

# School of Pharmacy

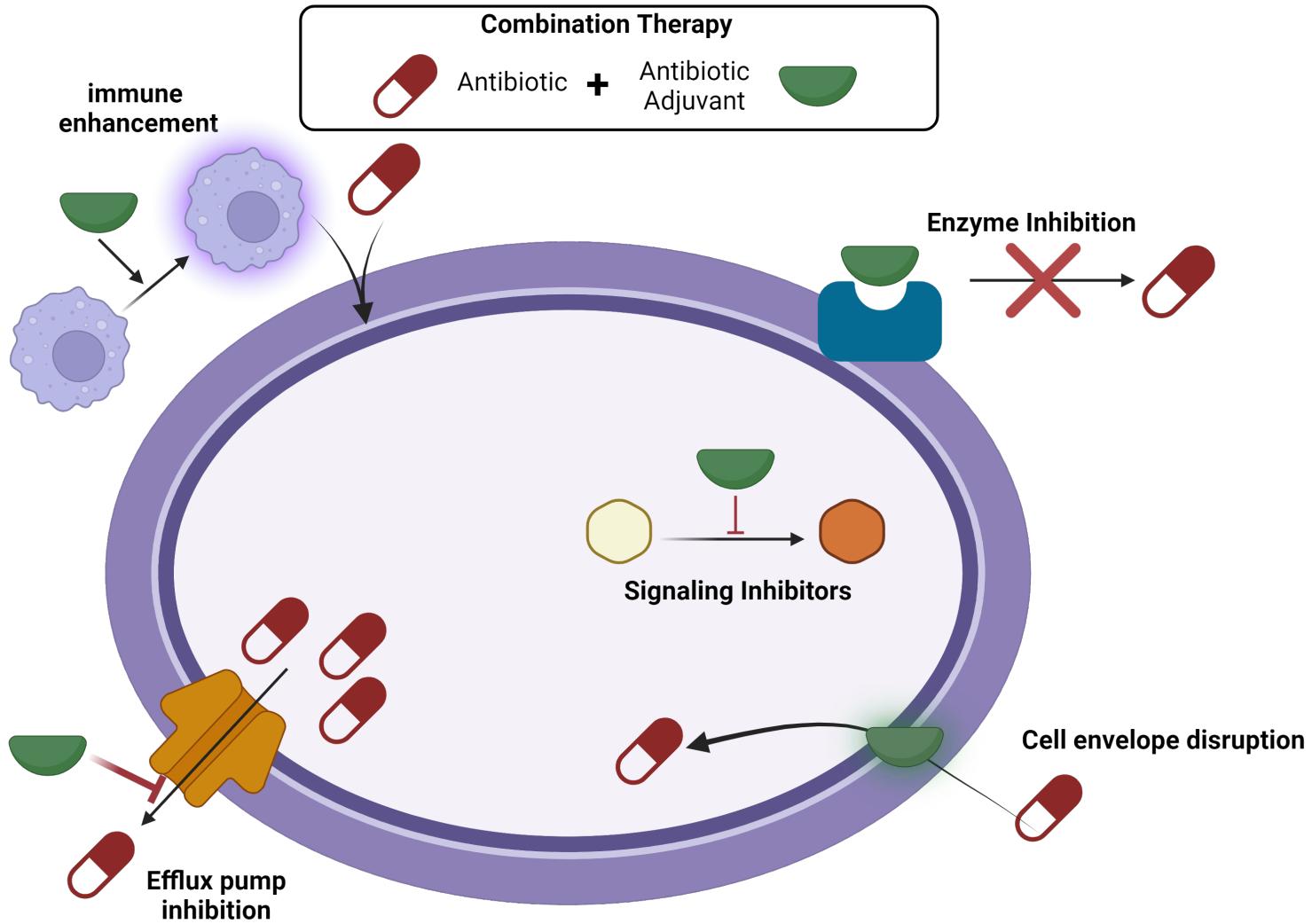
UNIVERSITY OF WISCONSIN-MADISON

## Novel Drug Combinations *and Mechanisms* to Address Antimicrobial Resistance

Warren Rose, PharmD, MPH, FIDSA  
Associate Professor



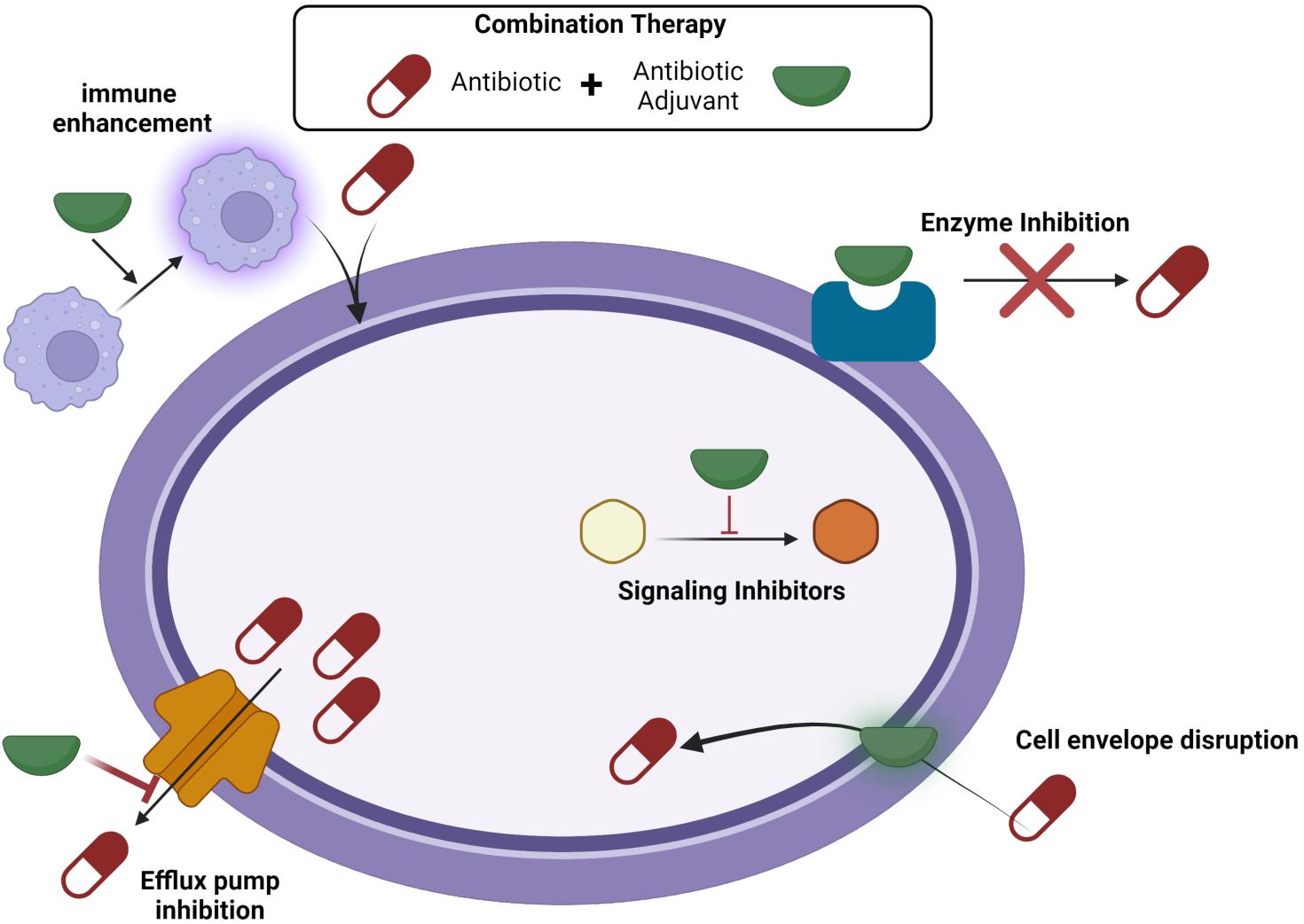
# Combination Therapy Approaches for AMR



Drug Repurposing e.g.  
Fosfomycin  
Minocycline  
Daptomycin  
Oritavancin  
 $\beta$ -lactams  
Ketoconazole  
Zidovudine  
Azithromycin



# Combination Therapy Approaches for AMR

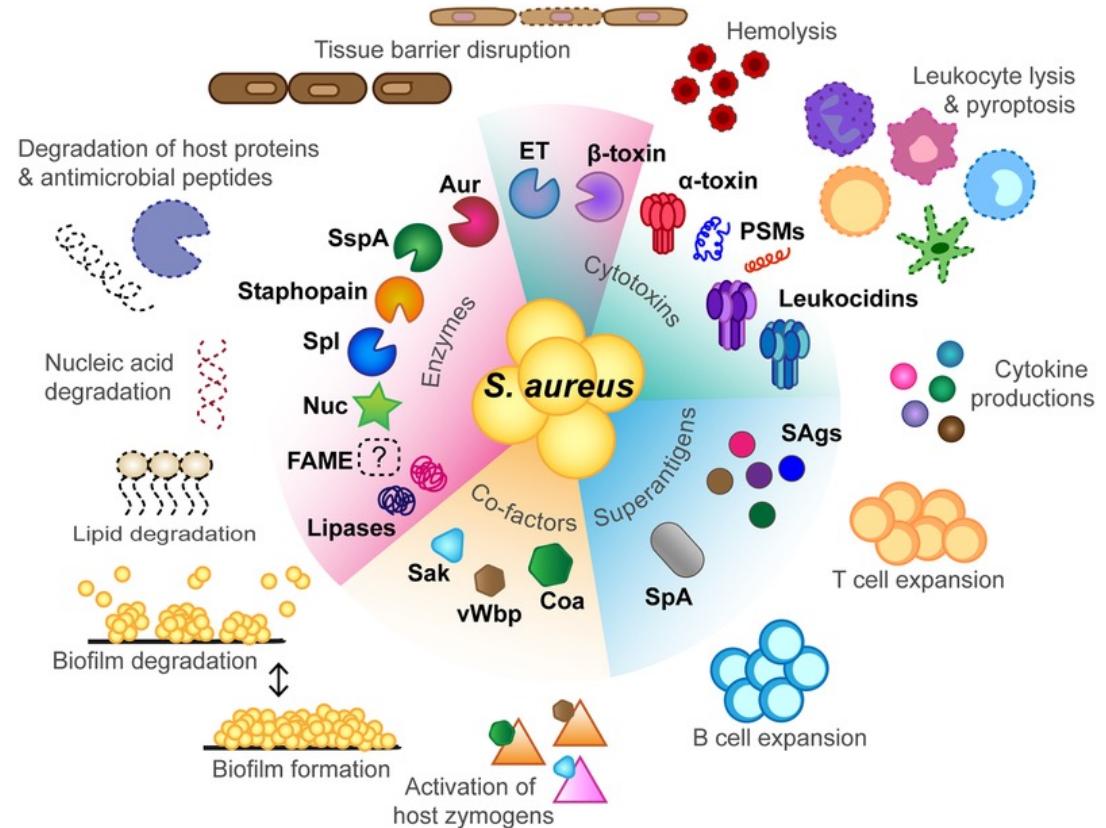


Drug Repurposing e.g.  
Fosfomycin  
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Oritavancin  
**β-lactams**  
Ketoconazole  
Zidovudine  
Azithromycin



# *Staphylococcus aureus*: the complicated pathogen

- Resistance documented to every class of antimicrobials
- Treatment failure persistently high for complicated infections
- Consistently changing epidemiology and patient risks
- Immune evasion...the non-commensal commensal



Tam and Torres. *Microbiol Spectr*. 2019 7(2)

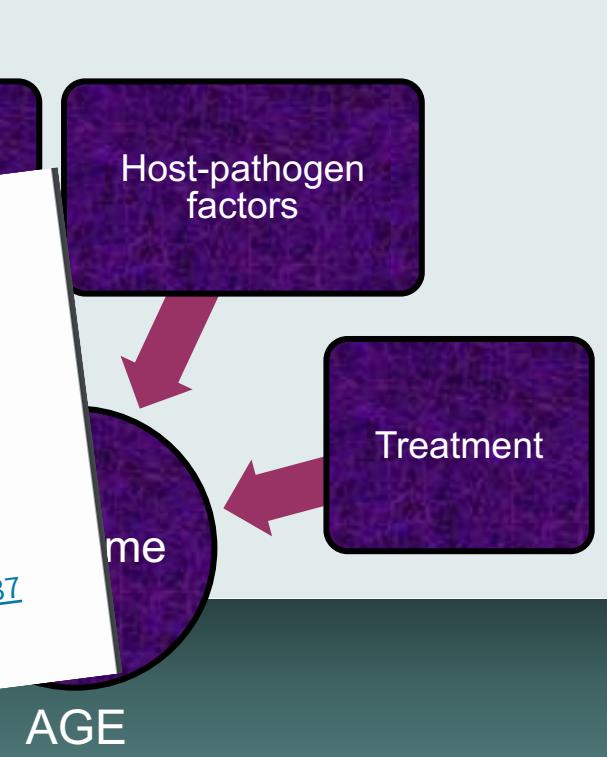
# MRSA Bacteremia

## Can We Improve Upon the Standard of Care?

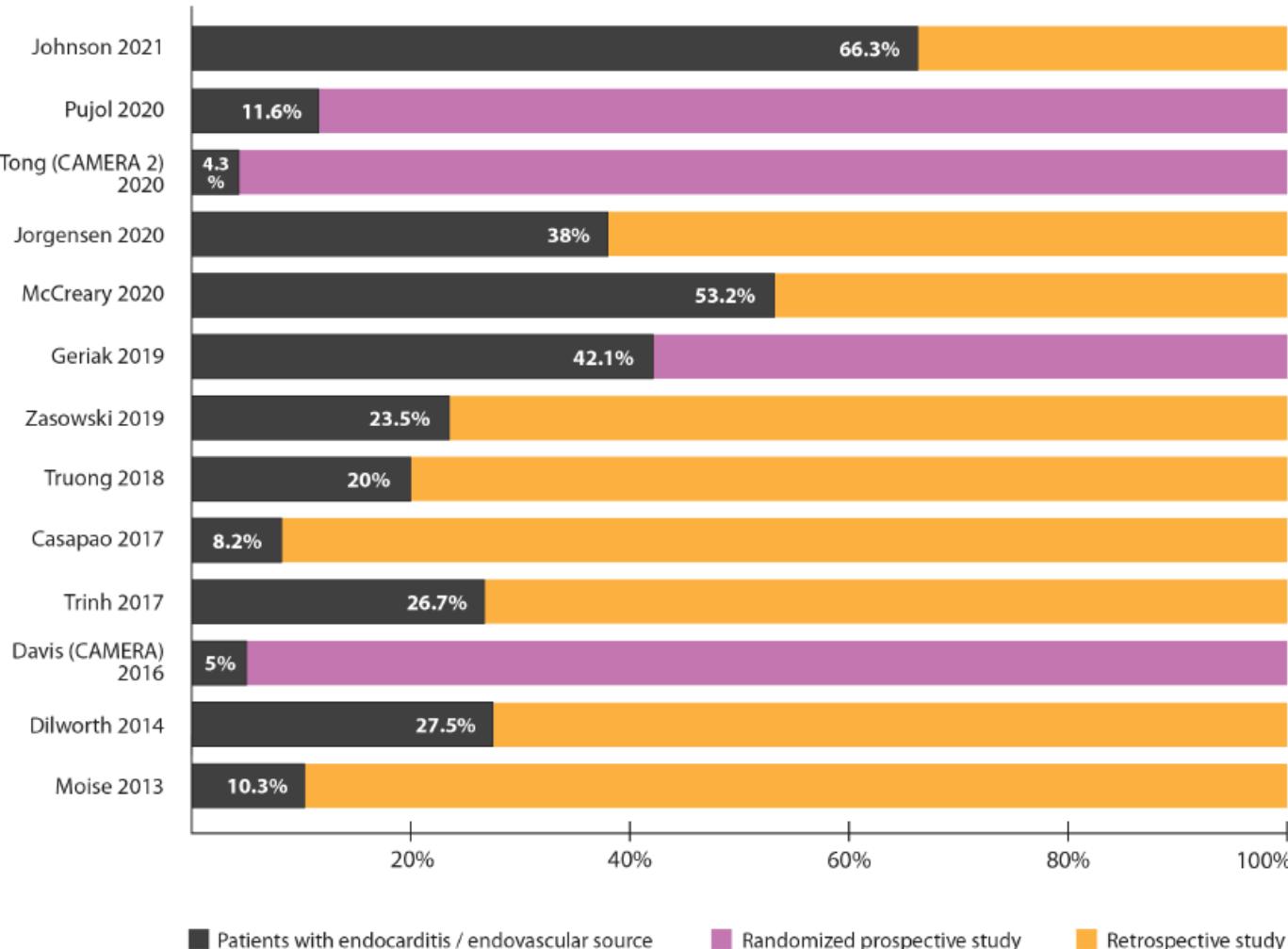


- Assessing
- Mortality
- Treatment
- Bundled care and ID consult

Approaching 65 Years: Is It Time to Consider Retirement of Vancomycin for Treating Methicillin-Resistant *Staphylococcus aureus* Endovascular Infections?  
Warren Rose,<sup>1</sup> Cecilia Volk,<sup>1</sup> Thomas J. Dilworth,<sup>2</sup> and George Sakoulas<sup>3</sup>  
<sup>1</sup>School of Pharmacy, University of Wisconsin–Madison, Madison, Wisconsin, USA, <sup>2</sup>Department of Pharmacy Services, Advocate Aurora Health, Milwaukee, Wisconsin, USA, and <sup>3</sup>Division of Host-Microbe Systems and Therapeutics, Center for Immunity, Infection and Inflammation, University of California, San Diego School of Medicine, La Jolla, California, USA  
<https://doi.org/10.1093/ofid/ofac137>



# Difficulty in applying combination clinical study results to your patients



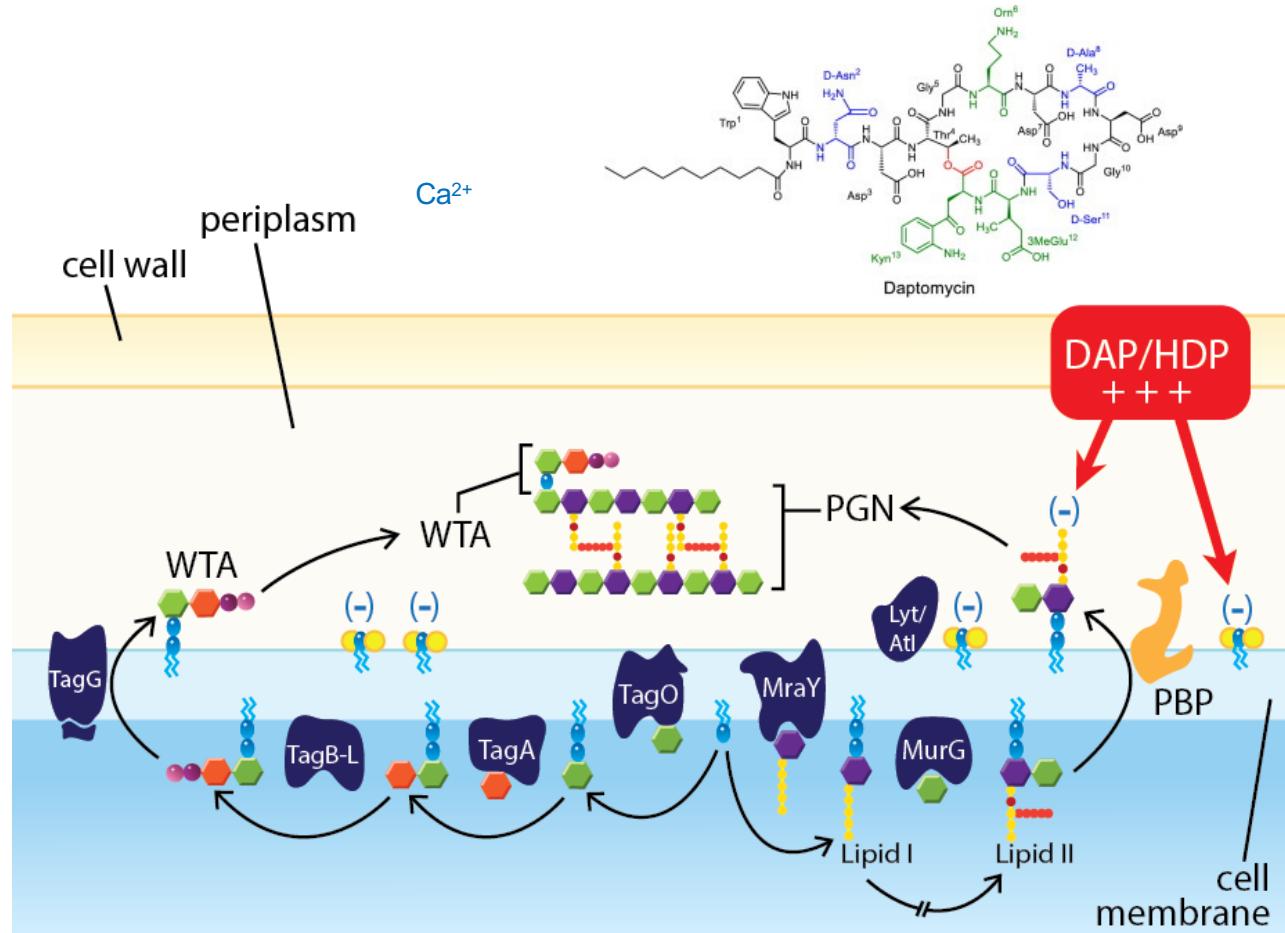
## Definitive IE among clinical studies of MRSA combination therapy:

Prospective clinical trials	Mean 15.8%
	Median 11.6%
Retrospective cohort studies	Mean 30.4%
	Median 26.7%



# Daptomycin (DAP) Mechanism and Resistance

lipopeptide antibiotic that is a functional cation

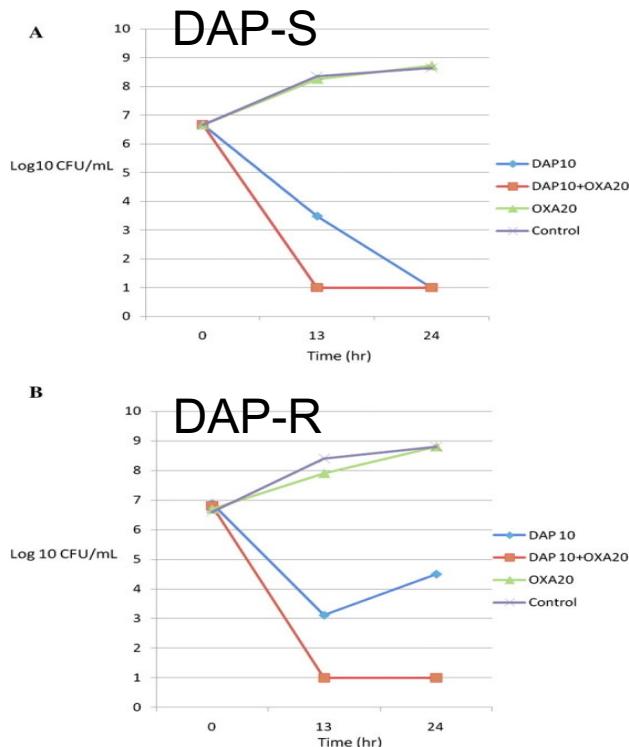
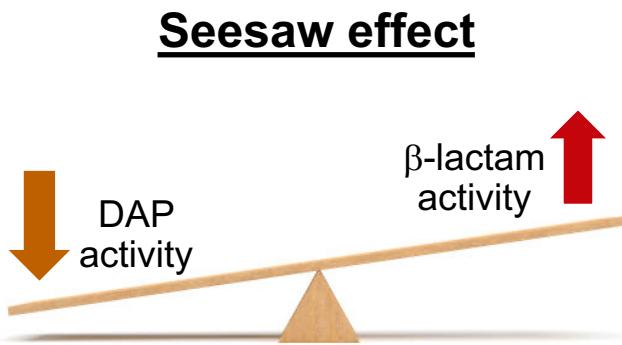


Host Defense Peptides (HDP): LL-37, tPMP (Thrombin-induced platelet microbial protein), HNP1 (human neutrophil peptide)

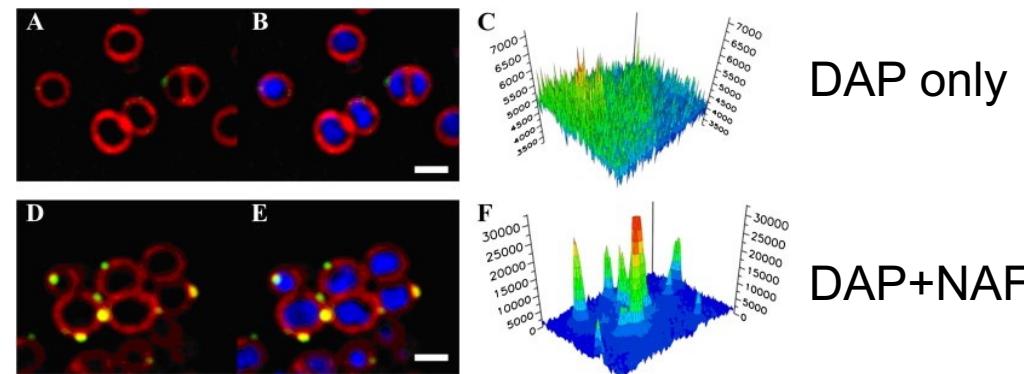


# B-lactam effects on DAP in MRSA

Use of antistaphylococcal  $\beta$ -lactams to increase daptomycin activity in eradicating persistent MRSA



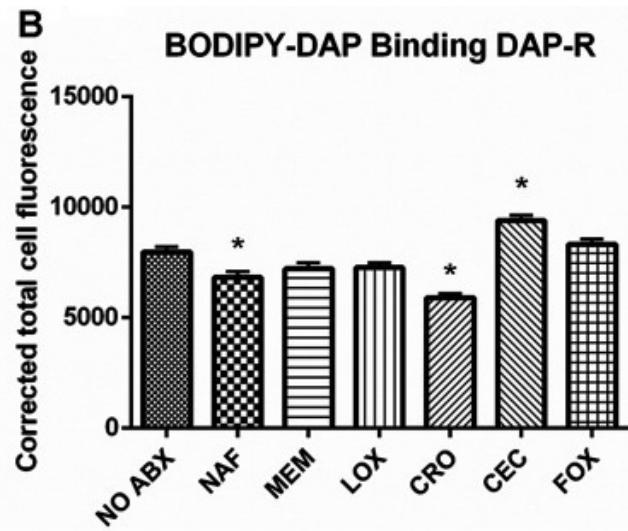
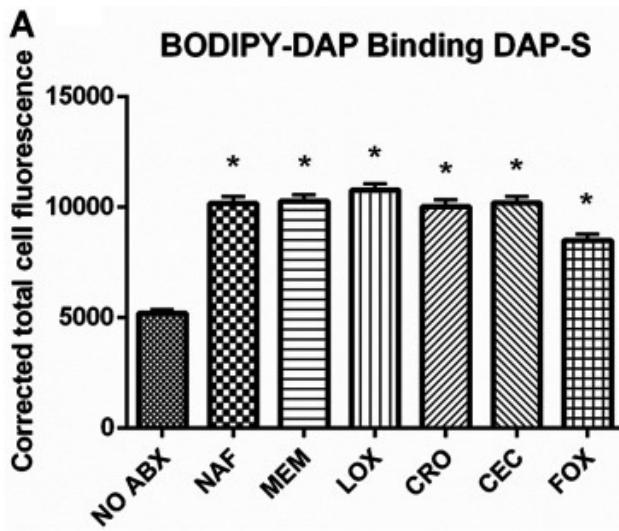
Incorporation of fluorescently-labelled DAP grown in the presence or absence of nafcillin



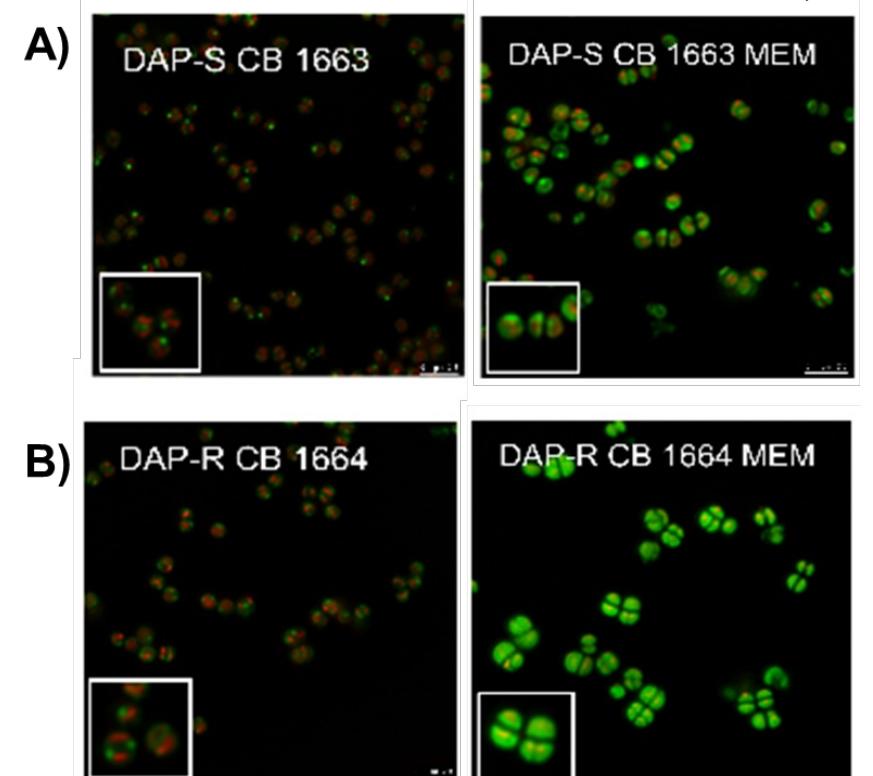
# But its more than just DAP binding



Cassandra Lew, PhD

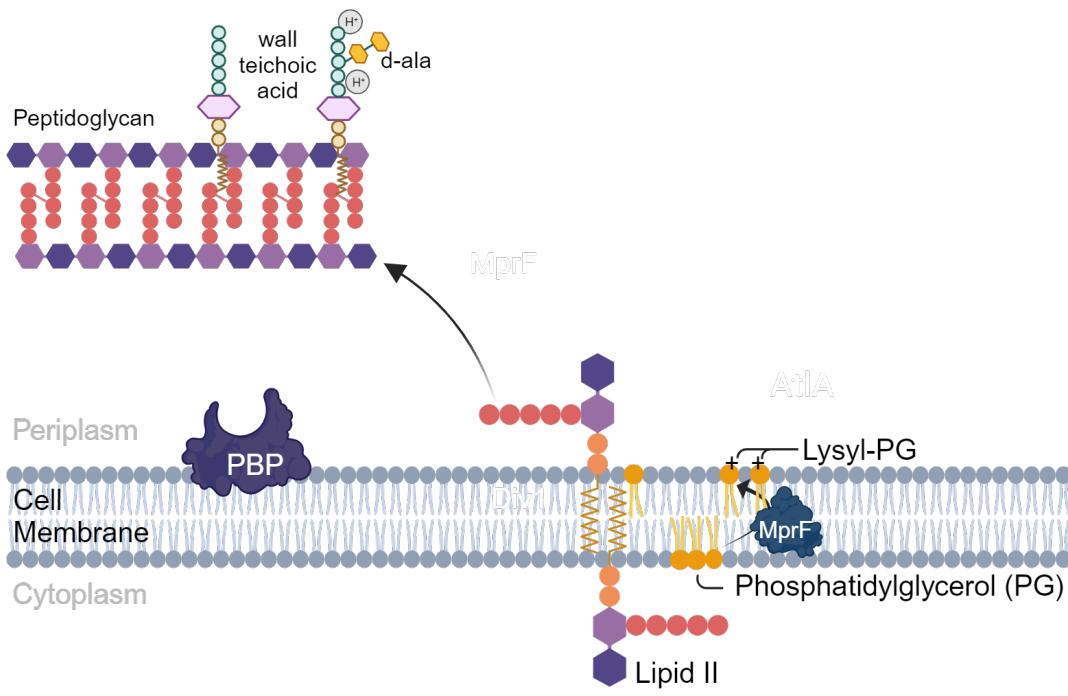


DAP binding NOT universally increased with  $\beta$ -lactams



$\beta$ -lactam conditioning increases cardiolipin distribution (NAO in green)



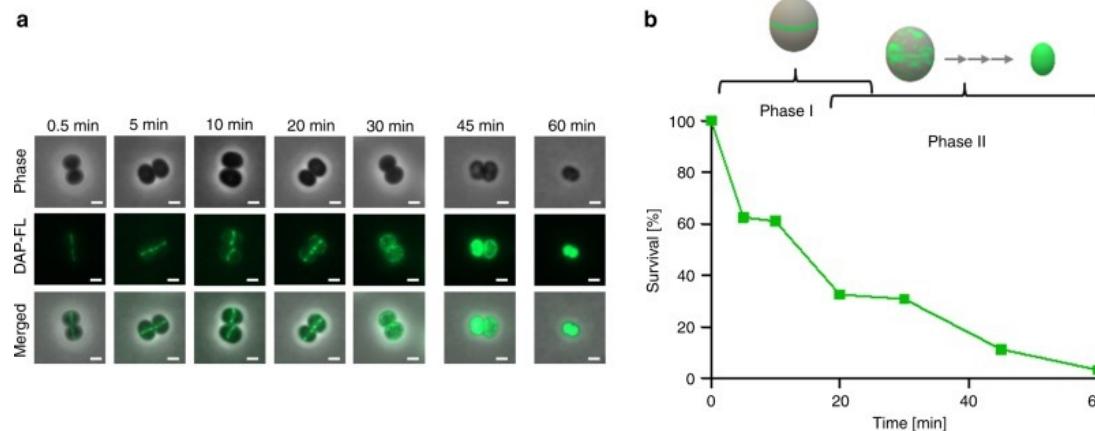


# DAP forms a tripartite complex with lipid II and PG and inhibits cell wall biosynthesis *in vitro*

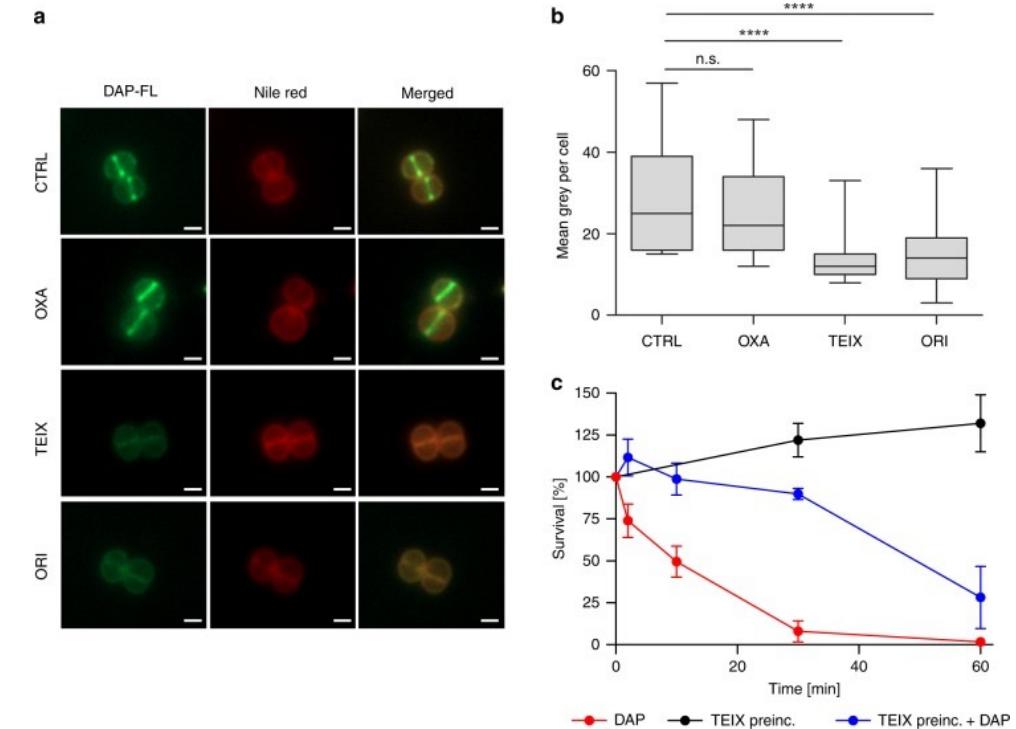


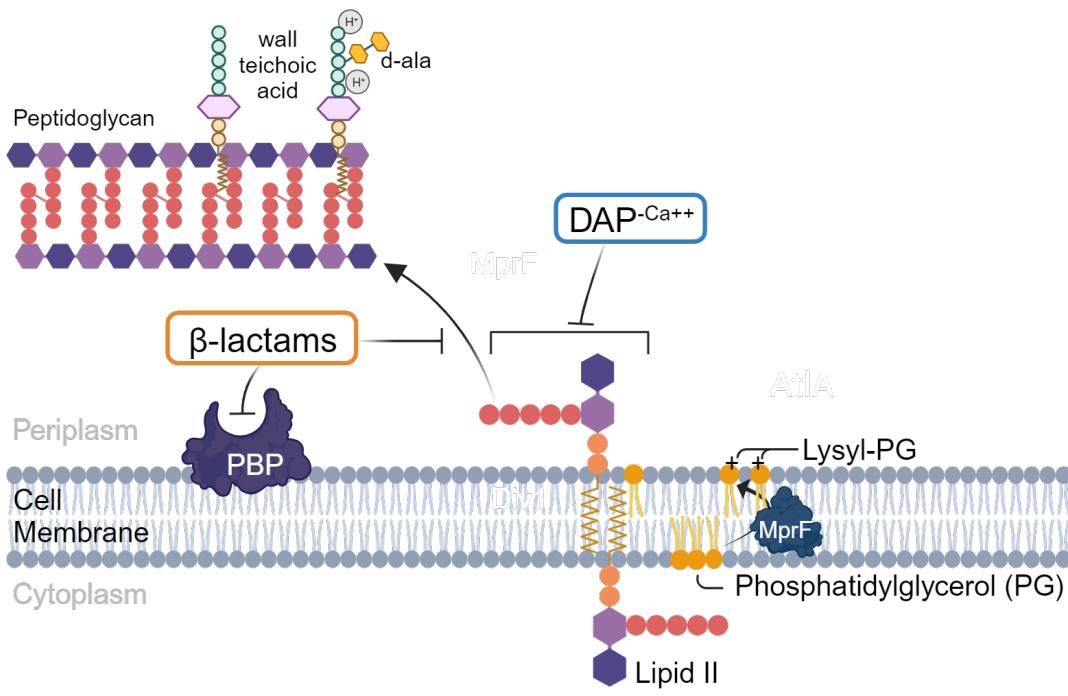
Tanja Schneider, PhD

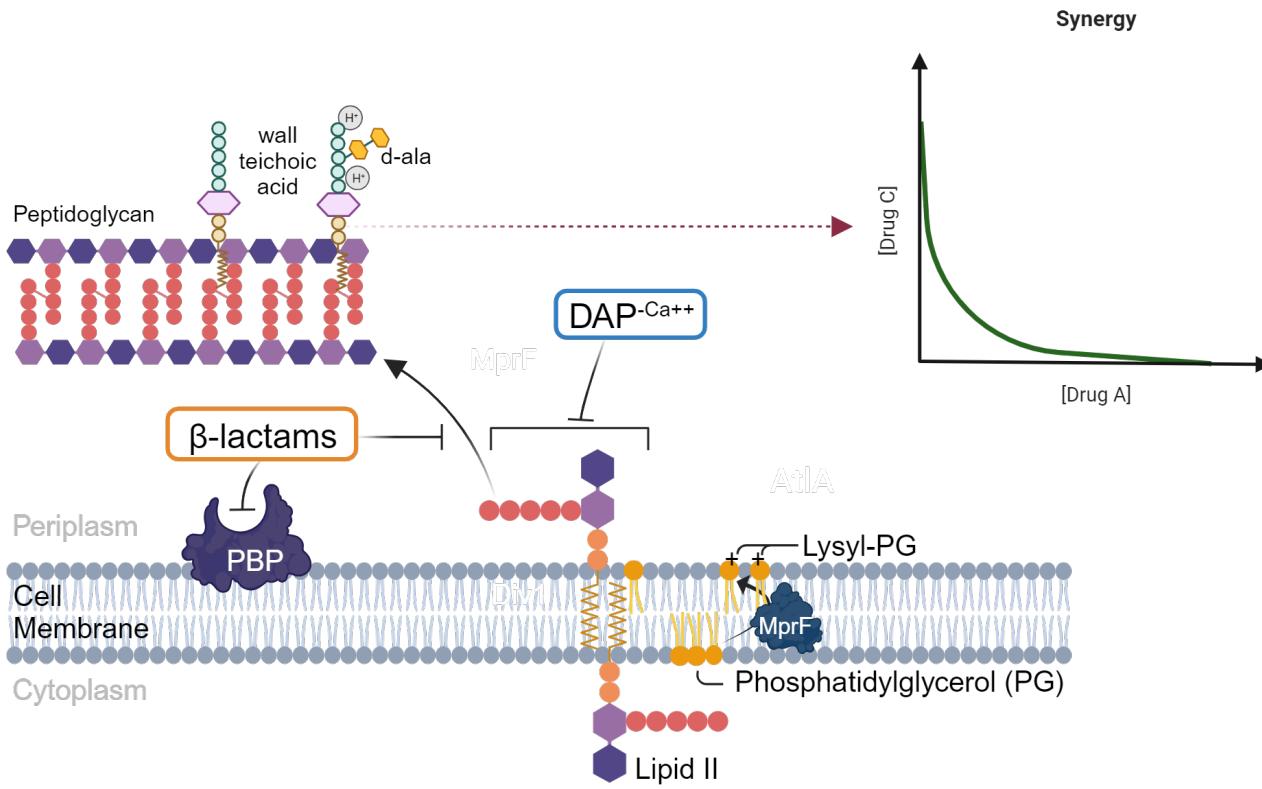
DAP binds to *S. aureus* in a biphasic manner.



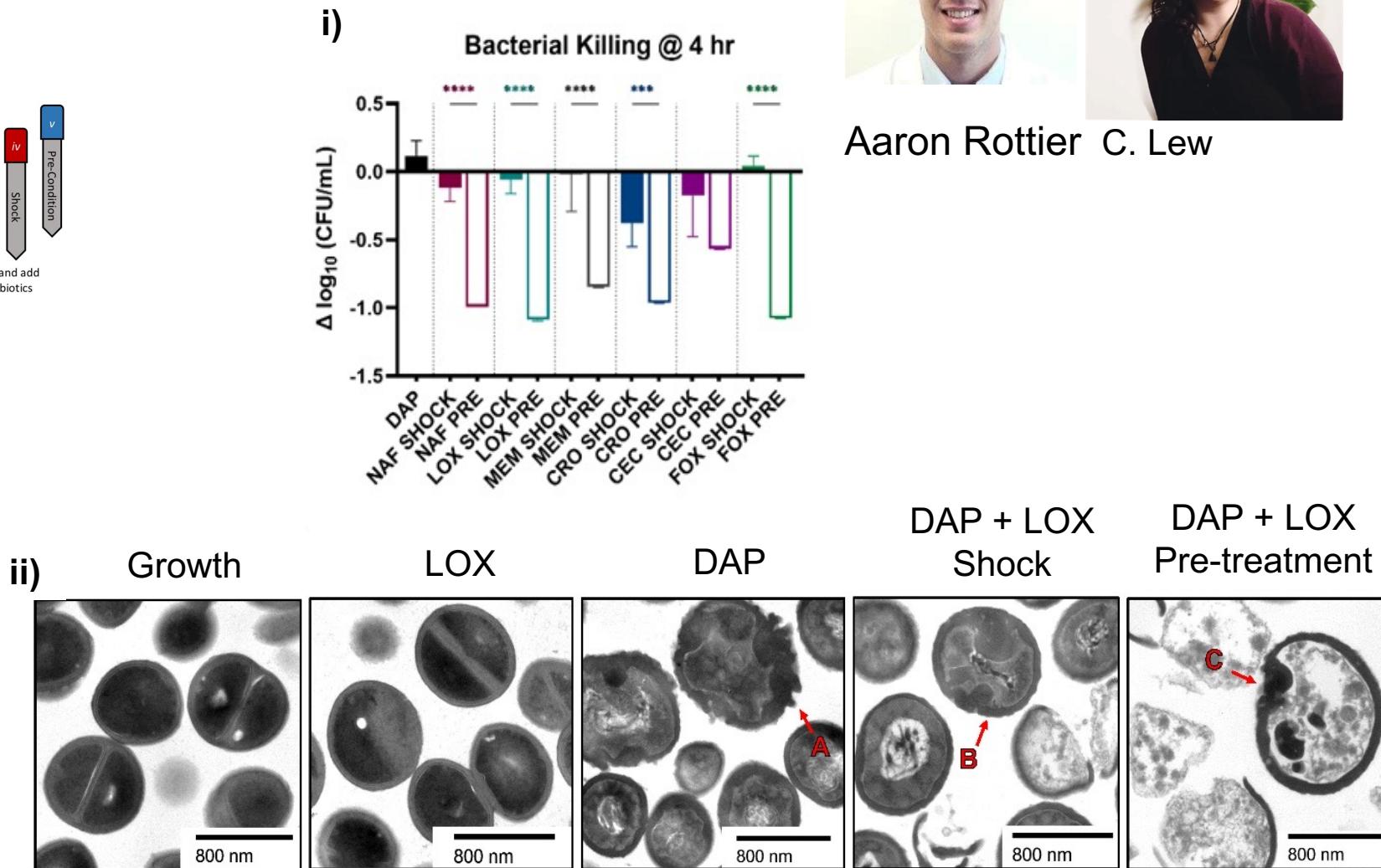
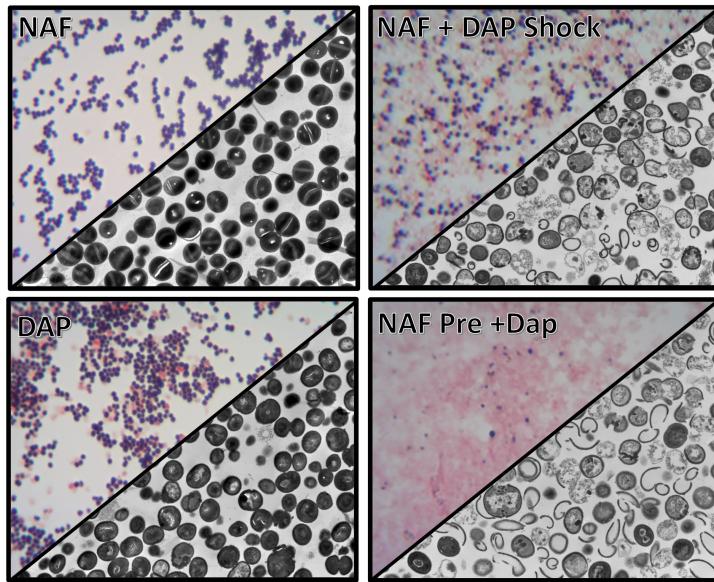
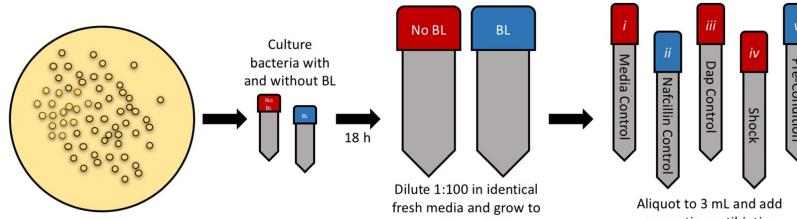
Cell wall precursor lipid II DAP binding.



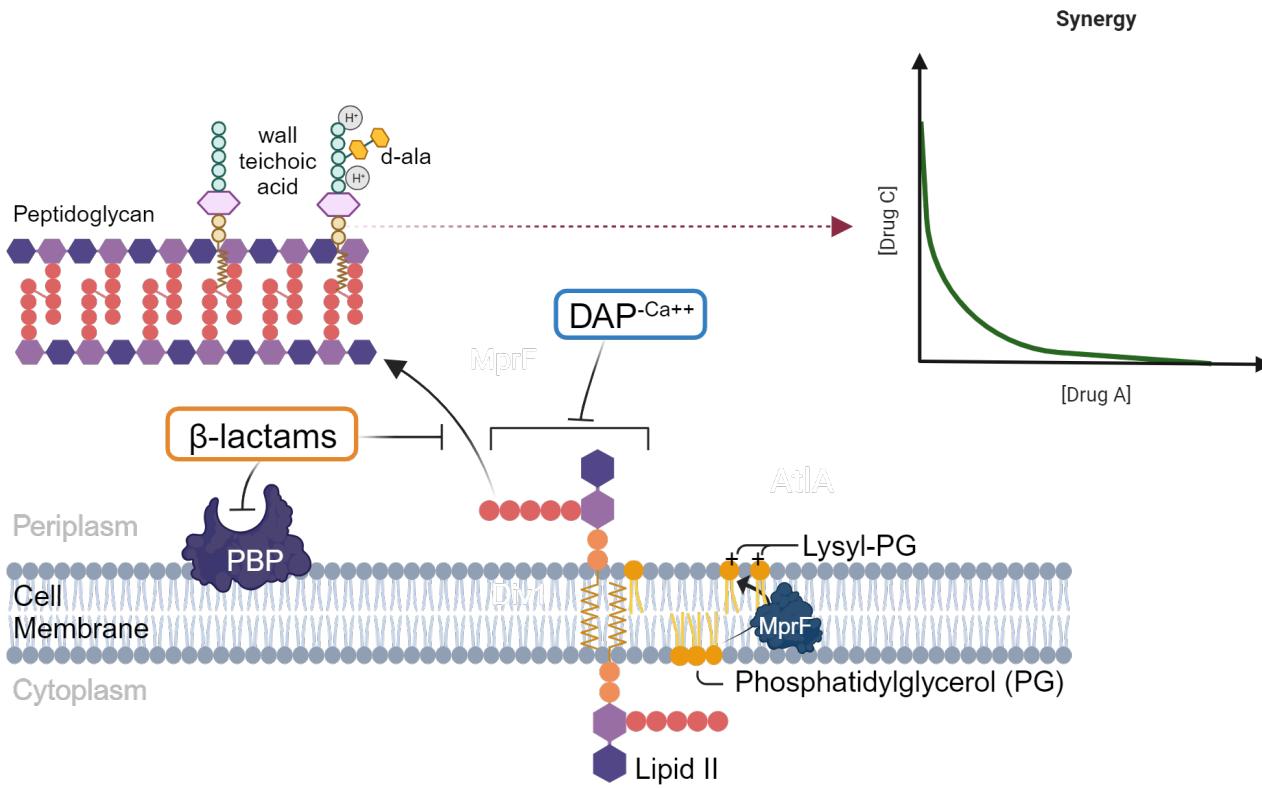


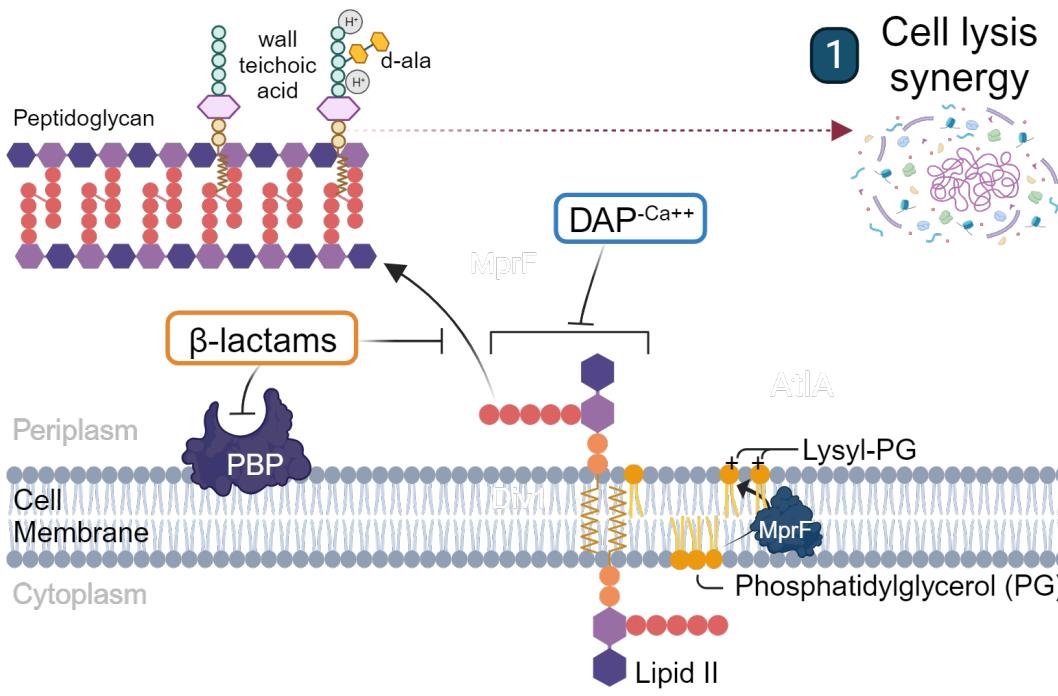


# B-lactam induced cell lysis with DAP vs MRSA



Aaron Rottier C. Lew

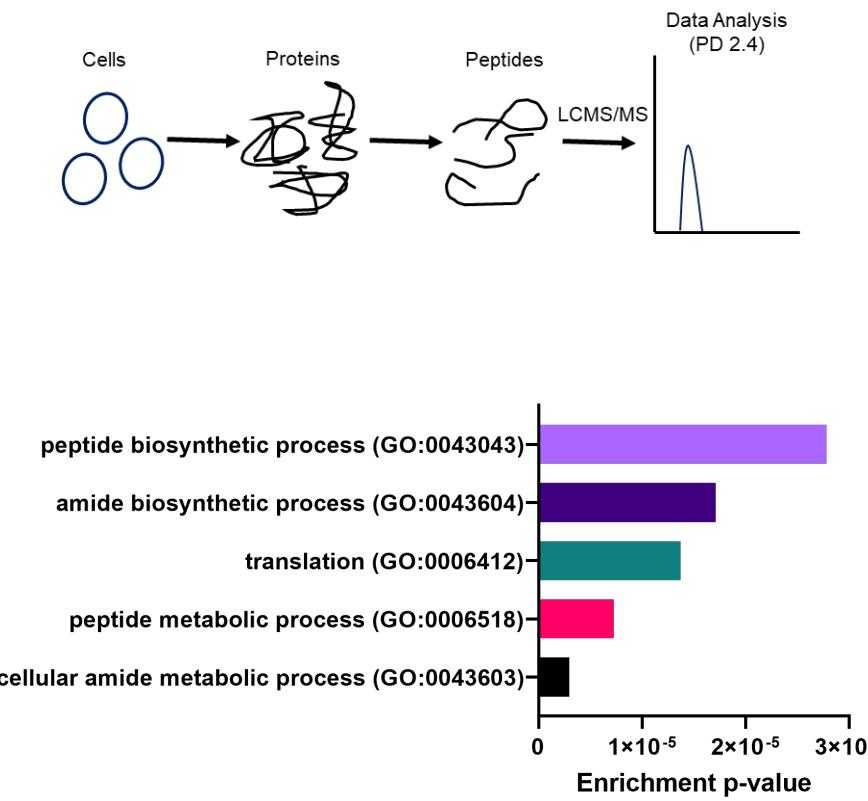




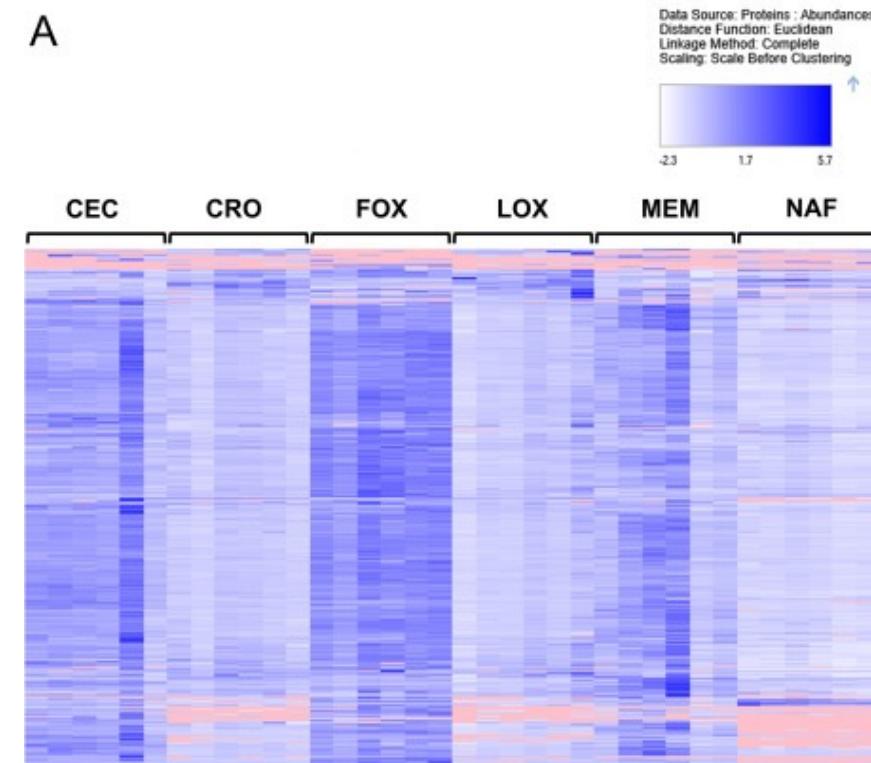
# Proteomic correlates of DAP+β-lactam synergy



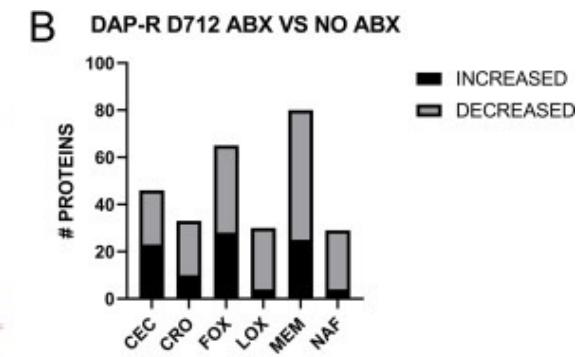
C. Lew



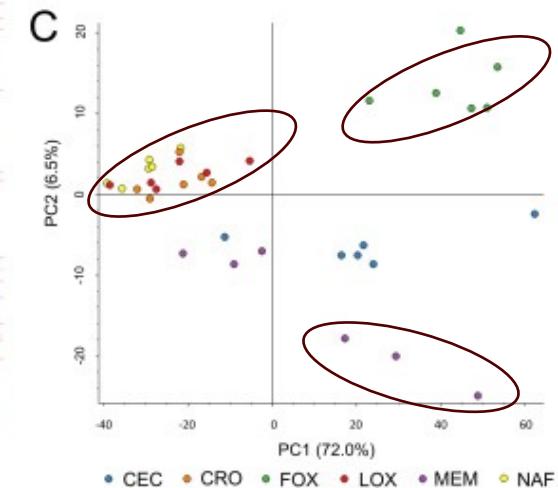
A



B



C

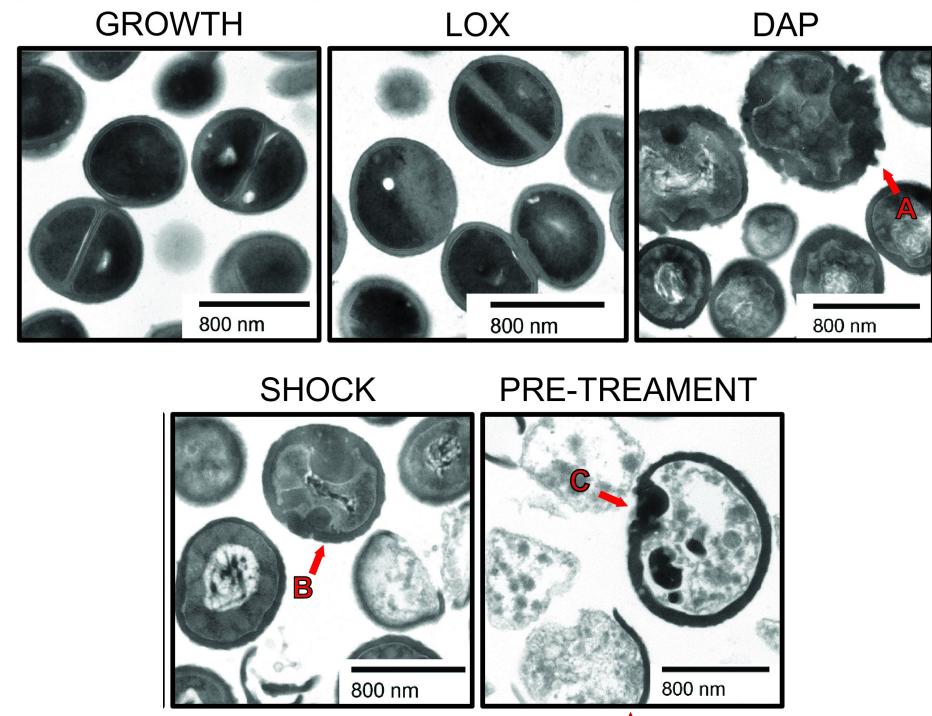


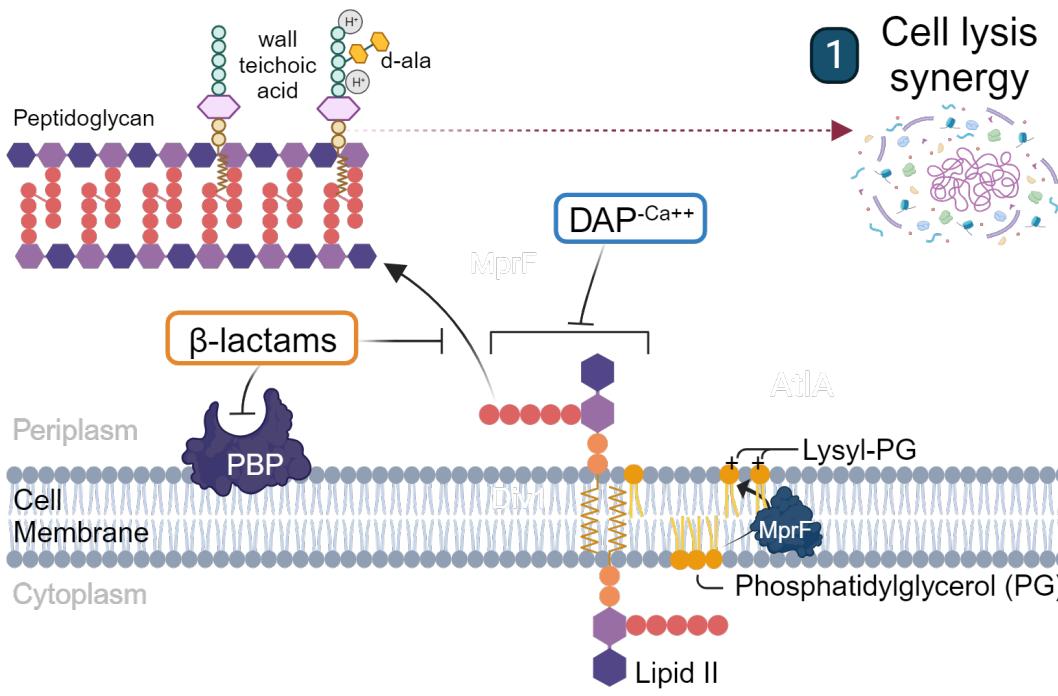
# Autolysins induced with $\beta$ -lactam preconditioning

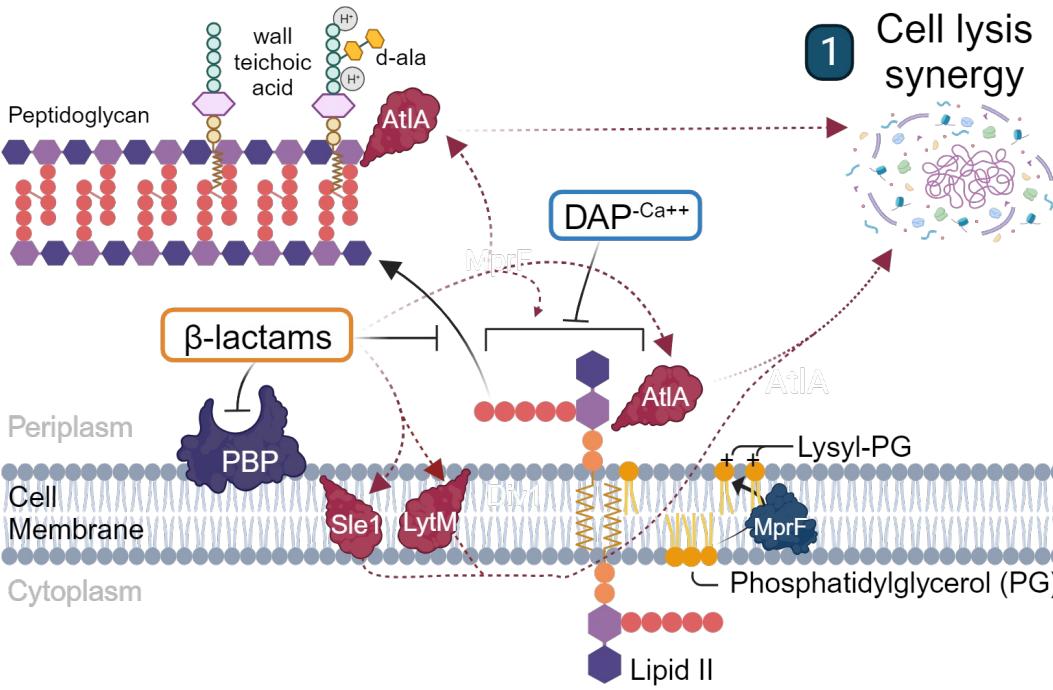
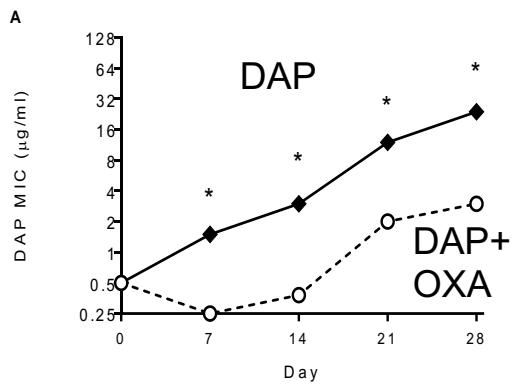
Identified autolysins with significant abundance ratios in different conditions<sup>a</sup>

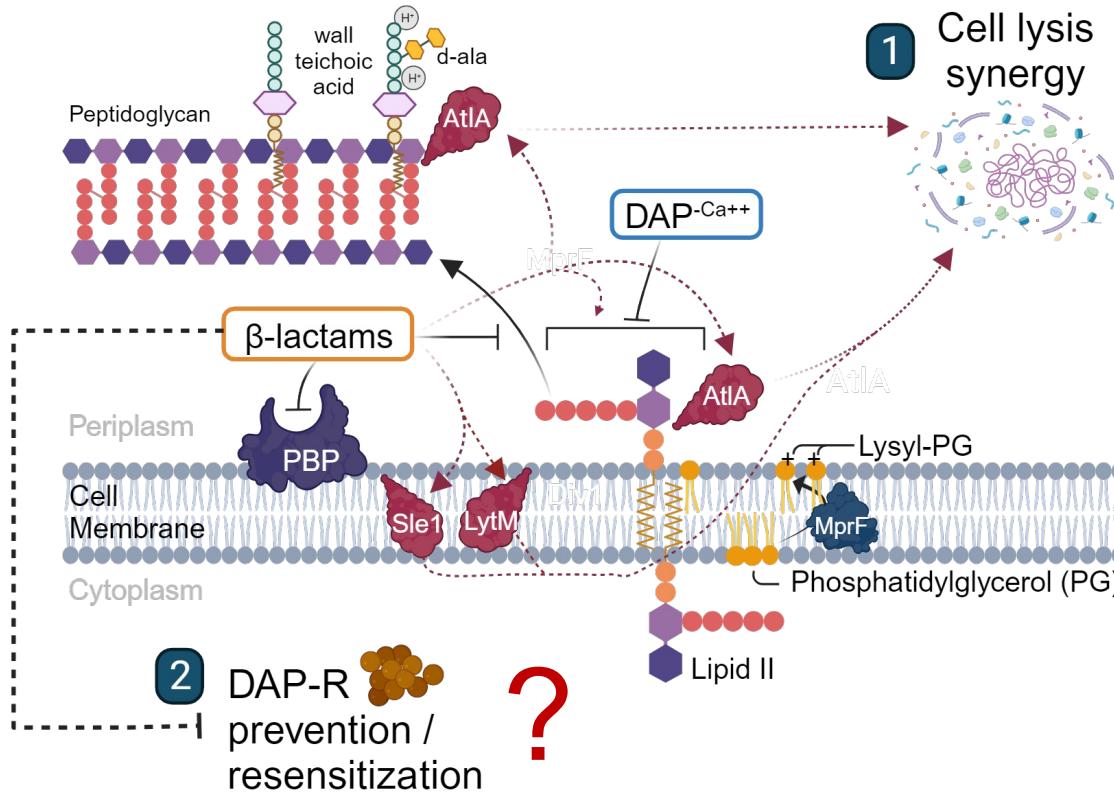
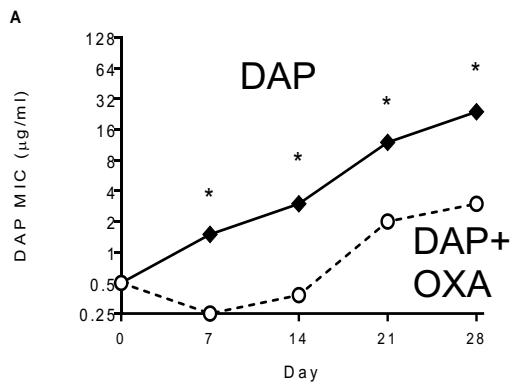
Condition with or without D712	ATLA	ISAA	LYTM	SCEd	sLE1	sSAa
NAF	0.764	<b>0.344</b>	0.060	<b>0.167</b>	0.764	0.257
MEM	<b>0.224</b>	<b>0.138</b>	0.010	<b>0.083</b>	<b>0.140</b>	<b>0.211</b>
LOX	0.933	0.693	—	0.881	0.663	0.664
CRO	0.587	0.413	0.682	0.563	0.452	0.393
CEC	0.271	0.224	0.160	0.184	0.209	0.215
FOX	<b>0.222</b>	<b>0.133</b>	0.010	<b>0.040</b>	0.284	<b>0.178</b>

<sup>a</sup>Values of <1 indicate increased abundance with  $\beta$ -lactam pretreatment (boldface indicates a significant ratio). —, protein not identified.

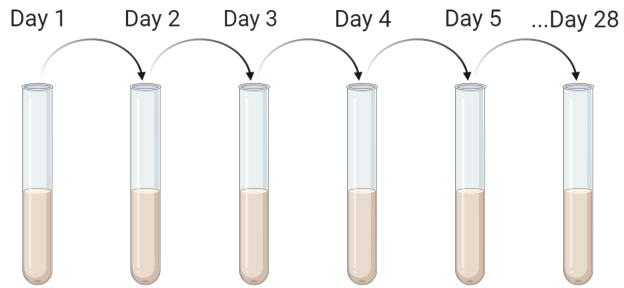




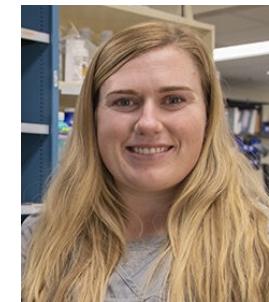
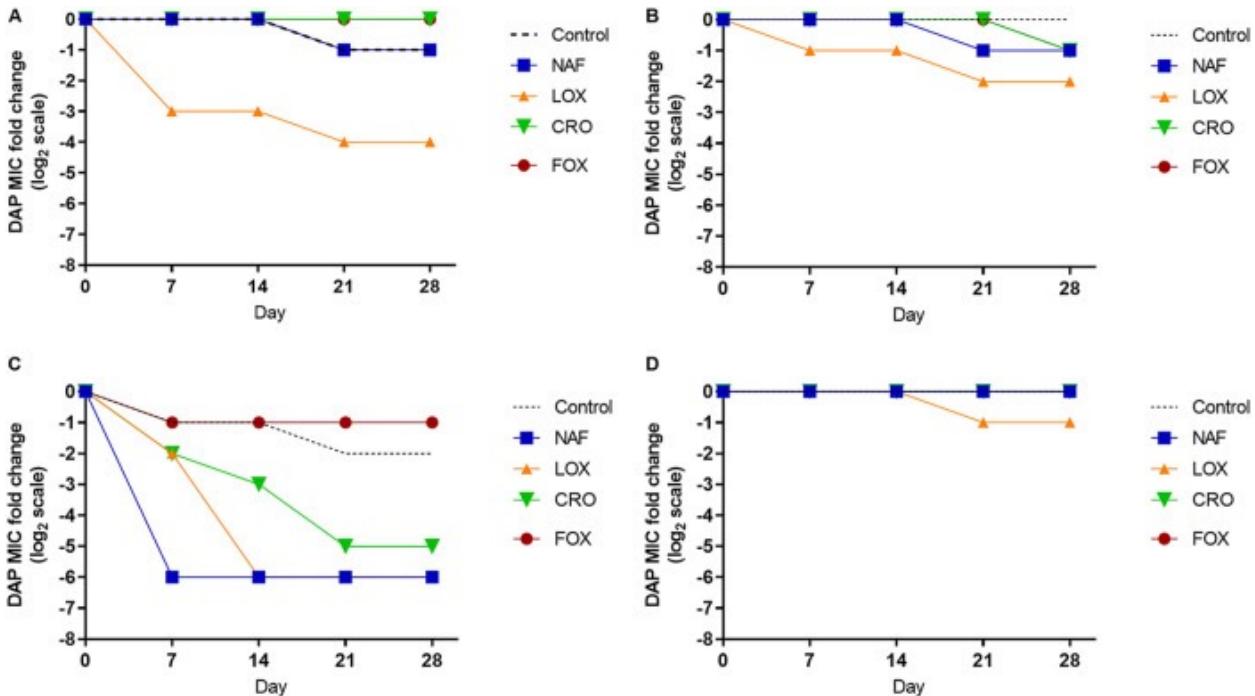




# B-lactam induced DAP-resensitization



Isolate	Passage	Replicate	DAP MIC (mg/liter)	$\beta$ -lactam MIC (mg/liter) <sup>b</sup>	<i>mprF</i> SNP	MprF domain	<i>div1b</i> mutation
J01	None		0.5				
J03	None		2		T <sub>345</sub> I	Bifunctional	
	Media	i	2			None	
	Media <sup>a</sup>	ii	1		Y <sub>325</sub> H	Bifunctional	
	Media <sup>a</sup>	iii	1		R <sub>437</sub> P	Synthase	
CRO <sup>c</sup>	ii		0.75	512	V <sub>152</sub> G	Translocase	
LOX <sup>c</sup>	ii		0.125	32	R <sub>788</sub> L	Synthase	Q <sub>425</sub> <sup>d</sup>
LOX <sup>c</sup>	iii		0.125	32	R <sub>788</sub> L	Synthase	Q <sub>415</sub> <sup>d</sup>



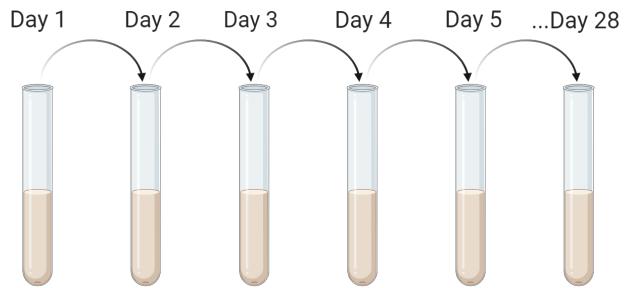
Rachel Jenson,  
PharmD



Benjamin Howden,  
MD, PhD,



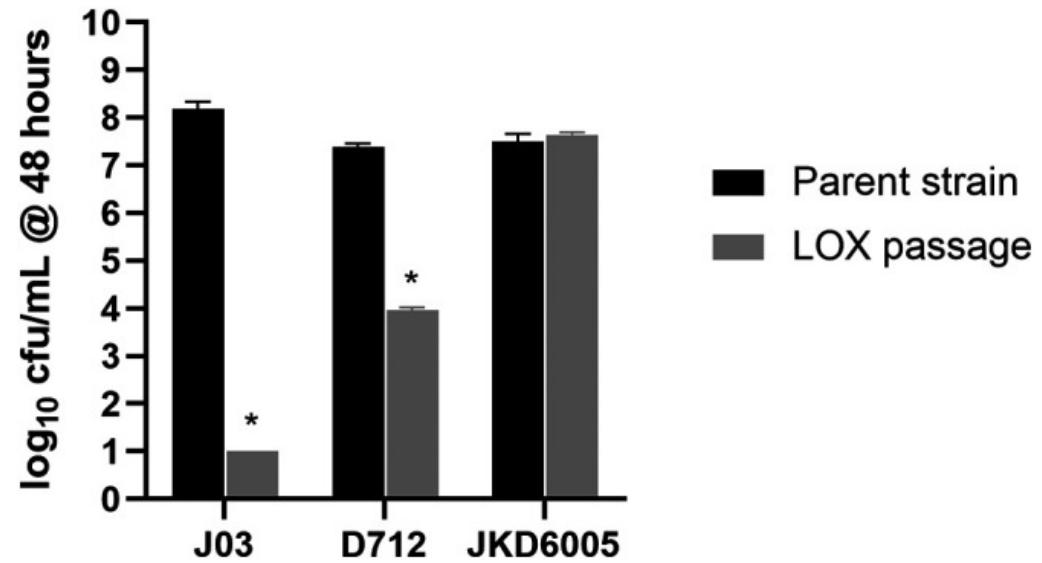
# B-lactam induced DAP-resensitization



## Rapid conversion to DAP-S

Isolate	Passage	Replicate	Additional <i>mprF</i> SNP frequency at day:			
			7	14	21	28
J03 <sup>b</sup>	LOX	i	0.58	0.74	0.11	0.00 <sup>c</sup>
J03 <sup>b</sup>	LOX	ii	0.11	0.08	0.78	0.98
J03 <sup>b</sup>	LOX	iii	0.51	0.61	0.79	0.99
D712 <sup>d</sup>	LOX	i	0.91	0.67	0.86	1.00
D712 <sup>d</sup>	LOX	ii	1.00	1.00	1.00	0.99
D712 <sup>d</sup>	LOX	iii	0.00	0.00	0.00	0.00

Highly susceptible to DAP killing



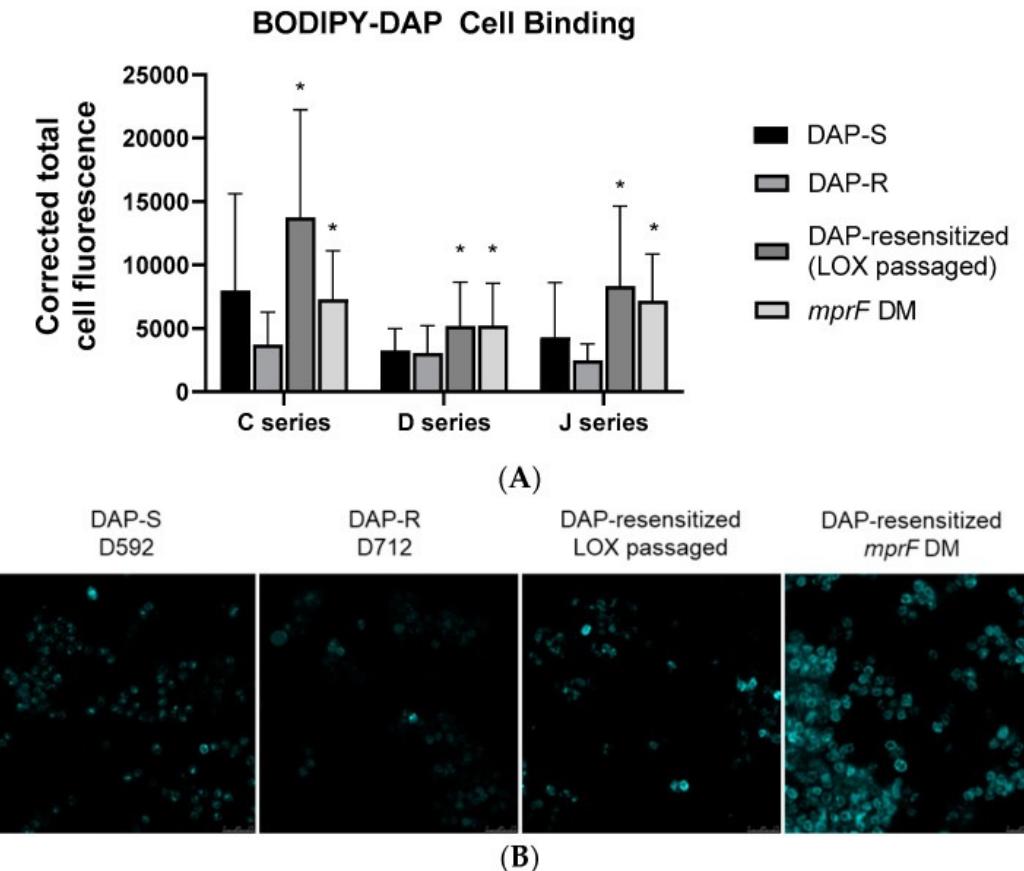
# Secondary mutations in *mprF* lead to DAP-resensitization

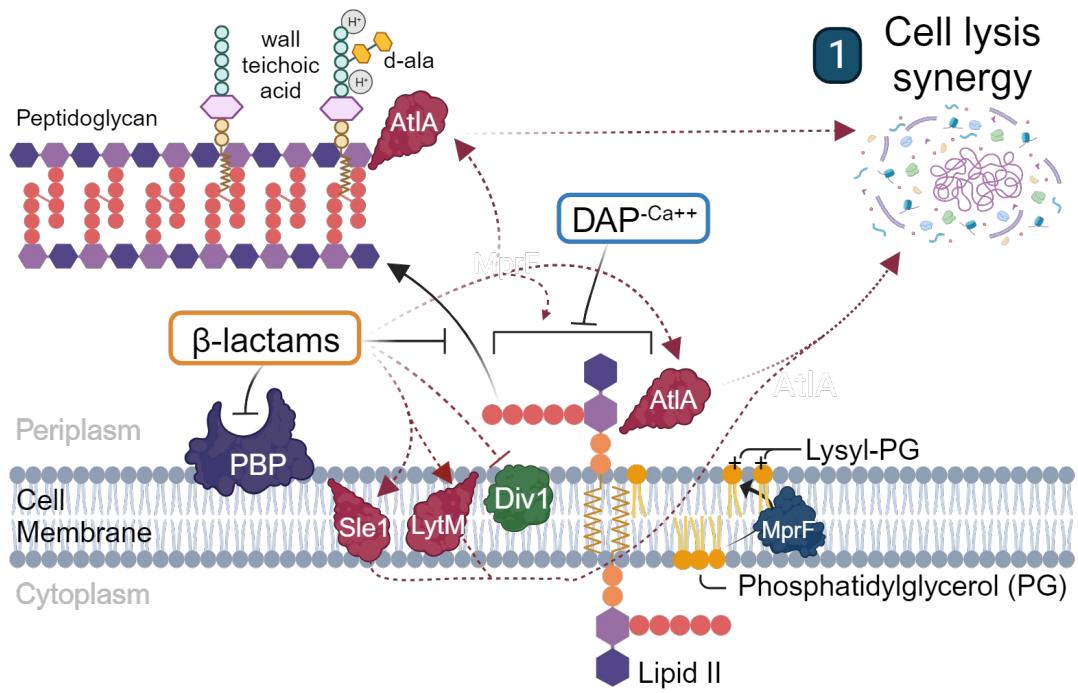


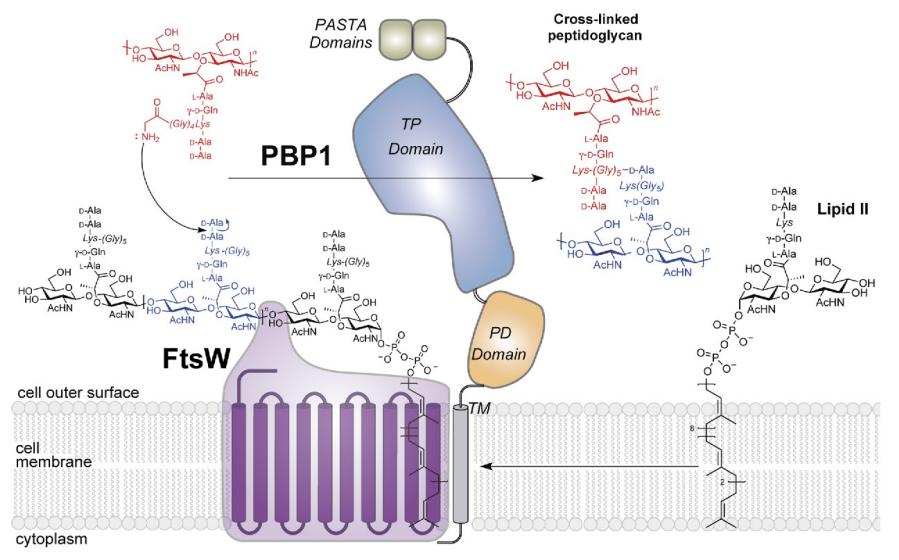
Nagendra Mishra PhD  
B. Howden PhD

Strain Set <sup>a</sup>	Strain Name	Strain Description	DAP MIC <sup>b</sup> ( $\mu\text{g/mL}$ )	LOX MIC <sup>c</sup> ( $\mu\text{g/mL}$ )	SNPs in <i>mprF</i> <sup>d</sup>
I	C24	DAP-S	0.5	8	WT
	C25	DAP-R	2	4	S295L
	C25-LOX	DAP-resensitized (LOX passaged)	<0.125	8	S295L + L84 (Translocase domain)
	C25, <i>mprF</i> DM	Secondary <i>mprF</i> mutation (L84 <sup>e</sup> ) introduced into C25	0.125	16	S295L + L84 <sup>e</sup>
II	D592	DAP-S	0.5	512	WT
	D712	DAP-R	2	512	L341S
	D712-LOX	DAP-resensitized (LOX passaged)	0.5	1024	L341S + S136L (Translocase domain)
	D712, <i>mprF</i> DM	Secondary <i>mprF</i> mutation (S136L) introduced into D712	0.5	1024	L341S + S136L
III	J01	DAP-S	0.5	16	WT
	J03	DAP-R	2-4	2	T345I
	J03-LOX	DAP-resensitized (LOX passaged)	0.125	32	T345I + R788L Synthase domain
	J03, <i>mprF</i> DM	Secondary <i>mprF</i> mutation (R788L) introduced into J03	0.125	16	T345I + R788L

<sup>a</sup> Sets of isolates are represented by alternative shading and no shading, with the first strain in each set being the DAP-S parental strain, the second in each set being the DAP-R or allelic exchange, respectively; <sup>b,c,d</sup> Data in this table have been previously published (41); <sup>e</sup> nonsense mutation (41).

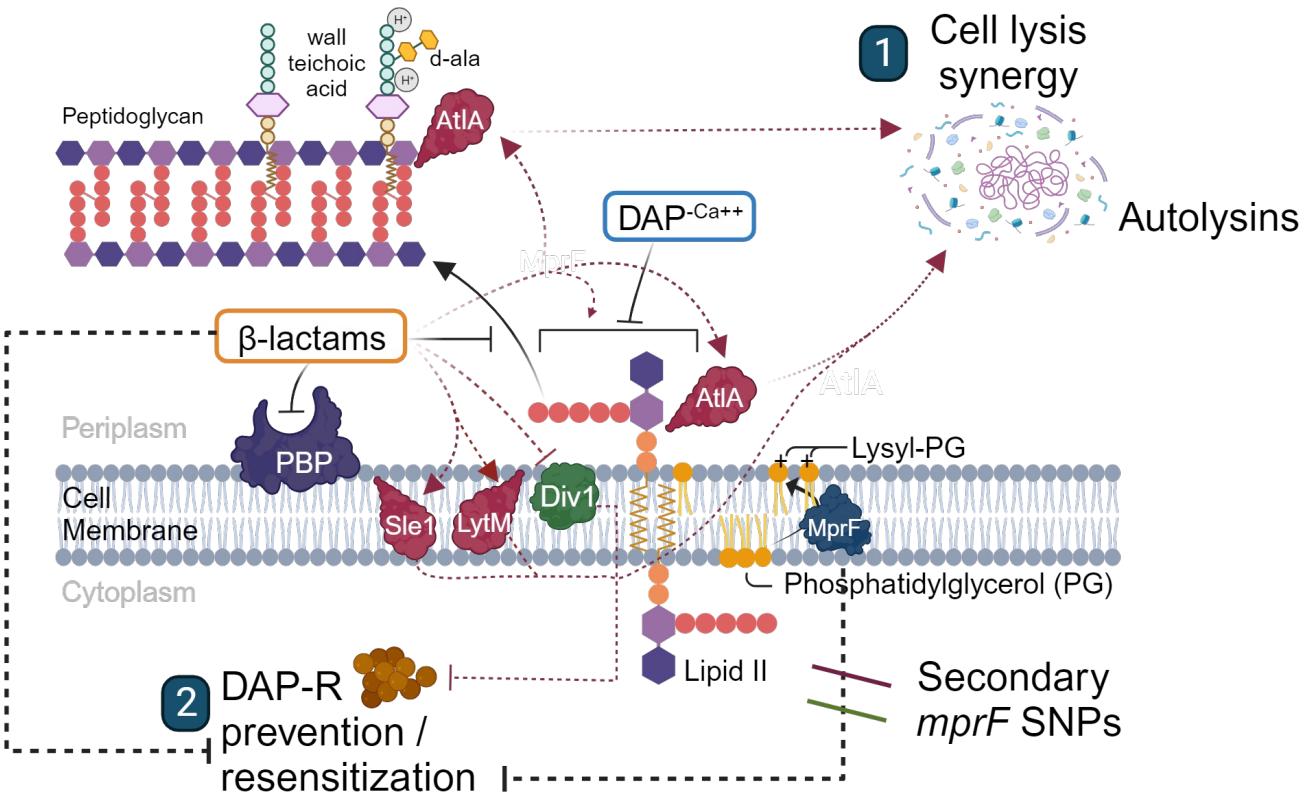




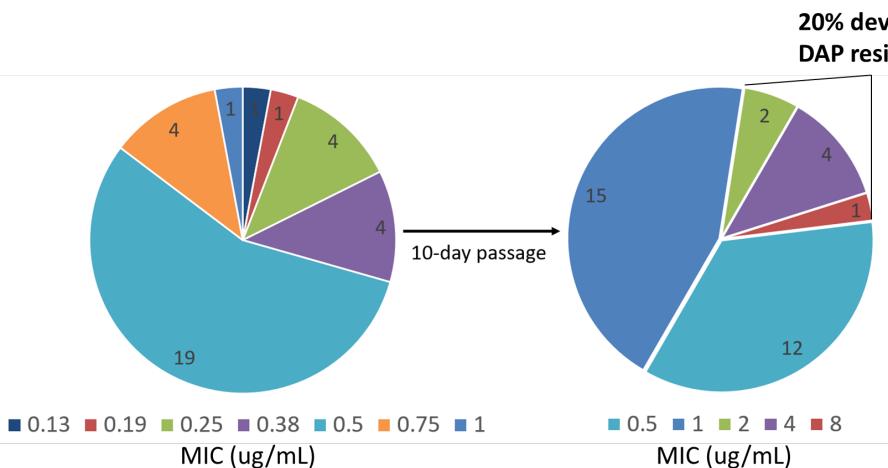
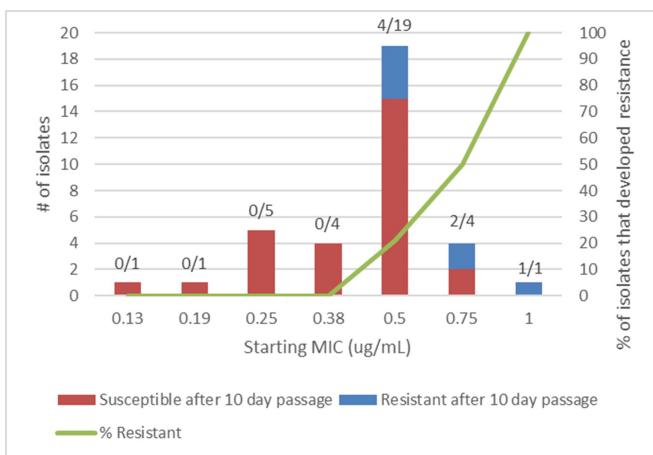
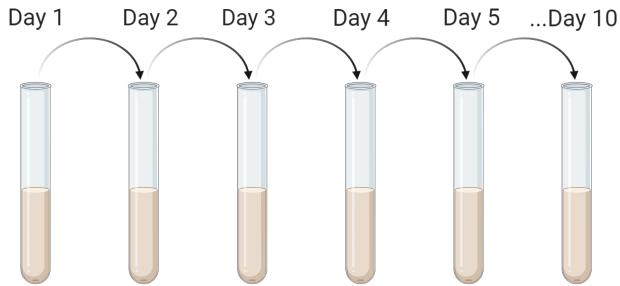


Martínez-Caballero, S et al. Comput Struct Biotechnol J. 2021 Sep 17:19:5392-5405.

### Divisome Proteins



# Select rapid DAP-R emergence: implications for combination $\beta$ -lactams therapy?



Article

## Membrane Phenotypic, Metabolic and Genotypic Adaptations of *Streptococcus oralis* Strains Destined to Rapidly Develop Stable, High-Level Daptomycin Resistance during Daptomycin Exposures

Nagendra N. Mishra <sup>1,2,\*</sup>, Rodrigo de Paula Baptista <sup>3,4,5</sup>, Truc T. Tran <sup>3,4,5</sup>, Christian K. Lapitan <sup>1</sup>, Cristina Garcia-de-la-Maria <sup>6,7</sup>, Jose M. Miró <sup>7</sup>, Richard A. Proctor <sup>8</sup> and Arnold S. Bayer <sup>1,2</sup>



Cecilia Volk, PharmD

### *mprF* mutations

In Vivo Mutation (occurrences)	In Vitro Mutation (occurrences)
L826F (10)	S295L (2)
L341S (4)	L341S (1)
S295L (3)	
T345I (3)	
M347R (2)	
S337L (2)	



# Future Directions



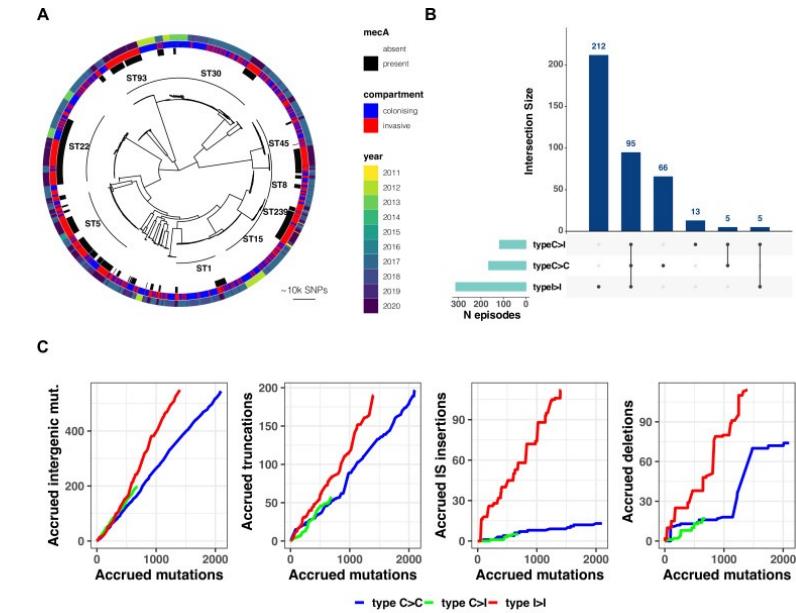
RESEARCH ARTICLE



B. Howden

## Niche-specific genome degradation and convergent evolution shaping *Staphylococcus aureus* adaptation during severe infections

Stefano G Giulieri<sup>1,2,3</sup>, Romain Guérillot<sup>1</sup>, Sebastian Duchene<sup>1</sup>, Abderrahman Hachani<sup>1</sup>, Diane Daniel<sup>1,4</sup>, Torsten Seemann<sup>4</sup>, Joshua S Davis<sup>5,6</sup>, Steven YC Tong<sup>6,7</sup>, Bernadette C Young<sup>8</sup>, Daniel J Wilson<sup>9</sup>, Timothy P Stinear<sup>1\*</sup>, Benjamin P Howden<sup>1,2,4\*</sup>

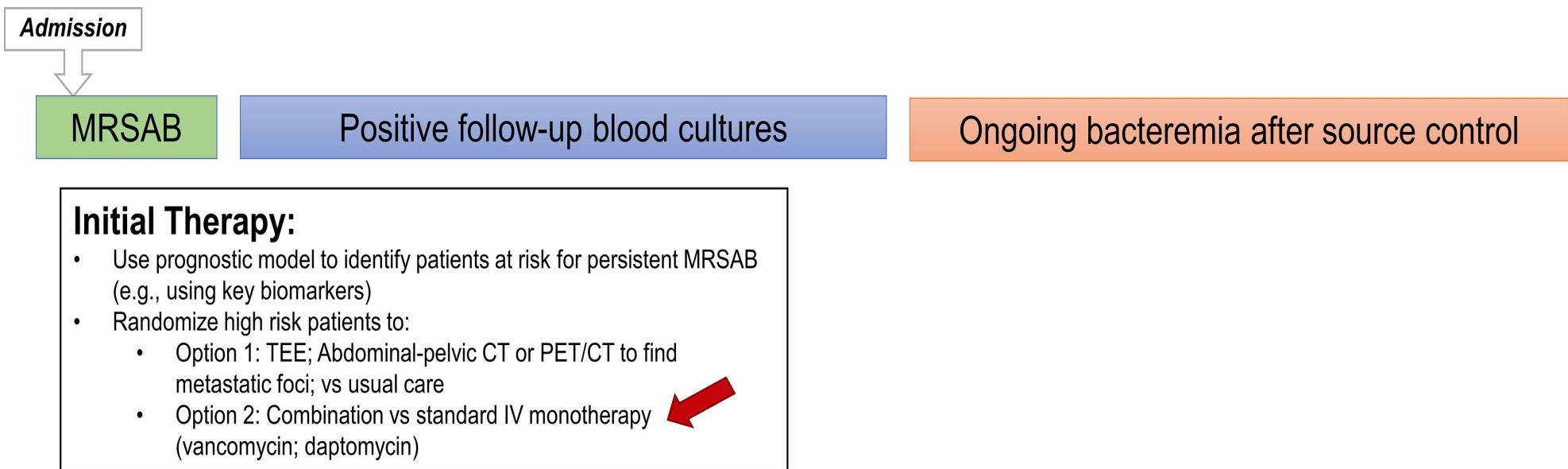


## DAP-β-lactam Combinatorial Effects

Determine the convergent genetic pathway(s) for PBP-1 targeting that can both temporally prevent DAP-R, as well as potentially revert DAP-R-to-DAP-S with β-lactam exposures.



# Considerations for Future Clinical Trials of MRSA Bacteremia



# Acknowledgements



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Infectious Diseases

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**UW ICTR**  
UW Institute for Clinical  
and Translational Research



University of Wisconsin-Madison

## ROSE Lab

Cassandra Lew, Rachel Jenson, Cecelia Volk,  
Dan Smelter, Aaron Rottier, Sue McCrone

### Collaborators

- Arnold Bayer, MD; Nagendra Mishra, PhD; UCLA
- Tanja Schneider, PhD; UBonnn
- George Sakoulas and Victor Nizet – Univ. CA San Diego
- Sanjay Shukla - Marshfield Clinic
- Benjamin Howden and Sarah Baines – Univ. of Melbourne



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School of Pharmacy