

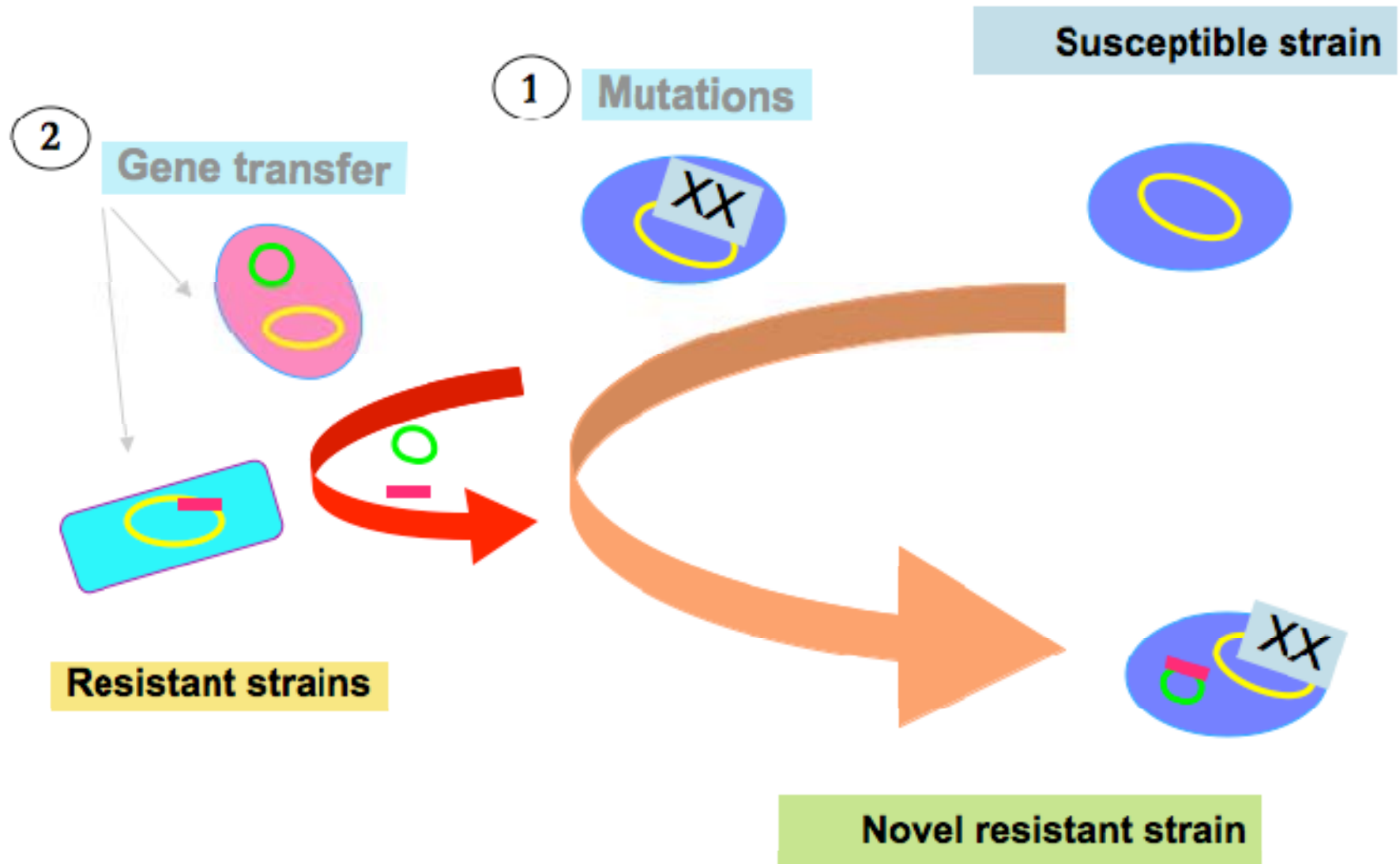


Génétique; résistances croisées, résistances associées chez les bacilles à Gram négatif

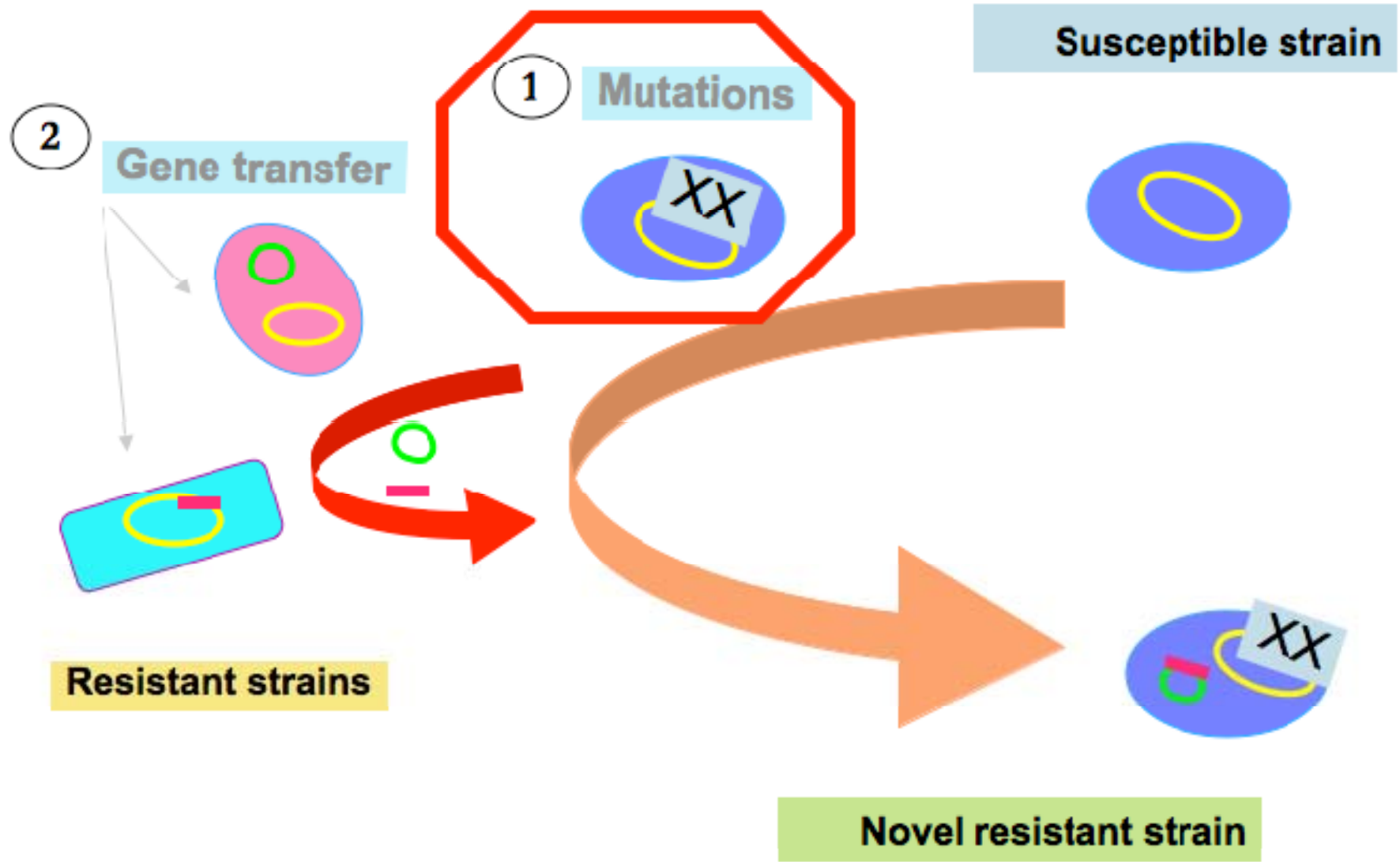
Pr. P. Nordmann
CHU de Bicêtre



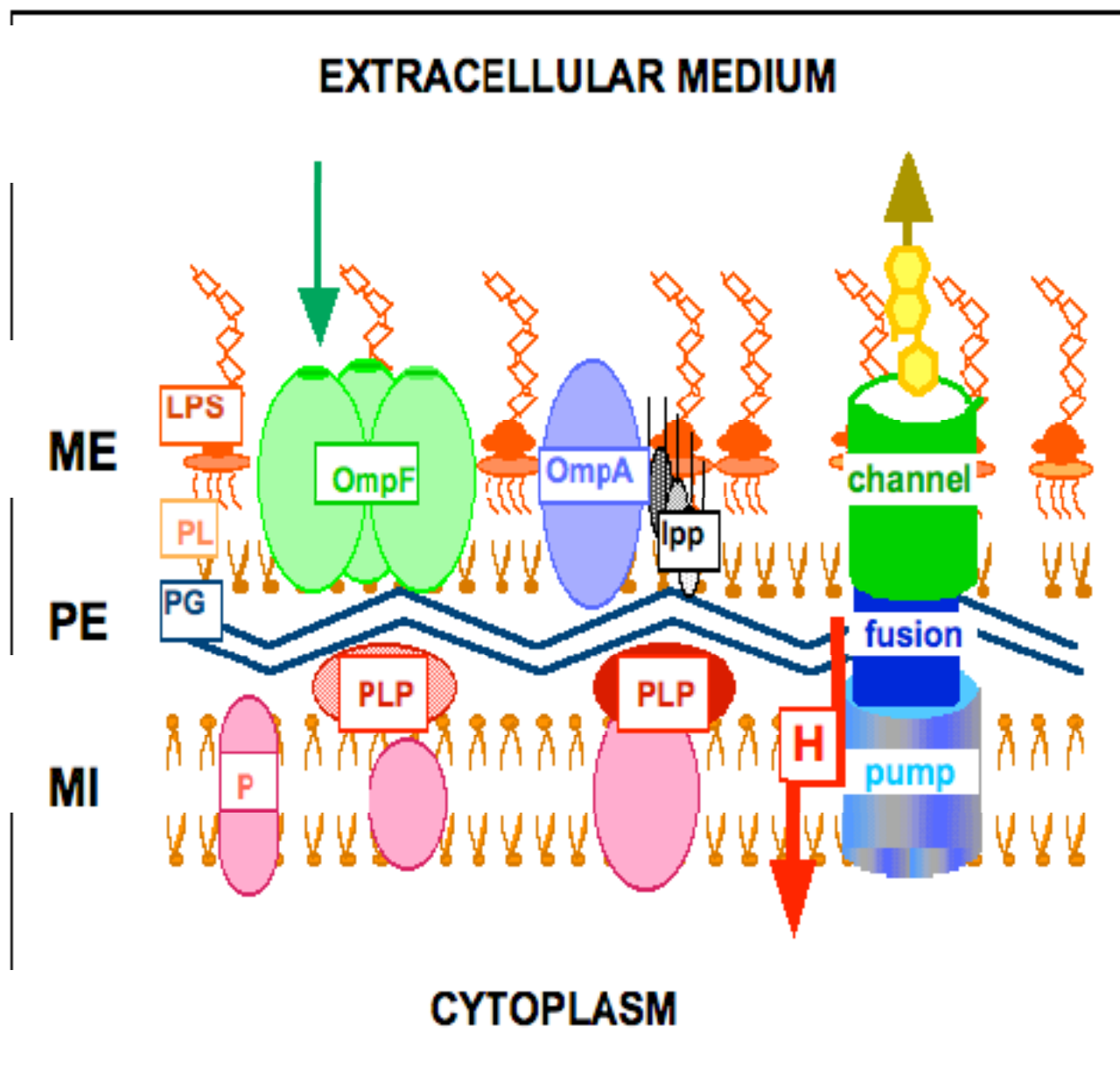
Genetics of Resistance

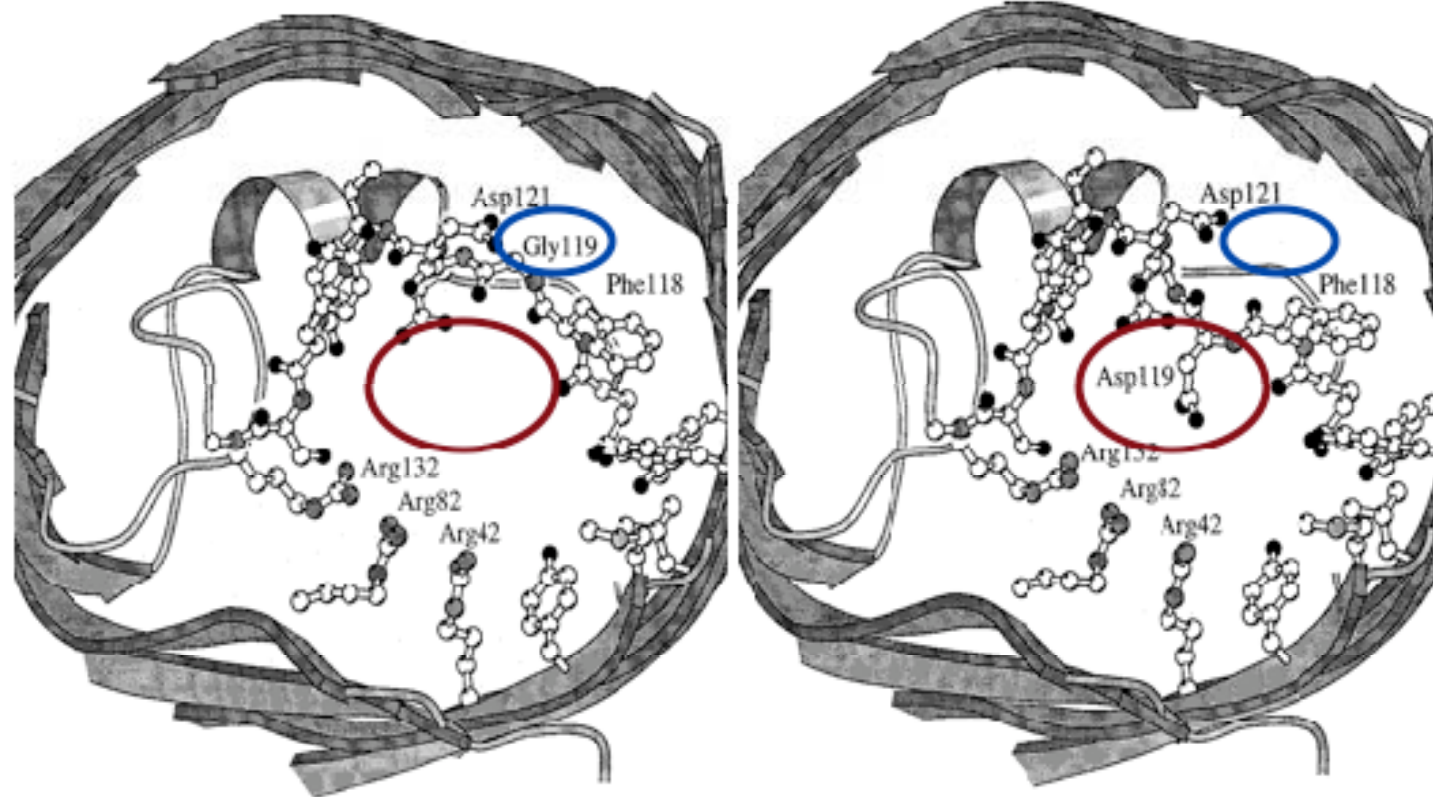


Cross Resistance



Mutations as a source of cross-resistance





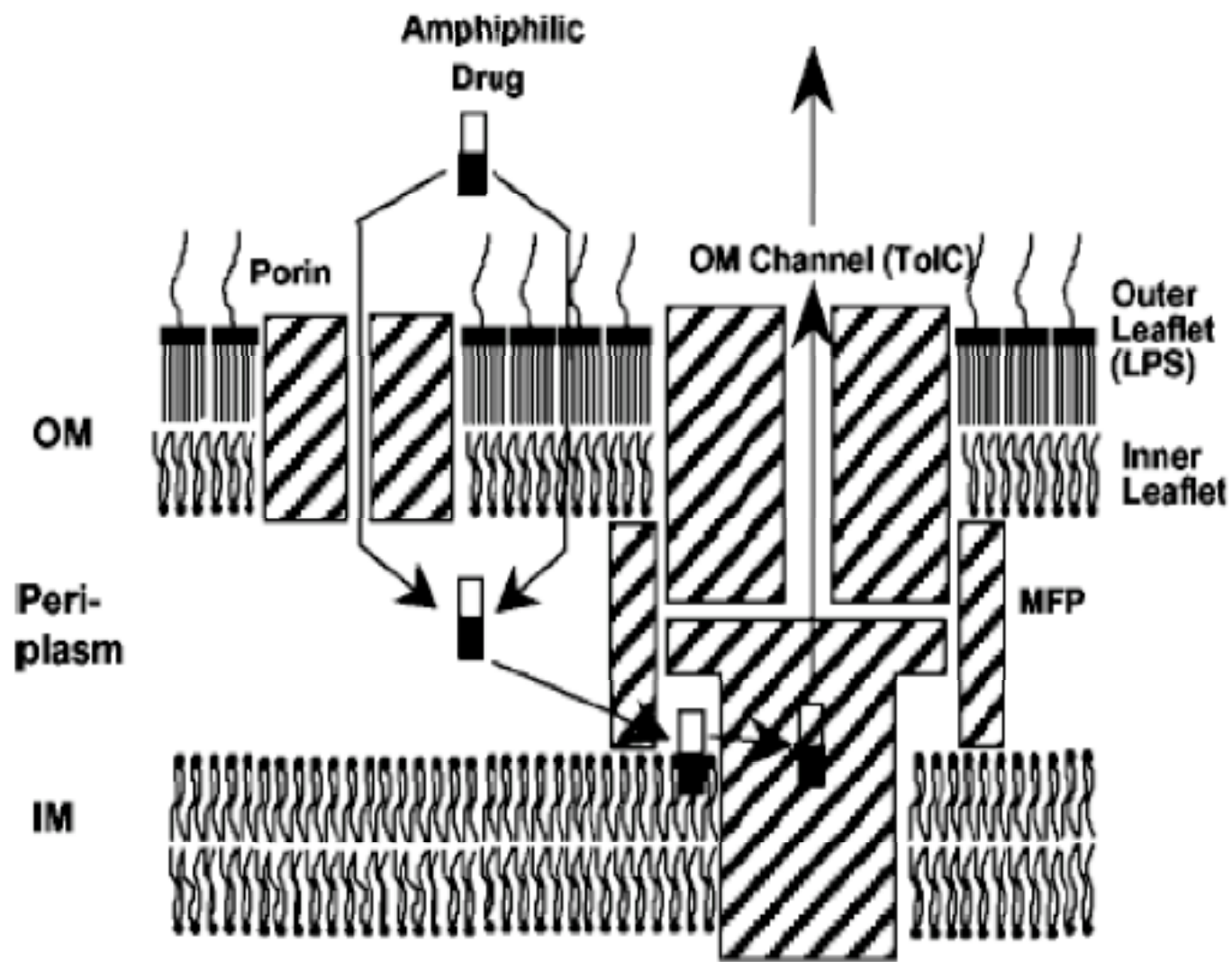
Wild-type porin

Mutated porin

J Infect Dis. 1985 Mar; 151(3):501-7.

Cross-resistance to nalidixic acid, trimethoprim, and chloramphenicol associated with alterations in outer membrane proteins of *Klebsiella*, *Enterobacter*, and *Serratia*.

Gutmann L, Williamson R, Moreau N, Kitzis MD, Collatz E, Acar JF, Goldstein FW



Efflux pump

RND efflux systems in *P. aeruginosa*

System	Operon	Antibiotics as substrates
MexAB-OprM	<i>mexAB-oprM</i>	FQ, β -lactam, Tmp, Cmp, Tet, Nov, Ery...
MexXY (OprM)	<i>mexXY</i>	FQ, AG, Fep, Cpo, Cbo, Tet, Ery...
MexCD-OprJ	<i>mexCD-oprJ</i>	FQ, Cpo, Fep, Tmp, Cmp, Tet, Ery...
MexEF-OprN	<i>mexEF-oprN</i>	FQ, Tmp, Cmp, Tet *
MexHI-OpmD	<i>mexGHI-opmD</i>	FQ, Tet
MexJK (OprM)	<i>mexJK</i>	Tet, Ery
MexMN (OprM)	<i>mexMN</i>	Cmp
MexPQ-OpmE	<i>mexPQ</i>	FQ, Ery
MexVW (OprM)	<i>mexVW</i>	FQ, Cmp, Tet, Ery
CzcBA-OpmN	<i>opmN-czcBA</i>	? *
TriABC-OpmH	<i>triABC</i>	?
YegMNO-OpmB	<i>yegMNO-opmB</i>	?

FQ: (fluoro)quinolones; β -lactam (except imipenem); Tmp: trimethoprim; Cmp: chloramphenicol; Tet: tetracycline; Nov: novobiocin; Ery: erythromycin; AG: aminoglycosides; Fep: cefepime; Cpo: ceftiofime; Cbo: ceftobiprole; * imipenem resistance due to porin OprD down-regulation.

Courtesy P. Fléziat

Wild-type phenotype



Courtesy P. Plésiat

MexAB-OprM mutant



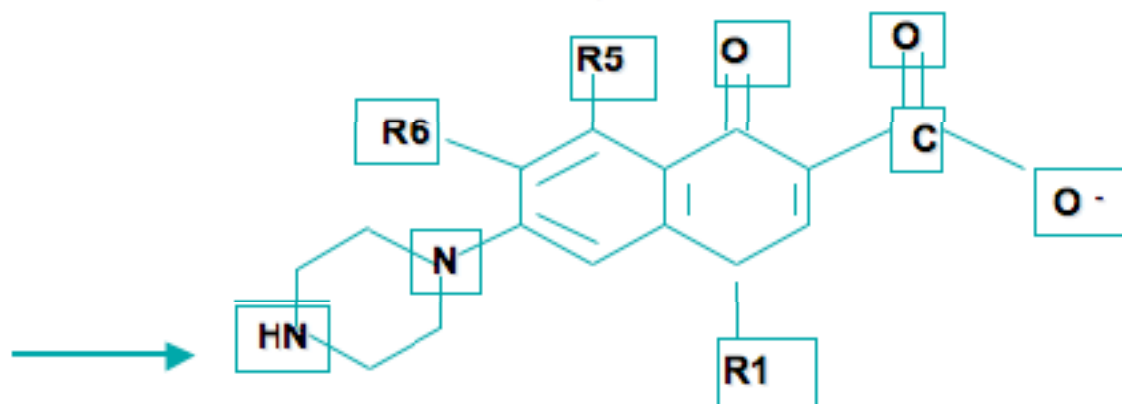
Courtesy P. Plésiat

Enzymatic inactivation of fluoroquinolones:

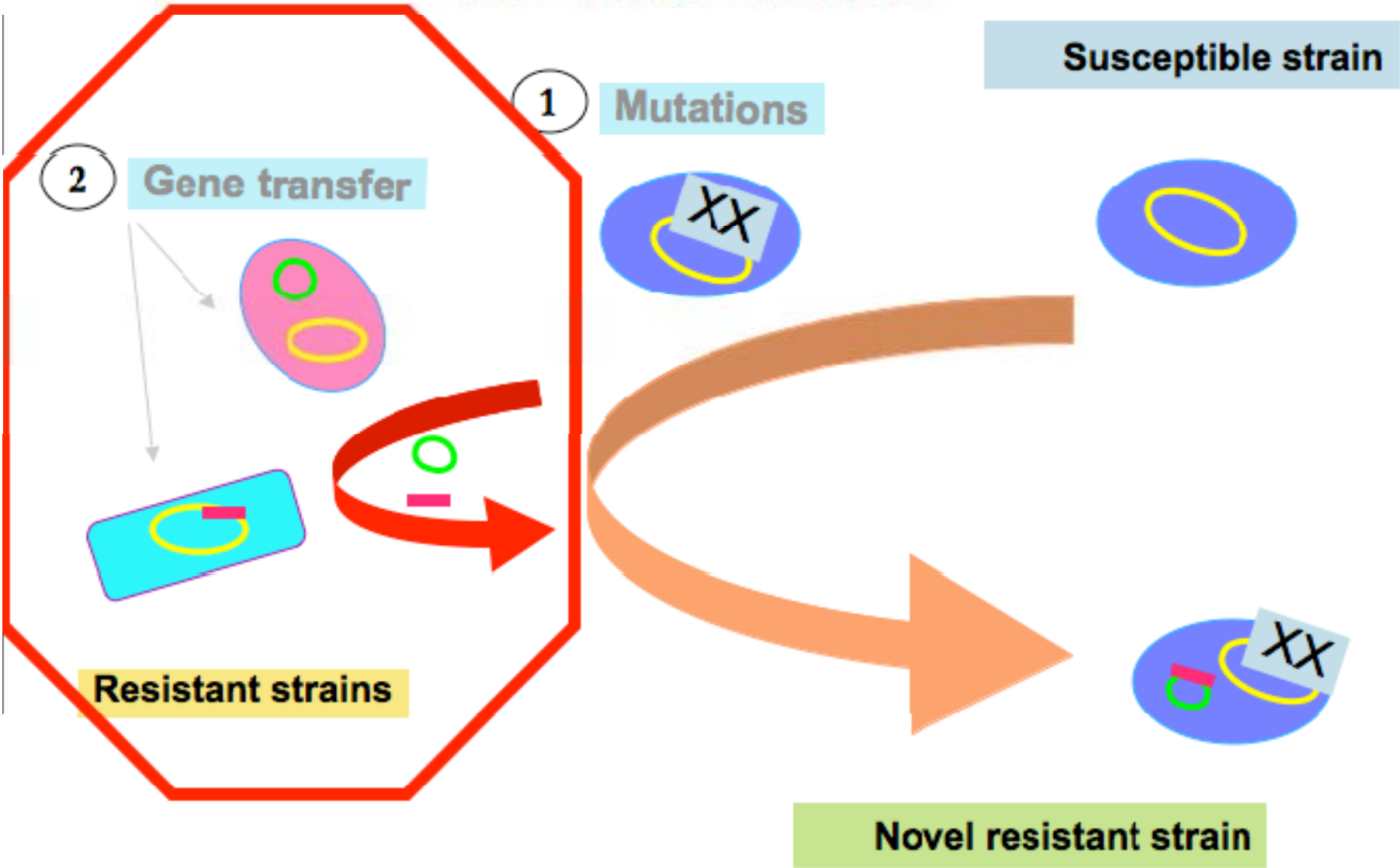
Bifunctional acetyltransferase : **AAC(6')-Ib-cr**

- Two substitutions: Trp102Arg and Asp179Tyr
- *N*-acetylation of fluoroquinolones
- Confers reduced susceptibility to aminoglycosides and fluoroquinolones (MIC of ciprofloxacin increased 3 fold)

(Robicsek *et al.* Nat Med 2006)

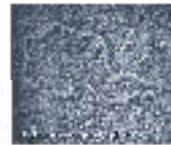


Co-Resistance

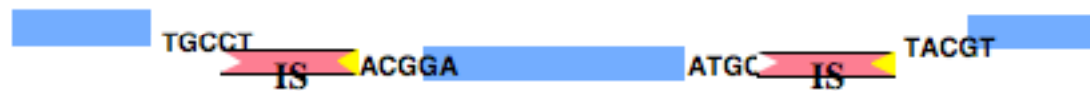


Co-Resistance

- Plasmid



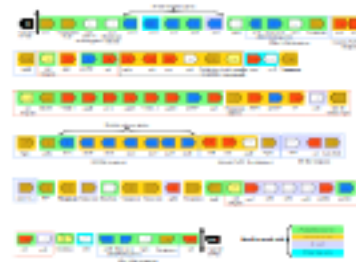
- Transposon



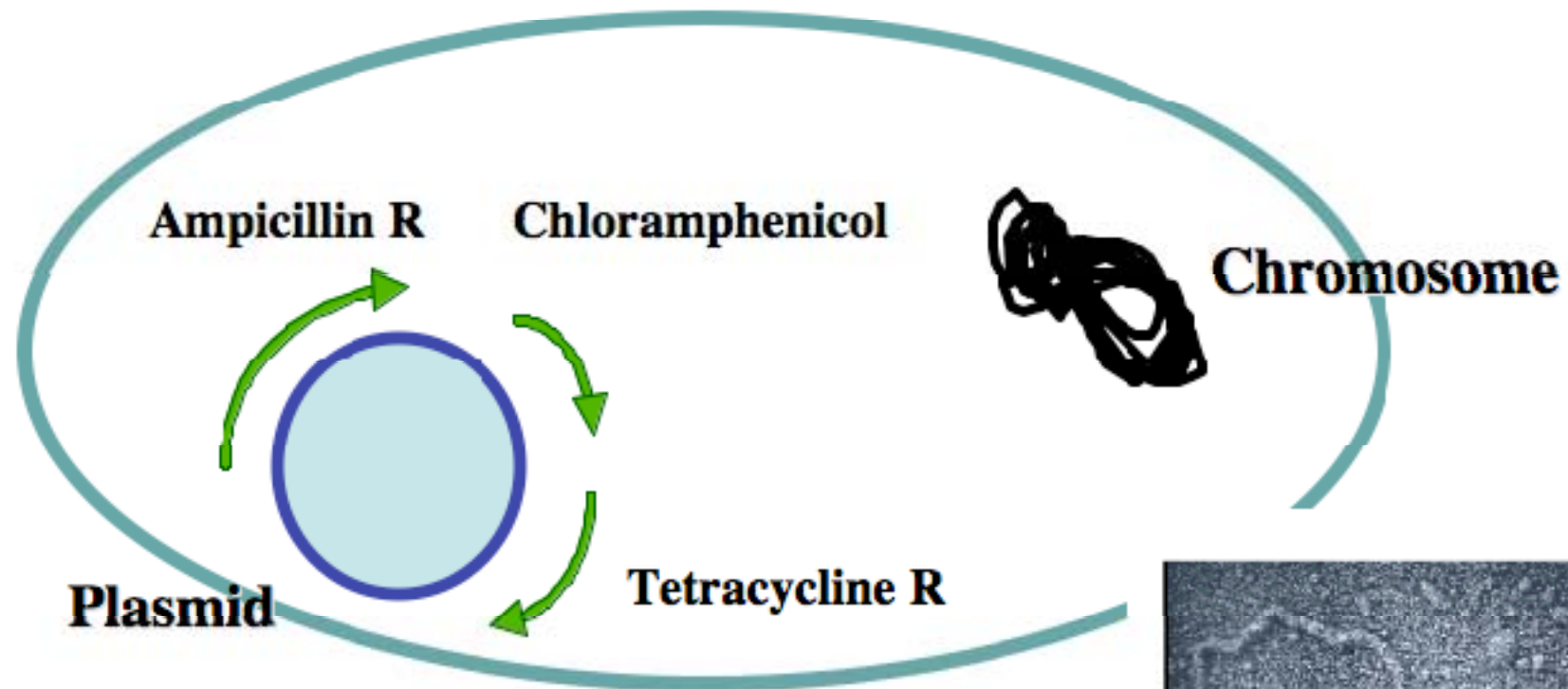
- Integron



- Resistance island

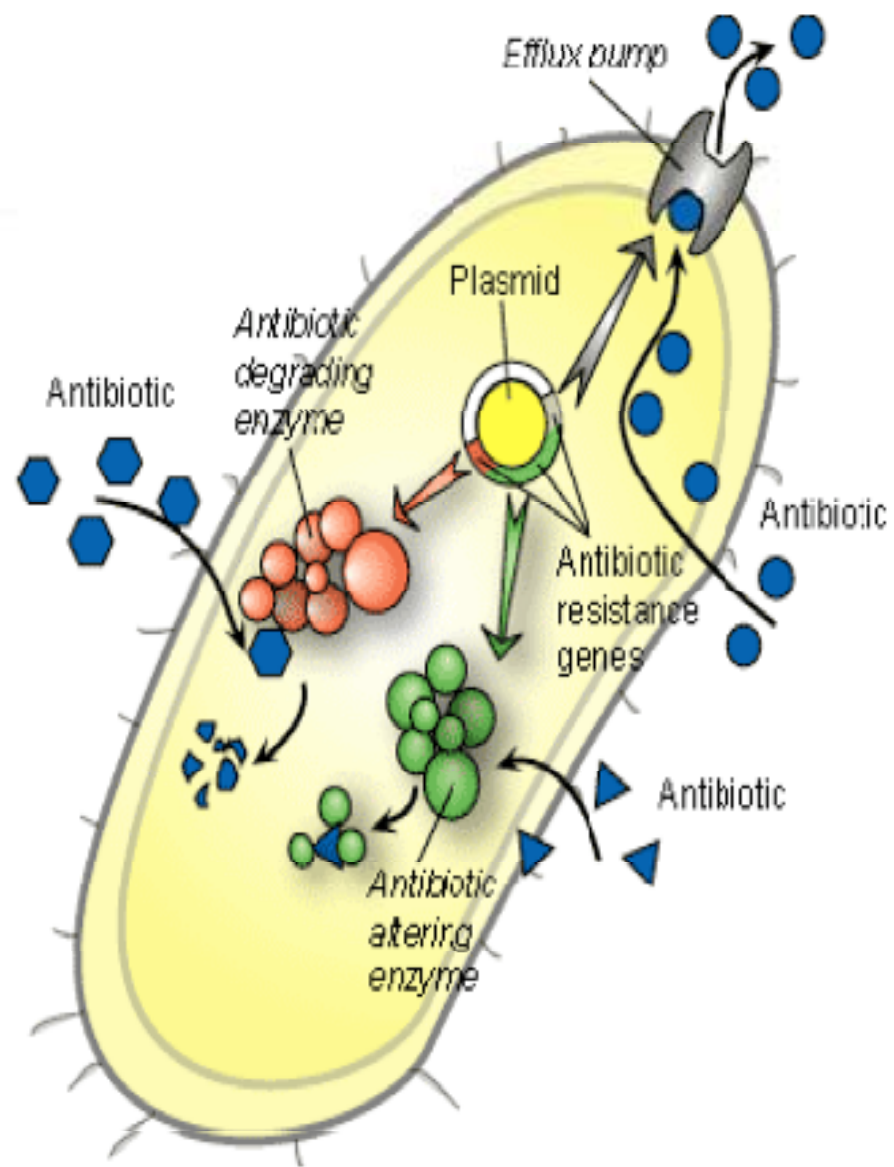


Plasmid



- Interspecies transfer - 5- > 500 kb
- Conjugative, non conjugative mobilisable....

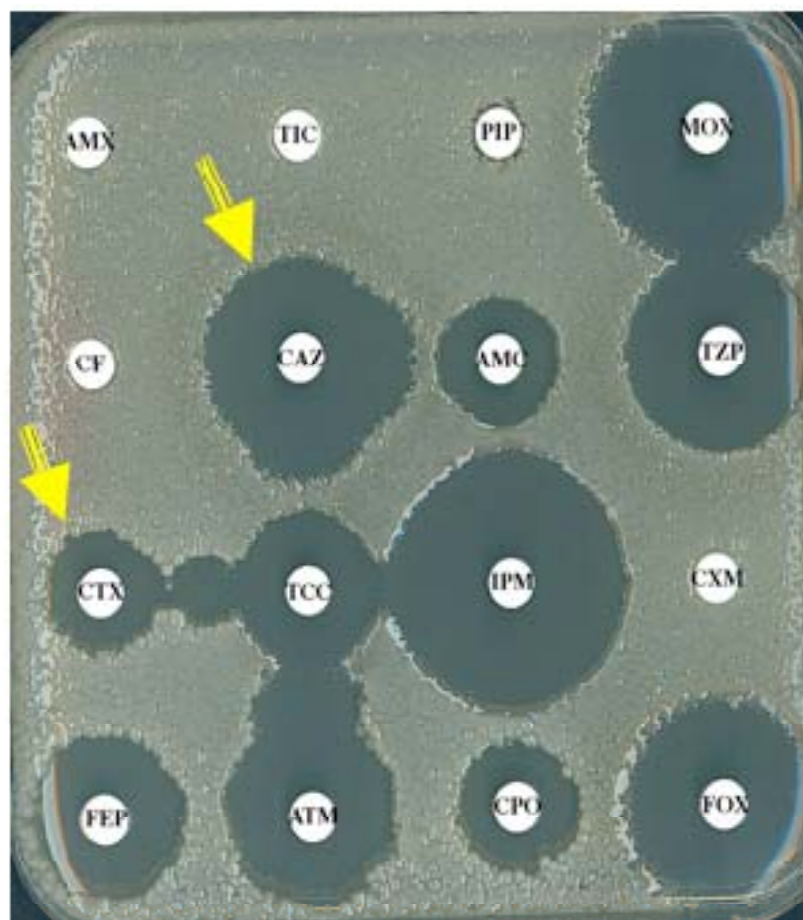




Biochemical mechanisms of antibiotic resistance and their genetic determinants

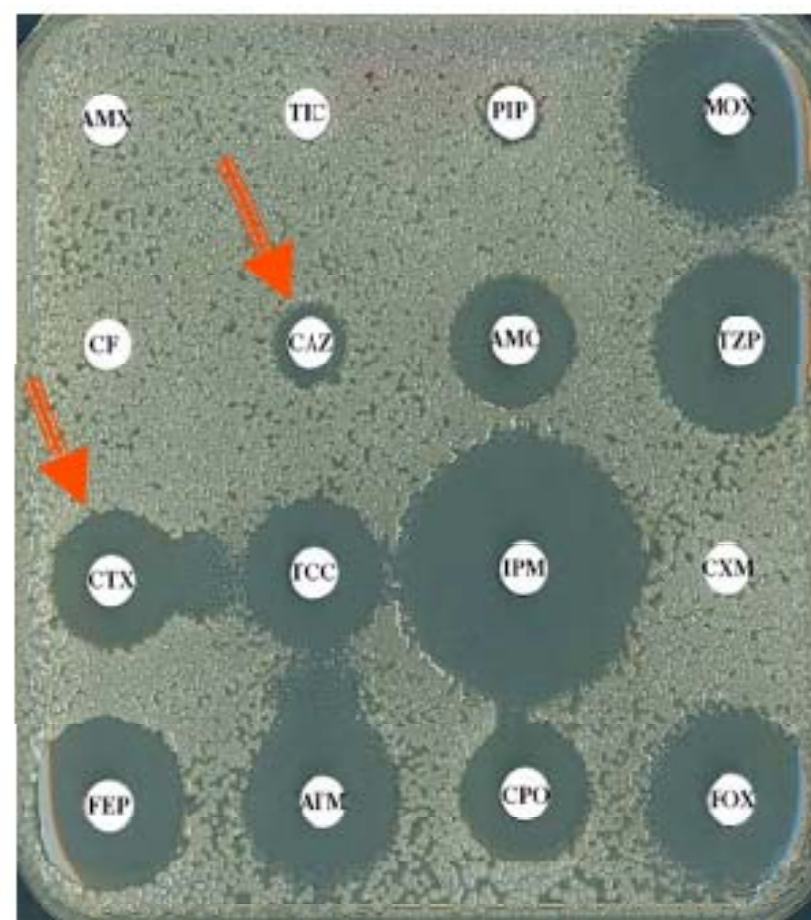
<i>Mechanism</i>	<i>Examples</i>	<i>Genetic determinants</i>	
		<i>Mutation</i>	<i>Gene acquisition</i>
Reduced permeability	Aminoglycosides	+	+
Pro-drug not activated	Isoniazid	+	
Active efflux	Tetracycline		+
	Fluoroquinolones	+	
Alteration of drug target	Erythromycin		+
	Fluoroquinolones	+	+
	Rifampicin	+	+
	Tetracycline		+
Inactivation of drug	Aminoglycosides		+
	Chloramphenicol		+
	β -lactams	+	+
'By-pass' inhibited step	Sulfonamides		+
	Trimethoprim		+
Immunity protein	Bleomycin		+
Amplification of target	Trimethoprim	+	+
	Sulfonamides	+	+
Sequestration of drug	β -lactams	+	+

CTX-Ms: hydrolyse du cefotaxime... et de la ceftazidime



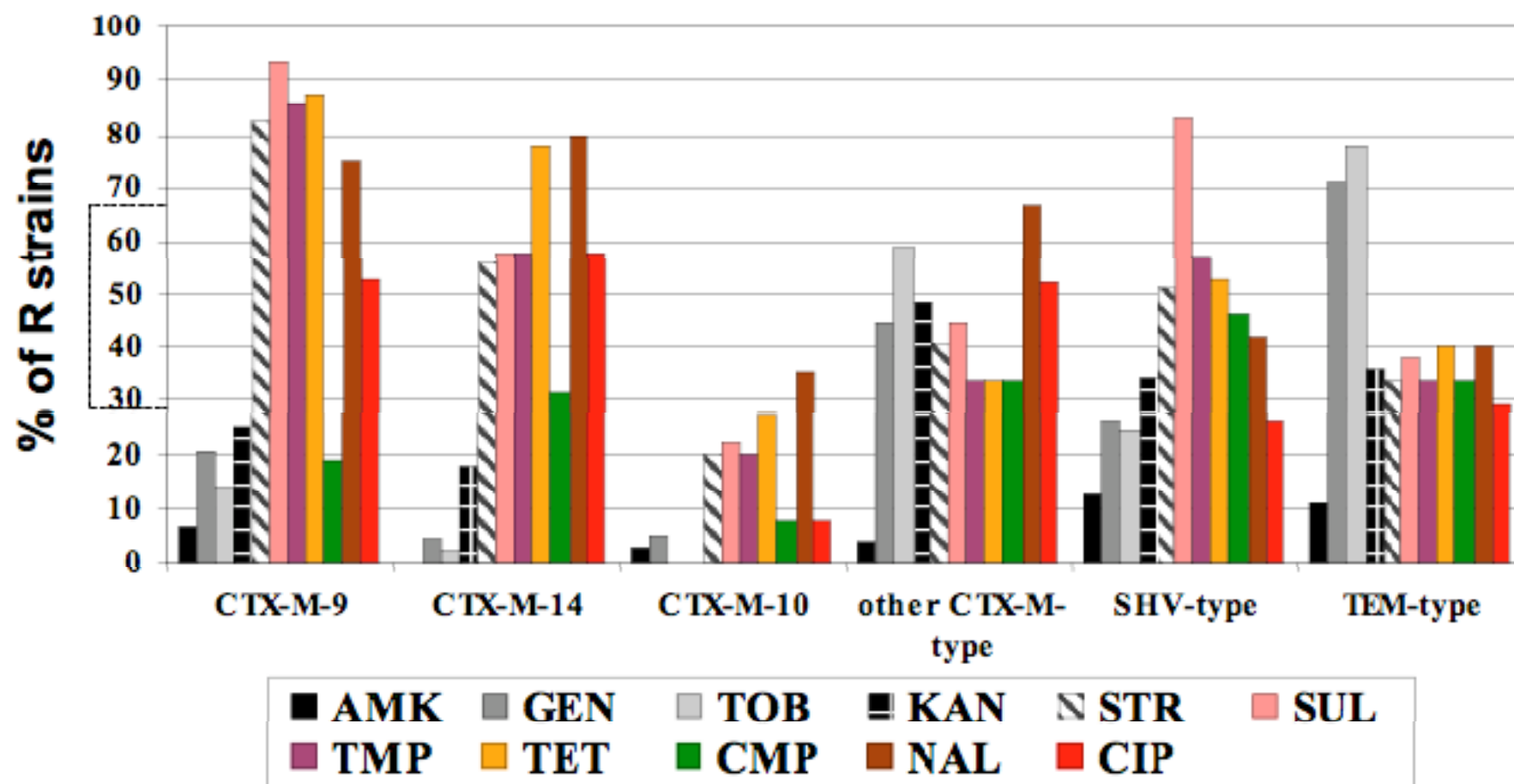
CTX-M-3

E. coli



CTX-M-15

Co-resistance of the CTX-M producers



Multidrug resistance among *Klebsiella* spp. collected in an European survey of *K. pneumoniae* isolates from ICUs

Resistance	ESβL Positive (n=110)	ESβL Negative (n=323)
Ceftazidime	75	25
Gentamicin	72	10
Amikacin	61	4
Ciprofloxacin	31	4

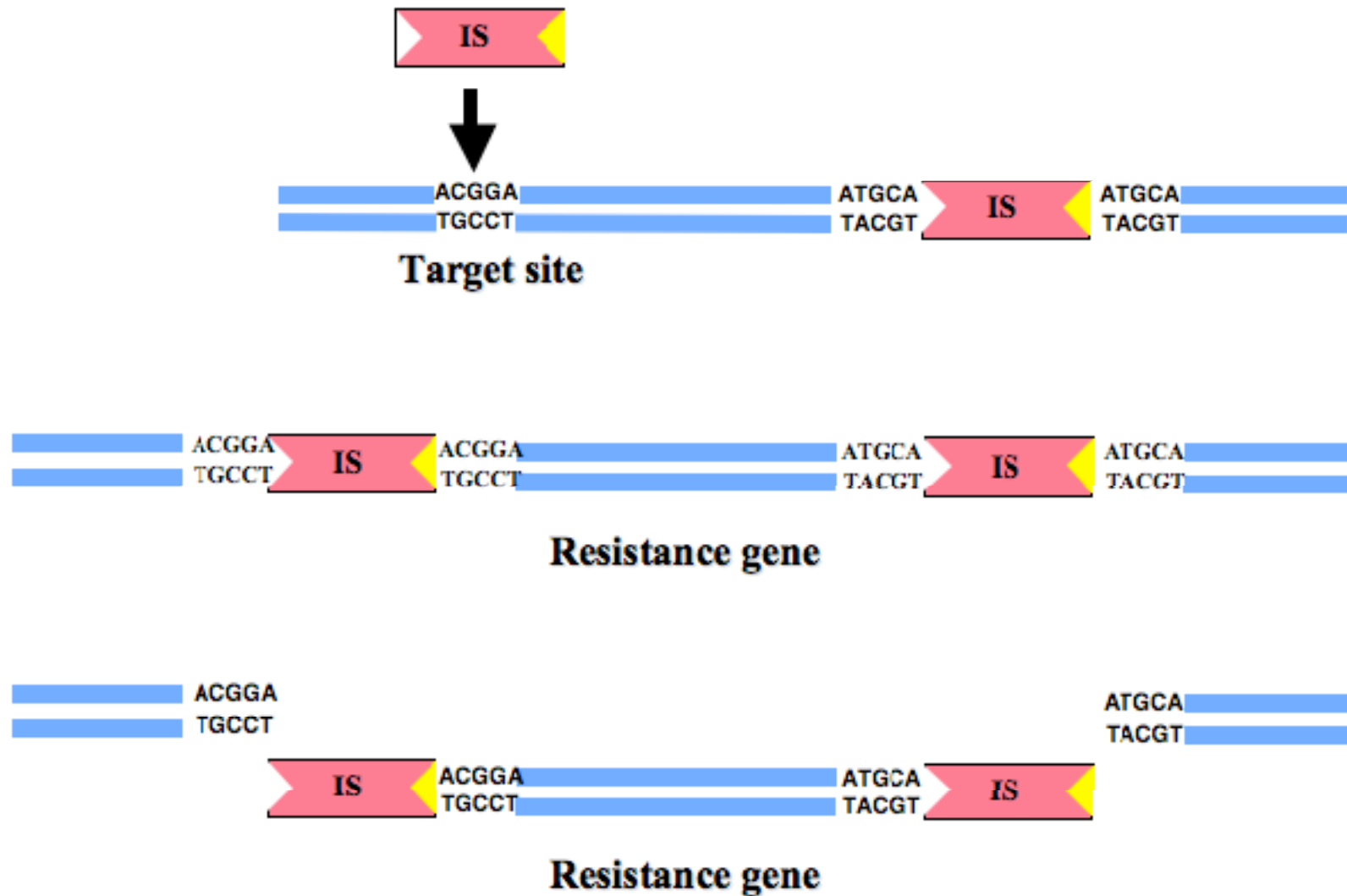
Co-resistances

ESBL - Bicetre - *E.coli* - 2007 (n=71)

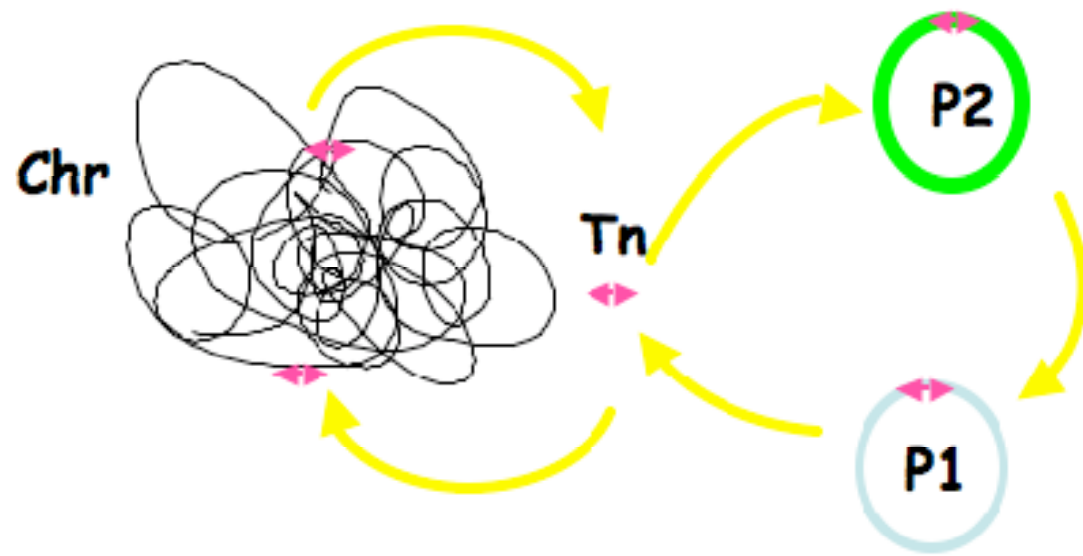
% résistance

Gentamicin	39
Tobramycin	51
Amikacin	27
Netilmicin	47
Tetracycline	58
Trimethoprim/Sulfamethoxazole	63
Nalidixic acid	79
Ciprofloxacin	72
Colistin	0
Fosfomycin	0
Ertapenem/Imipenem	0
Tigecycline	0

Transposons



Transposons

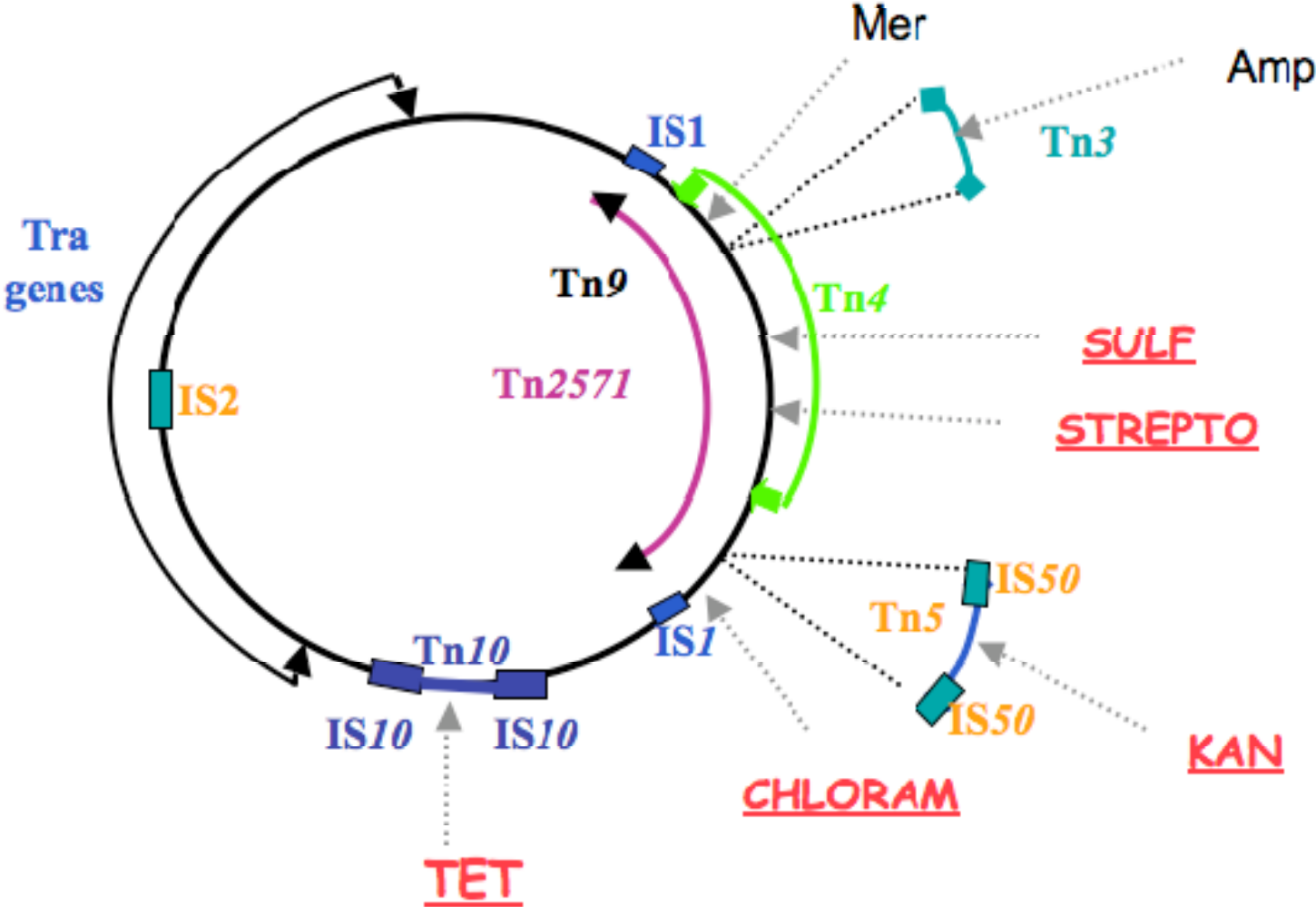


Resistance genes

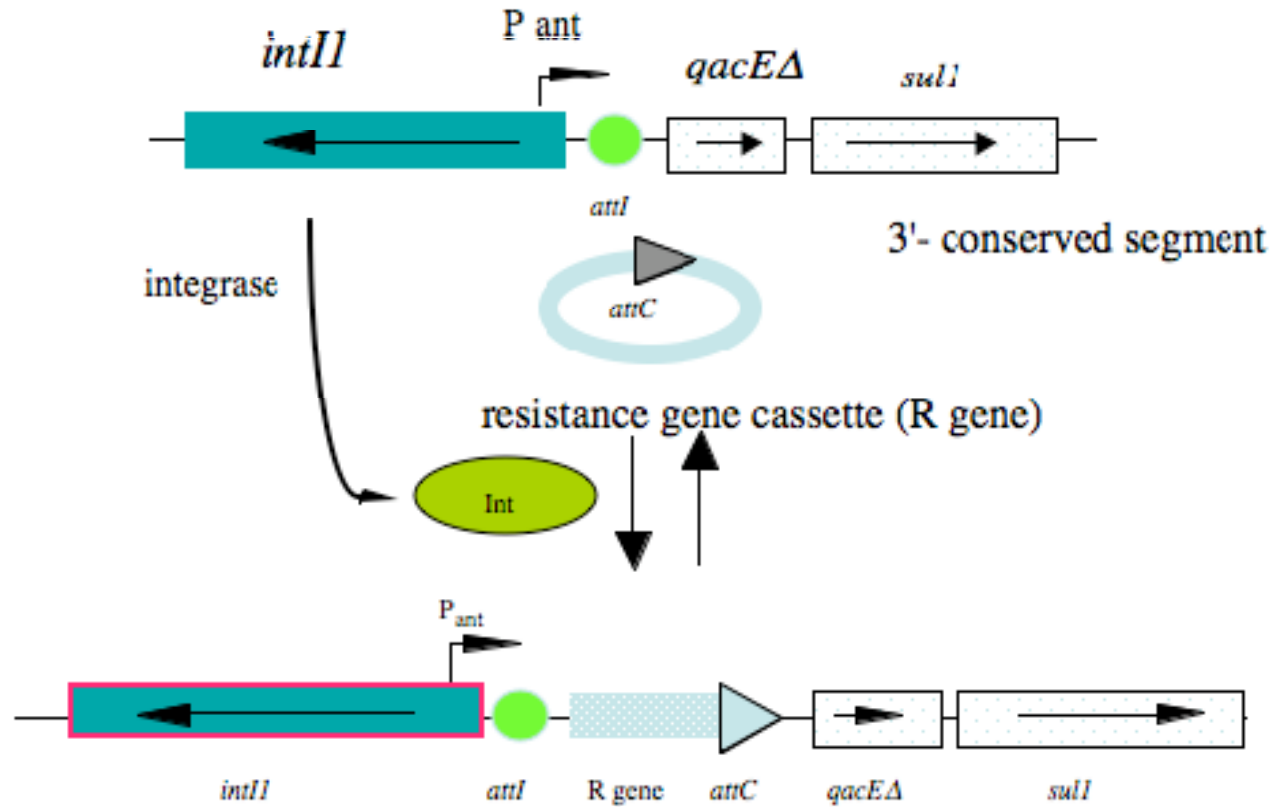
- Aminoglycosides
- β -lactamines
- Sulfamides
- Rifampicine
- Cyclines

- Chloramphenicol
- Trimethoprim

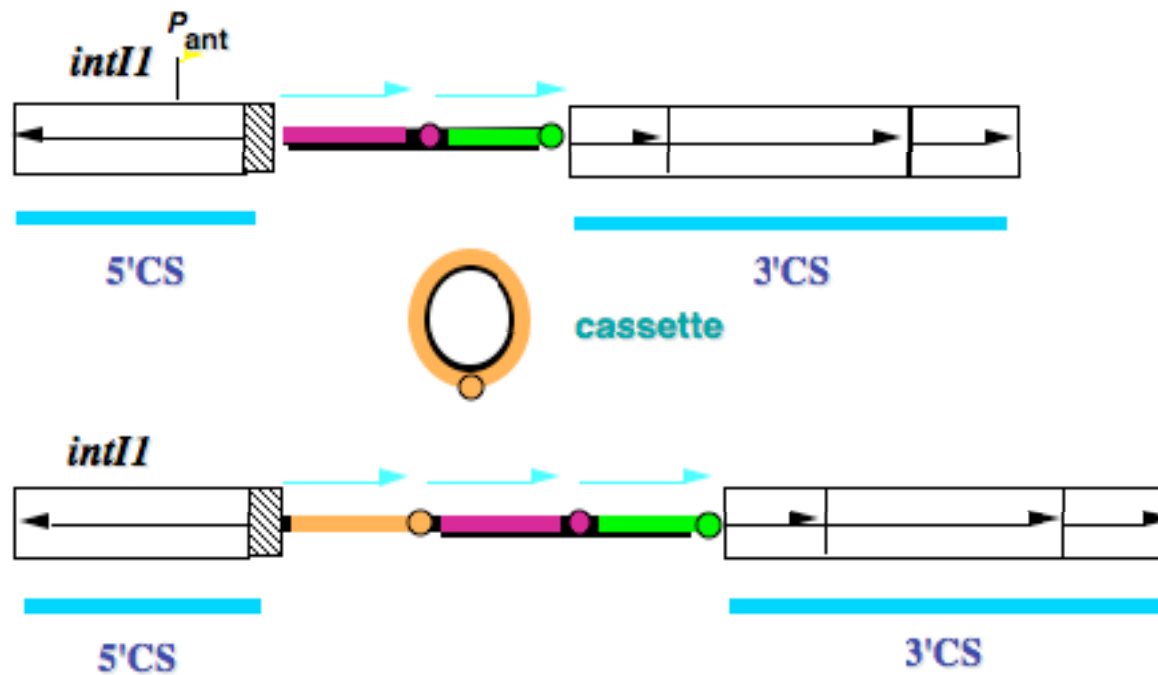
R1 plasmid (100-kbp): an example of resistance accumulation mediated by transposable elements, and especially composite transposons



Integrations

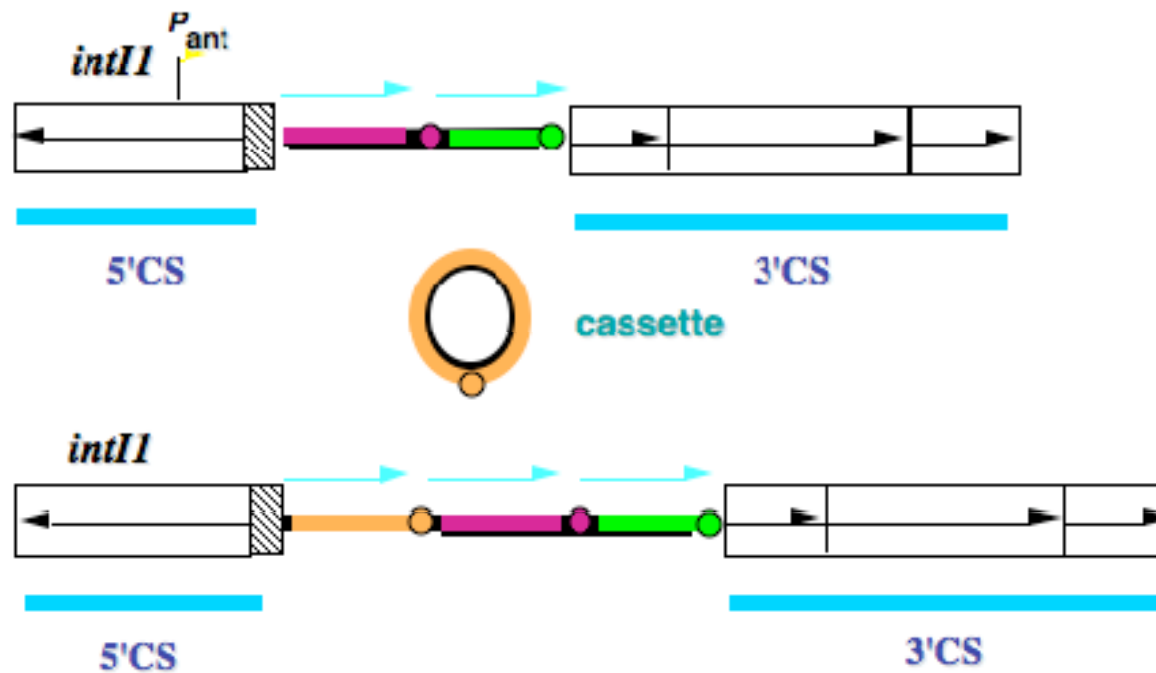


Integrations



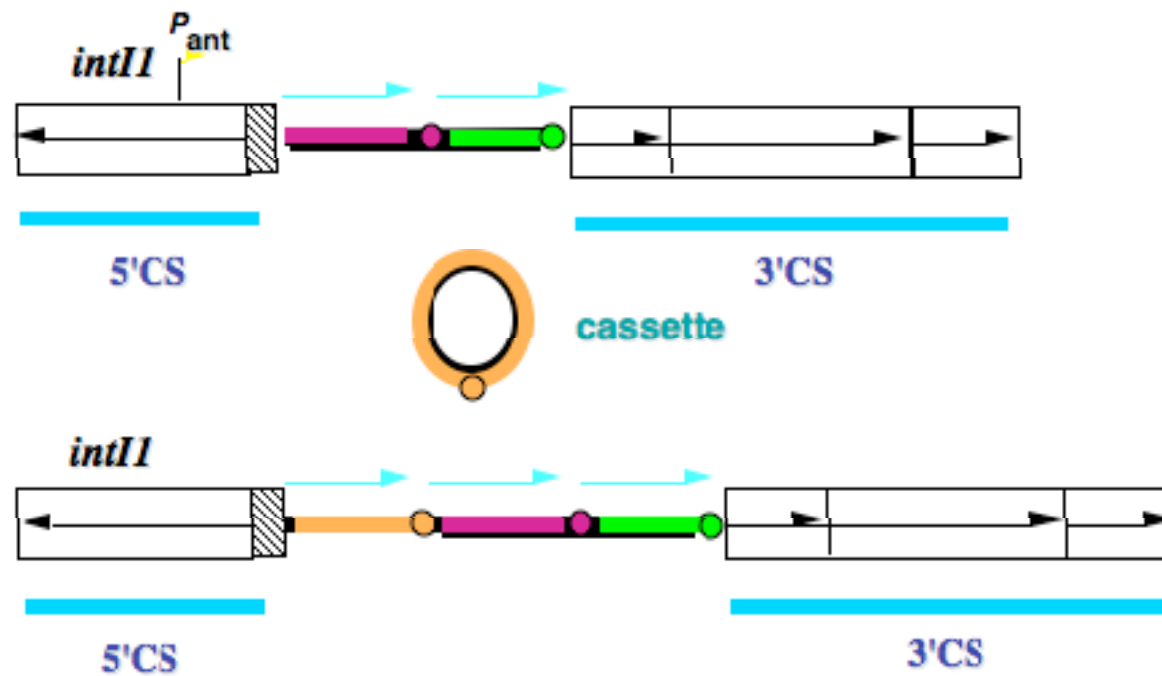
- **Consequences:** co-resistance

Integrations



- **Consequences** : **co-resistance** ; **co-expression**

Integrans



- **Consequences** : co-resistance ; co-expression ; co-selection

*bla*_{VEB-1} as a gene cassette identified
in In53 in *E. coli*

In53



Integrans

- Extensively found in Gram clinical isolates
- Over 80 different antibiotic-resistance-gene cassettes :

β-Lactam resistance		Aminoglycoside resistance	
<i>bla_{CCR-1}</i>	Extended-spectrum β-lactamase	<i>aadA4</i>	Adenyltransferase
<i>bla_{GES-1}</i>	Extended-spectrum β-lactamase	<i>aadA5</i>	Adenyltransferase
<i>bla_{GES-2}</i>	Extended-spectrum β-lactamase	<i>aadA6</i>	Adenyltransferase
<i>bla_{TBC-1}</i>	Extended-spectrum β-lactamase	<i>aadA8</i>	Adenyltransferase
<i>bla_{TBC-2}</i>	Extended-spectrum β-lactamase	<i>aadA10</i>	Adenyltransferase
<i>bla_{IMP-2}</i>	Carbapenemase	<i>aac(3)-Ib/aac(6')-Ib</i>	Acetyltransferase
<i>bla_{IMP-3}</i>	Carbapenemase	<i>aac29a</i>	Acetyltransferase
<i>bla_{IMP-4}</i>	Carbapenemase	<i>aac29b</i>	Acetyltransferase
<i>bla_{IMP-6}</i>	Carbapenemase	<i>aacA1b/orfG</i>	Acetyltransferase
<i>bla_{IMP-7}</i>	Carbapenemase	<i>ant(3'')-II-aac(6')-IIIa</i>	Adenyltransferase/ acetyltransferase
<i>bla_{VBS-1}</i>	Extended-spectrum β-lactamase	<i>aphA15</i>	Phosphotransferase
<i>bla_{VM-1}</i>	Carbapenemase	Chloramphenicol resistance	
<i>bla_{VM-2}</i>	Carbapenemase	<i>catA4</i>	
<i>bla_{VM-3}</i>	Carbapenemase	<i>catA5</i>	Efflux pump
<i>aca-10/aadA1</i>	Extended-spectrum β-lactamase/adenyltransferase	Rifampicin resistance	
<i>aca-11</i>	Extended-spectrum β-lactamase	<i>arr-2</i>	ADP-ribosylation
<i>aca-13</i>	Extended-spectrum β-lactamase	Trimethoprim resistance	
<i>aca-15</i>	Extended-spectrum β-lactamase	<i>dhfr13</i>	Dihydrofolate reductase
<i>aca-16</i>	Extended-spectrum β-lactamase	<i>dhfr17</i>	Dihydrofolate reductase
<i>aca-17</i>	Extended-spectrum β-lactamase	<i>dhfrA17</i>	Dihydrofolate reductase
<i>aca-18</i>	Extended-spectrum β-lactamase	Quaternary ammonium compound resistance	
<i>aca-19</i>	Extended-spectrum β-lactamase	<i>qacF'</i>	Efflux pump
<i>aca-28</i>	Extended-spectrum β-lactamase	<i>qacI</i>	Efflux pump
<i>aca-31</i>	Extended-spectrum β-lactamase		
<i>aca-32</i>	Extended-spectrum β-lactamase		

adapted from
Fluit &
Schmitz,
CMI 2004

Increased Prevalence of Class I Integrons in *Escherichia coli*, *Klebsiella* Species, and *Enterobacter* Species Isolates over a 7-Year Period in a German University Hospital

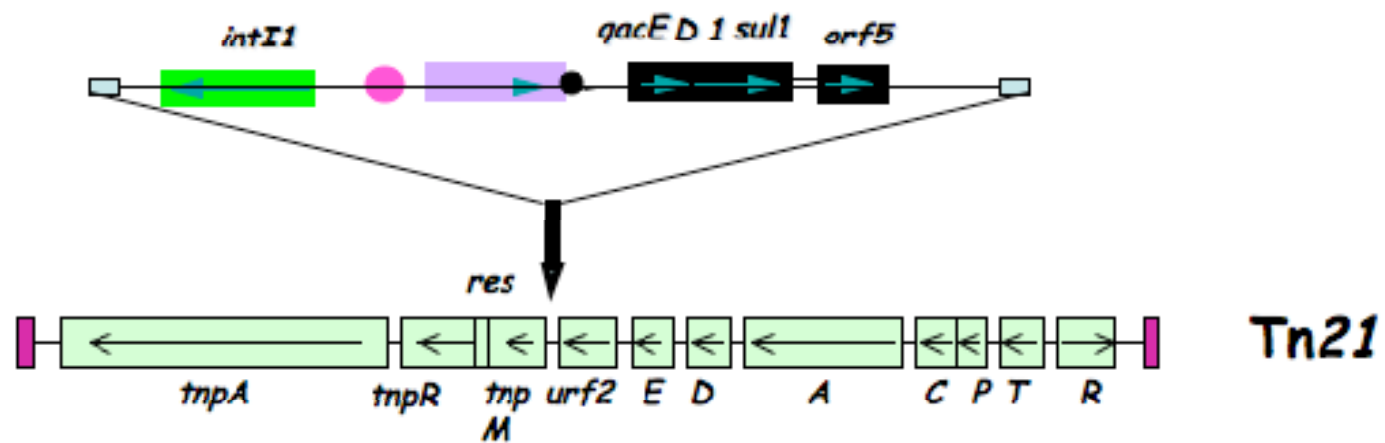
FRANZ-JOSEF SCHMITZ,^{1,2*} DIETER HAFNER,³ ROLAND GEISEL,¹ PATRICK FOLLMANN,¹
CHRISTIAN KIRSCHKE,¹ JAN VERHOEF,² KARL KÖHRER,¹ AND AD C. FLUIT²

*Institute for Medical Microbiology and Virology*¹ and *Institute of Pharmacology and Clinical Pharmacology*,³
Heinrich-Heine University Düsseldorf, Düsseldorf, Germany, and Eijkman-Winkler Institute for
*Medical Microbiology, University Medical Center Utrecht, Utrecht, The Netherlands*²

Received 28 December 2000/Returned for modification 31 May 2001/Accepted 18 July 2001

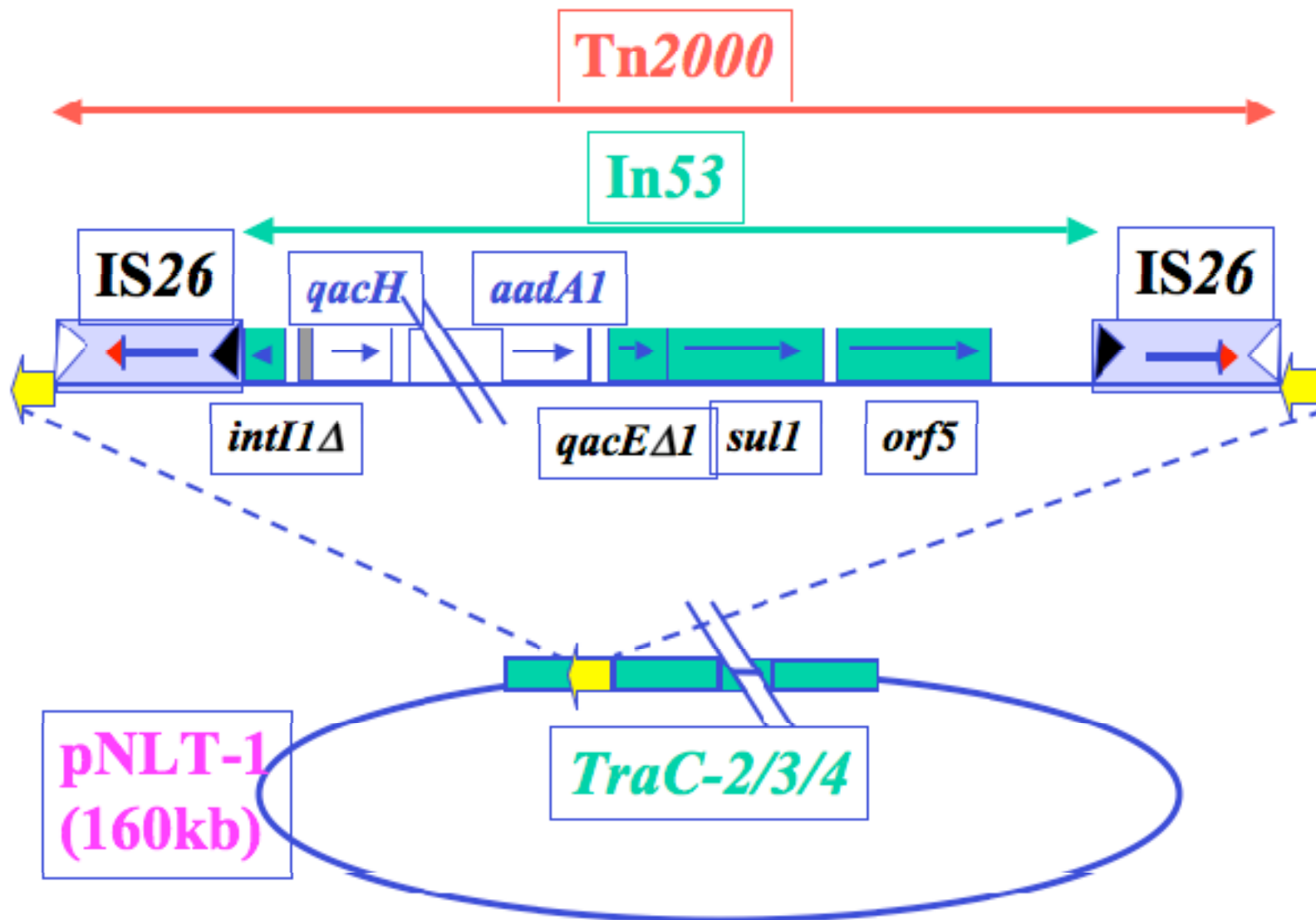
The prevalence of integrons in five enterobacterial species was analyzed in 900 blood culture isolates from 1993, 1996, and 1999. Remarkably, the prevalence increased from 4.7% in 1993 to 9.7% in 1996 and finally to 17.4% in 1999 ($P < 0.01$). Within 7 years the combined percentage of P1 strong promoters and P1 weak plus P2 active promoters with high transcription efficacies has increased from 23.1 to 33.3 and finally 60% ($P < 0.05$).

Integrations are carried by transposons

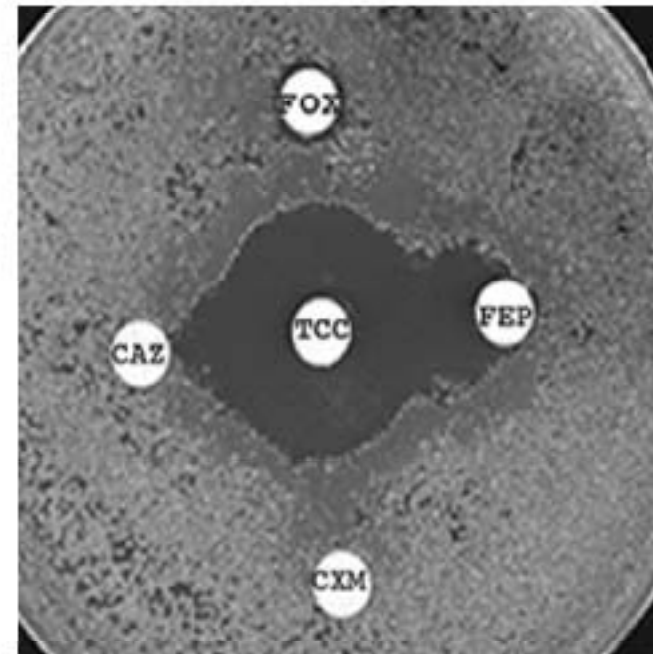
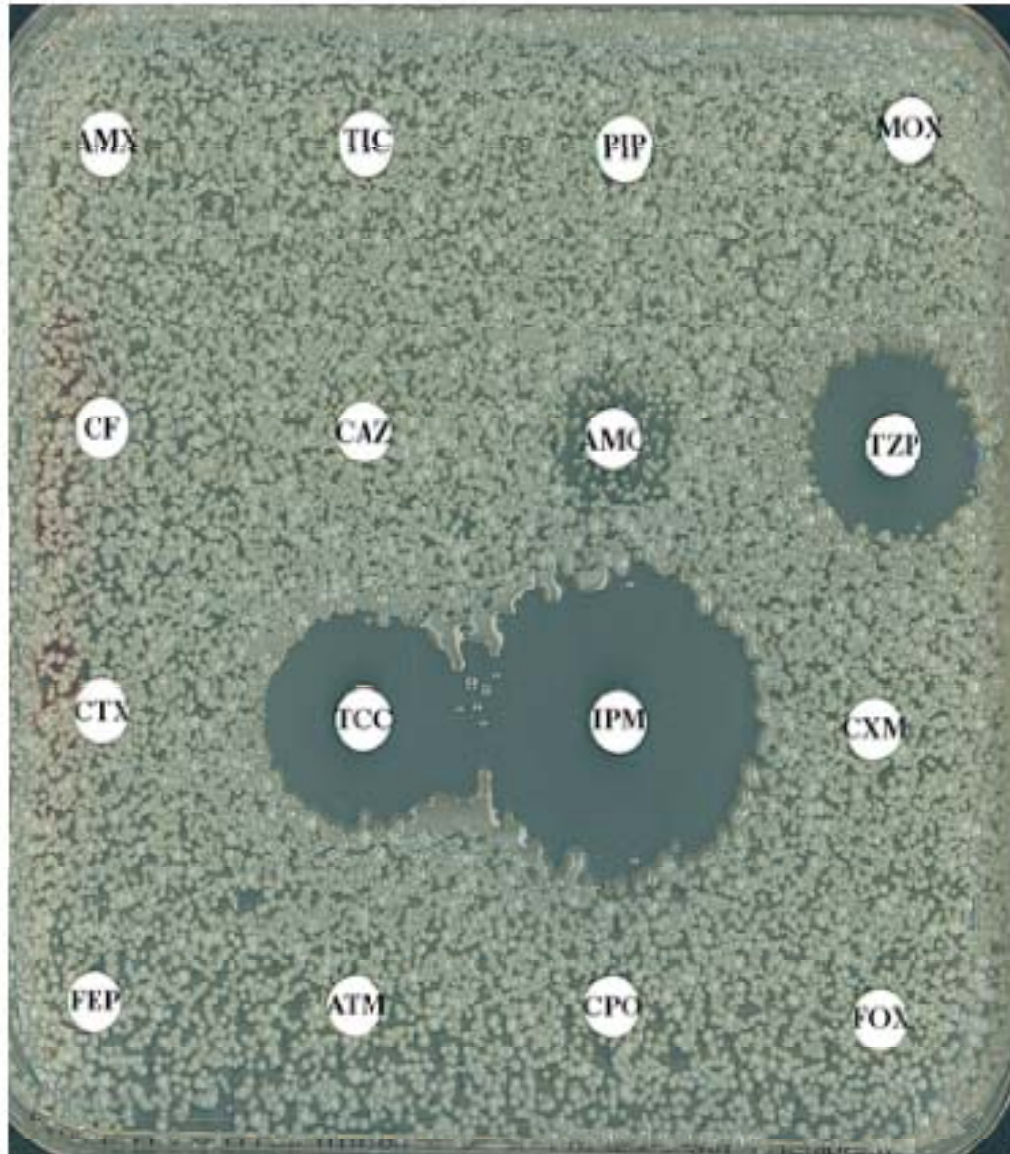


The integron is not self-mobile, it is mobilized by the transposon

Integrans, transposons, plasmids

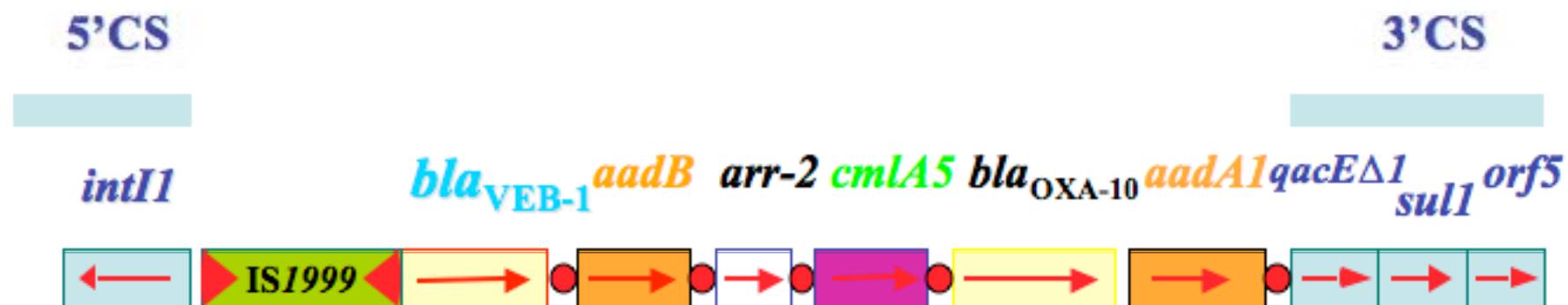


***Acinetobacter baumannii*: VEB-1**

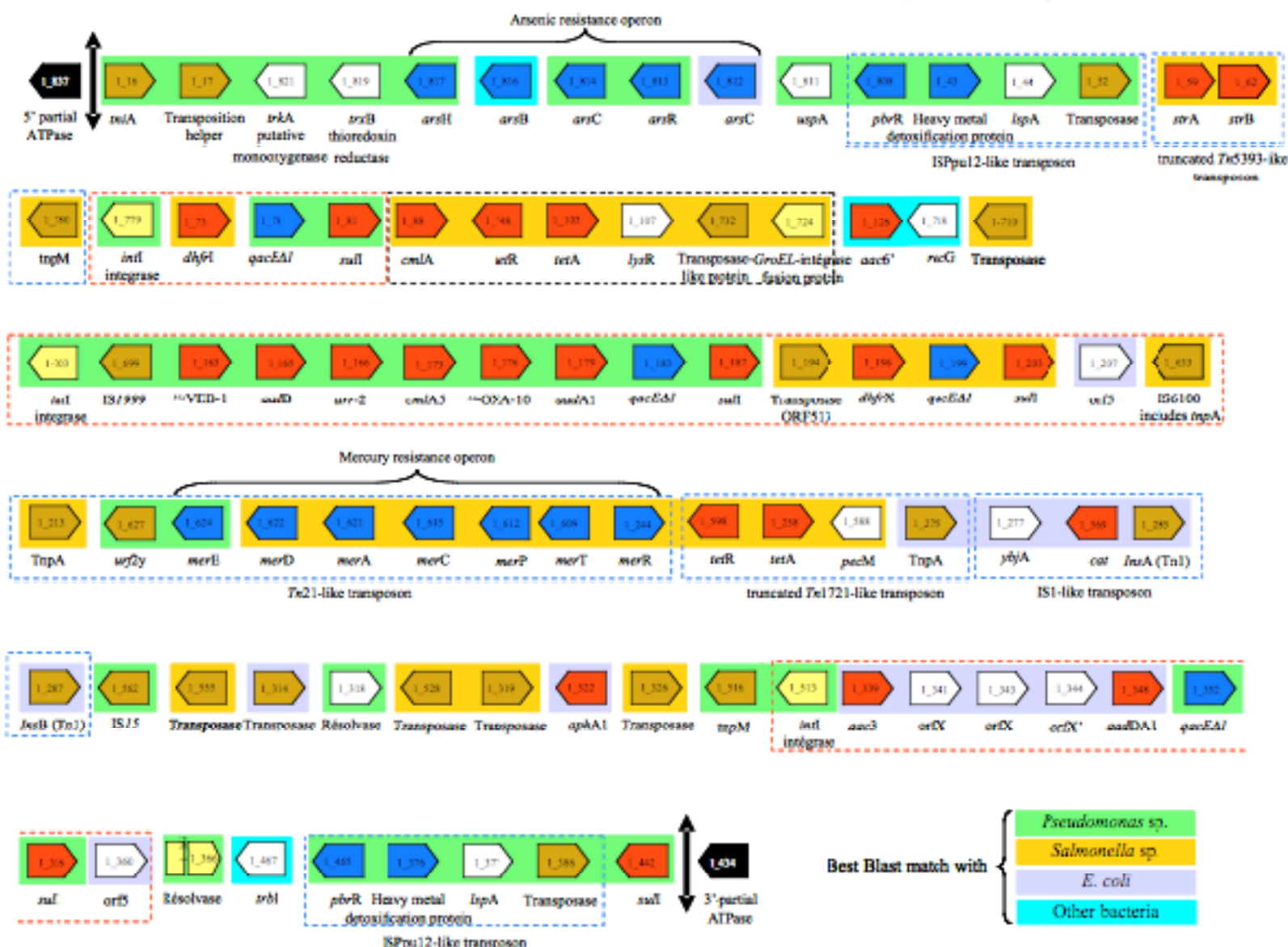


Poirel, Nordmann J. Clin. Microbiol. 2003

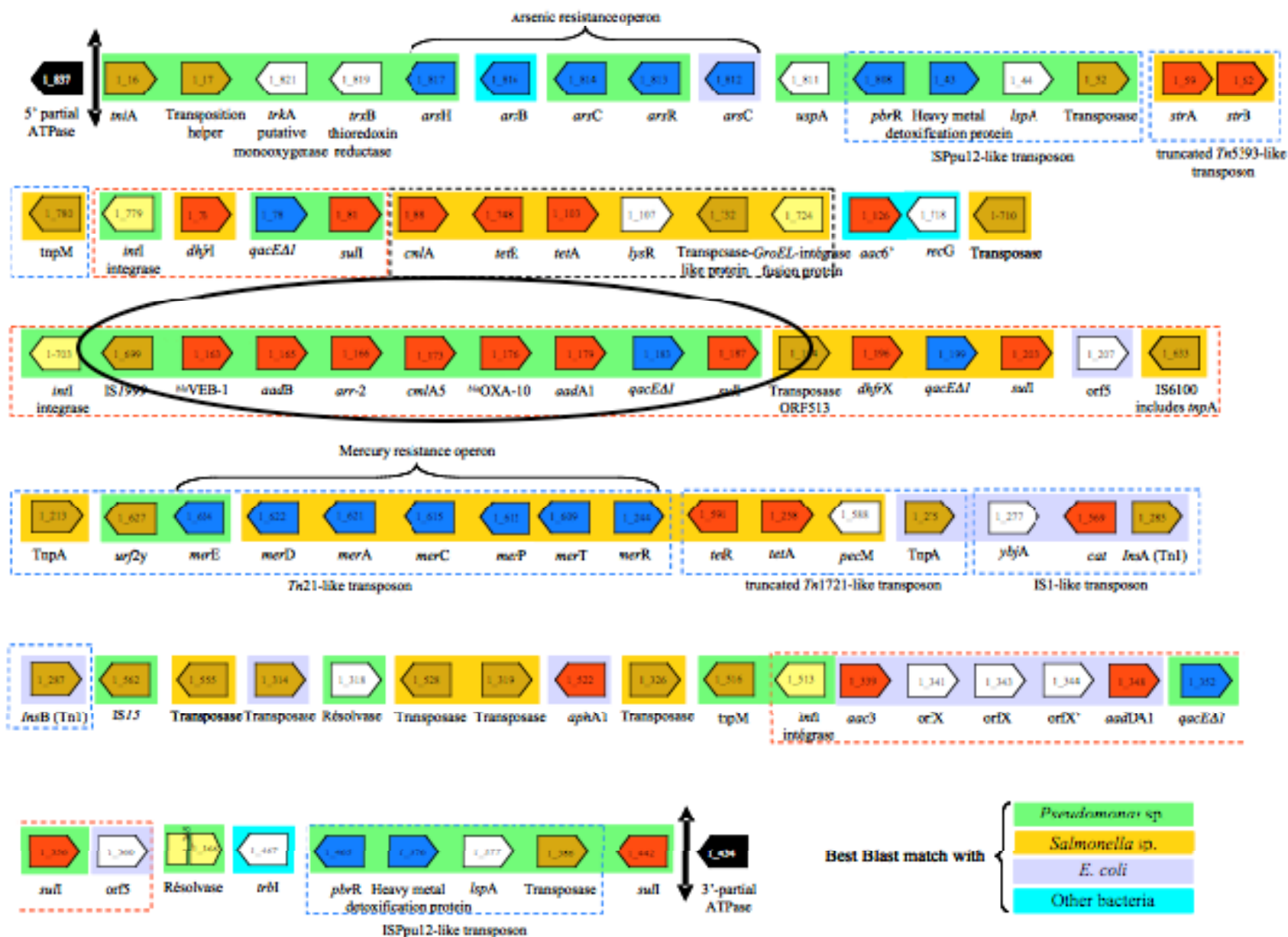
Structure of the *bla*_{VEB-1}-containing integron identified in the epidemic clone of *A. baumannii* in France



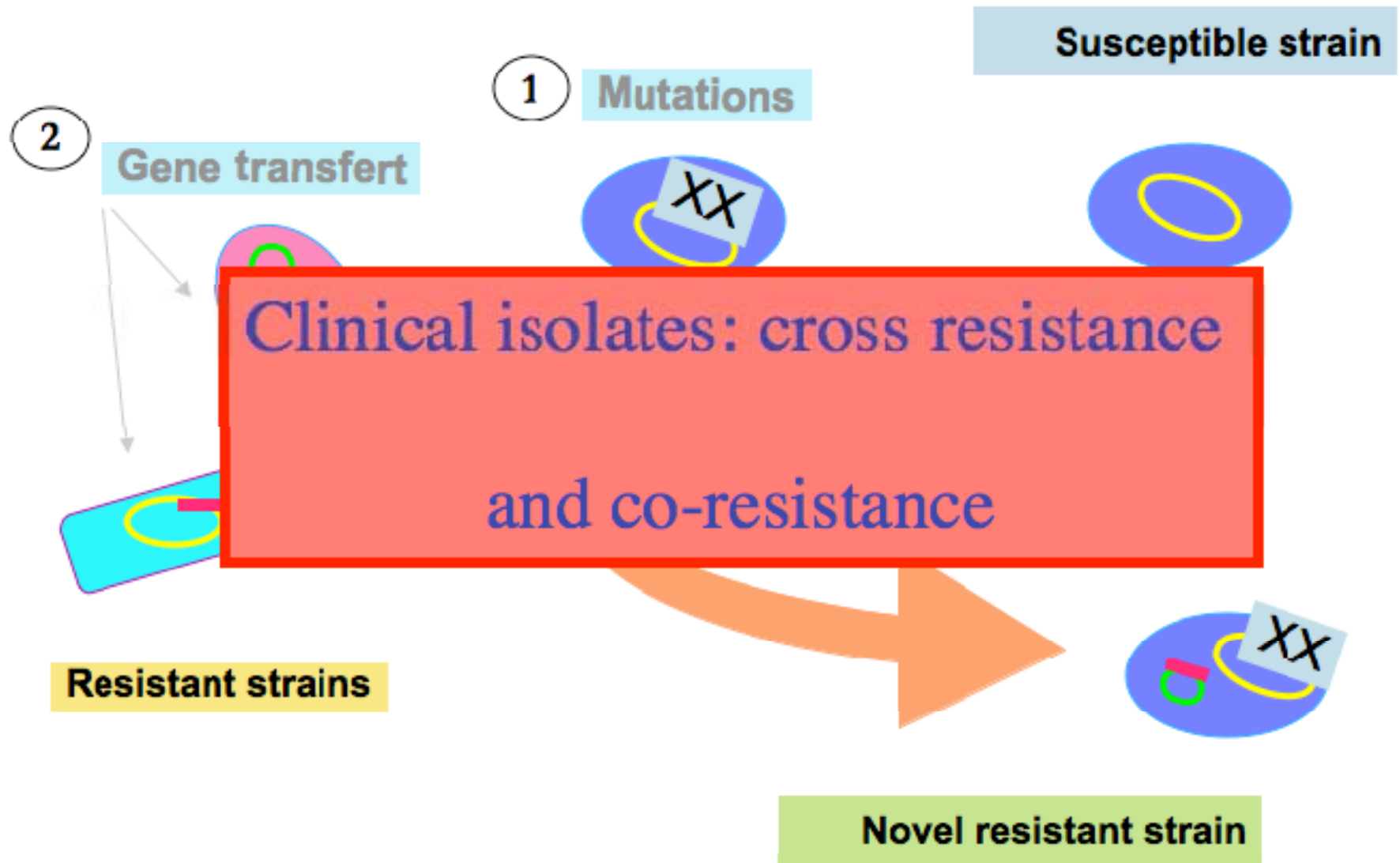
An 86-kb resistance island in *A. baumannii*

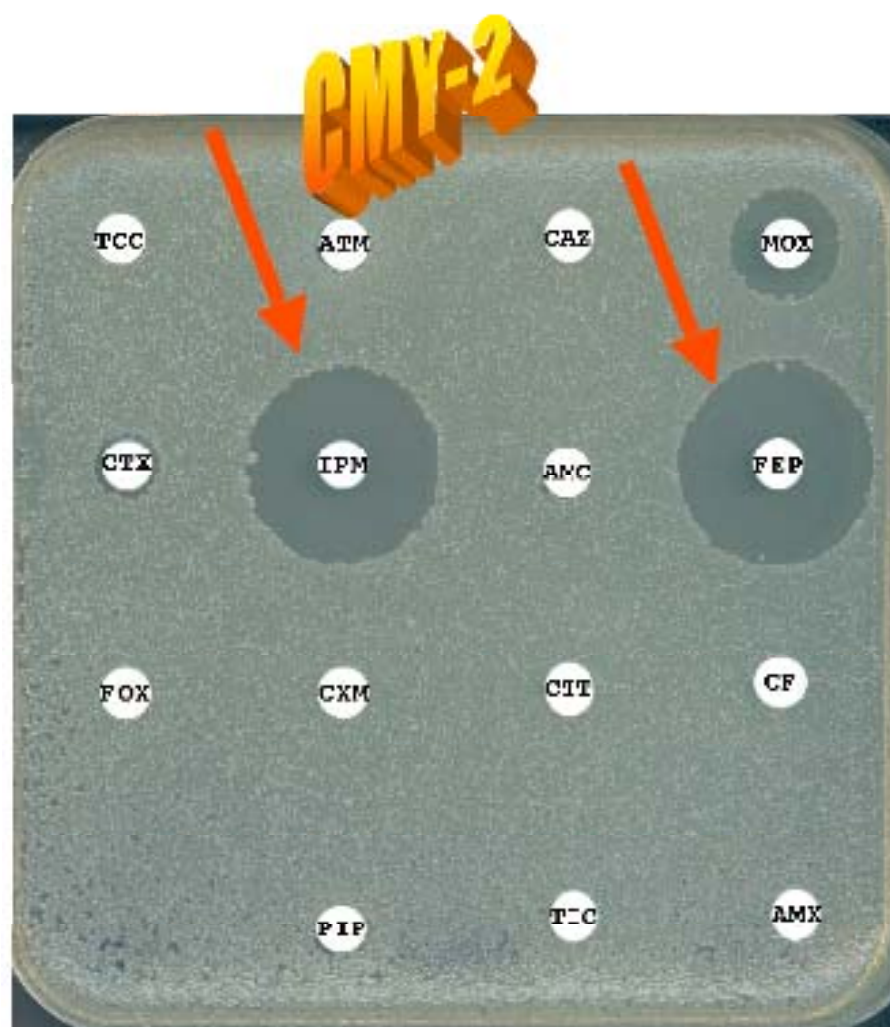


An 86-kb resistance island in *A. baumannii*

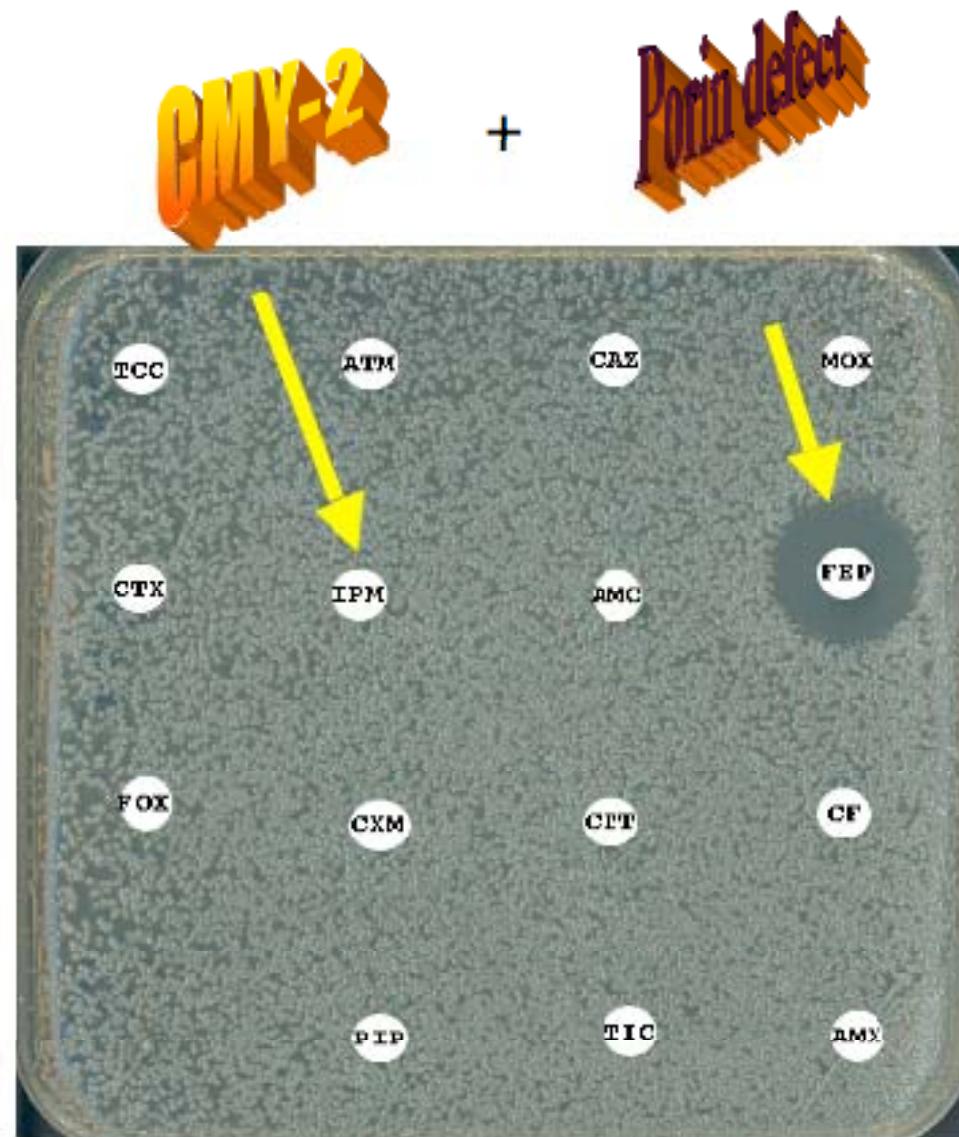


Genetics of Resistance





E. coli ND1



E. coli ND2

Superbacteria NDM-1

Experții avertizează că un nou tip de superbacterie rezistentă la antibiotice ar putea face victime în întreaga lume. Rezistența superbacteriei la antibiotice este dată de enzima New Delhi metallo-beta-lactamase-1.

■ **NDM-1:** Bacteria poate deveni rezistentă la gama de antibiotice carbapenem, folosită doar în cazuri de urgență.



■ **Origine:** Este răspândită în India, Pakistan și Bangladesh. Acum, NDM-1 a ajuns și în SUA, Canada, Australia și Olanda.

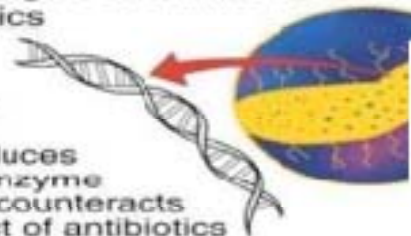
■ **Gazde:** Enzima NDM-1 a fost găsită în două tipuri de bacterii: *E.coli* - ce atacă intestinele - și *Klebsiella* - ce atacă plămânii.

Superbug gene

Discovered in pneumonia and E.coli bugs resistant to last-line antibiotics

NDM-1

- Produces an enzyme that counteracts effect of antibiotics
- Found in India, Pakistan



LE FIGARO santé

Alerte aux nouvelles bactéries ultrarésistantes

Venues d'Inde, elles survivent à presque tous les antibiotiques et circulent déjà dans plusieurs pays.



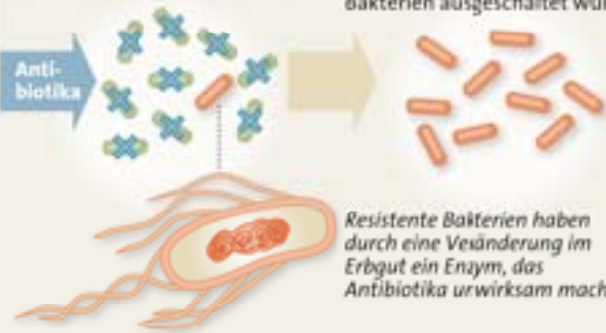
NDM-1, nouvelle molécule antibiotique d'une efficacité remarquable dans les hôpitaux. Les bactéries ultrarésistantes ont été découvertes en Inde en 2008. Elles sont capables de résister à presque tous les antibiotiques. Elles ont été découvertes en Europe en 2011. Elles sont capables de résister à presque tous les antibiotiques. Elles ont été découvertes en Europe en 2011. Elles sont capables de résister à presque tous les antibiotiques.

Gefährliche „Superbakterien“

- „Normale“ Bakterien, werden von Antibiotika abgetötet
- Resistente Bakterien, Antibiotika wirken nicht

Beispiel Patient erhält Antibiotika gegen eine Infektion mit dem Bakterium Escherichia coli

Resistente Keime können sich ungehindert ausbreiten, auch weil „Konkurrenz“ durch andere Bakterien ausgeschaltet wurde



Resistente Bakterien haben durch eine Veränderung im Erbgut ein Enzym, das Antibiotika unwirksam macht

Gründe für Entwicklung resistenter Keime

- ▶ Häufiger, oft unnötiger Einsatz von Antibiotika (z.B. ge...
- ▶ Zu kurze Antibiotika-Einnahme durch Patienten
- ▶ Besonders gefährliche resistente Keime entstehen in 5

Neue Keime

- ▶ Entstehung in Indien/Pakistan
- ▶ Durch Gen NDM-1 sind auch Peneme unwirksam (Antibiotika, die bisher bei mehr...

Grafik: © APA, Quelle: AJ

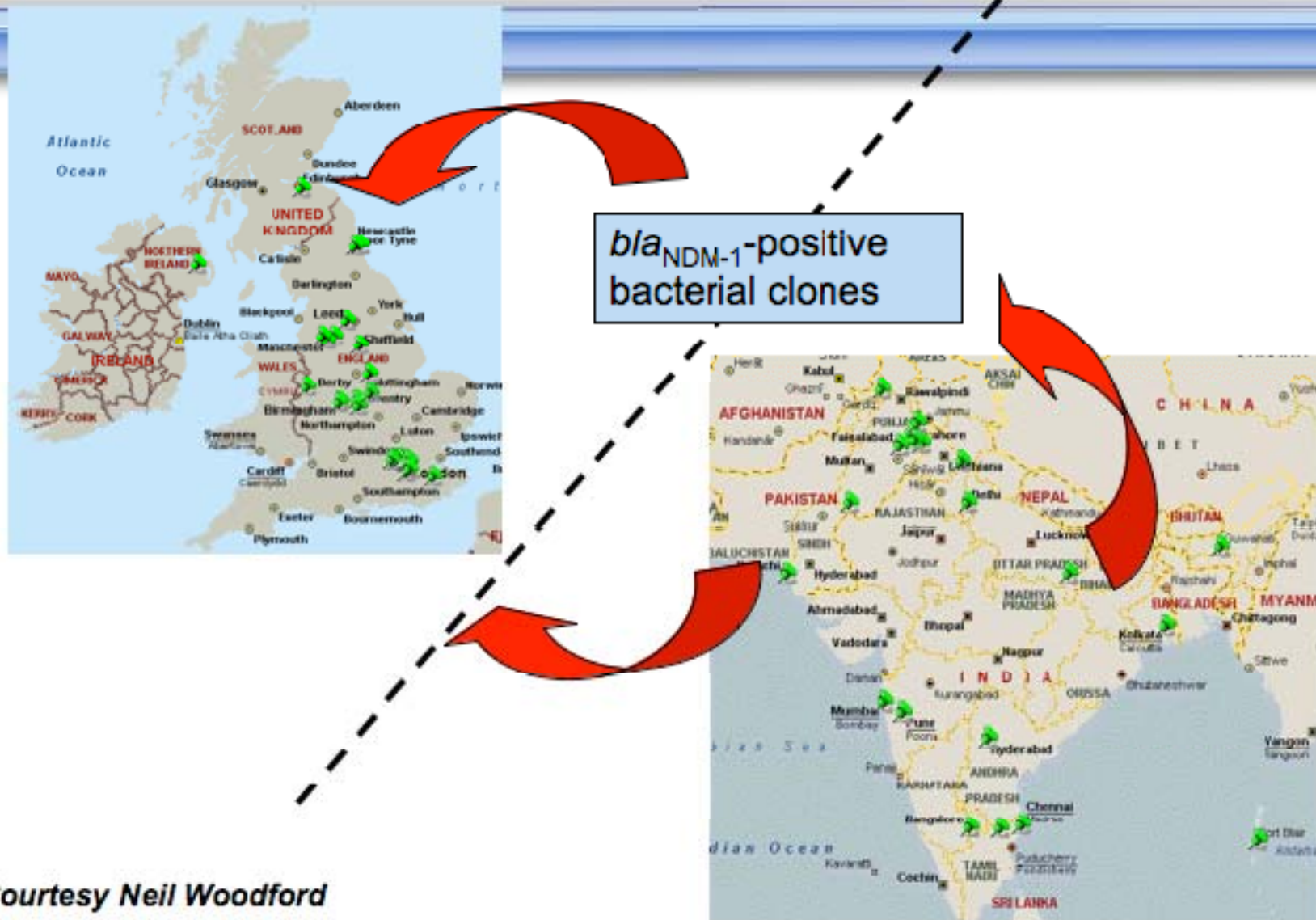


Daily Mail 24 HOURS A DAY

theguardian

THE HINDU

Spread of NDM-1 from India/Pakistan to the UK



Courtesy Neil Woodford

E. coli NDM-1, Australia

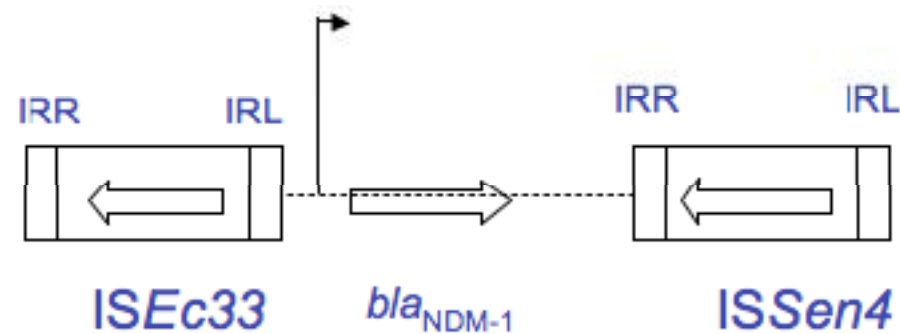


Poirel L, Lagrutta E, Taylor P, Pham J, Nordmann P

Emergence of metallo- β -lactamase NDM-1-metallo- β -lactamase producing multidrug resistant *Escherichia coli* in Australia. *Antimicrob Agents Chemother*, 2010, 54, 4914-6

E. coli NDM-1, Australia

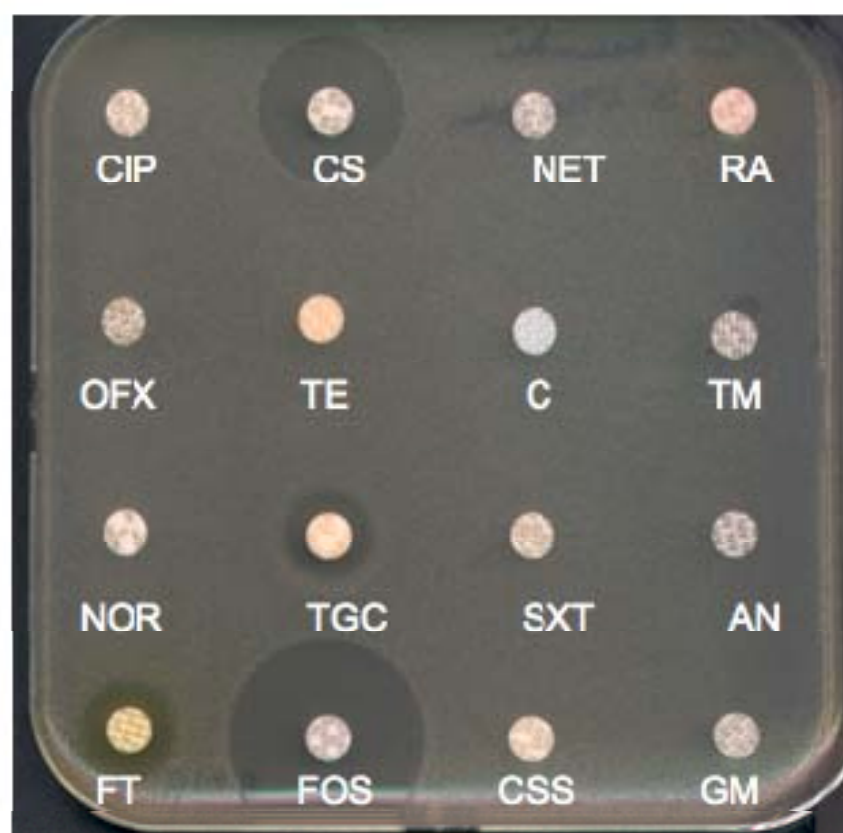
- Plasmids: 160, 130, 80, 30 kb
- Resistance genes:
 - *bla*_{NDM-1}
 - *bla* CTX-M-15, *bla* TEM-1
 - Methylases *armA*, *rmtB*
 - mutations in gyrase genes.....



Poirel L, Lagrutte
Emergence of
Escherchia coli in Australia. Antimicrob Agents Chemother, 2010, 54, 4914-6

stant

C. freundii NDM-1, France



Poirel L, Ros L, Carricajo A, Berthelot P, Pozetto B, Bernabeu S, Nordmann P
Extremely drug-resistant *C. freundii* in a patient returning from India and producing NDM-1 and other Carbapenemases. *Antimicrob. Agents Chemother.* 2010, in press

C. freundii NDM-1, France

Plasmids: 200, 160, 70, 65, 40 kb

Resistance genes:

- *bla* NDM-1
- *bla* OXA-181
- *bla* VIM-4
- *bla* CTX-M-15
- *bla* OXA-1, *bla* OXA-9, *bla* OXA-10
- *bla* TEM-1
- Methylase ArmA
- Mutations in gyrase genes....

Poirel L, Ros L, Carricajo A, Berthelot P, Pozetto B, Bernabeu S, Nordmann P
Extremely drug-resistant *C. freundii* in a patient returning from India and producing NDM-1 and other Carbapenemases. *Antimicrob. Agents Chemother.* 2010, in press

Conclusion

Class of pieces	Genes and sequences		Function
Operative	Resistance genes	<i>mecA</i>	PBP2a (methicillin resistance in <i>S. aureus</i>)
	Regulatory genes	<i>mecR</i>	<i>mecA</i> regulation expression
Translocative	Integrases	<i>int</i>	<i>attC</i> recognition in class 1 integrons
	Resolvases	TnpR	<i>res</i> sequence recognition in Tn 3-like transposons
	Invertases	<i>inv</i>	Specific recombination sequence
	Transposases	ORF513	CTX-M and AmpC capture (?)
Dispersive	Conjugation genes	<i>oriT</i>	Plasmid conjugation
	Incompatibility genes	<i>inc</i>	Plasmid interference
	Integration sequences	<i>res</i>	Recombination in Tn3-like transposons
	Insertion sequences (IS)	IS10	Recombination and integration (<i>tet</i> genes)



F. Baquero. Nature Rev Microbiol 2004; 2:510-17