed into sterile Petri dishes. Five grams were taken under aseptic conditions to sterile homogenizer containing 45ml of sterile alkaline peptone water (3% NaCl and pH 8).

Screening of Vibrio spp.:

It was done according to **FDA (2004)** as follows:

Enrichment broth:

The previous prepared tubes were incubated overnight at 37°C.

Isolation:

Loopfuls from each previous cultured tube were separately streaked onto Thiosulfate citrate bile and sucrose agar (TCBS) then incubated at 37°C for 24hrs.

2.3. Identification of total Vibrio specie:

The suspected isolates were further identified according to the protocol

recommended by **ISO/ TS 21872-1 (2007)** and **ISO/ TS 21872-2 (2007)**.

1. Morphological examination:

Films were prepared from a pure culture of the isolated microorganism stained with Gram's stain and then examined microscopically. Vibrio species were short, rigid, Gram -ve, single flagellum and comma - shape.

2. Biochemical identification:

2.1. Oxidase test:

Oxidase test was done by streaking of the pure culture onto filter paper moistened with oxidase reagent. The test is positive if the colour turns to mauve, violet or deep purple within 10 seconds. All vibrio species are oxidase positive.

2.2. TSI agar test:

The agar slope was inoculated by stabbing to the bottom of the agar butt and streaking longitudinally along the slope. Then it was incubated at 37°C for 24hrs.

2.3. Detection of Ornithine decarboxylase (ODC):

Suspected colonies were inoculated into ornithine decarboxylase medium just below the surface. One ml of sterile mineral oil was added to the top of the medium and incubated at 37°C for 24 hrs. Turbidity and violet color after incubation indicate a positive ODC.

2.4. Detection of L- lysine decarboxylase (LDC):

Suspected colonies were inoculated into L- lysine decarboxylase medium just below the surface. One ml of sterile mineral oil was added to the top of the medium and incubated at 37°C for 24 hrs. Turbidity and violet color after incubation indicate a positive LDC.

2.5. Detection of Arginine decarboxylase (ADH):

Suspected colonies were inoculated into arginine decarboxylase medium just below the surface. One ml of sterile mineral oil was added to the top of the medium and incubated at 37°C for 24 hrs. Turbidity and violet color after incubation indicate a positive ADH.

2.6. Detection of β -galactosidase (ONPG):

The suspect colonies were inoculated into a sterile tube containing 2.5% NaCl solution and mixed. Drop of toluene was added and the tube was shaken. The tube was placed in the water bath adjusted at 37°C and left to stand for approximately 5 min. About 0.25 ml of the reagent was added for detection of β -galactosidase (2-ortho-Nitrophenyl- β -D-galactopyranoside) "ONPG" and mixed. The tube was replaced in the water bath set at 37°C and left to stand for 24 hrs then examined from time to time. A yellow color indicates positive β -galactosidase.

2.7. Indole test:

A tube containing 5 ml of the tryptone-tryptophan saline medium was inoculated with the suspected colony; Incubated at 37°C for 24 hrs, One ml of the Kovacs' reagent was added. Formation of red ring indicates a positive production of indol.

2.8. Halotolerance test:

Series of sterile peptone water was produced with increasing salt (NaCl) concentration: 0 %, 2 %, 4 %, 6 %, 8 % and 10 %. Suspension with the suspected colony was prepared to be identified and each of the tubes with loopful was lightly inoculated and incubated at 37°C for 24 hrs. The turbidity indicates that the suspected colony could grow at a concentration of

sodium chloride present in the tube of sterile saline peptone water.

Polymerase Chain Reaction (PCR)

1. Materials used for PCR:

1.1. Reagents used for agarose gel electrophoresis:

1.1.1. Agarose powder, Biotechnology grade (Bioshop^R, Canda inc. lot No: OE16323).

It prepared in concentration 2% in 1× TAE buffer.

1.1.2. Tris acetate EDTA (TAE) electrophoresis buffer (50×liquid concentration) (Bioshop^R, Canda inc. lot No: 9E11854).

The solution diluted to 1× by adding 1 ml stock solution to 49 ml double dist. Water to be used in the preparation of the gel or as a running buffer.

1.1.3. Ethidium bromide solution (stock solution) biotechnology grade (Bioshop[®] Canda Inc, Lot No: 0A14667):

The stock solution was diluted by 25 μ l /200ml double distilled water and stored covered at 4°C. It was used for staining of PCR products that electrophoreses on agarose gel to be visualized by UV light.

1.2. Gel loading buffer (6×stock solution) (Fermentas, lot No: 00056239).

The components were dissolved in sterile double distilled water and stored covered with aluminum foil at room temperature.

1.3. DNA ladder (molecular marker): 100 bp (Fermentas, lot No: 00052518).

1.4. 5X Taq master (Fermentas):

Containing polymerase enzyme, Magnesium chloride (Mg Cl₂), Deoxy nucleotide triphosphate (dNTP) and PCR grade water.

1.5. Primer sequences of *V. parahaemolyticus* for virulence factors:

Molecular characterization of virulence factors including **thermostable direct hemolysin (tdh) and thermolabile hemolysin (tlh) genes of *V. parahaemolyticus*.**

2. DNA Extraction using QIA amp kit (Shah et al., 2009):

After overnight culture on nutrient agar plates, one or two colonies were suspended in 20 ml of sterile distilled water, and the suspension was then heated at 100°C for 20 minutes. Accurately, 50-200 µl of the culture were placed in Eppendorf tube and the manufacturer's instructions (Qiagen, Hilden, Germany) were applied according to the following steps:

1. Equal volume from the lysate (50-200 µl) was added, addition of 20-50µl of proteinase K, then incubation at 56 °C for 20-30 min. After incubation, 200 µl of 100% ethanol was added to the lysate.

2. The solution was added to the column and centrifuged at 8000 rpm for 1 min. then the filtrate was discarded.

3. The sediment was washed using AW1 buffer (200 µl), the column was centrifuged at 8000 rpm / 1 min, and the filtrate was discarded.

4. Washing was applied by using the AW2 buffer (200µl), the column was centrifuged at 8000 rpm / 1 min. and the filtrate was discarded.

5. The column was placed in a new clean tube and 25-50 µl from the elution buffer was added, centrifuged at 8000 rpm/1min.

6. The column was discarded. The filtrate was put in clean tube containing the pure genomic DNA. Nucleic acid was eluted with 100 µl of elution buffer provided in the kit.

3. DNA amplification:

3.1. Amplification of *V. parahaemolyticus* virulent genes (Khalil et al., 2014):

The amplification was performed on a Thermal Cycler (Master cycler, Eppendorf, Hamburg, Germany). The final volume of the reaction mixture was adjusted to 50 µL with sterile deionised distilled water. Amplification of DNA segment was performed) with the following temperature cycling parameters; initial denaturation at 94 °C for 2 min followed by 30 cycles of denaturation at 94 °C for 30 sec, primer annealing at 57°C for 30 sec, primer extension at 72 °C for 1 min and a final extension at 72 °C for 2 min. Ten µL of each amplified

product was loaded in 1.5% agarose gel in 1X Tris-boric acid-EDTA buffer [TBE: 0.089M Trisbase, 0.089M boric acid and 0.002 M EDTA (pH 8.0)] at 100 volts for 40 minutes. After electrophoresis, amplification products were captured and visualized on UV trans illuminator. A 100 bp plus DNA Ladder (Qiagen, Germany, GmbH) was used to determine the fragment sizes.

3- Results and Discussion

From the results clarified in Table (1) , the incidence of vibrio species in table (4) and figure (4) were 100% ,60% ,80%,80%,40% in the studied crab, shrimp, oyster, mantis and lobster, respectively The average percentage was 72% overall examined shellfish.

These results nearly agreed with those reported by **Waffa et al.(2011)**,who found that the total rate of isolation of *Vibrio* spp. isolated from shellfish was 88%. The low incidences of *Vibrio* in shellfish obtained by **Ripabelli et al. (1999)** they stated that *Vibrio* spp.

were isolated from shellfish with percentage of 48.4%.

The shellfish Contaminated by *Vibrio* spp because it is naturally present in aquatic environments. It is the primarily marine bacterial pathogens associated with illness. It is responsible for the third-highest number of shellfish-related illnesses, after non cholera *Vibrio* spp. and

Norwalk viruses. Toxigenic O₁ (epidemic biotype) of *Vibrio* spp. infections are associated with profuse, watery diarrhea whereas nontoxigenic, non-O₁ biotype (except O139) infections result in septicaemia and mild gastroenteritis (**Wittman and Flick,1995 ; Gnanambal and Patterson, 2005; DePaola et al.,2010 ; CDC,2010**).

Table 1:): Incidence of vibrio species in examined shellfish samples (No = 15 for each)

Type	Crab	Shrimp	S.mantis	Oyster	Lobster	Total
No. of positive sample	15	9	12	12	6	54
%	100	60	80	80	40	72

In the present study, The obtained results in table (2) and figure (1) showed that *Vibrio. fluvialis* were detected in 1(7%), 4(33%) and 3(25%) of examined crab, squilla mantis and oyster, respectively . The incidence of *V. fluvialis* was recorded by **Yücel and Balci (2010)** in blue crabs (>10%).

Vibrio mimicus was detected in 2(13%), 2 (17%) and 3(50%) of examined crab, mantis and lobster, respectively. A lower prevalence of finding obtained by **Wafaa et al. (2011)** who found 11% in some seafood.

Vibrio alginolyticus was detected in 3(20%), 3 (33%), 4(33%) and 3(25%) in examined crab, shrimp, Oyster and squilla mantis, respectively . A higher prevalence of *V. alginolyticus* had been

reported by **Gomathi et al. (2013)** who found 80% in sea food.

Vibrio vulnificus was detected in 3(20%), 1(11%) and 3 (25%) of examined crab, shrimp and oyster, respectively. Nearly similar finding obtained by **El Far et al. (2015)** who found 16.7% in Indian prawn. A higher prevalence of *V. vulnificus* was recorded by **Ripabelli et al. (1999)** in shellfish (32%).

Vibrio cholera was detected in 3(20%), 2(22%), 1(8%) and 1 (17%) of examined crab, shrimp, Oyster and Lobster, respectively.

Vibrio were the most commonly reported cause of seafood-associated outbreaks where toxigenic *V. cholerae* caused 3 outbreaks and 10 illnesses without deaths and non-O₁,

non-O139 *V. cholerae* caused 4 deaths Iwamoto et al.,(2010).
outbreaks and 12 illnesses without

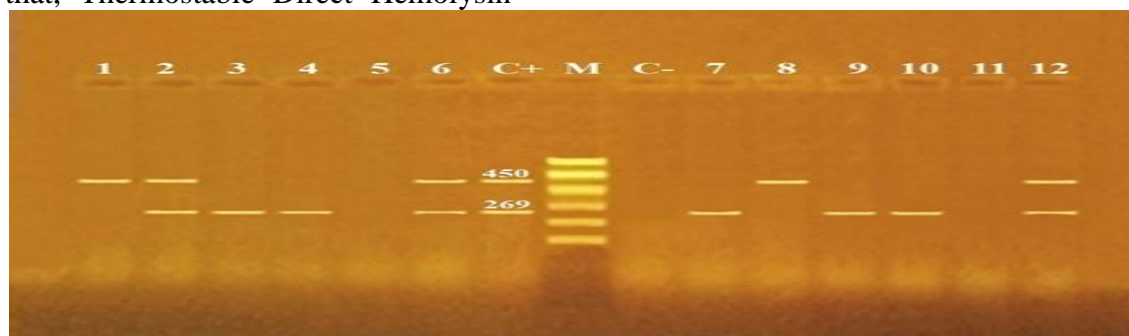
Table 2. Frequency distribution of isolated vibrio organisms recovered from examined shellfish sample:

Isolate	Crab		Shrimp		S.mantis		Oyster		Lobster	
	No.	%	No.	%	No.	%	No.	%	No.	%
<i>V.cholera</i>	3	20	2	22	0	0	1	8	1	17
<i>V.alginolyticus</i>	3	20	3	33	4	33	3	25	0	0
<i>V.fluvialis</i>	1	7	0	0	4	33	3	25	0	0
<i>V.parahaemolyticus</i>	3	20	3	33	2	17	2	17	2	33
<i>V.vulnificus</i>	3	20	1	11	0	0	3	25	0	0
<i>V.mimicus</i>	2	13	0	0	2	17	0	0	3	50

In the present study in photograph (1) showed that: Lane M:100 bp ladder as molecular size DNA marker. Lane C+: Control positive *V. parahaemolyticus* for *tdh* and *tlh* genes. Lane C-: Control negative. Lanes 3, 4, 7, 9 & 10: Positive *V. parahaemolyticus* strains for *tdh* gene. Lanes 1 & 8: Positive *V. parahaemolyticus* strains for *tlh* gene. Lanes 2, 6 & 12: Positive *V. parahaemolyticus* strains for *tdh* and *tlh* genes. Lanes 5 & 11: Negative *V. parahaemolyticus* strains for *tdh* and *tlh* genes. The obtained results of agarose gel electrophoresis virulence genes showed Positive *V. parahaemolyticus* strains for *tdh* gene and *tlh* gene. Our results supported the results obtained by Cook et al.(2002) and Okuda et. al.(1997a) who stated that, Thermostable Direct Hemolysin

(TDH) was the first recognized virulence factor for *V. parahaemolyticus* and has been used as an important marker for identifying virulent strains. In addition, The TDH encoding gene (*tdh*) is able to induce intestinal chloride secretion resulting secretory diarrhea (Raimondi et. al.,1995).

Similary, Bej et. al.,(1999) reported that out of 111 isolates of *V. parahaemolyticus* 60 (54%), showed amplification of *tdh* gene by using similar primer pair study which yielded desired amplified product of approximately 269 bp similar to the reference strain of *V. parahemolyticus* using the primer pair for *tdh* gene. Therefore proper chilling and handling of seafood's of marine water origin is essential.



Photograph (1): Agarose gel electrophoresis of multiplex PCR of *tdh* (269 bp) and *tlh* (450 bp) virulence genes for characterization of *V. parahaemolyticus*.

Lane M: 100 bp ladder as molecular size DNA marker. **Lane C+:** Control positive *V. parahaemolyticus* for *tdh* and *tlh* genes. **Lane C-:** Control negative. **Lanes 3, 4, 7, 9 & 10:** Positive *V. parahaemolyticus* strains for *tdh* gene. **Lanes 1 & 8:** Positive *V. parahaemolyticus* strains for *tlh* gene. **Lanes 2, 6 & 12:** Positive *V. parahaemolyticus* strains for *tdh* and *tlh* genes. **Lanes 5 & 11:** Negative *V. parahaemolyticus* strains for *tdh* and *tlh* genes.

4- Conclusion

It was concluded that examined shellfish samples were contaminated with *Vibrio* spp. which considered one of the main food poisoning microorganisms.

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مدى تواجد ميكروب الفيبريو في قشريات الاسماك المسوقه.

اسراء مصطفى عطيه زغل

قسم الأمراض المشتركة - كلية الطب البيطرى - جامعة الزقازيق- الزقازيق- 44511-مصر

الملخص العربى

تم تجميع عدد 75 عينة عشوائية من قشريات الاسماك المسوقه في محافظة الشرقية بواقع 15 عينة لكلا من (الجمبري ، الكابوريا ، الجاندوفلي ، الشيكال ، الاستاكوزا) خلال شتاء 2018 وتم ارسال هذه العينات تحت ظروف صحيه الي المعمل لفحصها بكتيريولوجيا وقد دلت النتائج علي الاتي :
نسبه تواجد ميكروب الفيبريو كان 100 % فى الكابوريا و 60% فى الجمبري و 80% فى الشيكال و 80 % فى الجندوفلي و 40% فى الاستاكوزا.

وتم تصنيف ميكروب الفيبريو فلافياس بنسب (7% ، 33% ، 25%) فى الكابوريا والشيكال الجندوفلي على التوالي و فيبريو يميكس بنسب (13% ، 17% ، 50%) فى الكابوريا والشيكال والاستاكوزا على التوالي وفيبريو الجينوليتيكس بنسب (20% ، 33% ، 33% ، 25%) فى الكابوريا والجمبري والجندوفلي والشيكال على التوالي وفيبريو فالينيفيكس بنسب (20% ، 11% ، 25%) فى الكابوريا والجمبري و الجندوفلي على التوالي وفيبريو كوليرا بنسب (20% ، 22% ، 8% ، 17%) فى الكابوريا والجمبري والجندوفلي والاستاكوزا. الفيبريو باراهيموليتيكس بنسب (20% ، 33% ، 17% ، 17% ، 33%) فى الكابوريا والجمبري والشيكال و الجندوفلي والاستاكوزا على التوالي.



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