Biofouling of FeNP-Coated SWRO Membranes with Bacteria Isolated after Pre-Treatment in the Sea of Cortez

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Abstract: Commercial seawater reverse osmosis (SWRO) membranes were coated with iron nanoparticles (FeNPs) and biofouled with a bacterium strain isolated from the Sea of Cortez, Mexico. This strain was selected and characterized, as it was the only cultivable strain in pretreated seawater. Molecular identification of the strain showed that it belongs to Bacillus halotolerans MCC1. This strain was Gram positive with spore production, and was susceptible to Fe^{2+} toxicity with a minimum inhibitory concentration of 1.8 g L^{-1}. Its biofouling potential on both uncoated and FeNP coated reverse osmosis (RO) membranes was measured via biofilm layer thickness, total cell count, optical density and organic matter. The FeNP-coated RO membrane presented a significant reduction in biofilm cake layer thickness (>90%), total cells (>67%), optical density (>42%) and organic matter (>92%) with respect to an uncoated commercial membrane. Thus, Bacillus halotolerans MCC1 shows great potential to biofoul RO membranes as it can pass through ultrafiltration membranes due to its spore producing ability; nonetheless, FeNP-coated membranes represent a potential alternative to mitigate RO membrane biofouling.

Keywords: Bacillus halotolerans; biofouling; reverse osmosis; model biofoulant; iron nanoparticles

1. Introduction

Halotolerant bacteria are of great biotechnological importance for industry, as they are easy to grow under limited nutritional requirements. Moreover, their tolerance to high salt concentrations minimizes laboratory contamination risks [1]. Several researchers investigating halotolerant bacteria have focused on their capacity to produce antibiotics and enzymes [2], while others have used them to design microbial inoculants to increase the salinity tolerance of crops [3]. For example, in India, a halotolerant bacillus has been used for the biosynthesis of enzymes (L-Glutaminase) in bioethanol production [4]. In Algerian wetland ecosystems, a study found that halotolerant strains belonging to the genera Haloferax, Halococcus and Haloarcula showed a high production of molecules with important biotechnological applications, such as in the coastal agriculture, pharmaceutical and environmental fields [5]. In Spain, the dominant halotolerant genera isolated from the Bras del Port salt basins included Salinivibrio, Pseudomonas, Alteromonas, Alcaligenes, Acinetobacter and Flavobacterium. These halotolerant microorganisms have exhibited potential for several applications in the food, pharmaceutical, medical and environmental fields [6].

In the seawater reverse osmosis (RO) desalination industry, halotolerant bacteria have gained great interest because they are involved in membrane biofouling, one of the biggest problems faced
by membrane-based desalination [7]. Biofouling is an operational problem in SWRO that can reduce production efficiency, leading to severe economic impacts [8]. Biofouling involves microorganism attachment and growth on the membrane surface, and is very difficult to treat due to the ability of microorganisms to multiply even if most are eliminated in pretreatment [9]. As biomass accumulates in the feed channel of membrane modules, this leads to technical problems (increases in the working pressure or reduction of permeate flux, as well as the need of more frequent chemical cleaning, among others), decreasing the process productivity, reducing the life span of membranes and increasing process costs [10,11]. To reduce the impact of biofouling problems, some research efforts have focused on coating the membrane surface with various nanoparticles (NPs) that are known to have antimicrobial properties, such as Ag, CuO, ZnO and TiO$_2$ [12–15]. In addition, NPs have been used as antifouling coatings in different marine industrial environments in order to prevent biofouling [16]. However, these NPs are expensive and difficult to synthetize; therefore, their use as a membrane coating increases desalination costs. Iron nanoparticles (FeNPs) are a promising coating agent, as they cost less than other NPs, are easy to synthesize and have a biocide effect by promoting Fenton or Fenton-like reactions, producing oxidative stress by generating reactive oxygen species (ROS), disrupting lipids, proteins and DNA, and eventually causing bacterial death [17–19].

Interestingly, most biofouling studies of NP coatings have been carried out with strains that are not native from seawater, such as Escherichia coli and Pseudomonas aeruginosa [20,21]. Those studies are not very useful for the desalination industry, since those strains are not ones that commonly cause RO membrane biofouling. Hence, the studying and understanding of seawater halotolerant bacteria is paramount for the RO desalination industry due to their negative effect on membrane durability and productivity. Moreover, the shoreline of the Sea of Cortez is home to 32 RO desalination plants with capacities ranging from 100 to 17,280 m$^3$ d$^{-1}$ [22] and one under construction [23]. Further, the number of RO desalination plants in that region has been increasing in recent years (up by 68% in three years) [22] due to frequent water scarcity. For these reasons, the aim of this paper is to study the effect of FeNP coating of RO membranes on the extent of biofouling caused by bacteria native to the Sea of Cortez. This is achieved by isolating and characterizing the only bacterium strain found after a typical pretreatment of seawater sampled from the Sea of Cortez, Mexico (Bacillus halotolerans MCC1), and using it as a model biofoulant.

### 2. Materials and Methods

#### 2.1. Sampling

Seawater samples were obtained from the Sea of Cortez, at Cochiorit beach near Empalme, Sonora, Mexico at a depth of 4 m and at a distance of 3.76 km from the coast (27° 53’ 1.28” N, 110° 47’ 5.56” W) (Figure 1). The samples were collected following the procedure detailed in the field manual for water quality sampling [24]. Five samples were taken throughout the year (spring, summer, autumn and two in winter) and at different times of the day. The physicochemical parameters measured at the time of sampling are shown in Table 1. Three hour after sampling, the collected water was pretreated by 5 µm cartridge filtration and 0.1 µm ultrafiltration (PURIKOR, PKUF-M) with a constant working pressure of 0.2 MPa and a permeate flow rate of 3.7 L min$^{-1}$, as a representative analogue of the pretreatment that takes place in typical large-scale desalination plants [25].

| Temperature ($^\circ$C) | Electrical Conductivity (µS cm$^{-1}$) | Total Dissolved Solids (mg L$^{-1}$) | Salt Fraction (%) | Dissolved Oxygen (mg L$^{-1}$) | pH      |
|-------------------------|---------------------------------------|-------------------------------------|------------------|-----------------------------|---------|
| 24.1 ± 0.3              | 48,100 ± 350                          | 34.99 ± 0.1                         | 3.56 ± 0.0       | 8.42 ± 0.1                  | 8.1 ± 0.2 |
2.2. Bacterial Isolation and Molecular Identification

Petri dishes containing nutrient agar (NA) supplemented with sterilized pretreated seawater (34.99 g L\(^{-1}\) of salinity) were used as a culture medium. The pretreated seawater was previously sterilized in an autoclave (Felisa FE-399), for 15 min and 0.103 MPa at 121 °C. Pretreated seawater was used to provide nutritional conditions for bacterial growth, similar to those observed in their natural environment (Table 1).

To determine the cultivable bacterial population and diversity of the unsterilized pretreated seawater, the serial dilution (1:10) method was used, preparing dilutions up to 10\(^{-3}\)\[^{[26]}\]. Finally, 1 mL of each dilution was dispersed, onto the NA prepared in the Petri dishes, and incubated for 24 h at 28 °C\[^{[27]}\]. The bacterial isolation was done by triplicate.

Macro and microscopic bacterial characterization, spore production and Gram staining were carried out using the Wirtz–Conklin method\[^{[28]}\] and the Hycel stain kit (Hycel, Cat 541), respectively. All obtained isolates were preserved in the Native Endophytes and Soil Microorganisms Collection (COLMENA)\[^{[29]}\].

Molecular identification was carried out for the isolated bacterium strain that grew in the previously mentioned Petri dishes with culture medium, which was isolated from the pretreated seawater. Genomic DNA was extracted using the methodology of Reader and Broda\[^{[30]}\]. Bacteria molecular identification was carried out amplifying the 16S ribosomal gene (16S rRNA)\[^{[31]}\]. The PCR protocol was carried out according to the procedure described by Villa-Rodriguez et al\[^{[32]}\]. The DNA sequences were processed using the FinchTV 1.4.0 software package (Geospiza Inc., Denver, CO, USA). The phylogenetic tree was created by Neighbor Joining methodology\[^{[31]}\] using MEGA 7.0.

2.3. Minimum Inhibitory Fe\(^{+2}\) Concentration

The minimum inhibitory concentration of Fe\(^{+2}\) was determined by the inoculation of 1 × 10\(^4\) colony forming units (CFU) onto Petri dishes containing NA supplemented with sterile pretreated seawater. Further, the culture medium was supplemented with Fe\(^{+2}\), at concentrations of 1.0, 1.2, 1.4,
1.6, 1.8 and 2 g L$^{-1}$. The inoculated Petri dishes, using three independent replicates, were incubated at 28 °C for six days. The bacterial population in each Petri dish was determined by CFU count every 24 h.

### 2.4. Accelerated Biofouling Test on RO Membranes

Accelerated biofouling on two commercial RO membranes (Dow Filmtec SW30HR) was carried out using a high concentration of the studied bacterial strain and providing it with sufficient nutrients in the feed water. One of these membranes was coated with the minimum inhibitory concentration of FeNPs (FeNP membrane) via immersion, ensuring the NP dispersion by sonication for 1 h, following the methodology of Armendariz et al. [18]. The membrane was immersed into fresh DI water for 10 min to activate the carboxyl groups and then the membrane was dipped into the aqueous suspension of FeNPs for 24 h, followed by rinsing with an ultrasonic bath for 5 min. The other membrane was not coated (uncoated membrane) and used as blank for reference and comparison. The FeNPs were synthetized and characterized [18,19] according to the method of Baltazar et al. [33] and Arancibia-Miranda et al. [34]. The chemical structures and morphology of the FeNPs were characterized by X-ray Diffraction (XRD) using a Shimadzu XRD-6000 diffractometer (Kyoto City, Japan) and Scanning Electron Microscopy (SEM) using a Nova NanoSEM-200 (FEI Company, Hillsboro, OR, USA).

The bacterial growth on the RO membrane surface took place in CF042 crossflow membrane cells (Sterlitech Corp., Kent, WA, USA) (Figure 2). The feed water comprised sterile pretreated seawater and a high amount (10$^9$ CFU mL$^{-1}$) of the studied strain. In addition, a sterile nutrient broth was added daily to the feed water, to guarantee optimal bacterial growth. The operating pressure was 6.38 MPa during the test, which ran for 90 h at an average temperature of 30 °C, while the pH remained in a range of 6–8. A low crossflow velocity of 0.15 m s$^{-1}$ was used in order to minimize shear stress and encourage a high biofouling rate. After the accelerated biofouling test, biofouled membranes were analyzed by scraping 1 cm$^2$ of biofilm. Then, the optical density of the scraped biofilm was determined by spectrophotometry, the biofilm cake layer was measured using inverted microscopy, and a Neubauer chamber using an optical microscope in 100× was used to carry out the total cell count in the biofilm. The amount of organic matter was determined by the catalytic combustion method [10].

**Figure 2.** Representative diagram of the experimental setup for accelerated biofouling of RO membranes.
2.5. Statistical Analysis

An analysis of variance of simple classification based on a linear model of fixed effects was performed for the CFUs found over time, and the statistical differences between the mean values were determined by the Tukey method for \( p \leq 0.001 \). For the biofilm cake layer, optical density (OD), total cell count (TCC) and organic matter (OM), the theoretical t-student probability distribution for quantitative continuous variables for significance levels of \( p \leq 0.001 \) was used. A simple linear regression analysis based on a linear model of fixed effects was performed. The degree of goodness of fit between the \( \text{Fe}^{2+} \) concentration and the final CFU count was determined. From these equations, the optimal inhibitory concentration of bacterial growth was determined using an ordinary least squares method. The software STATISTIC version 8.5 (StatSoft, Tulsa, OK, USA) was used for the statistical analyses.

3. Results and Discussion

3.1. Bacteria Growth, Isolation and Molecular Identification

After 24 h of incubation, only one bacterial isolate (MCC1) grew for each repetition in direct pretreated seawater. No growth of CFUs was observed for the \( 10^{-3}, 10^{-2}, 10^{-1} \) dilutions. This bacterial strain showed medium size colonies (5.0 ± 1.0 mm), convex, whitish such as wax, rounded edges, fusiform and circular shape (Figure 3a). In addition, the Gram stain showed a Gram positive cell with a size of 2.8 ± 0.2 µm (Figure 3b), and spore size of 18.0 ± 0.0 nm (Figure 3c). As the ultrafiltration membrane nominal pore size of 0.1 µm is smaller than the typical sizes of bacteria, this indicates that only spores could pass through the filter and grow on the Petri dishes, thus originating the CFU found [35]. The ultrafiltration membrane did not show integrity problems during the lapse of the testing.

![Figure 3](image-url) Figure 3. Macro and microscopic traits of the *B. halotolerans* MCC1 strain. (a) Bacterial colonies on nutritive agar; (b) bacterial Gram stain; (c) bacterial sporulation.

Molecular identification showed that this particular strain belongs to *Bacillus halotolerans* MCC1. This result confirms the macro and microscopic traits observed for the strain. Its Gram staining (Figure 3b) showing a Gram positive *Bacillus* strain, agrees with reports that most sea water *Bacillus* are Gram positive [36]. Moreover, this strain belongs to the specie *B. halotolerans*, which agrees with its source of isolation (ultrafiltered seawater). This finding indicates that the strain MCC1 is able to tolerate high electrical conductivity (48,100 ± 350 µS cm\(^{-1}\)) and is not retained by the ultrafiltration process, due to its ability to produce spores that are smaller than the UF membrane pore size (as seen in Figure 3c). Spore production is an adaptive strategy to compensate the osmotic stress generated by high salt concentrations in the environment [37], and is an ability that not all *Bacillus* possess [38,39]. These bacterial traits could be highly associated with the role of the strain MCC1 on biofouling RO membranes.
3.2. Minimum Inhibitory Fe\(^{+2}\) Concentration

Figure 4 shows the rate of bacterial growth in NA supplemented with various concentrations of Fe\(^{+2}\). The data shows that the biocidal effect of Fe\(^{+2}\) was greater from 24 to 48 h, at any concentration used. After 72 h, large bacterial populations were observed for the range of 1.0 to 1.6 g L\(^{-1}\) Fe\(^{+2}\). It is clear that Bacillus halotolerans MCC1 can easily grow in Fe\(^{+2}\) concentrations of 1 to 1.4 g L\(^{-1}\), as Fe is an essential element for life [40]. However, from 1.6 g L\(^{-1}\) the strain MCC1 is not able to maintain its optimal growth rate within the first 48 h. From that moment, the bacterium shows adaptive strategies (Fe\(^{+2}\) resistance) to be able to survive, because at 72 h of incubation it is possible to observe CFUs. This could be attributed to MCC1 having adaptive abilities under extreme conditions, also evidenced by the sporulation capacity reported in this work (Figure 3c) [37]. Finally, the biocidal effect of Fe\(^{+2}\) at higher concentrations (1.8 to 2 g L\(^{-1}\)) is greater, since there was negligible bacterial growth.

![Figure 4](image_url)

**Figure 4.** Effect of Fe\(^{+2}\) concentration on the growth rate of B. halotolerans MCC1.

A significant decrease in the final bacterial population for the strain MCC1 was observed with increasing Fe\(^{+2}\) concentration. There was a negative correlation with a highly significant (95% confidence) dependence between these variables \((r = 0.95, p < 0.001)\). The degree of goodness of fit was high \((R^2 = 0.91)\) for a linear regression model (Figure 5). The greatest decrease in CFUs for MCC1 was observed when the Fe\(^{+2}\) concentration was increased from 1.4 to 1.6 g L\(^{-1}\), presenting 50% of the total inhibition by Fe\(^{+2}\). At Fe\(^{+2}\) concentrations of 1.8 g L\(^{-1}\) and higher, no bacterial growth was observed. Despite the previously mentioned capability of Bacillus halotolerans MCC1 to adapt to extreme conditions, there is a clear adaptability limit after which it cannot survive. Therefore, the minimum inhibitory Fe\(^{+2}\) concentration was 1.8 g L\(^{-1}\), as the data suggests that this bacterium cannot grow at higher concentrations. At concentrations above that level, the accumulation of Fe\(^{+2}\) can stop cell growth or cause death due to DNA damage [40].

![Figure 5](image_url)

**Figure 5.** Final bacterial population of the strain MCC1 at different Fe\(^{+2}\) concentrations. \(C\) (Fe\(^{+2}\)): concentration of Fe\(^{+2}\). \(R\): correlation coefficient. \(p\): probability. \(R^2\): coefficient of determination without adjustment. **: highly significant.
3.3. Accelerated Biofouling Test on RO Membranes

The SEM image (Figure 6) depicts the FeNPs, showing a mean particle size around 14.6 ± 1.2 nm. Figure 7 shows the XRD peaks for the FeNPs. The crystalline structure of the FeNPs was confirmed by XRD analysis (2θ = 44.5°, 65.0° and 82.4°) [41]. The accelerated biofouling test showed that the bacteria can exhibit growth under conditions of very high salinity, in the range of 3%–8% salt concentration by weight (see Figure 8). This result suggests that the bacteria indeed exhibits sporulation, as endospore production allows the bacteria to survive in extreme environmental conditions [42,43]. In addition, the ability of Bacillus halotolerans MCC1 to biofoul RO membranes was also demonstrated in this test. A biofilm cake layer was formed on the membrane surface during the experiment, resulting in a decrease in permeate flux. This strongly suggests that this bacterium strain can be useful as a model biofoulant, especially for desalination plants in the Sea of Cortez. Further, as can be seen in Figure 8c, the FeNP coated membrane showed significantly less biofouling compared to the uncoated membrane (Figure 8b), evidencing the antimicrobial effect of FeNPs already demonstrated in the medical field [44].

Figure 6. SEM image of FeNPs.

Figure 7. XRD of the FeNPs showing the peaks associated with the Fe (red arrows) and iron oxides (black arrows) present on the FeNPs.
The FeNP coating reduced the biofilm thickness by 90% (Figure 9a), and the bacterial density on the membrane showed a highly significant reduction of 67% compared to the uncoated membrane, which can be attributed to the FeNPs (see Figure 9b). The optical density also showed a highly significant reduction related to the FeNP coating, as it was 42% lower than for the uncoated membrane (Figure 9c). In addition, the organic matter content in the biofilm was lower (about 92%) for the FeNP-coated membrane compared to the uncoated membrane (Figure 9d). All these results can be related to the *Bacillus halotolerans* MCC1 being unable to produce iron-chelating metabolites (siderophores), which limits its ability to tolerate and grow under high Fe concentration [45]. Therefore, the direct exposure of this strain to the FeNPs located on the membrane affects its reproduction and growth rates. This can also be associated with the oxidative stress that has been shown to be caused by the generation of ROS by FeNPs, leading to damage of cellular DNA, lipids and proteins, causing bacterial death [46,47].

The existence of this latter mechanism also suggests that the Fenton reaction facilitated by the FeNPs can cause the death of different bacteria, not only of *Bacillus halotolerans* MCC1 [19,46], since the FeNPs in the presence of dissolved oxygen and water form hydrogen peroxide and Fe$^{+2}$ ions. The hydrogen peroxide and Fe$^{+2}$ then, in turn, initiate the Fenton reaction cycle, which produces several ROSs with strong biocidal effects. However, in this reaction the FeNPs also are oxidized to Fe$^{+2}$, which eventually limits the FeNPs’ effectiveness as a biocide due to the leaching of Fe$^{+2}$ ions [48]. Nevertheless, the FeNPs can still hinder biofilm growth and decrease biofouling problems in desalination plants.
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