

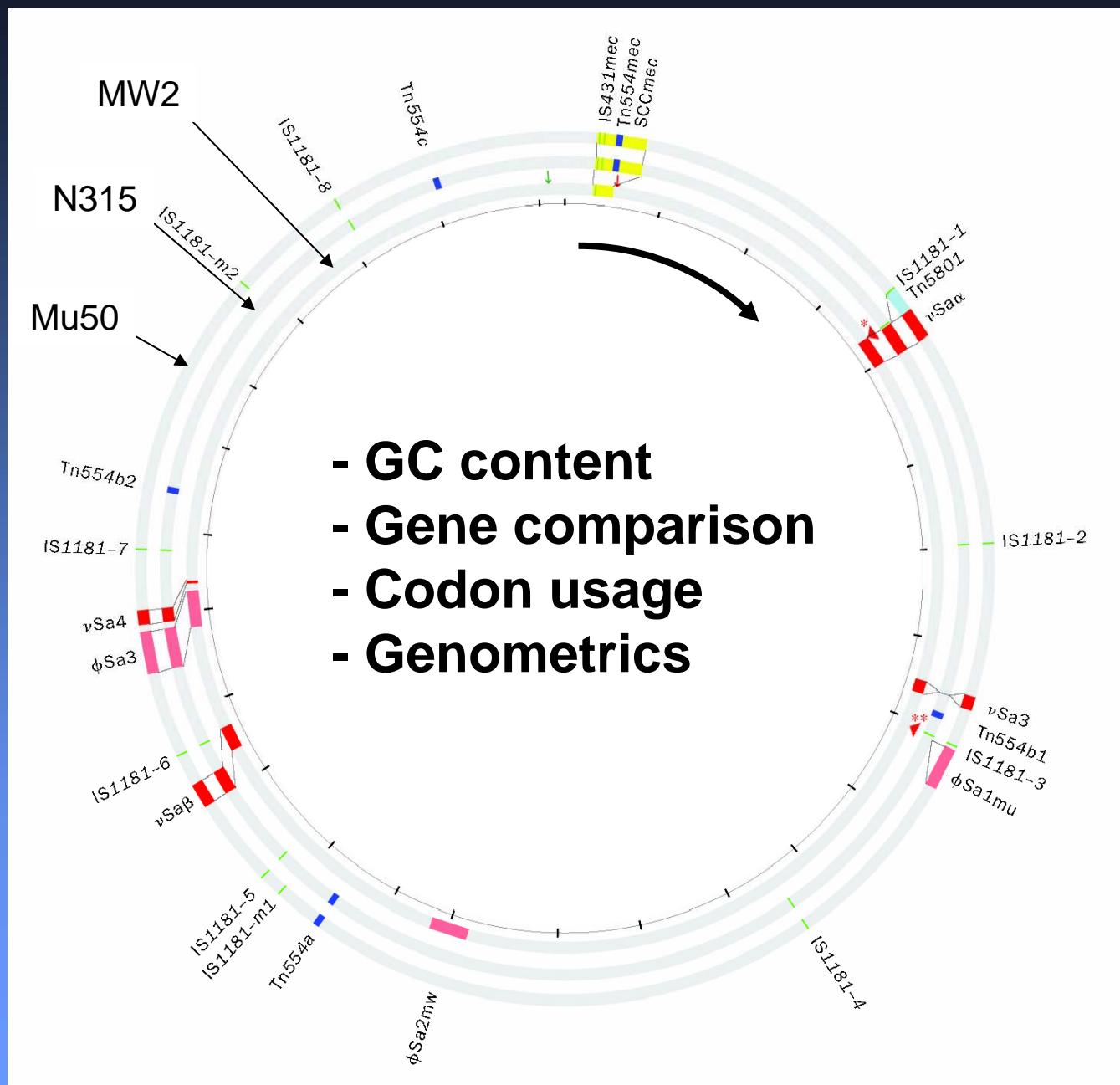
Staphylococcus aureus Pathogenesis

- Gene exchanges
- Gene regulation
- Gene products
- Gene product cooperation

Some Staphylococcal Species from Mammals

Host	Species	Coagulase	Clumping factor	Virulence
Human and other primates	<i>S. aureus</i>	++	++	+++
	<i>S. epidermidis</i>	-	-	+
	<i>S. capitis</i>	-	-	(+)
	<i>S. caprae</i>	-	-	(+)
	<i>S. saccharolyticus</i>	±	-	-
	<i>S. warneri</i>	-	-	-
	<i>S. pasteurii</i>	-	-	-
	<i>S. haemolyticus</i>	-	-	+
	<i>S. hominis</i>	-	-	(+)
	<i>S. lugdunensis</i>	-	+	+
	<i>S. auricularis</i>	-	-	(+)
	<i>S. saprophyticus</i>	-	-	+
	<i>S. cohnii</i>	-	-	-
	<i>S. xilosus</i>	-	-	-
	<i>S. simulans</i>	-	-	-
	<i>S. schleiferi</i>	±	+	+
Carnivores	<i>S. intermedius</i>	+	-	++
	<i>S. felis</i>	-	-	++

Whole genomes of *S. aureus* MW2, N315 and Mu50



From Linguistics to Genometrics

theicaacisagreatconférenceparcequellegathermanygreatscientists

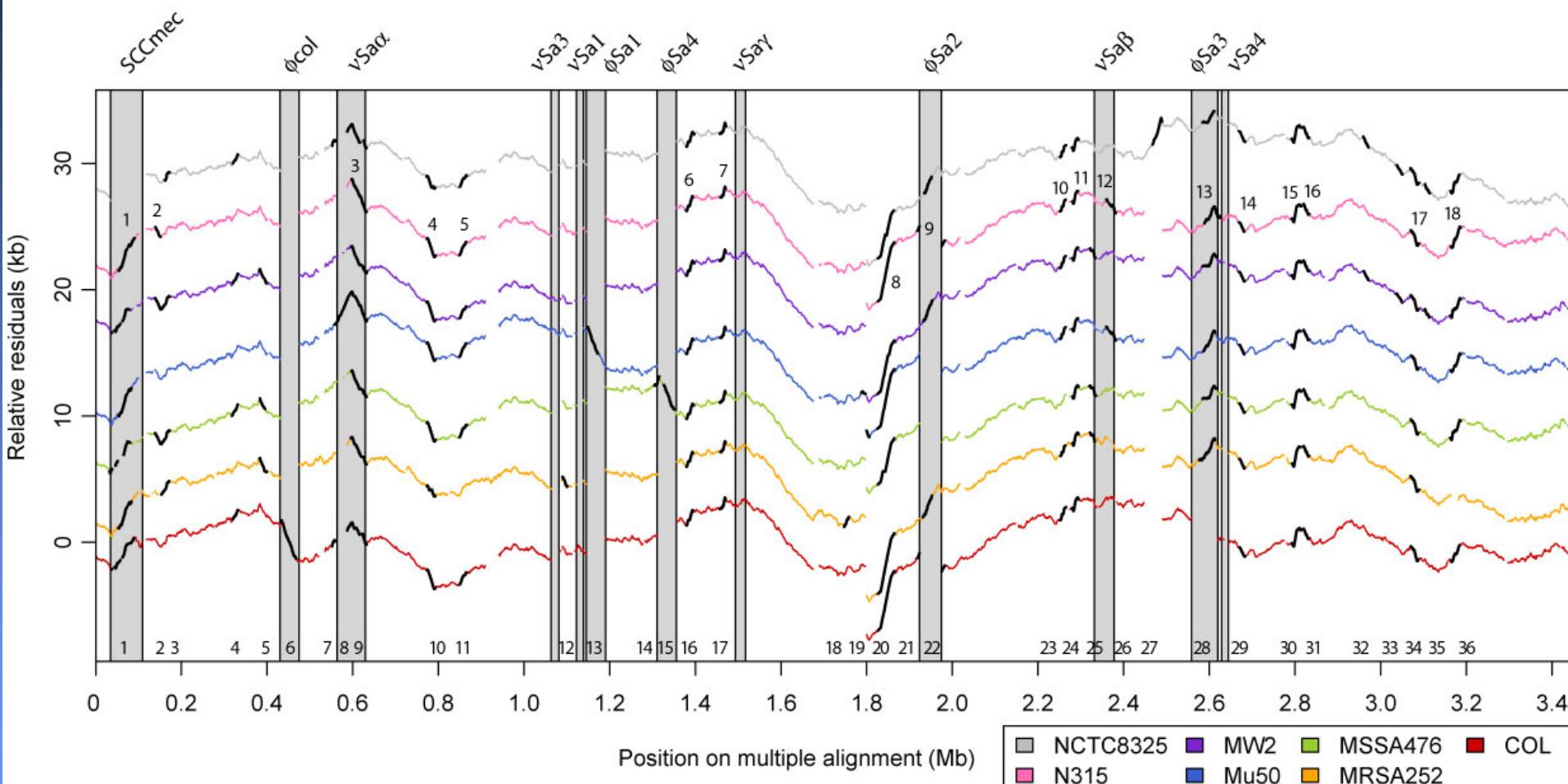
theicaacisagreatconférenceparcequellegathermanygreatscientists



<http://www2.unil.ch/comparativegenometrics/>

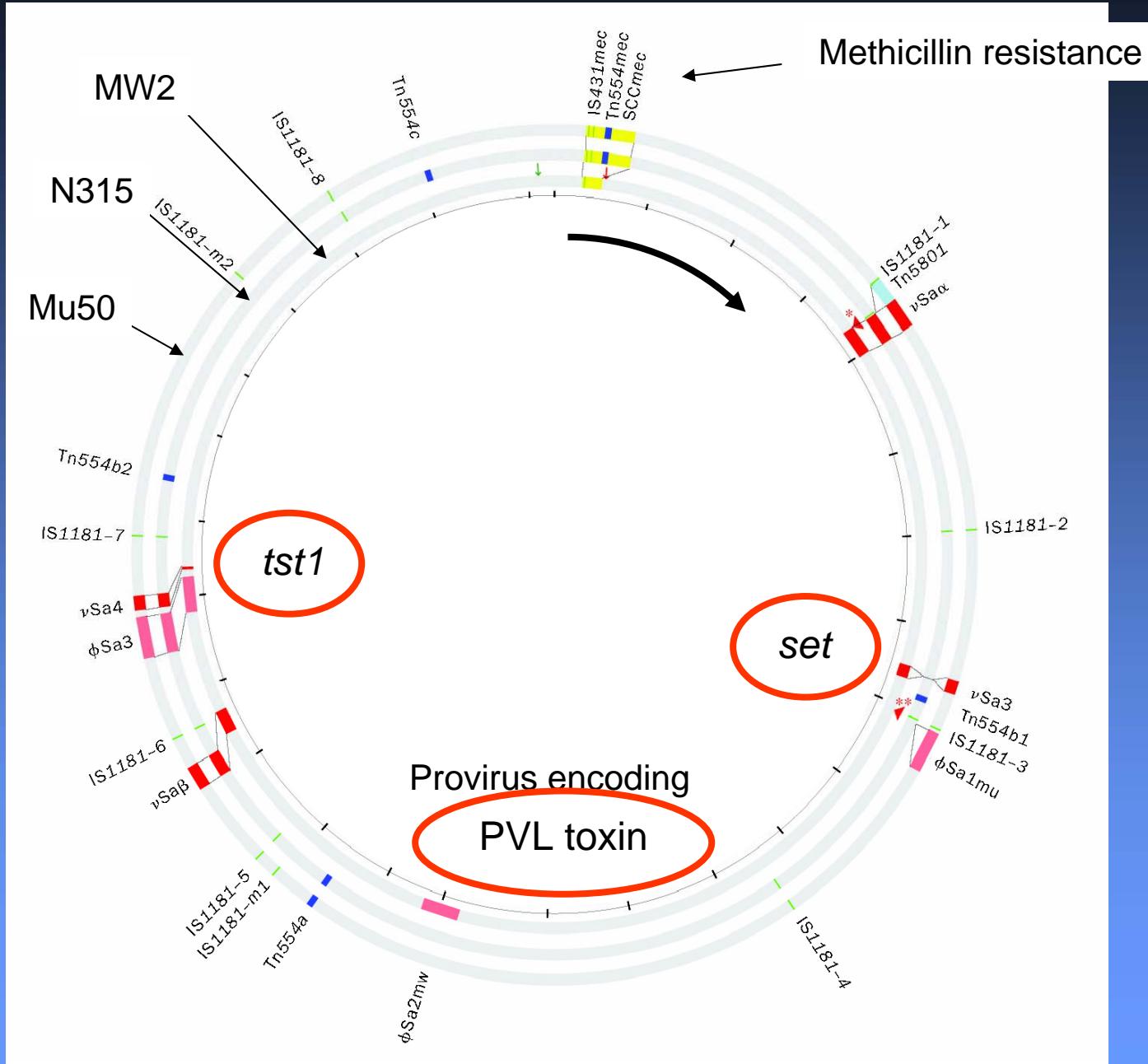
CA Roten and L. Guy

S. aureus Genometric Analysis



Guy Lionel, Thesis 2007

Whole genomes of *S. aureus* MW2, N315 and Mu50



Define Pathogenicity Islands

- DNA region (up to 200 kb) of foreign origin
- Contain pathogenic genes
- May have a different GC content
- Flanked by direct repeats of IS
- Associated with tRNA genes
- Contain relics of integrase/excisase genes or of phage genes

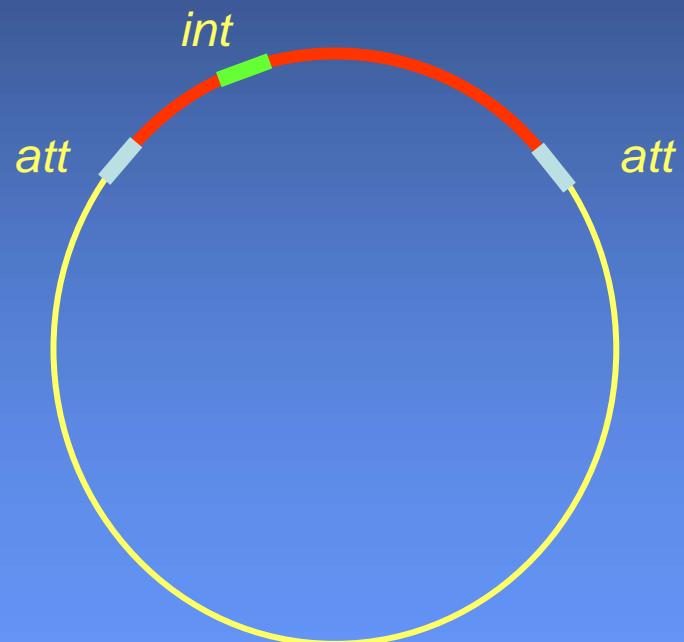
Hacker J et al. *Mol Microbiol* 1997; 23:1089-97

How do they move ?

- Transformation
- Conjugation
- Transduction (generalized)

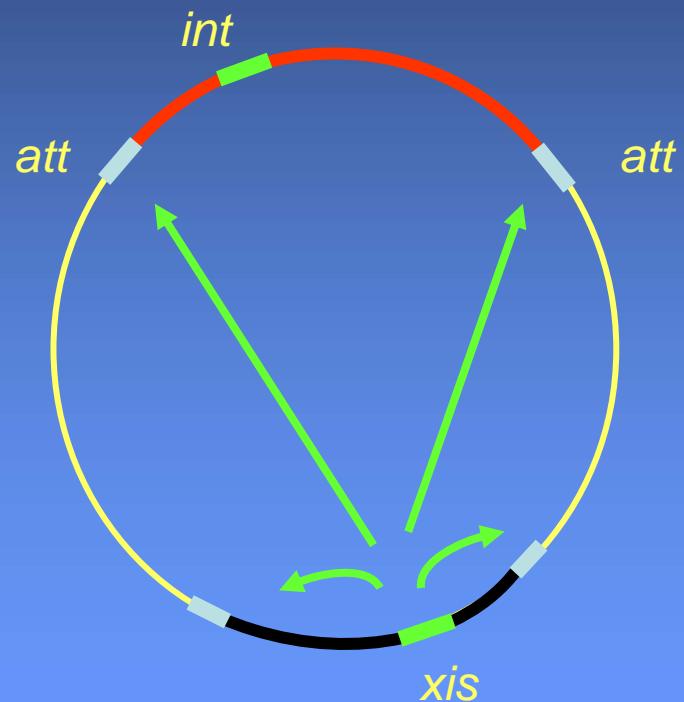
Moving Genes Around

The example of SaPI1



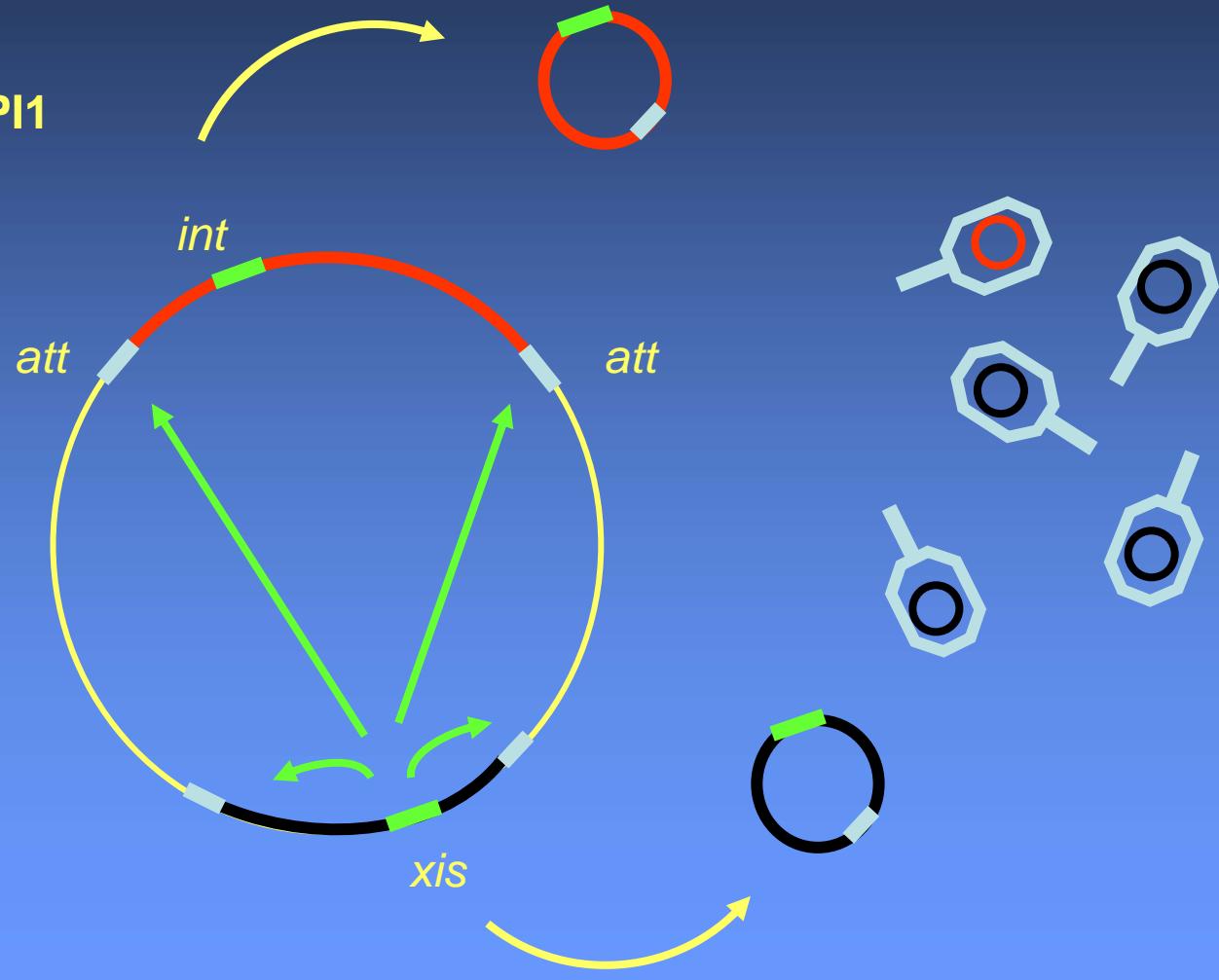
Moving Genes Around

The example of SaPI1



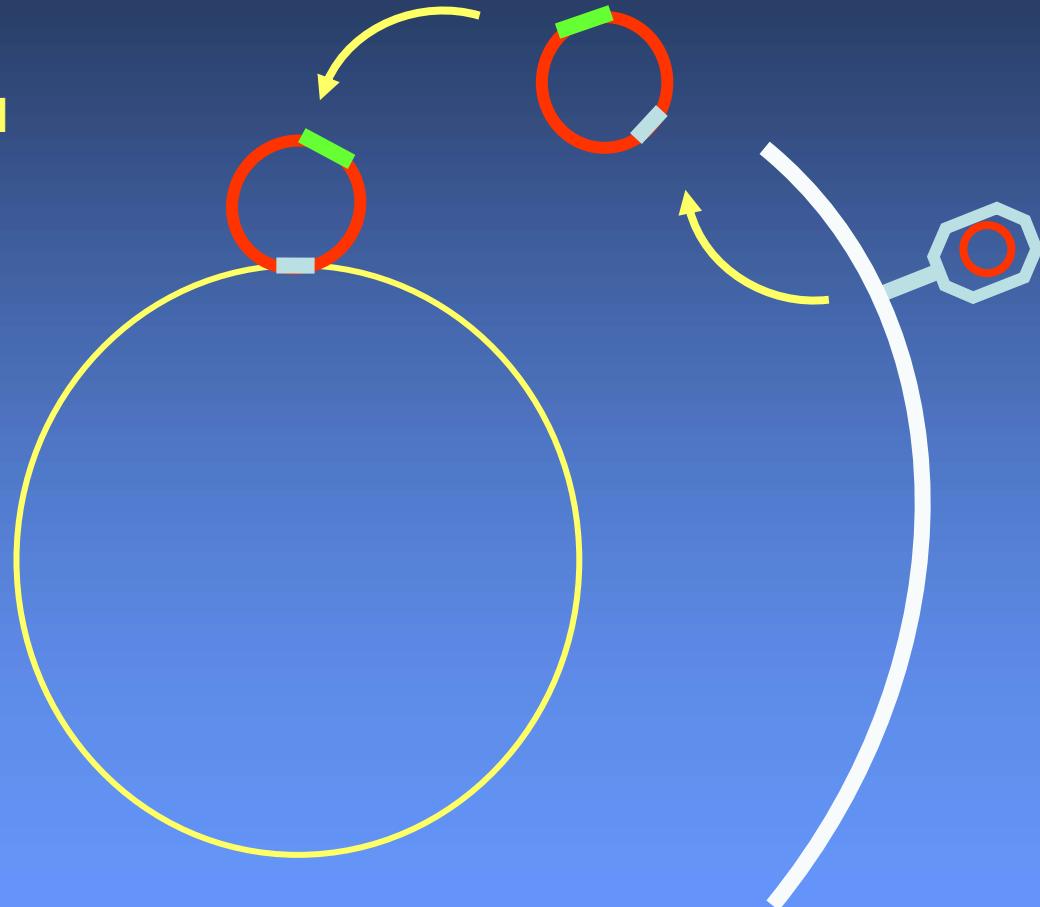
Moving Genes Around

The example of SaPI1



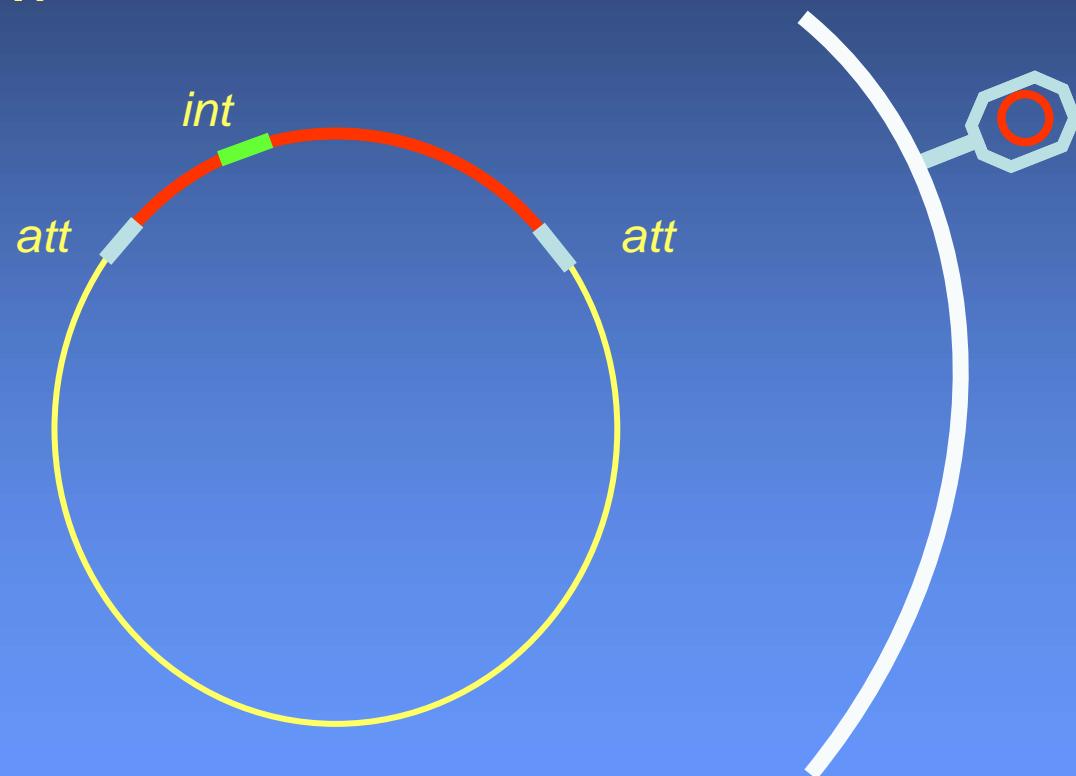
Moving Genes Around

The example of SaPI1

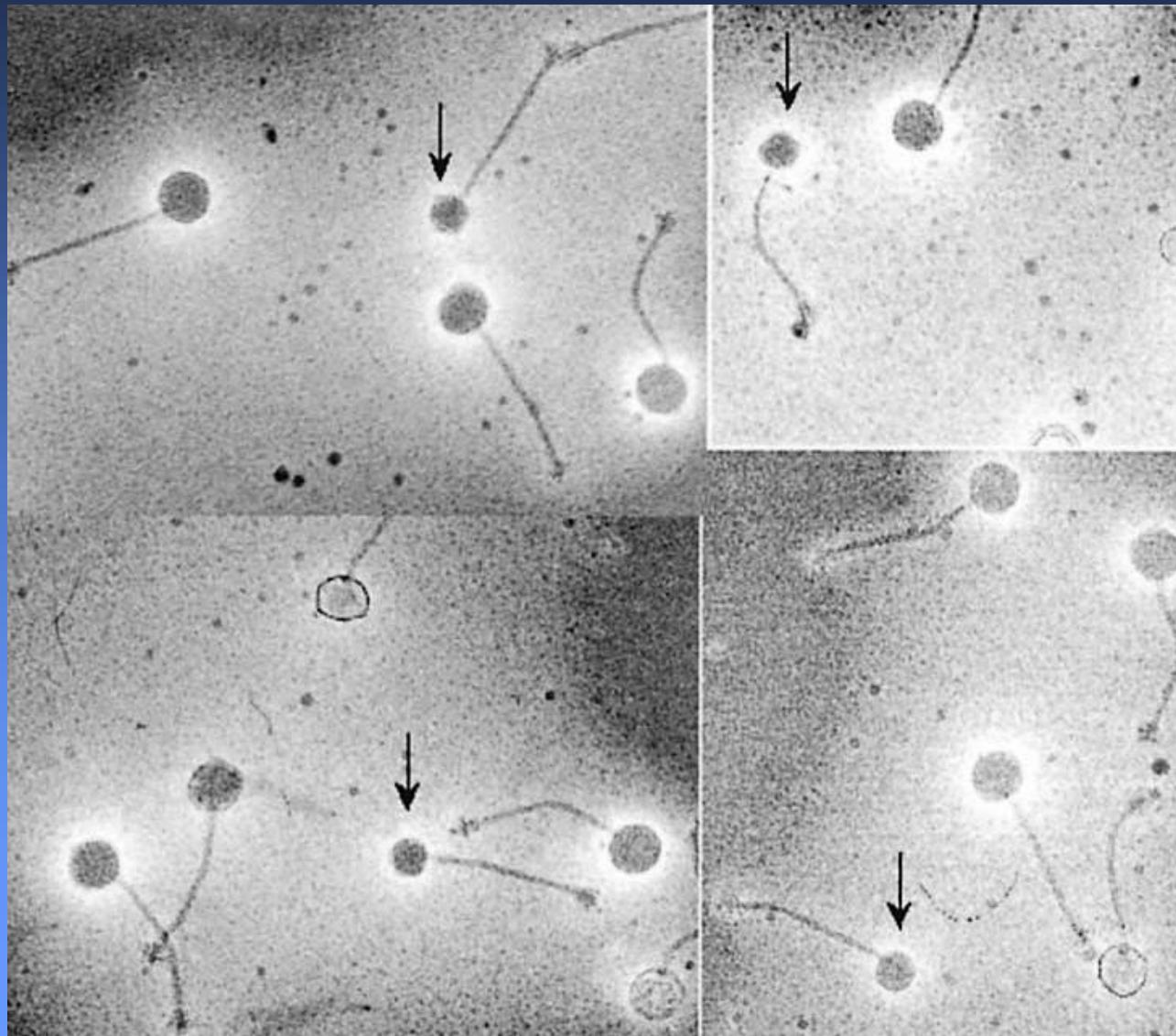


Moving Genes Around

The example of SaPI1



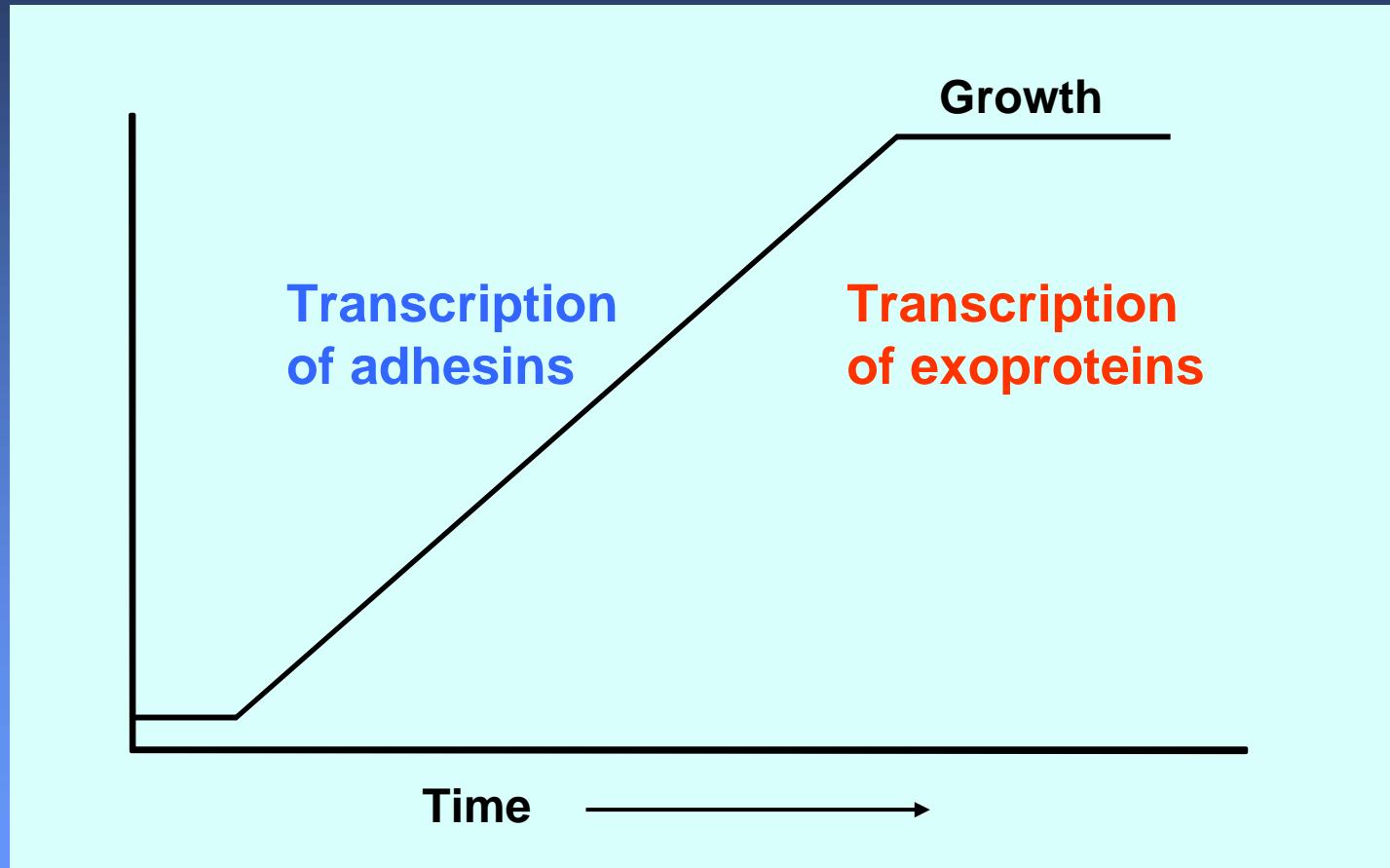
Moving SaPI1 with Phage 80 α



Staphylococcus aureus Pathogenesis

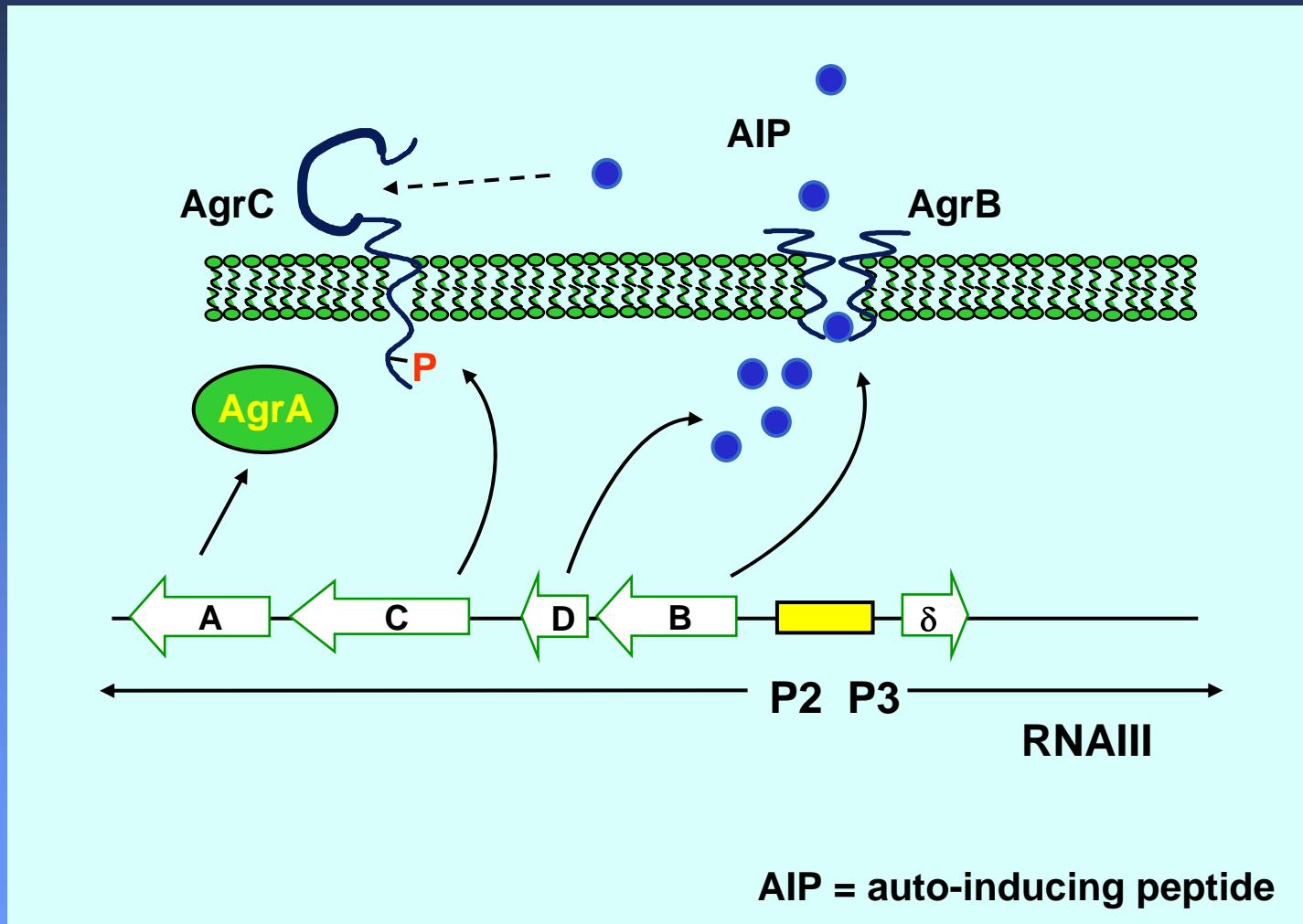
- Gene exchanges
- Gene regulation
- Gene products
- Gene product cooperation

agr-Regulation of Virulence Determinants During in vitro Growth of S. aureus (1)



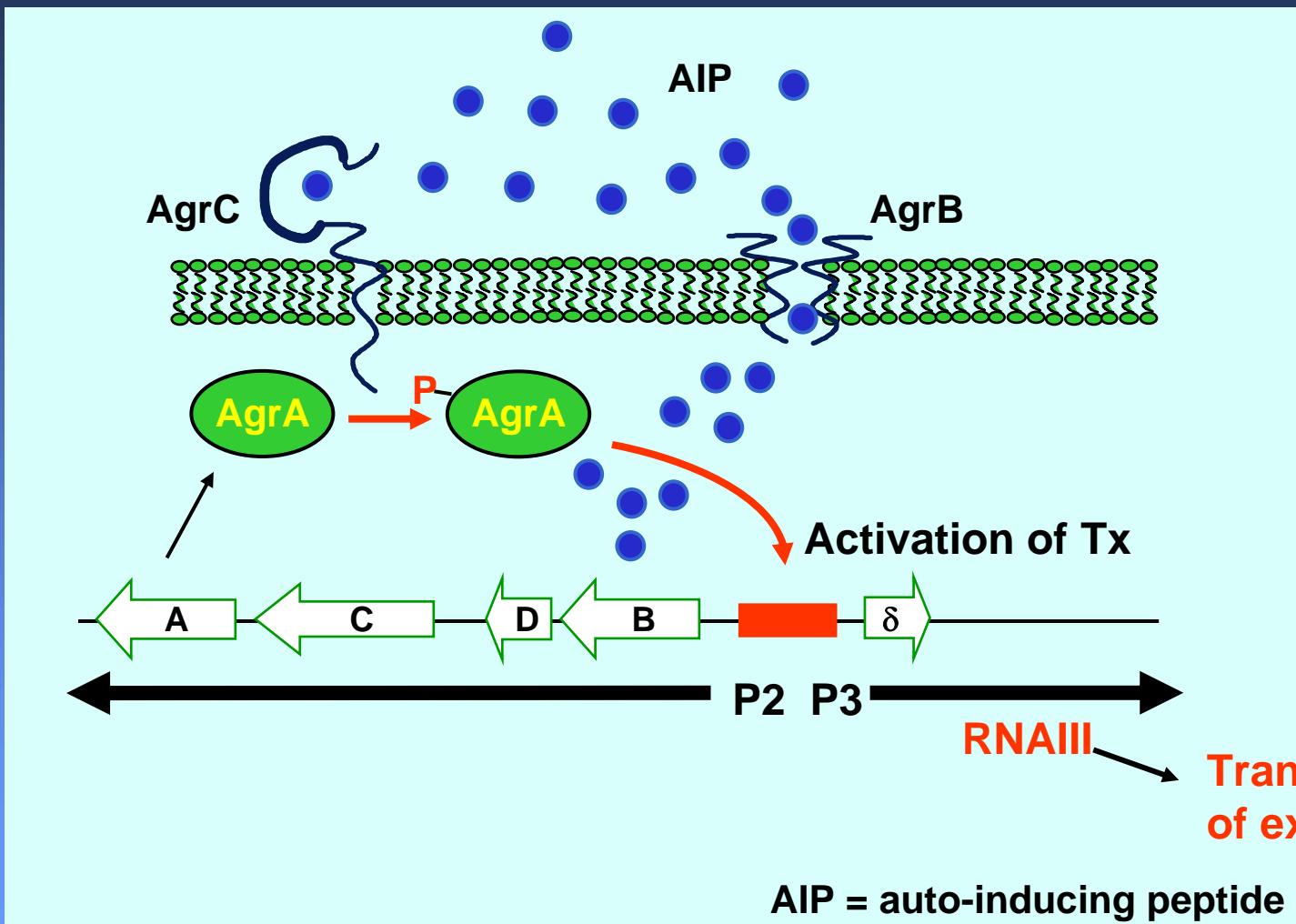
Adapted from R.P. Novick ASM press, 2000

Virulence Gene Regulation by Peptides in *S. aureus* (1)



Novick et al. *Curr. Op. Microbiol.* 1999;2:40-45

Virulence Gene Regulation by Peptides in *S. aureus* (2)



Novick et al. *Curr. Op. Microbiol.* 1999;2:40-45

Regulation of *S. aureus* super-antigens

Gene	Product	Activity/function	Action of regulatory genes						
			<i>agr</i>	<i>Sae-RS</i>	<i>rot</i>	<i>sarA</i>	<i>sarS</i>	<i>sarT</i>	<i>tst</i>
Superantigens									
<i>sea</i>	Enterotoxin A	Food poisoning, TSS	0						0
<i>seb</i>	Enterotoxin B	Food poisoning, TSS	+			b			-
<i>sec</i>	Enterotoxin C	Food poisoning, TSS	+						
<i>sed</i>	Enterotoxin D	Food poisoning, TSS	+						
<i>eta</i>	Exfoliatin A	Scalded skin syndrome	+						
<i>etb</i>	Exfoliatin B	Scalded skin syndrome	+						
<i>tst</i>	Toxic shock toxin-1	Toxic shock syndrome	+	-		b			-

0, no effect of gene on expression; +, upregulated; -, downregulated.

b. Controversial.

Conditions for Developing Toxic Shock Syndrome

Host:

- Lack of anti-TSST antibodies

Environment:

- High protein concentration
- Neutral pH
- High pCO₂
- High pO₂

Altered
Ecology !

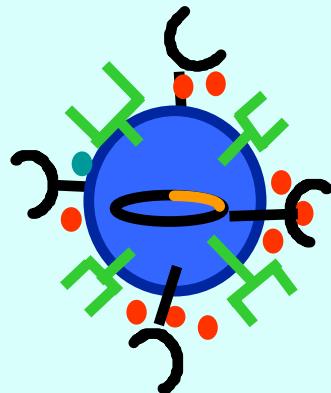
Staphylococcus aureus Pathogenesis

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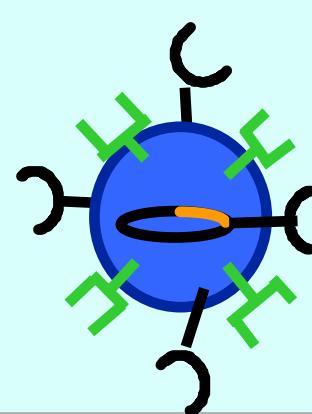
Gene	Protein	Function	Implication in disease
<i>spa</i>	Protein A	Binds Fc fragment	Sepsis, osteoarthritis
<i>cifA</i>	Clumping factor A	Binding to fibrinogen	Endocarditis
<i>cifB</i>	Clumping factor B	Binding to fibrinogen	-
<i>cna</i>	Collagen binding protein	Binding to collagen	Osteomyelitis
<i>fnA</i>	Fn-binding protein A	Binding to fibronectin	Endocarditis, persistence
<i>fnB</i>	Fn-binding protein B	Binding to fibronectin	-
<i>sdrC</i>	Serine-aspartate repeat prot	Binding to fibrinogen	-
<i>sdrD</i>	Serine-aspartate repeat prot	Binding to fibrinogen ?	-
<i>sdrE</i>	Serine-aspartate repeat prot	Binding to fibrinogen ?	-
<i>pls</i>	Plasmin-sensitive protein	Binding to nasal mucosa	Nasal colonization?
<i>fmtB</i>	Methicillin resistance	Putative cell-wall building	Expression of methi-R
<i>sasA</i>	<i>S. aureus</i> surface protein A	Undetermined	-
<i>sasB</i>	<i>S. aureus</i> surface protein B	Undetermined	-
<i>sasC</i>	<i>S. aureus</i> surface protein C	Undetermined	-
<i>sasE</i>	<i>S. aureus</i> surface protein E	Undetermined	-
<i>sasF</i>	<i>S. aureus</i> surface protein F	Undetermined	-
<i>sasG</i>	<i>S. aureus</i> surface protein G	Binding to nasal mucosa	Invasive disease
<i>sasH</i>	<i>S. aureus</i> surface protein H	Undetermined	Invasive disease
<i>sasI</i>	<i>S. aureus</i> surface protein I	Undetermined	-
<i>sasJ</i>	<i>S. aureus</i> surface protein J	Undetermined	-
<i>sasK</i>	<i>S. aureus</i> surface protein K	Undetermined	-

Understanding Pathogenic Factors

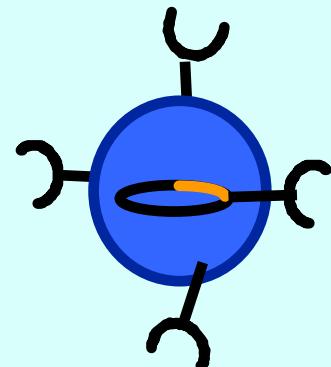
Wild Type



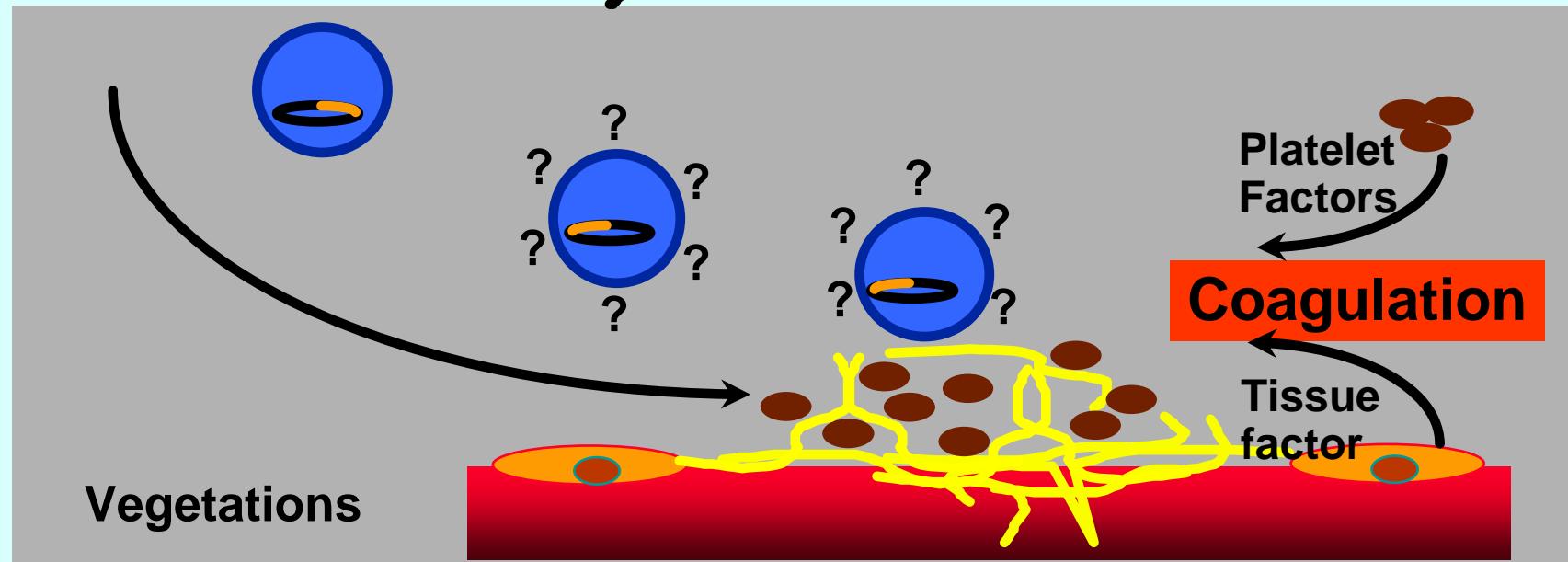
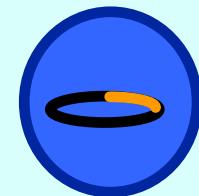
Coa⁻



Coa⁻/ClfA⁻



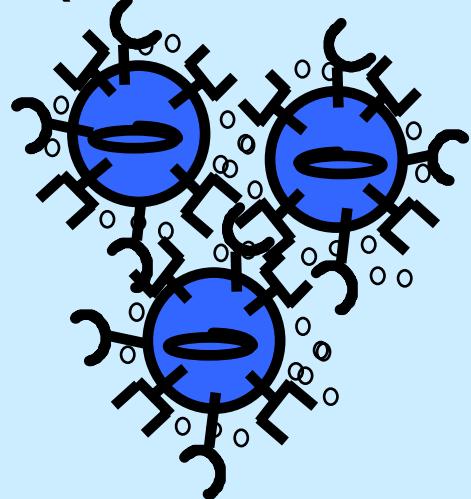
Coa⁻/ClfA⁻/Fnb⁻



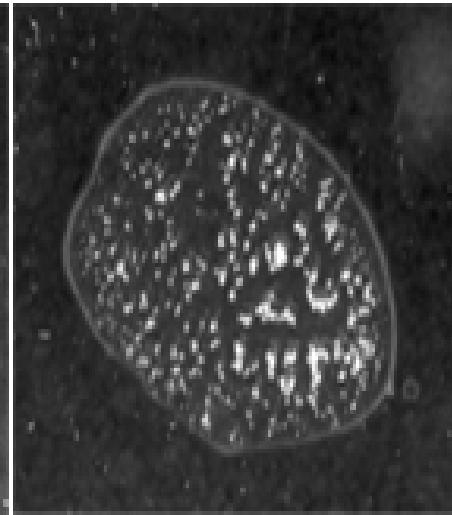
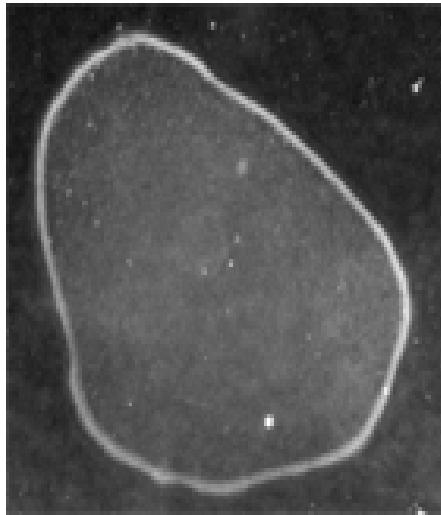
Transfer or Staple genes into *lentil*

nic

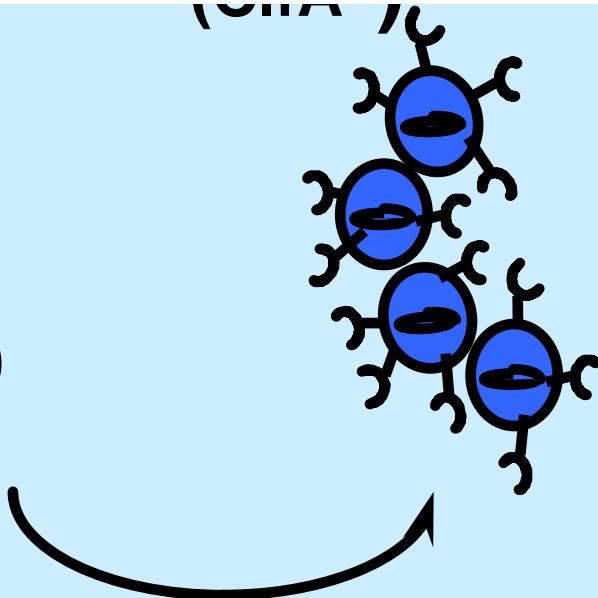
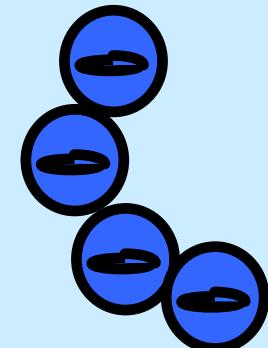
S. aureus donor
(Coa⁺/ClfA⁺/Fnb⁺)



Lac⁺
re

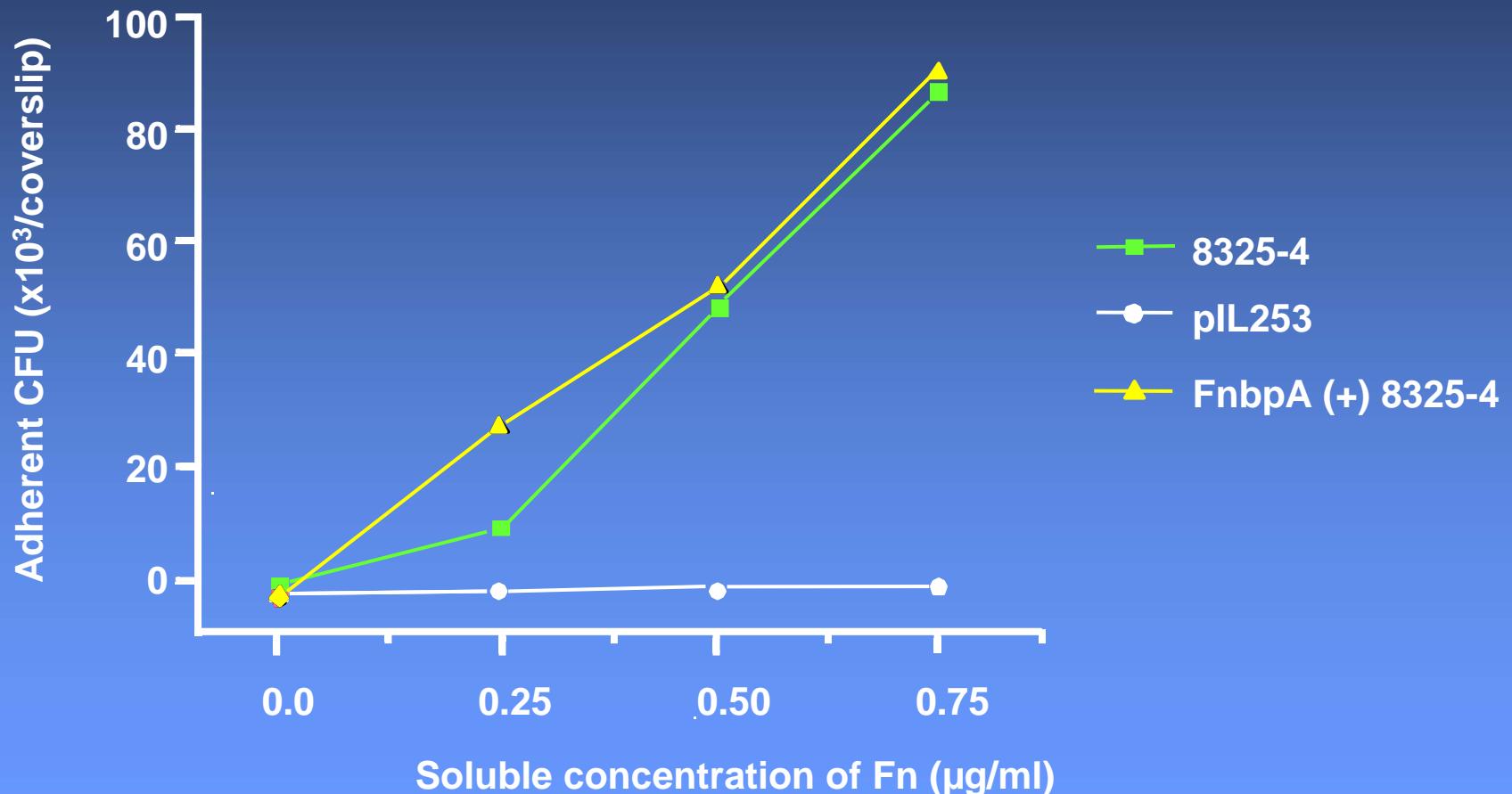


1: Transfer of *S.aureus clfA* gene
into viridans group streptococcus

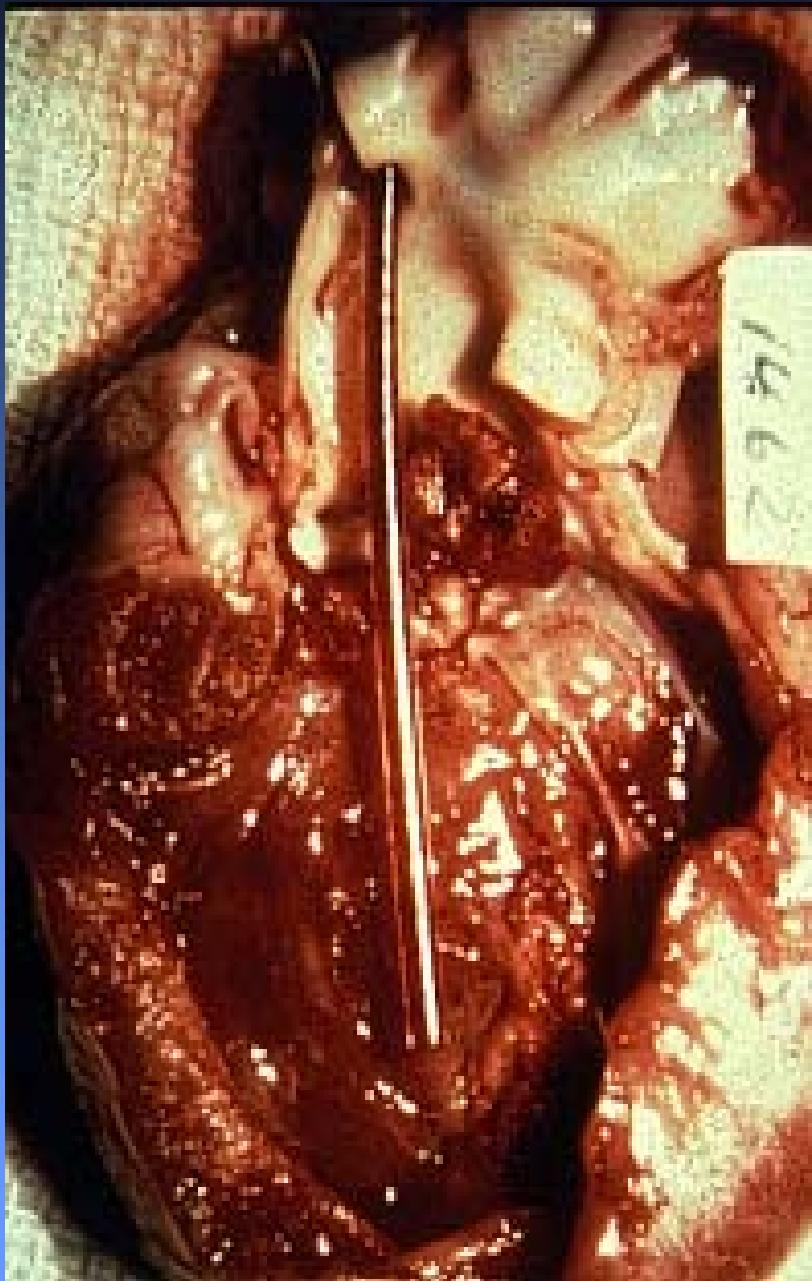


2: Select clumping-positive
recombinants

Adherence of *L. lactis* to PMMA
coverslips coated with fibronectin

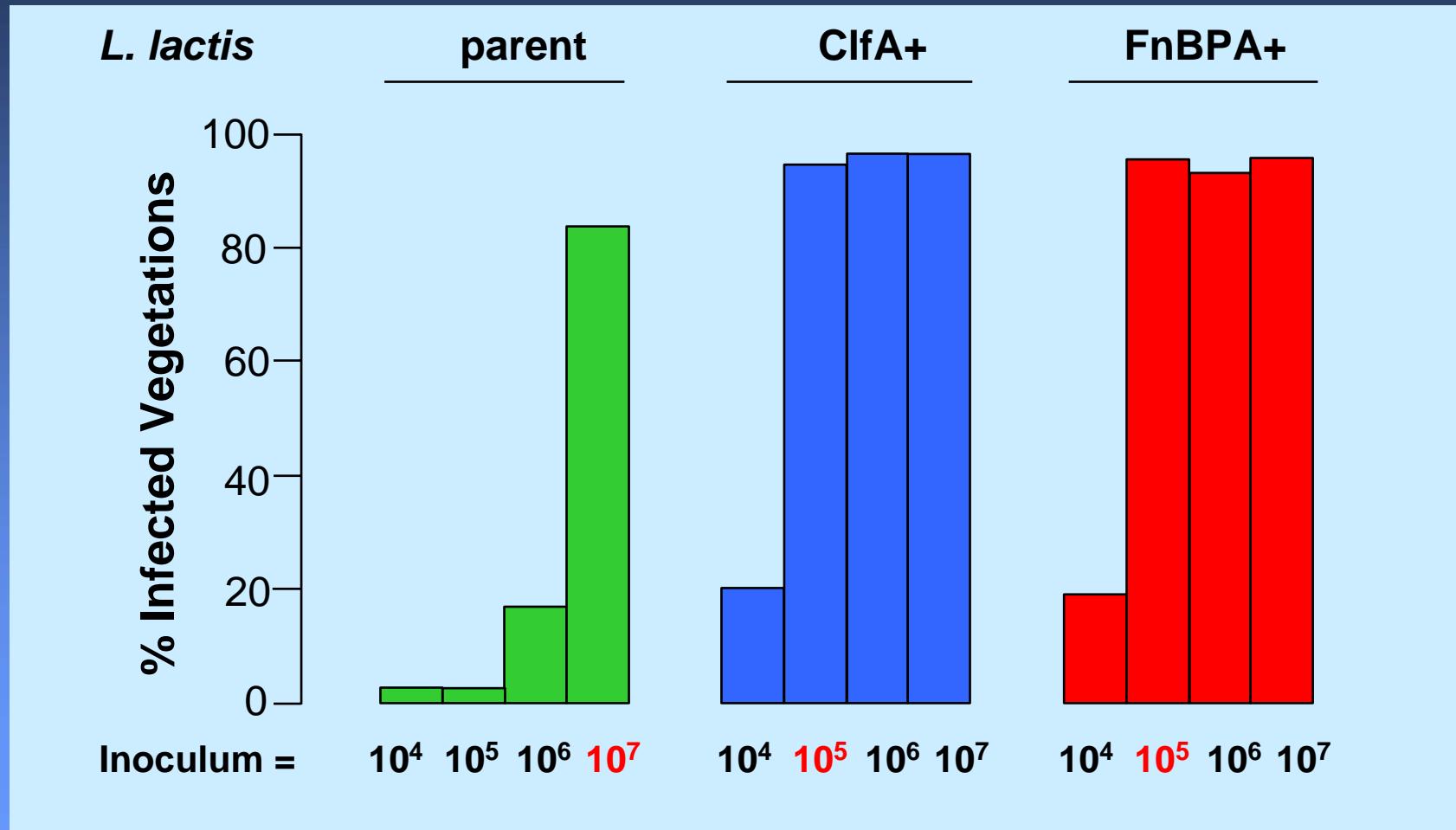


Experimental Endocarditis



Infectivity of *L. lactis* Recombinants

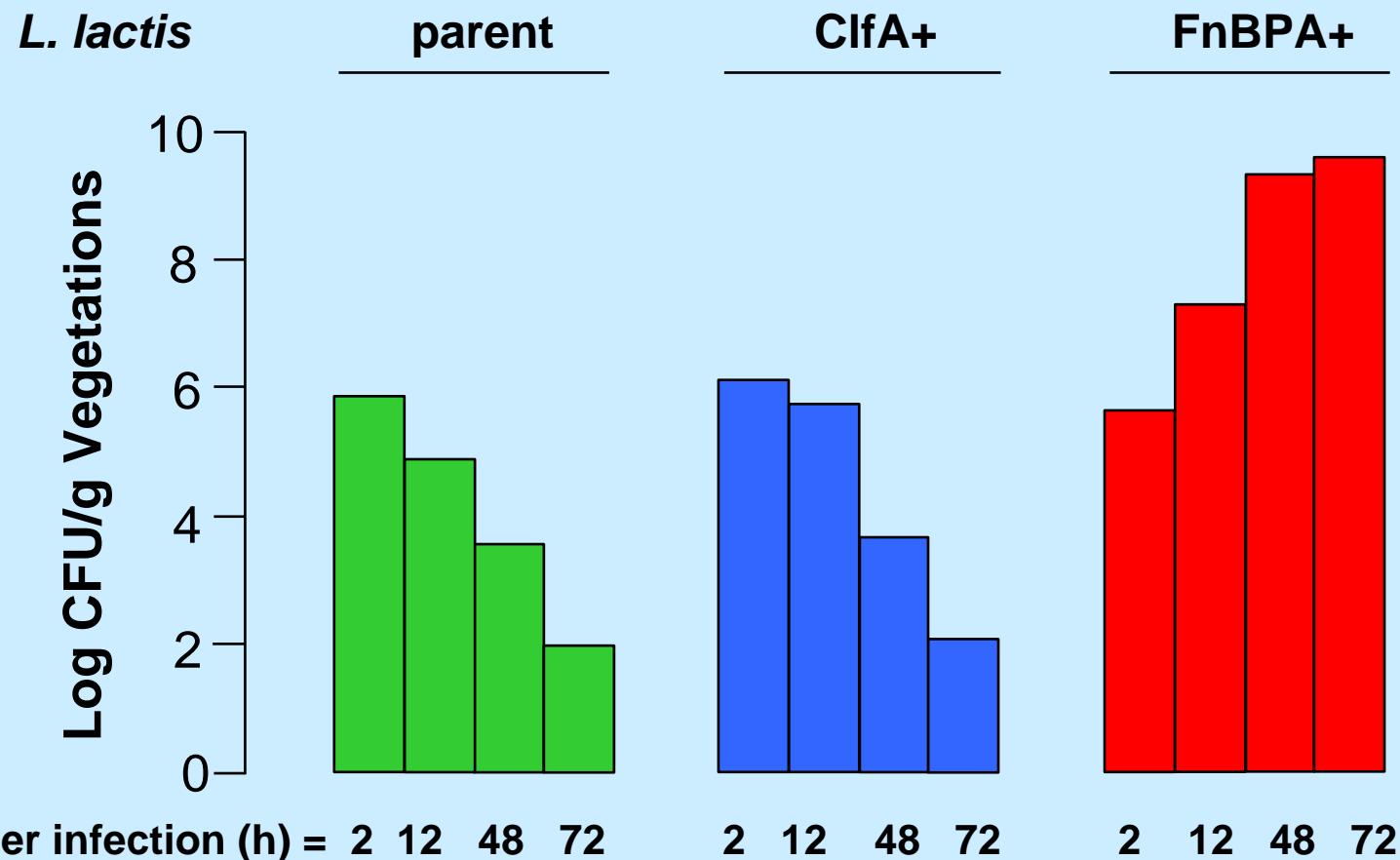
Binding to Fibrinogen or Fibronectin



Que et al. Infect. Immun. 2001

Brussels 07

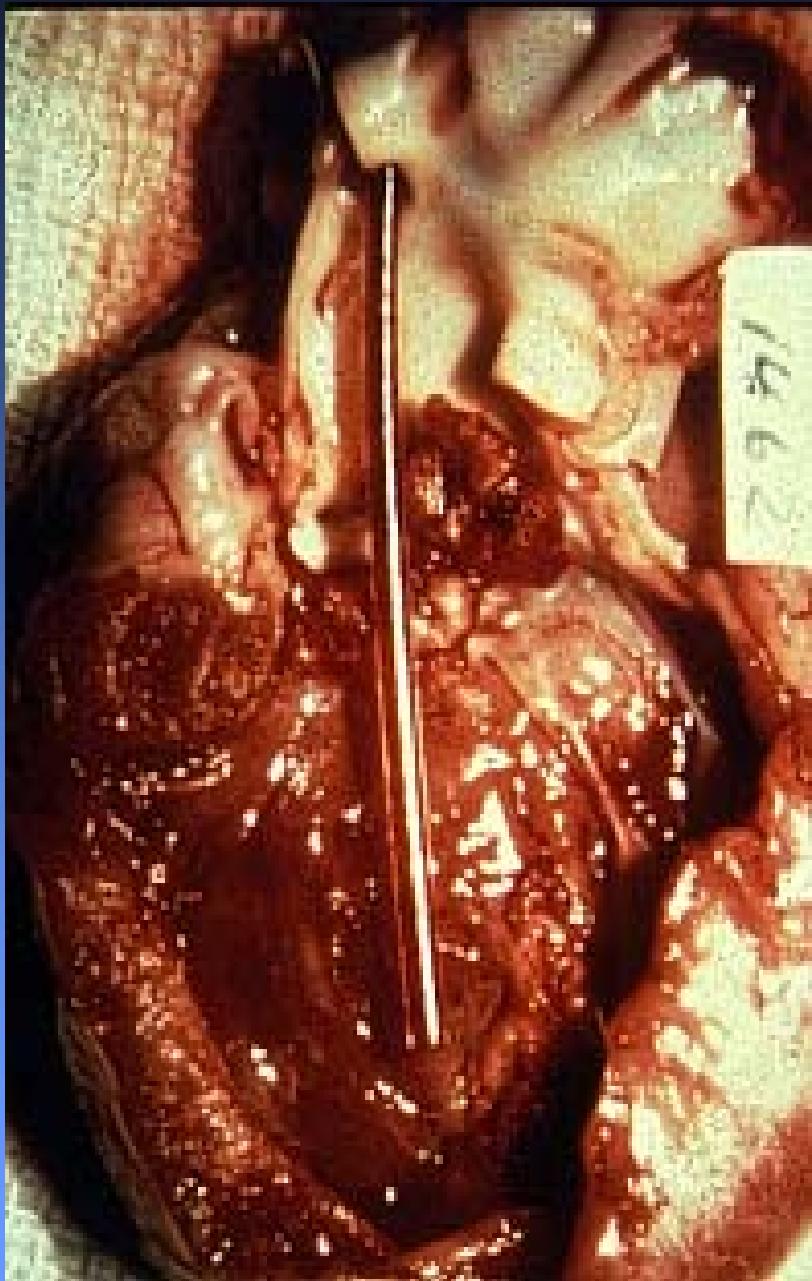
Dynamic of *L. lactis* Endocarditis



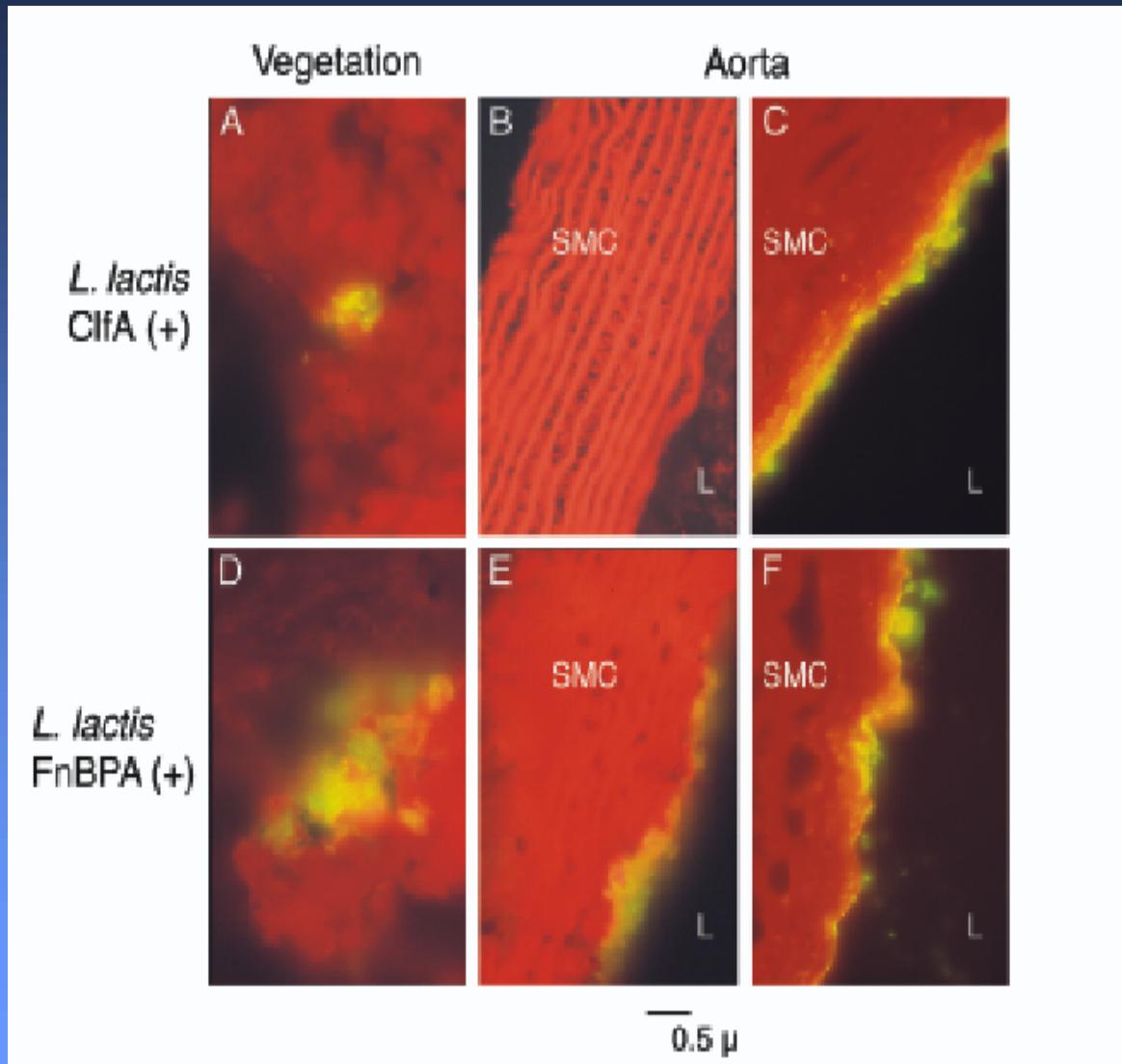
Que et al. J. Exp. Med., 2005

Brussels 07

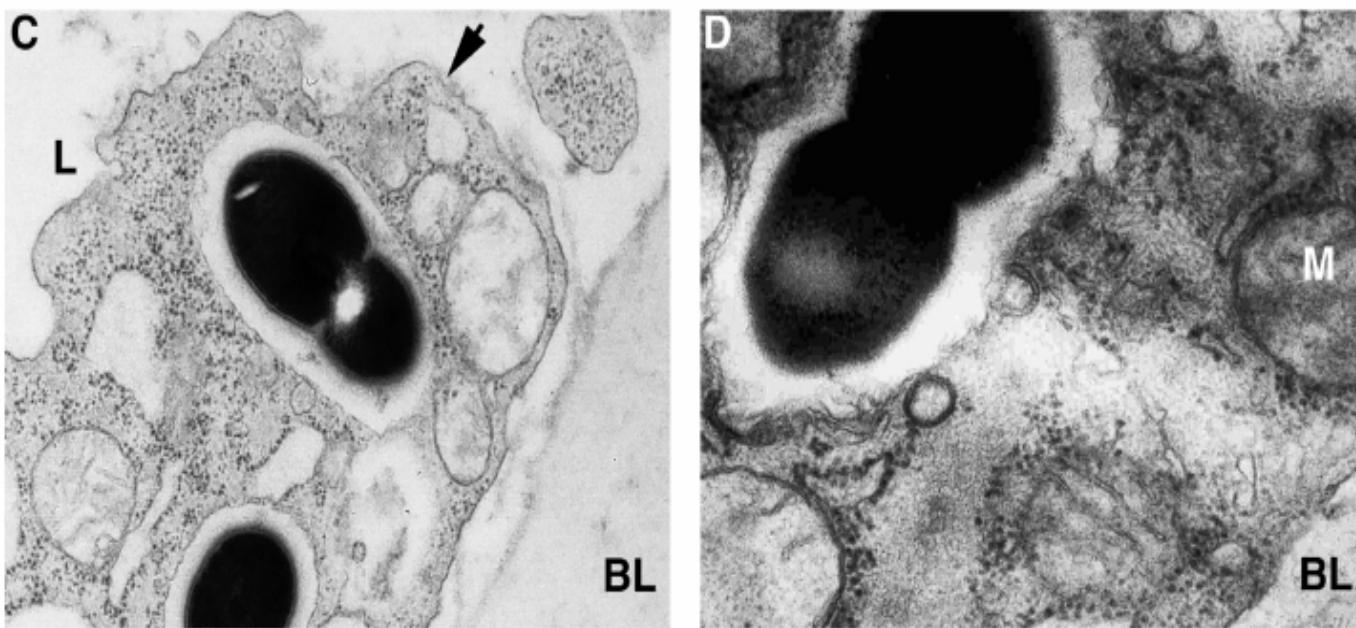
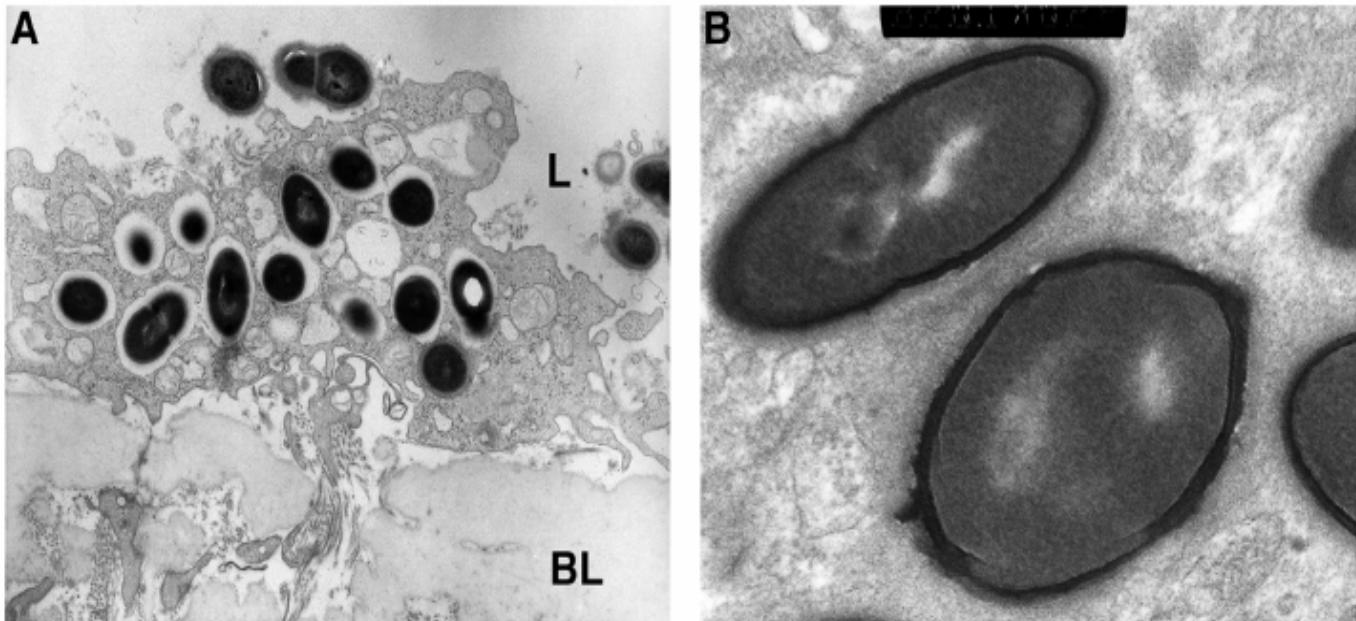
Experimental Endocarditis



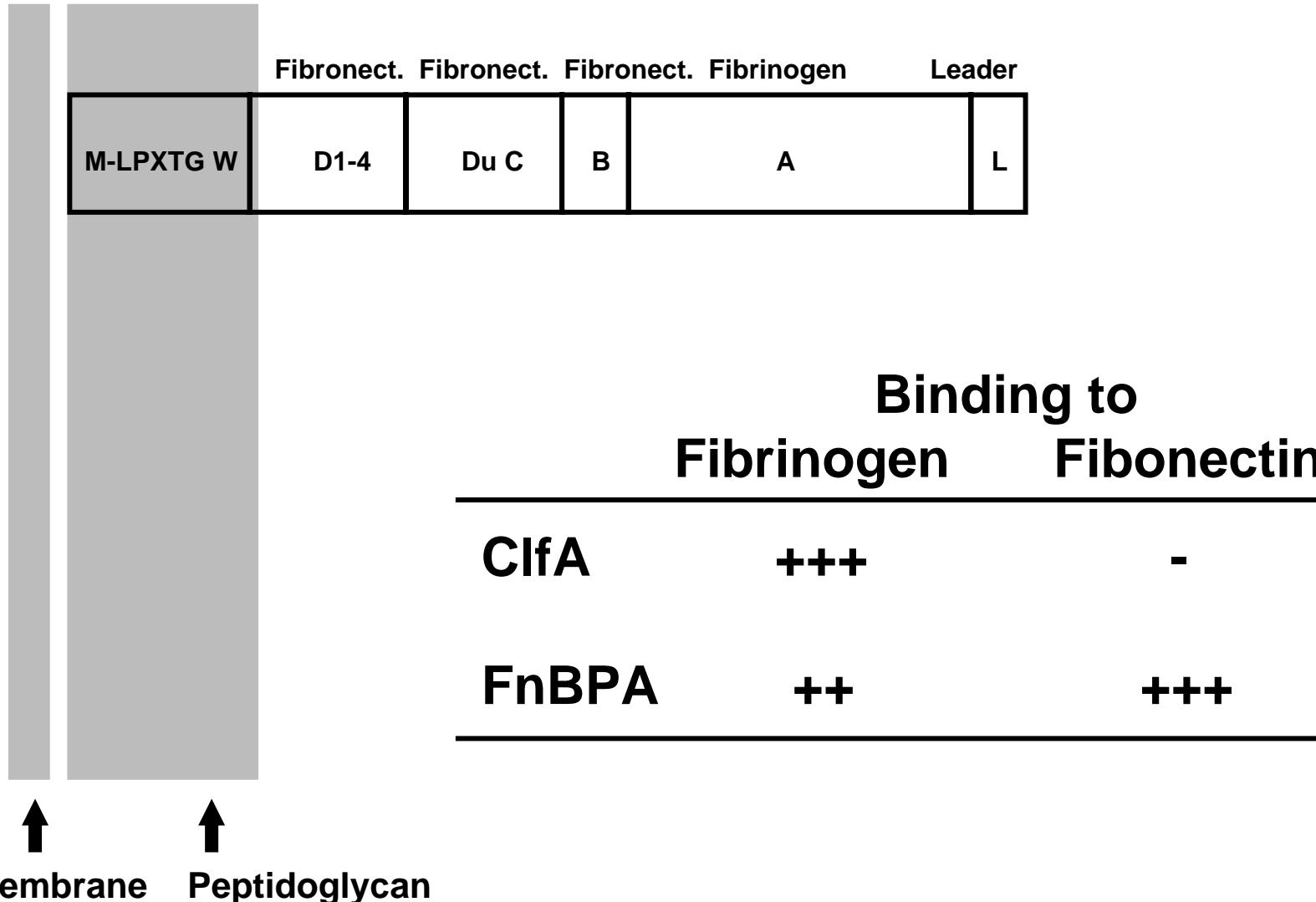
Neighboring Aorta of Valves Infected with ClfA-positive Lactococci /Gram Stain



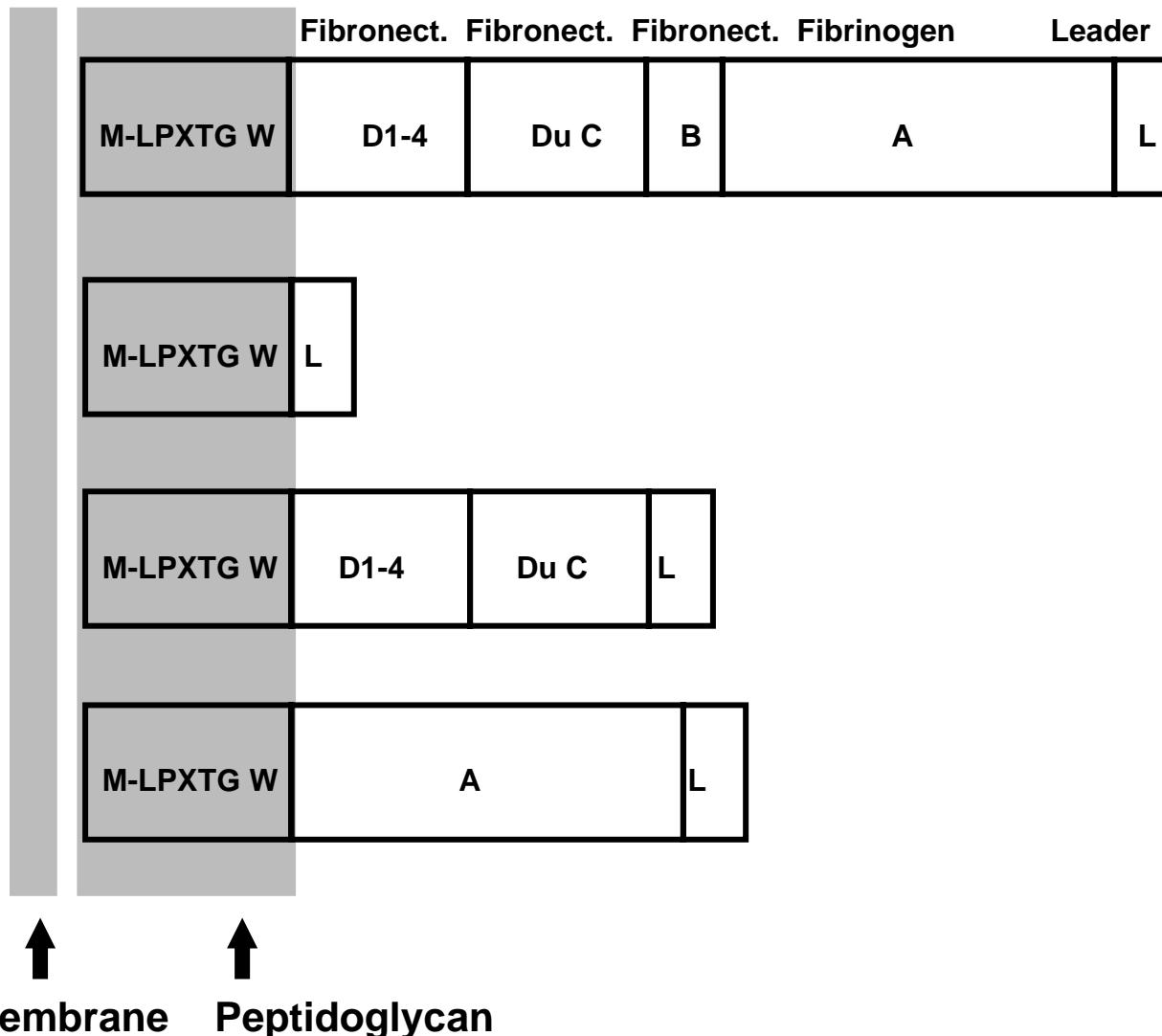
L. lactis FnBPA (+)



Binding Domains of FnBPA

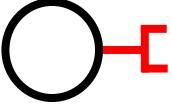
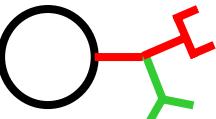
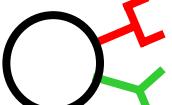


Binding Domains of FnBPA

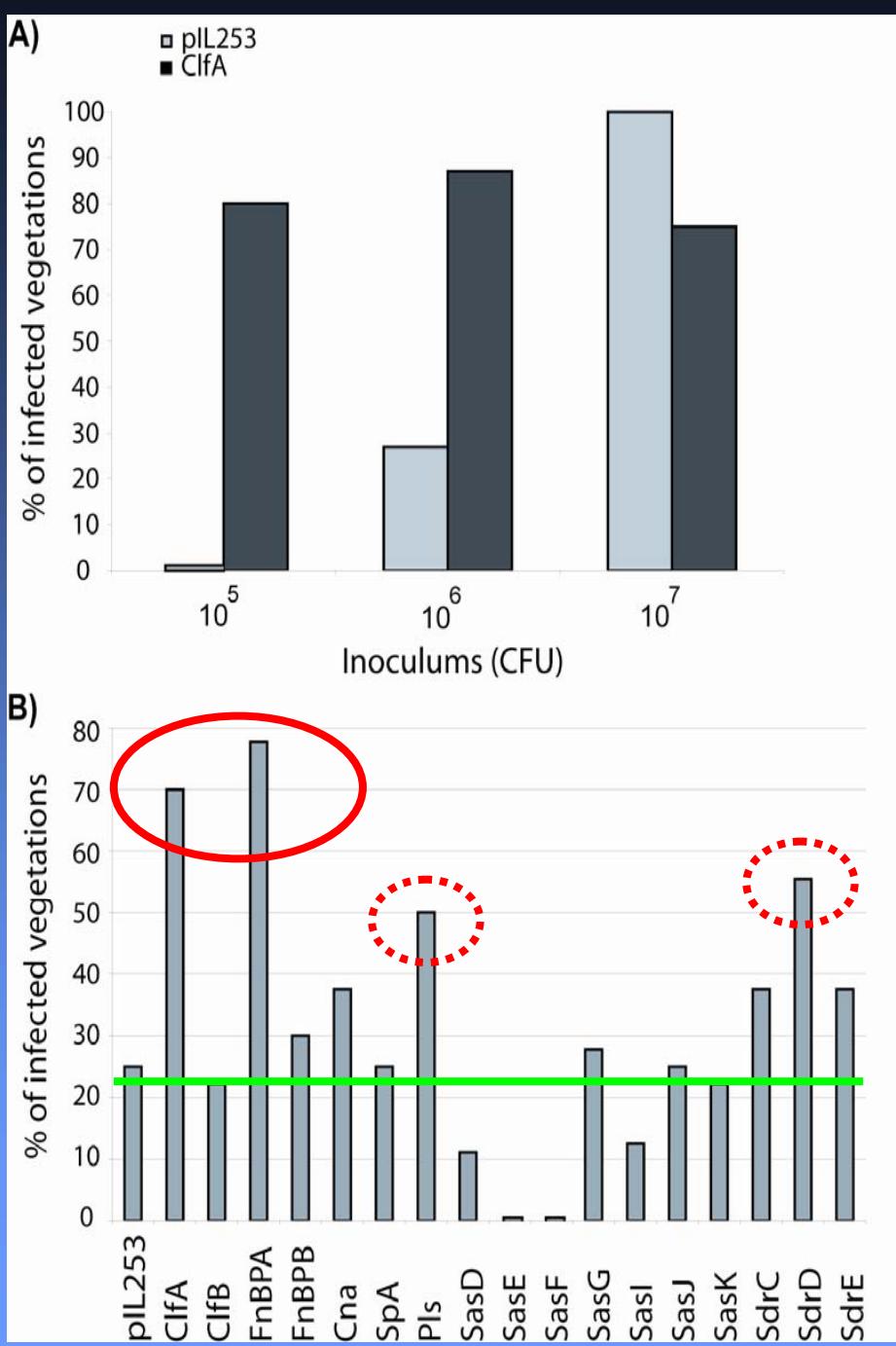
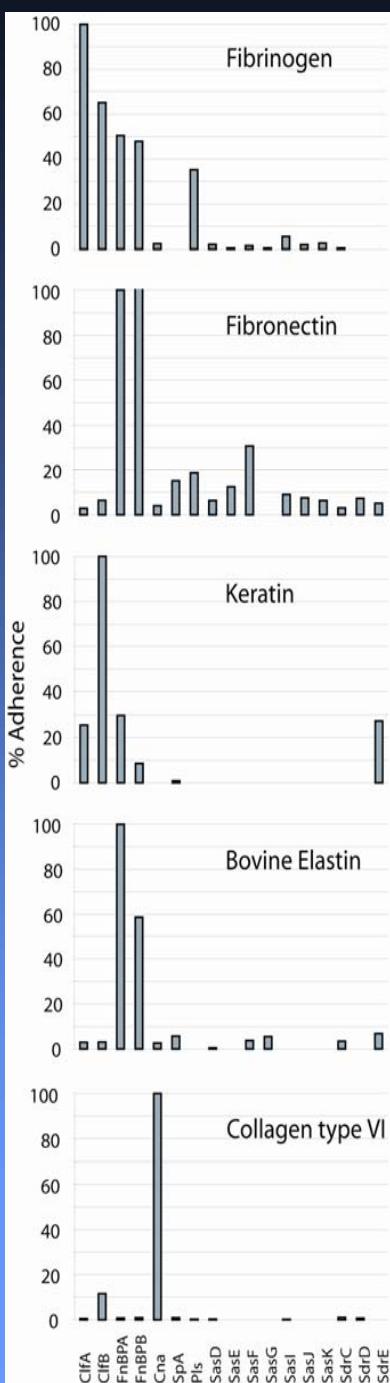


Membrane Peptidoglycan

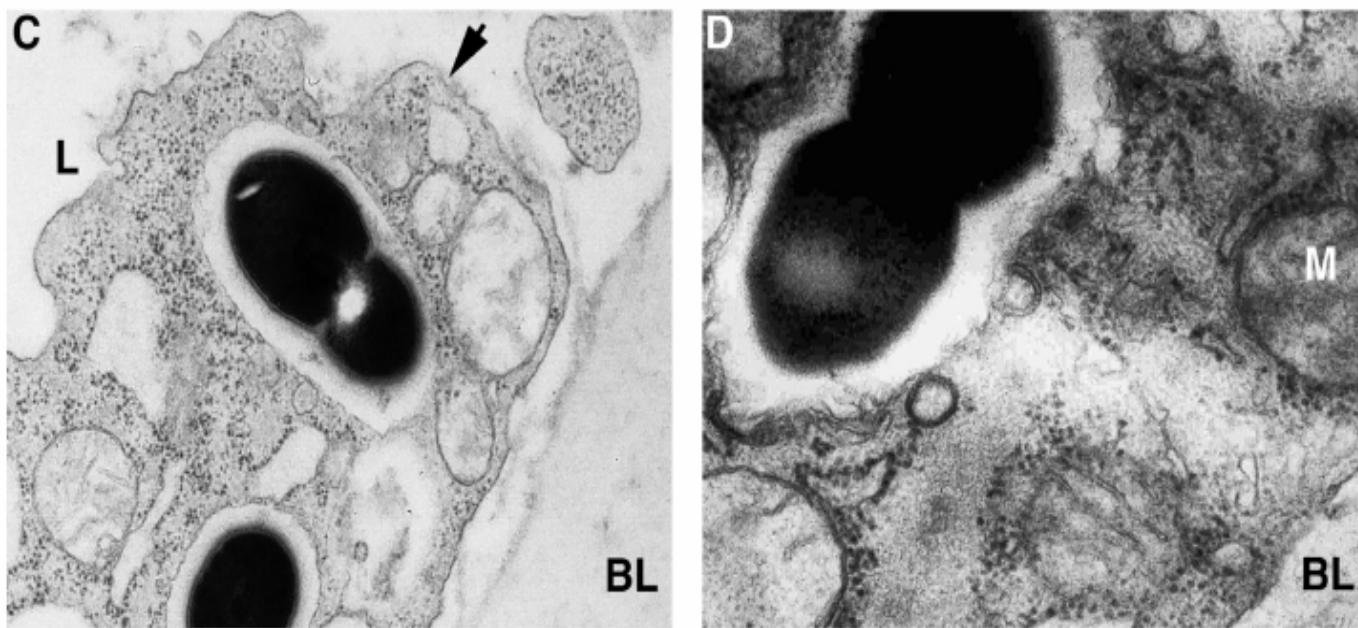
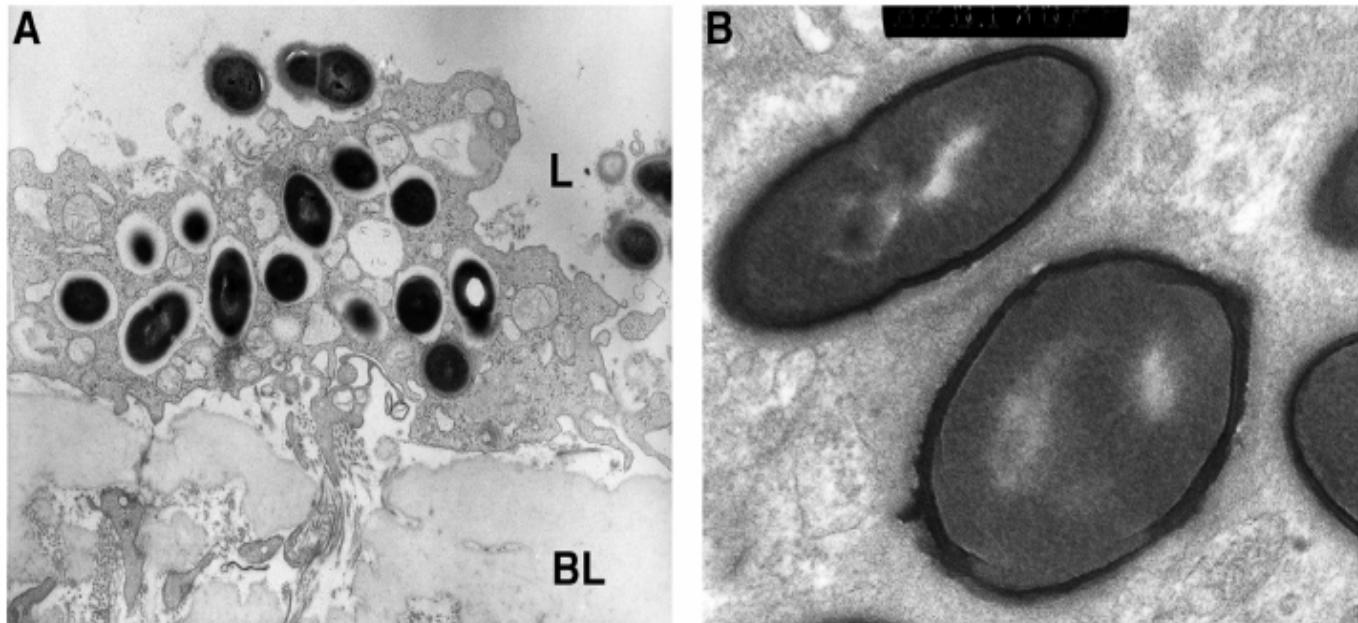
Binding Domains of FnBPA

Surrogate recombinant	Binding	Infection	Internalization	Persistence
	Fg	+++	---	---
	Fn	---	+++	---
	<i>cis</i> Fn/Fg	+++	+++	+++
	<i>trans</i> Fn/Fg	+++	+++	+++

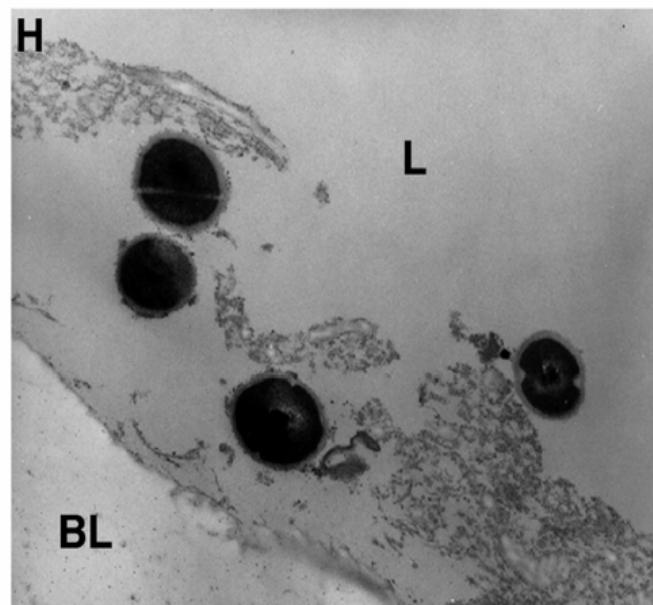
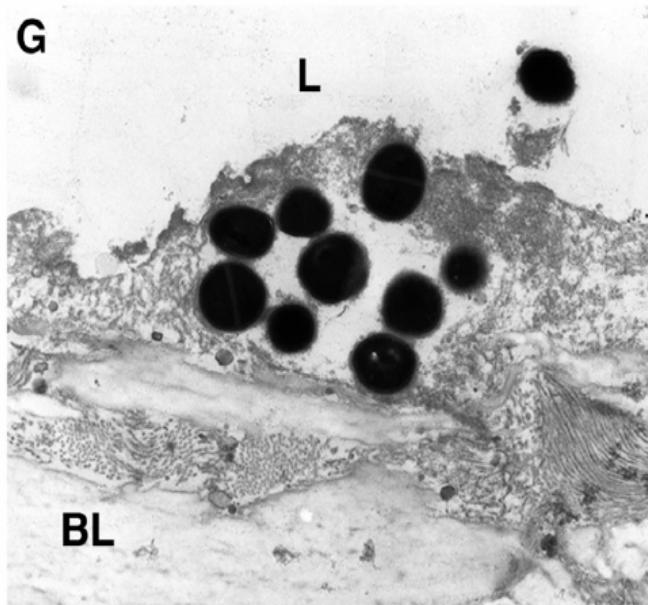
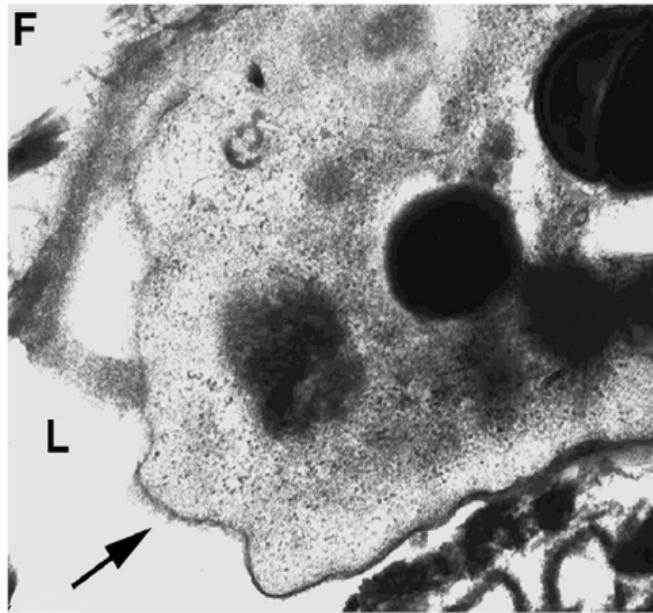
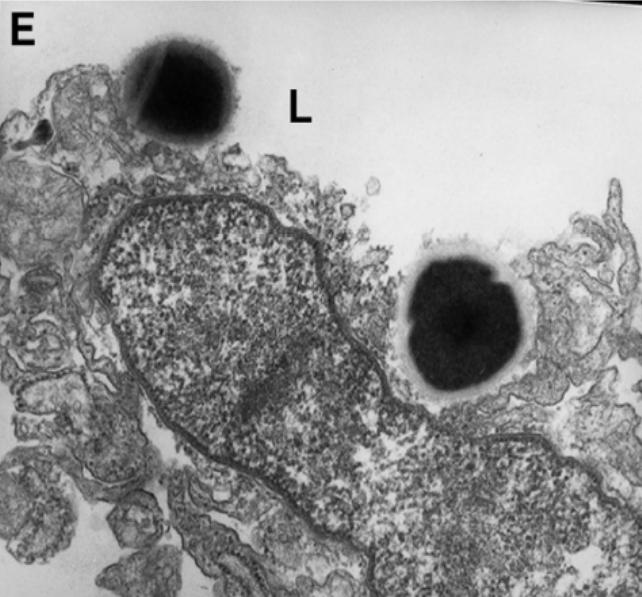
- *S. aureus* needs both fibrinogen- and fibronectin-binding domains to be fully infective
- Fibronectin-binding protein represents an example of *intra-molecular* domain cooperation
- Fibronectin- and fibrinogen-binding represent an example of *inter-molecular* domain cooperation

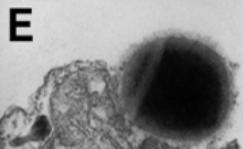


L. lactis FnBPA (+)

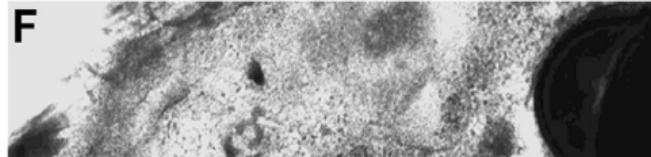


S. aureus Cowan





L



LPXTG-proteins

Toxins

<i>S. aureus</i>	21	> 30
------------------	----	------

<i>S. epidermidis</i>	> 10	No
-----------------------	------	----

<i>S. haemolyticus</i>	6	No
------------------------	---	----

<i>S. saprophyticus</i>	1	No
-------------------------	---	----

But all have prophages and genomic islands, including the SCC staphylococcal K7 chromosome



Staphylococcus aureus a paradigm of host-parasite evolution

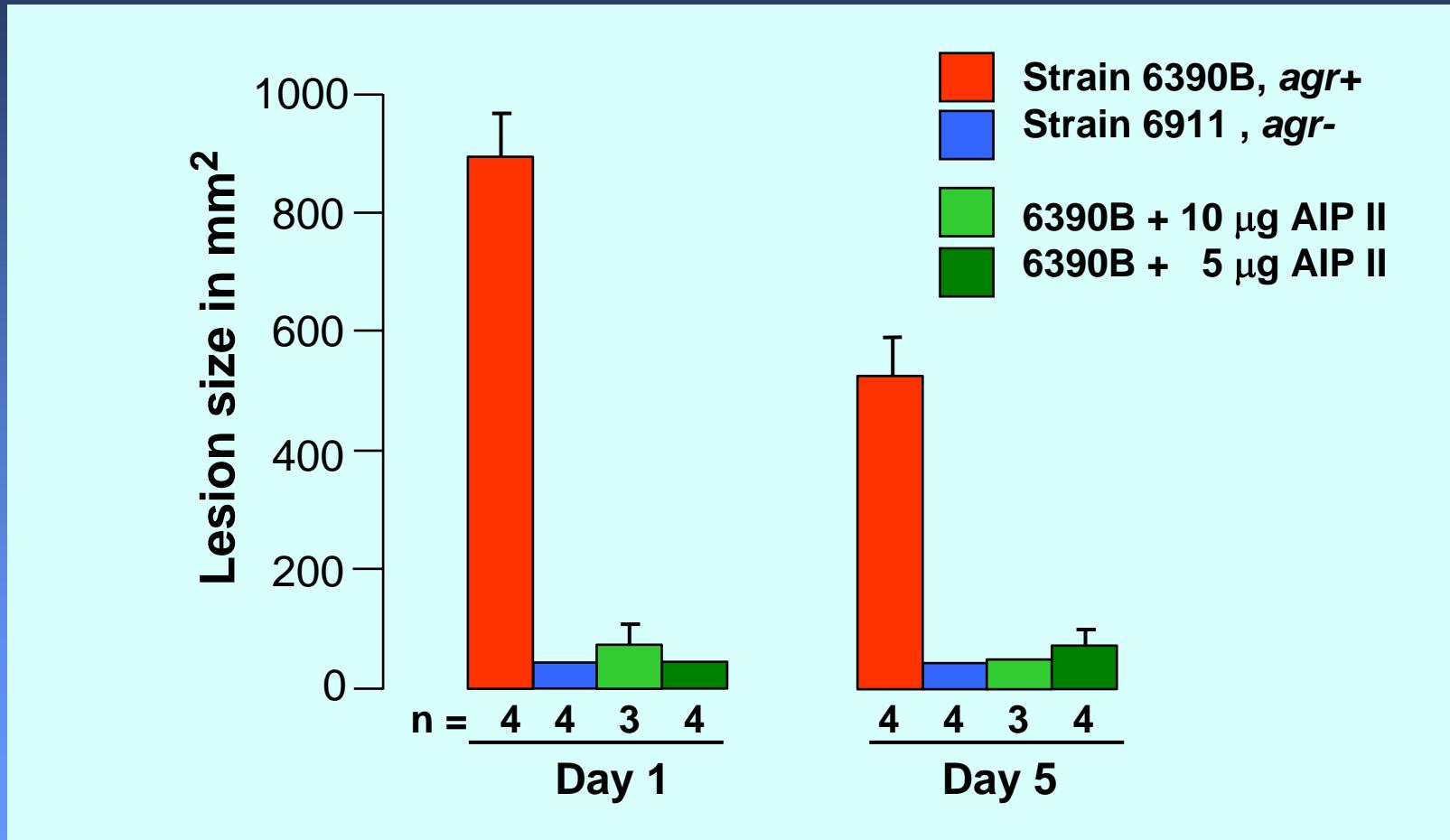
N. Widmer
M. Giddey
J. Entenza
S. Piu
J. Vouillamoz

YA Que
L Piroth
CA. Roten
L. Guy
F. Collyn



Subcutaneous Infection of Mice with 10^8 CFU of AIP I

S. aureus supplemented or not with AIP II



Mayville et al. PNAS 1999; 96:1218-1223

Is *agr* a Virulence Factor ?

- No, not *sensu stricto*
- All *Staphylococcus* spp. carry an *agr*
- *agr* is also required for regulation of house-keeping genes