Efficacy of virginiamycin for the control of periodontal disease in calves

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Periodontal diseases are multifactorial infectious processes caused by complexes of microorganisms, with damage to health, production, and animal welfare. The aim of the present study was to evaluate the efficacy of virginiamycin in the prevention and control of two early forms of periodontal disease: gingivitis and necrotizing gingivitis. Ten weaned calves, aged four to six months, were permanently kept in a single lot under the same rotational grazing regime in a newly reformed area of Panicum maximum. Five of the calves were orally administered 340mg of virginiamycin (Virginiamycin Group) daily for a period of 18 weeks, while the remaining five calves (Control Group) remained under the same food management but did not receive virginiamycin. During this period, animals underwent 18 weekly evaluations regarding periodontal health, with monitoring and recording of clinical parameters of the eight deciduous incisor teeth on the labial and lingual faces. At approximately two-week intervals, nine collections of subgingival sulcus material from five sites of the four right incisor teeth of each animal were performed and subjected to microbiological evaluation using polymerase chain reaction with primers of 25 microorganisms considered potentially pathogenic. After 1440 periodontal clinical evaluations of incisor teeth of the 10 calves, a total of 395 episodes of gingivitis were recorded, of which 267 occurred in the Control Group and 128 in the Virginiamycin Group. Similarly, 89 episodes of necrotizing gingivitis were recorded; 58 in the Control Group and 31 in the Virginiamycin Group. Comparison of between-group means found significant differences for teeth with gingivitis and necrotizing gingivitis (t test; p<0.05). The total number of teeth with gingivitis (p<0.01) and necrotizing gingivitis (p<0.01) in Control Group was significantly higher than that of gingivitis (p<0.01) and necrotizing gingivitis (p<0.05) in the Virginiamycin Group. There was a positive correlation between total occurrence of gingivitis and necrotizing gingivitis in the Virginiamycin Group by Pearson's test. Virginiamycin had a protective effect on treated animals compared with the Control Group (OR = 0.36: CI (95%) = 0.27-0.43). In the Control Group, Actinomyces israelli (4.74%), domain Archaea (1.58%), Eikenella corrodens (1.05%), Fusobacterium nucleatum (27.37%), Porphyromonas endodontalis (5.26%) and Porphyromonas gulae (0.53%).
Prevotella buccae (6.32%), Prevotella loescheii (3.68%), Prevotella nigrescens (8.42%), Prevotella oralis (1.58%), Tannerella forsythia (0.53%), and Treponema denticola (4.21%) were detected at healthy sites, and gingivitis or necrotizing gingivitis samples. In the Virginiamycin Group, A. israelii (3.41%), domain *Archaea* (0.98%), *F. nucleatum* (9.27%), class *Mollicutes* (4.39%), *P. endodontalis* (4.39%), *P. gulae* (0.49%), *P. buccae* (8.29%), *P. loescheii* (6.83%), *P. nigrescens* (15.61%), *P. oralis* (1.46%), *Selenomonas sputigena* (0.49%), *T. forsythia* (0.49%), and *T. denticola* (2.44%) were detected. In conclusion, virginiamycin administered at a dosage of 340mg/animal/day significantly reduced the occurrence of gingivitis and necrotizing gingivitis in cattle maintained on reformed pastures, and was revealed to have action against periodontal bacterial microbiota considered to be potentially pathogenic.

INDEX TERMS: Virginiamycin, control, periodontal disease, gingivitis, necrotizing gingivitis, ruminants, cattle, microbiology, clinics.

INTRODUCTION

Periodontal diseases are a group of diseases that affect the tissues associated with protection and support of teeth. Among the reversible forms of periodontal disease are gingivitis and necrotizing gingivitis, which are caused predominantly by aggression of gingival biofilm (Konradsson et al. 2007, Kistler et al. 2013).

Untreated gingivitis can progress to periodontitis, with consequent compromise of periodontal ligaments and alveolar bone, culminating in tooth loss (Page 1986, Lyon 2005, Kinane & Bartold 2007, Herrera et al. 2014). This way, the health of the periodontium depends on the balance between bacterial composition of dental biofilm and its interaction with the host immune system (Hajishengallis 2015).

In cattle, gingivitis has been reported in calves aged 5 to 60 days old, characterized as a physiological manifestation resulting from eruption of teeth (Döbereiner et al. 1974). Additionally, necrotizing gingivitis has been described in cattle with leukocyte adhesion deficiency (Nagahata et al. 1993). However, in this animal species, little is known about gum diseases, most likely due to the difficulties in evaluating the oral cavity of the animals and by the fact that these diseases do not present evident clinical changes as seen in periodontitis. However, in humans and adolescents, these diseases have been well described (Kirian et al. 2011, Marshall et al. 2014).

In cases of gingivitis, the gingival border is noted to be or with probing, may be observed and in more severe
cases, the gingiva may have ulcerations (Diehl & Rosychuk 1993, Lyon 2005, Riggio et al. 2011, Newman et al. 2012, Antiabong et al. 2013, Kutasi et al. 2016). In cases of necrotizing gingivitis, spontaneous bleeding or bleeding after probing as well as the presence of a layer of yellowish-white or grayish-white fibrin on the necrotic gingival border may be observed (Klokkevold 2012, Rodriguez-Pulido et al. 2016).

Periodontal diseases occur by the modification of the microbiota and constituents of the oral cavity; in humans, several modifying factors have been associated with the occurrence of gingivitis and necrotizing gingivitis, including hormonal changes and immunodeficiencies (Stamm 1986, Rowland 1999, Dufty et al. 2016). However, in cattle, modifying factors associated with the occurrence of periodontal diseases are unknown and the suspicions are related to soil management and diet (Dutra et al. 1993, Döbereiner et al. 2000).

In a recent study, oral microbiomes of healthy cattle and those with periodontitis revealed 72.1% dissimilarity; however, the diversity of bacteria found in healthy and diseased sites was similar, with a predominance of the genera Prevotella, Fusobacterium, and Porphyromonas in periodontal lesions (Borsanelli et al. 2018). Thereby, it can be evidenced that bovine oral microbiota is rich and diversified, composed of 395 genera or higher taxa and that microorganisms considered as potential periodontal pathogens are associated with the occurrence of periodontitis in these animals. These include several species of Porphyromonas, Prevotella, and Treponema, as well as Fusobacterium nucleatum, Fusobacterium necrophorum, and Actinomyces naeslundii (Dutra et al. 2000, Borsanelli et al. 2015a, 2015b, Borsanelli 2017).

In previous studies, virginiamycin has been shown to be efficient for recovery of calves with periodontitis (Tims et al. 1992), as well as periodontitis prevention (Dutra et al. 1993). In this context, the present study aimed to evaluate the efficacy of virginiamycin for the control of gingivitis and necrotizing gingivitis in cattle, in view of the need to develop strategies for the control of periodontal diseases in production animals and characteristics of the antibiotic.

**MATERIALS AND METHODS**

**Animals.** Ten male Jersey or Jersey cross calves, weaned and healthy, aged four to six months, were used. They were weighed and randomized into two groups of five animals each. The animals remained under rotational grazing and even single-plot zootechnical management in 24 paddocks (approximately 3 hectares) that had previously been reformed in order to simulate the potential representative situation for the occurrence of periodontal disease in cattle. Mixed pastures of Massai grasses (Panicum maximum cv. Massai) and Mombasa (Panicum maximum cv. Mombasa) were reformed following conventional farming practices such as soil analysis, liming, and fertilization. Water and mineral salt were supplied ad libitum over the total period of approximately 24 weeks of experimentation, including adaptation of the animals to the diet. Fresh animals (Virginiamycin Group) received daily, oral (pour dressing) administration of virginiamycin (340mg/day/animal) for 18 consecutive weeks while the other five (Control Group) did not receive the antibiotic.

**General and periodontal clinical evaluation.** At the beginning of the experiment, the animals had good body condition scores for their age and were apparently healthy. In the initial oral cavity clinical examination, they had deciduous dentition, with dental units (tooth and periodontium) normal. Weekly periodontal clinical evaluation of

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**Extraction of microbial DNA.** Each sample for bacterial DNA detection in sterile ultrapure water was extracted by boiling. First, the sample was homogenized for 20 seconds; 500μl aliquots were then removed and stored in Eppendorf tubes. The samples were then boiled for 15 minutes, centrifuged at 10.36×g for 5 minutes and aliquots of 400μl. The samples were then stored at -80°C until DNA extraction.


**Amplifications of target DNA by PCR were performed in 25μl volumes containing 1X PCR / Mg + 2 buffer (Boehringer Mannheim, Indianapolis, IN, USA), 0.2μl of each dNTP (Pharmacia Biotech, Piscataway, NJ, USA), 0.5U Taq DNA polymerase (Invitrogen do Brasil, São Paulo/SP, Brazil), 0.4μl of each primer pair (Invitrogen), and 10ng template. The amplifications were performed in a thermocycler (Perkin Elmer, GeneAmp PCR System 9700, Norwalk/CT, USA) programmed at 94°C (5 minutes), 30 to 40 cycles at 94°C (30 to 60 seconds,
according to the primers used), specific annealing temperature for each primer pair; 72°C (30 to 60 seconds), followed by a 5 minute period at 72°C for the final extension of the amplified DNA strands. The PCR amplification products PCR were subjected to agarose electrophoresis in 1% TBE buffer (1M Tris, 0.9M boric acid, 0.01M EDTA, pH 8.4), stained with ethidium bromide (0.5mg/ml) and photographed in ultraviolet light transilluminator (UV Light Transilluminator; Eastman Kodak Co., NY, USA).

DNA samples of reference strains were used for control of detection procedures, as well as clinical samples positive for the target microorganisms (Gaetti-Jardim Junior et al. 2012). Ultrapure water was used as a negative control.

Statistical analysis. The means of the cases with gingivitis and necrotizing gingivitis were compared between the Virginiamycin Group and the Control Group using Student’s t test and Pearson’s test, with a significance level of p <0.05. For the statistics regarding the detection of microorganisms, a comparison was made between the dichotomous variables, in which the presence of microorganisms between the Virginiamycin and Control Groups was compared; the correlation between the microorganisms, and their relation to sites with periodontal or healthy lesions, was evaluated using the chi-Square test of maximum-likelihood, with a significance level of p<0.05. Odds ratios (OR) and confidence intervals (CI) for the OR were calculated to verify whether the use of virginiamycin would be a protective factor against gingivitis and necrotizing gingivitis.

Research ethics commission. The experiment was approved by the Ethics Committee on Animal Use (CEUA) of the Faculty of Agrarian and Veterinary Sciences - Unesp, Campus Jaboticabal/SP (Process FCAV/Unesp No. 15.207/16).

RESULTS

At the end of 18 weeks, the animals that did not receive virginiamycin had unsatisfactory body condition scores. Furthermore, at different moments of the study, the control animals presented symptoms of diarrhea and nasal secretion, and showed an apparent increase in susceptibility to intercurrent diseases such as endoparastisos. In contrast, among the five animals in the Virginiamycin Group, only one had episodes of diarrhea. The mean weight of calves receiving virginiamycin (188.2kg) was significantly higher (p=0.02) than that observed in the Control Group (123.5kg).

Periodontal evaluation of incisor teeth of the calves in the Control and Virginiamycin Groups kept in a single lot, under rotational grazing in a newly reformed area and even feeding management, revealed episodes of gingivitis and same necrotizing gingivitis, initial forms for progression of periodontitis in the animals of the two experimental groups.

During the study period, according to the weekly clinical examination of the eight deciduous teeth (labial and lingual faces), 395 episodes of teeth (dental unit) with gingivitis were recorded in the 1440 evaluations performed. Of these episodes, 267 occurred in Control Group calves and 128 in the Virginiamycin Group (Fig.1). In the same evaluation, of a total of 89 records of teeth with necrotizing gingivitis, 58 occurred in the Control Group and 31 in the Virginiamycin Group (Fig.2).

The clinical manifestations of gingivitis included changes in the color of the gingival mucosa, ranging from moderate to severe red, edema, and bleeding (spontaneous or with probing). In necrotizing gingivitis, the marginal gingival epithelium presented necrotic ulcerations or epithelial areas covered with a white-gray/yellowish-white membranous layer, and bleeding (spontaneous or with probing) (Fig.3 and Fig.4).

In the comparison of the means of the experimental groups, the differences found for teeth (dental unit) with gingivitis and necrotizing gingivitis were significant according to the t test (p<0.05). Thus, the total number of teeth with gingivitis (p<0.01) and necrotizing gingivitis (p<0.01) in the Control Group was significantly higher than the total number of gingivitis (p<0.01) and necrotizing gingivitis (p<0.05) in the Virginiamycin Group. There was a positive correlation between the total occurrence of gingivitis and necrotizing gingivitis in the Virginiamycin Group according to Pearson’s test. Virginiamycin was considered a protective factor against the development of gingivitis and necrotizing gingivitis (OR = 0.36; CI (95%) = 0.27-0.43).
In the microbiological evaluation, 395 samples were evaluated by PCR (Table 1) to detect the presence of 25 microorganisms considered indicators and with pathogenic potential. The sites with the lowest number of microorganisms detected were those associated with necrotizing gingivitis (Table 2 and 3). The amount of microorganism samples at the different sites collected differed among the animals and between the groups (Fig.5). In addition, there was a difference in the types of microorganisms detected between the sample collections.

*Fusobacterium nucleatum*, class *Mollicutes*, *Porphyromonas endodontalis*, *Prevotella loescheii*, *Prevotella nigrescens*, *Prevotella oralis*, and *Treponema denticola* were detected in five or more collections, whereas *Actinomyces israelii*, *Archaea domain*, *Eikenella corrodens*, *Porphyromonas gulae*, *Tannerella forsythia*, and *Selenomonas sputigena* were rarer. *Prevotella buccae* was detected in all collections. *Actinomyces naeslundii*, *Campylobacter spp.*, *Fusobacterium necrophorum*, *Parvimonas micra*, *Porphyromonas assacharolytica*, *P. gingivalis*, *Prevotella intermedia*, *P. melaninogenica*, *Treponema amylovorum*, *T. maltophilum*, and *T. pectinovorum* were not identified in either of the two groups. *S. sputigena* was negative in the Control Group animals and *E. corrodens* was negative in the Virginiamycin Group.

In the analysis of the variables using the chi-square test of maximum-likelihood, there was a significant difference in the frequency of detected microorganisms (*p*<0.05). The presence of *Fusobacterium nucleatum* (*p*<0.01) was more significant in Control Group, while *Prevotella nigrescens* (*p*<0.04) and other bacteria of the genus *Prevotella* (*p*<0.02) was more frequent in the Virginiamycin Group. *Prevotella loescheii* showed association with *Prevotella buccae* (*p*<0.01) and *P. nigrescens* (*p*<0.01); *P. buccae* was associated with *P. nigrescens* (*p*<0.01) and
Fig. 5. Number of microorganisms detected in the 395 samples collected from four deciduous teeth of each animal from the Virginiamycin Group (n=5) and Control Group (n=5). * P < 0.01, ** P < 0.05.

### Table 1. Microorganisms detected by polymerase chain reaction (PCR) in the samples of 395 subgingival sulcus sites of 10 calves of groups Control (n=190) and Virginiamycin (n=205)

<table>
<thead>
<tr>
<th>Microorganism</th>
<th>Control group</th>
<th>Virginiamycin group</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>Actinomyces israeli</td>
<td>9 (4.7)</td>
<td>7 (3.4)</td>
<td></td>
</tr>
<tr>
<td>Archaea domain</td>
<td>3 (1.6)</td>
<td>2 (1.0)</td>
<td></td>
</tr>
<tr>
<td>Eikenella corrodens</td>
<td>2 (1.1)</td>
<td>0 (0.0)</td>
<td></td>
</tr>
<tr>
<td>Fusobacterium nucleatum</td>
<td>52 (27.4)</td>
<td>19 (9.3)</td>
<td>&lt;0.01*</td>
</tr>
<tr>
<td>Porphyromonas endodontalis</td>
<td>10 (5.3)</td>
<td>9 (4.4)</td>
<td></td>
</tr>
<tr>
<td>P. gulae</td>
<td>1 (0.5)</td>
<td>1 (0.5)</td>
<td></td>
</tr>
<tr>
<td>Prevotella buccae</td>
<td>12 (6.3)</td>
<td>17 (8.3)</td>
<td></td>
</tr>
<tr>
<td>P. loescheii</td>
<td>7 (3.7)</td>
<td>14 (6.8)</td>
<td></td>
</tr>
<tr>
<td>P. nigrescens</td>
<td>16 (8.4)</td>
<td>32 (15.6)</td>
<td>&lt;0.05*</td>
</tr>
<tr>
<td>P. oralis</td>
<td>3 (1.6)</td>
<td>3 (1.5)</td>
<td></td>
</tr>
<tr>
<td>Selenomonas Sputigena</td>
<td>0 (0.0)</td>
<td>1 (0.5)</td>
<td></td>
</tr>
<tr>
<td>Tannerella forsythia</td>
<td>1 (0.5)</td>
<td>1 (0.5)</td>
<td></td>
</tr>
<tr>
<td>Treponema denticola</td>
<td>8 (4.2)</td>
<td>5 (2.4)</td>
<td></td>
</tr>
</tbody>
</table>

*Significant values of P by the Chi-square test of M-L.

### Table 2. Microorganisms identified by PCR in four healthy incisor teeth (n=119) with gingivitis (n=63) or necrotizing gingivitis (n=8) of five calves in the Control Group

<table>
<thead>
<tr>
<th>Microorganism</th>
<th>Healthy</th>
<th>Gingivitis</th>
<th>Necrotizing gingivitis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Actinomyces israeli</td>
<td>6 (5.0)</td>
<td>3 (4.8)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Archaea domain</td>
<td>3 (2.5)</td>
<td>0 (0.0)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Eikenella corrodens</td>
<td>1 (1.0)</td>
<td>1 (1.6)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Fusobacterium nucleatum</td>
<td>31 (26.1)</td>
<td>19 (30.2)</td>
<td>2 (25.0)</td>
</tr>
<tr>
<td>Mollicutes class</td>
<td>6 (5.0)</td>
<td>4 (6.4)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Porphyromonas endodontalis</td>
<td>9 (7.6)</td>
<td>1 (1.6)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>P. gulae</td>
<td>1 (1.0)</td>
<td>0 (0.0)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Prevotella buccae</td>
<td>8 (6.7)</td>
<td>3 (4.8)</td>
<td>1 (12.5)</td>
</tr>
<tr>
<td>P. loescheii</td>
<td>4 (3.4)</td>
<td>1 (1.6)</td>
<td>2 (25.0)</td>
</tr>
<tr>
<td>P. nigrescens</td>
<td>13 (11.0)</td>
<td>1 (1.6)</td>
<td>2 (25.0)</td>
</tr>
<tr>
<td>P. oralis</td>
<td>1 (1.0)</td>
<td>2 (3.2)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Tannerella forsythia</td>
<td>0 (0.0)</td>
<td>1 (1.6)</td>
<td>0 (0.0)</td>
</tr>
<tr>
<td>Treponema denticola</td>
<td>4 (3.4)</td>
<td>4 (6.4)</td>
<td>0 (0.0)</td>
</tr>
</tbody>
</table>

Archaean domain (p<0.04). Likewise, there was an association between the frequency of *P. nigrescens* and *Actinomyces israeli* (p<0.04), *E. corrodens* (p<0.01), and *Mollicutes* class (p<0.03); while *A. israeli* also showed association with *Mollicutes* class (p<0.04). Concerning the association of microorganisms with clinical signs, only *Prevotella oralis* had a level of significance with gingivitis (p<0.02).

### DISCUSSION

The occurrence of gingivitis and necrotizing gingivitis in cattle kept on reformed pastures is an original description in the literature about this important complex of diseases of ruminant’s oral cavity. In fact, although gingivitis and necrotizing gingivitis are precursors of necrotizing periodontitis and periodontitis, their description in farm animals is rare. In general, existing studies deal with the final event (periodontitis) that is possible to be evaluated by extensive periodontal lesions such as gingival recession, clinical loss of insertion level, and damage to animal health and welfare (Döbereiner et al. 2000, 2004, Borsanelli 2017). In this context, the present longitudinal study and clinical monitoring of deciduous teeth of calves has enabled an unprecedented documentation of a natural occurrence, in successive episodes followed by remission, of two precursor forms of periodontitis. No less important, results of virginiamycin use in periodontal disease control in a longitudinal study with clinical monitoring of the gingival condition of the incisor teeth for four consecutive months and the presence of microorganisms indicative of potentially pathogenic microbiota are also unprecedented.

Progression of gingivitis in periodontitis is associated with individual, environmental, and etiological factors (Lang et al. 2009). It is known that pasture reform, through liming and fertilization, can favor the occurrence of bovine periodontitis in previously endemic areas (Dutra et al. 2000). However, it is unknown whether this frequent association is valid for all periodontal diseases; it should be added that...
It is worth noting that the succession of episodes of these two forms of periodontal disease precursors in cattle is paralleled throughout the literature with respect to this group of diseases in humans and other animals (Kinane & Bartold 2007, Marshall et al. 2014). In an objective and conclusive manner, it is possible to attribute a lower frequency of episodes of these two periodontal diseases in calves treated with virginiamycin. The difference between the two experimental groups was significant, with significant benefit to the group that ingested 340mg of virginiamycin daily for the study period. Tims et al. (1992) previously reported the effect of the antibiotic on the recovery of animals with the aggressive form of periodontitis even when they continued under the same epidemiological condition that triggered the outbreak of the disease.

As multifactorial infectious diseases, periodontal diseases are non-linear processes, external to the organism but associated with the bacterial biofilm adhered to the tooth and planktonic microbiota (Socransky & Haffajee 2005, Borsanelli et al. 2018). In this approach, the action of virginiamycin probably has been in the promotion of bacterial homeostasis favorable to the maintenance of periodontal health or in the prevention of dysbiosis, which causes the onset of periodontal diseases.

In periodontal diseases, gingival biofilm plays an essential role in its etiology (Löe 1994). In the calves included in the present study, there was practically no biofilm accumulation or dental calculus visible in the incisors.

A widely evidenced concept in the literature is that each tooth can have its own microbial complex, which, when analyzed, show differences in their constitution among teeth, such as supragingival and in periodontal pockets in cattle (Borsanelli 2017). It is interesting to note that the role of bacteria in periodontal infections must meet Sokransky’s postulate, since they are not conventional exogenous infectious diseases, but diseases of an endogenous nature, associated with the microbial biofilm, so that the interaction between the different taxa is important for the development of these diseases (Socransky & Haffajee 2010). In this sense, a statistical correlation of co-occurrence was observed between the different members of the genus Prevotella, which are part of the so-called “orange complex”, associated with the first manifestations of inflammatory
periodontal diseases, in both gingivitis and necrotizing gingivitis (Socransky et al. 1998, Larsen & Fiehn 2017). *Porphyromonas* spp., *Tannerella* spp., *Campylobacter* spp., *Eikenella* spp., *Parvimonas* spp., *Treponema* spp., and *Selenomonas* spp. were detected in monkeys with gingivitis, but in greater quantity in monkeys with periodontitis and periodontal abscesses (Gaetti-Jardim Junior et al. 2012). Thus, the use of amplification of the target DNA by conventional PCR in the present study made it possible to detect several microorganisms considered potentially pathogenic. However, it is a qualitative test, which requires caution in interpreting the results, especially in the face of endogenous disease or dysbiosis. On the other hand, Gram-positive bacteria, and some Gram-negative bacteria such as *F. nucleatum* and spirochetes, detected in sites with gingivitis and necrotizing gingivitis, can be associated as agents involved in the progression of these diseases in cattle. This is similar to observations in humans with these diseases (Harvey 2017).

In the present study, there was a trend in the frequency of occurrence of microorganisms detected in necrotizing gingivitis. In the Control Group, *P. nucleatum*, *P. buccae*, *P. loescheii*, and *P. nigrescens* were detected, while in the Virginiamycin Group, the *Archaea* domain and *Mollicutes* classes were observed. In the samples of animals with necrotizing gingivitis, no spirochetes of the genus *Treponema* were detected, which diverges from studies in humans, in which this genus has a recognized role (Herrera et al. 2014). *T. denticola* was identified in samples from healthy or gingivitis sites, similar to that observed in the oral microbiota of healthy adult bovines (Borsanelli 2017).

In ruminants and other animals it is known that *Porphyromonas* spp. is present in the periodontal pockets of animals with periodontitis (Senhorinho et al. 2011, Borsanelli et al. 2015b, Agostinho 2017, Campello 2017). Based on the results of the present study, it can be stated that besides being related to periodontitis, *P. endodontalis* and *P. gulae* are present in samples from healthy calves and calves with gingivitis. *Mollicutes* class and *Archaea* domain, in humans and animals, are present in healthy patients with gingivitis, necrotizing gingivitis, or periodontitis (Yamabe et al. 2008, Faveri et al. 2011, Griffen et al. 2012, Chen et al. 2015, Harris et al. 2015). In cattle with periodontitis, *Mollicutes* class is more often associated with the disease (Borsanelli et al. 2018). In the calves in the present study, class *Mollicutes* was more often detected in sites with gingivitis and necrotizing gingivitis, and it can be noted that this microorganism was present in the different phases of periodontal diseases. *Archaea* domain to date has not been reported in this species; this is the first study to detect *Archaea* domain in healthy sites of both groups and in calves with necrotizing gingivitis that ingested virginiamycin. This difference indicates that the promoter leads to dysbiosis, favoring the growth of these microorganisms.

*Prevotella nigrescens* is associated with the development of necrotizing gingivitis and periodontitis in humans (Loesche et al. 1985, Stingu et al. 2013) and was also identified in healthy sites and periodontal pockets of bovines (Borsanelli et al. 2015a). In the present study, the frequency of *P. nigrescens* was significant, being detected in greater number in treated animals. This genus has evidence of resistance to this class of streptogramin, which would explain the higher number of positive samples in the group that took the antibiotic (Chung et al. 2002). Another factor that makes the detection of the genus *Prevotella* important in this work is due to the greater number of samples detected at the same time as the increase of cases of necrotizing gingivitis in the Virginiamycin Group, suggesting that in the treated animals this bacterium possibly has some function in evolution of necrotizing gingivitis.

Among the microorganisms associated with gingivitis, *F. nucleatum* presents great importance, since it is one of the organisms responsible for the inflammatory process, the beginning of the development of periodontitis, and the recruitment of other periodontopathogens. Although *F. nucleatum* is commonly found in microbiological studies of periodontal diseases, it is also detected in healthy periodontal sites, as well as in sites with gingivitis and periodontitis in humans and animals (Kolenbrander 2000, Kolenbrander et al. 2002, Senhorinho et al. 2011, Signat et al. 2011, Harris et al. 2015). Species of *Fusobacterium* genus are susceptible to the action of virginiamycin (Araújo et al. 2016), which is related to the results of this present study, in which the highest number of positive samples were in the Control Group (fig.3 and fig.4). Consistent with these events, more episodes of gingivitis and necrotizing gingivitis were observed in the Control Group animals. Antibiotic treatment causes changes in the structure of the microbial community (Antiabong et al. 2013), which reflects the condition between health and disease. In the calves, it was possible to visualize the difference caused by the treatment between the groups and the presence of certain microorganisms in the analyzed sites. This is consistent with observations of the bacterial composition of humans with periodontitis treated with amoxicillin and metronidazole, in which individuals with a good response to treatment had reduced periodontopathogens such as *P. endodontalis*, *Prevotella* spp., and *Fusobacterium* spp. (Colombo et al. 2012).

In a previous study, Tims et al. (1992) reported the benefit of virginiamycin in the recovery of cattle kept in an endemic area. In the present study, the results show the benefit of the use of the antibiotic in the control of periodontal diseases, which are considered complex, polymicrobial, and dependent on relationships among the bacterial, host, and environment communities. Employed in growth promoter dose in this study, virginiamycin was effective in controlling two precursor forms of periodontitis, gingivitis and necrotizing gingivitis, and it can be considered a form of protection against the development of periodontal disease. In this context, it represents an excellent alternative to the control of periodontal disease in bovines, especially as it does not leave residues in the meat and does not pose any risks to public health (Menzies-Gow & Young 2011, Bessegatto et al. 2017).

**CONCLUSION**

Virginiamycin administered at a dosage of 340mg/animal/day reduced significantly the occurrence and was effective in control and prevention of periodontal diseases (gingivitis and necrotizing gingivitis) in calves kept in reformed pastures and with a potentially periodontal pathogenic bacterial microbiota.

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