Winter dysentery is an epizootic acute infectious and contagious disease with a seasonal trend that affects adult bovines worldwide, with a higher prevalence in dairy cows, caused by Bovine Coronavirus (BCoV), with cases of association with salmonella sp, rotavirus, Bovine Viral Diarrhea Virus, Cryptosporidium parvum and Eimeria bovis (Campbell and Cookingham, 1978). The disease was first reported in Brazil in 2002 in a dairy farm in São Paulo State (Brandão et al., 2002).

The annual outbreaks occur during the winter and last up to 10 days, most times affecting 100% of cows in a farm; affected animals present an acute and severe dysentery, dehydration, loss of weight and agalacty leading to an average 90% fall in milk production.

Besides winter dysentery, BCoV also causes diarrhea in newborn calves and respiratory illness in older calves (Heckert et al., 1990; Tegtmeier et al., 1999).

Surveys on the occurrence and etiology of enteric diseases in cows are highly valuable to the control of ongoing outbreaks and to develop more efficient preventive measures to the coming ones, once they allow the depiction of the relations between the occurrence of these diseases and their causes.

This paper reports a survey for BCoV in stool samples from cows collected during outbreaks of winter dysentery using a nested RT-PCR assay targeted to the RdRp gene of the virus and rotavirus using PAGE.

Stool samples were collected from 13 adult dairy cows with dysentery from dairy farms in São Paulo State, Southern Brazil, a region under tropical temperate weather, in May 2000 (1 sample), June 2001 (8 samples) and July 2002 (4 samples). These are fall (May and June) and winter (June and July) months in Brazil. No outbreaks have been reported in these farms in any other month.

Samples were prepared as suspensions in phosphate buffer solution 0.01M/BSA 0.1% pH 7.2 (PBS) and clarified by centrifugation (12,000xg/30 minutes), the supernatant stored at -80º until analysis.

The samples were tested for the presence of BCoV with a group II-coronaviruses-specific RT-PCR assay targeted to the RNA-dependent RNA-polymerase gene (RdRp) with a 136-bp predicted product (Brandão et al., 2005). BCoV Kakegawa strain (Akashi et al., 1980) was used as positive and PBS as negative controls.

Ten microliters of the nested PCR product were analyzed in 1.5% agarose gel electrophoresis stained with 0.5µg/ml ethidium bromide.
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The 13 samples were searched for rotavirus 11-segmented RNA in PAGE (polyacrylamide gel electrophoresis) according to Herring et al. (1982).

All stool samples were found positive to BCoV in the PCR targeting the RdRp gene as described above, evidenced by the appearance of the predicted 136-bp fragment, as seen to the BCoV Kakegawa positive control. Neither the nested negative controls nor the reactions relative to PBS showed bands. No non-specific band or laboratory contaminations were detected. Furthermore, none of these samples tested positive for rotavirus in PAGE.

The nested PCR used herein was proven to be a useful tool to detect BCoV in stool samples, allowing a fast and reliable diagnosis of this virus, mainly because the RdRp gene is a highly conserved region in genus Coronavirus (Stephensen et al., 1999), where low variation in nucleotide sequence is expected.

The finding of BCoV in stool samples of cows with dysentery, in which no rotavirus was found, suggests that this virus may be a primary agent involved in the etiology of the cases here reported and strengthens the already indicated role of BCoV in winter dysentery. Also, the disease described here has the same seasonal pattern already reported in similar outbreaks (Alenius et al., 1991; Cho et al., 2000), occurring in winter, cold months.

Furthermore, virus-shedding cows, whether with or without symptomatic enteritis, may serve as reservoirs for calves, in which BCoV is known as a cause of severe neonatal diarrhea (Clark, 1993) which, in their turn, may serve as reservoirs to the adult.

Thus, besides allowing preventive measures against winter dysentery itself, surveys on etiological agents of diarrhea in cows may also provide useful information to the control and prevention of diarrhea in calves.

As a conclusion, winter dysentery was detected during three consecutive outbreaks with the classical clinical and seasonal patterns of the disease and in association with Bovine Coronavirus. Continued studies including other agents, as well as phylogenetic analyses of the BCoV strains found, are being carried out with samples from the surveyed area.

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Keywords: bovine, winter dysentery, coronavirus, PCR, diagnosis.

RESUMO

Descreve-se a pesquisa de BCoV e rotavírus em 13 amostras fecais de vacas de surtos de disenteria utilizando uma nested PCR dirigida ao gene RdRp e PAGE, respectivamente. Todas as amostras fecais foram positivas para BCoV e nenhuma delas apresentou-se positiva para rotavírus em PAGE. O encontro de coronavírus bovino em amostras fecais de vacas com disenteria sugere que este vírus possa ser o agente primário envolvido na etiologia dos casos aqui relatados.

Palavras-chave: bovino, disenteria de inverno, coronavírus, PCR, diagnóstico

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REFERENCES


BRANDÃO, P.E.; BIRGEL Jr., E.H.; GREGORI, F. et al. Bovine coronavirus...


