



Research Article

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Effect of Prostaglandin F_{2α} on Growth of *Streptococcus uberis* associated with Bovine Mastitis



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Abstract

Certain fatty acids have been shown to inhibit the growth of mastitis pathogens. Moreover, prostaglandin in F_{2α} (PGF_{2α}) inhibits the growth of *Staphylococcus aureus* *in vitro*. The objective was to determine the antibacterial effects of PGF_{2α} (in the form of dinoprost tromethamine) on growth of *Streptococcus uberis* (*Strep. uberis*) *in vitro*. Flasks containing tryptic soy broth were inoculated with *Strep. uberis*, and subsequently treated with PGF_{2α} at concentrations of 0 (control), 0.6, 1.2, 2.4, and 4.8 mg/ml. Cultures were sampled every 4h over 24h to determine bacterial growth. The experiment was repeated 3 times in duplicate. Bacterial growth was assessed by counting colony forming units (CFU) in triplicate. Data were analyzed by repeated measures analysis of variance and reduced and full dummy variable regression models to determine the effect of PGF_{2α} concentrations on growth patterns of *Strep. uberis* over time. There was an effect of treatment and treatment by time interaction on mean log CFU for *Strep. uberis* (P < 0.05). At the time of inoculation (0h), mean log CFU values were not different among treatments. At 12 and 24h of growth, mean log CFU for all PGF_{2α} concentrations were smaller and differed (P < 0.05) from the control, in a dose dependent manner. The predicted growth curve pattern over 24h for each treatment was different (P < 0.05) when compared with the control, and the growth rate over time for treatments 2.4 and 4.8 mg/ml was slower and different from the control (P < 0.05). These results provide evidence that PGF_{2α} has inhibitory effects on growth of *Strep. uberis* *in vitro*.

Keywords: Fatty acids, mastitis, prostaglandin F_{2α}, *Streptococcus uberis*

Abbreviations: PGF_{2α}: Prostaglandin F_{2α}; *Strep. uberis*: *Streptococcus uberis*; CFU: Colony Forming Units

Introduction

The spread of mastitis pathogens causes large economic losses to the dairy industry. These losses include increased involuntary culling rate, reduced milk production, increase somatic cell count, and discarded milk [1,2]. Costs associated with mastitis infections in the U.S. dairy industry have been estimated at nearly US \$2 billion per year [3]. *Streptococcus uberis* is one of the most common gram-positive causes of clinical mastitis and contributes to a large proportion of subclinical mastitis cases [4,5]. *Streptococcus uberis* infections are often difficult to cure with traditional intra-mammary antibiotic preparations, especially in older animals [6,7]. Similar to treatment trials focused on *Staphylococcus aureus* (*S. aureus*) intra-mammary infection, increased somatic cell counts before treatment was associated with a decreased probability of cure [6]. This may explain why the response of *Strep. uberis* mastitis to treatment can be poor, even after extended therapy [7]. If cure

rates are low, it is generally not considered cost-effective to treat cows with chronic cases infections [8].

The bactericidal activities of various fatty acids as an alternative to antibiotics have been studied and reviewed [9,10]. Kelsey et al. [11] demonstrated that lauric acid, capric acid, myristic acid and linoleic acid inhibited growth of two different mastitis strains of *S. aureus*. Arachidonic acid, a fatty acid derived from linoleic acid [12] inhibited gram-positive bacteria such as *S. aureus* and *S. pyogenes* [13]. Considering that both linoleic and arachidonic acid have been shown to affect bacterial growth, we speculated that PGF_{2α}, synthesized from these fatty acids, may have similar antibacterial properties. In fact, the results from a recent study in our laboratory [14, 15]; indicate that PGF_{2α} inhibits the growth of *S. aureus* and *Mycoplasma bovis*. Given that *Strep. uberis* is one of the most common gram-positive causes of clinical mastitis, our hypothesis was that PGF_{2α} would

inhibit the growth of *Strep. uberis*. The objective of this study was to determine the effect of PGF_{2α} on *Strep. uberis* in vitro by characterizing the growth response of *Strep. uberis* to PGF_{2α} in the form of dinoprostromethamine.

Materials and Methods

Experimental design and treatment

Bacterial cultures were prepared by inoculating a single colony into 3ml of tryptic soy broth (TSB) (EMD Chemicals Inc., Darmstadt, NJ) followed by overnight incubation at 37 °C with shaking at 250rpm. In order to obtain a sufficient culture for the experiment, culture tubes containing 10ml of fresh TSB were inoculated at 1:100 with 3ml of overnight *Strep. uberis* culture, and once more incubated overnight at 37 °C with shaking at 250rpm. Prostaglandin F_{2α} in the form of dinoprostromethamine (Zoetis, Florham Park, NJ) was added to flasks for a final concentration of 0, 0.6, 1.2, 2.4 and 4.8 mg/ml (2 flasks/treatment). Flasks, which included both treatment and controls, were inoculated with the 10 ml overnight culture of *Strep. uberis* at a concentration of 1:100. Flasks were incubated at 37 °C and shaken at 250rpm for 24h; at 0h, and every 4h thereafter, 1ml samples were taken from each flask to determine bacterial growth. The entire experiment was repeated three times, in duplicate, in different days to account for variation associated with a day effect, categorized as run.

Determination of bacterial growth

To determine colony forming units (CFU), samples of 0.5ml were taken from flasks for plating. Serial dilutions were performed before samples were placed on agar plates (EMD Chemicals Inc., Darmstadt, NJ). The CFU counts were done in duplicate per sample from each flask. Plates were incubated at least 12h, or until colonies were apparent, at a constant temperature of 37 °C. The CFU counts from each of the two agar plates were averaged for each of the corresponding flasks.

Statistical analysis

The number of live cells, as measured by log CFU, was determined by averaging the number of cells for the duplicate concentrations of both plates at each 4h time point. An analysis of variance (repeated measures) was carried out using the mixed procedure of SAS (SAS Institute, Cary, NC) where the model included treatment, time (repeated factor) and their interaction. To further analyze the effect of treatments over time on the growth pattern and growth rate of *Strep. uberis*, a full model dummy variable regression procedure was also performed. The coincidence or equality of the estimated regression lines, the rate of bacterial growth over time, and the point at which the inflection of the growth curve occurred (an indication of maximum bacterial growth) were determined. The estimation of the reduced models for each treatment was carried out using PROC REG procedures of SAS, and that of the full model was carried out using PROC GLM procedures of SAS. The fitted reduced model for each treatment took the form of

$$Y = \beta_0 + \beta_1 x + \beta_2 x_2 + \varepsilon_1$$

Where Y was the logarithmic value of the number of live cells (log CFU/ml), x represented time, β_0 was the intercept (estimated log CFU/ml at time 0), β_1 was the rate of increase for bacterial growth, β_2 was the point of inflection, and ε_1 represented the random error under the classical regression assumptions. The adequacy of the fit was determined by the significance of the parameter estimates (declared at $P < 0.05$), their corresponding magnitudes and signs, and the examination of the estimated residuals.

Results

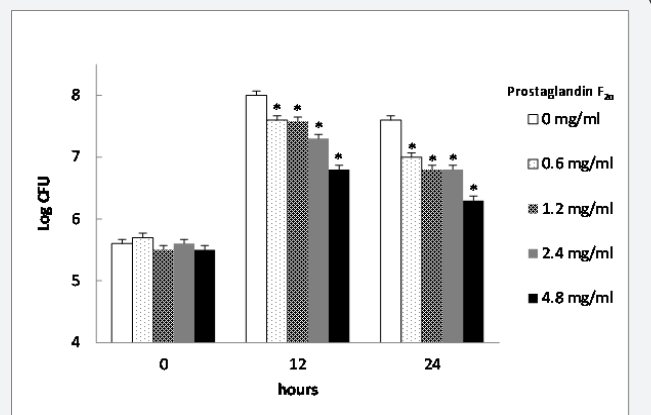


Figure 1: Growth of *Strep. uberis* treated with various concentrations of PGF_{2α} at 0, 12 and 24h of culture. *Means of each treatment at 12 and 24h differ ($P < 0.05$) from control (0 mg/ml of PGF_{2α}).

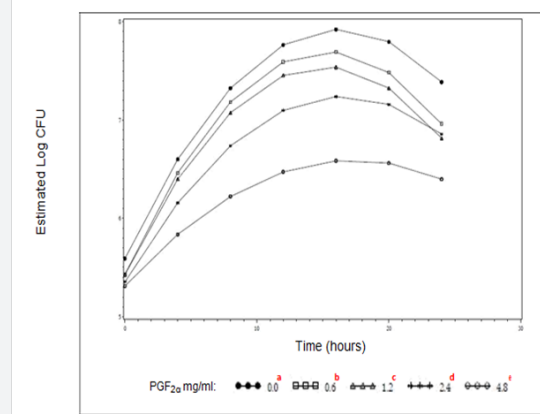


Figure 2: Full regression model ($Y = \beta_0 + \beta_1 x + \beta_2 x_2 + \varepsilon_1$) fit to growth of *Strep. uberis*, as measured by estimated log colony forming units (CFU), treated with PGF_{2α} at concentrations of 0 (black circles), 0.6 mg/ml (squares), 1.2mg/mL (triangles), 2.4mg/ml (stars) and 4.8 mg/ml (open circles) where Y was the logarithmic value of the number of live cells (log CFU/ml), x represented time, β_0 was the intercept (estimated log CFU/ml at time 0), β_1 was the rate of increase for bacterial growth, β_2 was the point of inflection, and ε_1 represented the random error under the classical regression assumptions. ^{a, b, c, d, e} Pattern of bacterial growth with different letters are different from control from each other ($P < 0.05$). The rate of bacterial growth over time (β_1) was less ($P < 0.08$) for 2.4 ($P < 0.08$) and 4.8 mg/ml ($P < 0.001$) PGF_{2α} treatment when compared with the control.

Bacterial growth curves were evaluated in growth media containing PGF_{2α} at concentrations of 0, 0.6, 1.2, 2.4mg/ml and 4.8, with 0 mg/ml referring to the control with no PGF_{2α}. Based on bacterial growth curves, PGF_{2α} in the form of dinoprosttromethamine, has inhibitory effects on growth of *Strep. uberis* (Figure 1). Overall, growth of *Strep. uberis* decreased with increasing concentrations of PGF_{2α} with 4.8 mg/ml of PGF_{2α} being the most inhibitory (Figure 2).

There was an effect of treatment and treatment by time interaction on log CFU/ml (P < 0.05), providing evidence that bacterial growth of *Strep. uberis* over time was not similar among PGF_{2α} treatments. Preplanned contrasts were conducted to compare the mean log CFU/ml values between treatments at 12 and 24h. At 0h the mean log CFU/ml values were not different among treatments and control, and averaged 5.6±0.02 log CFU/ml (Figure 1). Mean log CFU/ml values at 12 and 24h for each PGF_{2α} treatment dose, however, were different (P < 0.05) from the control (Figure 1). At 12h after incubation, the bacterial growth for each PGF_{2α} treatment reached its maximum (Figure 1). Mean log CFU/ml for 0.6 mg/ml (7.7±0.06 log CFU/ml), 1.2 mg/ml (7.6±0.06 log CFU/ml), 2.4 mg/ml (7.3±0.06) and 4.8 mg/ml (6.7±0.06) were all different (P < 0.05) from 0mg/ml (control, 8.0±0.06). At 24h, mean logs CFU/ml for 0.6 mg/ml (6.9±0.06 log CFU/ml), 1.2 mg/ml (6.8±0.06 log CFU/ml), 2.4 mg/ml (6.8±0.06) and 4.8 mg/ml (6.4±0.06) were also different (P < 0.05) when compared with 0 mg/ml (control, 7.5±0.06) (Figure 1). Interestingly, PGF_{2α} at the greatest dose (4.8 mg/ml) had the greatest effect on bacterial growth as log CFU/ml never reached above 6.4 log CFU/ml.

The reduced and full dummy variable models were carried out to evaluate the effects of different PGF_{2α} concentrations on the growth pattern of the bacteria concentration on the growth pattern of the bacteria over time. The parameter estimates of the reduced model for all treatment doses of PGF_{2α} were significant (P < 0.05), indicating that the reduced model fit the data well for each of those treatments and that all parameters are required (Table 1, Figure. 2). The preplanned contrasts carried out using the dummy variable regression model (Table 2) indicated that the overall line of growth over a 24h period was different (P<0.05) for treatments 0.6, 1.2, 2.4, and 4.8 mg/ml, when compared with the control (0 mg/ml), implying that the bacterial growth pattern for these treatments were different from control and lending support to results found through the repeated measure analysis. The rate of bacterial growth over time (β₁) was less (P < 0.08) for 2.4 (P < 0.08) and 4.8 mg/ml (P < 0.001) PGF_{2α} treatment when compared with the control, providing evidence that the rate of bacterial growth over time was different between those treatments in a dose dependent manner (Table 2). In addition, the rate of bacterial growth was slower in 4.8 than with 1.2, 2.4mg/ml PGF_{2α} (data not shown). Each growth curve had an estimated point of inflection, where the estimated maximum log CFU/ml was reached at a specific time. The estimate parameter point of inflection (β₂, the estimate

of maximum bacterial growth) for 4.8 mg/ml PGF_{2α} treatment was different from the control (P < 0.05, Table 2) and all other PGF_{2α} treatment groups (P < 0.05, data no shown).

Table 1: Parameter estimates, standard errors, and the associated P-values for the reduced regression model of *Strep. uberis* treated with various doses of PGF_{2α}.

| PGF _{2α} Dose (mg/ml) | Parameter | Parameter Estimate | Standard | P valuea |
|--------------------------------|----------------|--------------------|----------|----------|
| 0 | β ₀ | 4.3 | 0.21 | <0.001 |
| 0 | β ₁ | 1.43 | 0.12 | <0.001 |
| 0 | β ₂ | -0.14 | 0.01 | <0.001 |
| 0.6 | β ₀ | 4.41 | 0.21 | <0.001 |
| 0.6 | β ₁ | 1.35 | 0.12 | <0.001 |
| 0.6 | β ₂ | -0.14 | 0.01 | <0.001 |
| 1.2 | β ₀ | 4.16 | 0.21 | <0.001 |
| 1.2 | β ₁ | 1.41 | 0.12 | <0.001 |
| 1.2 | β ₂ | -0.15 | 0.01 | <0.001 |
| 2.4 | β ₀ | 4.33 | 0.21 | <0.001 |
| 2.4 | β ₁ | 1.13 | 0.12 | <0.001 |
| 2.4 | β ₂ | -0.11 | 0.01 | <0.001 |
| 4.8 | β ₀ | 4.65 | 0.21 | <0.001 |
| 4.8 | β ₁ | 0.73 | 0.12 | <0.001 |
| 4.8 | β ₂ | -0.06 | 0.01 | <0.001 |

Significance of parameter estimates represents appropriate fit of model to data; β₀ represents y-intercept at time of inoculation; β₁ represents the rate at which bacterial growth increases; β₂ represents point of inflection where growth reached its maximum at a specific time.

Table 2: Degrees of freedom (df), and associated P-values for the pre-planned contrasts of estimated parameters for growth of *Strep. uberis* reacted with various doses of PGF_{2α}.

| Parametersa | Pre-planned contrasts | df | P value |
|-------------------|-----------------------|----|-----------------|
| ^a Line | 0 vs. 0.6 mg/ml | 3 | 0.0203 |
| | 0 vs. 1.2 mg/ml | 3 | 0.001 |
| | 0 vs. 2.4 mg/ml | 3 | <0.001 |
| | 0.6 vs. 4.8 mg/ml | 3 | <0.001 |
| β ₁ | 0 vs. 0.6 mg/ml | 1 | NS ^b |
| | 0 vs. 1.2 mg/ml | 1 | NS |
| | 0 vs. 2.4 mg/ml | 1 | 0.0804 |
| | 0 vs. 4.8 mg/ml | 1 | 0.001 |
| β ₂ | 0 vs. 0.6 mg/ml | 1 | NS |
| | 0 vs. 1.2 mg/ml | 1 | NS |
| | 0 vs. 2.4 mg/ml | 1 | 0.1401 |
| | 0 vs. 4.8 mg/ml | 1 | 0.001 |

^aLine = is the estimated line of growth over time, β₁ is the rate at which bacterial growth increases, and β₂ = represents the point of inflection where the estimated growth reached its maximum at a specific time. ^bNS = not significant.

Discussion

This research addresses the question whether the fatty acid PGF_{2α} is inhibitory to growth of *Strep. uberis*. The results support

the hypothesis that PGF_{2α'} in the form of dinoprosttromethamine has inhibitory effects on growth of *Strep. uberis*. These findings support the results from our previous research in which PGF_{2α'} inhibited growth of gram positive bacteria, *S. aureus* [14] as well as *Mycoplasma bovis* [15]. The antimicrobial properties of fatty acids on bacteria have been studied for years. The effectiveness of fatty acids in inhibiting growth of several gram-positive bacteria have also been demonstrated and reviewed [11,13,16].

Arachidonic acid, a fatty acid originally derived from linoleic acid, has been shown to inhibit gram-positive bacteria such as *Streptococcus faecalis* and *Staphylococcus epidermidis*, and *S. aureus* [13]. The authors hypothesized those bactericidal effects on *S. aureus* mediated by peroxidation of arachidonic acid. Because both linoleic acid and arachidonic acid are precursors to PGF_{2α'}, it is plausible that PGF_{2α'} synthesized from these fatty acids, has similar antibacterial properties. The results supported our hypothesis that commercially available PGF_{2α'} (dinoprosttromethamine) inhibited the growth of *Strep. uberis* in vitro in a dose dependent manner (Figure 3), resembling the actions of linoleic acid on growth of *S. aureus* Novel as previously described [11].

The mechanism by which PGF_{2α'} affected *Strep. uberis* cannot be determined from the current study. The inhibitory properties of fatty acids were more noticeable with increased chain length and degree of un-saturation [10,13,17]. Zheng et al [13] found differences in antibacterial activity between unsaturated fatty acids and saturated fatty acids in that saturated fatty acids had less or no antibacterial activity against *S. aureus* and *S. pyogenes*. Dinoprosttromethamine contains two double bonds and consists of 24 carbons. These features may be important factors in its antibacterial properties.

One potential mechanism of action centers on the ability of fatty acids to penetrate and disrupt the phospholipid bi-layer of the plasma membrane of bacteria and ultimately increases the negative charge on the bacterial membrane surface [10]. Zheng et al. [13] proposed that antibacterial action of unsaturated fatty acids is mediated by inhibition of bacterial enoyl-acyl carrier protein reductase which is an essential component of bacterial fatty acid biosynthesis. Another proposed mechanism involves the hindering of bacterial growth via an interaction with lipid bi-layer of the cell membrane, resulting in a change in membrane permeability, or the interference with transduction cascades leading to cell lysis [10, 18]. In summary, the current in vitro results provide evidence that the fatty acid PGF_{2α'} in the form of dinoprosttromethamine, has inhibitory effects on the growth of *Strep. uberis* in a dose dependent manner. The potential use of PGF_{2α'} as an anti-bacterial fatty acid, for treatment of mastitis requires more research.

Acknowledgments

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