





Pathogenic Vibrios in shellfish

1 November 2007 **By George J. Flick, Jr., Ph.D.**

Understand risks associated with consuming raw seafood



Studies of mussels in Italy and elsewhere have found various *Vibrio* species, including several potentially pathogenic strains.

The microflora in the estuarine and marine environments include various members of the family Vibrionaceae, some of which are pathogenic to humans and constitute a potential health threat for consumers of raw or partially cooked mussels.

It is essential to understand the distribution of bacterial species – especially opportunistic pathogens – in the marine environment in order to predict and prevent bacterial gastroenteritis transmitted by seafood. Although improvements in postharvest treatments such as depuration, high-pressure processing, irradiation, and thermal pasteurization reduce the potential health hazard, it is still important for compromised individuals to understand the risks associated with consuming raw seafood.

Vibrio research

The vibrios constitute a considerable part of marine halophilic bacterial populations, which require high concentrations of salt for growth and performance. In the past 20 years, research on medical, environmental, and taxonomic aspects of *Vibrio* species has expanded exponentially.

Environmental research has indicated that ecological parameters such as nutrient availability, temperature, and salinity influence the presence and persistence of different *Vibrio* species in the sea. Vibrios are found less frequently when the temperature drops and salinity increases. However, temperature, dissolved oxygen, and salinity are interactive, so the effects of a single environmental parameter are difficult to assess.

More than 35 species of *Vibrio* microorganisms have been identified, and more species are being added as new scientific information becomes available. In addition to *V. cholerae*, the most widely known pathogenic vibrio, a number of *Vibrio* species can cause disease.

Vibrio parahaemolyticus, V. minicus, and *V. vulnificus* are food-poisoning bacteria frequently isolated from seawater and shellfish. Among *halophilic vibrios, V. alginolyticus, V. fluvialis,* and *V. metschnikovii* are also pathogenic to humans, while *V. anguillarum* is a pathogen that affects fish and other aquatic animals.

Many human-pathogenic species have distinct clinical features, pathogenic mechanisms, epidemiological characteristics, and ecological positions. However, in recent years, other microorganisms now classified as *Aeromonas* and *Plesiomonas* have often been responsible for enterotoxic episodes.

Vibrios and shellfish

V. parahaemolyticus has been reported responsible for one-fourth of all gastrointestinal pathologies caused by food. In the United States, it caused 14 food-poisoning outbreaks between 1971 and 1978. Though *Vibrio vulnificus* does not cause outbreaks, it is responsible for very severe syndromes that are fatal in 50-60 percent of cases.

In 1998, shellfish were incriminated in 7.1 percent of the 84 reported foodborne disease outbreaks in Italy. European Directive 492 per 91 prescribed that bivalve mollusks marketed for human consumption should be raised in waters that meet certain microbiological requirements defined as Type A or, if

originating elsewhere, should be run through a depuration procedure.

In order for bivalve mollusks to be deemed fit for human consumption, legislation requires that they contain less than 300 fecal coliforms (or less than 230 *Escherichia coli*) per 100 grams of shellfish meat and intervalve water, and no *Salmonella* species in 25 g of meat. In the United States, the fecal coliform median (or geometric mean MPN) of the water sample results shall not exceed 14 per 100 millimeters, and the estimated 90th percentile shall not exceed 43 MPN per 100 millimeters for a five-tube decimal dilution test or 49 MPN per 100 millimeters for a three-tube decimal dilution test.

However, it has been noted repeatedly that in these products, there is no close correlation between the presence of microorganisms of fecal origin and the presence of vibrios that are potentially pathogenic for humans.

In coastal environments, a direct relationship exists between the consumption of raw or partially cooked shellfish and the occurrence of episodes of human intestinal and extraintestinal infections attributed to some *Vibrio* species. Mollusks are theoretically an excellent transmission vehicle because they are filter-feeding organisms and can concentrate bacteria.

It has been estimated that one in 2,000 meals of raw shellfish results in disease, giving these shellfish a reputation as one of the most hazardous foods. The most-studied mollusk species are the Pacific oyster, (*Crassostrea gigas*), and Eastern oyster, (*C. virginica*).

Italy studies

In studies in Italy, samples of (*Mytilus galloprovincialis*) mussels were collected in central Italy from approved coastal sites located on the Adriatic Sea to be examined for the presence of *Vibrio parahaemolyticus* and the occurrence of pathogenic strains. Thirty-five *V. parahaemolyticus* strains were isolated out of 144 samples, and four of the isolates were considered pathogenic after performing a genetic analysis.

Another Italian study of the same mussel species obtained from retail stores in the Puglia region revealed that *V. parahaemolyticus* and *V. vulnificus* were found in 7.83 percent and 2.83 percent, respectively, of 600 samples. There were no significant differences among vibrios present in the mussels during the three-year survey.

A third study of mussel farms in Italy showed that a total of 125 vibrios from 152 isolates were identified as *Vibrio fluvialis* (55 strains), *V. alginolyticus* (40 strains), *V. parahaemolyticus* (11 strains), and *V. mimicus* (nine strains). The remaining 27 isolates were not identified.

A fourth study of mussels and water collected in 30 sampling sites in the Ionian Sea revealed that *V. alginolyticus* constituted the predominate component of the total culturable vibrios. Some *Vibrio* species such as *V. mediterranei, V. parahaemolyticus, V. diaztrophicus, V. nereis*, and *V. splendidus* were present in both the water and the mussels. Selective retention in mussels, however, was demonstrated for *V. vulnificus, V. cincinnatiensis, V. orientalis, V. anguillarum, V. marinus*, and *V. hollisae*.

A fifth study, performed over a two-year period in which 726 bacterial strains were isolated, showed that 46.9 percent belonged to the *Vibrio* genus, 29.0 percent was of the *Aeromonas* genus, and the remaining 23.3 percent was made up of the *Pseudomonas, Flavobacterium, Pasteurella, Agrobacterium*, and *Ochrobacterium* genera.

Norwegian study

A Norwegian study of 885 samples of blue mussels, (*Mytilus edulis*), showed that *V. parahaemolyticus*, *V. cholerae*, and *V. vulnificus* were isolated at 10.3 percent, 1.0 percent, and 0.1 percent, respectively. Four of the *V. parahaemolyticus* samples contained one of the major virulent genes.

Netherland studies

A study of *Campylobacter* contamination of unspecified mussels species in the Netherlands revealed that over 90 percent of the contamination was *C. lari*. Other *Campylobacter* species were *C. jejuni*, *C. coli*, *C. upsaliensis*, and *C. hyointestinalis*.

The study was initiated because *Campylobacter* is a common bacterial pathogen that causes enteritis in humans worldwide. Specifically, *Campylobacter jejuni* and *C. coli* account for the majority of enteric infections in humans. *Campylobacter lari* appears to be widely present in the environment, but is rarely reported as a human pathogen. During recent years, a very limited number of cases of *C. lari* have been described.

(Editor's Note: This article was originally published in the November/December 2007 print edition of the Global Aquaculture Advocate.)

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