Reaching Fish Farms Through AquaFish CRSP Technology Transfer: Elimination of MT from Intensive Masculinization Systems Using Bacterial Degradation

Mitigating Negative Environmental Impacts/Study/09MNE07UA

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OBJECTIVES

The general objective of this project is to transfer A&F CRSP-generated technology to the private sector, ensuring the development of sustainable tilapia aquaculture in Southeastern Mexico and Central America. The specific objectives are:

- 1. Evaluate combination of bacteria that best degrade MT in tilapia masculinization systems.
- 2. Transfer MT degrading technology using bacteria to a large hatchery.
- 3. Develop written materials for the dissemination of these protocols and train tilapia farmers in the use bacterial bio-flocks as MT-degrading entities and probiotics.
- 4. Continue evaluation of bacterial combinations that improve degradation of MT and that can serve as probiotics

Introduction

Aquaculture industry considers different technologies in order to improve food production for the increasing human needs. One alternative for male monosexual population production is the utilization of synthetic steroids. The aquaculture industry commonly uses steroids including the 17 $\alpha-$ methyltestosterone (MT), miberolone, fluoxymesterone, nortestosterone acetate, 17 $\alpha-$ methyldihydrotestosterone and norgestrel. MT is the most common form in worldwide use (Hurtado, 2005; López et al., 2007). Specifically, MT is a strong androgen widely used for tilapia masculinization. Around 90% of MT used during masculinization treatments is degraded during the first 24 h and only 1% is remaining 3 weeks later in the fish body (Raisz et al., 2006; Contreras, 2001).

Other natural and synthetic chemicals present in the environment like 17α - estradiol, estroil, estrone, progesterone, testosterone and phytoestrogens mimic hormonal actions. These chemicals actually are considered a worldwide issue because they could disturb sexual characteristics in fish, amphibians, reptiles and mammals (Toft et al., 2004; Argemi et al., 2005; Bracho, 2007). These compounds are consider Endocrine Disrupting Chemicals (or Compounds) or EDCs because are exogenous chemicals which could promote endocrine system alterations and adverse effects on individuals, broods and populations health (Anonymous, 2002). This phenomenon includes reproductive system dysfunction, neoplasia, malformations, neurotoxicity, and decreasing immune response, promoting difficulties to detect possible causes and effects (Swan et al., 2003; Rivas et al., 2004).

In order to decrease environmental impacts of these compounds, Best Management Practices for aquaculture include the use of Recirculation Aquaculture Systems, (RAS). RAS decreases the total waste production, saving water and decreasing waste in the fish culture systems improving water quality for reutilization. Another method for chemical elimination of hormones is the use of granular activated charcoal to absorb the compounds. Finally, phytosteroids have been tested from vegetal

origin; these compounds mimic fish steroids and sometimes have similar impacts on gonadal development (Contreras, 2001; Campos, 2004).

Microbiological degradation processes have been widely studied in recent years; microorganisms have biodegrading features for many pollutants. Microorganisms accumulate, transform or degrade different pollutants including petroleum hydrocarbons. In order to decrease risks for human and environment health the main scope for this study was to determine bacterial strains able to degrade $17 \, \alpha$ – methyltestosterone (MT). These bacteria were isolated from a biofilm obtained from a RAS used for fish chemical sex reversion.

MATERIALS AND METHODS

Bacterial strains selection

Strains were obtained from the Microbiology Laboratory at the División Académica de Ciencias Biológicas at UJAT. These strains were originally isolated from biofilms obtained from biological filters used in a tilapia (*Oreochormis niloticus*) masculinization system. Strains of *Pseudomonas aeruginosa, Pseudomonas fluorescens, Bacillus cereus* and *Bacillus subtilis* were selected according to previous experimental data showing potential to degrade MT under culture conditions where these steroids were used as carbon and energy source.

Bacterial strains adaptation

The four bacteria strains were subjected to adaptation to MT as main source of carbon and energy during 10 days-period, previous to experimental procedure. Each bacterial strain was inoculated in a glass beaker containing 100 mL of liquid mineral culture media containing 40 mg of MT (Sigma-Aldrich, Inc.). according to the procedure presented by Pérez *et al.*, (2006), Yamanaka *et al.*, (2007) and Montpas *et al.*, (1997) respectively (Table 1). All treatments were evaluated in triplicate, including control treatments containing MT and no bacteria, and without MT and bacteria present. Cultures were incubated at 30 ± 1 °C in an orbital shaker incubator 175 rpm, (PolyScience®) (Perez *et al.*, 2006; Yamanaka *et al.*, 2007).

During the experimental phase, the number of viable cells was estimated by conventional plate counting technique. MT-enriched liquid mineral medium was used adding 15 g per liter of bacteriological agar (BIOXON®). Each sample was serially diluted (1:10) in a sterile 0.85% saline solution to obtain a 10^{-5} final dilution. For each diluted sample, $100 \, \mu$ l were inoculated in the center of each Petri dish and extended using a sterile Drigalski spreader. Inverted Petri plates were incubated for 48 h to 30 °C \pm 0.5 °C and the count were estimated using a Darkfield Quebec colony counter (Bracho *et al.*, 2004).

Table 1.- Chemical composition of the culture media used g/L. (-) indicated the absence of the mineral compound.

Minerals	Bacillus	Pseudomonas	Modificated
NH4Cl	-	0.1	0.005
$MgSO_4$. $7H_2O$	0.1	0.05	0.1
K_2HPO_4 . $3H_2O$	6.3	0.1	6.3
KH_2PO_4 . $3H_2O$	1.83	0.05	1.83
FeSO ₄ . 7H ₂ O	0.1	0.001	0.1
CaCl ₂	0.1	0.001	0.1
(NH ₄)2SO ₄	1	-	1
KCl	-	0.01	0.01

Determination of MT to evaluate degradation capacity

The experiment had 4 treatments with controls: 1) *Pseudomonas aeruginosa and Pseudomonas fluorescens;* 2) *Bacillus subtilis and Bacillus cereus;* 3) *Pseudomonas aeruginosa and Bacillus cereus; and* 4) *Pseudomonas aeruginosa, Pseudomonas fluorescens, Bacillus subtilis and Bacillus cereus* mineral culture media and MT, without bacteria; and 2) A flask with mineral culture media without MT and with the corresponding bacteria according to the treatment. Each treatment was performed in triplicate, but the treatment control had only one experimental unit.

Each treatment was inoculated with 1mL of a bacterial suspension equal to the Turbidimetry McFarland scale of 0,5 and corresponding to a population of 1,5 x 10^8 UFC/ 100 mL. The inoculum was incubated at 30° C \pm 1 $^{\circ}$ C with rotary agitation at 175 rpm (PolyScience®) during 22 days.

The number of microorganisms in each treatment was measured daily through viable counting with the superficial extension technique. The media used in this phase was a mineral media enriched with MT with the addition of 15 g of bacteriological agar (BIOXON®) by liter of media. Serial dilutions were made from 1 mL of sample in 9 mL of sterile saline solution at 0,85% (10-1) up to a dilution of 10^6 . From each dilution $100~\mu L$ was inoculated in the center of a Petri dish, extending over the media surface with a sterile Drigalski dipstick. The Petri dishes were placed in an inverted position for incubation at $30~^{\circ}C~\pm 0.5~^{\circ}C$. The dish readings were performed at 48~h using a Darkfield Quebec counter.

Elisa test or Immunoassay method (r-biopham)

Samples from each treatment were collected in order to determine MT concentrations at 0, 2, 6, 10, 16 and 20 days. All samples were stored in a freezer (-20 °C) in Eppendorf tubes until analysis. Samples were diluted in methanol at 40% up to a dilution of 10⁶, in order to determine the maximum and minimum limits of MT in the calibration curve of the kit.

The quantitative measurement of MT was performed by the Elisa test or Immunoassay method (r-biopham): 50 μ l of sample of each treatment were inoculated by duplicate. Each analysis had six standards in order to obtain the calibration curve of MT. Once the samples inoculated, 50 μ l of the conjugate and 50 μ l of the antimetiltestosterone (antibody) solution was added. Incubation was performed for 12-16 h at 2-8 ° C. After incubation two distilled water rinses were completed, using a semiautomatic washer.

Following the wash step, $50~\mu l$ of substrate and $50~\mu l$ of chromogen were pipetted into each well and they were incubated in darkness for 30~minutes. Finally $100~\mu l$ of stop solution was added to each well and we proceeded to the reading with a spectrophotometer at an absorbance of 450~nm. The obtained data was analyzed with the software RIDA® SOFT Win in order to obtain the MT concentration in each sample. After obtaining the raw data the number was multiplied by the dilution factor.

RESULTS FOR OBJECTIVES 1 AND 4

The treatment which showed the greatest growth after 24 h and a greater number of cells was *P. aeruginosa* and *P. fluorescens*. With a growth curve of 30 days, the treatment with *B. cereus* and *B. subtilis* showed growth at 48 h with a duration of 22 days. Figure 1 shows the growth curves of the two finalized treatments.

DISCUSSION

Objectives 1 and 4

The results obtained with respect to the two treatments demonstrated bacteria growth in MT media (40 mg/100 ml) as the only carbon source. Yomokito et al. (2004) and Ke et al. (2007), observed bacteria growth and biodegradation from 100 mg/ 100 ml of 17 β – estradiol and 17 α -methylestradiol and 3-methoxy-17b-hydroxy-1,3,5 (10),8(9)-d-4-estren (MHE) respectively. In both studies the strains came from activated sludge from a treatment plant with estrogen residues. Jiang et al. (2010), isolated *B. subtilis* as degrading of 17 β – estradiol at a concentration of 1 mg/100 mL which differs with our project. Due to this the strain was isolated in a mineral media with 50 mg/100 mL of

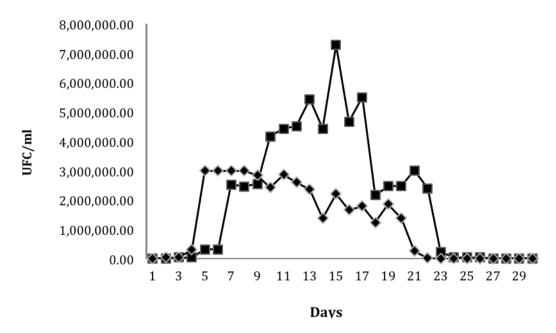


Figure 1. Average value of the microbial growth kinetics in mineral media enriched with 17α -methyltestosterone (MT), the values correspond to mean of triplicate measurements. Pseudomonas aeruginosa and Pseudomonas fluorescens; * Bacillus cereus and Bacillus subtilis

estrogen plus 15 mL of activated sludge from sewage of a manufacturer of oral contraceptives in Zhejiang, China, performing successive cycles and changing the concentration up to 1 mg of estrogen.

Steroid biotransformations are important in the manufacture of drugs and hormones; anti-inflammatories, diuretics, steroids, anti-androgenic and anti-cancer agents and with some other applications (Moschet *el al.*, 2009; Walaa *et al.*, 2009). The steroids biotransformation by bacteria and fungi have been suggested by Smith et al. (1998) and Walaa et al. (2009) investigating the transformation of progesterone by *Apiocrea chrysosperma*, *Botryosphaerica obtusa*, *Bacillus subtilis* and *Aspergillus niger*. Phelps and Popma (2000) mentioned that bacteria and fungi could be responsible of steroids metabolization and they assumed the existence of an androgen microbial degradation in the biofilters of the recirculation systems.

It has been proved that the genera *Nocardia, Bacillus subtilis, Mycobacterirum, Bacterium* and *Arthrobacter* act as estrone degrading. Lee and Liu, (2001), Weber et al. (2007); Shulan et al. (2007) and Jiang et al. (2010), have studied the steroids microbial degradation identifying: *Achromobacter xylosoxidans, Ralstonia sp, Raoultella thalpophilim* and *B. subtilis* as responsible for estrogens biodegradation, although the conditions of our experiment were different, the percentages of biodegradation were between 80 to 85%, which are similar to the above investigations. The *Pseudomonas sp.* and *Bacillus sp.* have been used in the biodegradation of hydrocarbons, obtaining a good elimination rate, thus it is recommended that both microorganisms can be used for biodegradation of these compounds (Mohamad et al., 2004). Duarte and Gomes (2000) mentioned that degradation increased with a bacteria conjugate due to better degradation capacity with a wider range of action.

Objective 2. Transfer MT degrading technology using bacteria to a large hatchery

Work on this objective is still underway. During the extreme flooding in mid-2011, the hatchery at our demonstration farm partner was lost and the biofiltration system was destroyed. The operation has been moved to a second location that is significantly higher and should be above the flood stage that was breached in each of the last two years. We are collecting data in late 2011 and early 2012.

Objective 3. Develop written materials for the dissemination of these protocols and train tilapia

farmers in the use bacterial bio-flocks as MT-degrading entities and probiotics.

Work on this objective is nearing completion. A biofloc workshop was held in September 2010 at UJAT for hatchery managers who would be working with biofloc systems. Some were interested to use bioflocs to remove residual MT and other more specifically to produce bioflocs as a feed source. A Power Point presentation has been prepared and shared with the farmers.

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