

Effect of *H. contortus* infection on parasitological and local cellular responses of Creole kids

J.-C. Bambou, T. Larcher, W. Ceï, P.-J. Dumoulin, N. Mandonnet

INRA-URZ Guadeloupe, FWI





Introduction

Small ruminants (goats and sheep) are constantly exposed to gastrointestinal nematode infections

>Major cause of morbidity and mortality: profitability

Anthelmintics resistance is a widespread problem

Need for alternative control strategies: To reach a favorable equilibrium for animal production between nematode parasites and their host





Genetic variability of resistance to GIN in goats and sheep

Intra and inter-breed genetic variability :

➢Goats: 0.15 < h² < 0.2</p>

Sheep: 0.3 < h² < 0.4</p>

Closely associated with the host immune response

Mechanisms underlying genetic resistance?
Most of the studies in sheep
Limited knowledge of the goat response



The experimental model

Creole goats of Guadeloupe

Local crossbred meat breed maintained in a production flock



➢Grazed all year on pasture under natural mixed infection (*H. contortus, T. colubriformis*)

➤Genetic variability available for selection

➢Underlying mechanisms under investigation







To evaluate the relationships of cellular changes in the abomasal mucosa and parasitological parameters by comparing resistant and susceptible kids after experimental infection with *Haemonchus contortus*









Animals and experimental design

- Growing female Creole kids at 8-month old (n=28)
 - Maintained nematode-free (naïves)
 - Average predicted breeding values on FEC distant of 1.04 genetic standard deviation
 - Experimental H. contortus infection (10,000 L3)



At slaughter: Nematode burden, Histopathological analysis



PCV



Anaemia observed after challenge 1 but not challenge 2
No difference between resistant and susceptible kids





FEC higher after challenge 1
FEC higher in susceptible kids after both challenges





>ADG lower in susceptible kids after both challenges



Nematode burden

	Challenge 1			Challenge 2			- 50
	¹ R	² S	³ P value	¹ R	² S	⁴ P value	³ P value
⁶ Male	234±66	120±14 7	0.49	203±66	195±65	0.93	0.81
⁷ Female	241±59	118 ± 72	0.19	234 ± 59	178 ± 59	0.51	0.45
⁸ Immature male	5 ± 7	31±8	0.65	15 ± 8	36±9	0.21	0.02
⁹ lmmature female	19 ± 18	24 ± 22	0.56	46 ± 18	71 ± 18	0.53	0.05
¹⁰ Female length	18.8±0. 5	21.0±1. 0	0.07	18.0±5	18.2±0. 6	0.8	0.02

No effect of the genetic status:

Iow number of animal/group?

 After challenge 2: higher number of immature nematode reduction of female length

Protective immune response?



Histopathological analysis

Antrum









Cellular infiltration more pronounced after challenge 1

No effect of the genetic
status for eosinophils and
mononuclear cells

Globule leucocytes
infiltration higher in resistant

Cellular functionality?

st chalenge

2nd chalenge

Conclusion

Analysis of the global response resulting from the interaction nematode-host: FEC and ADG

Significant differences are evidenced

BUT

 Analysis of « finer » responses : Histopathological changes at targeted time point

differences are less marked

Characterisation/understanding of the complex cross-talk between two organisms:

Dynamic of finer responses more pertinent than targeted time point ?

> Need for development of alternatives experimental strategies





This study was supported by:



Sustainable Solutions for Small Ruminants







SEVENTH FRAMEWORK PROGRAMME

EAAP 2013, Nantes, FRANCE