

Genetic support of Extended-Spectrum β -Lactamases

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ESBLs

- TEM-like (*Enterobacteriaceae* + *P. aeruginosa*)
- SHV-like (*Enterobacteriaceae* + *P. aeruginosa* + *A. baumannii*)
- CTX-M-like (*Enterobacteriaceae* + *A. baumannii*)
- Other enzymes
(*Enterobacteriaceae* + *A. baumannii* + *P. aeruginosa*)
 - PER-1, PER-2
 - VEB-1
 - GES-like (IBC)
 - BEL-1

Variety of genetic elements

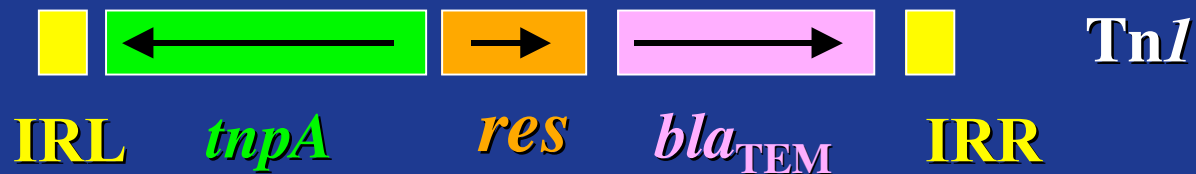
- Transposons
 - Class II transposons
 - Composite transposons
- Integrons
 - Class 1 integrons
 - *sulI*-type integrons
- Insertion sequences
- Other features

The *bla*_{TEM}-like genes

- Point mutant derivatives leading to ESBL enzymes but with similar genetic contexts
- Reservoir of *bla*_{TEM} genes ?

Tn3-type transposons

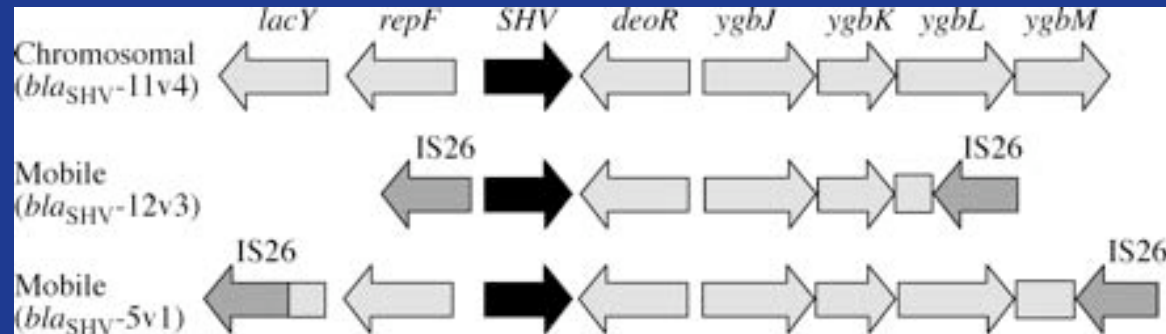
- 2 extremities corresponding to inverted repeat sequences (IR)
- 1 transposase (TnpA) => recognizes IR extremities in the transposition process and cuts target DNA
- 1 resolvase (TnpR) : negative regulator of TnpA expression and recombinase mediating the resolution of the transposon integration process



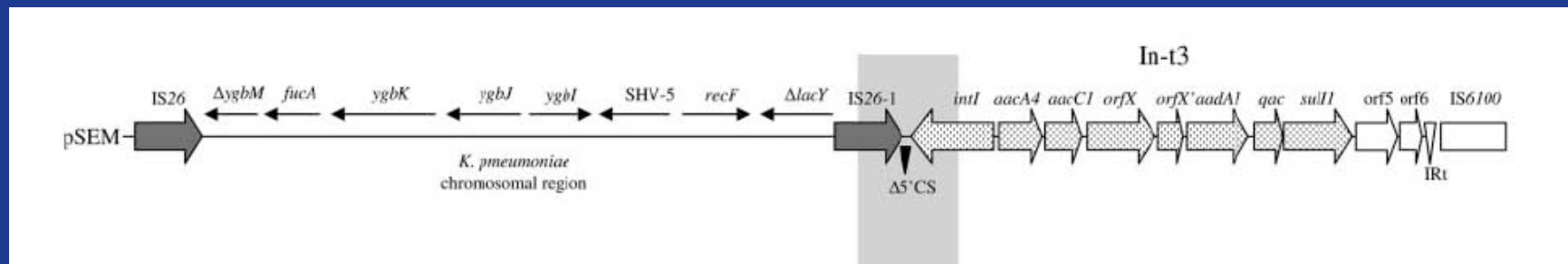
The *bla*_{SHV}-like genes

- Point mutant derivatives leading to ESBL enzymes
 - Originating from *Klebsiella pneumoniae*

IS26-mediated mobilization of *bla*_{SHV}



Ford & Avison, JAC 2004



Miriagou, Carattoli et al., AAC 2005

The *bla*_{CTX-M}-like genes

- Five main groups made of
 - CTX-M-1
 - CTX-M-2
 - CTX-M-8
 - CTX-M-9
 - CTX-M-25
- Originate from *Kluyvera* spp.

Main genetic structures for *bla*_{CTX-M} acquisition

ISEcp1

CTX-M of different groups

sul1-type integrons
(including CR1 element)
CTX-M-2 and CTX-M-9

Phage-related elements
CTX-M-10 (CTX-M-1 group)

A peculiar insertion sequence : *ISEcp1*

- *ISEcp1*, insertion sequence responsible for mobilization and expression of β -lactamase genes by transposition using imperfect right inverted repeats

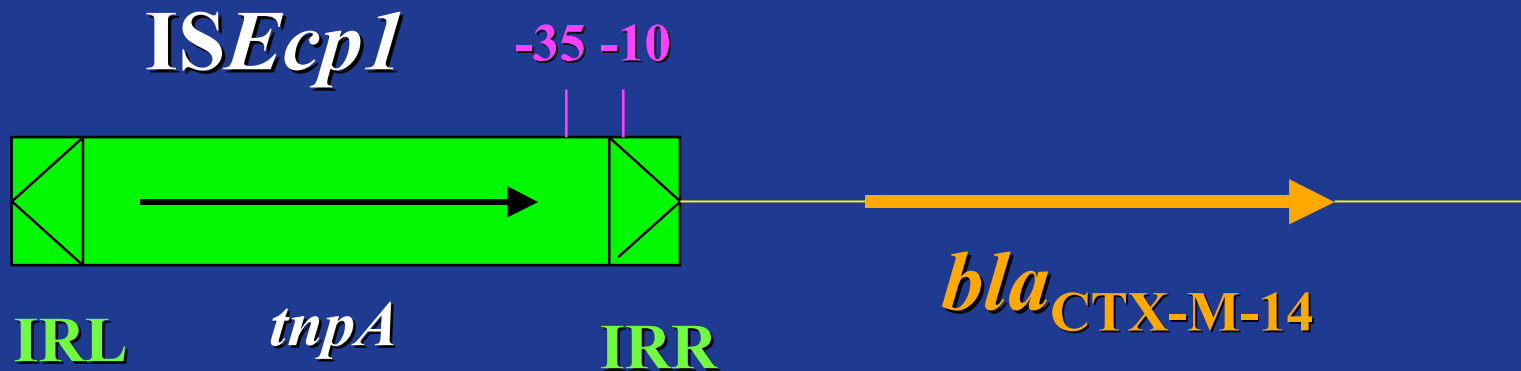


- Implications in term of mobilization and dissemination of genes coding for ESBL

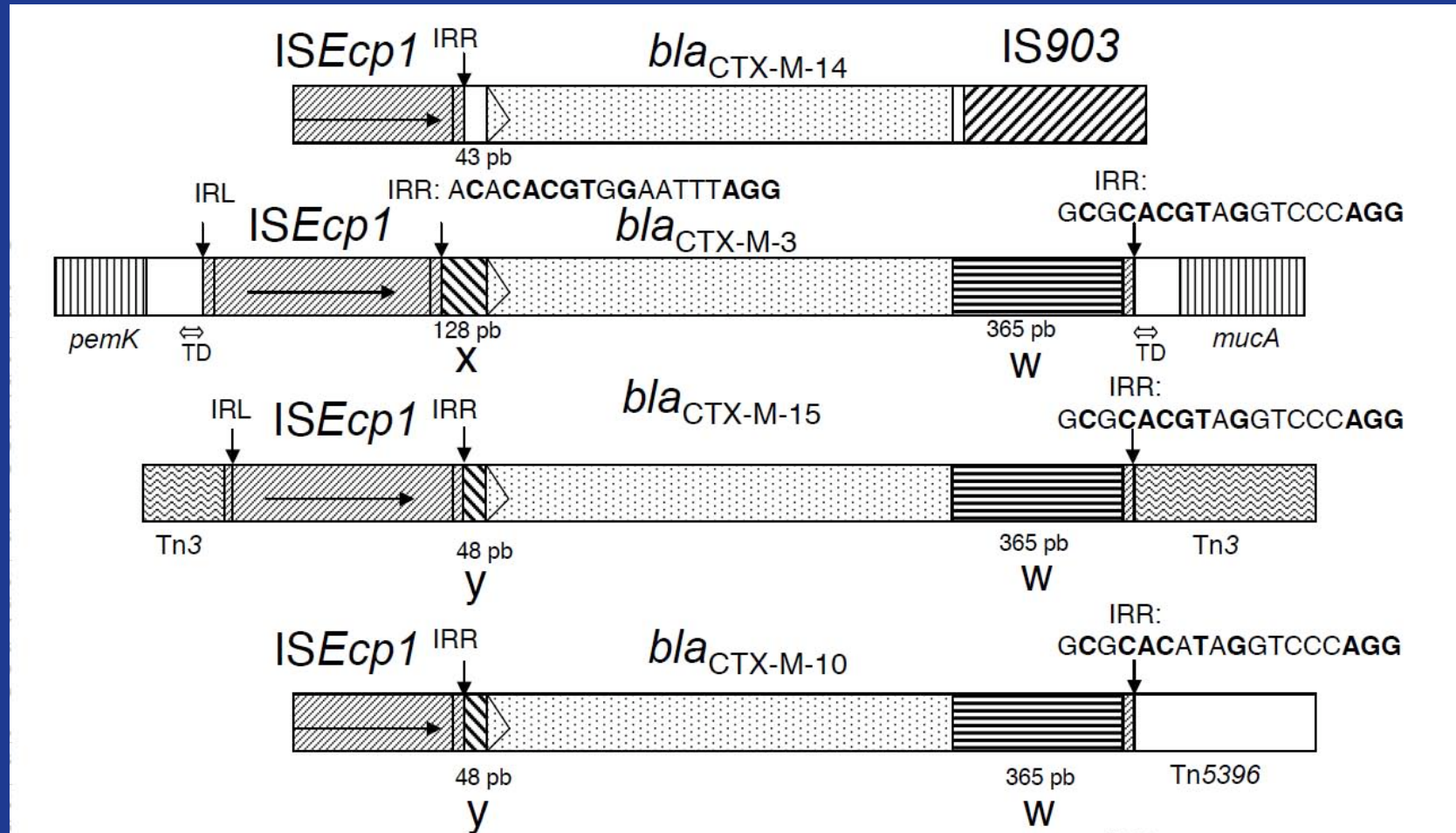
Poirel et al., AAC 2003;47:2938

Poirel et al., AAC 2005;49:447

Role of *ISEcp1* in expression of *bla*_{CTX-M-14}



Distinct mobilization events

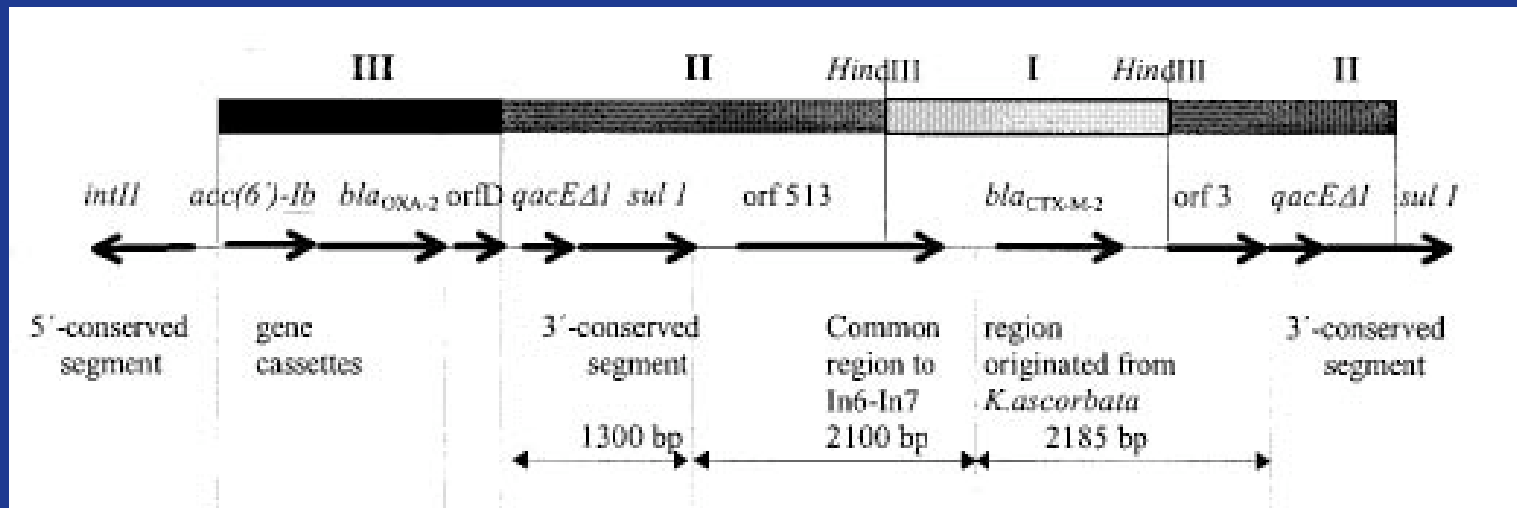


Lartigue, Poirel, Nordmann, FEMS Microb. Lett. 2004

***bla*_{CTX-M} genes associated to
sul1-type integrons**

A « complex » *sul1*-type integron

Example; In35 in a *Proteus mirabilis* isolate



↑
CR1

Arduino et al., AAC 2002, 2303

CR (Common region)

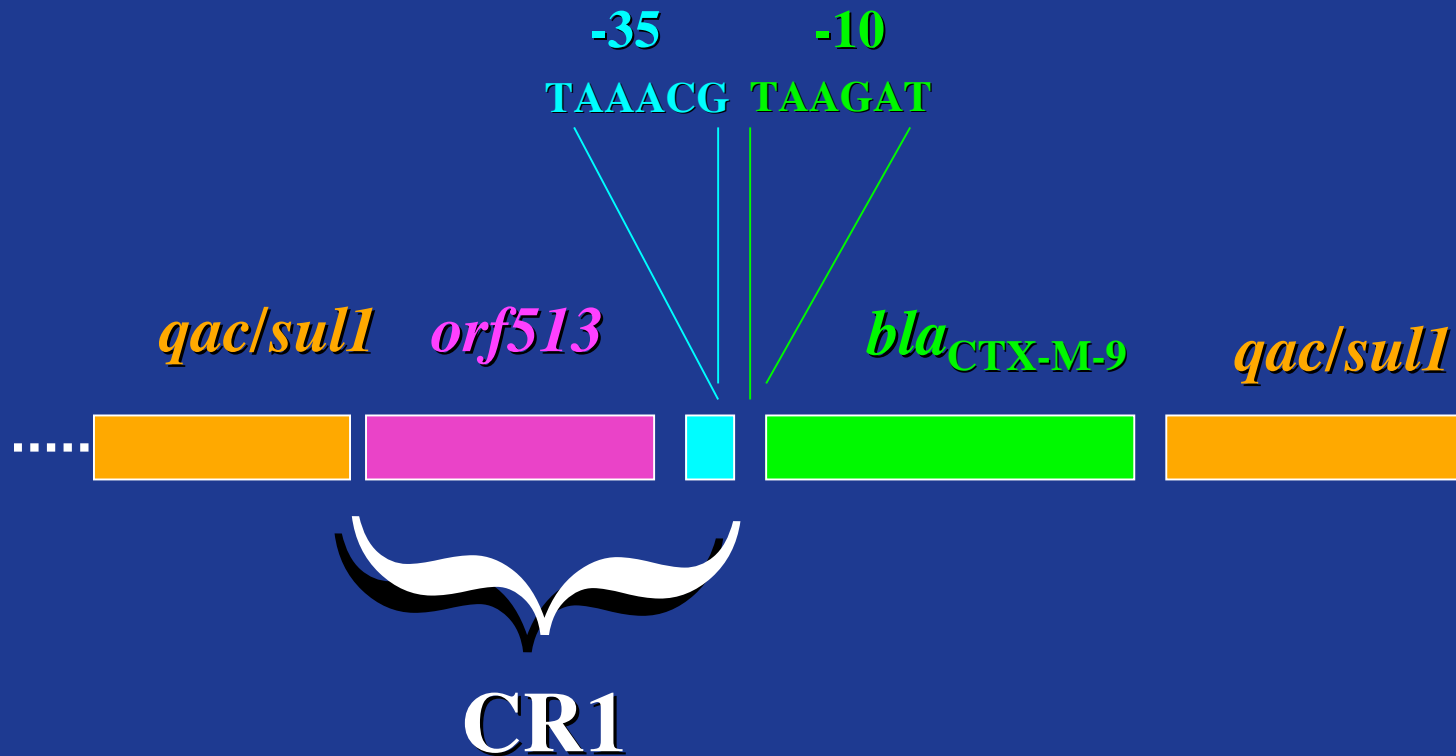
First Definition

- A transposase-like protein
- One extremity serves as a recombination site



(Partridge and Hall, AAC 2003)

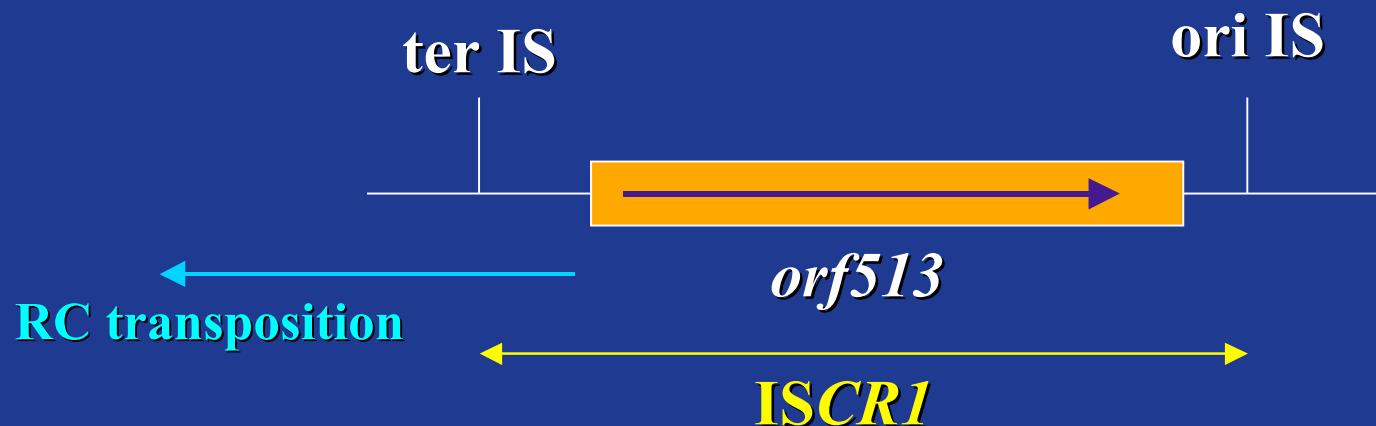
CR1 enhances the expression of *bla*_{CTX-M-2} and *bla*_{CTX-M-9}



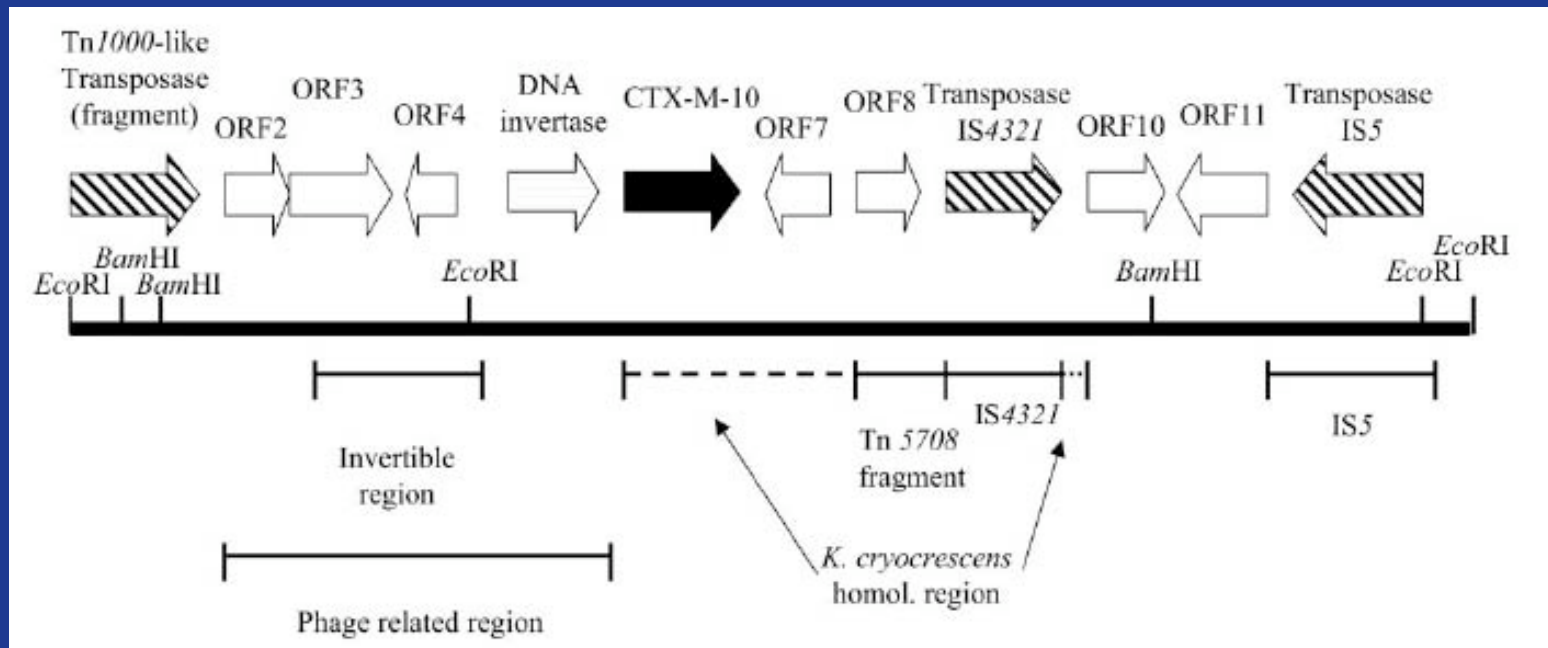
Update on the definition of CR1

T. Walsh, personal communication

- CR1 is an IS91-like transposable element
- *ISCR1* lack the terminal IRs typical of most IS elements
- *ISCR1* moves by rolling-circle (RC) transposition
- Orf513 is a transposase
- No target site duplication is generated by RC transposition



Phage-associated structure at the origin of *bla*_{CTX-M-10} acquisition in *Klebsiella pneumoniae* in Spain

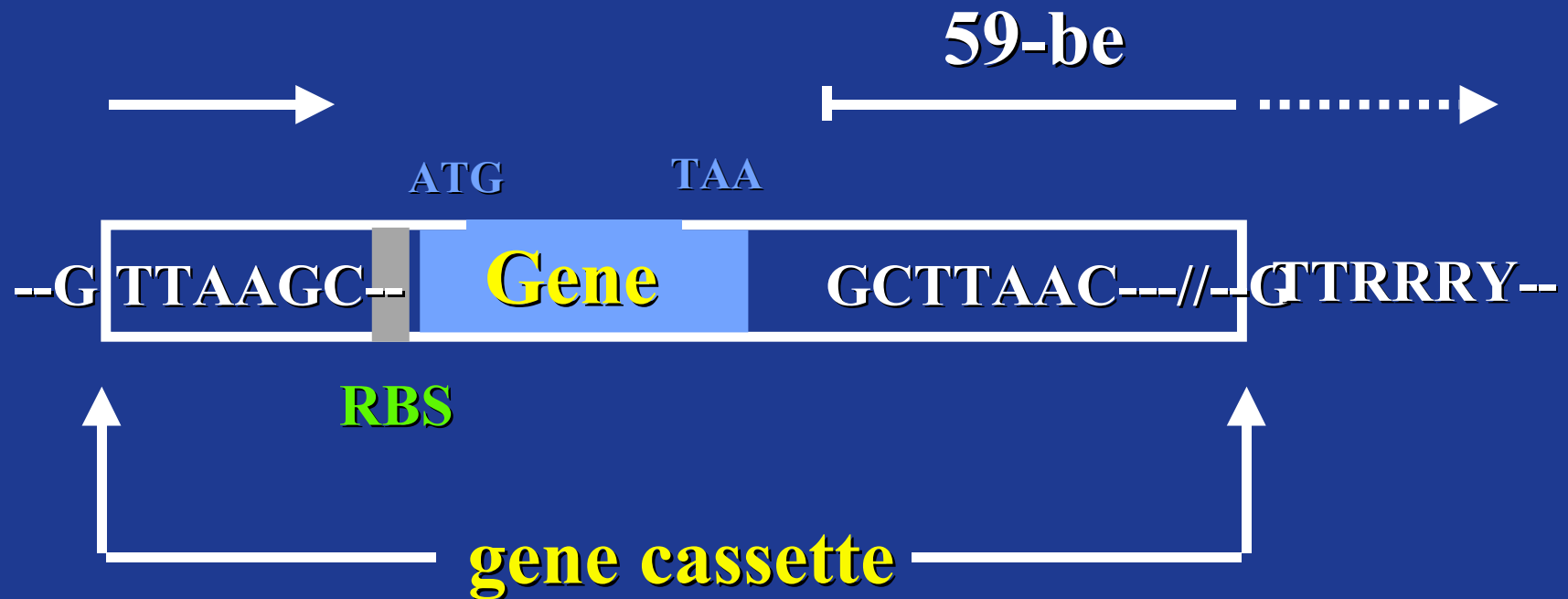


Other ESBL-encoding genes

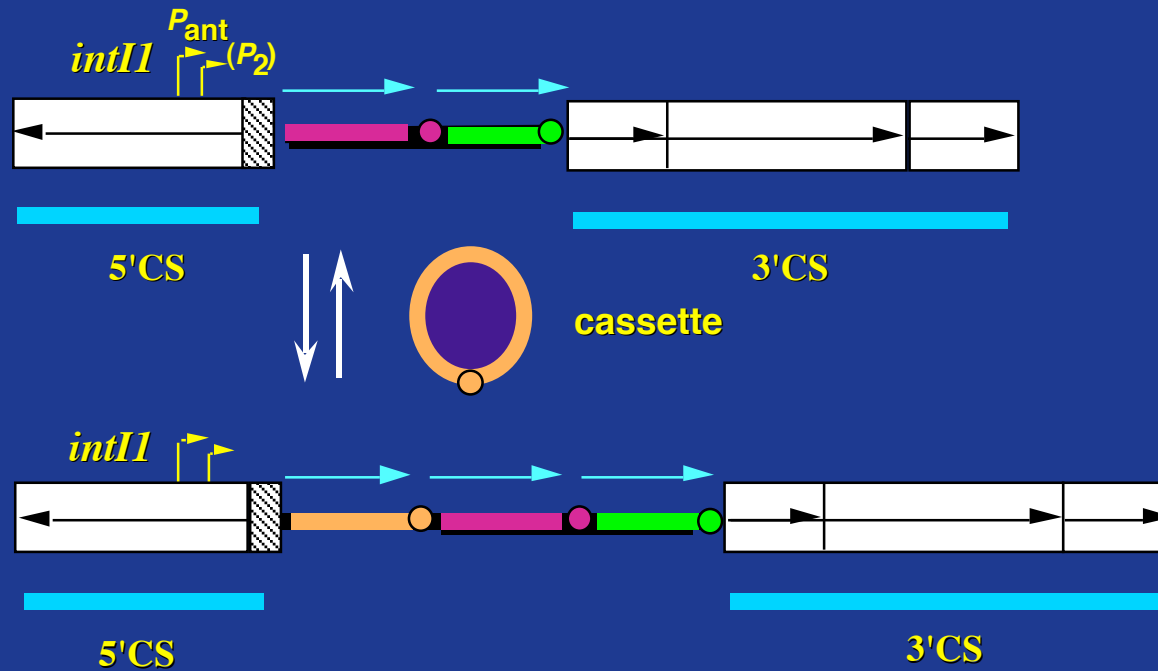
Integrans as a source of β -lactamase gene acquisition

The ESBL gene is part of a gene cassette

Structure of a gene-cassette

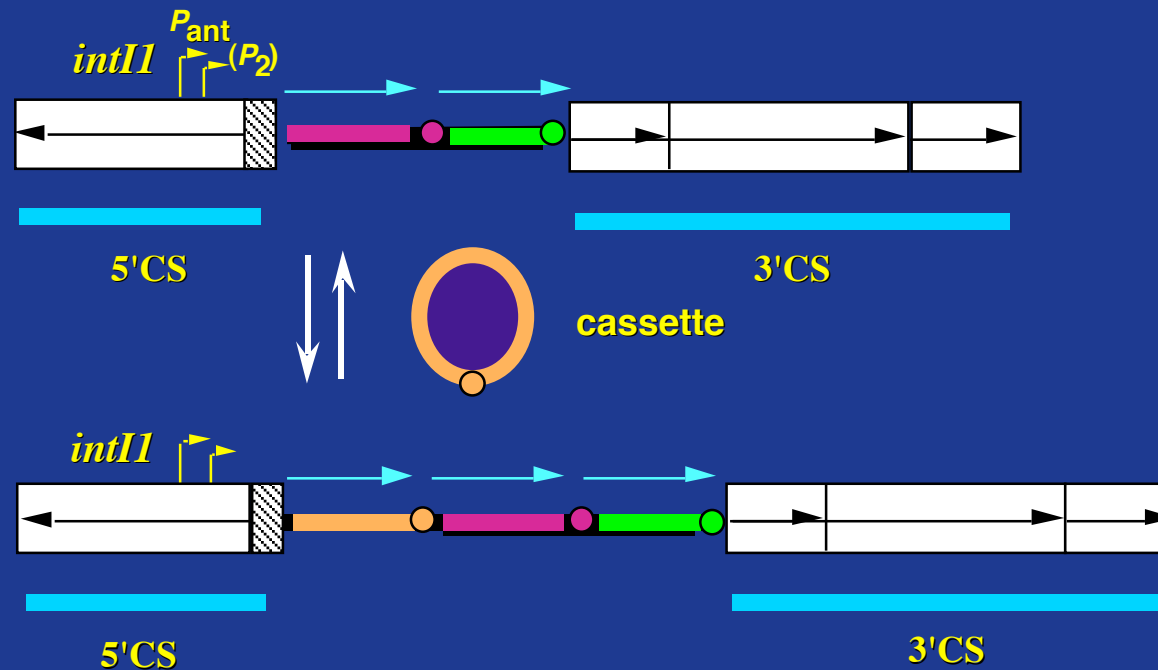


Class 1 integrons



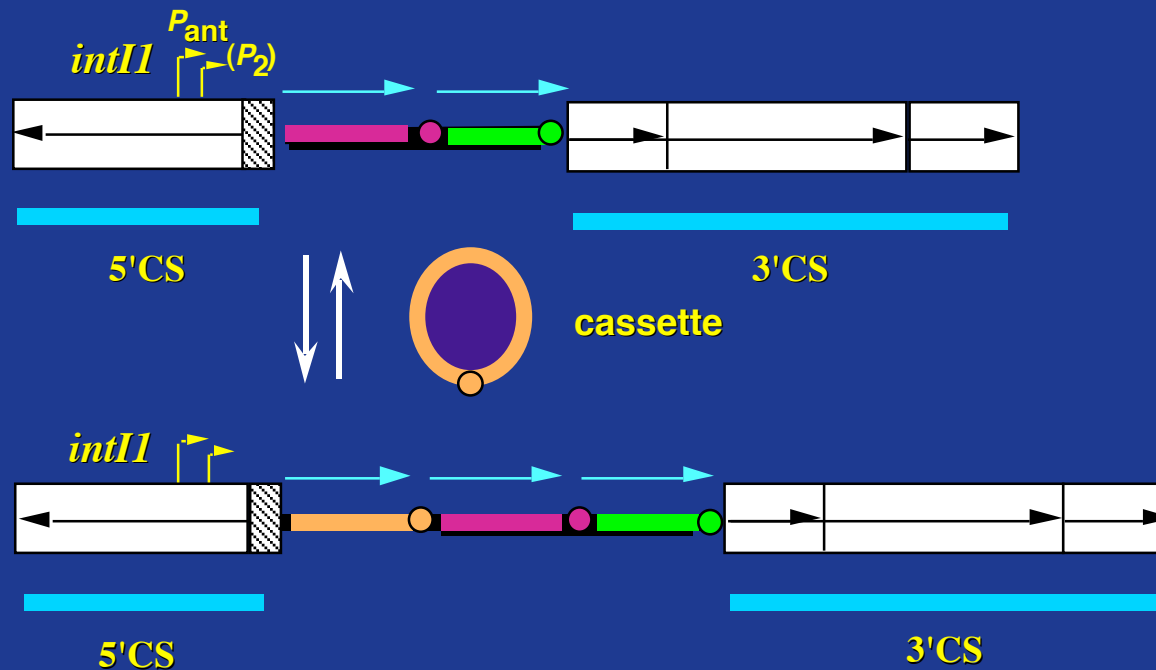
- **Consequences : co-resistance**

Class 1 integrons



- Consequences : co-resistance ; co-expression

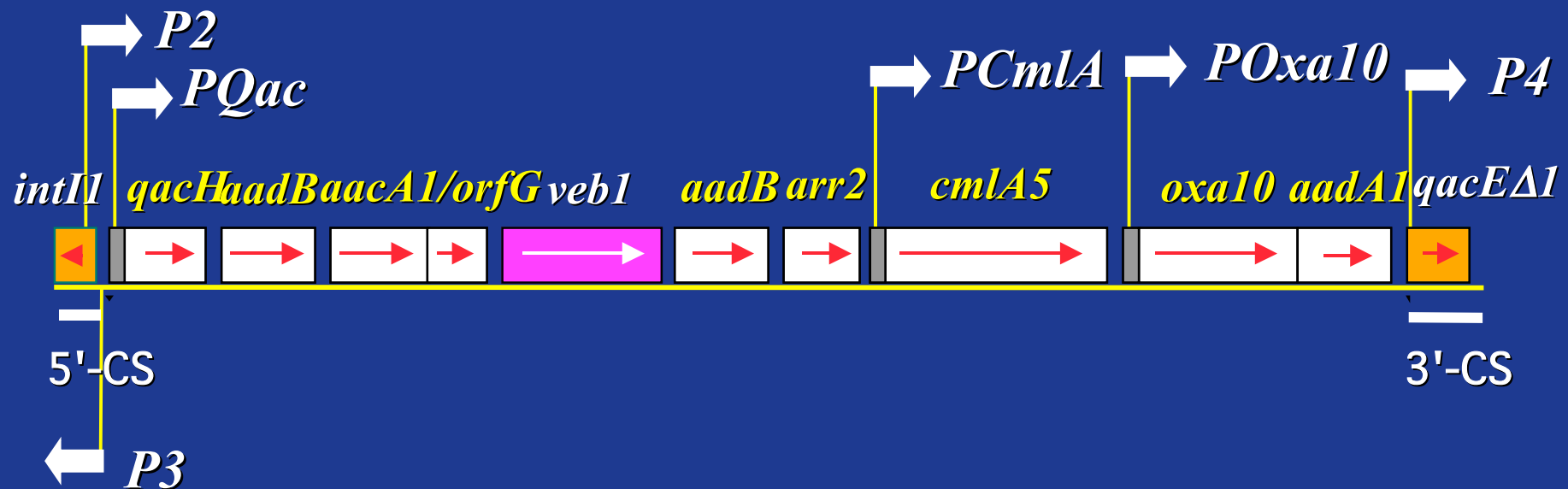
Class 1 integrons



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

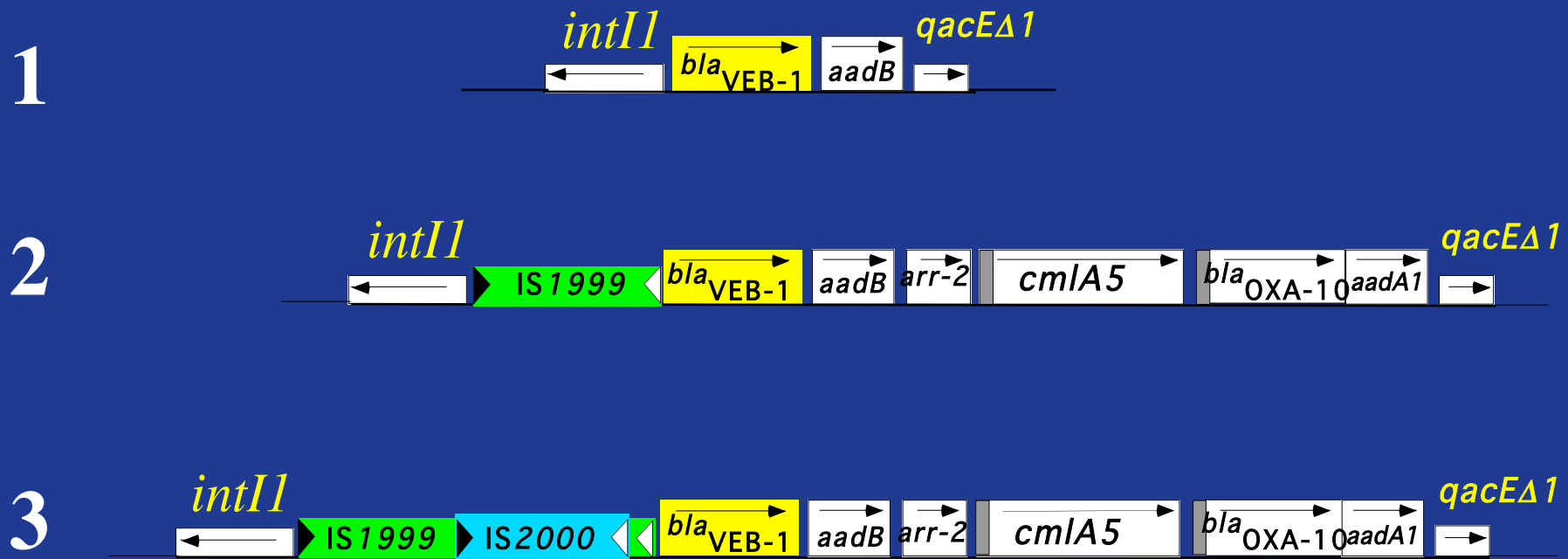
*bla*_{VEB-1} as a gene cassette identified in In53 in *E. coli*

In53



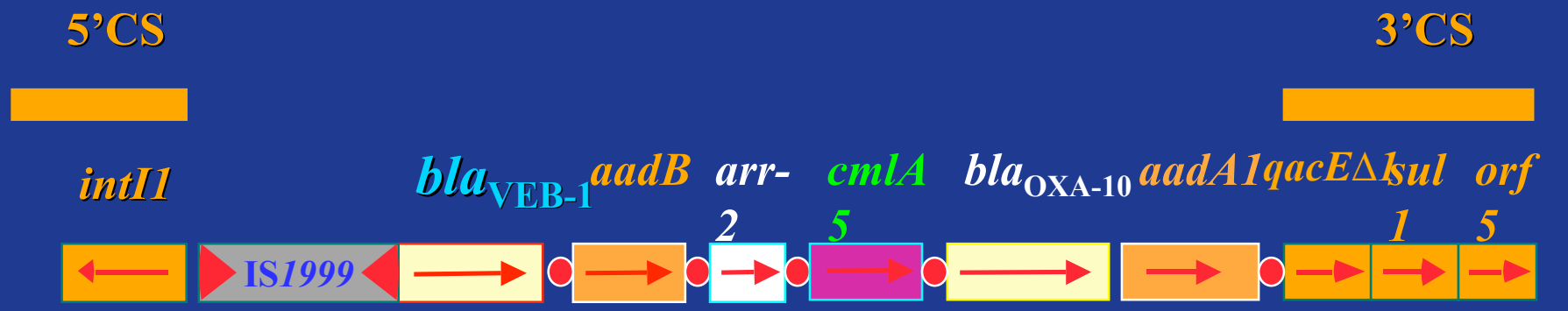
Naas et al., J Bact 2001

*bla*_{VEB-1} in *P. aeruginosa*



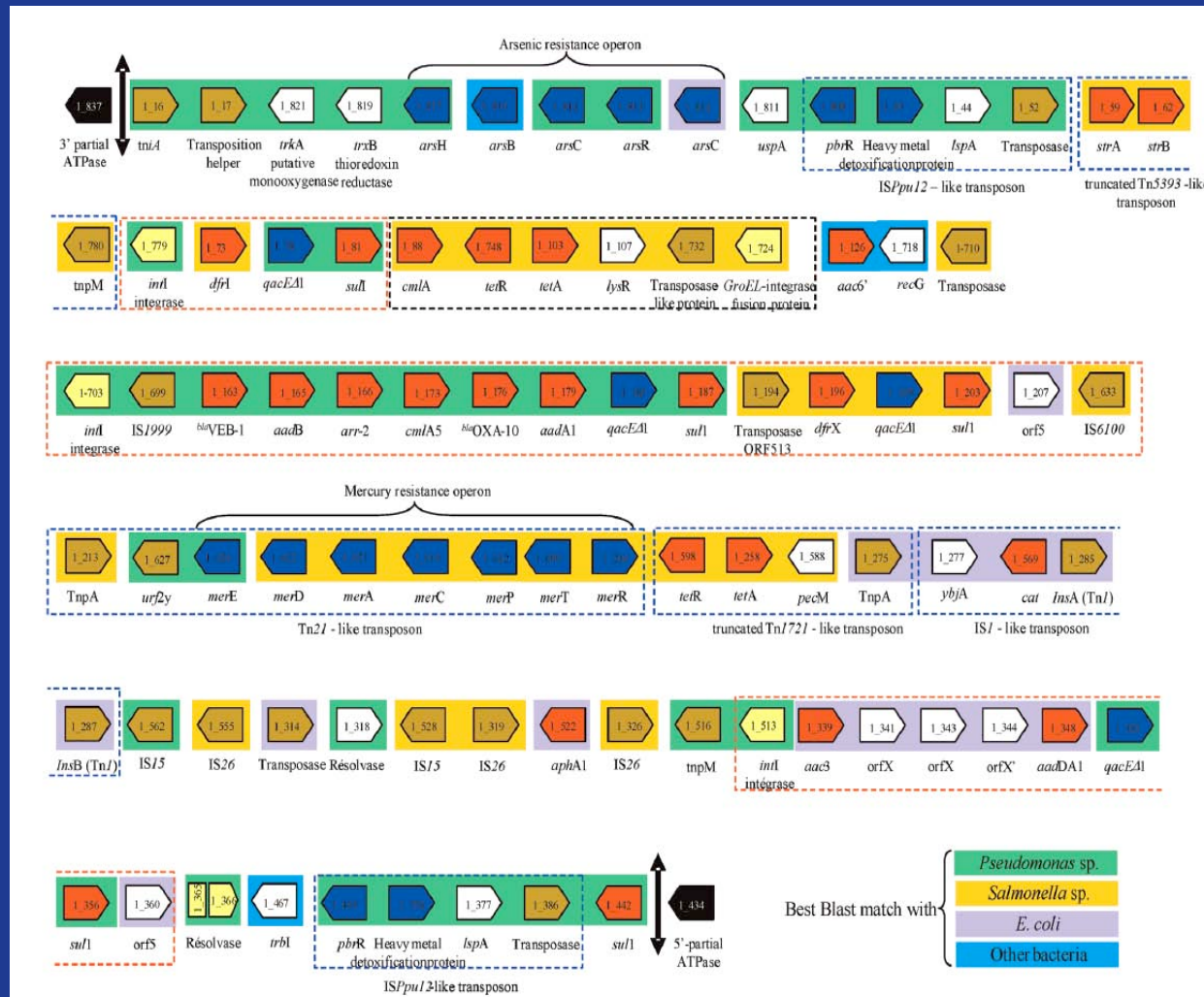
Girlich *et al.*, CID 2002

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



Poirel *et al.*, JCM 2003;41:3542

An 86-kb resistance island

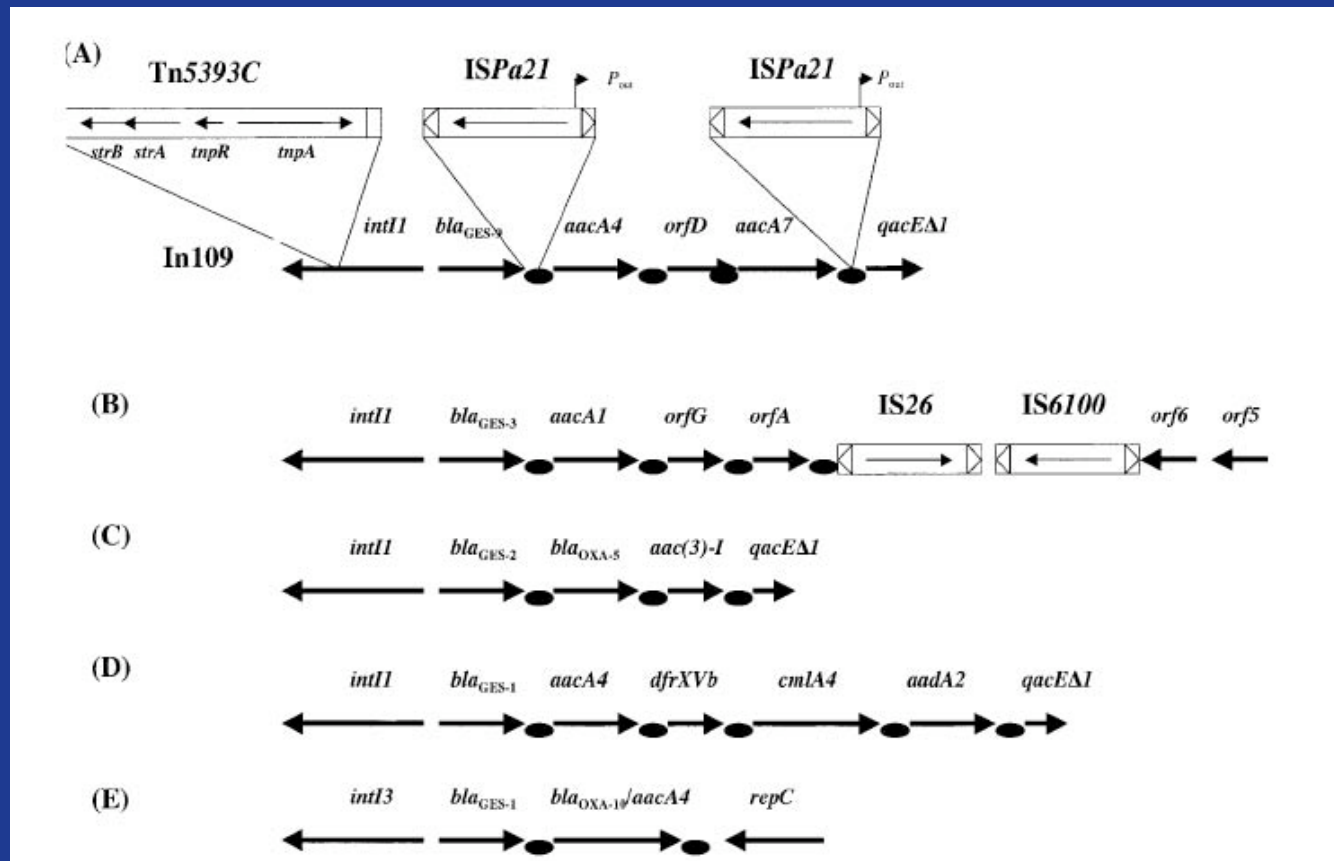


Fournier et al., PLOS Genetics 2006

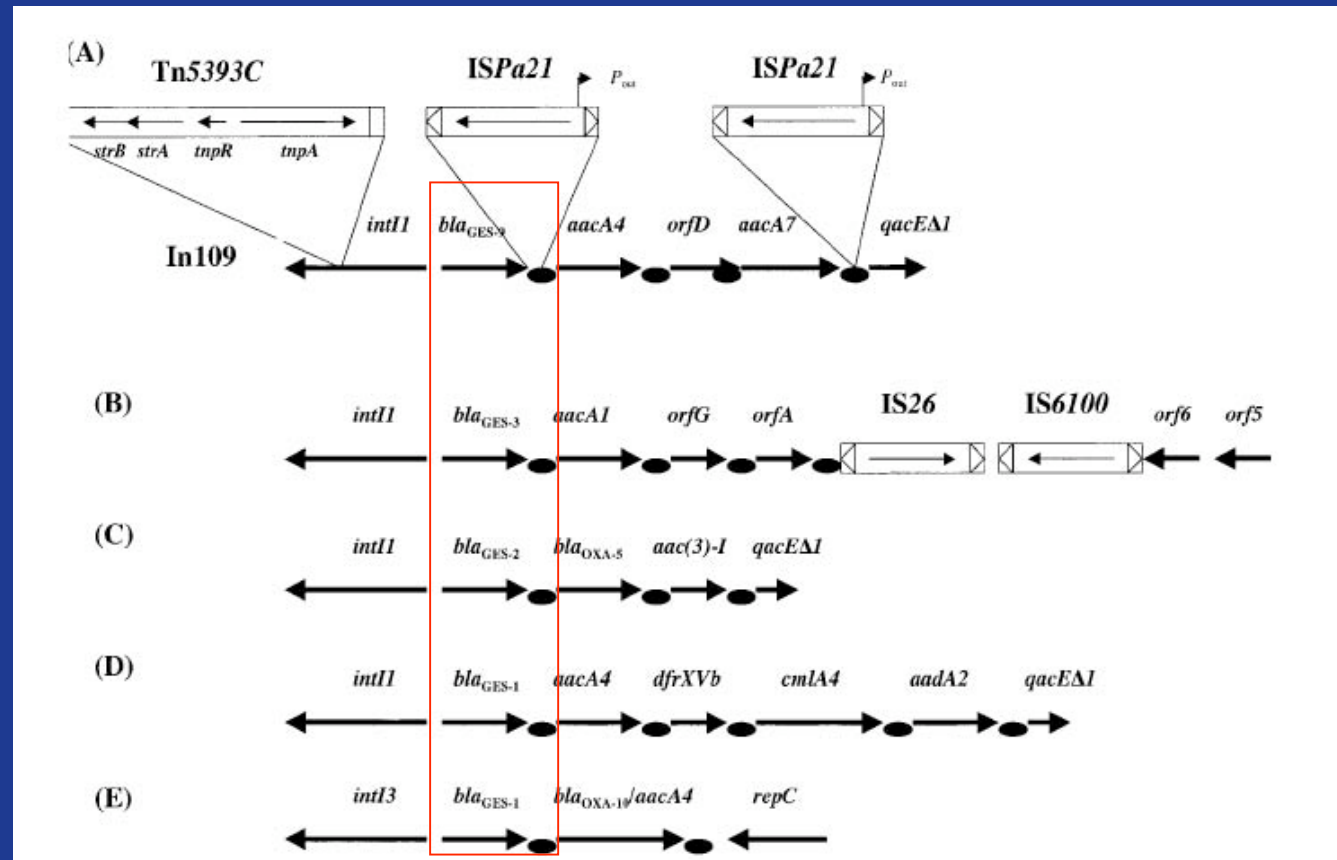
Other ESBLs encoded by gene cassettes located into class 1 integrons

- GES-type enzymes (GES-1 to -9)
(France, Greece, South Africa, Asia, South America)
- BEL-1
(Belgium)

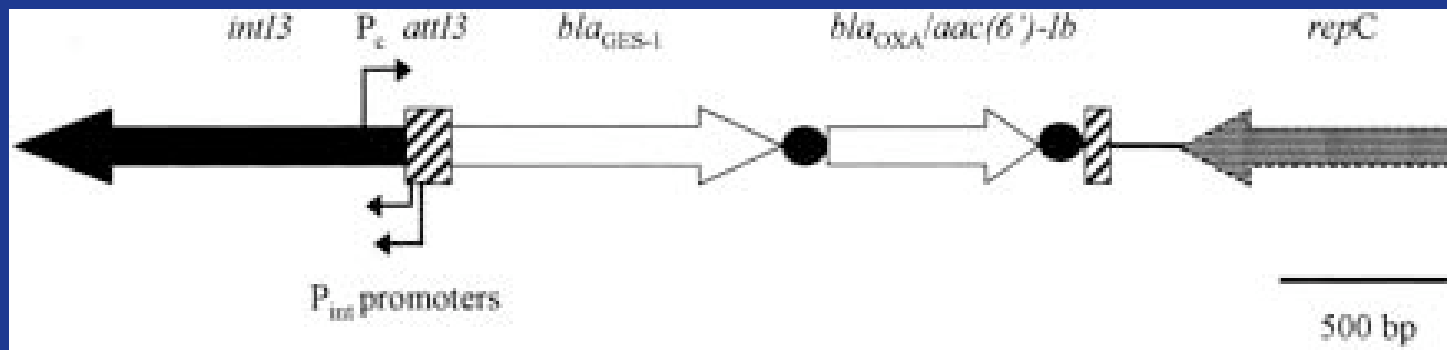
Structures of integrons harboring *bla*_{GES-like} gene cassettes



Structures of integrons harboring *bla*_{GES-like} gene cassettes

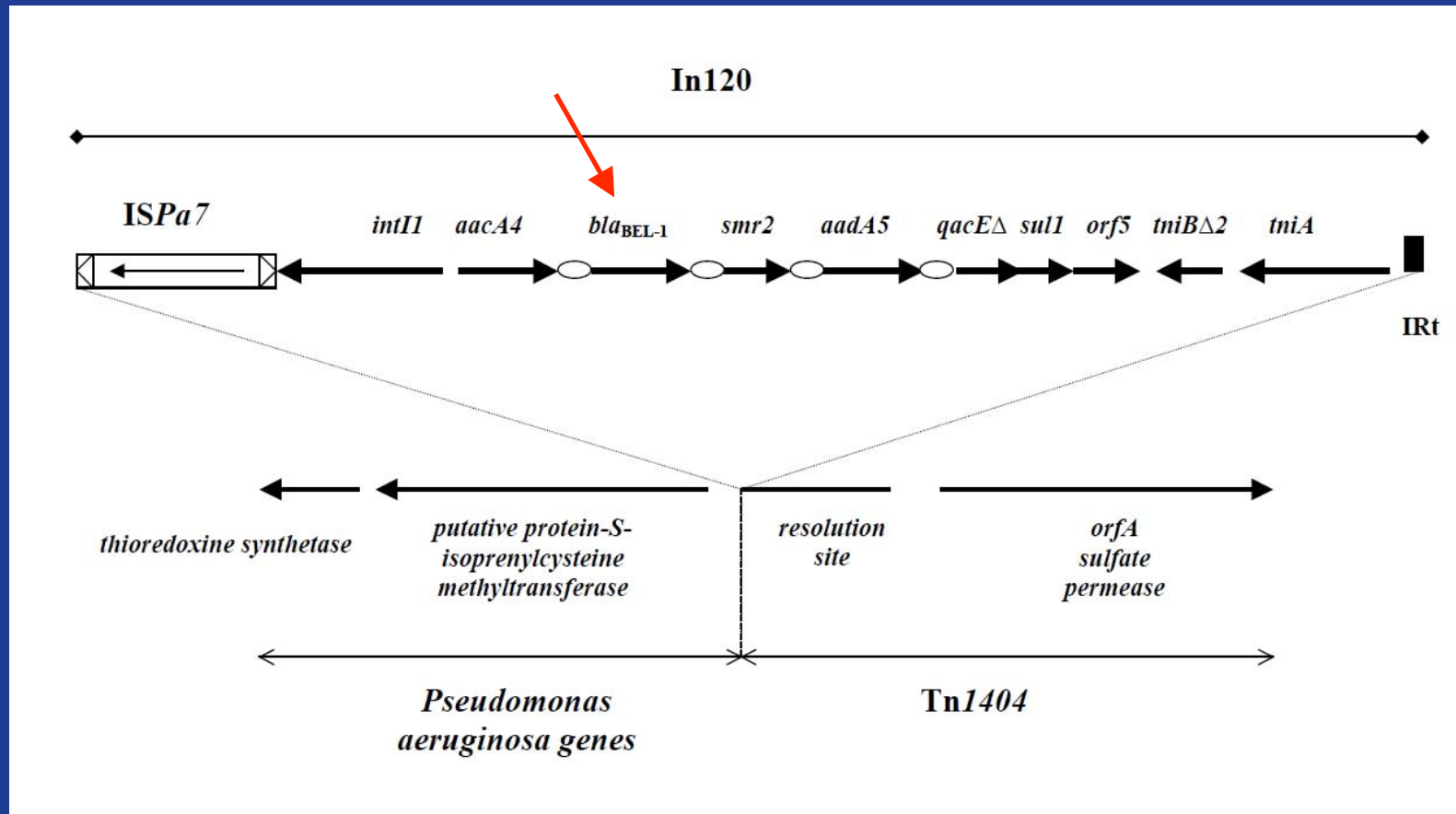


A class 3 integron at the origin of *bla*_{GES-1} acquisition

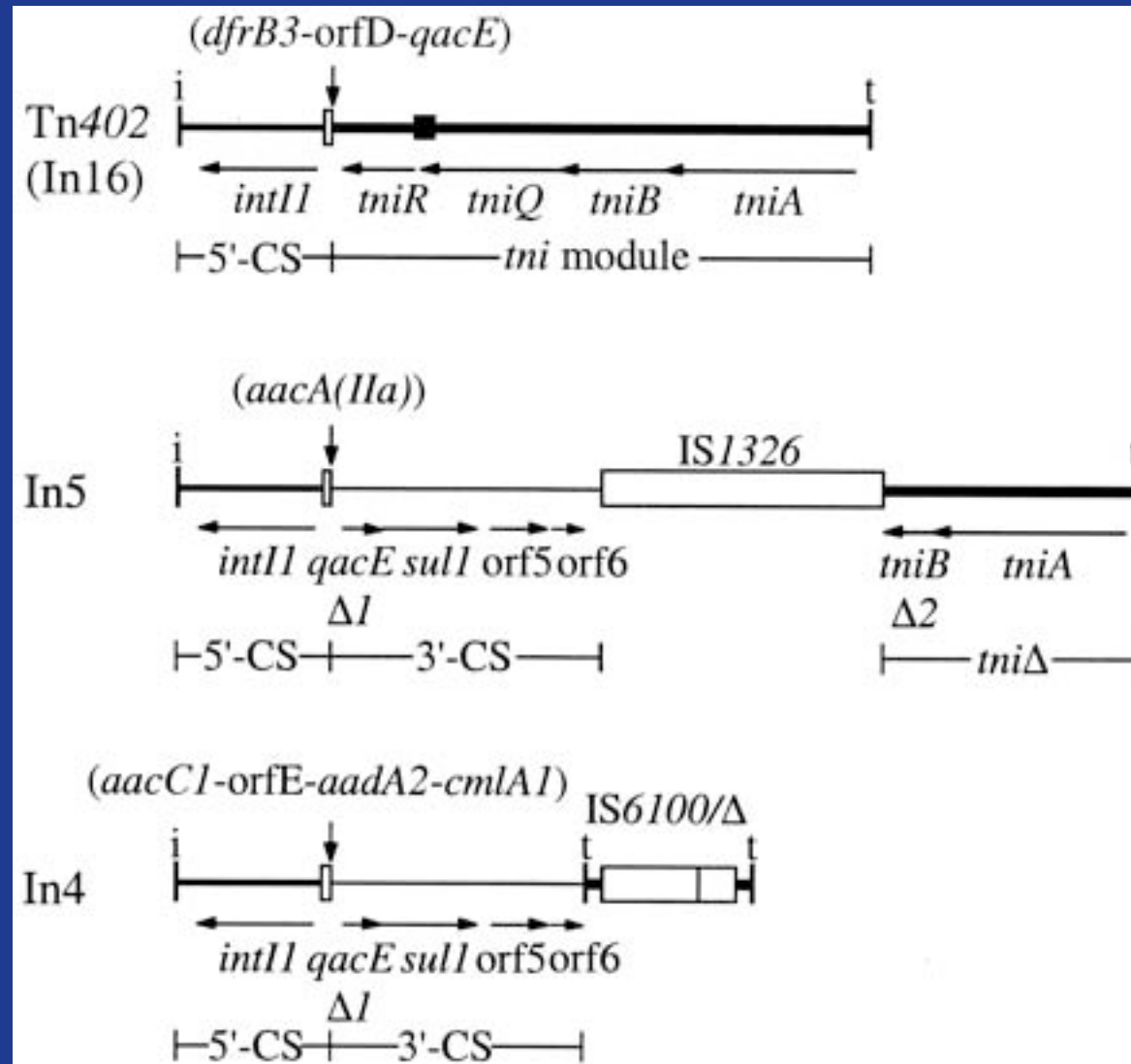


Correia *et al.*, AAC 2003

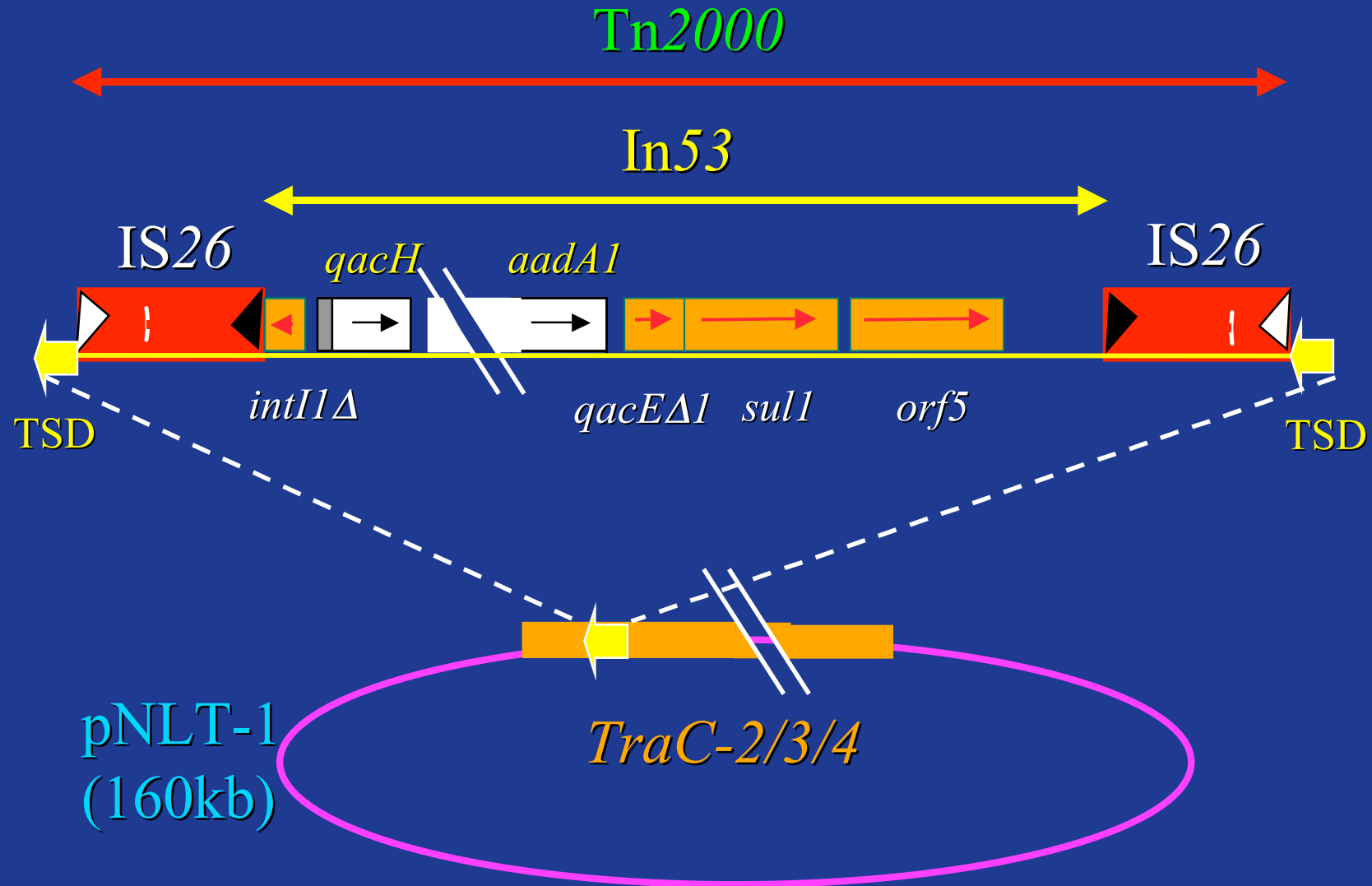
*bla*_{BEL-1} and its integron In120



The integron-borne transposons



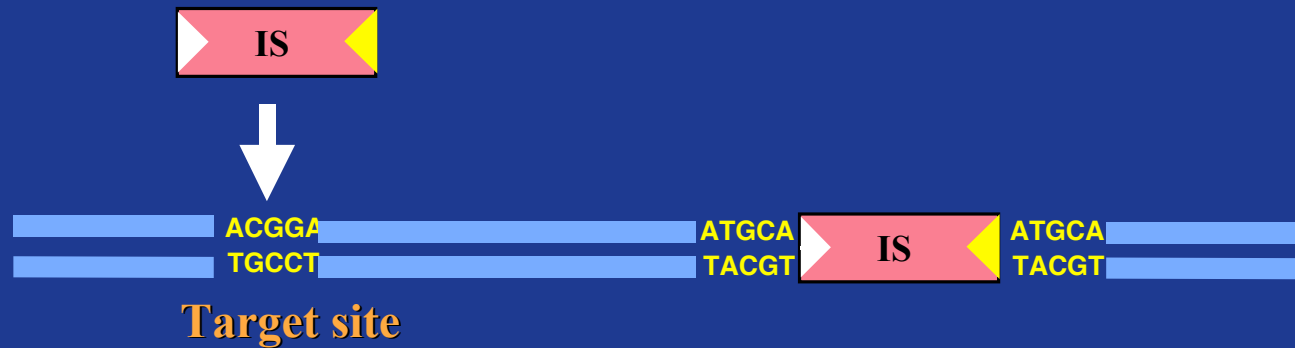
Structure of Tn2000



Naas *et al.*, J Bacteriol 2001

Mobilisation of resistance genes

- Composite transposon



Mobilisation of resistance genes

- Composite transposon = two IS bracketing the transposed fragment



Resistance gene

Mobilisation of resistance genes

- Composite transposon = two IS bracketing the transposed fragment

ACGGA
TGCCT

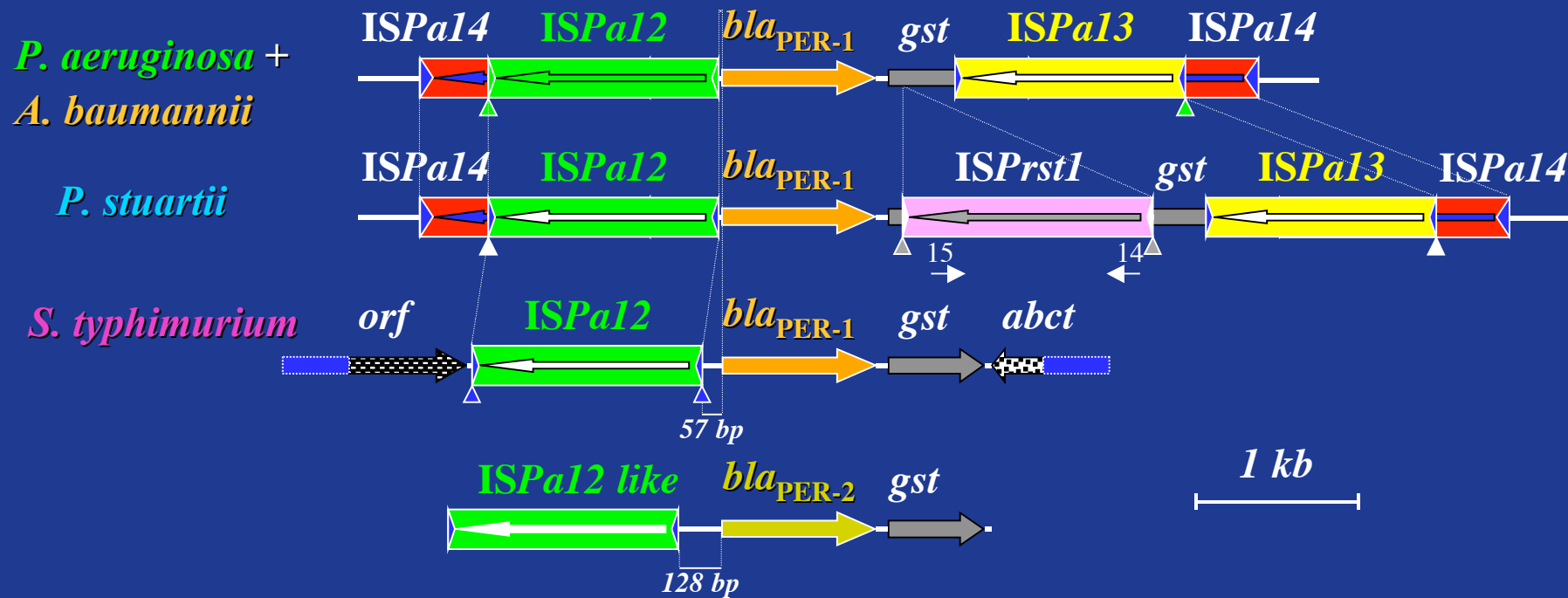
ATGCA
TACGT



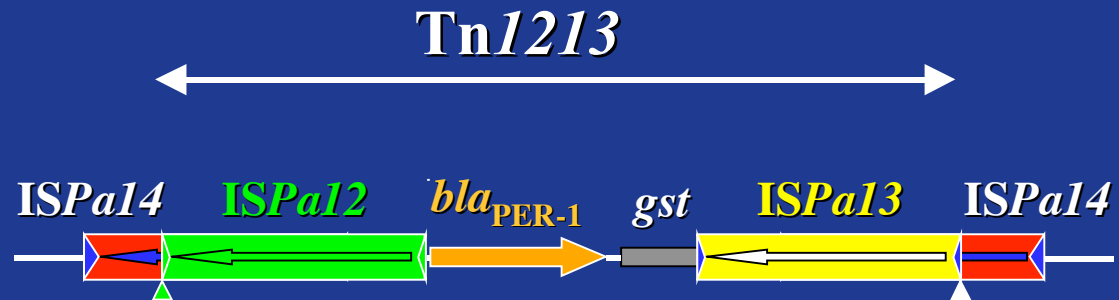
Resistance gene

**A peculiar composite transposon at
the origin of *bla*_{PER-1} acquisition**

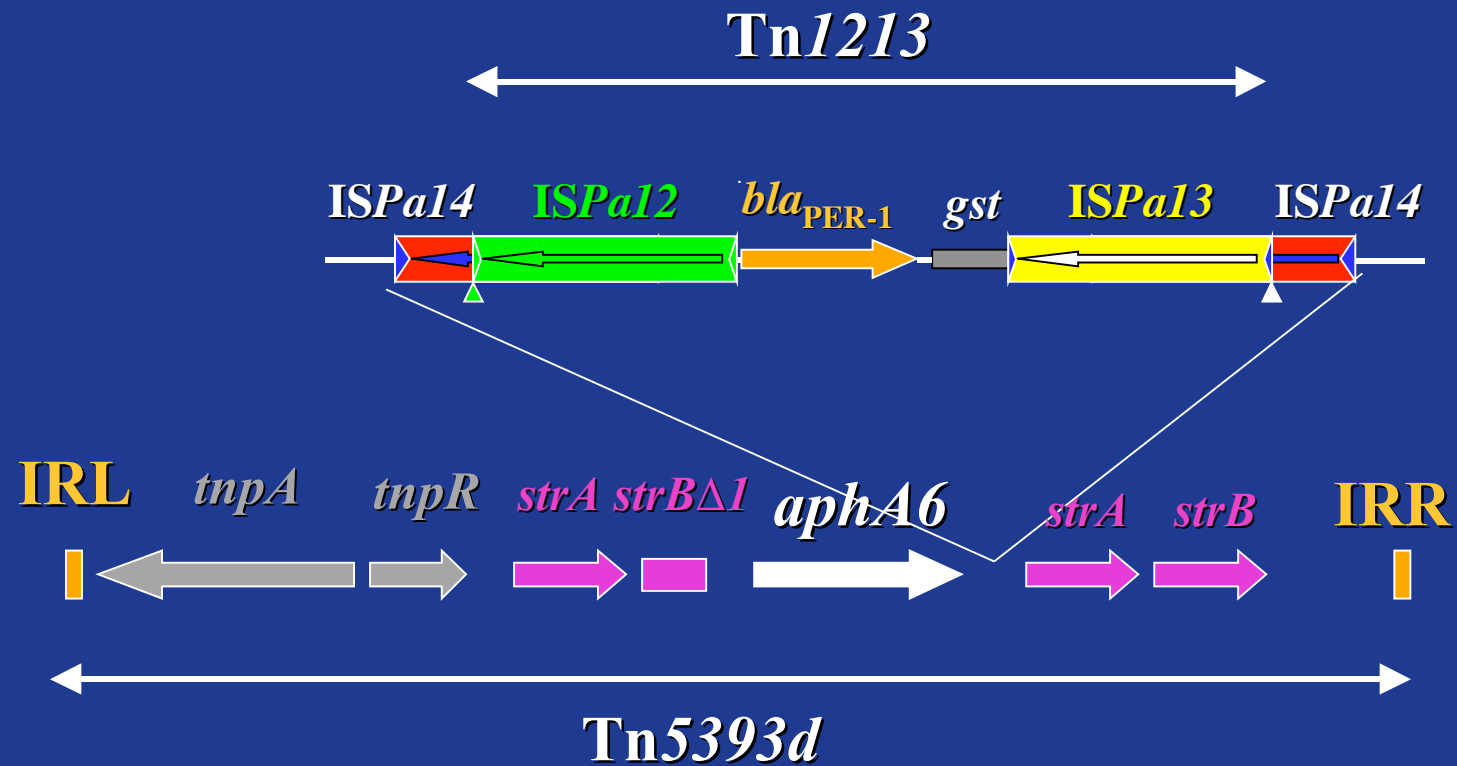
Genetic environment of *bla*_{PER-1} (1)



Genetic environment of *bla*_{PER-1} (2)



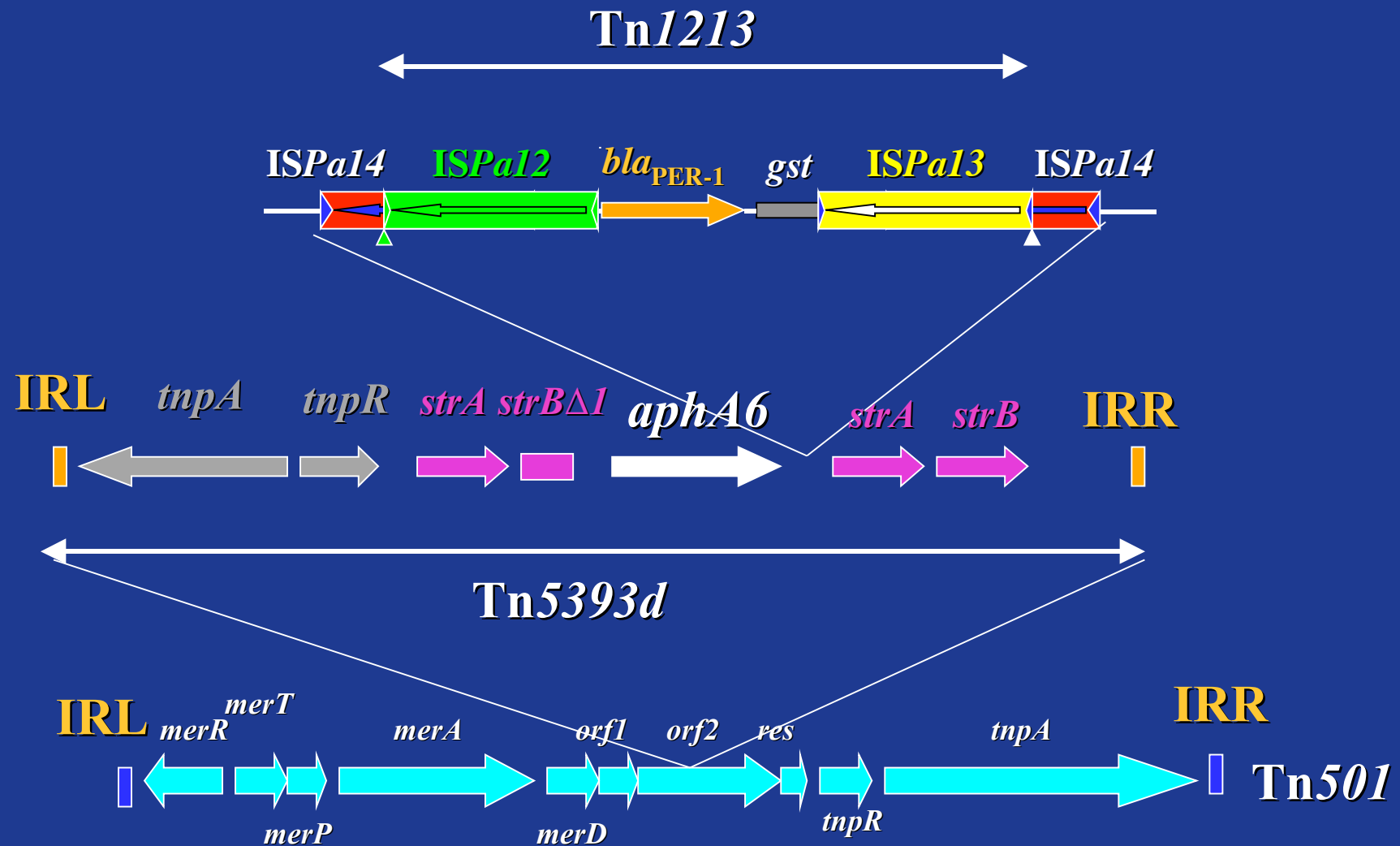
Genetic environment of *bla*_{PER-1} (2)



Alcaligenes faecalis, Italy

Mantengoli & Rossolini, AAC 2005

Genetic environment of *bla*_{PER-1} (2)

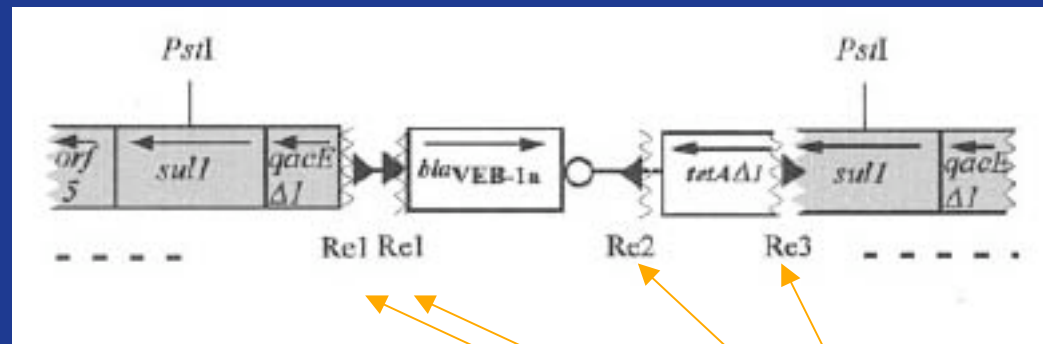


Alcaligenes faecalis, Italy

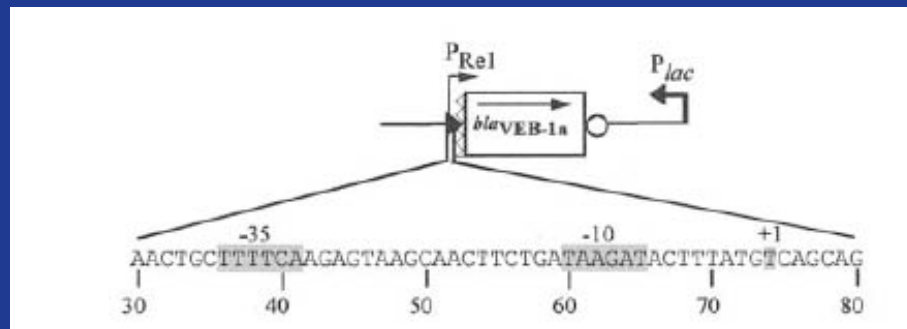
Mantengoli & Rossolini, AAC 2005

Other features

Re elements and *bla*_{VEB-1} acquisition

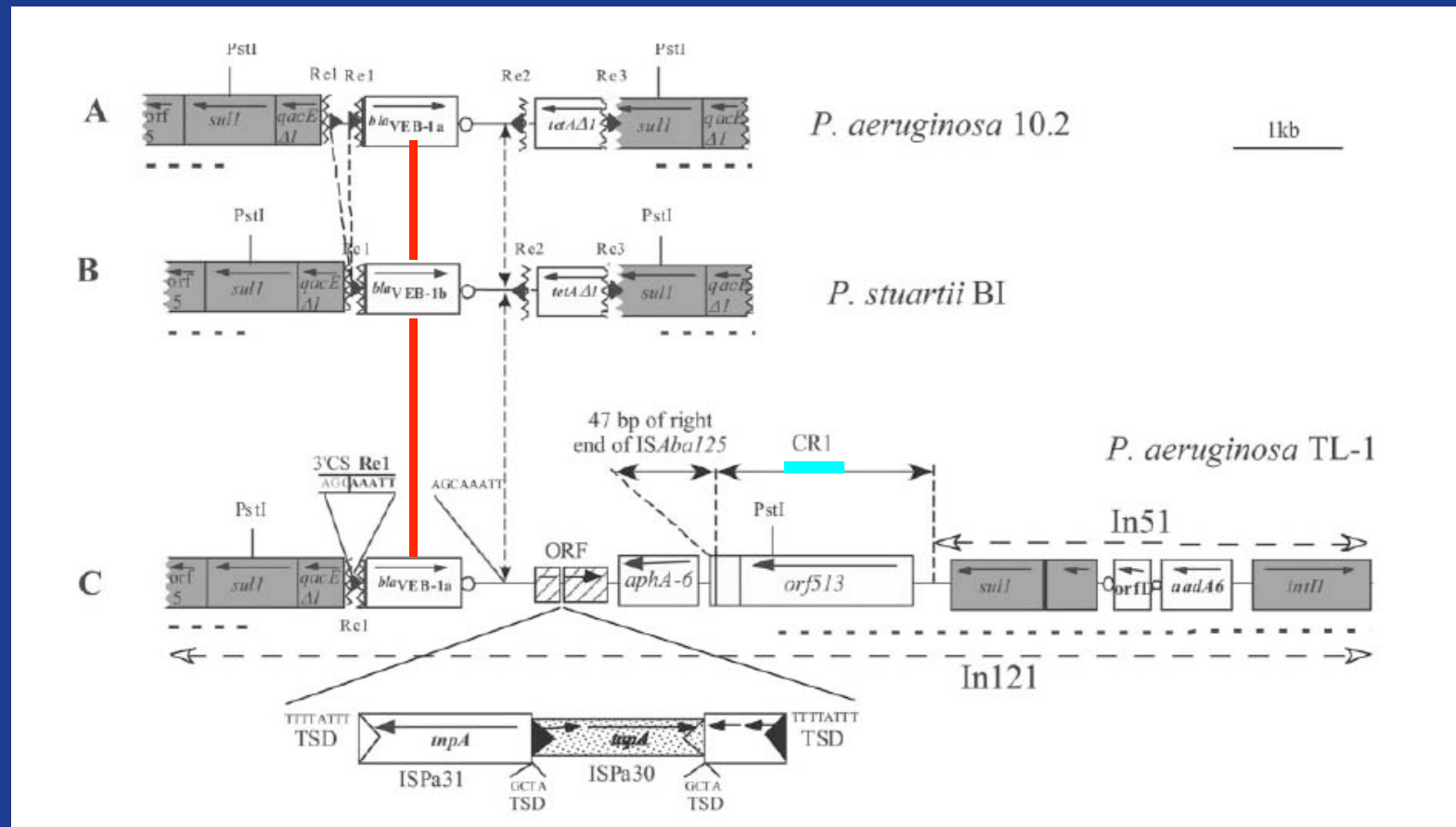


Re = 135 bp



Aubert et al., AAC 2004

***bla*_{VEB-1} may be part of a *sulI*-type integron**

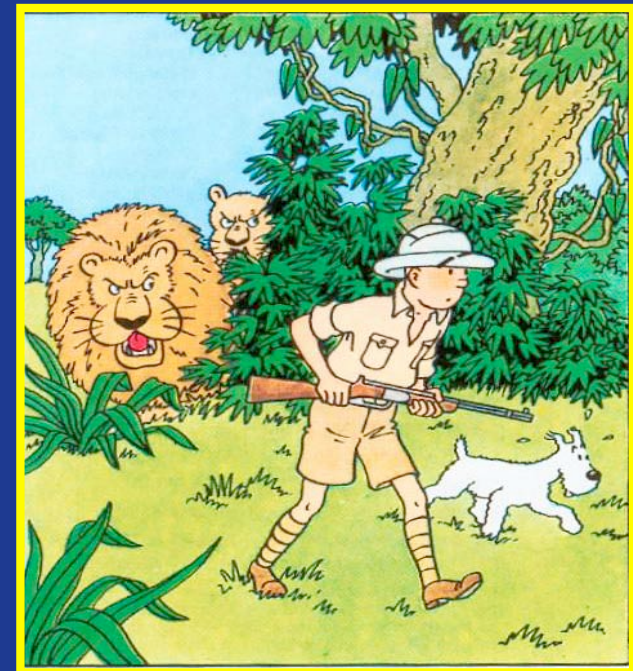


Genetic elements involved in mobilization of genes encoding class A β -lactamases

- Transposons for *bla*_{TEM}
- IS26 element for *bla*_{SHV}
- Class 1 integrons for *bla*_{VEB}, *bla*_{GES}
- CR1 element for some *bla*_{CTX-M} genes (*sul1*-type integrons)
- *ISEcp1* for other *bla*_{CTX-M} genes
- Peculiar composite transposons for *bla*_{PER-1}

Conclusions

- More and more vehicles for ESBL genes acquisition and dissemination are identified
- Genetic plasticity at the origin of ESBL gene diffusion => increasing spread
- Searching for the reservoirs
 - Mobile elements
 - Resistance genes by themselves



Acknowledgments

P. Nordmann

T. Naas