

The capacity of *Listeria monocytogenes* mutants with in-frame deletions in putative ATP-binding cassette transporters to form biofilms and comparison with the wild type

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Abstract

Listeria monocytogenes (Lm) is a food-borne pathogen responsible for human listeriosis, an invasive infection with high mortality rates. Lm has developed efficient strategies for survival under stress conditions such as starvation and wide variations in temperature, pH, and osmolarity. Therefore, Lm can survive in food under multiple stress conditions. Detailed studies to determine the mode of action of this pathogen for survival under stress conditions are important to control Lm in food. It has been shown that genes encoding for ATP-binding cassette (ABC) transporters are induced in Lm in food, in particular under stress conditions. Previous studies showed that these genes are involved in sensitivity to nisin, acids, and salt. The aim of this study was to determine the involvement of some ABC transporters in biofilm formation. Therefore, deletion mutants of ABC transporter genes (LMOf2365_1875 and LMOf2365_1877) were created in Lm F2365, and then were compared to the wild type for their capacity to form biofilms. Lm strain F2365 was chosen as reference since the genome is fully sequenced and furthermore this strain is particularly involved in food-borne outbreaks of listeriosis. Our results showed that ALMOf2365_1875 had an increased capacity to form biofilms compared to the wild type, indicating that LMOf2365 1875 negatively regulates biofilm formation. A deeper knowledge on the ability to form biofilms in these mutants may help in the development of intervention strategies to control Lm in food and in the environment.

Introduction

Listeria monocytogenes (Lm) can cause listeriosis, a severe invasive disease with high hospitalization (>90%) and mortality rates (20 to 30%), especially in immunocompromised people, elderly individuals, and pregnant women. Therefore listeriosis is an infection of great concern to public health despite its low incidence (0.4 cases per 100,000 population) (EFSA, 2011). Lm is a food-borne pathogen of significant concern also to the food processing industry because of its ability to grow in food under multiple stress conditions (Nair et al., 2000). A better understanding of the mechanisms of Lm for survival under stress conditions is important to control this pathogen in food. In response to changes in the natural environment, bacteria undergo a complex program of differential gene expression. A number of transcriptional regulators important for stress response gene expression have been identified in Lm (Hanawa et al., 2000; Leimeister-Wachter et al., 1990; Nair et al., 2000). ATP-binding cassette (ABC) transporters genes have been shown to be induced in Lm subjected to high pressure and under stress conditions (Liu and Ream, 2008; Liu et al., 2012b). Previous studies showed that these genes are involved in sensitivity to nisin, acids, and salt (Liu et al., 2012a). All ABC transporters are either exporters or importers. There are more than 30 copies of different ABC transporters in the genome of Lm. Some ABC transporters have been shown to be involved in biofilm formation (Zhu et al., 2008, 2011; Vanderlinde et al., 2010; Seaton et al., 2011). ABC transporters The manganese LMOf2365_1875 and LMOf2365_1877 were highly induced in milk (Liu and Ream, 2008). Manganese is involved in a number of cellular functions such as virulence and oxidative stress (Papp-Wallace and Maguire, 2006). Since the ABC the transporter operon was induced with a number of treatments such as high pressure and nisin (Liu et al., 2011), it may be supposed that it is also involved in Lm ability to form biofilm. Therefore, the in-frame deletion mutants, *\DeltaLMOf236_1875* and LMOf2365_1877 were constructed, and tested for their capacity to form biofilms in comparison with the wild type.

Materials and Methods

Lm strain *F2365* isolated from Mexican-style soft cheese that had been implicated in an outbreak of listeriosis in California in 1985 (Linnan *et al.*, 1988) was used in this study since its genome is fully sequenced and annotated (Nelson *et al.*, 2004).

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The construction of in-frame deletion mutants ΔLMOf2365_1875 and ALMOf2365 1877 in Lm F2365 was performed according Liu et al. (2012a). Glycerol stock cultures of Lm F2365 and isogenic mutants of this parent strain stored at -80°C were streaked onto Brain Heart Infusion (BHI) (Sigma-Aldrich, St. Louis, MO, USA) agar plates and grown at 37°C prior to each experiment. Five milliliters of Mueller-Hinton broth overnight cultures for Lm strains Lm 2365, Lm 1875, and Lm 1877 were initiated from plate grown cultures. The overnight cultures were incubated at 32°C with agitation (200 rpm), and the he next day, the overnight cultures were diluted 1:100 into fresh Mueller-Hinton broth. Flat bottom cell culture 96-well microtiter plates (Greiner Bio-one, Monroe, NC, USA) were washed with 100% Ethanol (ETOH) and allowed to air dry in a biological hood until all residual ETOH had evaporated. For each strain, 100 uL of the freshly diluted culture were placed in 8 different wells. Additionally, as a negative control, 100 µL of sterile Mueller-Hinton broth were also placed into 8 additional wells. The plates were incubated statically at 32°C for 48 h. The wells were then observed to see if visible biofilms were present in the *Lm* inoculated wells. The medium was then removed from the microtiter plate wells, and the individual wells were washed 5 times with 150 µL of sterile distilled water. The plates were then allowed to air dry for 45 min, and then 150 µL of a 1% crystal violet solution were added to each of the wells. After 45 min, the stain was removed and the wells were washed 5 times with 150 µL sterile distilled water. The wells were then destained of the residual biofilm bound crystal violet with 200 µL of 95% ETOH. One hundred microliters of the destain mixture were then transferred to a new microtiter plate, and the OD 590nm was measured for each well. The resulting data for

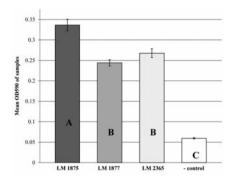


Figure 1. The mean optical density (590 nm) values produced by multiple biofilm density assays. No letters in common between columns denoted significant differences (P<0.05) in the mean values generated by the L. monocytogenes strains. Error bars represent the standard error of the means.

three separate trials was subjected to an analysis of variance. The individual trials were considered as a block when performing the mean separations using the least significant difference technique at a P<0.05 level (Miller, 1981).

Results

Resultsdemonstratedthat $\Delta LMOf2365_1875$ formed more biofilm thanthe wild type whereas biofilm formation by $\Delta LMOf2365_1877$ was similar to the wild type(Figure 1).

Discussion and Conclusions

Functional genomics research on Lm allows a better understanding of the genes related to stress responses, and this knowledge may help in the development of intervention strategies to control this food-borne pathogen. In L. monocytogenes, one ABC transporter (lm.G_1771) encoding a putative ABC transporter permease has been identified to be involved in the negative regulation of biofilm formation since deletion of this gene resulted in increased capacity in biofilm formation (Zhu et al., 2008). In our study, ALMOf2365_1875 also showed increased capacity in biofilm formation compared to the wild type, although LMOf2365_1875 showed very little homology to Lm G_1771. These results suggest that these two genes may both be involved in the capacity of Lm to form biofilms. Since ATP-Binding Cassette transporters have been shown to be involved in nisin resistance and sensitivity to acids and salt, it may be hypothesized that these genes could be used as targets for the development of new antimicrobials in food, but not to prevent the biofilms formation. EC Regulation 2073/2005 on the microbiological criteria for foodstuffs, contains provisions for Lm, and the competent authority has to verify compliance with the rules and criteria laid down in this Regulation. The results of this study, related to the molecular basis of virulence and stress responses of *Lm*, may help in the development of targeted intervention strategies and treatments for the control of the pathogen in foods.

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