Recovery of *Mollicutes* from the reproductive tract of dairy cattle in the state of Pernambuco, Brazil

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The aim of the present study was to report the occurrence of members of the *Mollicutes* class in the reproductive system of dairy cattle in Brazil. Five farms containing dairy cattle were visited in January of 2012. In total, 100 cows of different ages, breeds and stages of lactation were examined in the present study. The cows were part of intensive or semi-intensive management systems and were submitted to mechanical milking or hand milking. The samples were collected after washing the vulvar region with water and soap, and then drying it with paper towels and disinfecting the area with alcohol (70°GL). Vaginal mucous was collected using a sterile alginate cotton swab, which was rubbed on the vagina, as well as the lateral and internal walls. Vulvovaginal mucous samples were cultured in both liquid and solid modified Hayflick’s medium, for mycoplasmas, and UB medium, for ureaplasmas. The PCR assays for *Mollicutes* and *Ureaplasma* spp. were performed according to the standard protocols described in the current literature. During isolation, the frequency of *Mycoplasma* spp. was of 13.0% (13/100) and for *Ureaplasma* spp. was of 6.0% (6/100). In the PCR assays the frequency of *Mollicutes* was of 26.0% (26/100) and for *Ureaplasma* spp. was of 13.0% (13/100) in the dairy cattle studied. This is the first report of these agents in reproductive system of bovine of the Pernambuco state. Further studies are necessary to determine the pathogenic potential and species of these field isolates.

**INDEX TERMS:** Reproduction diseases, mycoplasmosis, dairy cows, vaginal mucous, *Ureaplasma* spp.
reprodutivo de bovinos no Estado de Pernambuco. Estudos adicionais são necessários para determinar as espécies e o potencial patogênico destes isolados de campo.

TERMOS DE INDEXAÇÃO: Doenças da reprodução, micoplasmoses, vacas leiteiras, muco vaginal, Ureaplasma spp.

INTRODUCTION

Mollicutes are bacteria that differ from other prokaryotes due to the lack of a cell wall and their genome size, which is the smallest reported (Herrmann 1992, Dybvig & Voelker 1996). These bacteria inflict a wide range of diseases on livestock and are generally associated with clinical manifestations such as pneumonia, conjunctivitis, polyarthritis, mastitis, agalactia, abortion and infertiltiy (Nicholas 2002, McAuliffe et al. 2005). In cattle, the species of most relevance are in the group known as the Mycoplasma mycoides Cluster (MMC): M. mycoides subsp. mycoides Small Colony (MmmmSC); the agent of contagious bovine pleuropneumonia (CBPP); and other mycoplasmas species such as Mycoplasma bovis, Mycoplasma canadense, Mycoplasma arginini, Mycoplasma alkalescens, Mycoplasma bovirisin, Mycoplasma bovigetalium, Mycoplasma conjunctivae, M. mycoides mycoides capri, as well as Ureaplasma diversum and Acholeplasma laidlawaii (Landford 1975, Mulira et al. 1992, Razin et al. 1998, Frey 2002, Tenk 2005, Buzinhani et al. 2007, Marques et al. 2009, Santos et al. 2009). Some of these species are pathogenic, whereas others are considered ubiquitous of the normal flora in the mammary gland, as well as the respiratory and reproductive tracts (Whitford et al. 1994, Razin et al. 1998).

Mycoplasma spp. has often been associated with arthritis, pneumonia and otitis in neonatal calves (Yeruham et al. 1999, Gagea et al. 2006). M. bovis is one of the most economically important species of Mycoplasma in cattle herds, causing outbreaks of otitis, pneumonia and arthritis in calves, as well as highly infectious mastitis in cattle (Ghadersohi et al. 1999, Yeruham et al. 1999, Maeda et al. 2003, Francoz et al. 2004, Lamm et al. 2004, Radaelli et al. 2011). Economic losses caused by the syndrome (otitis-pneumonia-arthritis) in calves, as well as mastitis-agalactia in dairy cattle, have been previously studied in the Europe, United Kingdom, USA and Canada (Jasper 1981, Woldehiwet et al. 1990, Walz et al. 1997, Tenk 2005, Foster et al. 2007, Maunsell & Donovan 2009) and in the Brazil (Mettifogo et al. 1996, Pretto et al. 2001, Nascimento et al. 2005). M. bovirisin and M. alkalescens have also been isolated from cattle with mastitis (Jasper 1979, Jasper et al. 1979, Jasper 1981). Infected milk has been the source of Mycoplasma spp. in calves causing otitis and pneumonia. This agent is more prevalent in calves from herds with Mycoplasma mastitis than in calves from herds without Mycoplasma mastitis, and is considered a risk factor for outbreak infections in suckling calves (Jasper et al. 1979, Pfutzner 1990, Lamm et al. 2004).

Ureaplasma diversum colonizes the urogenital region, where the infection has been associated with different clinical manifestations. In cattle, the main disorders are vulvitis, granular vulvovaginitis, endometritis, salpingitis, placentitis, fetal alveolitis, abortions and the birth of weak calves. In bulls, the main disorders are the presence of seminal vesiculitis, epididymitis, balanoposthitis and alterations in spermatozooids (Panangala et al. 1981, Miller et al. 1983, Ruhrke et al. 1984, Piłaszek & Truszczyn Ski 1988, Cardoso et al. 2000, Nascimento et al. 2005, Oliveira Filho et al. 2005, Buzinhani et al. 2007, Rizzo et al. 2011). The agent is transmitted through organic secretions (semen, preputial and vaginal mucus) and direct contact during sexual intercourse, artificial insemination or embryo transfer (Kirkbride 1987, Hasso et al. 1993, Marques et al. 2009). In addition to reproductive diseases, Ureaplasma spp. has been reported in cases of destructive polyarthropathy (fever) and mastitis in cattle, although these cases are rare (Himsworth et al. 2009). The aim of the present study was to describe the occurrence of agents of the Mollicutes Class in dairy cattle from different geographical areas in northeastern Brazil.

MATERIALS AND METHODS

Geographic area. Five dairy cattle farms in the meso-region of the Agreste in the state of Pernambuco were visited in January of 2012. This geographical area is notable in the northeast of the country for its high milk production. The farms investigated were chosen at random in different cities, according to the following geographic coordinates: Garanhuns, municipal districts (Garanhuns: L 0773162 and UTM 9019050; Jupi: L 0565436 and UTM 9056244; Jurema: L 0813401 and UTM 9033624), Vale do Ipanema (Águas Belas: L 0716507 and UTM 8990870) Vale do Ipojuca (Alagoinha: L 0739156 and UTM 9059012), giving a total of five municipal districts (Fig.1).

Fig.1. Geographic area of the investigation for the presence of Mollicutes in dairy cattle in Brazil.
Cows and sampling reproductive tract. In total, 100 cattle of different ages, breeds and stages of lactation were examined in the present study. The cattle were part of intensive or semi-intensive management systems and were submitted to mechanical milking or hand milking. The sampling was performed using a stratified sampling farm, as described by Pereira (2003). Twenty, 15, 30, 23 and 20 cattle were randomly chosen from farms A, B, C, D and E, respectively.

The cattle were examined post milking. The vulvar and vaginal region of each animal was inspected for the presence of mucopurulent and purulent vaginal secretions or lesions. The medical history of the cattle herd was investigated for detected reproductive failure, endometritis, repeat breeding, infertility, the birth of weak calves and abortions. The samples were collected after washing the vulvar region with water and soap, and then drying it with paper towels and disinfecting the area with alcohol (70%GL). Vaginal mucous was collected using a sterile alginate cotton swab, which was rubbed on the vagina, as well as the lateral and internal walls. The samples were collected in sterile vials which were labeled with the number of the animal and farm. The samples were dipped into transport solution (phosphate buffered saline, pH 7.2), maintained at 4°C in a cooler with ice and taken to the Laboratory of Infectious Diseases (LDIC-DMV/UFRPE).

Bacteriological procedures. The vulvovaginal mucous samples were cultured for mycoplasmas and ureaplasmas, as described by Razin & Tully (1996) and Ruhmk & Rosendal (1994). Previously, 2mL of each sample were filtered through a sterile syringe holder coupled with a millipore membrane (0.45µm diameter). Then, 100µL of the filtrate was diluted up to 10⁻¹ at 10⁻⁵ and inoculated in both liquid and solid modified Hayflick’s medium, that optimizes the isolation of Mycoplasma classic and UB medium, selective for the propagation of ureaplasmas. Subsequently, all samples were incubated at 37°C for a minimum period of up to 21 days. The plates were placed in a jar to determine the microaerophilic conditions. The agar plates were checked daily for the presence of colonies using a stereomicroscope (40X). Blind sub cultures were used for up to 60 days, before disposal of the samples. Mollicutes isolates were confirmed by Dienes probe; Mycoplasma genera was used a digitonin sensitivity test and Ureaplasma spp. colonies were identified by the urease production in the U4 medium and their characteristic colony morphology (Whitford et al. 1994).

DNA extraction and PCR assays. The DNA was extracted from the samples by the method described in the commercially available kit (DNA Easy Blood and Tissues Kit®, Qiagen Biotechnology, guideline page 25). The primers (GPO-3 and MGSQ) that amplify the V6 and V7 conserved regions of the 16S RNA gene, specific to Mollicutes class was used as a procedure for screening of field samples (Van Kuppeveld et al. 1992). The samples were also submitted a PCR assays to detect Ureaplasma spp. using primers UGP-F’ (gene location 257-256) and UGP-R’ (gene location 862-881) conserved regions of the 16S RNA gene for Ureaplasma spp. (Lauerman 1998). The PCR assays for Mollicutes were performed in a mixture prepared with a volume of 25µL, containing 5µL of DNA template, 30pmol of each primer, MgCl₂ (1.5mM), buffer (10mM of Tris-HCl, pH 8.3), a mix of deoxynucleoside triphosphate (50µM), Taq DNA Polymerase (2.5U) and Milli-Q ultrapure water. For Ureaplasma spp., PCR assays were performed with primers at 30pmol, Milli-Q ultrapure water and 6.25µL of TopTa手工Mastermix, following the manufacturer’s instructions. In vitro amplifications were performed using thermovcycler model PTC-100 (MJ-Research®). Ultrapure water was used as a negative control. Mycoplasma mycoides (reference strain GM12) and Ureaplasma diversum (reference strain GMU132, Vup6) were used as positive controls in each reaction, respectively. The amplified PCR was visualized by electrophoresis in 1.5% agarose gel with 100bp molecular weight marker, colored with Bluegreen, viewed under ultraviolet light and photodocumented.

RESULTS

On the farms studied, the cattle examined were asymptomatic for mycoplasmosis and other reproductive disorders during the period of the research. In the swabs from their reproductive systems, the total prevalence for Mollicutes was 19.0% (19/100) in the isolation and 39.0% (39/100) in the PCR. In the samples cultivated in agar Hayflick’s medium, 13.0% growth was recorded and it was possible to visualize flask-shaped cells, which in the digitonin test was confirmed as Mycoplasma spp. In the agar UB medium, 6.0% of the samples exhibited characteristics of Ureaplasma spp. typical colonial morphology of brown gold colonies, confirmed by urease test. The subculture was made for growth confirmation of Mycoplasma spp. and Ureaplasma spp. The PCR confirmed 26.0% for Mollicutes the amplicon obtained was of 270bp and 13.0% for Ureaplasma spp. with amplicon of 644bp (Fig.2).

DISCUSSION

Mycoplasmas are ubiquitous in nature and frequently contain microbiota from different systems, such as the ear, eye, joints and mammary gland, as well as the respiratory and genitourinary tract in animals, humans and insects (Base- man & Tully 1997, Dybvig & Voelker 1996). The findings of the present study confirmed that Mycoplasma spp. e Urea- plasma spp. strains grew on both solid and liquid media and were confirmed at the genera level by the PCR. The presence of U. diversum in the reproductive tract of cattle

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**Fig.2. Results of the PCR for Ureaplasma spp. in the vaginal mucous of dairy cows, state of Pernambuco, Brazil.** Lane M, molecular size marker (Amresco® 1kb, ampliseize standard 500-10,000); Lanes 1,2,4,5,7,9,11 DNA Ureaplasma spp.; Lane (C-), negative control; Lanes (C+) Positive control.
with or without symptoms of mycoplasmosis has been reported in the literature by several authors (Oliveira Filho et al. 2005, Buzinhani et al. 2007, Razin et al. 1998). These agents in the reproductive tract of cattle may or may not be associated with outbreaks of reproductive disorders. However, the genus *Ureaplasma* spp. has often been associated with disease outbreaks (Albertsen 1955, Onoviran et al. 1975, Britton et al. 1987, Le Grand et al. 1995, Cardoso et al. 1997, Nascimento et al. 1998, Cardoso et al. 2000, Cardoso & Vascencelos 2004, Nascimento et al. 2005). In cattle, it can cause placentitis, fetal alveolitis, granular vulvovaginitis, abortion and the birth of weak calves and infection. In bulls, it may result in seminal vesiculitis, balanoposthitis and alterations in spermatozooids (Howard et al. 1976, Mulira & Saunder 1994, Himsworth et al. 2009, Marques et al. 2009). The occurrence of *Ureaplasma* spp. in cattle could be a potential risk for manifestations of reproductive diseases that have already been reported in the literature. The possible transmission route is through direct contact during sexual intercourse or by artificial insemination. This bacteria is shed in semen, milk, preputial and vaginal mucus (Kirkbride 1987, Britton et al. 1988, Cardoso & Vascencelos 2004).

In Brazil, there are very few studies reporting *Ureaplasma* infection in cattle herds. However, Nascimento et al. (1998) isolated *M. bovirhinis* in the vaginal mucus of heifers, highlighting the importance of this species as a cause of reproductive failure in this country, as well as a probable decrease in milk and meat production, similar to what occurs in other countries. *M. bovis* was diagnosed in 57% of the samples of vaginal mucus from cattle with reproductive disorders (vulvovaginitis, abortion, stillborn) and the authors highlighted the risk of venereal transmission of *Mycoplasma* spp. (Nascimento et al. 2005). In the northeast of Brazil, Santos et al. (2013) reported *Mollicutes* with frequencies of 65.6% and for *Ureaplasma diversum* of 15.6% in reproductive disease outbreak in cattle in Paraíba, the cows presented heat repetition, granular vulvovaginitis and abortions.

In the present study, the occurrence of *Mycoplasma* and *Ureaplasma* spp. in the vaginal mucus was low in comparison to the frequencies reported in other Brazilian states (Cardoso et al. 1997, Cardoso et al. 2000, Cardoso & Vascencelos 2004, Buzinhani et al. 2007, Santos et al. 2013). However, it was possible to prove the viability of these microorganisms through bacterial isolation, thus highlighting the possibility of these agents causing reproductive diseases in these cattle during opportunistic outbreaks. Further studies are required to prove the species and pathogenicity of these isolates. In the cattle assessed in the present study, there was no history of reproductive disorders or abortion, although the presence of these agents in the vaginal mucus of cows could be considered a risk factor for reproductive disease.

**CONCLUSION**

It was possible to verify the occurrence of *Mycoplasma* spp. and *Ureaplasma* spp. in dairy cattle from different geographical areas of the state of Pernambuco. This is the first report of these bacteria in this region of Brazil. Further studies are necessary to determine the species and pathogenic potential of these field isolates.

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**Conflicts of interest**- The authors have no conflicts of interest to declare.

**Research Ethics Committee**- This project was approved by the ethics committee of the UFRPE/DMV Process nº 23082.004146/2011-28.

**REFERENCES**


