## MOLECULAR EPIDEMIOLOGY OF CRYPTOSPORIDIOSIS ON CATTLE FARMS IN THE CANARY ISLANDS

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## INTRODUCTION

Cryptosporidiosis is a disease caused by different species of the genus Cryptosporidium, a protozoan parasite capable of causing large economic losses in ruminant livestock farms, as well as diarrhoea in humans. The objective of this study has been to determine the prevalence of Cryptosporidium species in cattle from farms in the Gran Canaria by both microscopic and molecular tools.

## MATERIALS AND METHODS

Survey: risk factors + economic impact (21 questions).

3 Microscopical analysis: sedimentation + Kinyoun carbol fuchsin stain.

2 Sampling: 8 faeces calves + 8 faeces adults ( 15 farms)
4. Molecular analysis:

DNA extraction with QIAamp Fast DNA Stool Mini Kit ${ }^{\circledR}$ and PCR (18S and PAR60).


Figure 1: calf with diarrhoea.


Figure 2: Kinyoun stain + for Cryptosporidium.

## RESULTS

- All farms were positive for Cryptosporidium by Kinyoun staining.
- The number of positives detected by PCR was higher than those detected by Kinyoun, especially in adults.
- A higher number of positives were detected with the 185 marker because PAR60 is specific to Cryptosporidium parvum.
- Survey results showed that $90 \%$ of the farmers were unaware of the parasite and did not apply any control measures.


Table 1: comparison of Kinyoun staining and PCR results.

## CONCLUSIONS

The results of this study show a high prevalence of cryptosporidiosis in cattle in the Canary Islands, with a high incidence of the zoonotic species Cryptosporidium parvum. Surprisingly, most farmers are not aware of the clinical and public heath implications of this fact.

ID: CRYPTO4

