





MINISTRY OF FOOD AND DRUG SAFETY

National Institute of Food and Drug Safety Evaluation



# WGS and antimicrobial resistance

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# WGS and antimicrobial resistance How useful is WGS? (Can <u>non-bioinformaticians use it?</u>)

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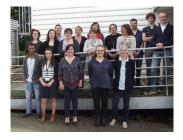
Antimicrobial Resistance NRL





#### Anses Ploufragan, MBA Unit: our activities on antimicrobial resistance





#### • Monitoring AMR in poultry, pigs, fish and their environment

- Zoonotic bacteria (Campylobacter, Decision 2013/652/EU)
- Commensal bacteria: *E. coli* from healthy animals
- Pathogenic bacteria: Resapath network (E. coli) and Mycoplasmas

#### Impact of agricultural or veterinary practices

- Antimicrobial administration, manure treatments...
- Impact on selection of resistant bacteria, on the microbiote, on the resistome

#### Alternatives to antimicrobials

Probiotics, bacteriocins, vaccines...



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• ESC: critically important antibiotics for human health

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- ESC-Resistance (ESCR): Increased incidence in humans since 2000s
- ESC-Resistance present also in bacteria of animal origin
- Resistance genes are mostly borne by conjugative plasmids and encode Extended Spectrum Beta-Lactamases (ESBL) or AmpC beta-lactamases











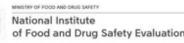
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# **Resistance to ESC: Questions**



- Diversity of the resistance mechanisms and their genetic supports, in commensal and pathogenic bacteria?
  - Characterization of isolates

- Origin, diffusion, persistence of resistance genes in animals, in farms, in the environment? →Longitudinal study in free-range broilers
- Diffusion or loss of resistance genes in animals? →Experimental studies in inoculated animals





• E. coli isolates resistant to ESC in poultry

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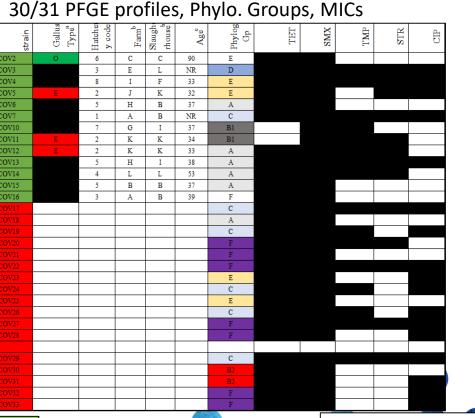
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- In 2010-2012: peak of the ESC resistance
- Diversity? Resistance genes? Virulence genes?
- ➢ Focus on 31 bla<sub>CTX-M-1</sub> isolates, most prevalent
  - Commensal (14) and pathogenic E. coli (17)
  - Various types of chickens (conventional, organic, free range)
  - Various geographic areas from France
  - Various hatcheries, farms, slaughterhouses





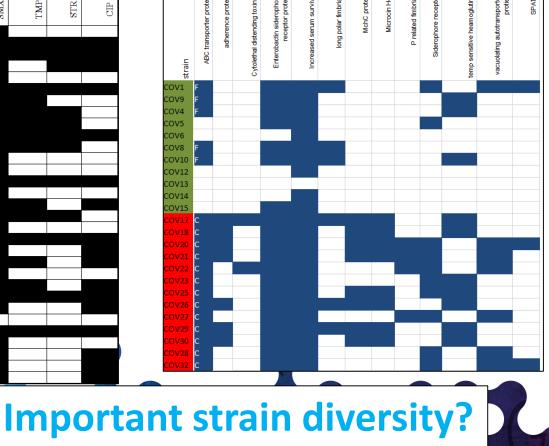
<u>Results 1/3: Conventional methods</u>



Fecal

Colibacillosis

#### Micro-array: virulence genes (Anjum et al.)









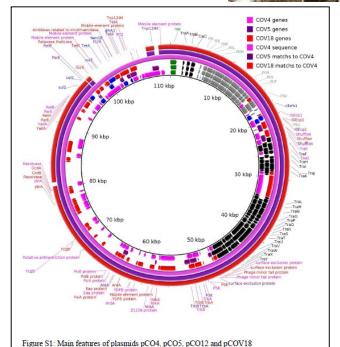
#### Results 2/3: Plasmids sequenced after transformation or conjugation

*bla*<sub>CTX-M-1</sub> on Incl1/pMLST3 plasmid in 30 isolates

#### **Resistance genes**

sul2: 27/30 tet(A): 20/30 dfrA17: 5/30 aadA5: 5/30 bla<sub>TEM-b</sub>: 1/30 dfrA1: 1/30

possible co-selection



**No APEC virulence genes** (*sitA-D,iucA-D , iutA, ompT, etsA-C, iss, iroB-E, iroN, cvaA-C, cvi*) although these genes were sometimes present in isolates

**Other genes**: toxin-antitoxin, conjugative transfer genes, genes involved in maintenance and stability regions, integrase...







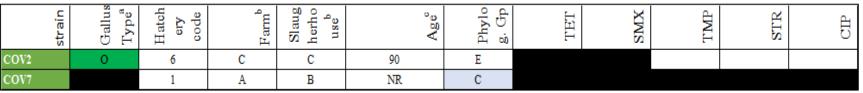


• Results 3/3: Comparison of sequences

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Comparison of the 30  $bla_{CTX-M-1}$  plasmids for the 114,313 common nt (but alignment could contain gaps), showed that overall they shared 67% identity.

Two plasmids from distinct commensal isolates showed <u>100% identity</u>



 One plasmid from a commensal strain and seven plasmids from colibacillosis shared more than 99% identity although the susceptibility/phylogenetic groups/virulence profile of their parental strains were different

### Diversity of strains but low diversity for ARG and plasmids in broilers isolates













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# Main results for ESCR pig isolates



- Mainly carried by highly similar *bla*<sub>CTX-M-1</sub> Incl1/ST3 plasmids.
- These plasmids very often bear sul2, dfrA17 and aadA5, and occasionally tet(A), mph(A), erm, floR or strA, strB. <u>Resistance genes</u> for quinolones or polymyxins were never detected.
- <u>Few virulence genes</u> (including colicins, heat-stable enterotoxins, adhesins or temperature-sensitive hemagglutinins).
- Occasionally: bla<sub>CTX-M-27</sub> and bla<sub>CTX-M-14</sub>, the latter being on an IncF plasmid which showed very close identity to a <u>human epidemic</u> plasmid.



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- Diversity of the resistance mechanisms and their genetic supports, in commensal vs pathogenic bacteria? → Characterization of isolates
- Origin, diffusion, persistence of resistance genes/plasmids in animals, in farms, in the environment?

### Longitudinal study in free-range broilers

Diffusion or loss of resistance genes in animals? → Experimental studies in inoculated animals







### Longitudinal study in free-range broilers



- Two hatcheries, 11 flocks, 9 farms
- Fecal samples collected in farms from D0 (before arrival) to D77
- Isolation of ESCR *E. coli*: characterization of isolates; plasmids sequenced from transformants or transconjugants





### <u>Results 1/4: *bla*<sub>CTX-M-1</sub></u><u>Incl1 pMLST3 plasmids</u>

15 *E. coli* (13 PFGE profiles) with*bla*<sub>стх-м-1</sub> Incl1 pMLST3 plasmids obtained from D0 to D77 from 7 flocks on 6 farms

Most contained the *tet*(A) and *sul2* genes.

Seven plasmids from isolates of different phylogenetic groups obtained from Day 2 to Day 41, from 5 flocks, 2 hatcheries shared sequence **identity higher than 99%.** 



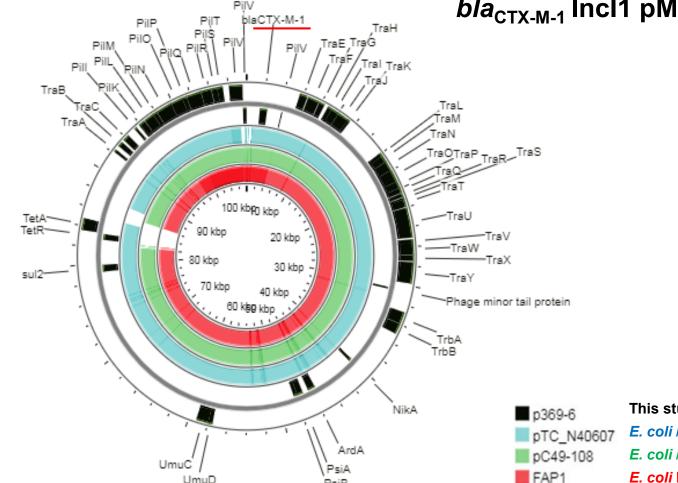
### <u>Results 2/4: *bla*<sub>CTX-M-1</sub></u><u>Incl1 pMLST3 plasmids</u>

But the  $bla_{CTX-M-1}$  Incl1 pMLST3 sequences were also very close (96% coverage, 99% identity) to:

- E. coli pC49-108 obtained from a chicken in Switzerland
- E. coli FAP1 from pig faeces in the Netherland
- *E. coli* pTC\_N40607 obtained from retail meat and cattle farms in the US...







*bla*<sub>CTX-M-1</sub> Incl1 pMLST3 plasmids

This study / *E. coli* / broiler / France *E. coli* / retail meat and cattle / USA *E. coli* / chicken / Switzerland *E. coli* FAP1 / pig faeces / Netherland



# Longitudinal study in free-range broilers



### Results 3/4: bla<sub>CMY-2</sub> isolates

10 isolates obtained from D0 to D7 with **bla<sub>CMY-2</sub>**, all from the same hatchery.

Six *E. coli* isolated from Day 0 to Day 7 from three flocks on F13 and F14, from chicks that had been hatched on the same day in the same hatchery were CTX-FOX-TMP-SMX-TET-CIP-CHL-resistant.





### Results 4/4: *bla*<sub>CMY-2</sub> plasmids

- 10 *bla*<sub>CMY-2</sub> Incl1 pMLST12, 110-117 kbp
- No resistance genes other than *bla*<sub>CMY-2</sub>, no virulence genes
- Very similar to each other (cover.98-100%, ident. 99-100%)
- And very similar to *E. coli* M63 plasmid from a pig in France (coverage 99%, identity 100%), but lacked a 13 kbp region (*aadA5, sul2 dfrA17*, IntI1, IS26, and mobile elements)
- And also very similar
  - pN13-01290 from S. Heidelberg from turkey meat in Quebec
  - pAMR588-04-00320 from S. Heidelberg from chicken faeces in Ontario
  - pSTY1-2010K-1587 from S. Typhimurium from human stool in the USA







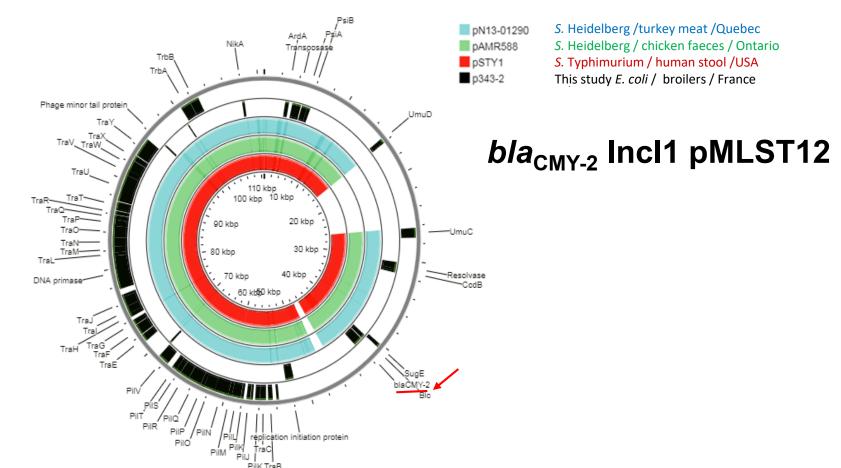


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## Longitudinal study in free-range broilers







- The Incl1 pMLST3 bla<sub>CTX-M-1</sub> plasmids were present in isolates obtained from Day 0 to Day 77 from both hatcheries, and were very similar both to each other and to plasmids very frequently described in French broilers, and in other animal species and regions. Their epidemic nature therefore makes it difficult to determine the origin of contamination.
- On the contrary, the Incl1 pMLST12 bla<sub>CMY-2</sub> plasmids are uncommon in French isolates from poultry. In this study, they were only obtained from the isolates of young birds from hatchery B, were very similar to each other, suggesting a probable vertical transmission.





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# **Resistance to ESC: Questions**

- Diversity of the resistance mechanisms and their genetic supports, in commensal vs pathogenic bacteria from different animal species and humans? Characterization of isolates
- Origin, diffusion, persistence of resistance genes/plasmids in animals, in farms, in the environment?
- Diffusion or loss of resistance genes in animals?
  Experimental studies in inoculated animals









# In vivo loss of multiresistance plasmid

- 4 different trials in controlled animal facilities, 51 SPF pigs
- Inoculated or contact pigs

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*E. coli* M63, rifampicin resistant; contains pESCR: Incl1 pMLST12 with *bla*<sub>CTX-M-1</sub>, *bla*<sub>CMY-2</sub>, *sul2*, *dfrA17*, and *aadA5* genes

TABLE 1 Design of animal experiments and culture results<sup>c</sup>

Trial	Group receiving M63 inoculation on day 0 (dose per pig)	Other inoculation(s) (dose per pig)	Fecal sample origin(s)	No. of samples	No. of rtfamptn- reststant <i>E. coli</i> tsolates	No. of CTX-susceptible E. coli isolates/total no. of isolates
04-12	Inoculated pigs (6.8 × 10 <sup>10</sup> CFU <sup>a</sup> )	Reinoculation with M63 on day 1	6 inoculated pigs sampled on days 1 to 11 and days 14, 16,18, 20, 22, <sup>b</sup> 24, <sup>b</sup> and 27 <sup>b</sup>	97	97	0/97
	Contact pigs (no inoculation)		4 contact ptgs sampled on days 1 to 11 and days 14, 18, 16, 20, 22, <sup>b</sup> 24, <sup>b</sup> and 27 <sup>b</sup>	66	55	2/55
10-12	All pigs (78 $\times$ 10 $^{9}$ CFU)	ED1a (days 8–12) (11.2 × 10 <sup>9</sup> CFU)	5 pigs sampled on days 2, 3, <sup>b</sup> 9, and 13 <sup>b</sup>	16	15	0/15
		EcN (days 8–12) (17 × 10 <sup>9</sup> CFU)	5 pigs sampled on days 2, 3, <sup>b</sup> 9, <sup>b</sup> 13, <sup>b</sup> and 16 <sup>b</sup>	14	13	0/13
01-13	All ptgs (78 $\times$ 10 $^{8}$ CFU)	ED1a (days 8–13) (4.4 × 10° CFU)	8 pigs sampled on days 1, 3, 7, 10, 13, 17, and 22	55	45	0/45
06-13	All pigs (73 $\times$ 10 $^{6}$ CFU)	No	8 pigs sampled on days 3, 10 <sup>b</sup> , 13, 16, 20, 23, 27, and 31 <sup>b</sup>	55	51	1/51
		ED1a (days 8–13) (74 × 10 <sup>6</sup> CFU)	7 pigs sampled on days 3, 10, <sup>b</sup> 16, 20, 23, <sup>b</sup> 27, and 31 <sup>b</sup>	42	42	0/42
		ED1a (days 8–13) (65 × 10 <sup>7</sup> CFU)	8 pigs sampled on days 3, 16, <sup>b</sup> 20, <sup>b</sup> 23, <sup>b</sup> 27, <sup>b</sup> and 31 <sup>b</sup>	36	36	0/36

" Pigs were inoculated on day 0 and day 1.

<sup>b</sup> Not all pigs were sampled.

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<sup>6</sup> There were 48 pigs inoculated with M63 that were not yet or not inoculated with ED1a or EcN with 200 samples, with 1 being a CTX-susceptible *E*. coli isolates; there were 4 contact pigs with 66 samples, with 2 being CTX-susceptible *E*. coli isolates; there were 28 pigs inoculated with M63 and then with ED1a with 19 samples, with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with ED1a with 19 samples, with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with ED1a with 19 samples, with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with ED1a with 19 samples, with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with M63 and then with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculated with 0 being CTX-susceptible *E*. coli isolates; and there were 4 pigs inoculates pigs inoculates on to 1353 Rif *E*. coli isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates ont 0 isolates; and there were 4 pigs inoculates



# *In vivo* loss of multiresistance plasmid

- Results
- 84% of 355 fecal samples from inoculated pigs and 83% of 66 from contact pigs contained *E. coli* M63
- Analysis of 353 Rif-R *E. coli* 
  - 350 positive by PCR for *bla*<sub>CTX-M-1</sub>, *sul2*, *lncl1*
  - 3 negative for  $bla_{CTX-M-1}$ , *sul2*, *lncl1*, no growth on CTX  $\rightarrow$  WGS







## In vivo loss of multiresistance plasmid

>pESCR covered 72 times on average for *E. coli* M63

8,

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- Average coverage depth of 1 for the three Rif-R CTX-S isolates
- $\blacktriangleright$  E. coli M63 and the three Rif-R CTX-S isolates share the same chromosomal sequences

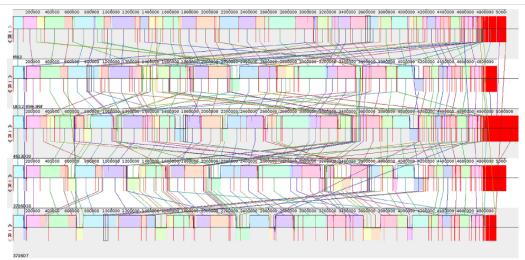


FIG 1 Chromosome assemblies of E. coli M63, UB12/059-3Rif, and the three rifampin-resistant cefotaxime-susceptible E. coli isolates reordered according to M63 with Mauve, aligned by using progressiveMauve. M63 is the inoculated strain; UB12/059-3Rif is the recipient strain used to prepare E. coli M63; and 3716D7, 3716D18, and 4613D20 are the rifampin-resistant cefotaxime-susceptible E. coli isolates obtained from pig 3716 on days 7 and 18 and pig 4613 on day 20, respectively.







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# Conclusion

### WGS is helpful for the study of:

- Diversity of resistant isolates, resistance genes, involved plasmids... in animals, humans and the environment
- > Detection of epidemic clones, plasmids
- Presence of different ARG (co-selection) or virulence genes on the same genetic support
- Origin, persistence, diffusion, loss of resistance, in vivo at different scales (animal, flock, population)



# Thank you to

e Bfr

- Mycoplasmology, Bacteriology and Antimicrobial Resistance Unit (S. Baron, C. de Boisséson, E. Jouy, E. Larvor, L. Le Devendec, G. Mourand)
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# Thank you for your attention

