

Systemic disease in turbot *Scophthalmus maximus* caused by a previously unrecognised *Cytophaga*-like bacterium

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ABSTRACT: A new group of *Cytophaga*-like bacteria has been recovered from healthy and diseased turbot in Scotland. Cultures produced orange colonies and contained short rods with distinctively thick cell walls. Gliding movement was observed and the guanine plus cytosine ratio of the DNA was 31.7 to 32.1 mols %. The organism caused gill hyperplasia and systemic haemorrhagic septicaemia in both naturally diseased and experimentally infected turbot. Control of the disease was possible by administering an injection or bath of furazolidone.

INTRODUCTION

In a previous study, suitable methods were developed for the recovery of bacteria from the gills of turbot *Scophthalmus maximus* L. (Mudarris & Austin 1988). A nutrient limited medium was formulated, designated as medium K, which was especially suitable for the isolation and maintenance of aerobic heterotrophic bacteria occurring on gill tissue. Using this medium, gill bacteria from turbot were enumerated and then identified using conventional diagnostic schemes (e.g. Starr et al. 1981). The bacterial cultures were also examined for possible deleterious effects on fish. During this investigation, cultures forming a single group of orange-pigmented Gram-negative rods with low DNA guanine plus cytosine ratios and gliding motility, i.e. *Cytophaga*, caused gill hyperplasia and a generalised haemorrhagic septicaemia in juvenile turbot. The significance of these findings was reinforced by the outbreak of a disease in farmed turbot in Scotland during 1987 caused by a similar bacterium. The results of bacteriological examinations of the disease are reported here. Details of the histopathology of the disease will be presented separately (Mudarris & Austin unpubl).

MATERIALS AND METHODS

Fish. Juvenile turbot (average weight = 100 g) were obtained from a commercial fish farm in Scotland. These were maintained within the laboratory in glass aquaria containing aerated re-circulating seawater at ca 18°C. Adult turbot (average weight = 2.5 kg) displaying clinical signs of disease (i.e. lethargy, haemorrhaging in the gills, distended abdomens, and an assortment of surface lesions) were obtained directly from a fish farm and from holding facilities within Heriot-Watt University. In addition, wild fish (and water samples) were obtained from the mouth of the River Forth (Firth of Forth) near Edinburgh, Scotland.

Isolation of bacteria from turbot. From healthy fish, gill tissue (1.0 g) was homogenised, and dilutions to 10⁻⁴ were prepared in 10 ml aliquots of sterile (121°C/15 min) seawater. Approximately 10 cm² areas of skin from the middle of the dorsal surface were swabbed (Exogen, cotton-tipped swabs) vigorously for ca 15 sec. With diseased turbot, samples of gill, skin, kidney, ascitic fluid, and intestinal contents were homogenised or swabbed, as appropriate. Diluted samples (0.1 ml) and swabbed material were inoculated onto triplicate plates of brain heart infusion agar (BHIA; Oxoid), glucose yeast extract agar (GYEA; Oxoid), marine 2216E agar (Difco), and medium K. Incubation was at 25°C for up to 14 d. Thereafter, colony counts were made, and pure cultures were prepared by streaking and re-streaking on fresh media. Purity was confirmed by the

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examination of Gram-stained smears (Hucker & Conn 1923). The pure cultures were maintained on slopes of medium K at 4 °C, with transfer every 6 to 8 wk. Stock cultures were also freeze-dried.

Examination of bacterial isolates. Cultures were examined for a wide range of phenotypic and genotypic characteristics. The phenotypic traits included all tests considered to have diagnostic and differential value for Gram-negative pigmented gliding bacteria (Starr et al. 1981). Methods were adapted from Cowan (1974), van der Meulen et al. (1974), and Gerhardt et al. (1981). The features examined included colonial (notably the presence of flexirubin pigments as determined by the production of a purple colour by colonies after the addition of 20% [w/v] potassium hydroxide) and cell-morphology; the presence of gliding on medium K; the oxidative or fermentative metabolism of glucose; the production of ammonia (from arginine), arginine dihydrolase, catalase, H₂S, indole, lysine and ornithine decarboxylases, methyl red reaction, nitrate reduction, oxidase, phenylalanine deaminase, phosphatase and the Voges-Proskauer reaction; the production of acid from arabinose, cellobiose, glucose, lactose, mannitol, raffinose, salicin, sucrose, and xylose; the degradation of aesculin, agar, blood (β -haemolysis), casein, cellulose, chitin, DNA, gelatin, starch, tributyrin, Tween 20, 40, 60, 80 and 85, tyrosine, and urea; the utilization of alginate, L-arabinose, L-arabitol, carboxymethyl cellulose, cellobiose, erythritol, ethanol, D-fructose, D-glucose, glycerol, heparin, inositol, inulin, L-leucine, lysine, maltose, mannitol, methanol, L-methionine, pectin, L-phenylalanine, polypectate, raffinose, D-ribose, sodium benzoate, sodium citrate, sodium succinate, sorbitol, sucrose, D-xylitol, and D-xylose as the sole source of carbon for energy and growth; the utilization of sodium nitrate, vitamin-free casamino acids and yeast extract as the source of nitrogen; growth in 0 to 5% (w/v) NaCl; growth at 4 to 42 °C; growth on MacConkey agar; and sensitivity to ampicillin (10, 25 μ g), carbenicillin (100 μ g), chloramphenicol (10, 50 μ g), chlortetracycline (10 μ g), cephaloridine (25 μ g), cloxacillin (5 μ g), colistin sulphate (10 μ g), co-trimoxazole (25 μ g), erythromycin (10 μ g), furazolidone (50 μ g), fusidic acid (10 μ g), gentamicin (10 μ g), kanamycin (30 μ g), lincomycin (2 μ g), methicillin (10 μ g), nalidixic acid (5, 30 μ g), neomycin (10 μ g), nitrofurantoin (200 μ g), novobiocin (5 μ g), oleandomycin (5 μ g), oxytetracycline (10 μ g), penicillin G (1.5 IU), streptomycin (10, 25 μ g), sulphadiazine (50 μ g), sulphafurazole (100, 500 μ g), and tetracycline (10, 50 μ g).

DNA base composition. DNA was extracted and purified by the method of Mandel et al. (1971). The base composition was determined from T_m values (Marmur & Doty 1962).

Fish experiments. Groups containing 6 juvenile turbot (average weight = 100 g) were maintained in aerated re-circulating seawater at 18 °C. In addition, rainbow trout *Salmo gairdneri* (average weight = 19 g) were kept in aerated fresh water, also at 18 °C. The fish were infected, via intraperitoneal injection with either 10⁷, 10⁵, or 10² cells of a representative strain, i.e. strain MM₁, suspended in either 0.5 ml volumes of 0.9% (w/v) saline or in 0.2 ml aliquots of distilled water for the turbot and rainbow trout, respectively. Negative and positive controls were also used and consisted of injections with saline and *Vibrio anguillarum* (NCMB 1873; National Collection of Marine Bacteria, Aberdeen, Scotland), respectively. Parallel experiments sought to examine the effect of immersion of groups of fish in suspensions containing 10⁷ bacterial cells ml⁻¹ for 15 min. The fish were examined daily for up to 10 d. Following challenge, bacteriological (as described above) and histological examinations were carried out on all dead and moribund fish. The effect of intraperitoneal injections with 0.5 ml volumes of 0.9% (w/v) saline containing culture supernatants (1 g dry wt), cell homogenates (5 mg dry wt), or lipopolysaccharide (LPS) (0.2 mg dry wt) preparations was also examined. Here, the relevant methods followed those of Schnaitman (1970), Saeed & Plumb (1986) and Neidhardt (1987). The findings of the histological examination will be reported separately (Mudarris & Austin unpubl.). Potential control of the disease was examined by administering furazolidone or sulphafurazole to infected turbot, either by single intraperitoneal injection (50, 100 mg/kg of body wt) or by bathing (50, 100 mg l⁻¹ for 30 min daily for 10 d) when the initial signs of disease were observed.

Transmission electron microscopy. Dense suspensions in 0.85% (w/v) saline were prepared from plate cultures. Suspensions were fixed overnight in 0.5% (v/v) buffered (pH 7.4) glutaraldehyde. To determine cell-morphology, preparations were stained with 1% (v/v) phosphotungstic acid. Thin sections were also prepared and stained according to Watson (1958), Millonig (1961), and Reynolds (1963). All prepared specimens were examined in an AEI EM6G transmission electron microscope (TEM).

RESULTS

Recovery of orange pigmented gliding bacteria

From the gills of healthy turbot a variety of orange pigmented Gram-negative bacteria, which demonstrated gliding movement, was recovered. These bacteria were consistently recovered in highest numbers on medium K. On the basis of their phenotypic traits, these organisms were equated with *Cytophaga* and

Flexibacter (Reichenbach & Dworkin 1981). Based on the results of initial experiments, attention focused on an organism possessing the key traits of the genus *Cytophaga*; it was different from any of the currently recognised species of the genus included in the 'Approved lists of bacterial names' (Skerman et al. 1980) or their supplements. Low numbers, i.e. ca 0.1 to 1.0% of the total number of colonies on medium K, of this organism were consistently recovered from the gills of all healthy juvenile and adult turbot. Moreover, an identical organism was recovered as dense pure culture growth from the gills and internal organs of clinically diseased turbot obtained from an external site, and from the wild. These fish displayed gill disease and a systemic haemorrhagic septicaemia. It should be emphasised, that as a result of routine histopathological and microbiological examinations, there was no evidence of infection by viruses, parasites or other bacteria in these fish. Additional cultures of the *Cytophaga*-like bacterium were obtained from the water of the Firth of Forth. Here, cultures accounted for ca 0.1% of the total number of colonies recovered on medium K. In total, 50 cultures, considered to represent a single taxonomic group of the genus *Cytophaga*, were examined in detail.

Characteristics of the *Cytophaga*-like bacteria

All isolates produced round, raised, shiny orange pigmented (flexirubin pigment) colonies of 2 to 3 mm diam. in 48 h of incubation at 25°C on medium K. Cultures contained uniformly shaped short fermentative Gram-negative rods of ca 2.0 × 0.8 µm in size, which showed gliding movement. Neither flagella nor resting stages were observed. Examination of thin sections by TEM revealed the presence of a thick (43.5 nm) cell envelope (Fig. 1) which was initially mistaken for an extracellular layer. In fact, the thickness was double (i.e. ~ 20 nm) that of *bona fide* *Cytophaga*, such as *C. aquatilis* (Strohl & Tait 1978), and other Gram-negative bacteria (Costerton et al. 1974). Additional characteristics of the bacterium are listed in Table 1. The guanine plus cytosine ratio of the DNA from a representative strain (MM₁) was 31.7 to 32.1 mols %.

Pathogenicity

Naturally diseased turbot, from which the *Cytophaga*-like bacterium was recovered, displayed swollen gill lamellae (hyperplasia) and haemorrhagic septicaemia. Externally, haemorrhaging was evident in the eyes, skin, and jaw. Internally, there was necrosis/haemorrhaging in the brain, stomach, intestinal tract,

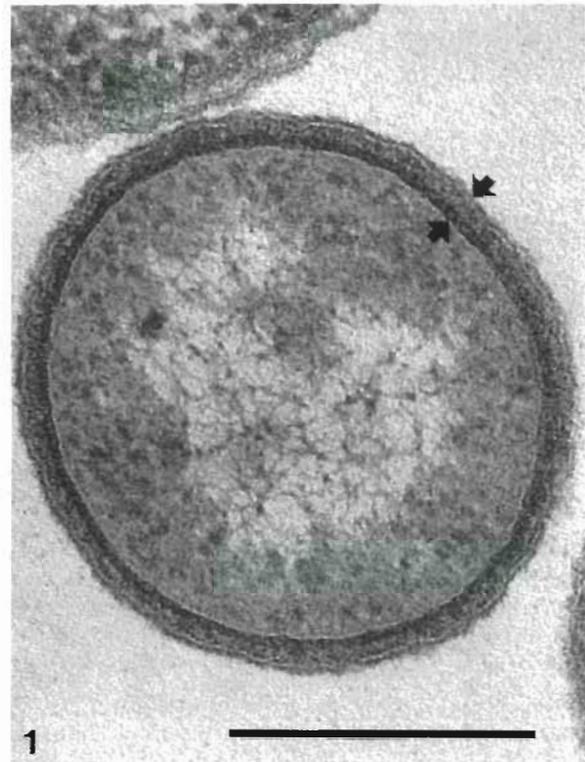


Fig. 1. *Cytophaga*-like bacterium. Transmission electron micrograph of thin section. Note presence of thick cell envelope (arrow). It should be emphasised that most preparations contained obviously rod-shaped rather than coccoid cells. This photograph probably represents a transverse section of a cell, and has been included because of the clear demonstration of the thick cell envelope. Bar = 2.25 µm

liver, and kidney. Ascitic fluid was present in the peritoneal cavity. The intestine was swollen and contained a yellowish fluid.

With juvenile turbot, there was 100% mortality within 2 d following infection with 10⁷ cells. With challenges of 10⁵ and 10² cells, mortalities were lower: 80% and 30%, respectively. By immersion, the disease signs developed after 5 d, with mortalities occurring 2 d later. Interestingly with trout, injection of 10⁷ cells resulted in 70% mortalities within 5 d. All dead fish displayed the same signs described above except that in trout there was no evidence of gill damage. Pure culture growth of the challenge organism was recovered from the internal organs. Interestingly, use of *Vibrio anguillarum* for control purposes produced 100% mortality among infected fish with haemorrhagic septicaemia but no involvement with the gills. Injection of fish with cell supernatant and LPS did not result in any disease signs. However, injection of cell homogenates produced clinical disease, notably the presence of swollen oedematous gill lamellae, and haemorrhaging in the stomach, intestinal tract, kidney, and liver.

Table 1. Characteristics (percentage positive response of 50 tested isolates) of the fish-pathogenic *Cytophaga*-like bacteria isolated from turbot *Scophthalmus maximus*

| Character | % positive response | Character | % positive response |
|--|---------------------|--|---------------------|
| Biochemical tests: | | | |
| Acid production from sugars (listed in text) | 0 | Inositol | 0 |
| Ammonia from arginine | 100 | Inulin | 0 |
| Arginine dihydrolase | 0 | L-leucine | 0 |
| Catalase production | 100 | Lysine | 0 |
| H ₂ S production | 0 | Maltose | 100 |
| Indole | 0 | Mannitol | 0 |
| Lysine decarboxylase | 0 | Methanol | 0 |
| Methyl red test | 0 | L-methionine | 0 |
| Nitrate reduction | 0 | Pectin | 100 |
| Ornithine decarboxylase | 0 | L-phenylalanine | 0 |
| Oxidase production | 100 | Polypectate | 82 |
| Phenylalanine deaminase | 82 | Raffinose | 0 |
| Phosphatase | 86 | D-ribose | 0 |
| Voges-Proskauer reaction | 0 | Sodium benzoate | 0 |
| Growth at/on: | | | |
| 15-25 °C | 100 | Sodium citrate | 16 |
| 4 °C | 80 (weak) | Sodium succinate | 0 |
| 35 °C | 0 | Sorbitol | 0 |
| 0-4 % (w/v) NaCl | 100 | Sucrose | 0 |
| 5 % (w/v) NaCl | 0 | D-xylitol | 100 |
| MacConkey agar | 0 | D-xylose | 0 |
| Degradation of: | | | |
| Aesculin | 88 | Utilization, as sole nitrogen source: | |
| Agar | 0 | Casamino acids, vitamin-free | 100 |
| Blood (β -haemolysis) | 84 | Sodium nitrate | 100 |
| Casein | 84 | Yeast extract | 100 |
| Cellulose | 0 | Susceptibility to: | |
| Chitin | 0 | Ampicillin (25 μ g) | 80 |
| DNA | 100 | Carbenicillin (100 μ g) | 78 |
| Gelatin | 96 | Chloramphenicol (50 μ g) | 100 |
| Starch | 0 | Chlortetracycline (10 μ g) | 0 |
| Tributyryn | 80 | Cephaloridine (25 μ g) | 16 |
| Tween 20 | 82 | Cloxacillin (5 μ g) | 0 |
| Tween 40 | 88 | Colistin sulphate (10 μ g) | 2 |
| Tween 60 | 82 | Co-trimoxazole (25 μ g) | 100 |
| Tween 80 | 92 | Erythromycin (10 μ g) | 90 |
| Tween 85 | 86 | Furazolidone (50 μ g) | 100 |
| Tyrosine | 96 | Fusidic acid (10 μ g) | 100 |
| Urea | 86 | Gentamicin (10 μ g) | 0 |
| Utilization, as sole carbon source: | | | |
| Alginate | 0 | Kanamycin (30 μ g) | 0 |
| L-arabinose | 0 | Lincomycin (2 μ g) | 0 |
| L-arabitol | 0 | Methicillin (10 μ g) | 0 |
| Carboxymethylcellulose | 0 | Nalidixic acid (30 μ g) | 18 |
| Cellulbiose | 88 | Neomycin (10 μ g) | 2 |
| Erythritol | 0 | Nitrofurantoin (200 μ g) | 96 |
| Ethanol | 100 | Novobiocin (5 μ g) | 100 |
| D-fructose | 100 | Oleandomycin (5 μ g) | 0 |
| D-glucose | 100 | Oxytetracycline (10 μ g) | 0 |
| Glycerol | 0 | Penicillin G (1.5 IU) | 0 |
| Heparin | 0 | Streptomycin (25 μ g) | 0 |
| | | Sulphadiazine (50 μ g) | 0 |
| | | Sulphafurazole (100 μ g) | 82 |
| | | Tetracycline (50 μ g) | 0 |

Control of the disease was possible by injection (83 % survival) and bathing (50 % survival) with furazolidone at 50 mg per kg body wt of fish and 50 mg l⁻¹, respectively. In contrast, with sulphafurazole, only 33 and 17 % of the population survived injection and bathing, respectively. All of the untreated fish in the control group died in 4 d.

DISCUSSION

An organism causing a gill and systemic disease has been recovered from turbot in the South of Scotland. In laboratory challenges the organism produced a disease which clearly resembled the natural infection. The organism may not be obligately marine because growth was recorded in 0 % (w/v) NaCl. The organism was equated with the genus *Cytophaga* because of its cell and colonial morphology, its gliding motility, and the guanine plus cytosine ratio of its DNA (Reichenbach & Dworkin 1981). It differed from any of the taxa of fish pathogens included in Austin & Austin (1987) and from any of the validly recognised species of *Cytophaga*. Although it appears likely that the organism represents a new species of *Cytophaga*, it should be emphasised that the genus is heterogeneous and may warrant sub-division into two or more genera (Reichenbach & Dworkin 1981). For this reason, the organism should be regarded as a *Cytophaga*-like bacterium until future improvements in the taxonomy of the genus *Cytophaga* permit a more definitive identification. It is noteworthy that the bacterium caused a haemorrhagic condition in fish, especially as haemorrhagic lesions are not usually associated with gliding bacteria (Austin & Austin 1987). There can be little doubt, however, that it was the responsible agent because other systemic microorganisms were not recovered from the experimentally or naturally infected fish.

From the results of this study, it is considered that the *Cytophaga*-like bacterium may well form part of the normal gill microflora of healthy turbot. This possibility agrees with the scenario for *Cytophaga* pathogens of penaeids, proposed by Lightner (1985). The change in status from harmless saprophyte to pathogen may indicate the presence of as yet unknown factors that serve to weaken the host or to enhance bacterial aggressiveness. The overall importance of the organism to fish pathology remains to be properly elucidated. Nevertheless, the disease has been associated with both farmed and wild fish, albeit not in epizootic proportions. At present, it is considered that virulence factors may involve exotoxins, as illustrated by the proteolytic activity. Certainly, further work should be directed at the unusually thick cell envelope, which may be involved with the pathogenic mechanism.

A culture of the *Cytophaga*-like bacterium, strain MM₁, has been deposited in the Czechoslovak Collection of Microorganisms, J.E. Purkyňe University, 662 43 Brno, Czechoslovakia, as CCM 1409.

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