Molecular detection of *Renibacterium salmoninarum* in rainbow trout (*Oncorhynchus mykiss*) from Poland

Leszek Guz, Krzysztof Puk

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Abstract. *Renibacterium salmoninarum* causes bacterial kidney disease mainly in salmonid fish. Oligonucleotide primers incorporating *R. salmoninarum* unique sequences were designed to amplify a 501 bp region of the gene encoding a 57 kDa soluble extra-cellular protein. The primers did not amplify other wide varieties of aquatic or piscine bacteria *Aeromonas salmonicida* or *Yersinia ruckeri*. This assay provides a molecular description and definitive identification of *R. salmoninarum* in Poland.

Keywords: *Renibacterium salmoninarum*, fish, bacterial kidney disease, infection

Renibacterium salmoninarum is a small, G(+), rod-shaped bacterium that causes bacterial kidney disease (BKD) mainly in salmonid species such as sockeye salmon, Oncorhynchus nerka (Walbaum), rainbow trout, O. mykiss (Walbaum), chinook salmon, O. tshawytscha (Walbaum), coho salmon, O. kisutch (Walbaum), cutthroat trout, O. clarkii (Richardson), masou salmon, O. masou (Brevoort), pink salmon, O. garbuscha (Walbaum), Danube salmon, Hucho hucho (L.), grayling, Thymallus thymallus (L.), brook trout, Salvelinus fontinalis (Mitchill), Atlantic salmon, Salmo salar L., brown trout, Salmo trutta L., and in non-salmonid species such as ayu, Plecoglossus altivelis (Temminck and Schlegel), burbot, Lota lota (L.), Pacific herring, Clupea pallasii Val., and sea lamprey, Petromyzon marinus L., in both freshwater and seawater environments (Eissa et al. 2006). Because a non-salmonid reservoir has not been identified, R. salmoninarum is currently considered an obligate pathogen of salmonids. The bacterium can be horizontally and vertically transmitted, and the disease is reported in many regions and countries worldwide including North and South America, Japan, and Europe (i.e., England, Denmark, Norway, France, Germany, and Poland) (Antychowicz and Grawiński 1994, Kozińska et al. 2001, Jansson 2002, Pękala-Safińska et al. 2017, Terech-Majewska et al. 2017).

Fish harboring *R. salmoninarum* can display various internal and external signs of the disease such as exophthalmia, skin blebs, cavitations in the musculature, distended abdomen, hemorrhagic areas in the skin, swollen kidney, and grayish nodules in the kidney, spleen, and liver. BKD is diagnosed by a combination of culturing and serological methods (Antychowicz and Grawiński 1994, Kozińska et al. 2001, Kozińska and Pękala 2005). The use of PCR has been advocated as a specific and sensitive means of diagnosing BKD (Brown et al. 1994, McIntosh et al. 1996). Several PCR and RT (reverse transcription) – PCR methods have been developed for *R*.

L. Guz [], K. Puk

Department of Fish Diseases and Biology, University of Life Sciences, 20-950 Lublin, Poland e-mail: leszek.guz@up.lublin.pl

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salmoninarum, most of which target the *msa* gene for amplification (McIntosh et al. 1996, Miriam et al. 1997, Vardić et al. 2007). A nested PCR detecting the 57-kDa major soluble antigen (msa, p57) is recommended by the OIE for screening purposes (Chase and Pascho 1998, OIE 2006, Arnason et al. 2013, Pękala-Safińska et al. 2017).

The aim of this work was to confirm the presence of *R. salmoninarum* in rainbow trout with clinical signs of BKD by PCR assay. Additionally, sequence analysis of the PCR product was performed to characterize the bacterium.

The diseased, moribund rainbow trout (n = 32) (60 - 150 g) were transferred alive to the Department of Fish Diseases and Biology, Faculty of Veterinary Medicine, University of Life Sciences in Lublin, Poland, during the fall of 2016. All the fish were measured, weighed, and dissected. Samples of liver, spleen, and kidney were collected aseptically and samples were streaked on selective kidney disease medium (SKDM) and incubated at 16°C for 6 weeks. The characteristic lesions and growth of *R. salmoninarum* were verified by Gram-staining smears. The predominant colonies that grew on the agar plates were purified and stored at -80°C after supplementation with 20% (v/v) glycerol.

R. salmoninarum was identified molecularly by isolating total DNA from freshly grown culture. DNA extraction was conducted using a Genomic Mini Kit according to the manufacturer's instruction (A&A Biotechnology, Gdynia, Poland). The DNA was eluted with 200 μ L of TB Buffer heated to 70°C. The samples were run immediately following elution.

The colonies were identified by amplifying a 501 bp region of the gene encoding a 57 kDa soluble protein (p57) using primers Rs1 (5'-CAAGGTGAAGGGAATTCTTCCACT-3') and Rs2 (5'-GACGGCAATGTCCGTTCCCGGTTT-3') (Brown et al. 1994). PCR reactions were performed in a thermal cycler (MJ-Mini, Bio-Rad, Warsaw, Poland) with the following amplification profile: initial denaturation at 94°C for 10 min, followed by 30 cycles at 94°C denaturing for 1 min, 48°C annealing for 1 min, 72°C extending for 1 min, and a final extension at 72°C for 5 min. To test the specificity and cross-reactivity of the PCR test, two other bacteria, Aeromonas salmonicida and Yersinia ruckeri, were tested. The PCR products were analyzed at a constant voltage of 7V cm⁻¹ on 1% agarose gel containing $(0.005 \,\mu L \,m L^{-1})$ Midori Green Advanced DNA Stain (Nippon Genetics, Tokyo, Japan) and DNA Marker 1 (A&A Biotechnology, Gdynia, Poland). DNA fragments of the expected length were purified using a Gel-Out Extraction Kit (A&A Biotechnology, Gdynia, Poland) according to the manufacturer's instructions and subjected to direct sequencing at a DNA sequencing core facility (Genomed). Bacterial identity was deduced by a BLAST search to ascertain its closest related sequences.

R. salmoninarum is a well-known intracellular pathogen and is capable of surviving and multiplying cytoplasm of infected macrophages in the (Gutenberger 1993, McIntosh et al. 1996) and salmonid eggs (Brown et al. 1994). In the present study, the BKD diagnosis was made by macroscopic, light microscopy, bacterial culture, and PCR (Table 1). The fish exhibited weak external clinical signs of bacterial disease, such as loss of appetite, lethargy, dark coloration, exophthalmia, and distended abdomen (Fig. 1A). The most obvious internal signs were swollen kidney and/or liver and grevish nodules in the kidney (Fig. 1B, Table 1). Samples were collected from the kidneys by scraping them with a scalpel blade. The material collected was examined as

Table 1

Prevalence and testing for bacterial kidney disease (BKD) in rainbow trout (O. mykiss)

Examined fish	Prevalence	Light microscopy	Positive culture fish pooled kidney samples	Positive PCR
With kidney lesions	20/32 (62.5%)	18/32 (56.2%)	1/1	1/1
Without kidney lesions	12/32 (37.5%)	8/32 (25.0 %)	1/1	1/1

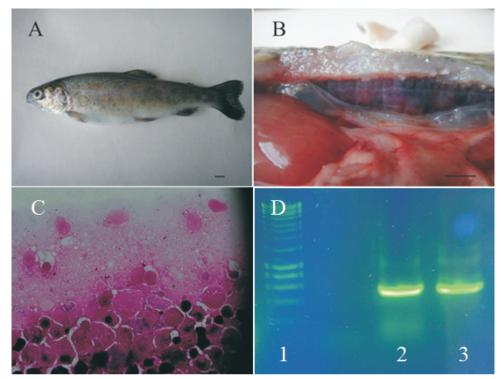


Figure 1. Rainbow trout with bacterial kidney diseases. Characteristic clinical signs observed: distended abdomen (A); enlarged kidney, grayish in color, with white nodular lesions (B); *R. salmoninarum* in kidney macrophages – Gram staining (C). Electrophoregram of 1.5% agarose gel stained with Midori Green Advanced DNA Stain showing amplification of the gene encoding a 57 kDa soluble protein (primers Rs1/Rs2). Lane 1, DNA Marker 1. Lane 2, *R. salmoninarum* DNA isolated from kidney with lesions, and 3, *R. salmoninarum* DNA isolated from kidney with lesions (D).

a wet-mount preparation using a compound microscope, and *R. salmoninarum* was noted in the macrophages (Fig. 1C, Table 1). Isolated colonies of bacteria were creamy, shiny, smooth, round, raised, and 2 mm in size on SKDM agar after 6 weeks of incubation at 16°C. Cells were Gram-positive and non-motile diplobacilli.

A fragment of the appropriate size (501 pb) was amplified from DNA extracted from *R*. *salmoninarum* grown in agar culture (Fig. 1D, Table 1). All positions containing missing data were eliminated. There was a total of 401 positions in the final dataset. Evolutionary analysis was conducted in the MEGA6 program. The sequence p57 gene of the clinical isolate of *R. salmoninarum* was deposited in the National Centre for Biotechnology Information Gene Bank database under accession number KU863577.

This study described a simple, DNA-based identification method of *R. salmoninarum* isolated from diseased rainbow trout in Poland.

ORCID ID

Leszek Guz (D) http://orcid.org/0000-0003-4978-5577

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