



First rabbit gut microbial gene catalog and metagenomic insights into the resistome: links with antibiotic use

Caroline Achard, Yuliaxis Ramayo Caldas, Amine Ghozlane, Mathieu M. Almeida, Samuel Boucher, Stéphane Chatellier, Adeline Chaubet, Sylvie Combes, Catherine Denis, Benoit Dile, et al.

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FIRST RABBIT GUT MICROBIAL GENE CATALOG AND METAGENOMIC INSIGHTS INTO THE RESISTOME : LINKS WITH ANTIBIOTIC USE

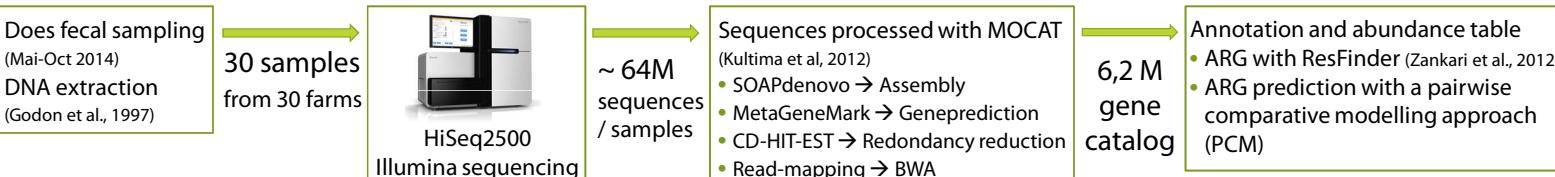
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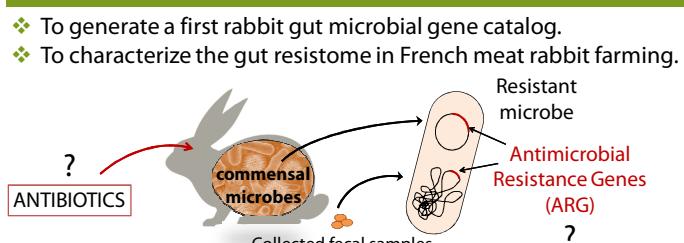
INTRODUCTION

Microbial antibiotic resistance genes emerging in livestock are a major public health concern. It is crucial to consider the prevalence of Antibiotic Resistance Genes (ARG) in commensal microbes which represent the vast majority of the gut microbiota of healthy individuals and can convey ARG between animal and human. ARG can be carried by mobile genetic element and transferred between bacterial species, ending up in human pathogens.

METHODS



OBJECTIVES



RESULTS

ANTIBIOTIC USES

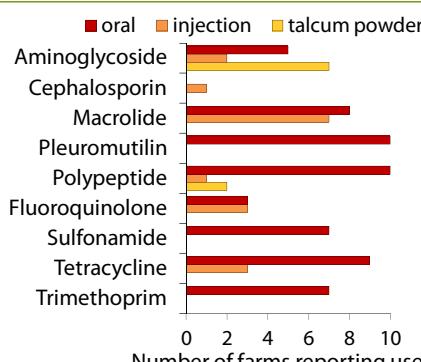


Figure 1 : Number of different antimicrobial molecules used in the rabbit farm panel per class and administration mode.

A global link between antibiotic uses and the sum of the relative abundance of all ARG detected.

Farm	# antibiotics used	# detected ResFinder ARG	Sum of ResFinder ARG abundance
02	0	17	5,49E+03
12	0	18	1,06E+04
05	0	23	1,09E+04
32	1	44	1,53E+04
26	0	43	2,22E+04
33	0	18	2,74E+04
01	0	23	4,58E+04
16	4	18	5,19E+04
18	0	49	5,24E+04
15	4	16	5,71E+04
13	1	22	5,94E+04
27	3	20	6,35E+04
17	3	24	6,51E+04
04	2	21	6,55E+04
24	2	18	6,73E+04
23	4	19	6,85E+04
14	1	61	6,95E+04
30	8	28	7,17E+04
11	4	21	7,53E+04
22	5	62	7,71E+04
03	6	20	7,71E+04
19	4	23	7,78E+04
25	5	24	8,11E+04
31	8	34	8,38E+04
10	5	49	8,74E+04
28 ^a	0	68	9,04E+04
08 ^b	1	50	9,13E+04
21	6	68	1,07E+05
20	7	77	1,94E+05

Oral exposure to antibiotics

Non oral exposure to antibiotics

^a oral exposure to antibiotics until end-2013

^b oral exposure to antibiotics until end-2012

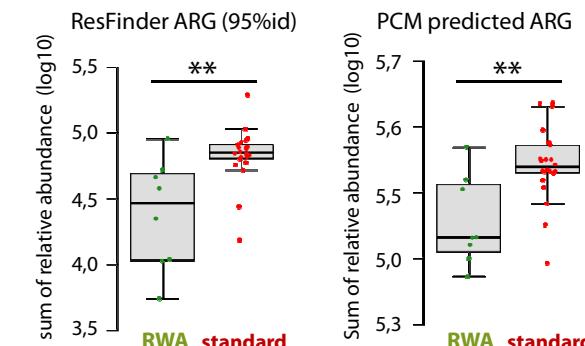
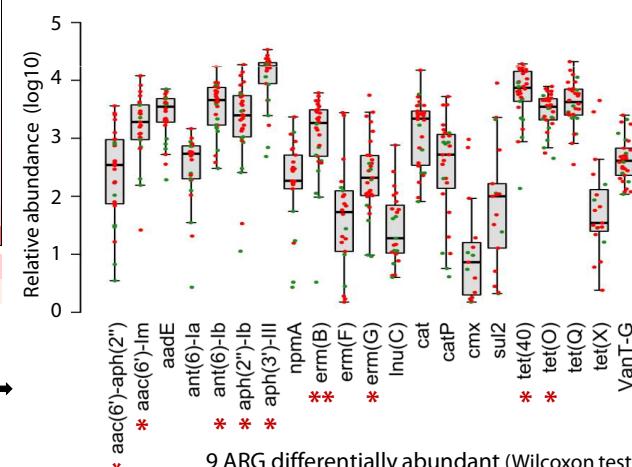


Figure 3 : Sum of relative abundances of all ARG in Raised Without Antibiotic (RWA) farms and in standard farm using antibiotics.



CONCLUSION

- First rabbit gut microbial gene catalog : 80 antibiotic resistance genes (ARG Resfinder) and 8235 ARG candidates (pairwise comparative modelling approach).
- Decreasing the antibiotic use decreases the global abundance of antibiotic resistance genes.
- Some antibiotic resistance genes persist in antibiotic farms reducing their antibiotic use : an active strategy is needed (see poster 328B)

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