Characterization of *Vibrio damsela* strains isolated from turbot *Scophthalmus maximus* in Spain

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ABSTRACT: During a 3 yr period several epizootics of vibriosis in turbot *Scophthalmus maximus* occurred in different marine farms located in northwestern Spain. Affected fish showed extensive haemorragic areas around the anus, eyes and mouth, as well as a characteristic accumulation of mucus and reddish fluid in the peritoneal cavity. Strains of *Vibrio damsela* were isolated from diseased fish, representing, to our knowledge, the first report of this bacterium causing problems in cultured turbot. Isolates were biochemically identical to collection strains isolated from both fish and humans, and exhibited the same pattern of drug-sensitivity. However, antigenic differences occurred among the strains, 4 distinct groups of *V. damsela* being recognized. These serological results were supported by lipopolysaccharide profiles and the outer-membrane protein patterns exhibited by the bacterial isolates. In addition, a high molecular weight plasmid band (approx. 90 to 100 MDa) was detected in all the turbot isolates and in some of the reference strains. Virulence tests showed that the *V. damsela* isolates were pathogenic for turbot and rainbow trout *Oncorhynchus mykiss*, the LD₅₀ ranging from 1×10^3 to 5×10^5 *V. damsela* cells per fish.

INTRODUCTION

The coast of Galicia (northwestern Spain) is an important location for the culture of turbot *Scophthalmus maximus*, a fish of high commercial value. One of the most important limiting factors in saltwater fish culture is the occurrence of disease outbreaks caused by *Vibrio* species. Although to date only *Vibrio anguillarum* (serotypes O1 and O2) (Devesa et al. 1985, Toranzo & Barja 1990) has been reported as the primary causative organism of turbot vibriosis, there are other members of the genus *Vibrio* in the estuarine environment that have been implicated in disease problems in mariculture (Lupiani et al. 1989, Toranzo et al. 1990).

Some of the marine vibrios associated with fish infections such as *Vibrio vulnificus* and *V. damsela* are also considered human pathogens. Within *V. vulnificus*, biotypes I and II have been associated with human infections and fish diseases, respectively (Tison et al. 1982). However, in *V. damsela* pathogenicity for fish still remains to be firmly established. *Vibrio damsela* is a common marine bacterium associated with wound infections in humans, but only a few reports have documented its isolation from diseased fish, e.g. damselfish, brown shark, lemon shark, yellowtail and seabream. This species has also been isolated from turtle, dolphin, octopus and uninfected fish (Table 1).

We have recently described the isolation of *Vibrio* damsela from turbot in different marine farms located in north western Spain. The isolates represented the first documentation of vibriosis in turbot caused by *V.* damsela (Fouz et al. 1991).

In this report the biochemical, serological, and virulence characteristics of isolates of *Vibrio damsela* are compared with those of reference strains. In addition, plasmid content and membrane protein patterns of *V. damsela* isolates were also studied.

MATERIAL AND METHODS

Development of the disease. A new infectious disease was observed in 2 turbot farms located in the Ría

| Species | Country | Source |
|------------------------|-------------|--------------------------|
| Fishes | | |
| Blacksmith | | |
| Chromis punctipinnis | USA | Love et al. (1981) |
| Brown shark | | |
| Carcharhinus plumbeus | USA | Grimes et al. (1984) |
| Undescribed seafish | Senegal | Schandevyl et al. (1984) |
| Lemon shark | | |
| Negaprion brevirostris | USA | Grimes et al. (1985) |
| Yellow tail | | |
| Seriola quingueradiata | Japan | Sakata et al. (1989) |
| Seabream | | |
| Sparus aurata | Spain | Vera et al. (1991) |
| Molluscs | | |
| Octopus | | |
| Octopus joubini | USA | Hanlon et al. (1984) |
| Reptiles | | |
| Turtle | | |
| Dermochelys coriacea | Australia | Obendorf et al. (1987) |
| Mammals | | |
| Dolphin | | |
| Tursiopp truncatus | Hawaii, USA | Fujioka et al. (1988) |
| Humans | | |
| Wound pathogen | USA | Love et al. (1981) |
| * 5 | | Morris et al. (1982) |
| Fatal wound infection | USA | Clarridge et al. (1985) |
| Necrotizing infection | USA | Coffey et al. (1986) |

Table 1. Vibrio damsela isolations reported to date in different countries

de Arosa (northwestern Spain). In the first one (Farm A) the problem was detected several times between summer 1987 and summer 1989, but only once in the second fish farm (Farm B) (Table 2). Size of the diseased fish ranged from 300 to 1500 g.

No abnormal swimming behaviour was observed in the diseased fish. The first deaths occurred when the water temperature increased suddenly from 18 °C to 22-24 °C. The mortality was low but continuous during the course of the outbreaks. Although the cumulative fish losses were lower than 5 % of the stock in each epizootic, the disease proved costly because of the size of affected fish.

The most remarkable clinical signs in moribund turbot were abdominal distension and haemorragic areas, especially in the eyes and mouth (including palate and jaws) and around the anus (Fig. 1). These external signs were quite similar to enteric redmouth disease of salmonids produced by *Yersinia ruckeri*. Internally, an accumulation of mucus and reddish fluid in the peritoneal cavity was observed and, in some cases, the liver was pale and with petechiae.

Bacterial isolation and identification. Samples were taken from the liver, kidney and muscle around the

haemorragic eyes of moribund turbot and cultured on tryptic soy agar (Difco) supplemented with 1 % NaCl (TSA-1) and on thiosulphate citrate bile sucrose (TCBS) agar (Oxoid) for bacterial isolation. Pure cultures of the

Table 2. Vibrio damsela. Origin of the strains used in this study. A and B are 2 turbot farms where the infectious disease occurred

| Strain | Host | Farm | Year |
|-------------------|-------------|------|------|
| Spanish isolates | | | |
| RG-91 | Turbot | А | 1987 |
| RG-151 | Turbot | А | 1988 |
| RG-153 | Turbot | А | 1988 |
| RG-191 | Turbot | А | 1988 |
| RG-192 | Turbot | А | 1988 |
| RG-193 | Turbot | А | 1988 |
| RG-214 | Turbot | А | 1989 |
| RM-71 | Turbot | В | 1988 |
| Reference strain: | 5 | | |
| ATCC 33539 | Damselfish | | 1981 |
| ATCC 35083 | Brown shark | | 1984 |
| CDC-2227-81 | Humans | | 1981 |
| CDC-1421-81 | Humans | | 1981 |

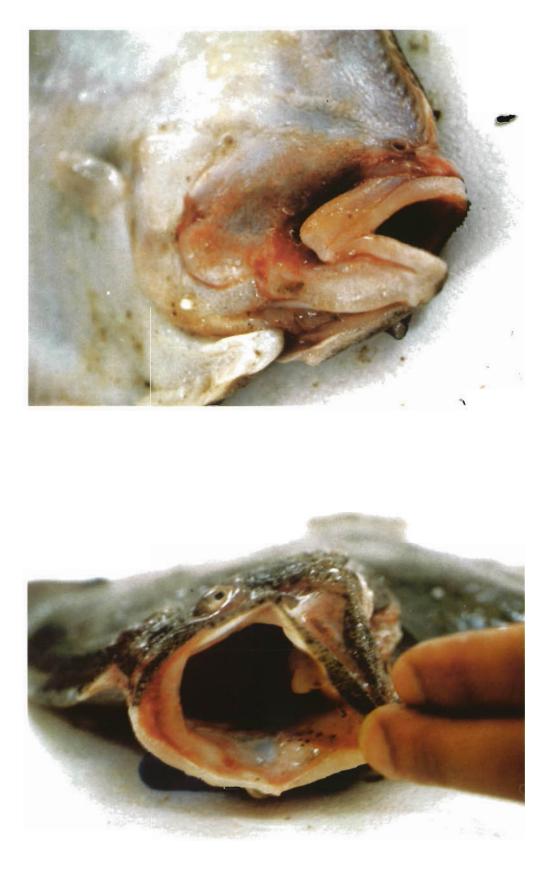


Fig. 1. Scophthalmus maximus. A diseased fish with haemorragic areas around the mouth as well as in the jaws and palate

isolated bacteria were identified using standard morphological, physiological and biochemical plate and tube tests (West & Colwell 1984, Fouz et al. 1990). Results were recorded after incubation at $22 \,^{\circ}$ C for 7 d.

Gram stain, oxidase test, morphology and motility, fermentation of glucose, and sensitivity to the vibriostatic agent 0/129 were used to identify the isolates as members of the genus *Vibrio*. Biochemical profiles of our isolates were compared with those of virulent strains of *V. anguillarum* isolated from turbot in Spain and with those of reference strains of *V. damsela* isolated from damselfish, brown shark and humans (Table 2).

Representative strains isolated from Farms A and B during the epizootics were maintained in tubes of soft MA (half-strength marine agar) and frozen at -70 °C in tryptic soy broth containing 1 % NaCl and 15 % (v/v) glycerol for long-term preservation.

Drug sensitivity of the isolates was assayed by the disc diffusion method on Mueller-Hinton Agar (Oxoid) supplemented with 1 % NaCl. The chemotherapeutic agents and their concentrations (μ g disc⁻¹) were: penicillin G (10), ampicillin (10), tetracycline (30), chloramphenicol (30), cephalothin (30), novobiocin (5), gentamycin (10), erythromycin (15), kanamycin (30), streptomycin (10), nalidixic acid (30), oxolinic acid (2), furazolidone (300), sulphafurazole (300), and trimethoprim-sulphametoxazole (25).

Serological assays. Source of antigens and antisera: We preparad antisera in rabbits against 6 representative strains of *Vibrio damsela* (3 of our isolates and 3 reference strains). Antisera were obtained as previously described by Sørensen & Larsen (1986). Briefly, rabbits were injected intravenously with formalinkilled cells twice weekly in consecutive doses of 0.2, 0.4, 0.8 and 1 ml (10^9 cells ml⁻¹). Rabbits were bled from the ear vein 1 wk after the last injection. The blood was allowed to clot and the sera were collected, separated and stored at -30 °C until used.

Agglutination tests: To examine the serological relationship among the Vibrio damsela isolates and other related vibrios, slide agglutinations tests were conducted according to the procedures of Sørensen & Larsen (1986) and Toranzo et al. (1987a). The reactions were performed using the heat-stable 'O' antigens of each strain. The 'O' antigens were prepared by heating suspensions of each strain in sodium-acetate buffered saline at 100 °C for 1 h. A strong and rapid agglutination was recorded as positive and no or only a weak agglutination occurring after 1 to 2 min as a negative reaction.

In order to determine the existence of possible serogroups within *Vibrio damsela* isolates, cross-quantitative agglutination tests were performed in microtitre plates using serial 2-fold dilutions of 25 μ l aliquots of the antisera. The agglutination titre was considered as the reciprocal of the highest dilution of the antiserum giving a positive reaction after incubation with the antigen overnight at 15 °C.

Virulence test. All our isolates were tested for pathogenicity in rainbow trout (5 to 8 g) and in turbot (5 g) at 20 °C (experimental water temperature). Virulence assays were performed by intraperitoneal inoculation of bacterial doses ranging from 10^2 to 10^6 cells per fish as previously described (Toranzo et al. 1983). Mortalities were recorded daily for a 7 d period and were considered to be due to the inoculated strain if it was recovered from the internal organs of dead fish in pure culture. The degree of virulence, expressed as the 50 % mean lethal dose (LD₅₀), was calculated by the method of Reed & Müench (1938).

Preparation of LPS and electrophoretic analysis. The preparation of lipopolysaccharides (LPS) was performed by the method of Hitchcock & Brown (1983). Bacteria were grown in TSA-1 at 25 °C for 24 h and then suspended in 3 ml of PBS to an optical density (O.D.) of 0.8 at 650 nm. Bacterial suspensions (1.5 ml) were centrifuged and the resulting cell pellets were resuspended in 50 μ l of 2× concentrated sample buffer (0.065 M Tris, pH 6.8, 2% SDS, 10% glycerol, 5% βmercaptoethanol, and 0.001 % bromophenol blue) and boiled for 10 min. Ten μ l of 2.5 mg ml⁻¹ proteinase K in 2× concentrated sample buffer were added and incubated at 60 °C for 1 h. Samples (10 µl) were run in sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) (Laemmli 1970) using 12% acrylamide in the resolving gel and 3 % acrylamide in the stacking gel. The separated LPS components were visualized by the silver staining method described by Tsai & Frasch (1982).

Immunoblot assays. After SDS-PAGE analysis, separated LPS components were transferred from the gel to 45 μ m nitrocellulose paper (NCP, Schleicher & Schuell) according to the procedure described by Towbin et al. (1979). The transblotted NCP sheet was incubated for 1 h with diluted (1:1000) rabbit antiserum against whole bacterial cells. LPS components recognized by the antiserum were visualized by reacting the rinsed NCP sheet with goat anti-rabbit IgG alkaline phosphatase conjugate (1:3000) for 1 h and the appropriate substrate (0.3 mg ml⁻¹ tetrazolium blue and 0.15 mg ml⁻¹ 5-bromo-4-chloro-3-indolyl phosphate toluidine salt in 0.1 M carbonate buffer, pH 9.8).

Analysis of cell-envelope proteins. Total and outermembrane proteins were prepared as previously described (Toranzo et al. 1983). The outer membranes were obtained by treatment of the cell envelopes with Sarkosyl 4 % w/v in 10 mM Tris-HCl (pH 8.0) at room temperature for 20 min to dissolve the inner membranes. Pellets from total cell envelopes or outer membranes were examined by SDS-PAGE as described for the LPS analysis.

Analysis of the plasmid content. Extrachromosomal DNA elements in our isolates and in the reference strains were compared. Plasmid DNA was isolated from small volumes (5 ml) of bacterial cultures, following basically the method of Kado & Liu (1981). DNA samples (15 μ l) mixed with 5 μ l of sample buffer (30 % glycerol, 1 mM EDTA, 0.1% bromophenol blue [Merck]) were electrophoresed through 0.7 % agarose in Tris-acetate buffer (40 mM Tris, 2 mM Na₂-EDTA, adjusted to pH 7.9 with glacial acetic acid) at 100 V for 2 h in a horizontal apparatus. Gels were stained in ethidium bromide (0.5 μ g ml⁻¹ of water), destained in water, and photographed at a wavelength of 254 nm. Plasmids from Escherichia coli 39R861 (it contained 4 plasmids of 4.6, 23.9, 42 and 98 MDa) and Vibrio anguillarum 775 (it contained the 47 Md plasmid pJM1) were used as reference standards.

RESULTS

Bacterial identification and characterization

A *Vibrio* was obtained in pure culture on TSA-1 and TCBS agar from the internal organs examined (kidney and liver) and from the muscle around the haemorragic eyes of affected turbot.

Preliminary screening showed the bacterial isolates to be Gram-negative motile rods that were relatively pleomorphic and occasionally formed long flexible chains. The additional physiological and biochemical tests listed in Table 3 allowed us to identify the present isolates from turbot as *Vibrio damsela* because they displayed the same phenotypic profiles as the reference strains from USA.

The strains were oxidase and catalase positive, sensitive to O/129 (both the 10 and 150 μ g disc⁻¹), arginine dihydrolase positive, producers of gas from glucose, and yielded positive Methyl-Red and Voges-Proskauer reactions. They also produced urease but failed to hydrolyze gelatin or to attack sucrose.

The isolates required salt for growth (they grew with 1 to 5 % NaCl) and grew over a very wide temperature range (from 15 to 37 °C), the latter property helping to explain their pathogenicity for both poikilothermic and homiothermic animals.

The strains of Vibrio damsela shared some biochemical characteristics in common with the species V. anguillarum, V. ordalii and V. vulnificus strains. However, they could readily be differentiated from these species of Vibrio on the basis of their reactions on TCBS-agar and with glucose, sucrose, mannitol, arginine, gelatin and urea (Table 4). With the chemotherapeutic agents tested, the *Vibrio* damsela isolates from turbot showed a drug-susceptibility pattern similar to that of the reference strains, all the isolates being resistant to penicillin G, ampicillin, erythromycin, streptomycin and sulphafurazol (Table 3).

Serological types and LPS profiles

The agglutination reactions with thermostable O antigens (Table 5) revealed that our isolates shared somatic antigens in common with all of the other strains of *Vibrio damsela* but not with *V. anguillarum* and *V. ordalii*. These results therefore supported the taxonomic findings reported above.

Serological differences among the *Vibrio damsela* isolates were detected, however, using cross-agglutination tests. The titres of 6 anti-*V. damsela* antisera against each of the 11 *V. damsela* strains tested are shown in Table 6. Although we found some cross-reactions between them, the 11 strains formed basically 4 recognizable groups: strains RG-91, RG-151, RG-153, RG-191, RG-192, RG-193, RG-214 and *V. damsela* ATCC 33539 (designated as Serogroup A); strain RM-71 (Serogroup B); *V damsela* ATCC 35083 (Serogroup C) and *V. damsela* CDC-2227-81 (Serogroup D).

Interestingly, the silver-stained LPS profiles of the *Vibrio damsela* isolates were similar in the 8 strains belonging to the same serological group and distinct for each of the other 4 cultures (Fig. 2a). In addition, the immunoblot assays using antisera from each group supported the serological differences detected by agglutination tests among the strains in that no cross-reactions were observed among the distinct serological groups. Fig. 2b illustrates this point for the reactions obtained with the antiserum prepared against *V. damsela* ATCC 33539, a Serogroup A strain: only the 8 Serogroup A strains reacted with the antiserum.

Pathogenicity tests

The virulence assays with *Vibrio damsela* strains demonstrated that practically all were pathogenic for turbot with an LD_{50} ranging from 1×10^3 to 3×10^5 CFU (colony forming units) per fish. Interestingly, rainbow trout proved to be highly susceptible to *V. damsela* under laboratory conditions: LD_{50} 's ranged from 9×10^3 to 1.3×10^5 CFU per fish. Only *V. damsela* ATCC 35083 (isolated from brown shark) was non-virulent for any of the fish species challenged.

The inoculated strains were reisolated in pure culture from the internal organs of all of the moribund and dead fish. Table 3. Comparison of the biochemical and physiological characteristics of the strains isolated from turbot with those of *Vibrio damsela* reference strains isolated from poikilotherms and homeotherms. +: Positive; (+): weak positive strain; -: negative; NC: no change, unreactive strain; F: fermentative; G: green colonies; R: resistant; S: sensitive; (S): weak sensitive strain

| Characteristics Luminescence Gram stain Motility Oxidase Catalase Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: 0 % NaCl | | Pre | sent isola | V. damsela reference strains | | | | | |
|---|-------|---------|------------|------------------------------|--------|---------------|---------------|----------------|----------------|
| Gram stain Motility Oxidase Catalase Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | RG-91 | RG-1513 | | | RM-71 | ATCC 33539 | ATCC 35083 | CDC 2227-81 | CDC 1421-81 |
| Motility Oxidase Catalase Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | | - | | | - | - | _ | | _ |
| Oxidase Catalase Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | _ | - | - | _ | | - | | _ |
| Catalase Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Growth on TCBS Resistance to: O/129 (10 µg disc ⁻¹) O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | ÷ | + | + | + | + | + | + | + |
| Resistance to: $O/129 (10 \ \mu g \ disc^{-1})$ $O/129 (150 \ \mu g \ disc^{-1})$ Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| O/129 (10 μg disc ⁻¹) O/129 (150 μg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | +, G | +, G | +, G | +, G | + , G | +, G | +, G | +, G | +, G |
| O/129 (150 µg disc ⁻¹) Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | | | | | | | | | |
| Voges-Proskauer Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | S | S | S | S | S | S | S | S | S |
| Methyl-Red Indole Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | S | S | S | S | S | S | S | S | S |
| Indole Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Moeller's arginine Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | ÷ | + | ÷ | + | ł |
| Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | | - | - | - | _ | _ | - | _ | _ |
| Moeller's lysine Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Moeller's ornithine Enzyme production: Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | _ | _ | - | _ | | - | NC | _ |
| Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | | - | - | _ | - | - | | |
| Gelatinase Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | | | | | | | | | |
| Urease Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | _ | - | | - | _ | - | _ | _ |
| Caseinase Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Alginase Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | _ | _ | - | _ | _ | _ | _ | _ |
| Chitinase Amylase Lipase Haemolysis of RBC Growth in: | _ | _ | _ | - | _ | _ | _ | _ | _ |
| Amylase Lipase Haemolysis of RBC Growth in: | + | + | + | 4 | + | (+) | + | (+) | + |
| Lipase Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Haemolysis of RBC Growth in: | + | + | + | + | + | + | + | + | + |
| Growth in: | | | | | | | | | |
| | + | + | + | + | + | + | + | + | + |
| 0% NaCl | | | | | | | | | |
| | _ | _ | | | | _ | _ | _ | _ |
| 3 % NaCl | + | + | + | + | + | + | + | + | + |
| 5 % NaCl | + | + | + | + | + | + | + | + | (+) |
| 8 % NaCl | _ | - | - | - | - | - | _ | _ | - |
| 10 % NaCl | _ | _ | - | - | - | _ | - | - | - |
| O/F (Leifson) | F | F | F | F | F | F | F | F | F |
| Gas from glucose | + | + | + | + | + | + | + | + | + |
| Acid production from: | | | | | | | _ | _ | |
| Sucrose | - | — | - | _ | | _ | - | | - |
| Arabinose | - | _ | - | _ | _ | _ | _ | _ | - |
| Glucose | + | + | + | + | + | + | + | + | + |
| Inositol | - | - | - | - | _ | _ | - | - | - |
| Mannitol | _ | _ | _ | _ | _ | _ | - | _ | - |
| Mannose | + | + | + | + | + | + | + | + | + |
| Maltose | + | + | + | + | + | + | + | + | + |
| Utilization of: | | | | | | | | | |
| γ-aminobutyrate | — | - | - | - | - | _ | - | _ | 22 |
| L-citrulline | | - | - | _ | _ | - | - | _ | |
| Gluconate | — | - | - | _ | _ | | | (+) | |
| Glucuronate | _ | - | - | - | _ | _ | - | - | |
| Heptanoate | _ | | - | - | _ | _ | - | - | |
| α-Ketoglutarate | _ | | _ | _ | - | _ | - | | |
| Putrescine | _ | - | - | _ | _ | _ | - | _ | 220 |
| Tyrosine | _ | | - | - | _ | _ | - | | - |
| Xanthine | _ | - | - | _ | - | - | - | - | |
| Growth at: | | | | | | | | | |
| 4 °C | | _ | | _ | - | _ | | - | - |
| 15°C | | | | | | | | | |
| 22°C | + | + | + | + | + | + | + | + | + |
| 37 °C | +++ | + + | + + | ++++ | + + | + + | + + | + + | + + |
| 44 °C | | | | | | | | | |

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Table 3 (continued)

| Characteristics | | Pre | V. damsela reference strains | | | | | | |
|-------------------------------------|-------|---------|------------------------------|--------|-------|---------------|---------------|----------------|----------------|
| | RG-91 | RG-1513 | RG-191 | RG-214 | RM-71 | ATCC 33539 | ATCC 35083 | CDC 2227-81 | CDC 1421-81 |
| Resistance/Sensitivity to: | | | | | | | | | |
| Penicillin G | R | R | R | R | R | R | R | R | R |
| Ampicillin | R | R | R | R | R | R | R | R | R |
| Tetracycline | S | S | S | S | S | S | S | S | S |
| Chloramphenicol | S | S | S | S | S | S | S | S | S |
| Cephalothin | S | S | S | S | S | S | S | S | S |
| Novobiocin | S | S | S | S | S | S | S | S | S |
| Gentamycin | S | S | S | S | S | S | S | S | S |
| Erythromycin | R | R | R | R | R | R | R | R | R |
| Kanamycin | (S) | (S) | (S) | (S) | (S) | (S) | (S) | (S) | S |
| Streptomycin | R | R | R | R | R | R | R | R | R |
| Nalidixic acid | S | S | S | S | S | S | S | S | S |
| Oxolinic acid | S | S | S | S | S | S | S | S | S |
| Furazolidone | S | S | S | S | S | S | S | S | S |
| Sulphafurazole | R | R | R | R | R | R | R | R | R |
| Trimethioprim- sulphamethosazole | S | S | S | S | S | S | S | S | S |

Analysis of the outer-membrane proteins in Vibrio damsela strains

Analysis of total cell envelopes as well as outer membranes showed that although all strains exhibited some common protein bands (40, 30, 21.5, 20.5), different patterns were observed (Fig. 3). All turbot isolates belonging to Serogroup A (Lanes B to H) shared at least 5 outer membrane proteins of 70, 31.5, 30, 27.5 and 26.5 kilodaltons (kDa). Only in the RG-91 strain (Lane B) could we not detect the 72.5 and 89 kDa proteins.

Reference strain ATCC 33539 (Lane I), also belonging to Serogroup A, showed a different pattern. This strain shared another major protein of 33.5 kDa with isolates of Serogroups B and C (Lanes J and K). However, the patterns of Serogroups B and C were more similar to each other than to the pattern of Serogroup A. The human isolates (Lanes L and M) also exhibited distinctive patterns.

Plasmid content of Vibrio damsela strains

Analysis of the plasmid content of *Vibrio damsela* strains revealed that all of the isolates from turbot carried a 90 to 100 MDa plasmid band similar in size to that found in the reference strains ATCC 33539 and CDC-2227-81 (Fig. 4). However, no plasmid of similar molecular mass was found in the fish strain ATCC 35083 (Lane H) or in the human isolate CDC-1421-81 (Lane K).

 Table 4. Vibrio spp. Main differential characteristics among V. damsela, V. anguillarum, V. vulnificus and V. ordalii. G: Green colonies; Y: yellow colonies; NG: no growth; V: variable

| Characteristics | V. damsela | V. anguillarum | V. ordalii | V. vulnificus |
|---------------------|------------|----------------|------------|---------------|
| Growth on TCBS-agar | +, G | +, Y | +, Y or NG | +, G |
| Gas from glucose | + | _ | _ | <u></u> |
| Fermentation of: | | | | |
| Arabinose | _ | Vª | _ | - |
| Sucrose | _ | + | + | - |
| Mannitol | - | + | - | _ |
| Moeller's arginine | + | + | _ | _ |
| Moeller's lysine | 1998 | _ | _ | + |
| Moeller's ornithine | _ | _ | _ | V |
| Gelatinase | - | + | + | + |
| Urease | + | _ | - | - |

| O Antigen | | | V. dai | | | Antisera to | | |
|--------------------------------------|-------|--------|----------------|---------------|---------------|-----------------|------|--------------|
| | | | V. anguillarum | V. ordali | | | | |
| | RG-91 | RG-191 | RM-71 | ATCC 33539 | ATCC 35083 | CDC- 2227-81 | R 82 | NCMB 2167 |
| RG-91 | ++ | ++ | (+) | ++ | + | (+) | _ | |
| RG-151 | ++ | ++ | (+) | + | (+) | (+) | | - |
| RG-153 | ++ | ++ | (+) | + | (+) | (+) | | - |
| RG-191 | + + | ++ | (+) | + | (+) | (+) | - | _ |
| RG-192 | ++ | + + | (+) | + | (+) | (+) | - | _ |
| RG-193 | ++ | + + | (+) | + | (+) | (+) | - | - |
| RG-214 | ++ | + + | (+) | + | (+) | (+) | - | - |
| RM-71 | + | + | ++ | (+) | + | (+) | - | |
| V. damsela ATCC 33539 | ++ | ++ | (+) | ++ | (+) | + | _ | _ |
| V. damsela ATCC 35083 | (+) | (+) | (+) | (+) | ++ | (+) | - | _ |
| V. damsela CDC-2227-81 | + | + | + | + | + | ++ | _ | _ |
| V. damsela CDC-1421-81 | AG | AG | AG | AG | AG | AG | AG | AG |
| V. anguillarum R 82 V. ordalii | - | | - | - | - | _ | ++ | - |
| NCMB 2167 | _ | _ | - | _ | _ | _ | _ | ++ |

| Table 5. Vibrio spp. Agglutination of thermostable O antigen of selected V. damsela strains using rabbit antisera prepared against |
|--|
| selected strains of V. damsela, V. anguillarum and V. ordalii. ++: fast, strong positive reaction; +: positive reaction in 20 to 30 s; |
| (+): weak and delayed reaction; -: negative; AG: agglutinating strain |

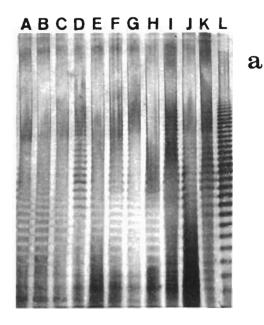
DISCUSSION

Turbot *Scophthalmus maximus* has been the main fish species cultured on the northwest coast of Spain during the last 5 yr. Perhaps as a result of this, diseases in these intensive culture facilities caused by bacteria, viruses, and parasites have increased, constituting a limiting factor in turbot production. Although until now the most important bacteriological problems were caused by *Vibrio anguillarum*, other *Vibrio* species were occasionally implicated in turbot diseases.

We recently reported the first occurrence of vibriosis

 Table 6. Vibrio damsela. Cross-agglutination titres among the selected V. damsela strains. Rabbit antisera were prepared against whole bacterial cells. Four distinct serogroups are indicated in boxes

| O Antigen | | Antisera | | | | | | | | |
|-------------|-------|----------|------------|-------|------------|-------------|--|--|--|--|
| | RG-91 | RG-191 | ATCC 33539 | RM-71 | ATCC 35083 | CDC 2227-81 | | | | |
| RG-91 | 1280ª | 1280 | 640 | <20 | <20 | < 20 | | | | |
| RG-151 | 640 | 1280 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| RG-153 | 640 | 5120 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| RG-191 | 640 | 5120 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| RG-192 | 640 | 5120 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| RG-193 | 640 | 5120 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| RG-214 | 640 | 5120 | 640 | 20-40 | 20-40 | 20-40 | | | | |
| ATCC 33539 | 1280 | 1280 | 1280 | < 20 | < 20 | < 20 | | | | |
| RM-71 | <20 | 40 | <10 | 160 | 40 | < 20 | | | | |
| ATCC 35083 | < 10 | < 20 | < 20 | 20 | 1280 | < 20 | | | | |
| CDC-2227-81 | 80 | 80 | 20-40 | 20-40 | < 10 | 640 | | | | |



ABCDEFGHIJKL

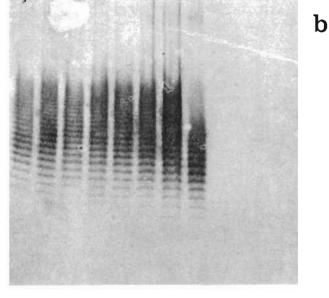


Fig. 2. (a) Silver-stained polyacrylamide gel of isolated lipopolysaccharides of *Vibrio damsela* strains. (b) Immunoblot assay of *V. damsela* strains using an antiserum prepared against the reference strain ATCC 33539. Lanes: (A) RG-91;
(B) RG-191; (C) RG-192; (D) RG-193; (E) RG-194; (F) RG-153;
(G) RG-214; (H) ATCC 33539; (I) RM-71; (J) ATCC 35083; (K) CDC-2227-81; (L) CDC-1421-81

caused by Vibrio damsela in 2 turbot farms along the Atlantic coast (Fouz et al. 1991). Interestingly, some of these isolates had been earlier reported to be members of the bacterial group *V. splendidus-V. pelagius*

(Lupiani et al. 1989), a group that is very abundant in the aquatic environment (Fouz et al. 1990). However, the taxonomic, serological, and molecular characterization of the turbot isolates indicates that they belong to V. damsela species.

The biochemical and physiological properties of the present bacterial isolates (Tables 3 & 4) are similar to those of the reference strains of *Vibrio damsela* isolated from poikilotherms and homiotherms, giving strong positive reactions for arginine dihydrolase, gas production from glucose, and hydrolysis of urea, and negative reactions for fermentation of sucrose and hydrolysis of gelatin. These features allowed us to differentiate *V. damsela* from other *Vibrio* species, such as *V. anguillarum, V. ordalii* and *V. vulnificus* which are considered to be the main causes of vibriosis in marine aquaculture.

Disease onset was correlated with a sudden increase in water temperature to about 22 to 24 °C. This condition is very favorable for the rapid growth and multiplication of *Vibrio damsela*, the optimal growth temperature of which under laboratory conditions is approximately the same.

The *Vibrio damsela* strains isolated from turbot, as well as the reference strains, were sensitive to the chemotherapeutic agents commonly used for controlling vibriosis outbreaks such as tetracycline, oxolinic acid and trimethoprim. No appearance of drug-resistance was observed in the isolates during the course of the disease.

The serological analysis supported the identification of our isolates as *Vibrio damsela*, because the 'O' antigens of our isolates reacted positively only with antisera raised against the reference strains of this species (Table 5). However, differences in the strength of the agglutination reactions indicated that the *V. damsela* strains used in this study were not totally homogeneous antigenically. Serological heterogeneity among these *V. damsela* isolates was confirmed by the cross-agglutination tests, which indicated the existence of 4 different Serogroups (A, B, C and D). These data suggest that *V. damsela*, like *V. anguillarum* (Sørensen & Larsen 1986), is likely to turn out to be a heterogeneous species.

The analysis of lipopolysaccharides (LPS) present in the bacterial cell wall supported the serological heterogeneity detected among the *Vibrio damsela* strains because all of the isolates belonging to Serogroup A exhibited the same LPS profile, a profile that was different from that of each of the other 3 groups. In addition, the immunoblot assays performed with LPS preparations confirmed these results. Furthermore, although the relationship between the electrophoretic patterns of the outer membrane proteins and the crossagglutination groups was not very strong, the patterns

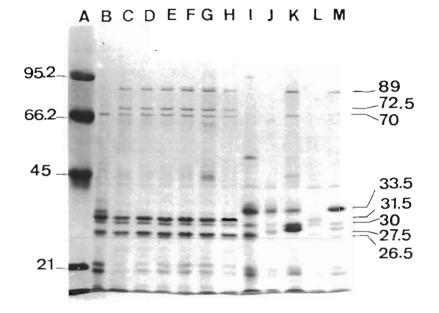


Fig. 3. SDS-PAGE of outer-membrane proteins of selected *Vibrio damsela* strains: (A) molecular weight standards; (B) RG-91; (C) RG-191; (D) RG-192; (E) RG-193; (F) RG-194; (G) RG-153; (H) RG-214; (I) ATCC 33539; (J) RM-71; (K) ATCC 35083; (L) CDC-2227-81; (M) CDC-1421-81. Numbers indicate molecular weight in kDa

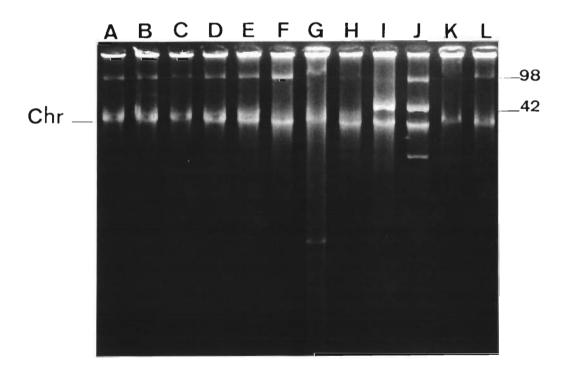


Fig. 4. Demonstration of plasmid profiles in selected strains of *Vibrio damsela* by the method of Kado & Liu (1981). Lanes: (A) RG-91; (B) RG-191; (C) RG-151; (D) RG-153; (E) RG-214; (F) RM-71; (G) ATCC 33539; (H) ATCC 35083; (I) *V. anguillarum* 775 (O1); (J) *E. coli* 39R861; (K) CDC-1421-81; (L) CDC-2227-81 Numbers indicate molecular weight in MDa. Chr: chromosomal band

tended to support our findings in the agglutination tests. Similar results have been reported by Nomura & Aoki (1985) and Pyle & Schill (1985) for several other Gram-negative fish pathogenic bacteria and for *V. anguillarum* by Toranzo et al. (1987b).

All of the turbot isolates harbored a high molecular weight plasmid band (90 to 100 MDa) but it is not known whether the plasmid coded for virulence of the isolates as does the 47 MDa plasmid of *V. anguillarum* serotype O1 (Crossa 1980, Tolmasky et al. 1985, Toranzo et al. 1987b).

Experimental infections in turbot with our isolates confirmed that the causative agent of these vibriosis outbreaks was *Vibrio damsela*. Our isolates were also highly pathogenic for rainbow trout, suggesting that *V. damsela* has a host range broader than that observed by Love et al. (1981). Among the reference strains, only ATCC 35083 was not pathogenic for both turbot and rainbow trout. All these features indicate that *V. damsela* does not possess host specificity.

Although *Vibrio damsela* has been more frequently associated with human wound infections that with fish mortalities, its importance as a pathogen in salt water aquaculture is being increasingly recognized. It seems likely, therefore, that it may become necessary to consider protecting fish against vibriosis with polyvalent vaccines made with strains of *V. anguillarum* serotypes O1 and O2 (the main virulent serotypes) and *V. damsela*.

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