

Teixobactin- Entdeckung und Wirkungsmechanismus

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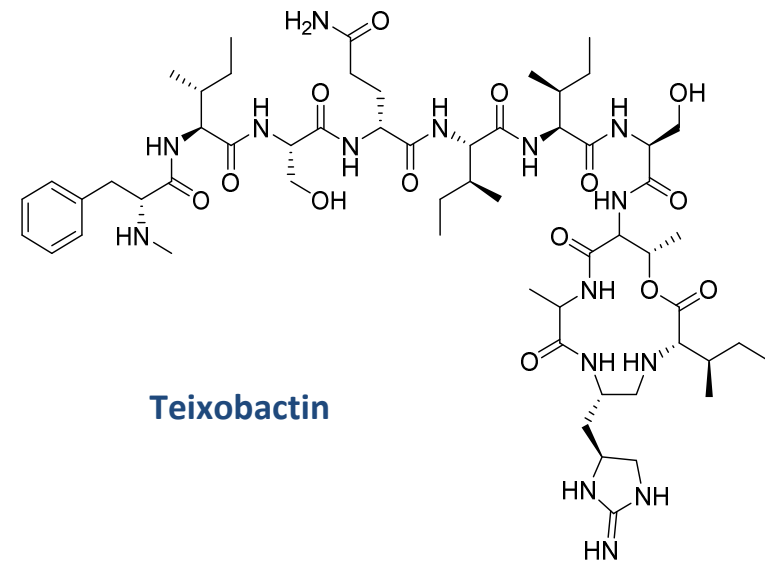
Institute for Pharmaceutical Microbiology
University of Bonn

Bad Honnef-Symposium 2015



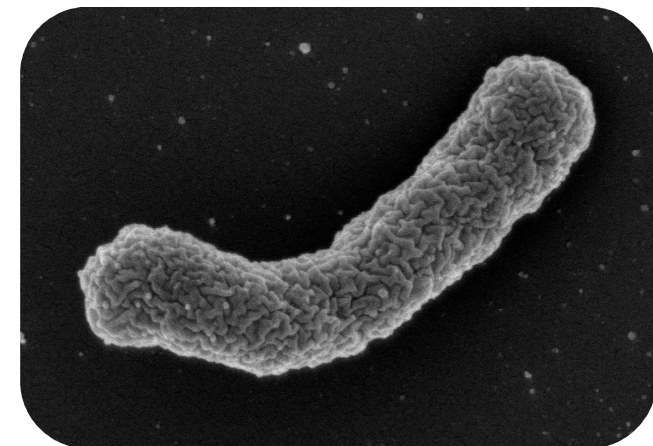
Teixobactin

- 11 amino acid depsipeptide
- enduracididine, methylphenylalanine, D-amino acids
- MW: 1.242 Da



Teixobactin

- new β -proteobacterium: *Eleftheria terrae*
- Gram-negative, related to Aquabacteria
- isolated from a grassy field in Maine



Eleftheria terrae

(William Fowley, Northeastern University)

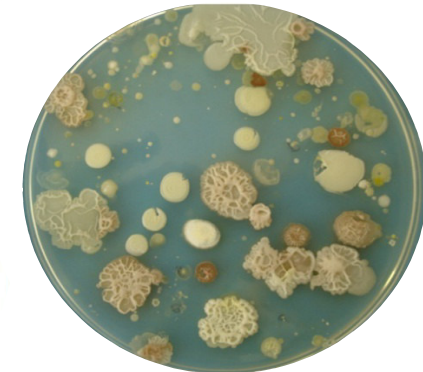
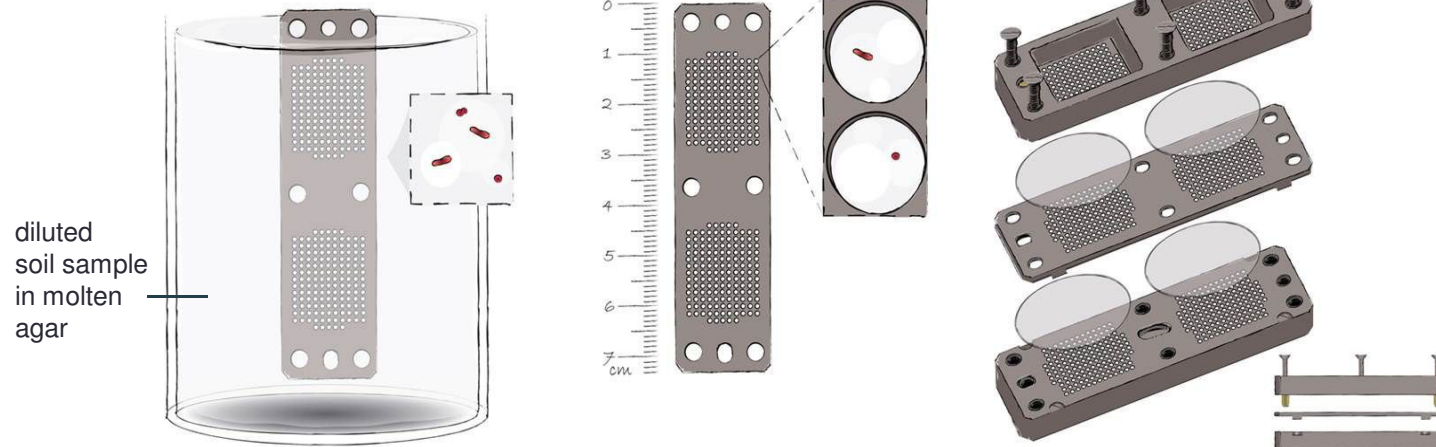
Growing the „uncultured“ – „Great plate count anomaly“

- most antibiotics are produced by environmental microorganisms
 - only 1% of all microbial species grow on nutrient media
 - ~ 99% constitute so far „uncultured bacteria“
 - „dark matter“: promising source for new antibiotics

(Stewart, 2012; Hongoh & Toyoda, 2001, Lewis *et al.*, 2010)

iChip

- grow and isolate „uncultured“ bacteria in one step in their natural environment
- high-throughput microbial cultivation

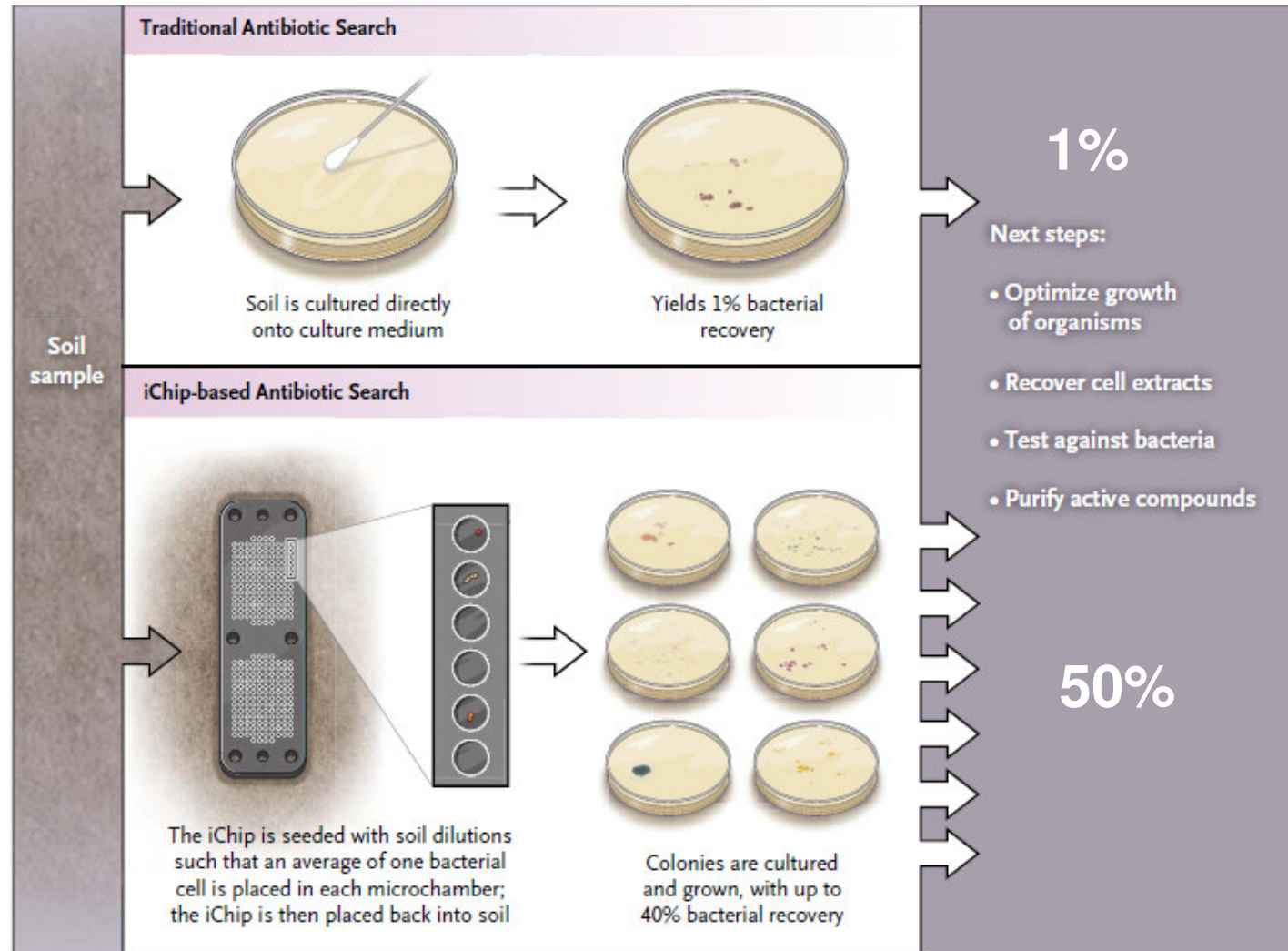


Slava Epstein, Northeastern University

(Nichols *et al.*, AEM, 2010)

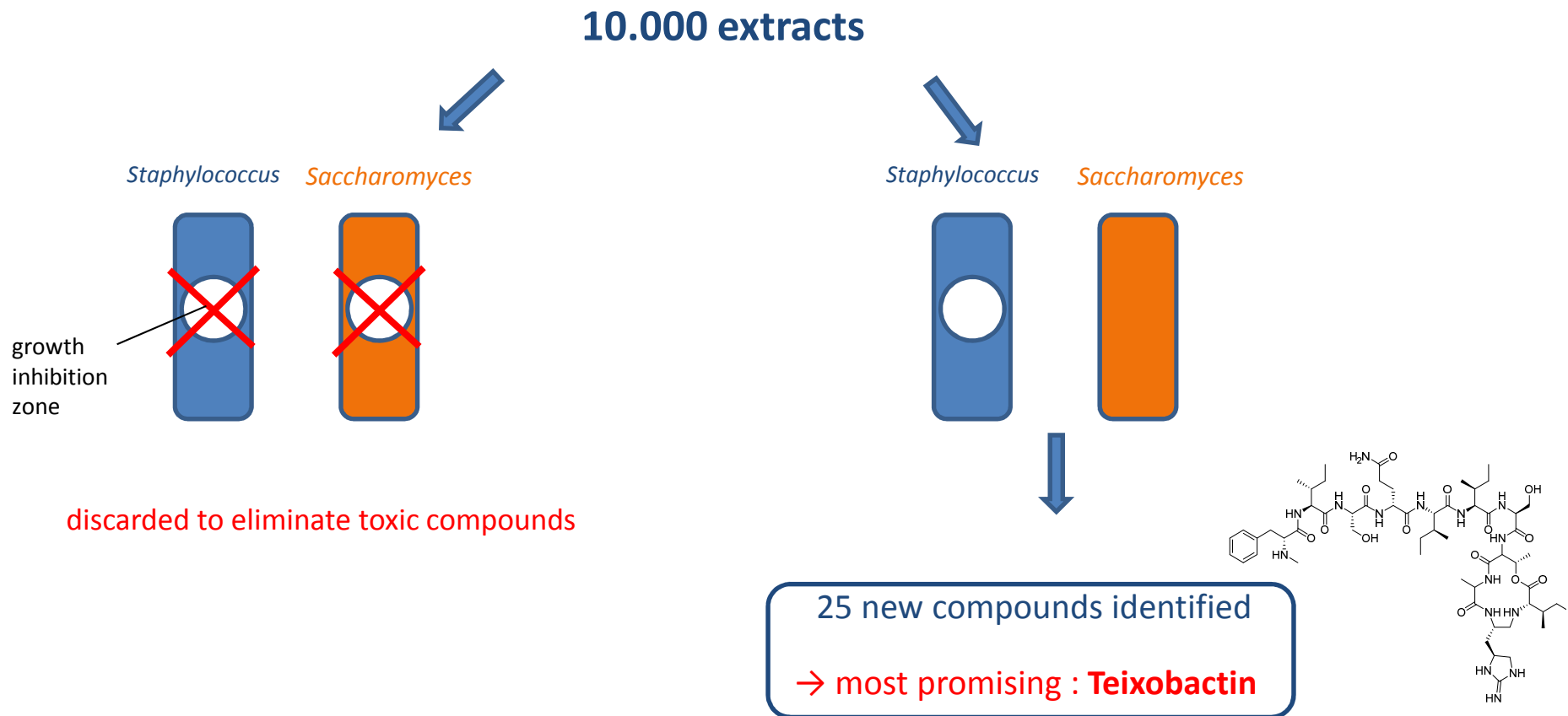
Traditional vs iChip based search

Growth recovery → iChip: 50% vs. petridish : 1 %



Growing the „unculturable“ - iChip (isolation chip)

- ~50.000 previously „uncultured“ bacteria were isolated from soil samples
- extracts of 10.000 isolates have been screened for antimicrobial activity
- 25 new compounds identified

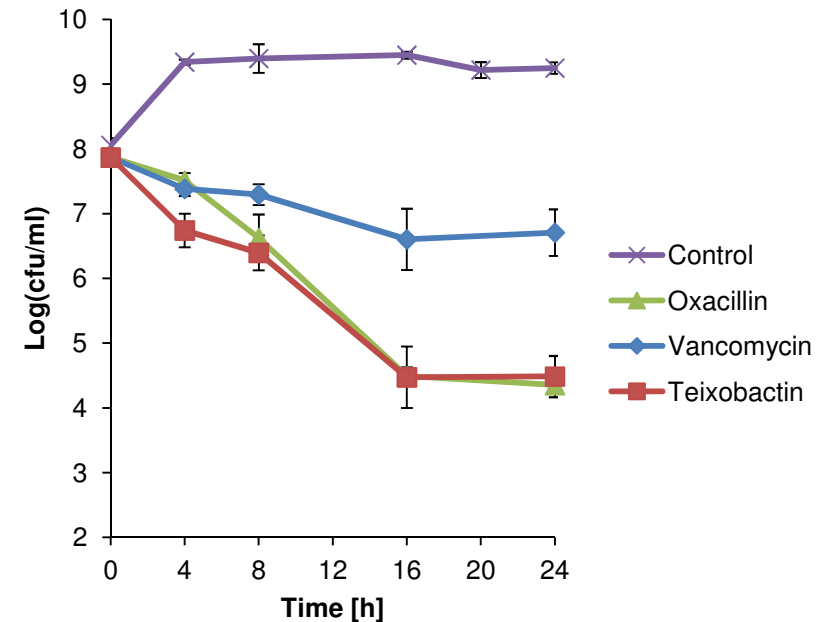


Activity of teixobactin against pathogenic microorganisms

- excellent activity against Gram-positive pathogens

pathogen	MIC (µg/ml)
<i>Staphylococcus</i>	
<i>S. aureus</i> ATCC 33591 (MRSA)	0.16
<i>S. aureus</i> ATCC 700699 (GISA)	0.32
<i>S. aureus</i> (DAP-R)	0.12
<i>S. aureus</i> (LIN-R)	0.12
<i>S. aureus</i> (VISA)	0.12
<i>Enterococcus</i>	
<i>E. faecium</i> (VRE)	0.31
<i>E. faecalis</i> (VRE)	0.31-0.63
<i>Mycobacterium</i>	
<i>M. tuberculosis</i> H37Rv	0.125
<i>M. tuberculosis</i> (clin. isolate 70)	0.125-0.25
<i>Streptococcus</i>	
<i>S. pneumoniae</i> ATCC 6303	0.02
<i>S. pyogenes</i>	0.31
 <i>Bacillus anthracis</i>	 <0.06
<i>Clostridium difficile</i> CD 196	0.005

Time dependent killing of *S. aureus*



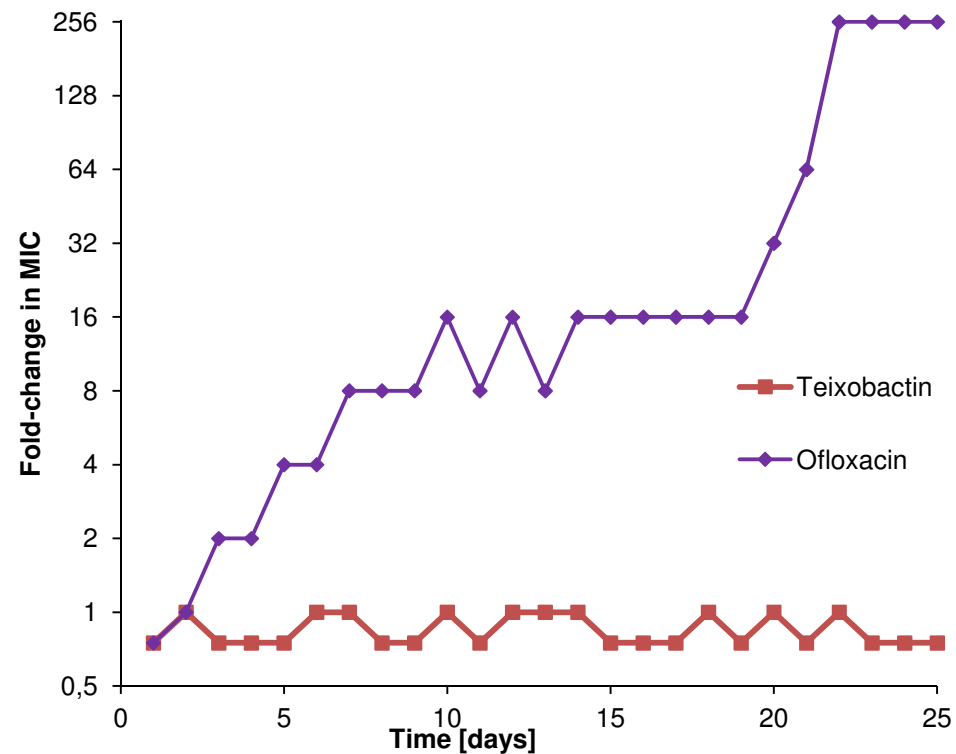
Control Oxacillin Vancomycin Teixobactin



- superior to vancomycin in killing *S. aureus*
- excellent bactericidal activity

No resistance acquisition detected during serial passaging

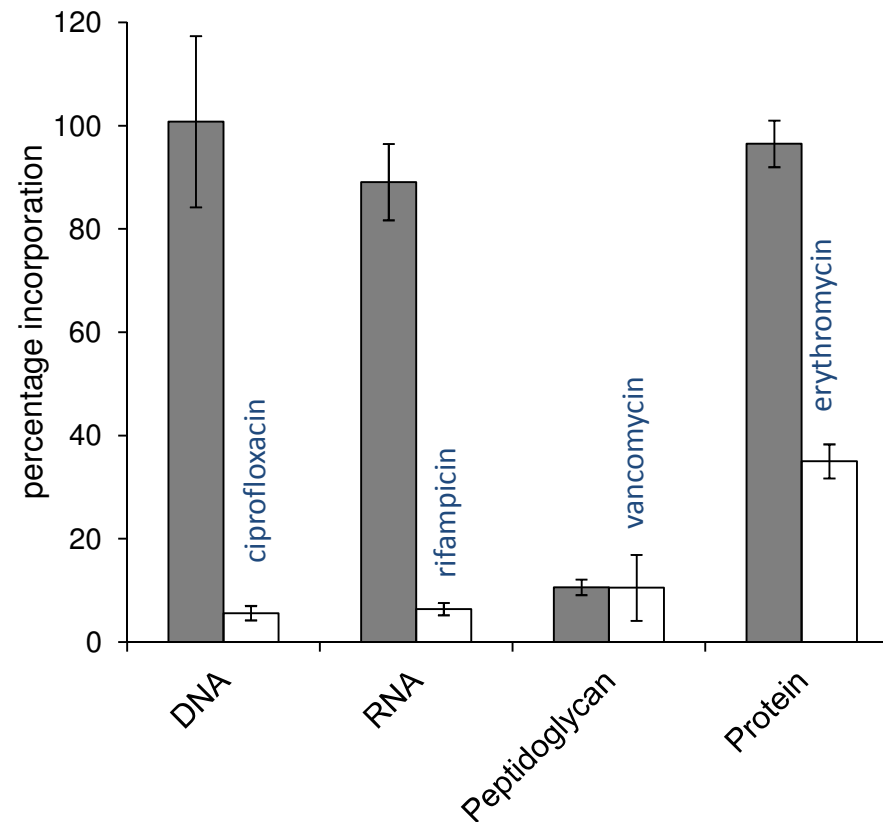
- no selection of resistant mutants of *S. aureus* oder *M. tuberculosis*, even at low doses
- serial passaging of *S. aureus* on sub-MIC levels of teixobactin
→ failed to produce resistant mutants



→ no toxicity against mammalian cells at 100 µg/ml (highest dose tested)

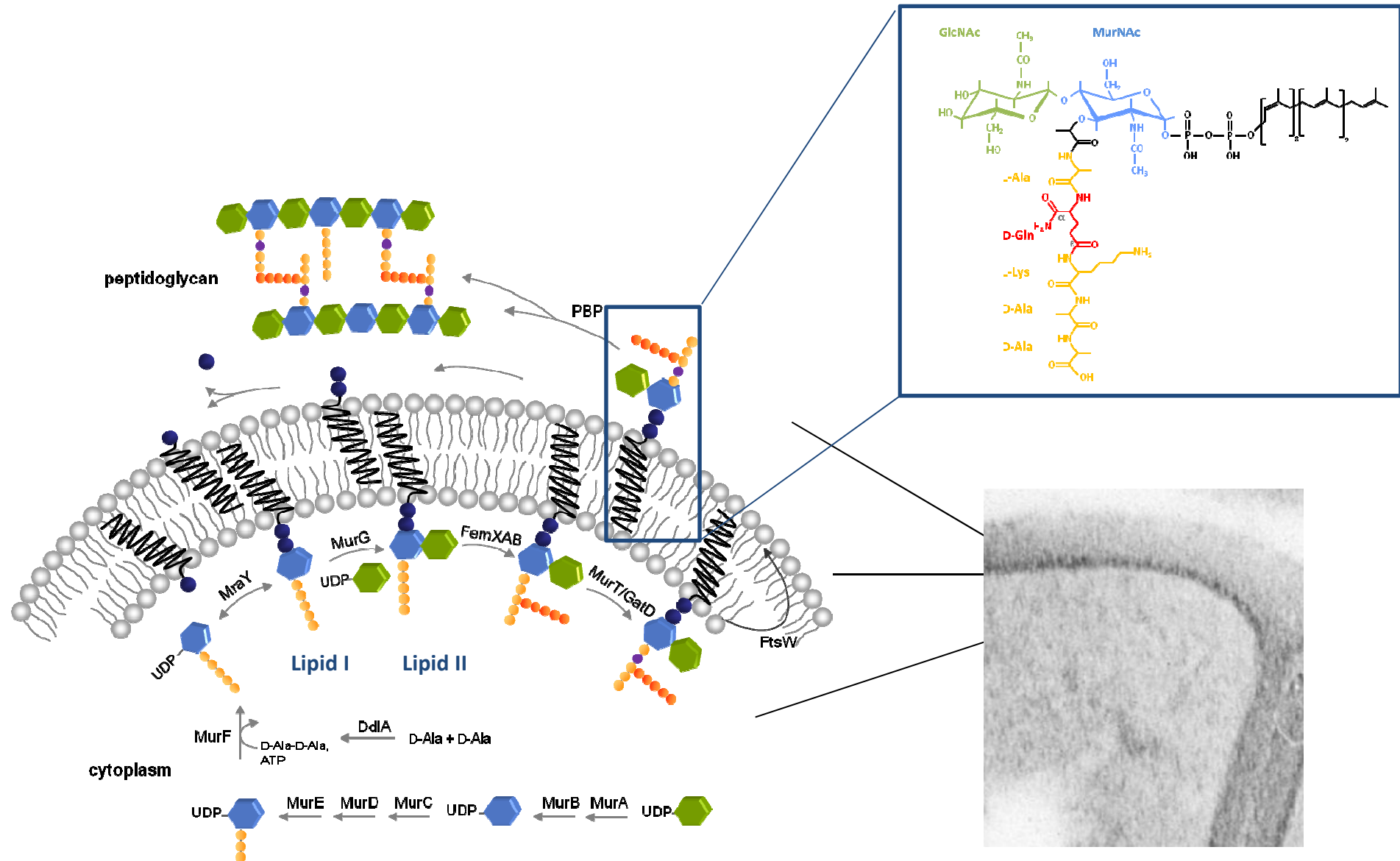
Mode of action analysis

Impact of teixobactin on macromolecular biosyntheses

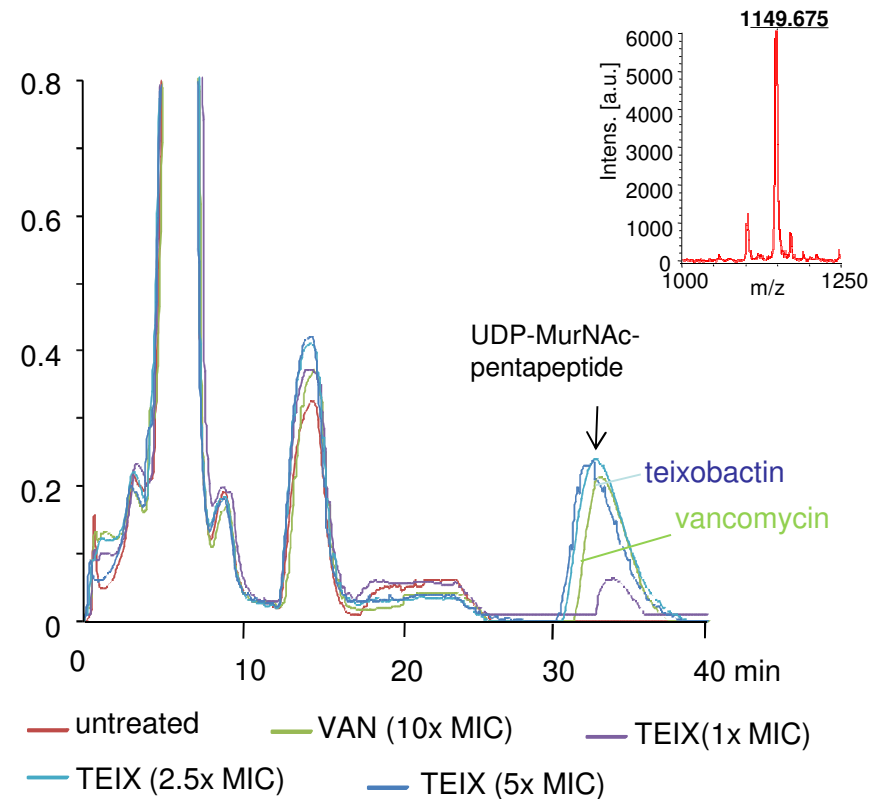
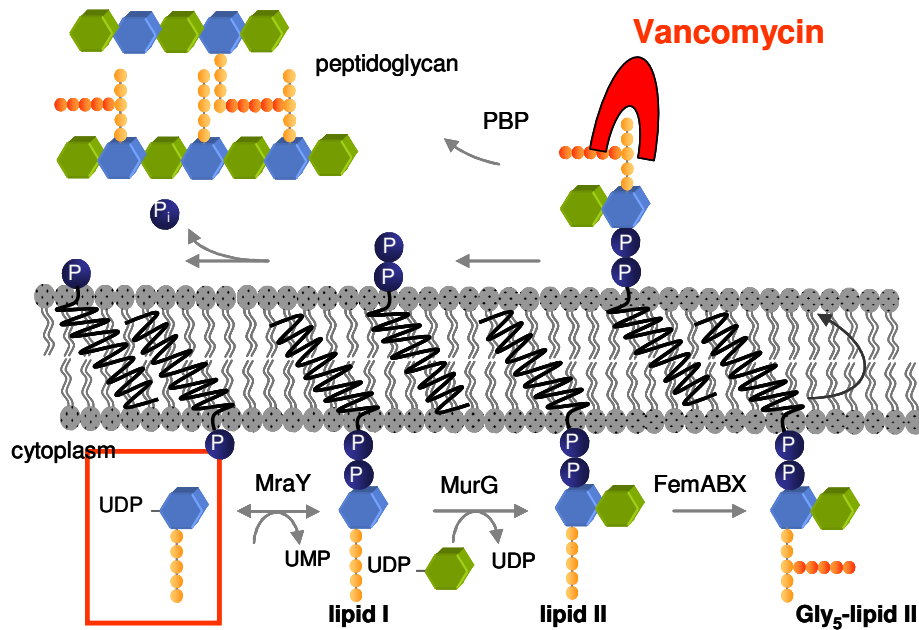


- incorporation of radiolabeled leucin (protein), thymidine (DNA) and uridine (RNA) was unaffected
- glucosamine was no longer incorporated – pointing towards cell wall biosynthesis as a target

Peptidoglycan biosynthesis in *S. aureus*

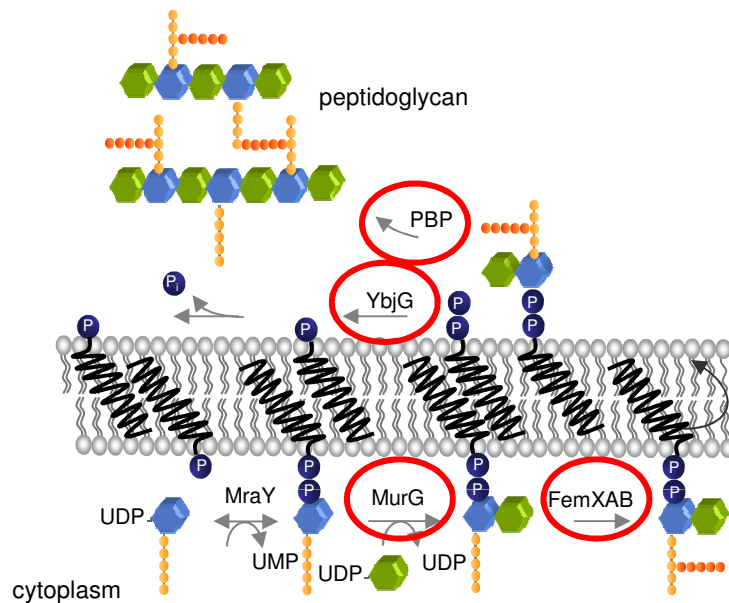


Teixobactin causes the accumulation of UDP-MurNAc-pentapeptide



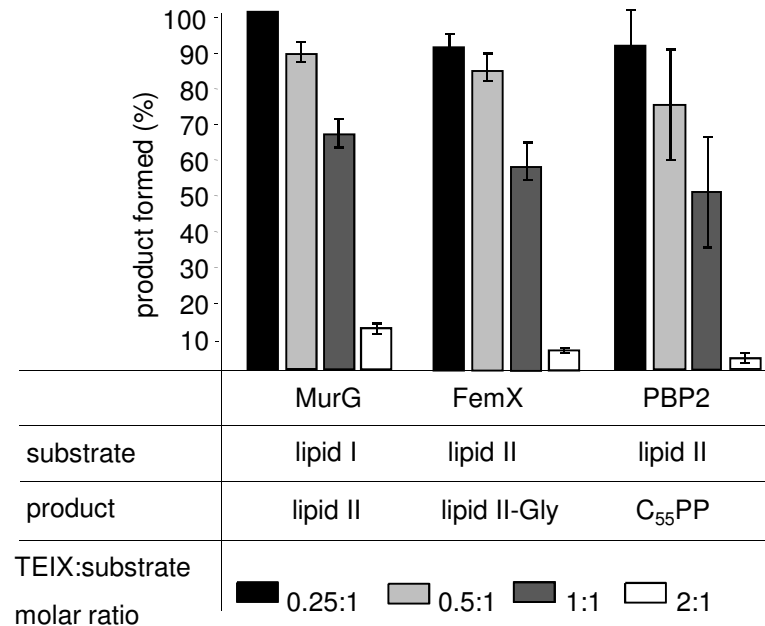
- teixobactin-treated cells accumulate the ultimate soluble cell wall precursor UDP-MurNAc-p₅, indicating that a later, membrane-associated step of cell wall biosynthesis is inhibited

Inhibition of membrane associated cell wall biosynthesis steps



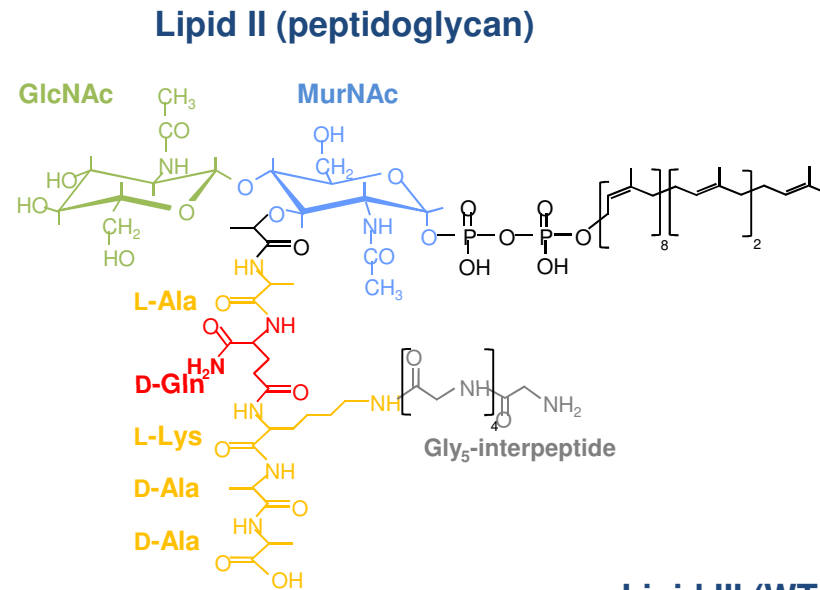
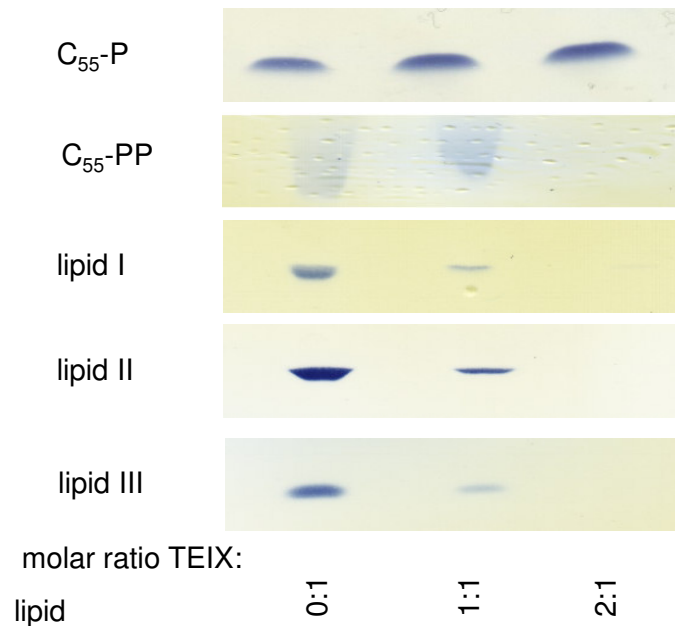
Lipid I Lipid II

- full inhibition is achieved at 2:1 molar ratio (Teixobactin:lipid)
- teixobactin forms a stoichiometric complex with the substrates lipid I and lipid II, rather than inhibiting the enzyme itself
- analogous WTA & capsule biosynthesis reactions are similarly inhibited
- secondary target: universal lipid carrier (un)decaprenyl(pyro)phosphate

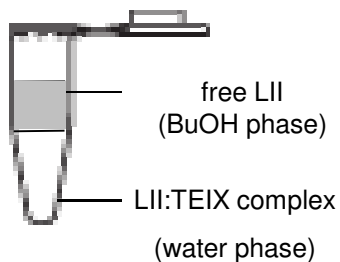


Complex formation with cell envelope precursors

- evaluate the minimal motif required for high affinity binding of teixobactin



BuOH extraction

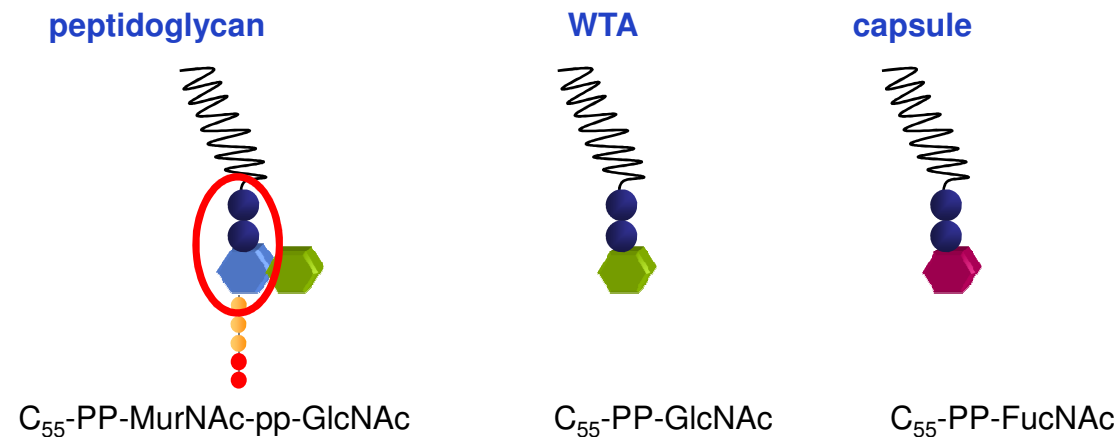


- pyrophosphate moiety is crucial for TEIX binding

Cell envelope precursors antagonize teixobactin antimicrobial activity

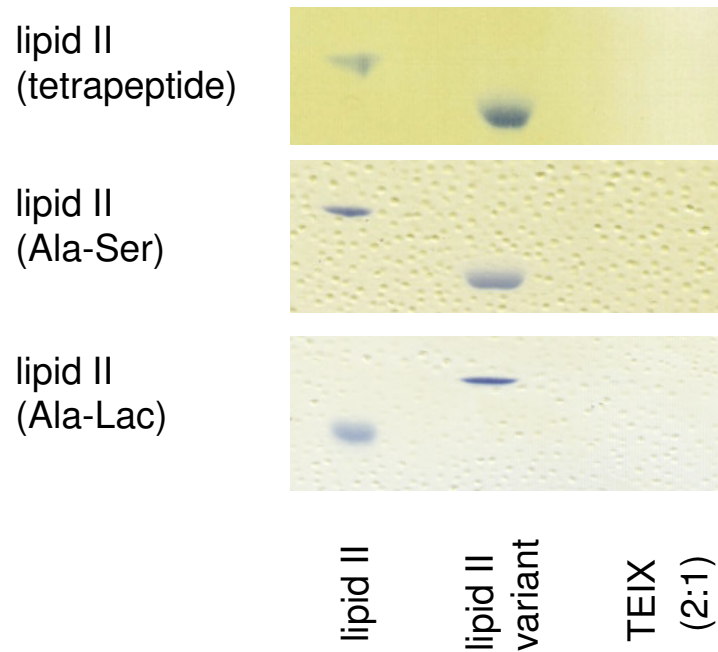
lipid intermediate	molar ratio of precursor to teixobactin						
	0 x	0.5 x	1 x	2.5 x	5 x	7.5 x	10 x
lipid II	-	+	+	+	+	+	+
C ₅₅ -PP	-	-	-	-	+	+	+

- high affinity binding relies on the interaction with the PP-sugar moiety
- the nature of the first sugar is less important



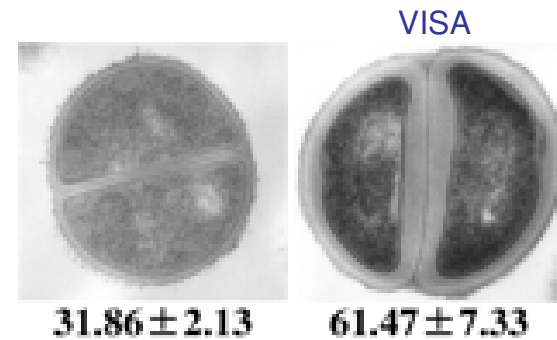
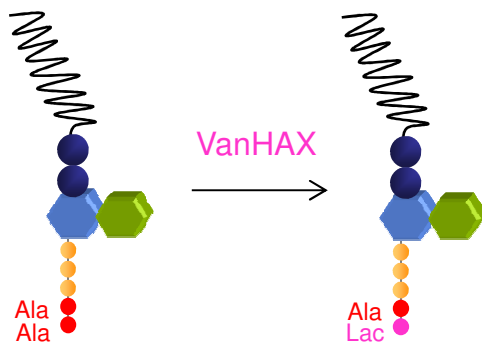
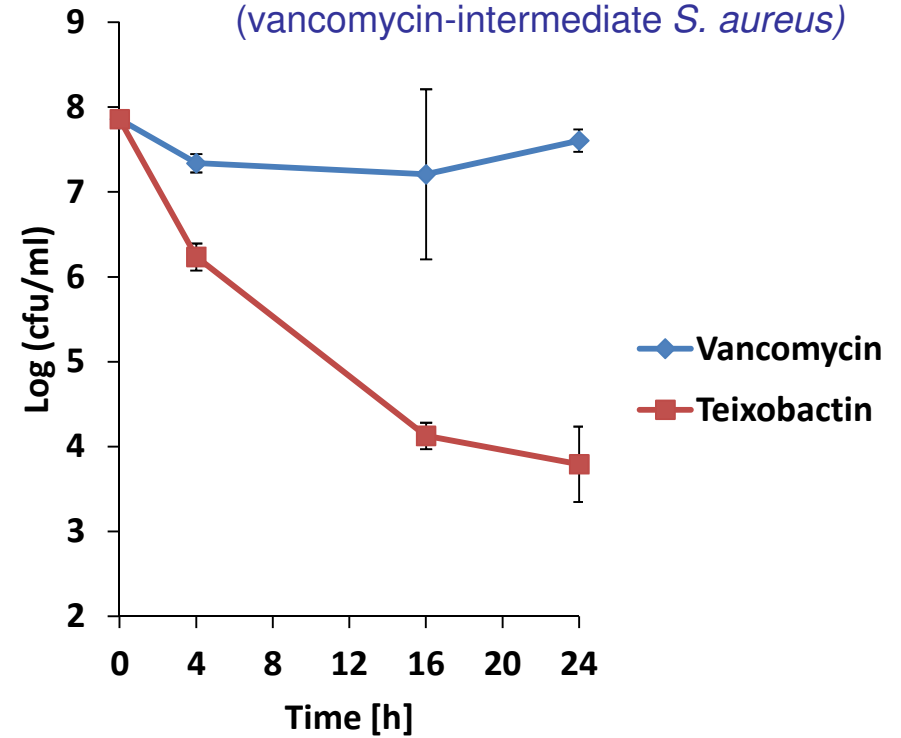
Teixobactin is active against Vancomycin resistant *S. aureus*

VRSA



Killing of VISA

(vancomycin-intermediate *S. aureus*)

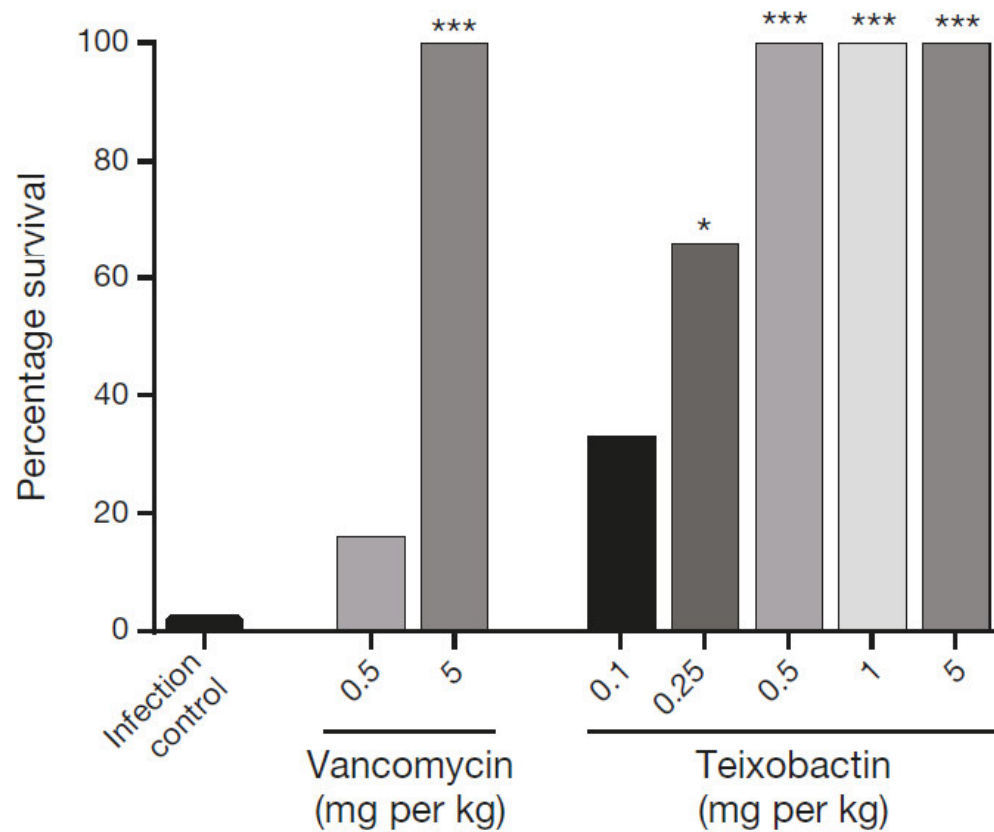


(Cui *et al.*, AAC, 2006)

Teixobactin is efficacious in mouse models of infection

- effective in 3 mouse models
- 100% survival of mice at 0.5 mg/kg

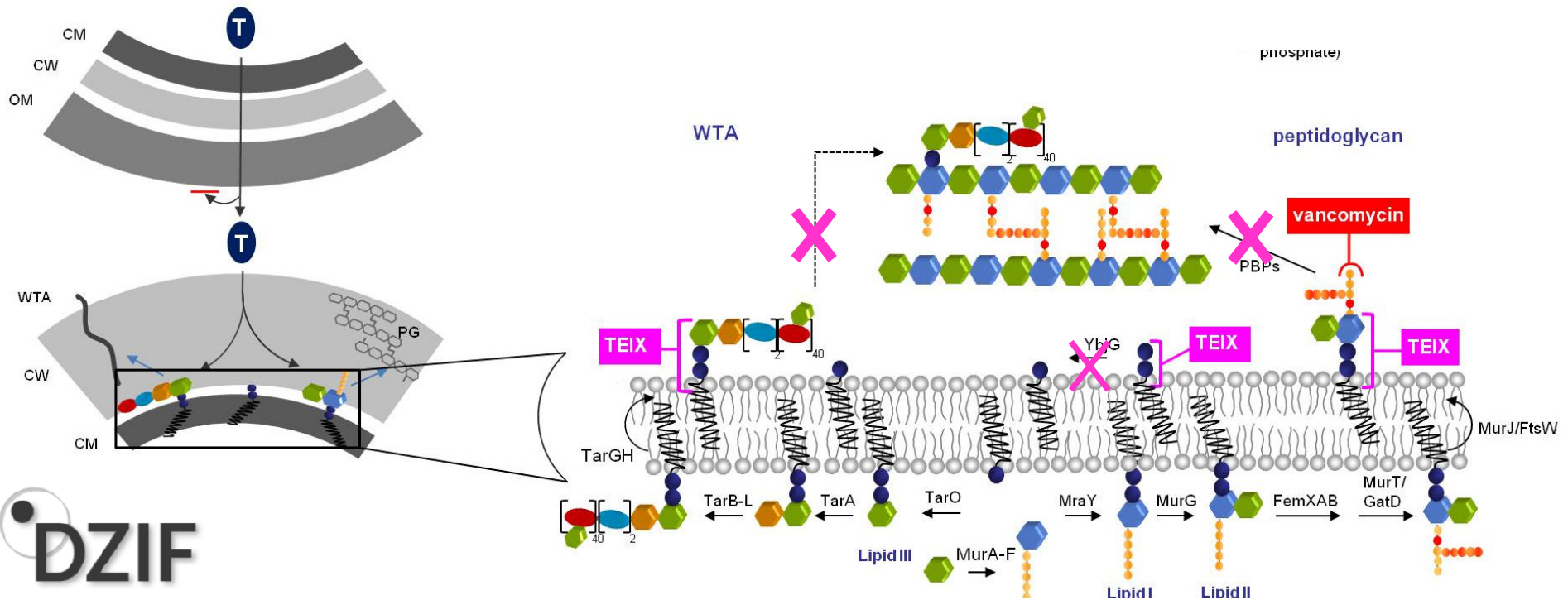
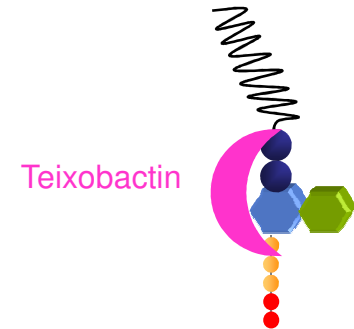
Mouse sepsis model



Mode of action model

- binds to multiple targets > polyprenol-coupled precursors of different cell envelope pathways
- highly conserved *non-protein* target structures
- binding sites are extremely hard to modify

→ strongly limits resistance development

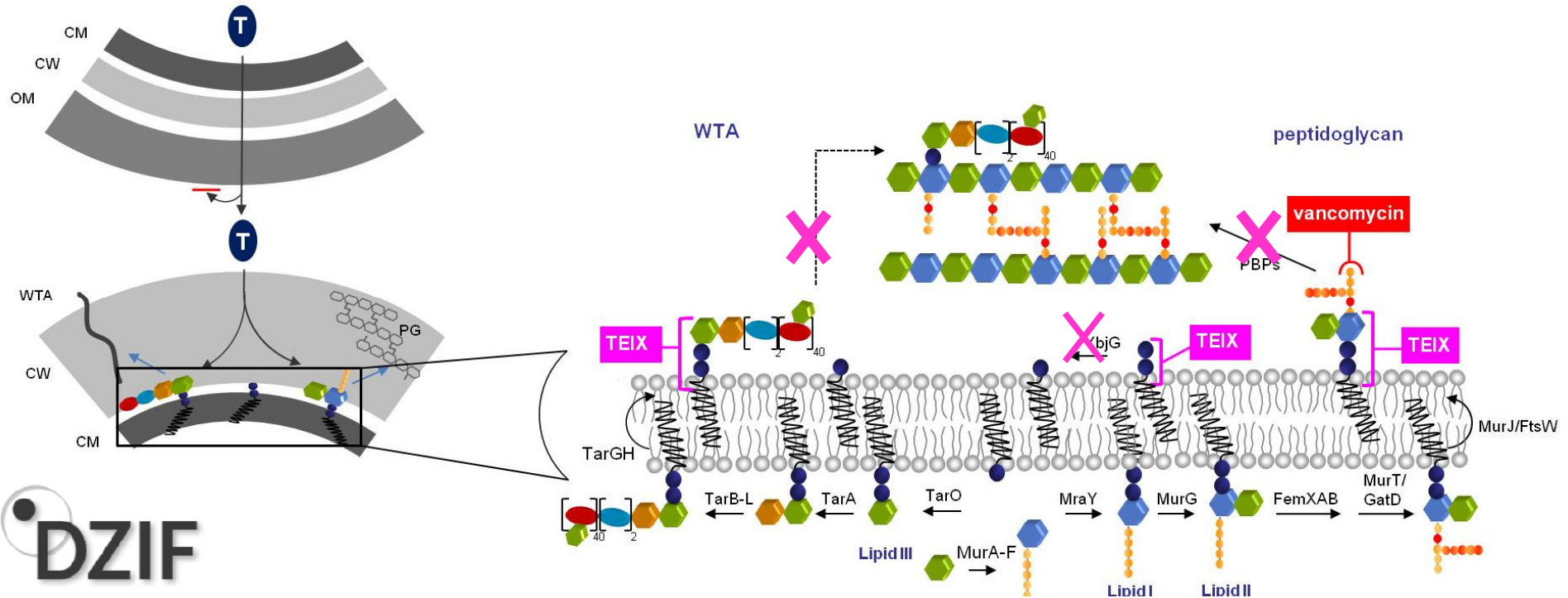


Mode of action model

- the producer is a gram-negative bacterium and the outer membrane protects it from re-entry of teixobactin → there appear to be no additional resistance mechanisms that other bacteria could borrow
- no cross-resistance to strains resistant to other „cell wall-active“ antibiotics, including VISA or DAP^R

BUT:

- antibiotic modifying or degrading enzymes might be produced by other microbes



Thanks ...

Ina Engels
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HG Sahl
M Josten

FOR 854



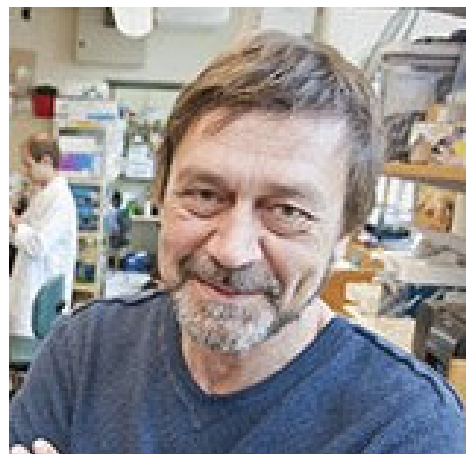
DFG Deutsche
Forschungsgemeinschaft

DZIF

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Kim Lewis



Slava Epstein

Lucy Ling, Novobiotic

Till Schäberle, University of Bonn

Brian Conlon, Northeastern University