RÉUNION INTERDISCIPLINAIRE DE CHIMICOHÉRAPIE ANTI-INFECTIEUS droitsteset

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LUNDI 13 & MARDI 14

Emergent aztreonam-avibactam resistance among NDM and OXA-48-Patrice Nordmann^{4,5,6}, Yancheng Yao^{1,2}, Linda Falgenhauer^{2,3}, Mustafa Sadek⁴, Can Imirzalioglu^{1,2}, and

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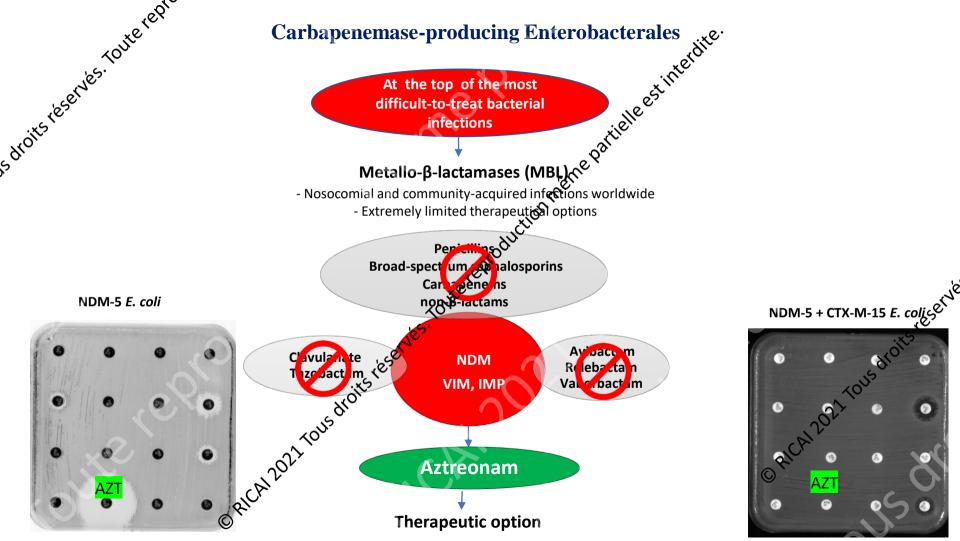
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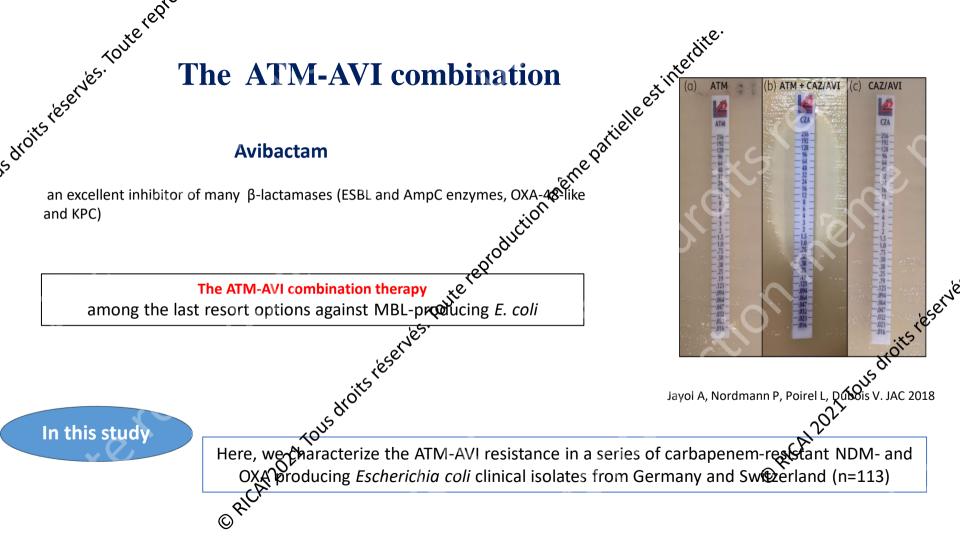
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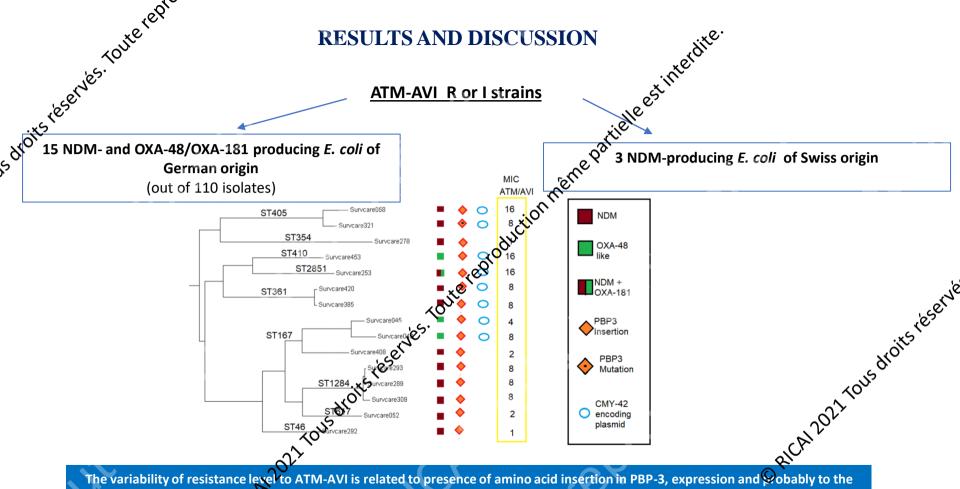
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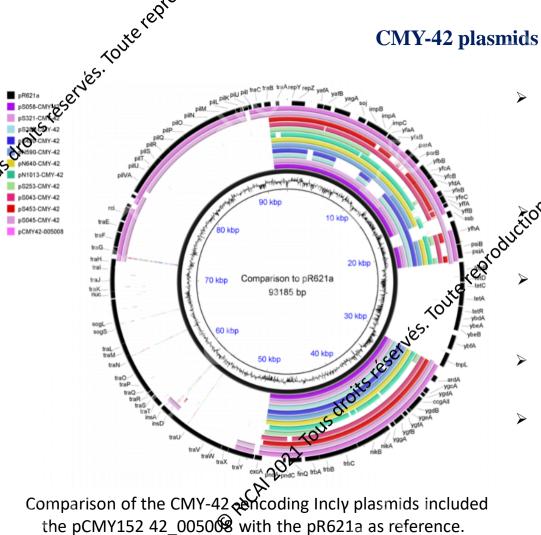






level of expression of CMY-4QMa, a specific plasmid-mediated cephalosporinase that compromises in part, the activity of aztreonam.

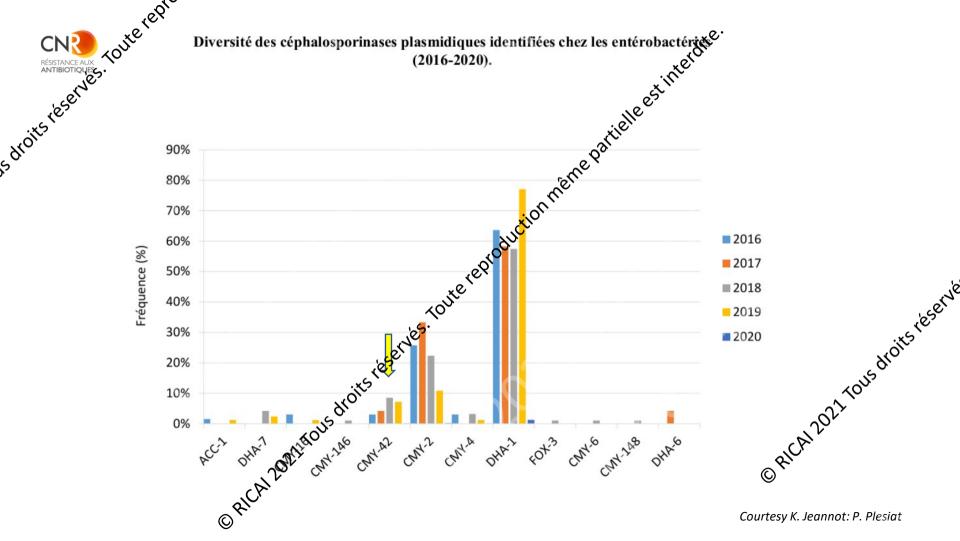
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The CMY-42 allefes are present at an identical location on Inclγ plasmight of 27 – 53 kb.

and the conjugative pilus (pill-V) (Figure) as previously also reported.

- t is unlikely that these truncated plasmids can directly promote transfer of CMY-42 (horizontal transfer)
- Plasmids were highly stable across a 7-day serial passage in the absence of antibiotics.
- Nevertheless, they all carry the Type I partition machinery 3 (ParAB) and a toxin-antitoxin PndBA post-segregational killing system, that can ensure stable presence and inheritance (vertical transfer) (Seen without antibiotic selection.



, Jun Stesenves. Toute rept Fourier Elevated MICs of ATM-AVI resulted from combination of differentic resistance mechanisms, including modification of PBP3 protein sequence through specific amino acid insertions, and production of a CMY-type enzyme, CMY-42.

- Emergent aztreonam-avibactam resistance may be identified with MBL producing strains but also with
- Their occurrence in carbapenem-resistant *E. coli*, that are associated with community spread, could light use of ATM-AVI combination as a first line therapy O_{C} RCA 2021