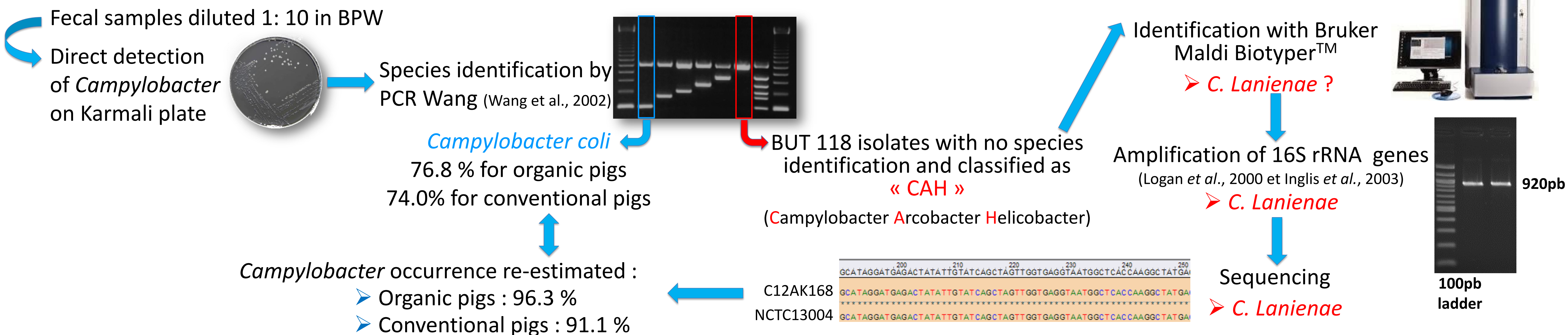


First description of *Campylobacter lanienae* from feces of organic and conventional pigs in France

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Context of the detection of *C. lanienae* isolates

In order to evaluate *Campylobacter* occurrence, antimicrobial resistance and genotypic diversity, fecal samples of 58 pigs from 31 conventional herds and 56 pigs from 31 organic herds, were collected in a slaughterhouse at evisceration step. The analysis of fecal samples was performed as following :



Characterization of *C. lanienae* isolates

Antimicrobial susceptibility

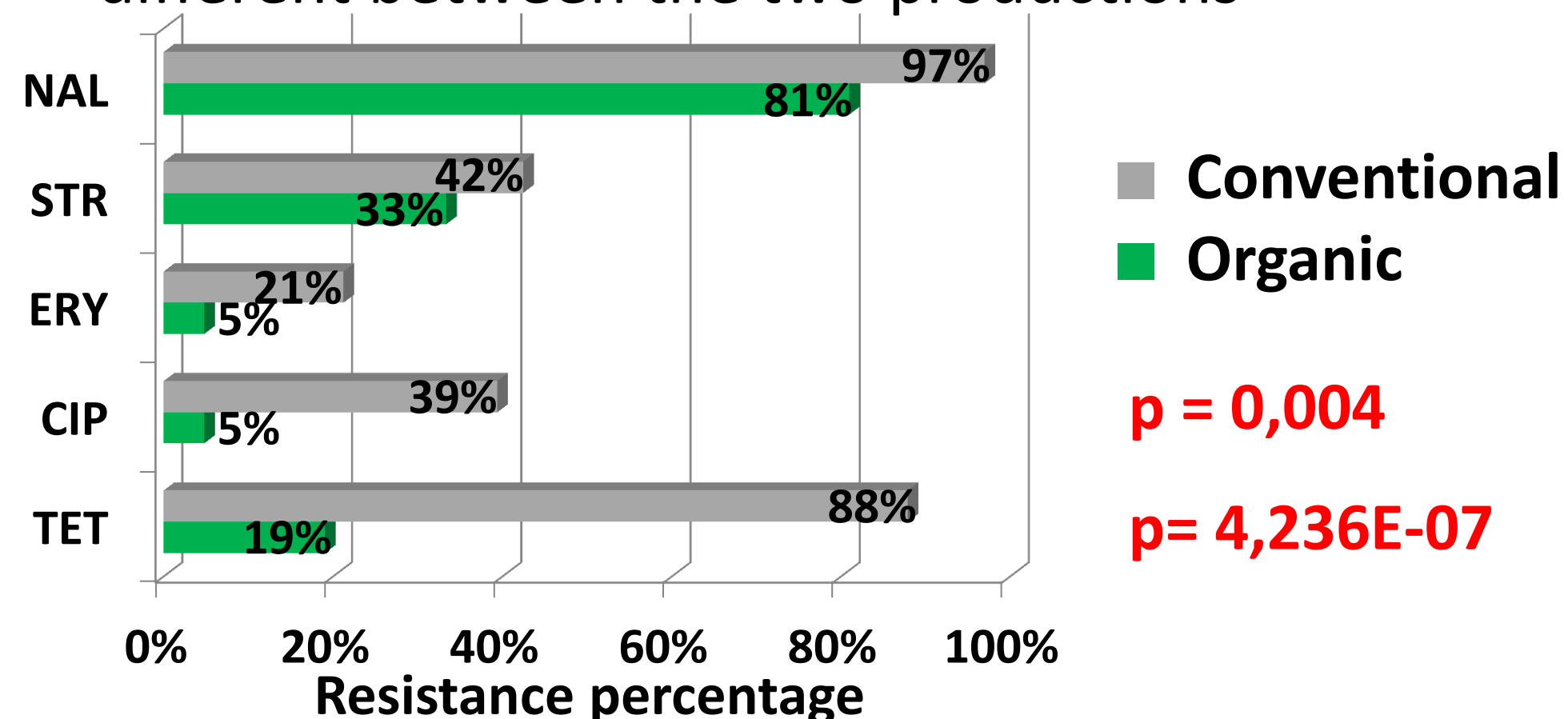
Method

- 55 *C. Lanienae* studied for their antimicrobial susceptibility by Minimal Inhibitory Concentration (MIC) using Sensititre® plates (Biocentric, Bandol, France)
- 7 antimicrobials tested : Gentamicin (GEN), Streptomycin (STR), Ciprofloxacin (CIP), Nalidixic Acid (NAL), Tetracycline (TET), Erythromycin (ERY), Chloramphenicol (CHL).
- Results analysed following ECOFFs from Eucast.

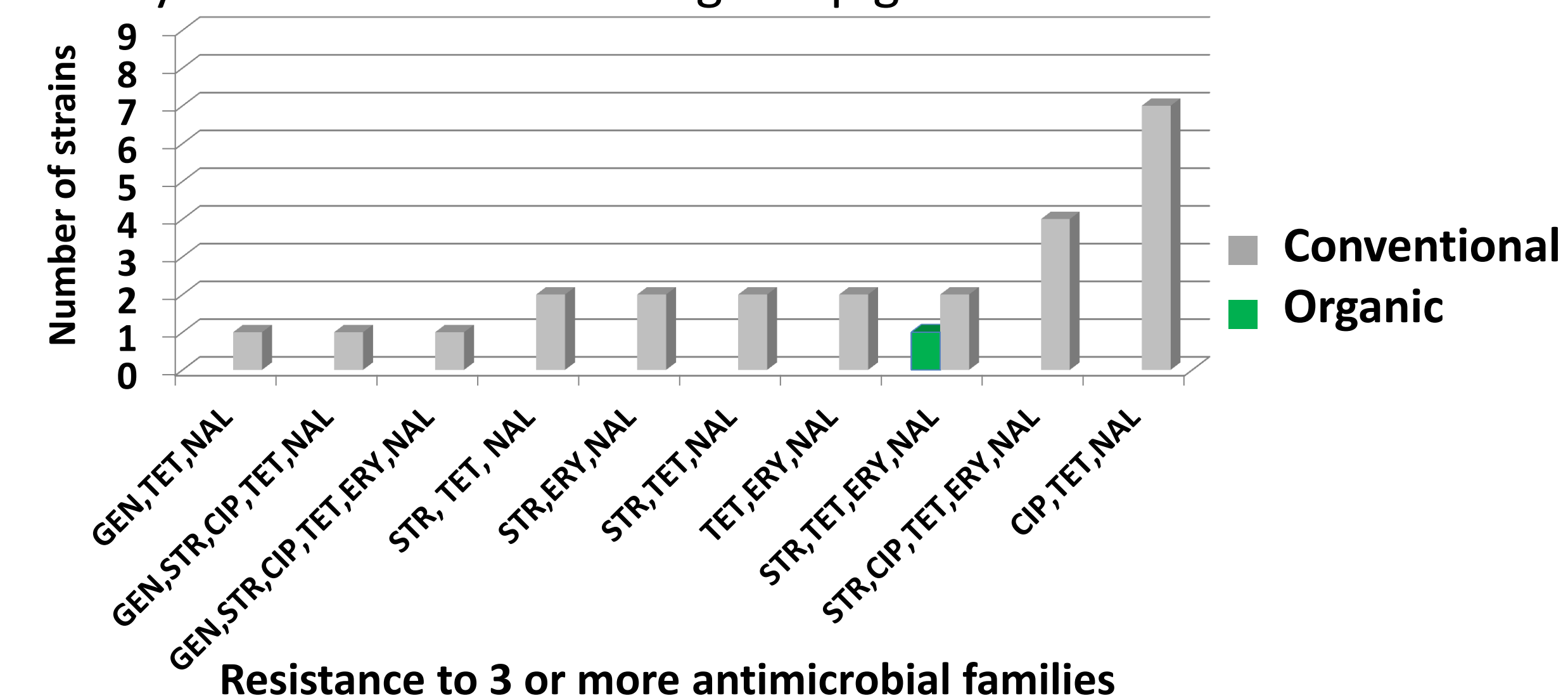


Results

- Only one isolate was pansusceptible (1.8%)
- All isolates were susceptible to Chloramphenicol and 94.5% susceptible to Gentamycin
- Resistance to Nalidixic acid (93 %) is very high : natural resistance
- Resistance to Tetracycline and Ciprofloxacin was significantly different between the two productions



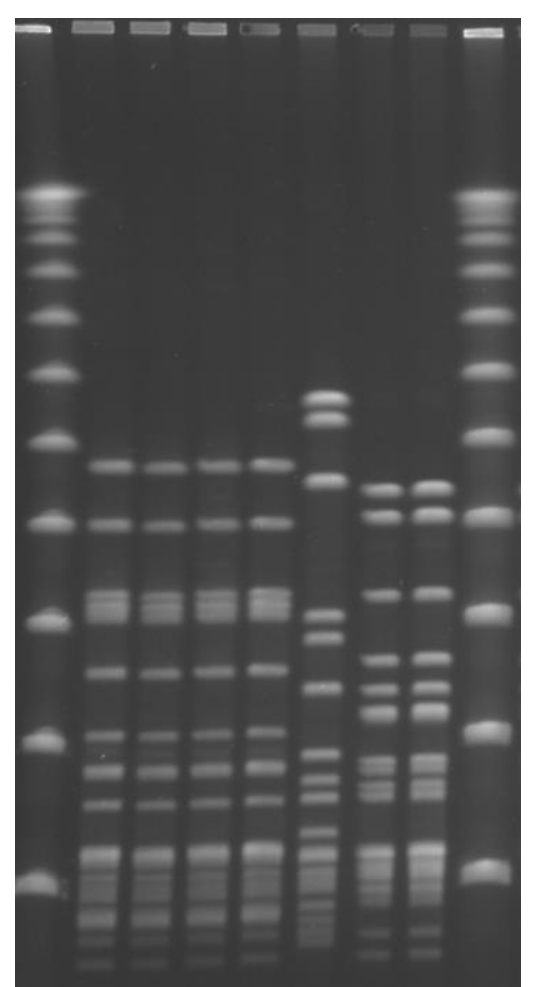
- 17 resistance patterns were identified
- Isolates from conventional pigs were mostly multiresistant (73%) vs only 5% of isolates from organic pigs



Genotypic diversity : Pulsed-field gel electrophoresis

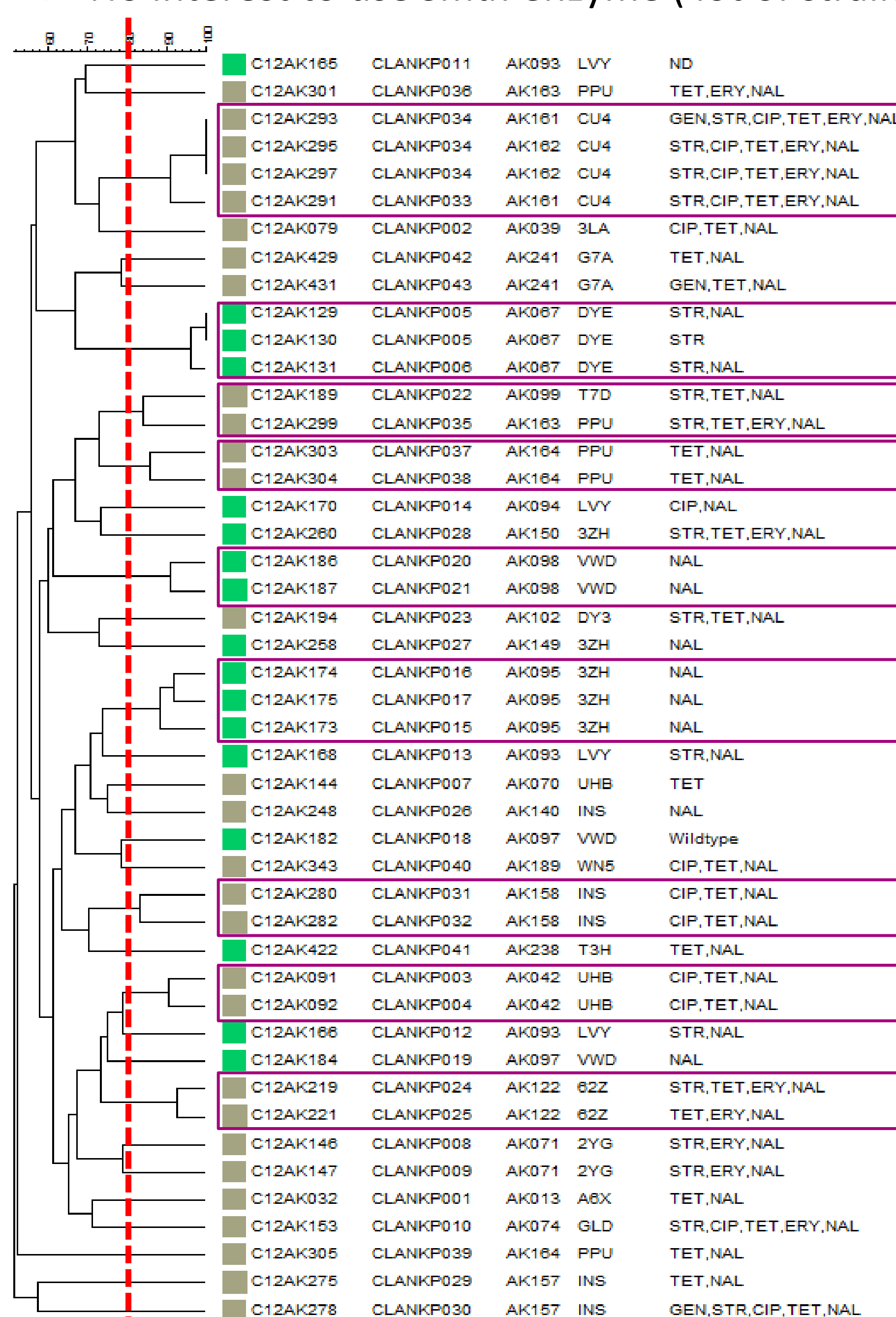
Method

- DNA preparation, restriction endonuclease digestion and PFGE carried out as described by the Campynet protocol
- DNA macrorestriction performed with *KpnI* and *SmaI* enzymes.
- Electrophoretic patterns compared using BioNumerics® (Applied Maths, Sint-Martens-Latem, Belgium).
- Simpson's index (D) used to assess the genetic diversity of the *Campylobacter* populations (Hunter & Gaston, 1988).



Results

- High diversity whatever the origin of strains, and the enzyme used (ID > to 0.98)
- No interest to use *SmaI* enzyme (lot of strains no typable)



At 80% of similarity :

- 9 clusters
- No evidence of genetic clusters linked to a type of production or to a resistance pattern
- When isolates showed a same PFGE pattern, they are from the same sample or same herd
- The patterns are distinct when they were compared with the others *Campylobacter* species patterns

Electrophoretic patterns compared using BioNumerics® software

Conclusion

This study allowed us to demonstrate for the first time in France that pigs, known to be a reservoir for *Campylobacter coli* may also carry in their feces *Campylobacter lanienae*, a species rarely highlighted. The species was present in conventional fecal samples as well as organic fecal samples. The lower level of antibiotic resistance and multiresistance of *C. Lanienae* strains for organic pigs may be related to the restricted use of antibiotics in this production and / or colonization of organic pigs with susceptible environmental strains. The genotypic diversity by RFLP-PFGE is very high, as generally observed for other more common species of *Campylobacter*.