



Transport across membranes:  
Multiple drug resistance, mechanisms and new tools  
*Summer School – Bremen – 7-14/07/2007*

# **Molecular bases for antibiotic resistance through efflux**

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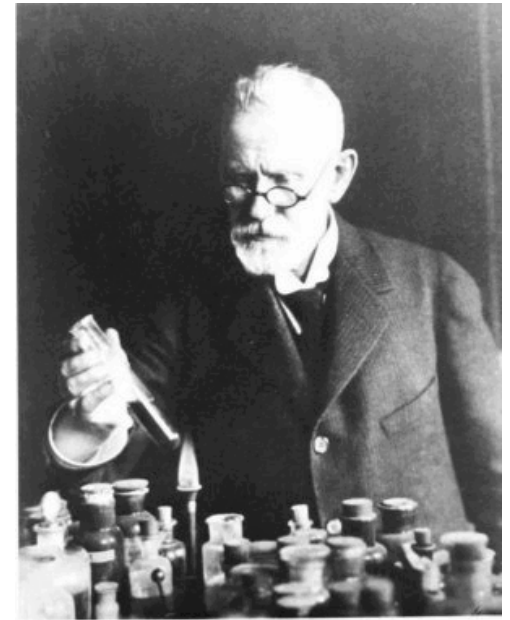
[www.facm.ucl.ac.be](http://www.facm.ucl.ac.be)



# P. Ehrlich, a father of modern chemotherapy

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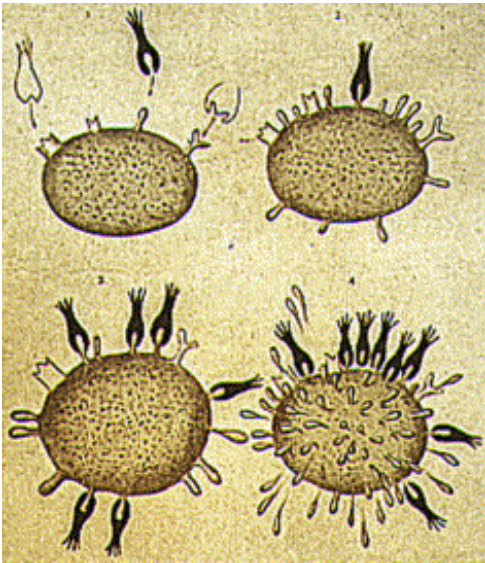
Paul Ehrlich (1854-1945)



*P. Ehrlich*

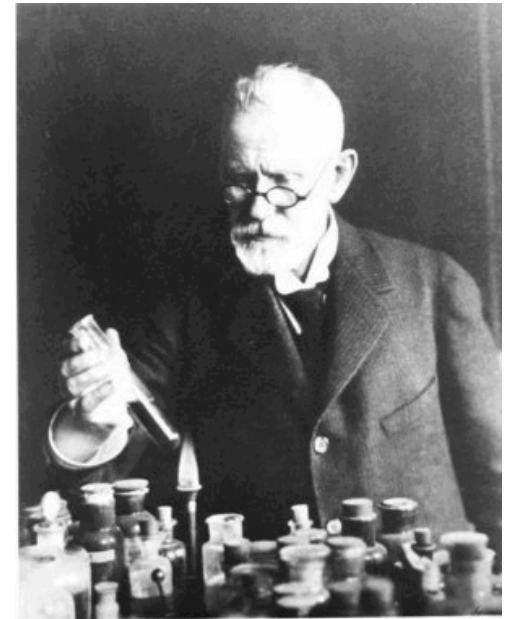
# Magic bullets need to reach their target

Paul Ehrlich (1854-1945)



*"corpora non agunt  
nisi fixata"*

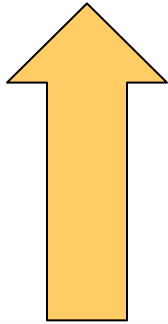
"The goal is ... to find  
chemical substances  
that have special affinities  
for pathogenic organisms  
and that,  
like magic bullets,  
go straight to their targets"



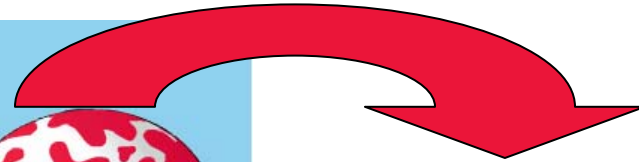
*P. Ehrlich*

# Chemotherapeutic agents exert toxic effects on specific target cells

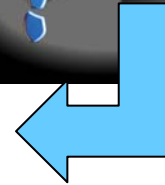
antibiotics



antifungals

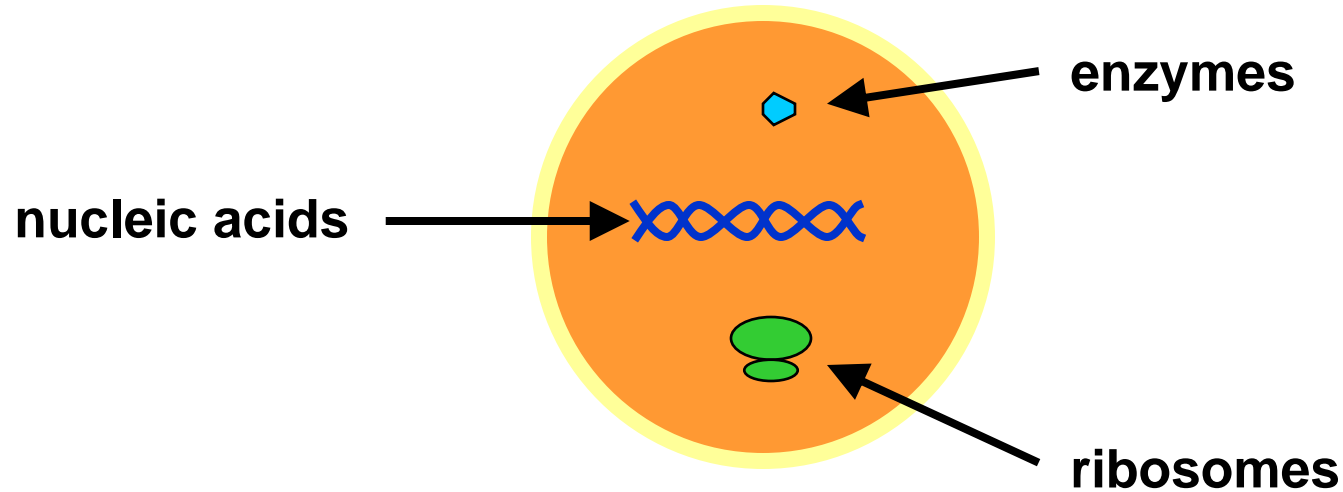


anticancer agents



# Chemotherapeutic agents exert toxic effects on specific target cells

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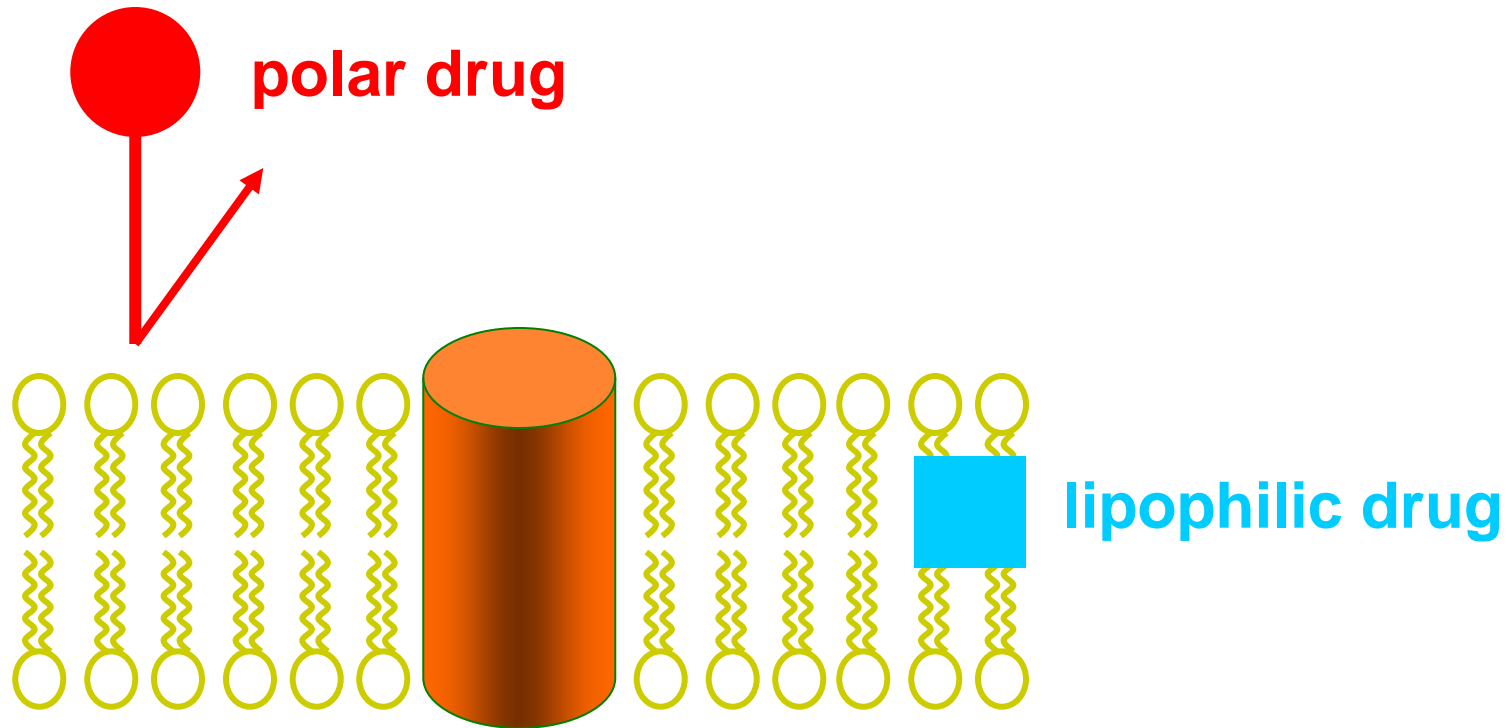


**How can these drugs reach their target inside the cells ?**

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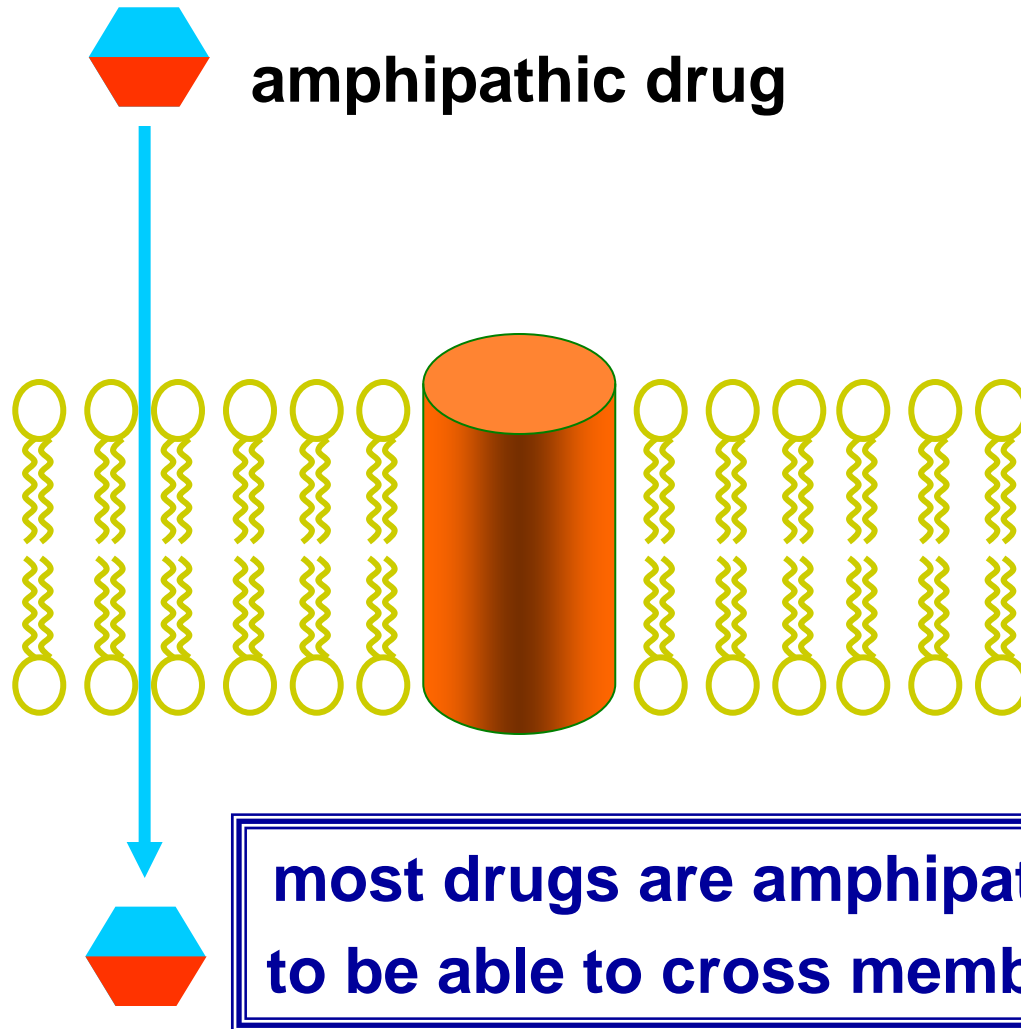


# Reaching an intracellular target ...

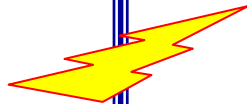
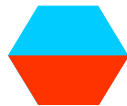
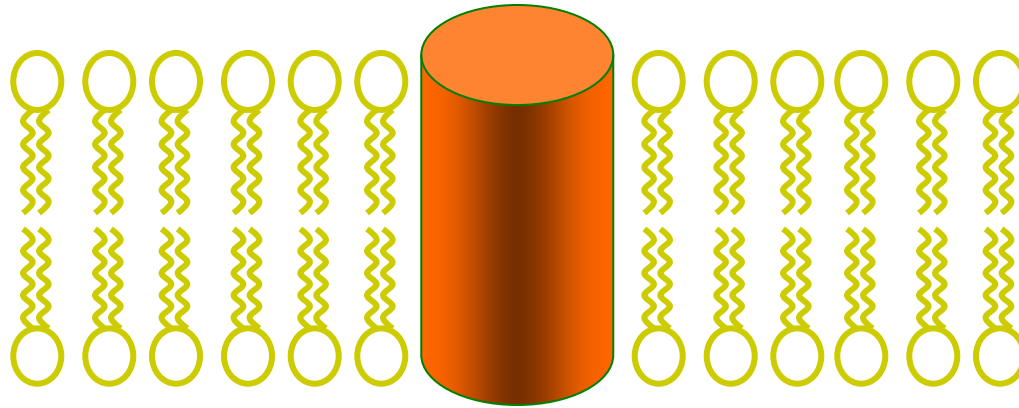


**physico-chemical properties are inadequate  
for reaching an intracellular target !**

# Reaching an intracellular target ...



# Intracellular chemotherapeutic agents



**But a diffusable compound  
may have  
potentially harmful effects !**





# Chemotherapeutic agents exert toxic effects on specific target cells

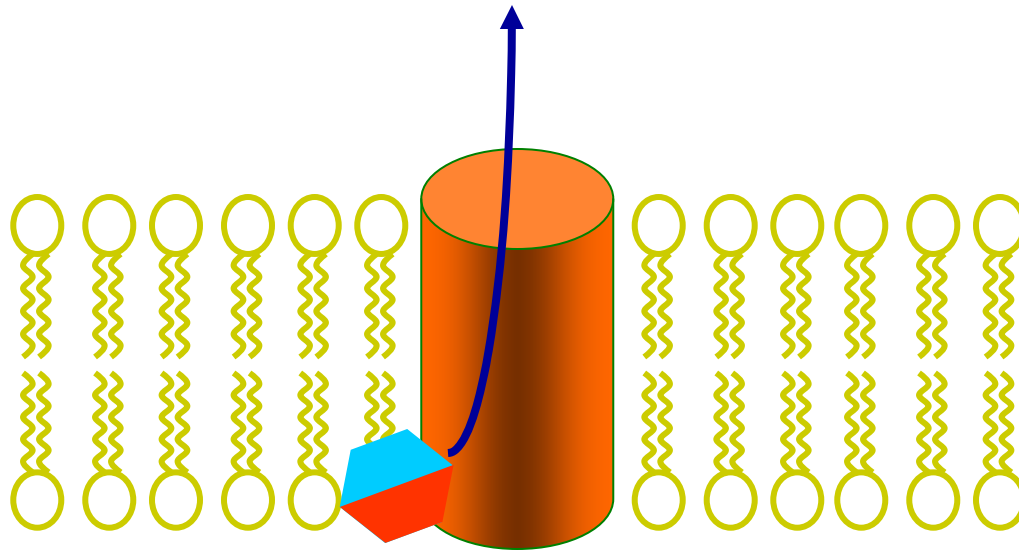
**How can cells protect themselves  
from these toxic substances ?**



# Why efflux transporters ?

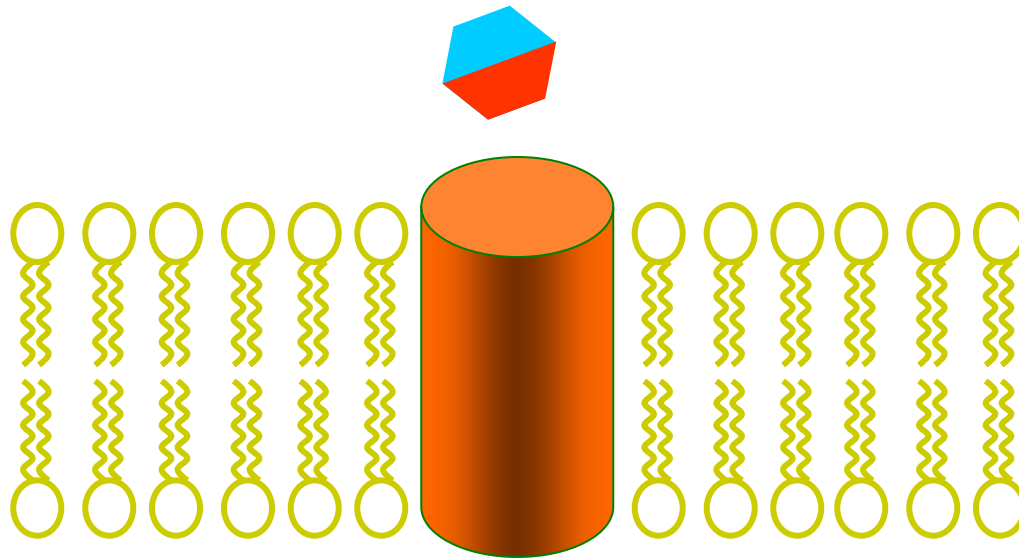
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## Extrusion by efflux pumps



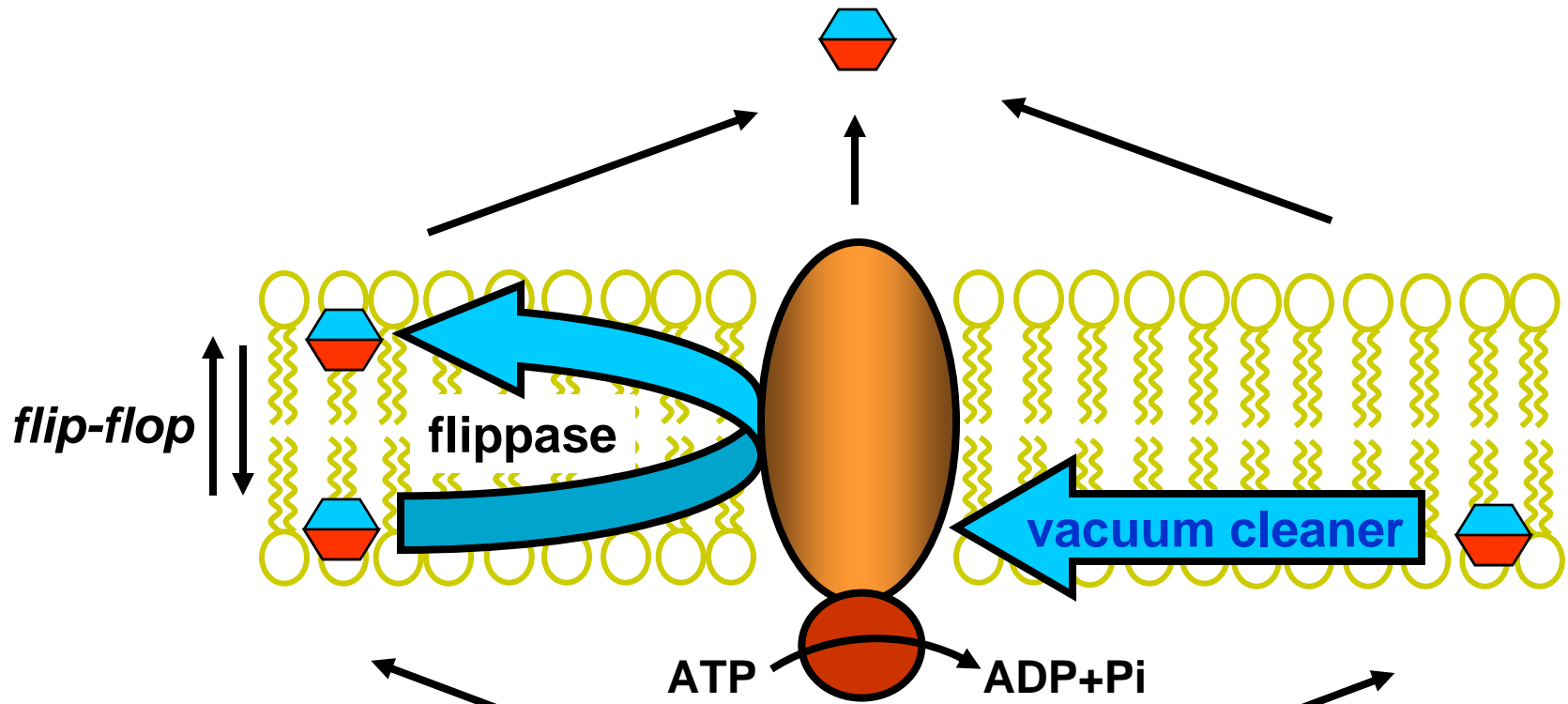
# Why efflux transporters ?

## Extrusion by efflux pumps



**general mean of protection  
against cell invasion by diffusible molecules**

# Mechanisms of active efflux

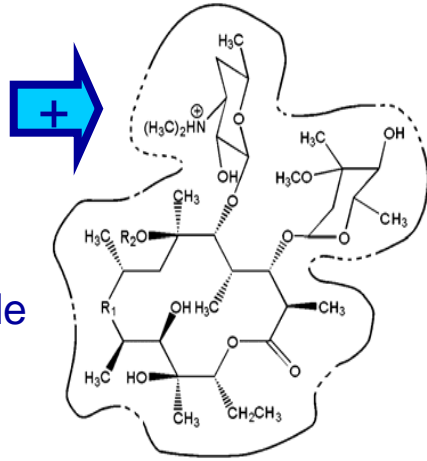


*membrane  
insertion / release*

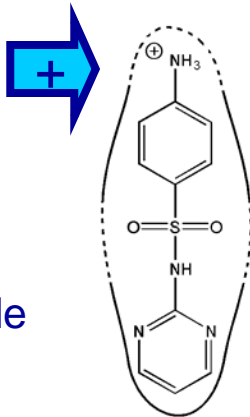
# Most antibiotics are amphiphilic !

## cationic amphiphiles

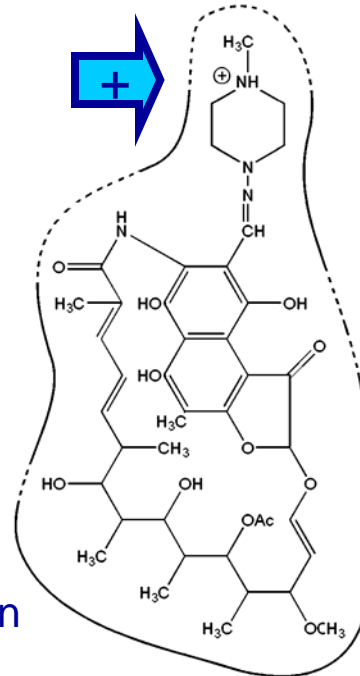
macrolide



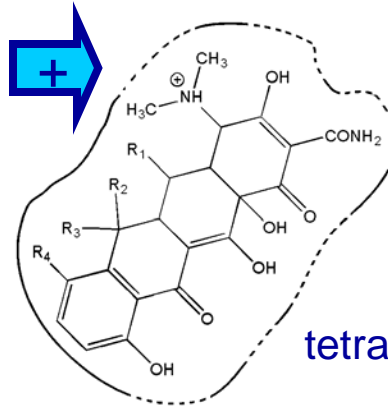
sulfamide



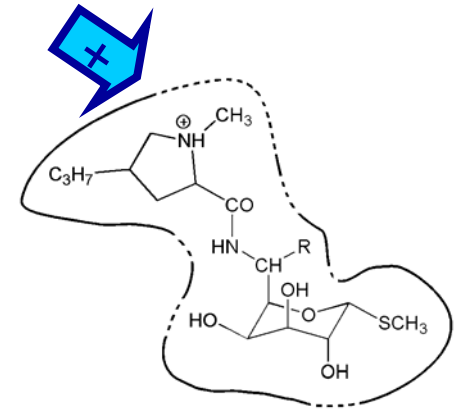
rifampicin



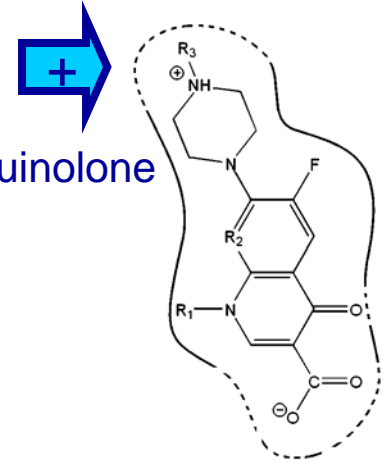
tetracycline



lincosamide

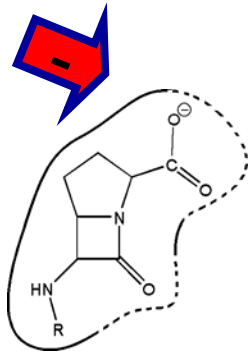


fluoroquinolone

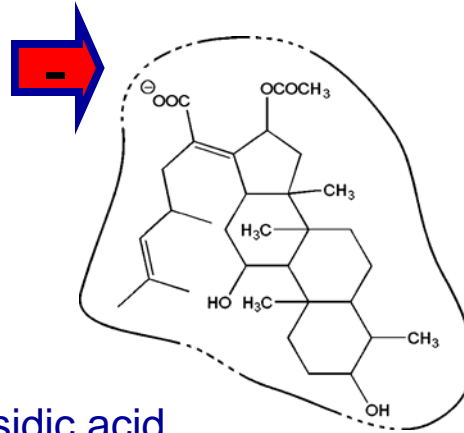


# Most antibiotics are amphiphilic !

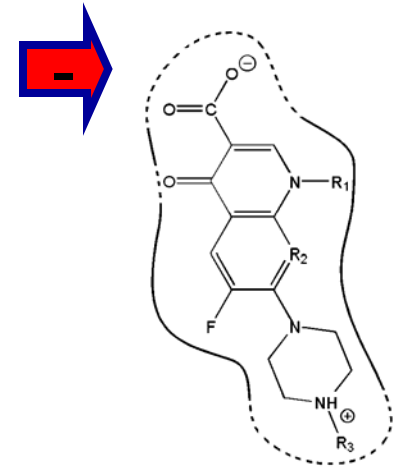
## anionic amphiphiles



$\beta$ -lactam

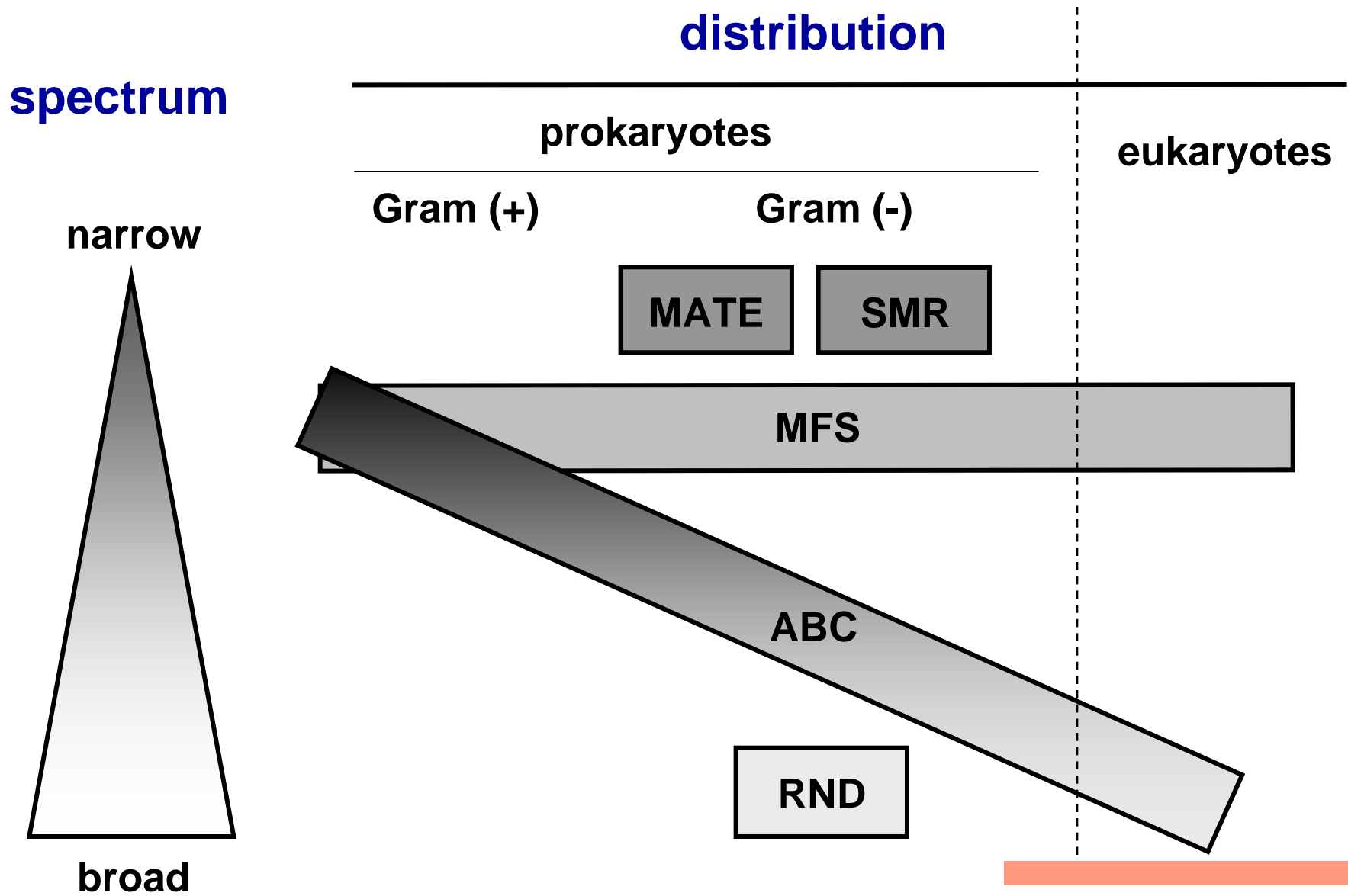


fusidic acid



fluoroquinolone

# Antibiotic efflux transporters are ubiquitous

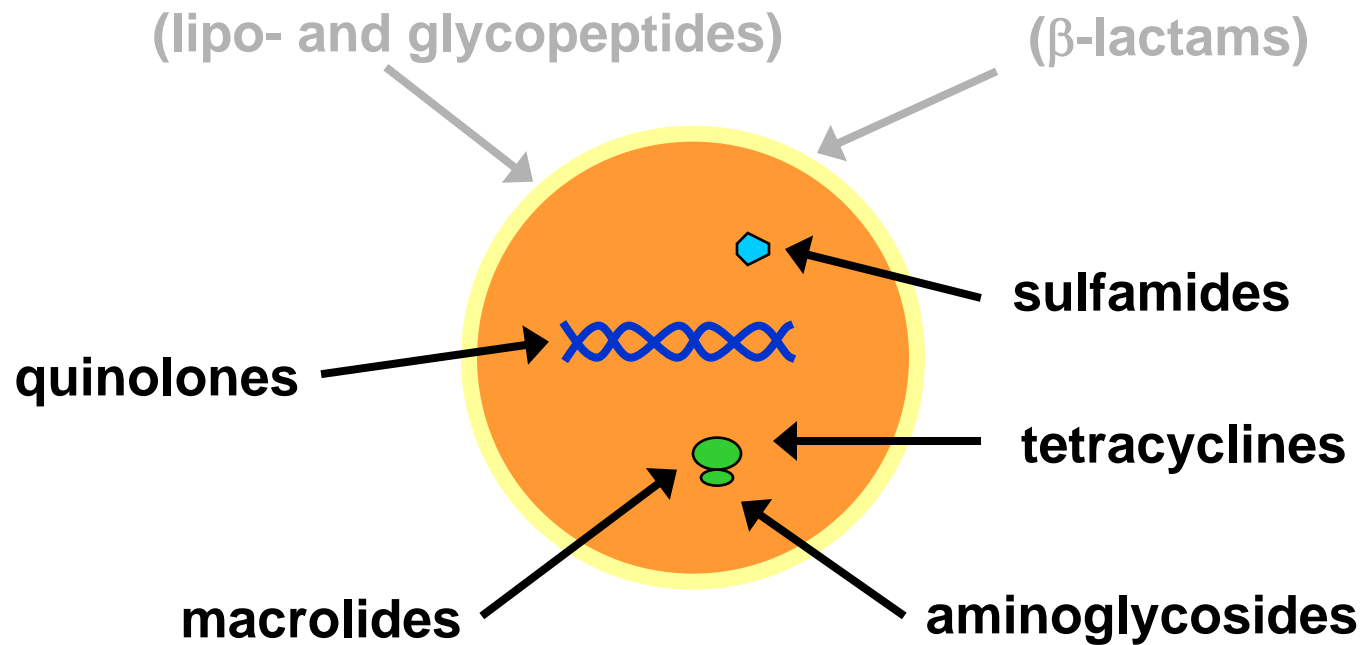




# Antibiotics as substrates of efflux pumps

Antibiotic class	bacteria		fungi	superior eucaryotes
	Gram (+)	Gram(-)		
$\beta$ -lactams	●	●	●	●
fusidic acid		●		
macrolides	●	●	●	●
streptogramins	●			●
tetracyclines	●	●	●	●
aminoglycosides		●	●	
chloramphenicol	●	●	●	
rifamycins				●
sulfamides			●	
trimethoprim		●		
fluoroquinolones	●	●		●

# Most antibiotics do act on intracellular targets

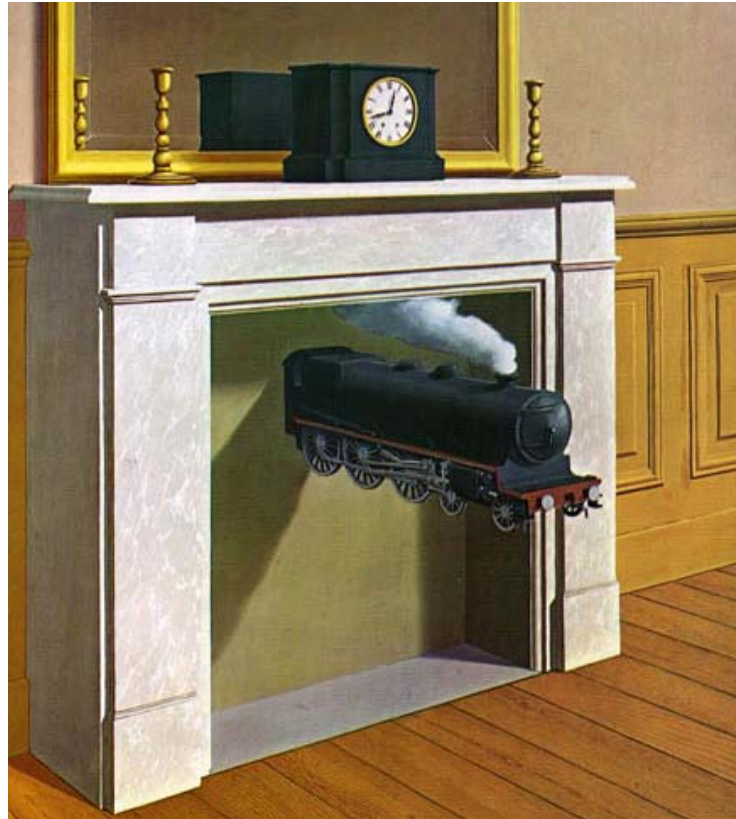


Efflux as a mechanism of resistance  
by reducing antibiotic concentration inside the bacteria

# Antibiotic efflux from bacteria as a mechanism of antibiotic resistance : molecular bases



# Efflux as a mechanism of export in antibiotic producers

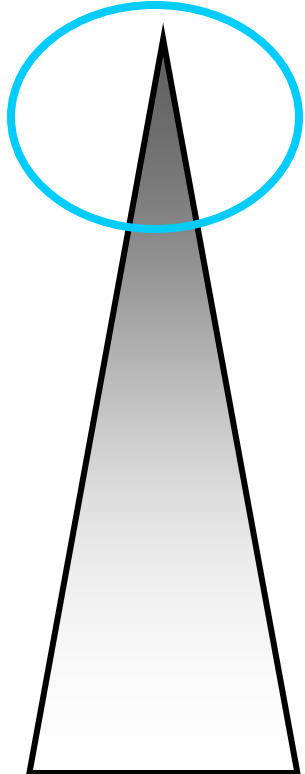


**Delivery of metabolites produced by the cell**

# Efflux as a mechanism of export in antibiotic producers

spectrum

narrow



specific for the produced antibiotic

ABC

OleB or C of *S. antibioticus*

→ oleandomycin

SmB of *S. ambofaciens*

→ macrolides

Tlrc of *S. fradiae*

→ tylosin

MFS

Ptr of *S. pristinaespiralis*

→ pristinamycin

LmrA of *S. lincolnensis*

→ lincomycin

RifP of *A. mediterranei*

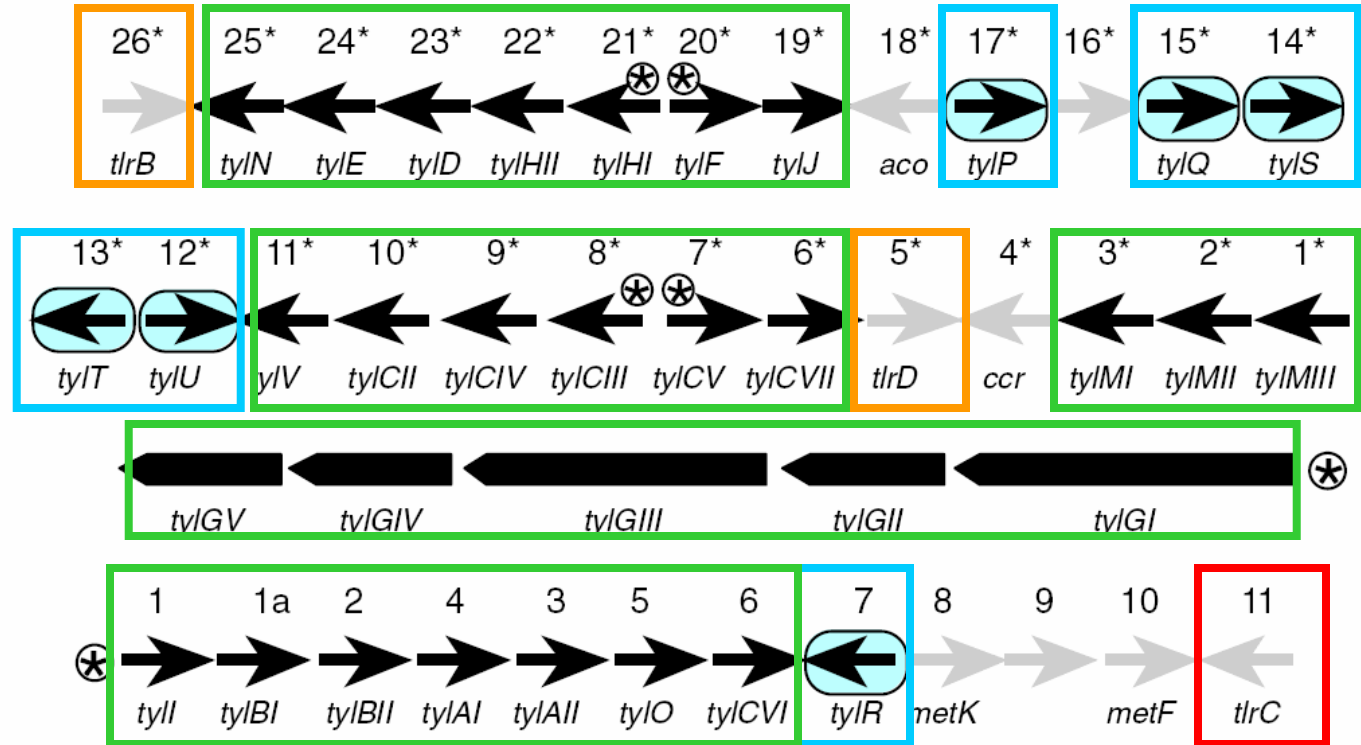
→ rifampicin

broad

# Efflux as a mechanism of export in antibiotic producers

## Tylosin-biosynthetic gene cluster in *S. fradiae*

**Fig. 1** The tylosin-biosynthetic gene cluster of *S. fradiae*. Not drawn to scale. The cluster occupies a contiguous portion of the genome (approximately 85 kb). Regulatory genes are outlined in *boxes*. Tylosin-biosynthetic genes are represented by *black arrows*. Resistance determinants (designated 'tlr'), ancillary genes and others that are unassigned are represented as *grey arrows*. The full complement of biosynthetic genes could, in principle, be expressed from three pairs of divergent promoters (*stars*) via operon control



tylosin biosynthesis  
regulation

resistance determinants  
efflux

# Efflux as a mechanism of antibiotic resistance in pathogenic bacteria



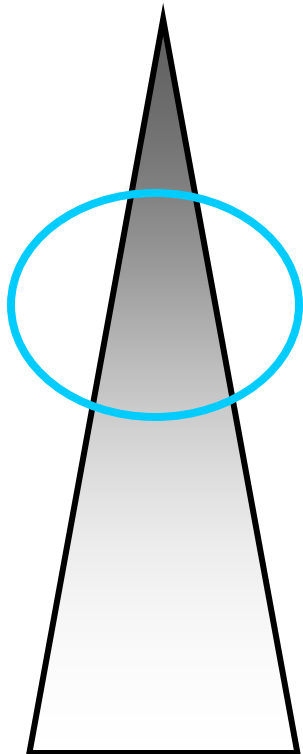


# Efflux as a mechanism of resistance

## in Gram-positive bacteria

### spectrum

narrow



broad

specific for one (or a few) families of drugs

ABC

PatA/PatB of *S. pneumoniae*

→ FQ, chl

MsrA of *S. epidermidis*

→ erythromycin

MFS

NorA of *S. aureus*

→ FQ, Tet, chl

MefE of *S. pneumoniae*

→ ML

PmrA of *S. pneumoniae*

→ FQ

MefA of *S. pyogenes*

→ ML

# Efflux as a mechanism of resistance

## in Gram-negative bacteria

### spectrum

narrow

specific for one (or a few) families of drugs

MFS

TetA of *E. coli*

→ Tet

MsrA of *S. epidermidis*

→ erythromycin

broad spectrum, conferring cross-resistance

RND

MexAB-OprM of *P. aeruginosa*

→  $\beta$ -lac, FQ, Tet, ML, chl, rif, sulf

AcrAB-TolC of *E. coli*

→  $\beta$ -lac, FQ, Tet, ML, chl, rif, sulf

broad

# Molecular determinants of drug efflux



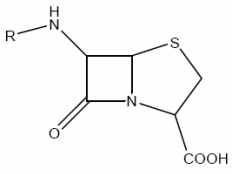
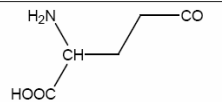
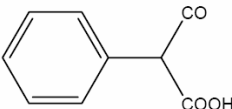
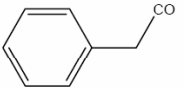
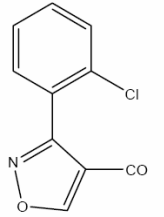
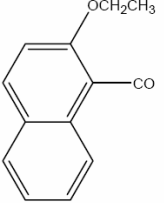
# Differences in transport

## between drugs within a class

usual difference in affinity of efflux pumps towards antibiotics			
antibiotic class	affinity for efflux pumps		
	high	variable <sup>a</sup>	low
penicillins	nafcillin, cloxacillin, penicillin G		carbenicillin
cephalosporins	cefalotin, cefotaxime, ceftriaxone		cefazolin, cephaloridin
carbapenems	meropenem	imipenem	
macrolides	14 - and 15 - membered		16 –membered, ketolides
tetracyclines	tetracycline	minocycline glycylcyclines	
quinolones	ciprofloxacin, norfloxacin	ofloxacin, levofloxacin	cinafloxacin, gatifloxacin, gemifloxacin, moxifloxacin, garenoxacin
<sup>a</sup> depending on the efflux pump			

# Differential efflux of $\beta$ -lactams

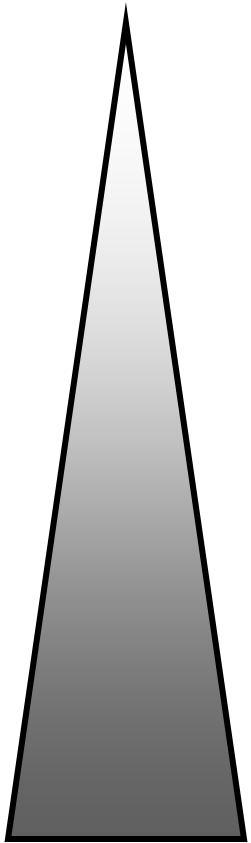
## by AcrB of *S. typhimurium*

	Relationship between structure, lipophilicity of the side chain of selected penicillins and MIC of bacteria resistant by efflux		
R (Side chain)	molecule	side-chain partition coefficient	MIC ratio <sup>a</sup>
	penicillin N	0	1
	carbenicillin	80	4
	penicillin G	270	32
	cloxacillin	890	256
	nafcillin	4200	128

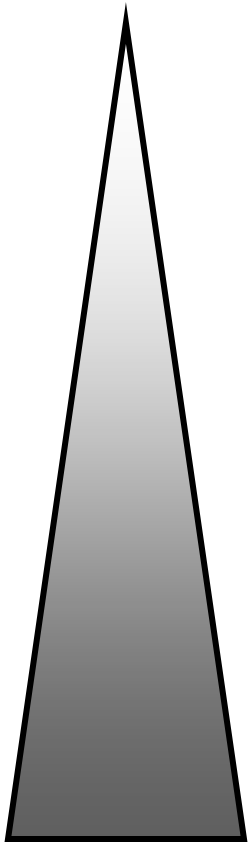
<sup>a</sup> ratio between the MIC of a *S. typhimurium* resistant mutant (AcrAB-ToIC overproducer) and of its susceptible parental strain

### $\beta$ -lactams & broad spectrum RND

Lipophilicity  
of the side chain

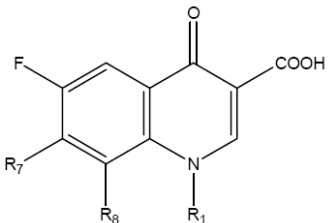

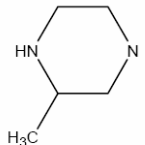

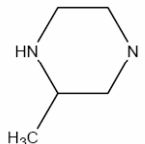

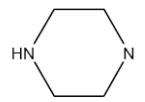
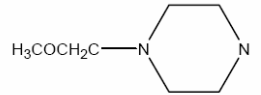
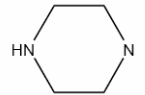


Resistance



# Differential efflux of quinolones

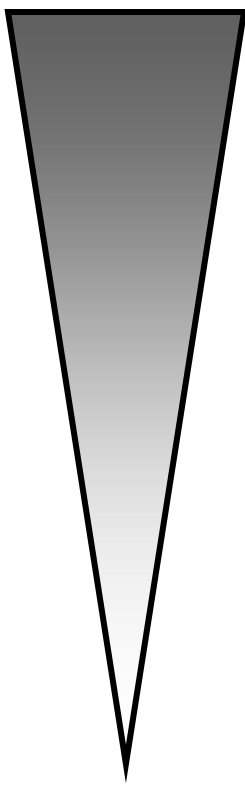
## by NorA of *S. aureus*

<div>  <div>Relationship between structure, bulkiness, and hydrophobicity of selected quinolone substituents and MIC of bacteria resistant by efflux.</div> </div>				
molecule	R <sub>1</sub>	R <sub>7</sub>	R <sub>8</sub>	MIC ratio <sup>a</sup>
(a)			OCF <sub>3</sub>	2
(b)			OCH <sub>3</sub>	4
ciprofloxacin			H	16
(c)	C <sub>2</sub> H <sub>5</sub>		H	16
norfloxacin	C <sub>2</sub> H <sub>5</sub>		H	64

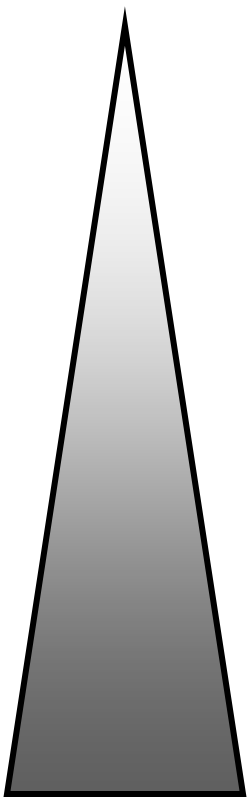
<sup>a</sup> ratio between the MIC of a *S. aureus* resistant mutant (NorA overproducer) and of its susceptible parental strain.

### Quinolones & narrow spectrum MFS

Bulkiness  
in 7 and 8



Resistance



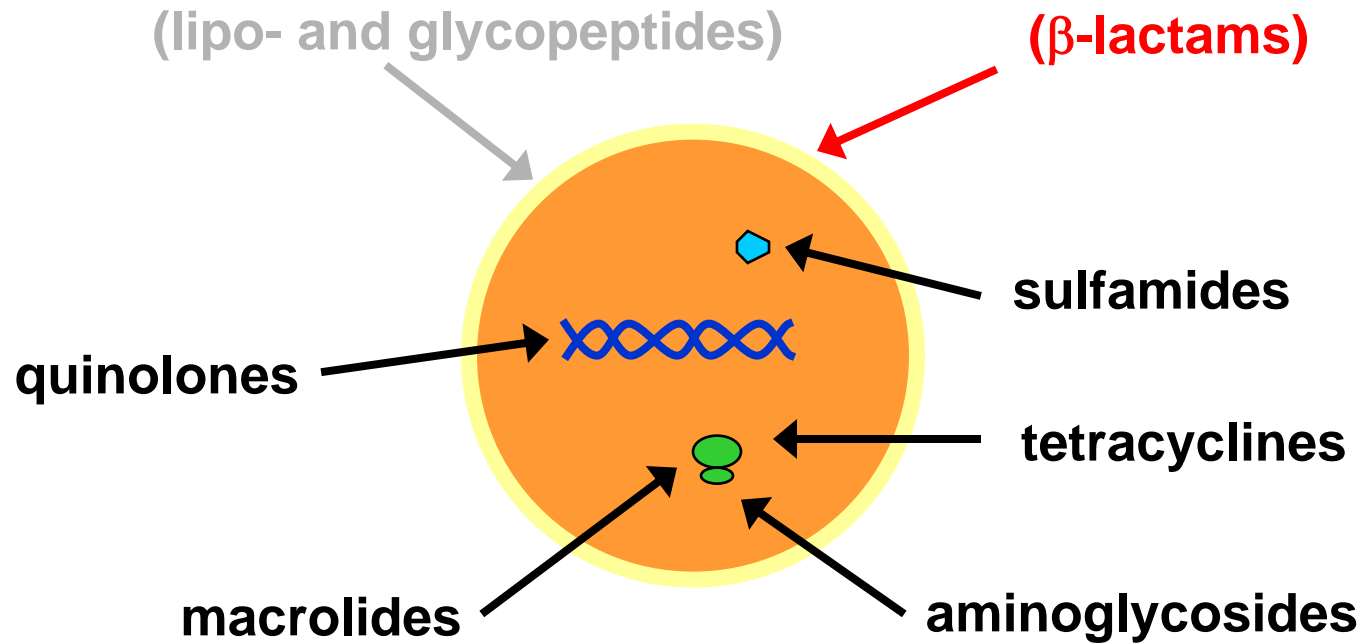
# Unexpected antibiotic substrates





# Two types of unexpected substrates

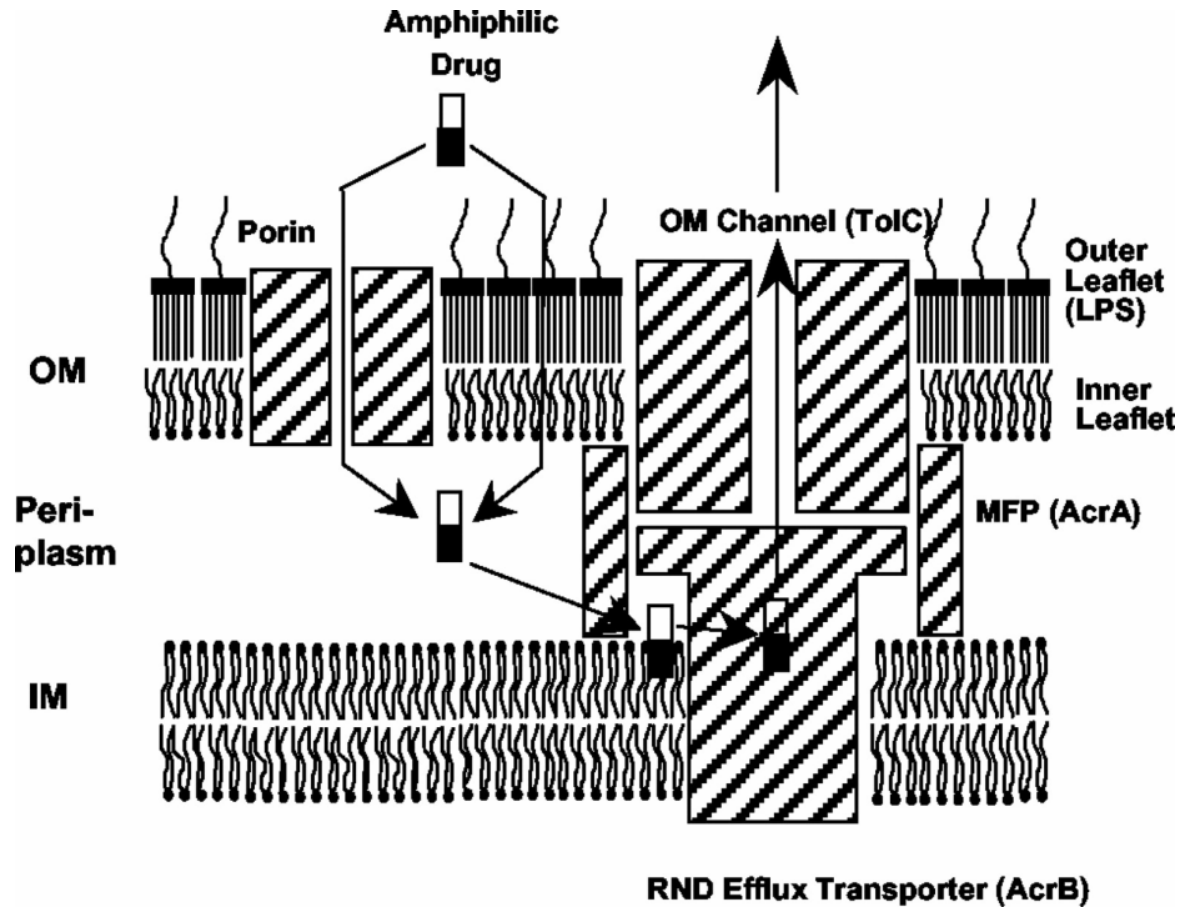
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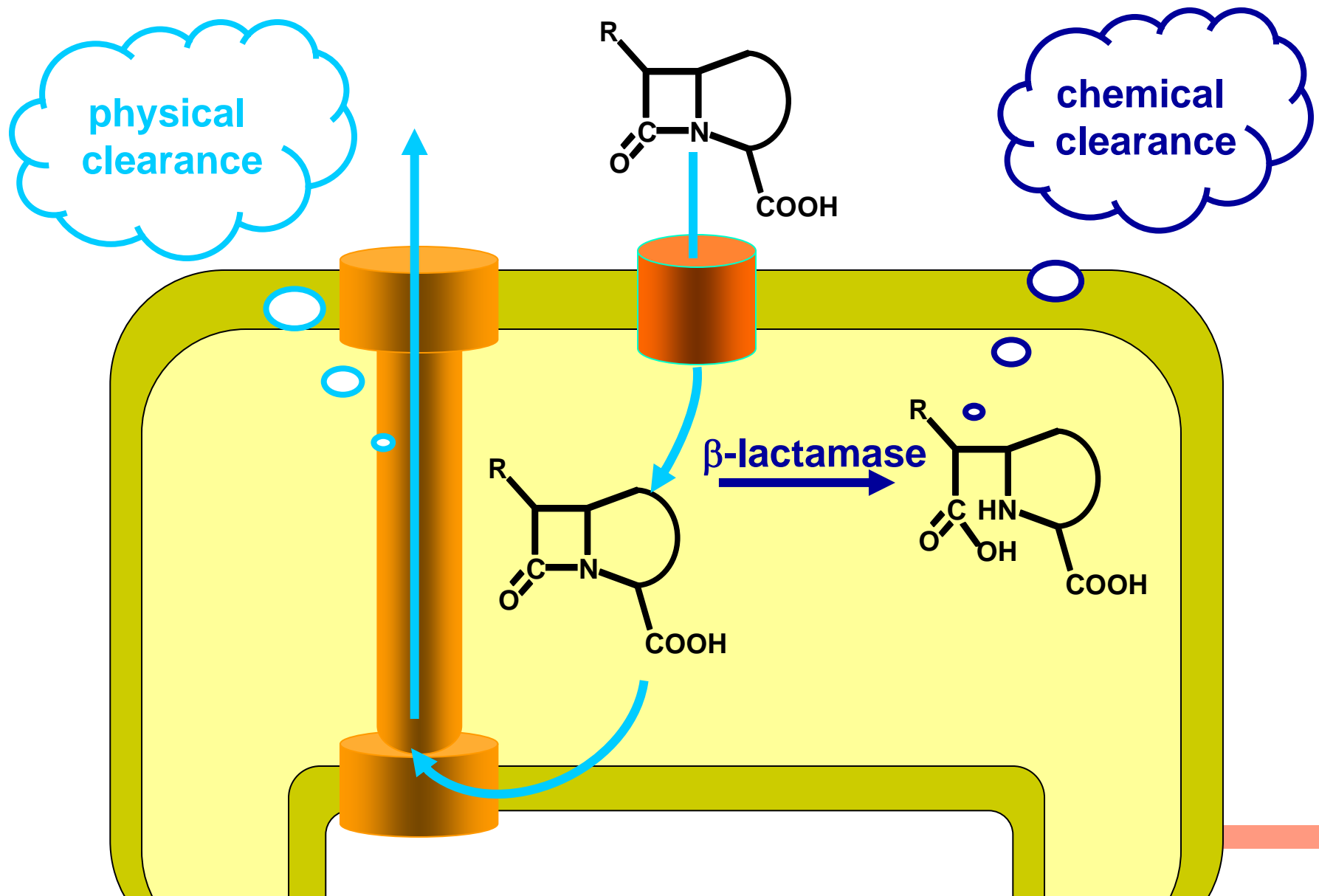
active on an extracellular target !

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# $\beta$ -lactams as unexpected substrates for efflux pumps



# $\beta$ -lactams as unexpected substrates for efflux pumps



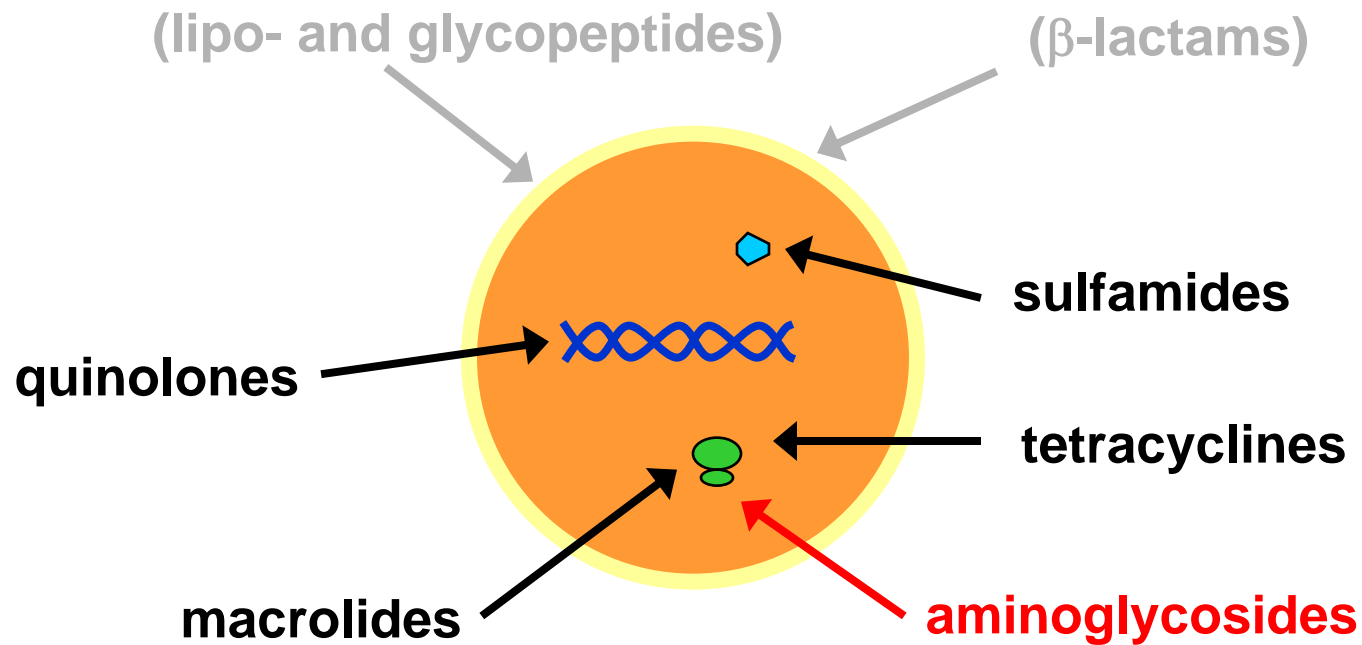
# $\beta$ -lactams as unexpected substrates for efflux pumps

## Contributions of the AmpC $\beta$ -lactamase and the AcrAB Multidrug Efflux System in Intrinsic Resistance of *E. coli* to $\beta$ -lactams

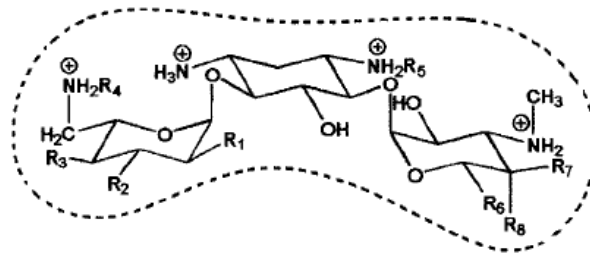
Efflux	$\beta$ -lactamase	CMI carbenicillin	CMI ofloxacin
-	-	0.2	0.05
+	-	12.5	0.2
+++	-	50	1.56
-	+	100	0.05
+	+	200	0.39
+++	+	400	1.56

WT:  
intrinsic  
resistance !

# Two types of unexpected substrates

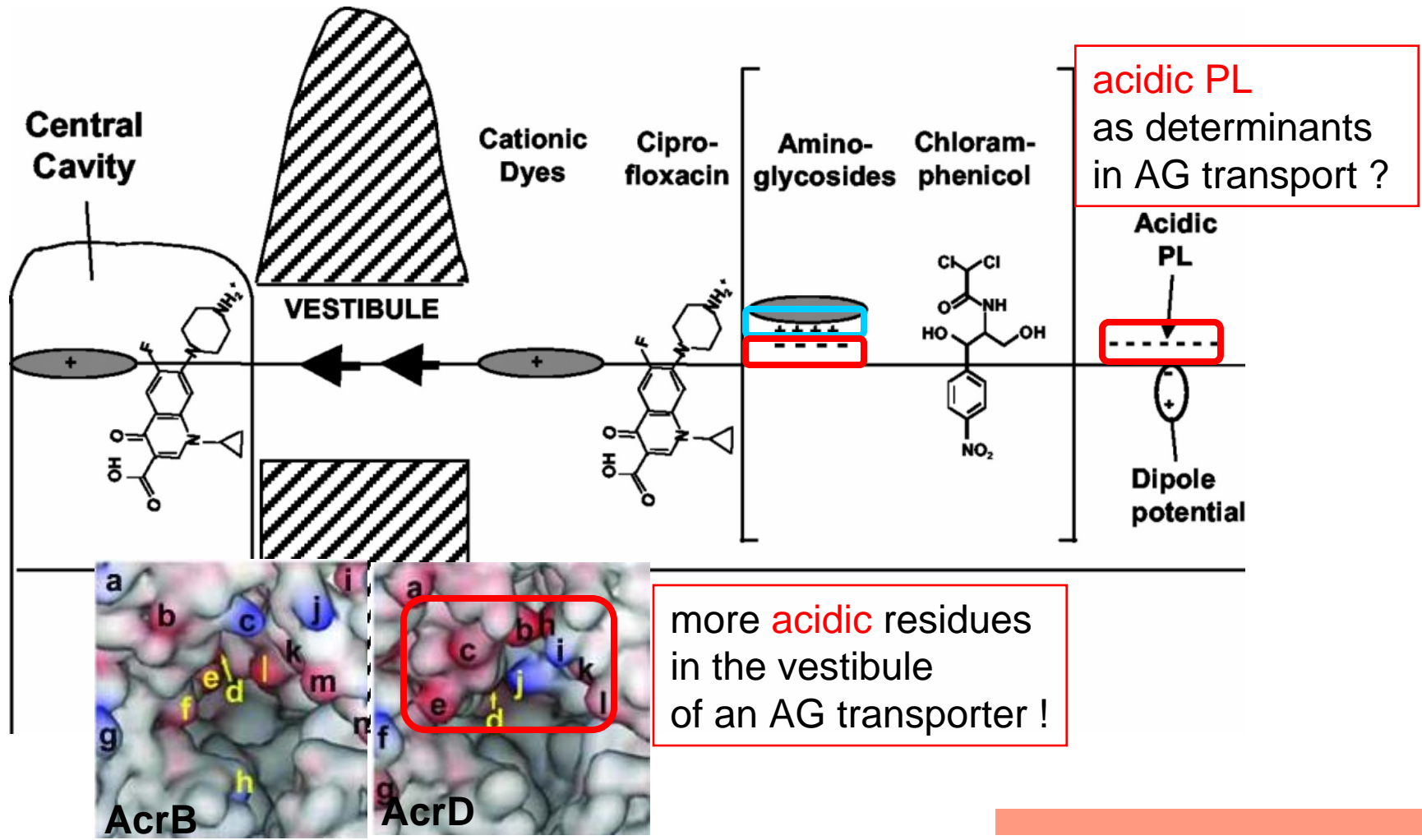


Highly hydrophilic molecules !

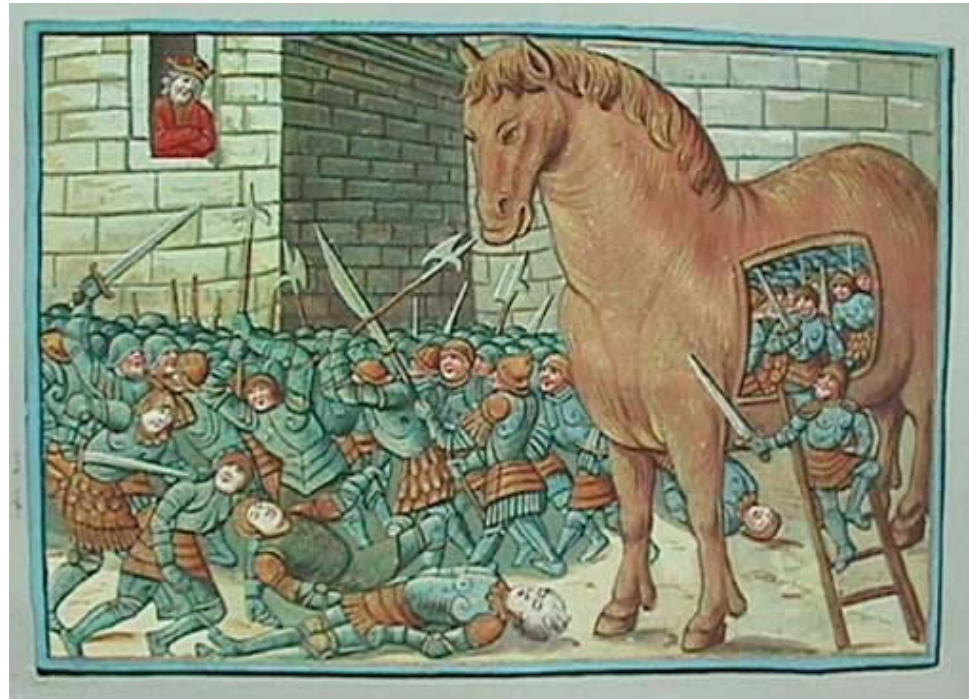


# Aminoglycosides as unexpected substrates for efflux pumps

## Composite binding site ?

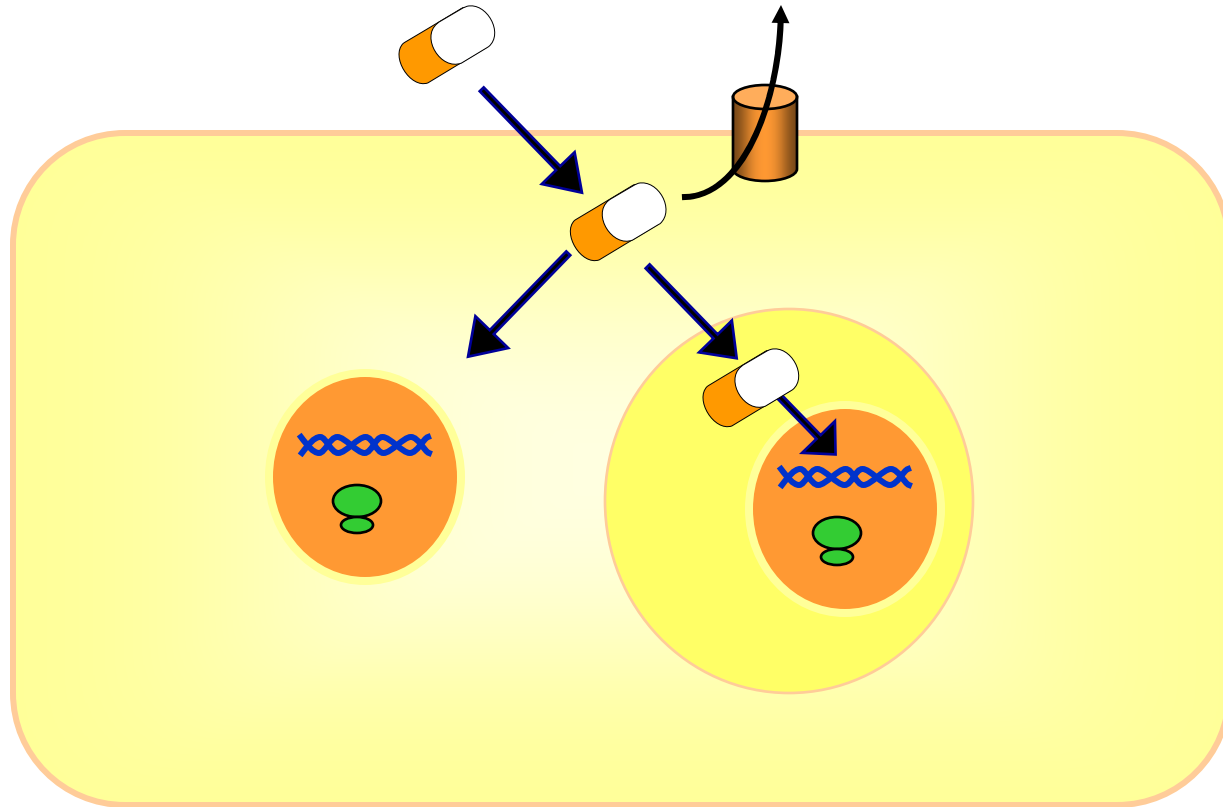


**efflux from eucaryotic cells  
as a mechanism of antibiotic 'resistance':  
molecular bases**





# Antibiotic efflux from eucaryotic cells and intracellular 'resistance' ?

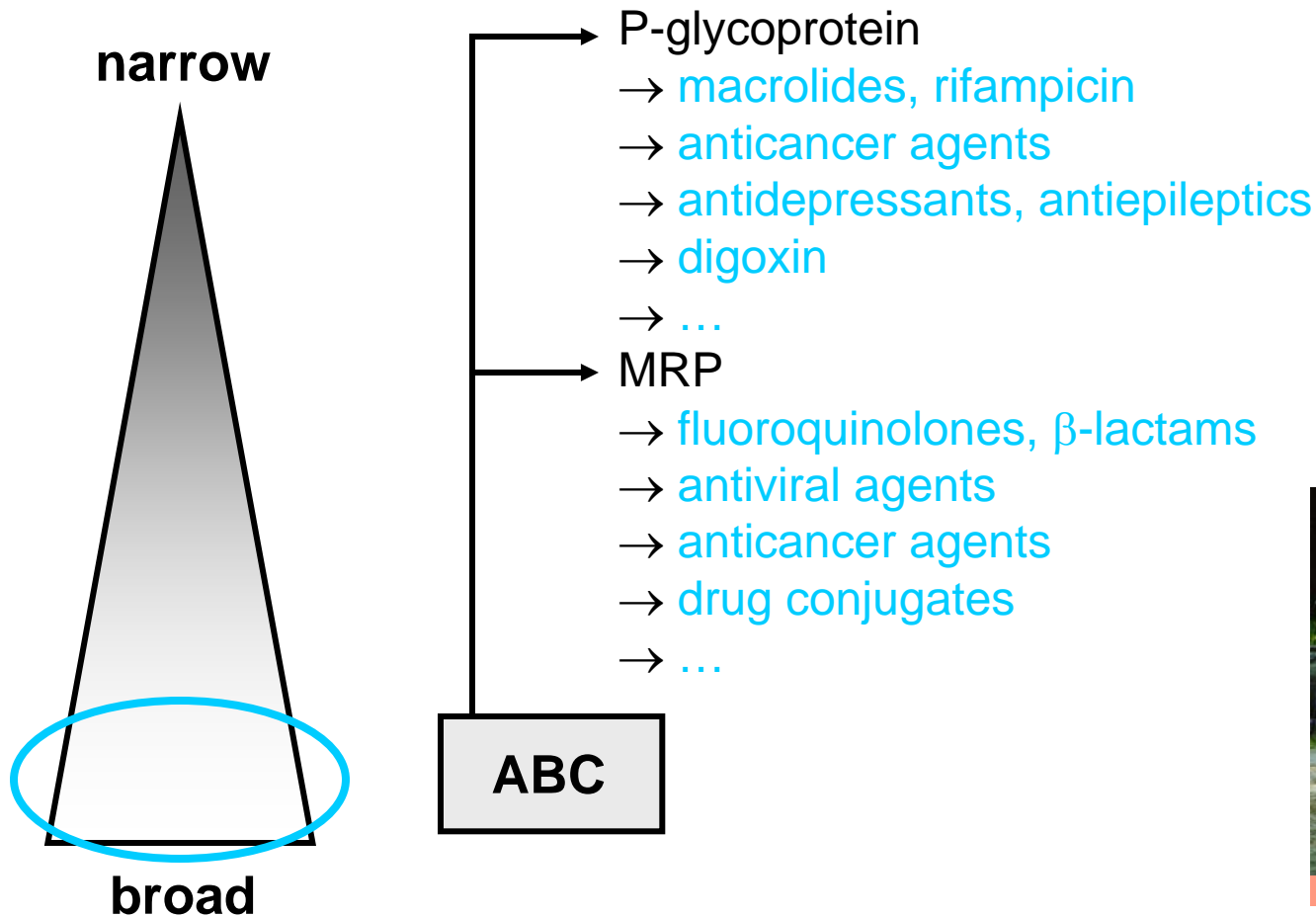


Reduction in intracellular drug concentration  
can result in inefficacy against intracellular bacteria

# Antibiotics as substrates of MDR efflux pumps

## spectrum

### Broad spectrum, several classes of drugs

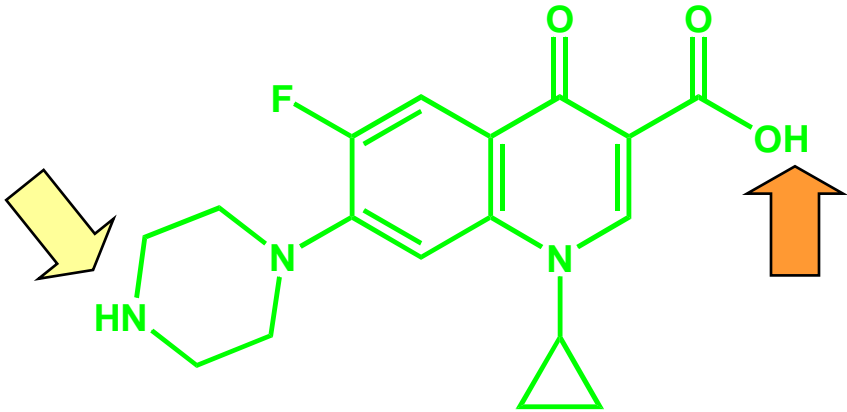
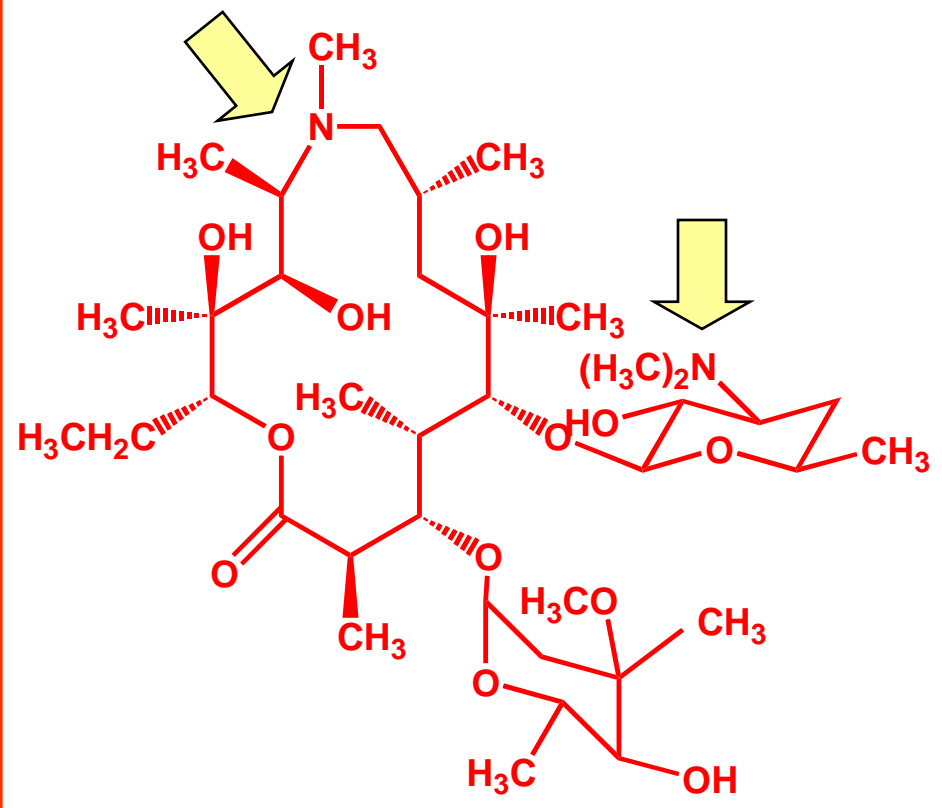


# Phenotypic description of antibiotic transport by MDR pumps

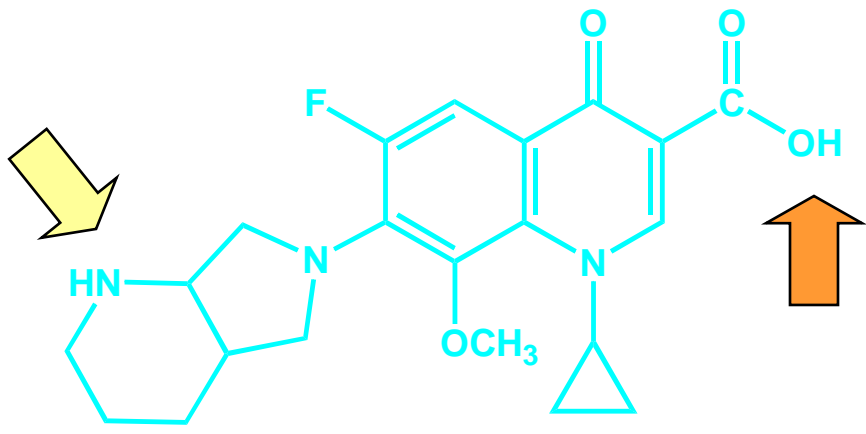


# Antibiotics as substrates of MDR efflux pumps

azithromycin



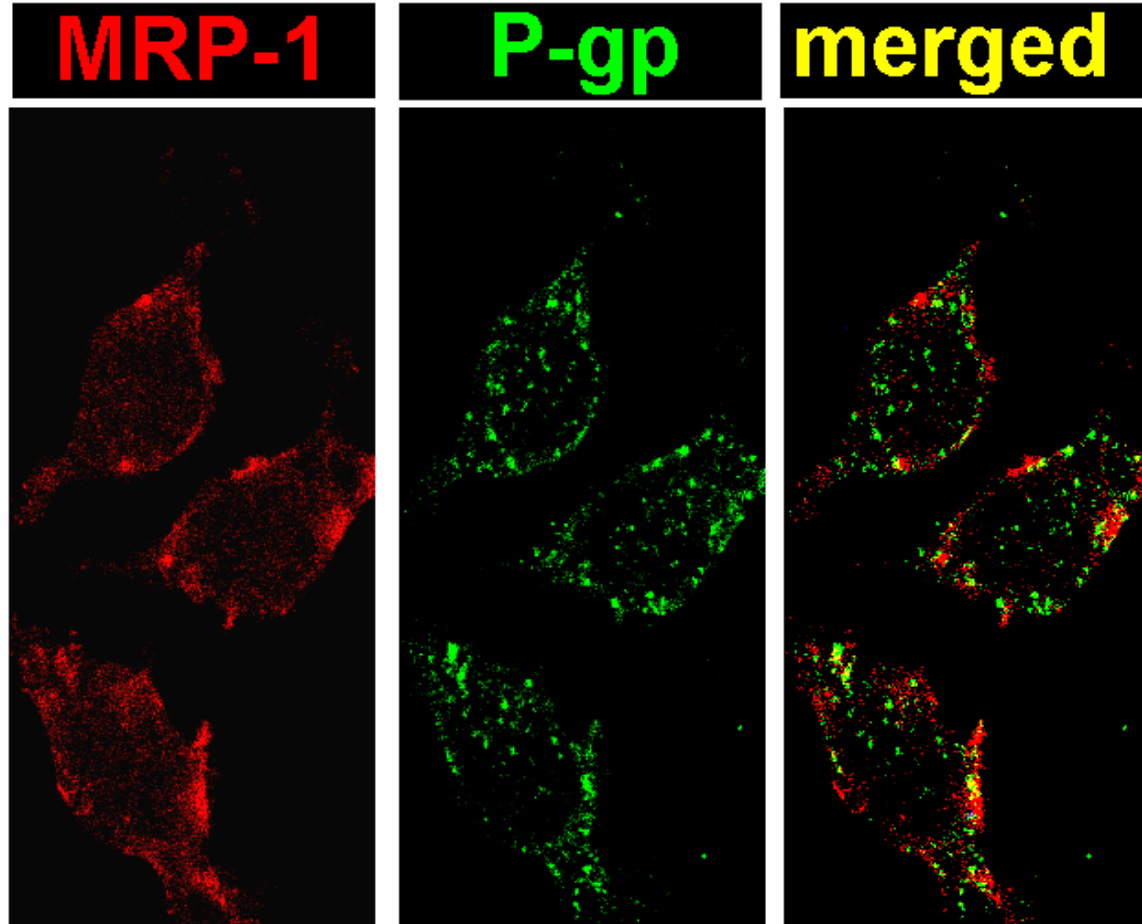
ciprofloxacin



moxifloxacin

# J774 macrophages do express MDR pumps

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# ABC multidrug transporters

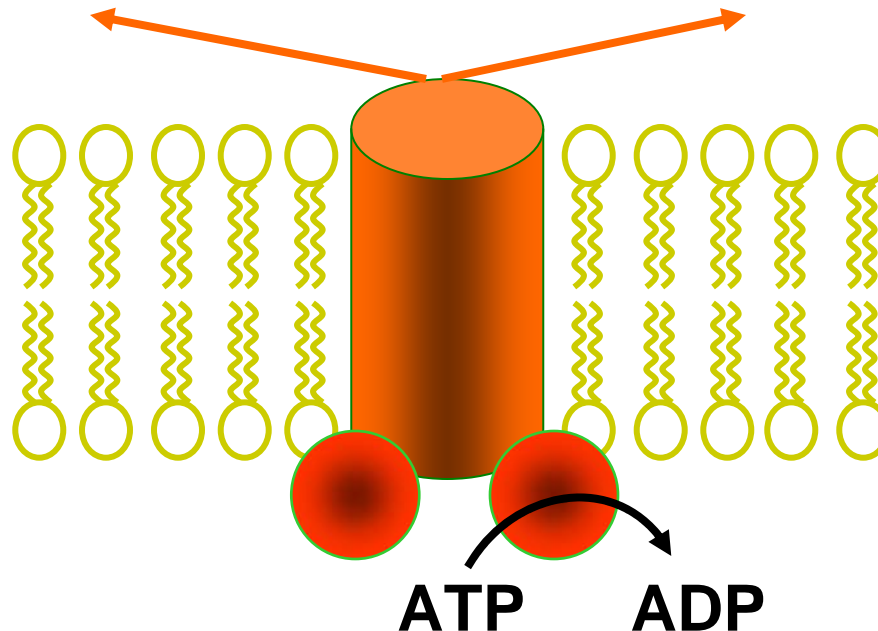
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**cationic  
amphiphiles**

**MDR-1 (P-glycoprotein)**

**anionic  
amphiphiles**

**MRP1-10**



# How to inhibit ABC transporters ?

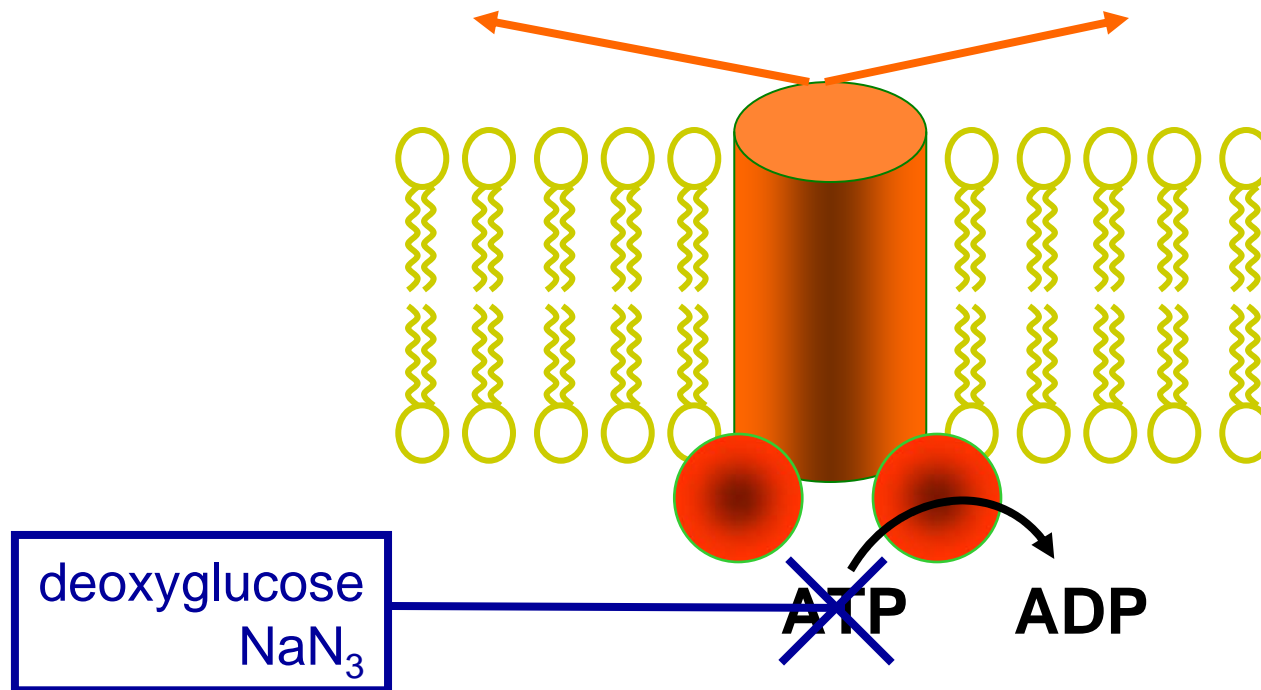
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**cationic  
amphiphiles**

**MDR-1 (P-glycoprotein)**

**anionic  
amphiphiles**

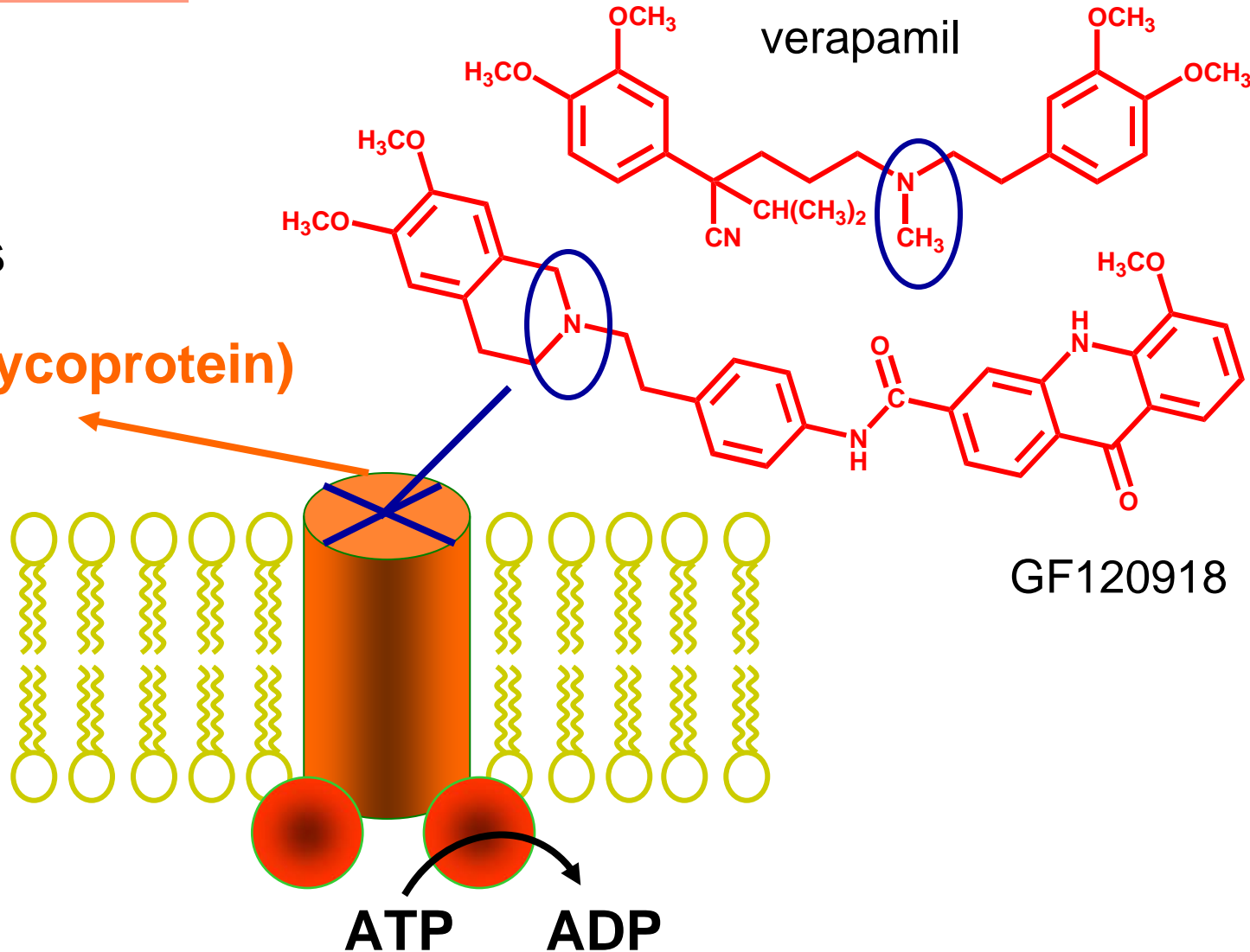
**MRP1-10**



# How to inhibit ABC transporters ?

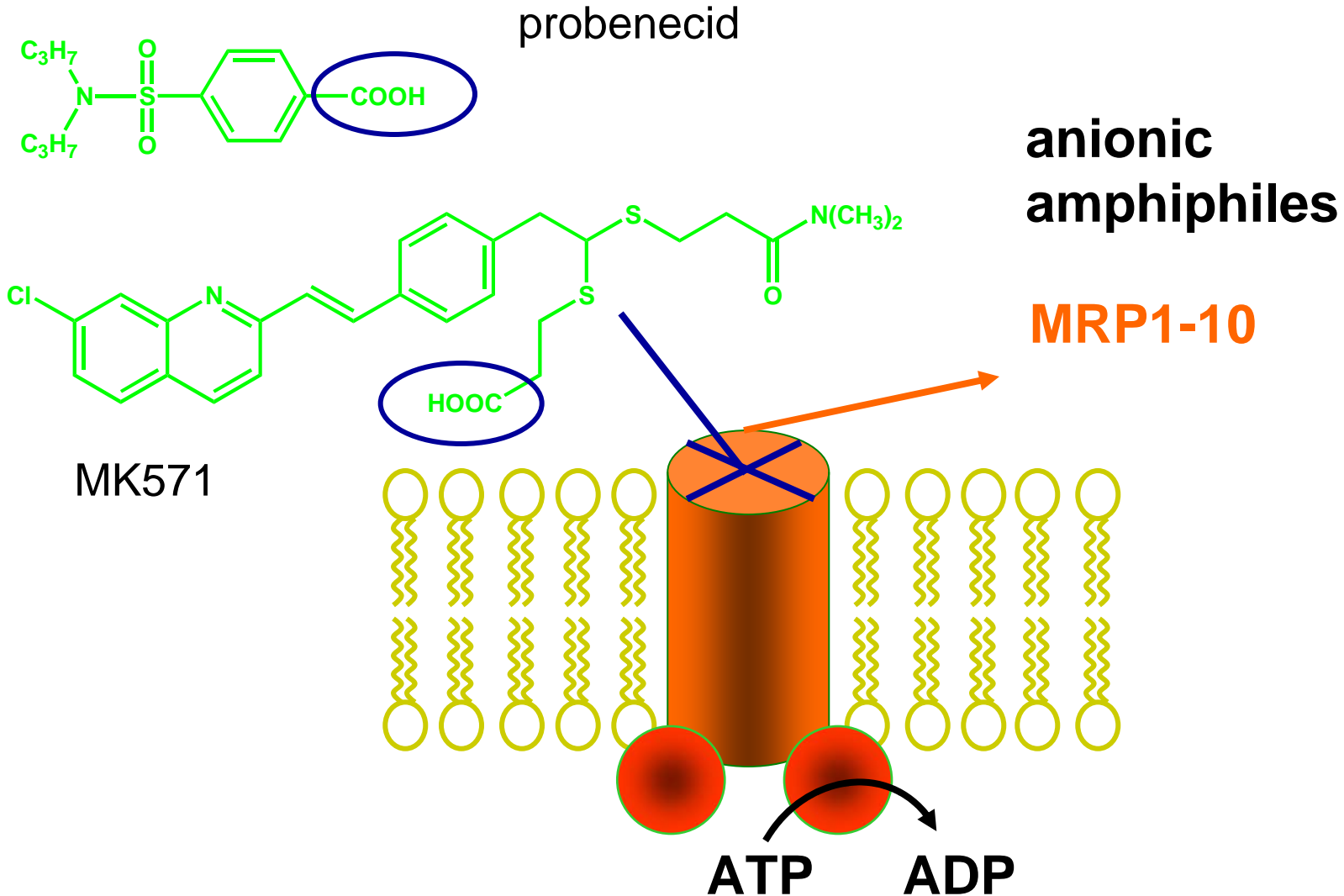
**cationic  
amphiphiles**

**MDR-1 (P-glycoprotein)**





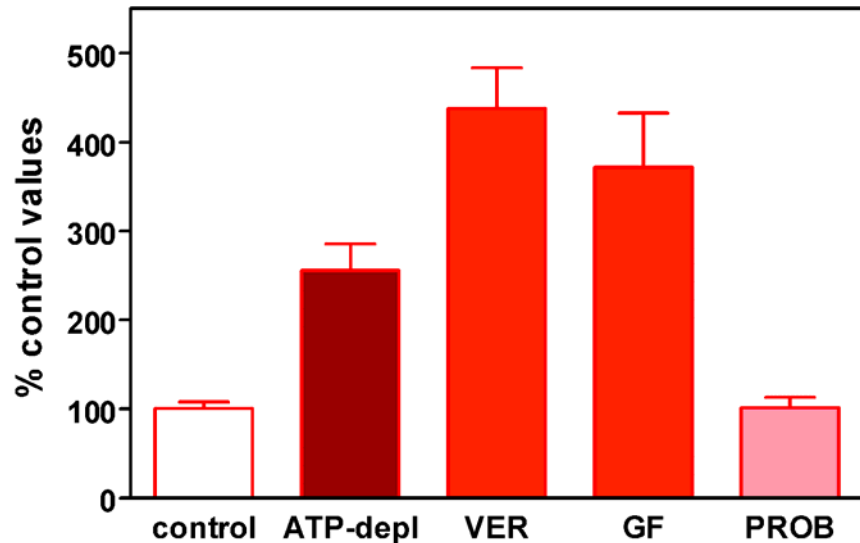
# How to inhibit ABC transporters ?



# Differential recognition of antibiotics by MDR pumps

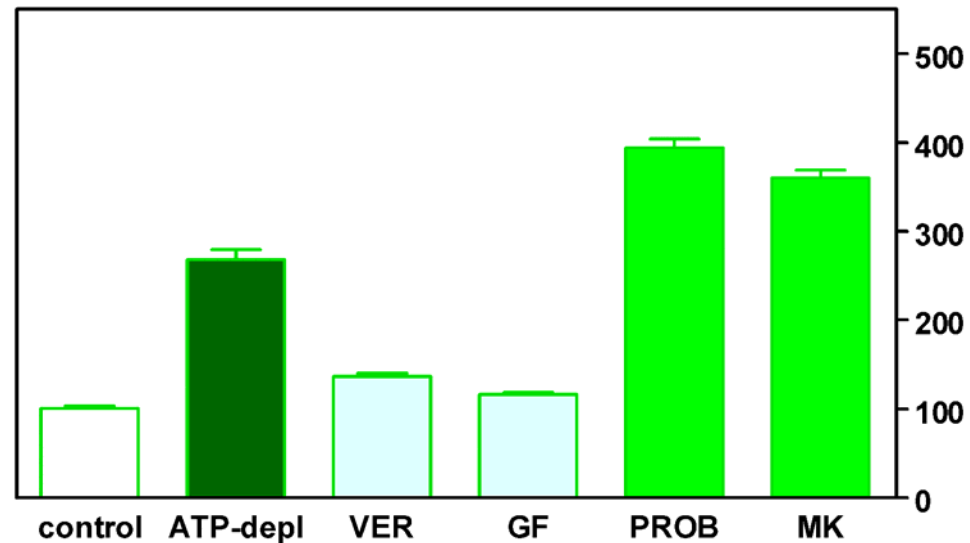
## Influence of ATP-depletion and pump inhibitors on accumulation at equilibrium

azithromycin



**azithromycin  
&  
P-glycoprotein**

ciprofloxacin

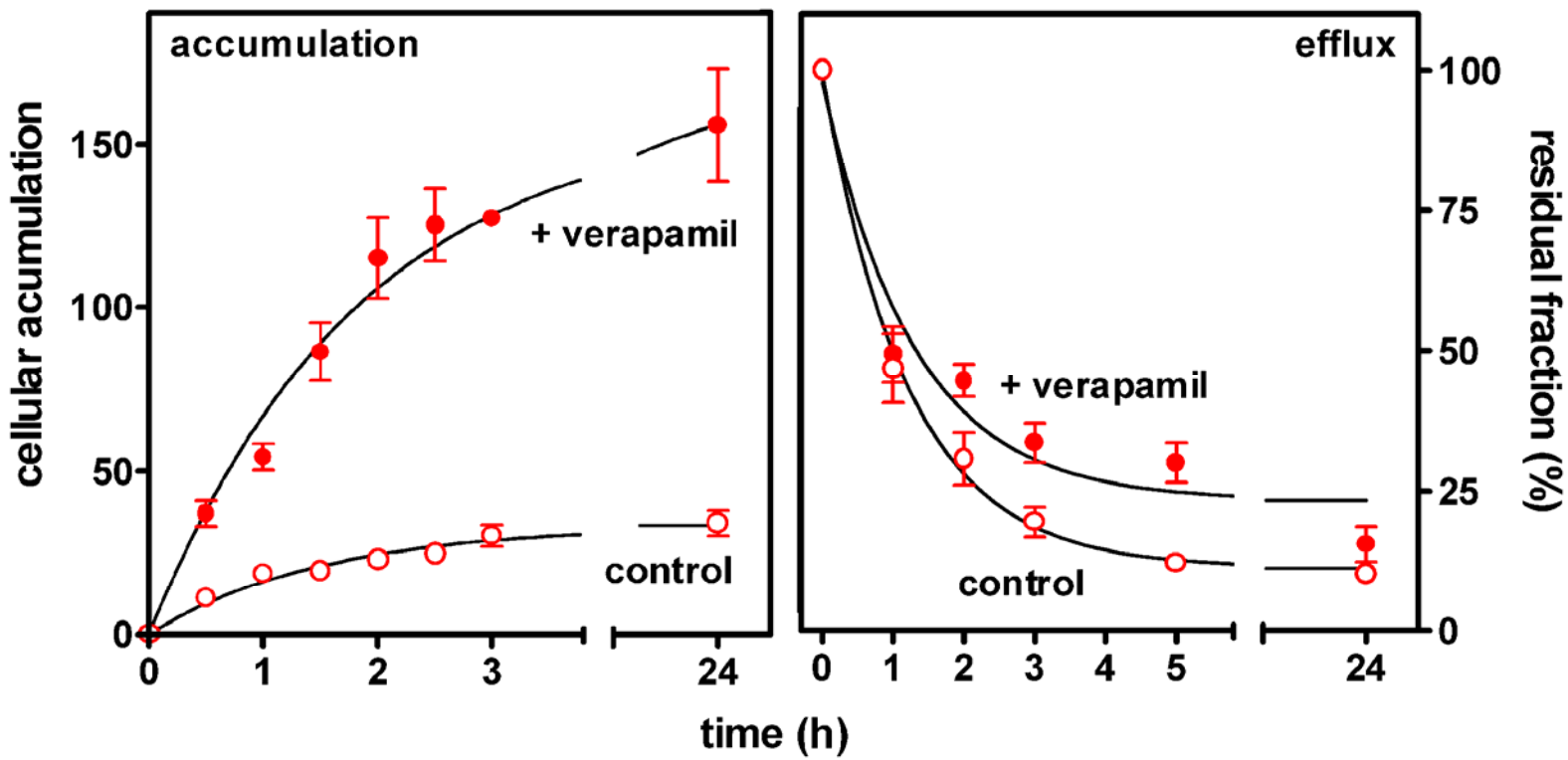


**ciprofloxacin  
&  
MRP**

extracell. conc. 5 mg/L;  
AZM 3 h; CIP 2 h

# Kinetics of accumulation and efflux for azithromycin

accumulation markedly increased; efflux marginally affected

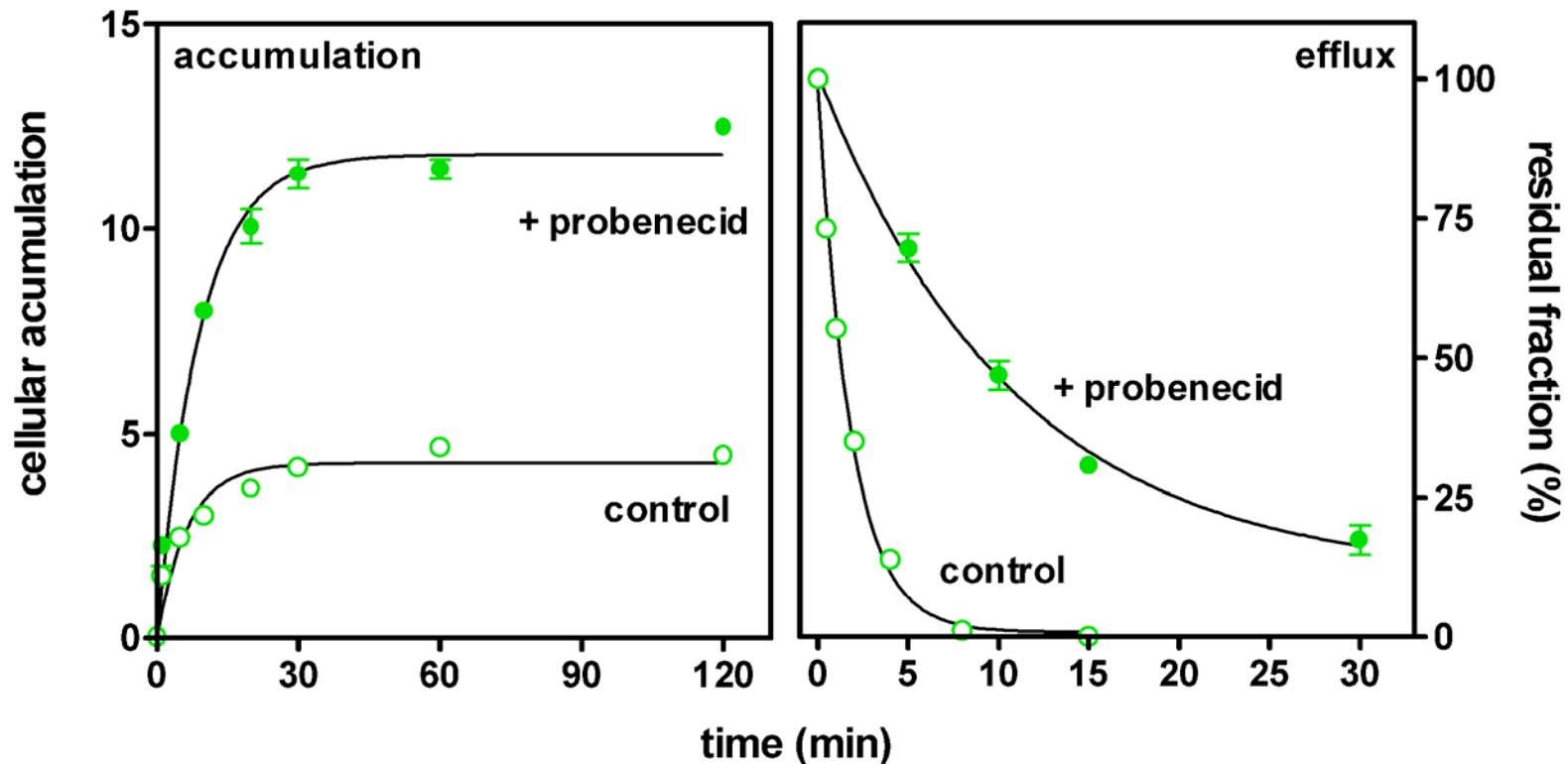


extracell. conc. 5 mg/L; verapamil 20  $\mu$ M

Seral et al. (2003) AAC 47:1047-51

# Kinetics of accumulation and efflux for ciprofloxacin

both accumulation and efflux markedly affected

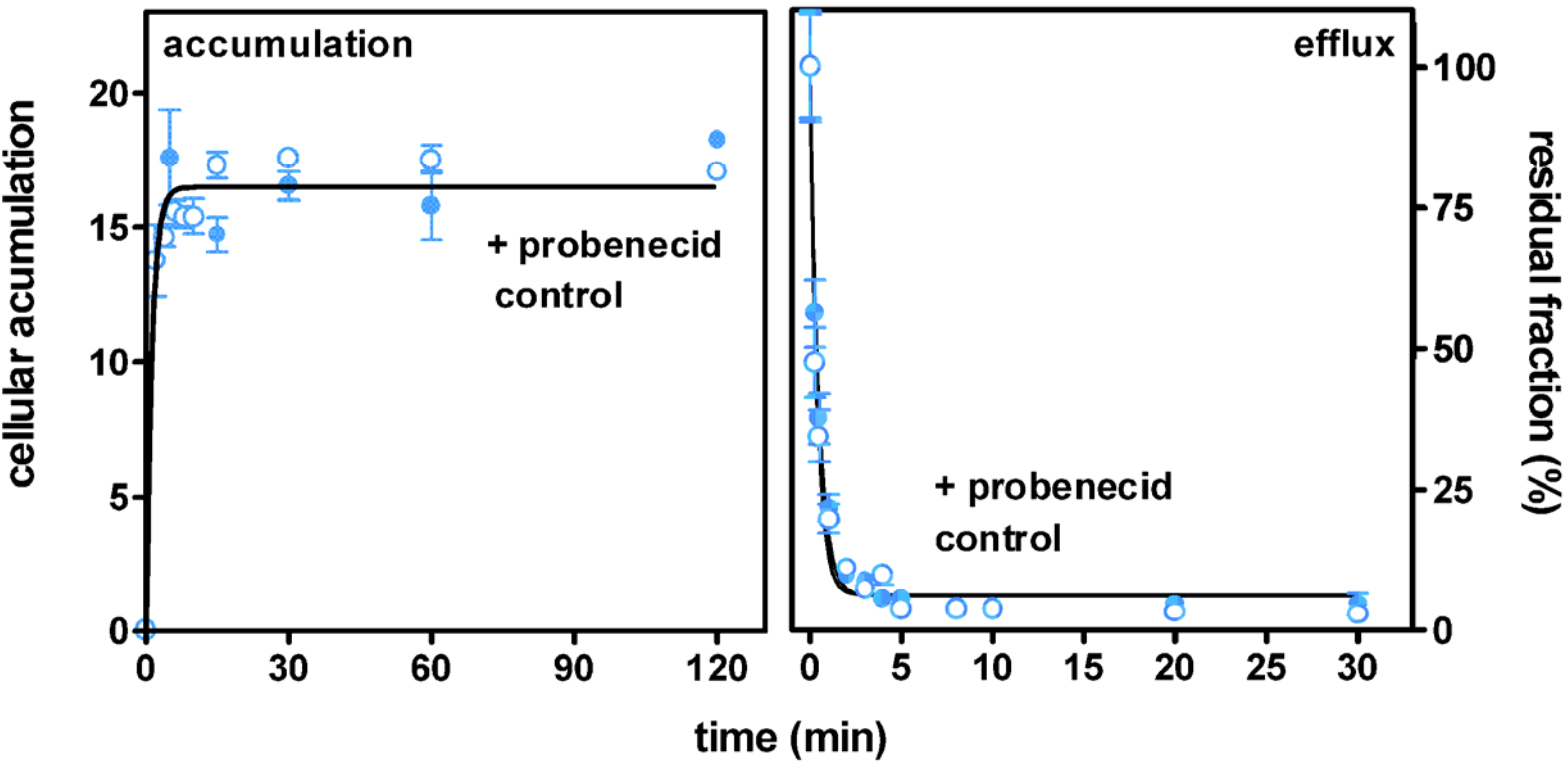


extracell. conc. 17 mg/L; probenecid 5 mM

Michot et al. (2004) AAC 48:2673-82

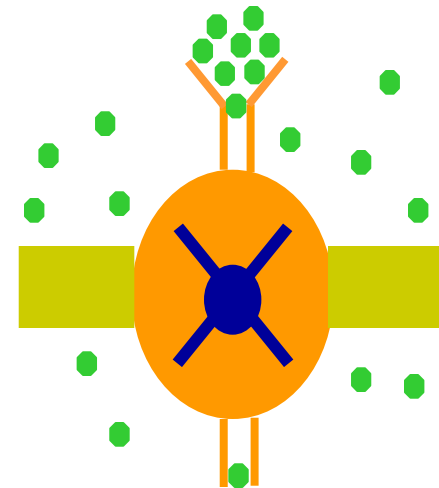
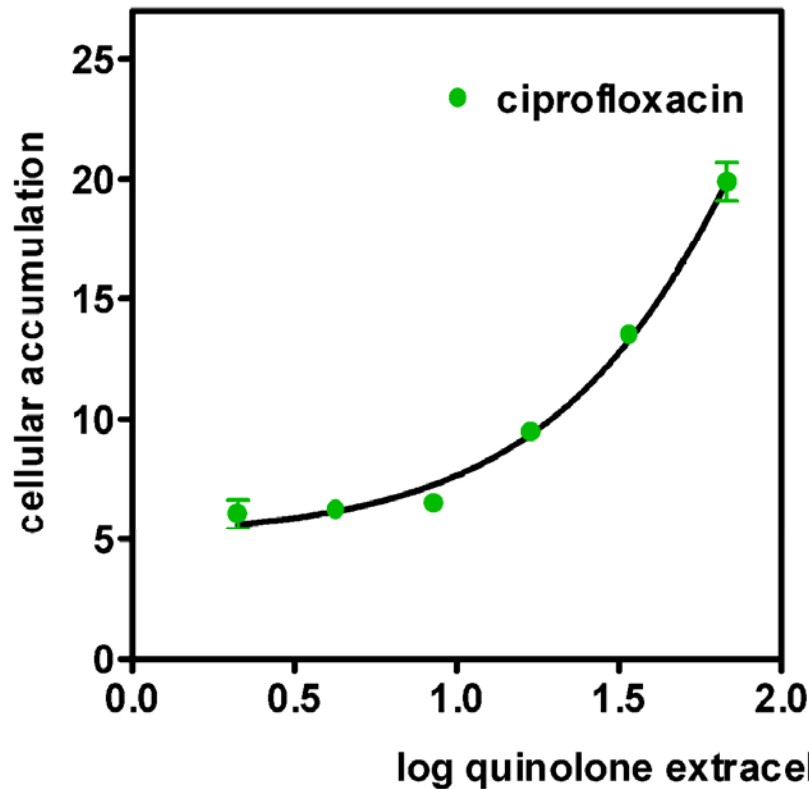
# Kinetics of accumulation and efflux for moxifloxacin

neither accumulation nor efflux affected



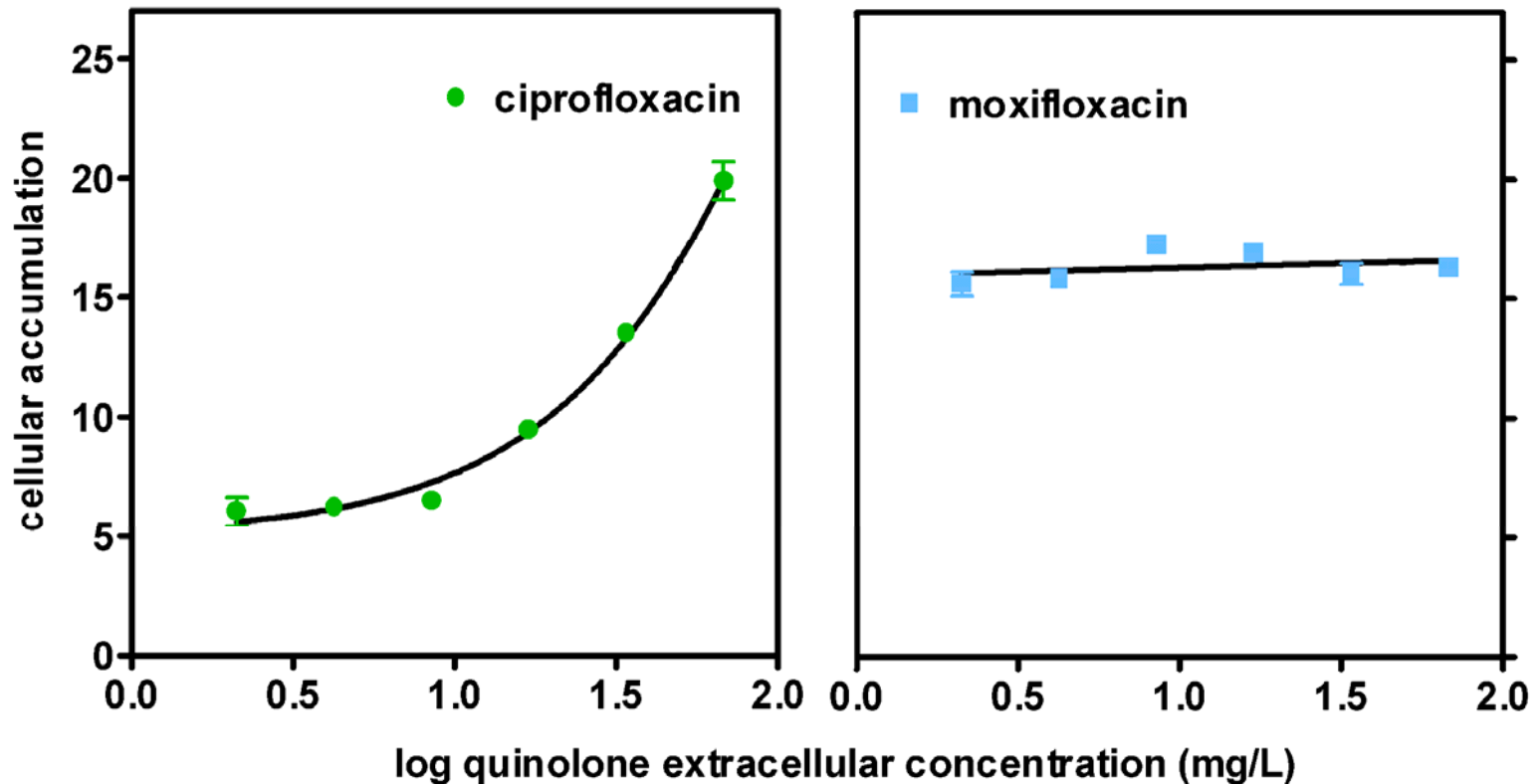
# Quinolones as inhibitors of ciprofloxacin efflux

- ciprofloxacin efflux inhibited by ciprofloxacin



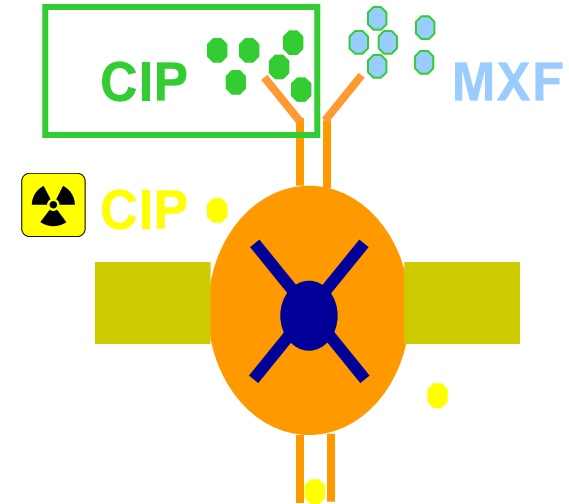
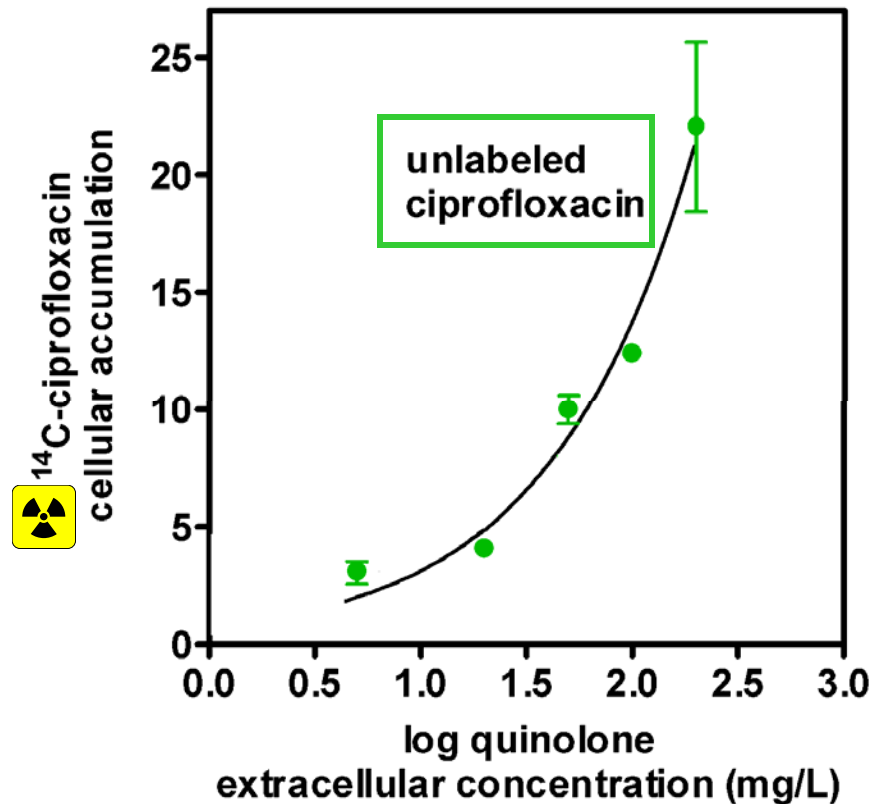
# Quinolones as inhibitors of ciprofloxacin efflux

- ciprofloxacin efflux inhibited by ciprofloxacin
- moxifloxacin not affected



# Quinolones as inhibitors of ciprofloxacin efflux

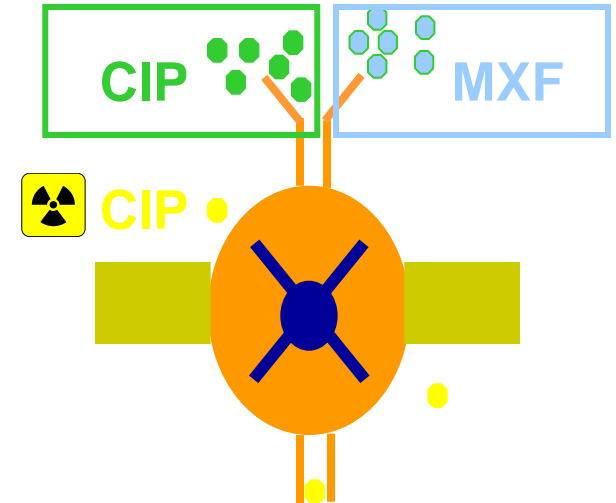
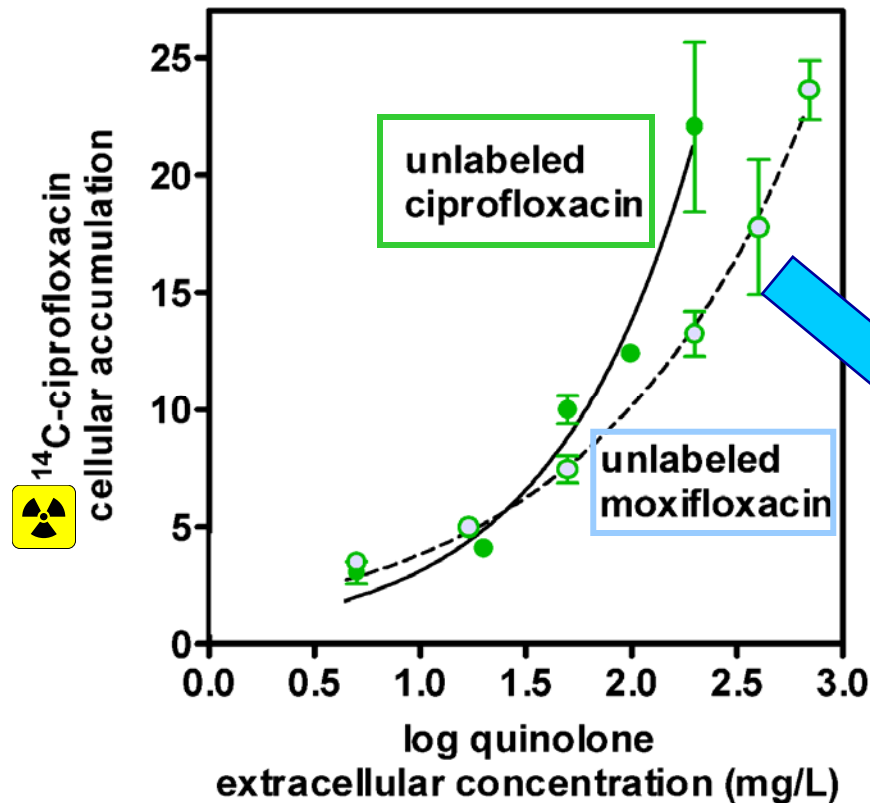
- ciprofloxacin efflux inhibited by ciprofloxacin  
moxifloxacin





# Quinolones as inhibitors of ciprofloxacin efflux

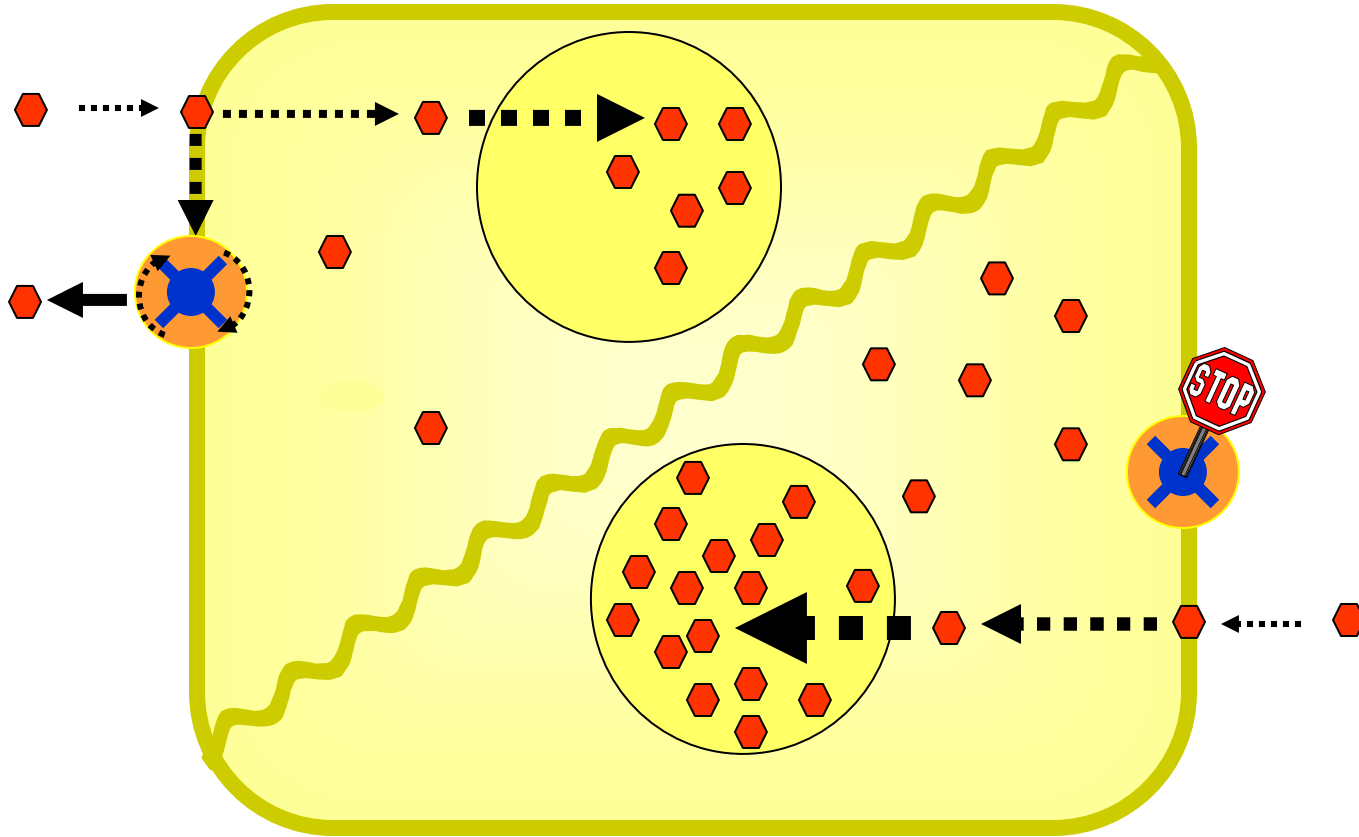
- ciprofloxacin efflux inhibited by ciprofloxacin  
moxifloxacin



**moxifloxacin  
also able  
to interact  
with the transporter !**

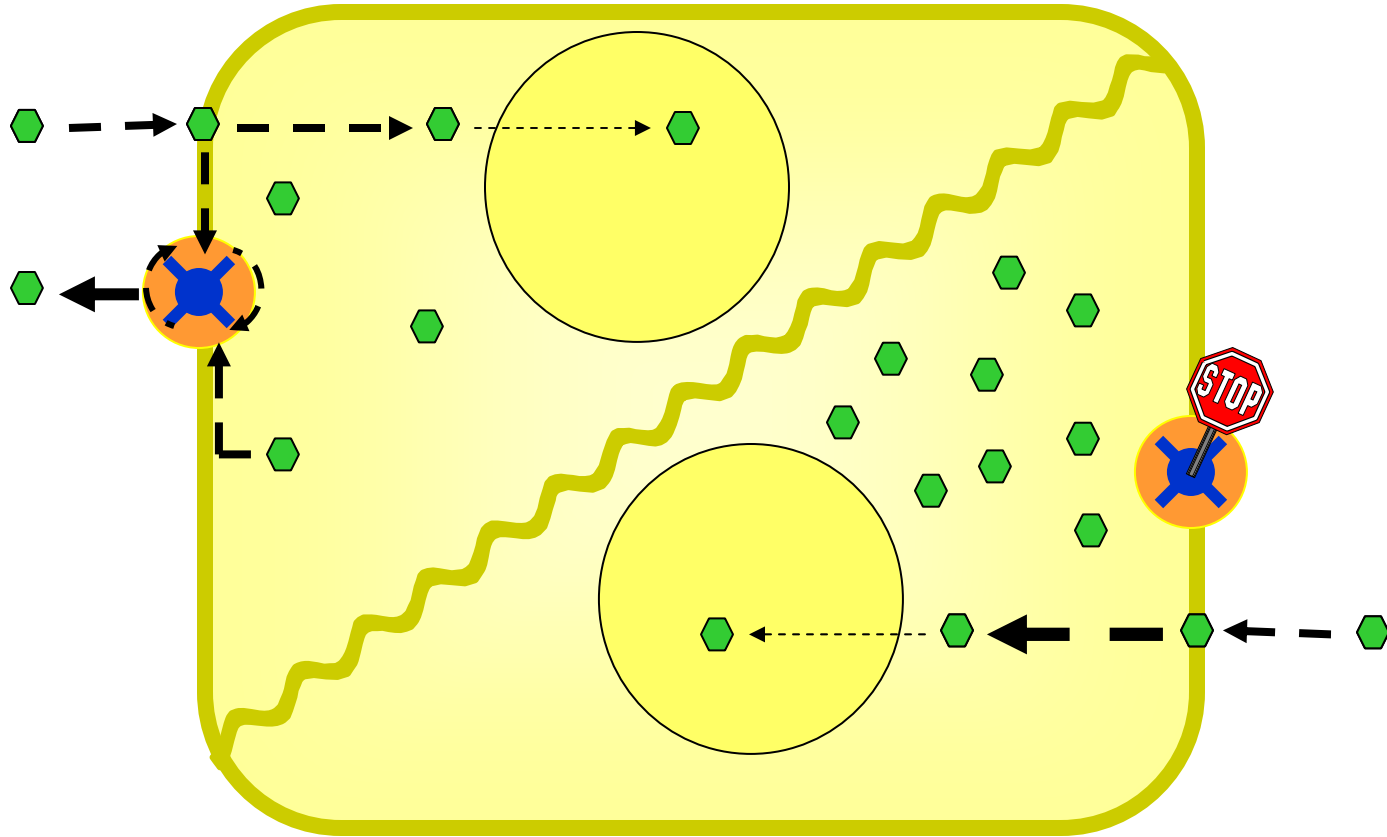
# Azithromycin, 'kick-back' model

Gaj *et al.* (1998) Biochem. Pharmacol. 55:1199-211



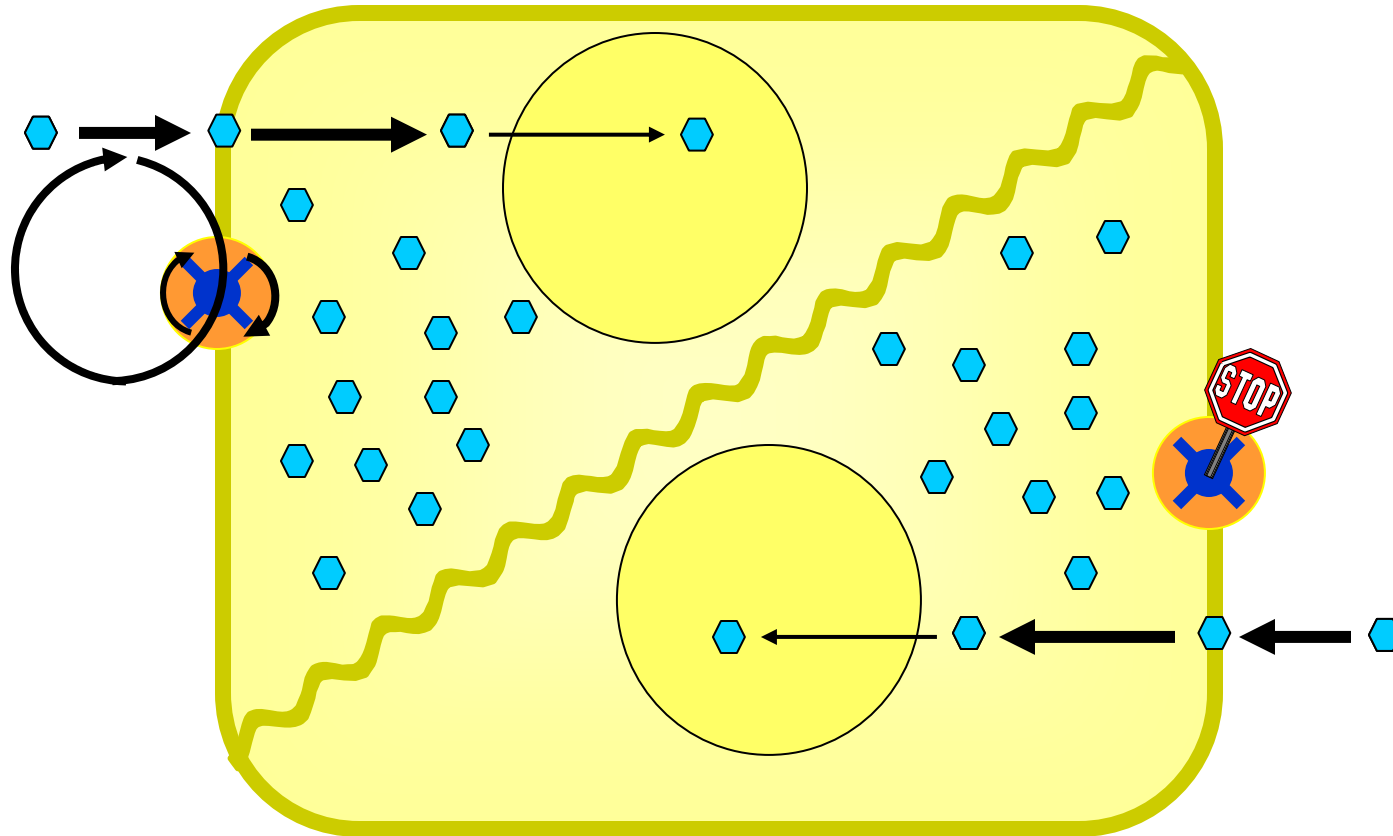
# Ciprofloxacin, classical model

Kolaczkowski & Goffeau (1997) Pharmacol. Ther. 76:219-42



# Moxifloxacin, 'futile-cycle' model

Eytan *et al.* (1996) JBC 271:12897-902



# Pharmacological consequences of antibiotic transport



# Models of intracellular infection

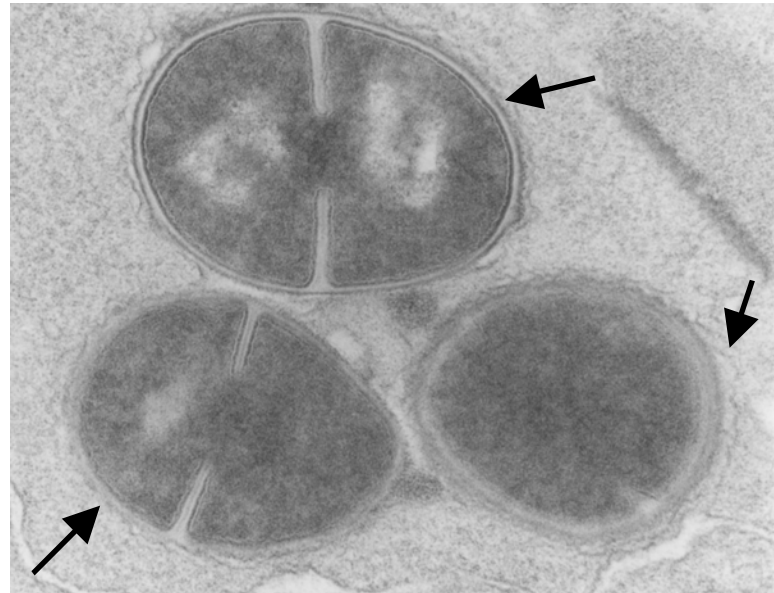
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*L. monocytogenes*



cytosol

*S. aureus*



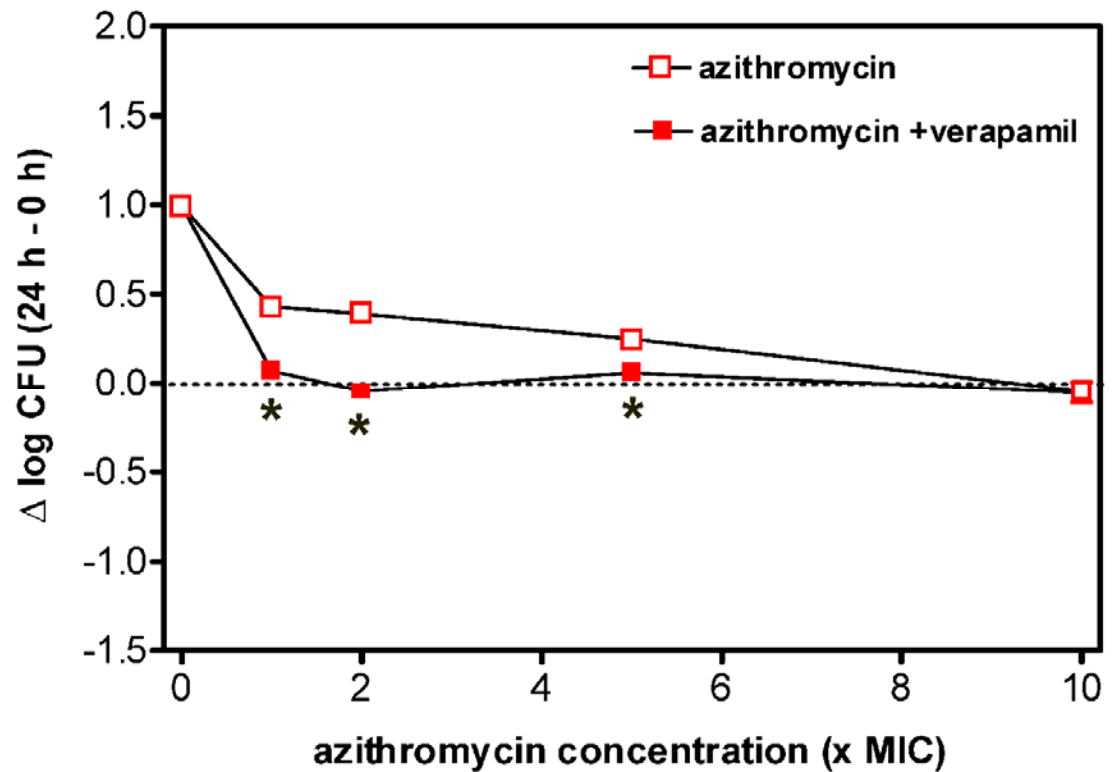
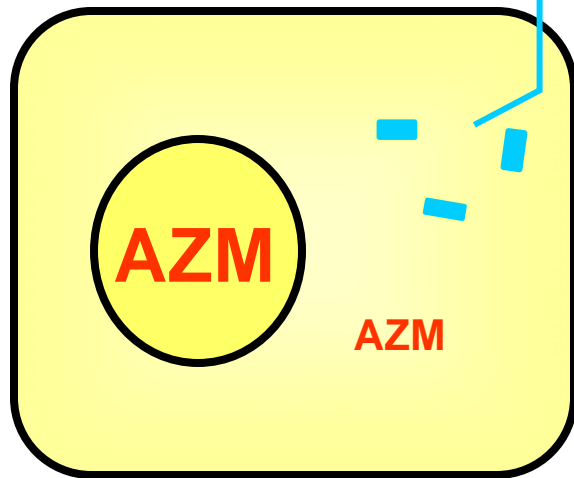
phagolysosomes

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# Influence of pump inhibitors on intracellular activity

## azithromycin and *L. monocytogenes*

*L. monocytogenes*

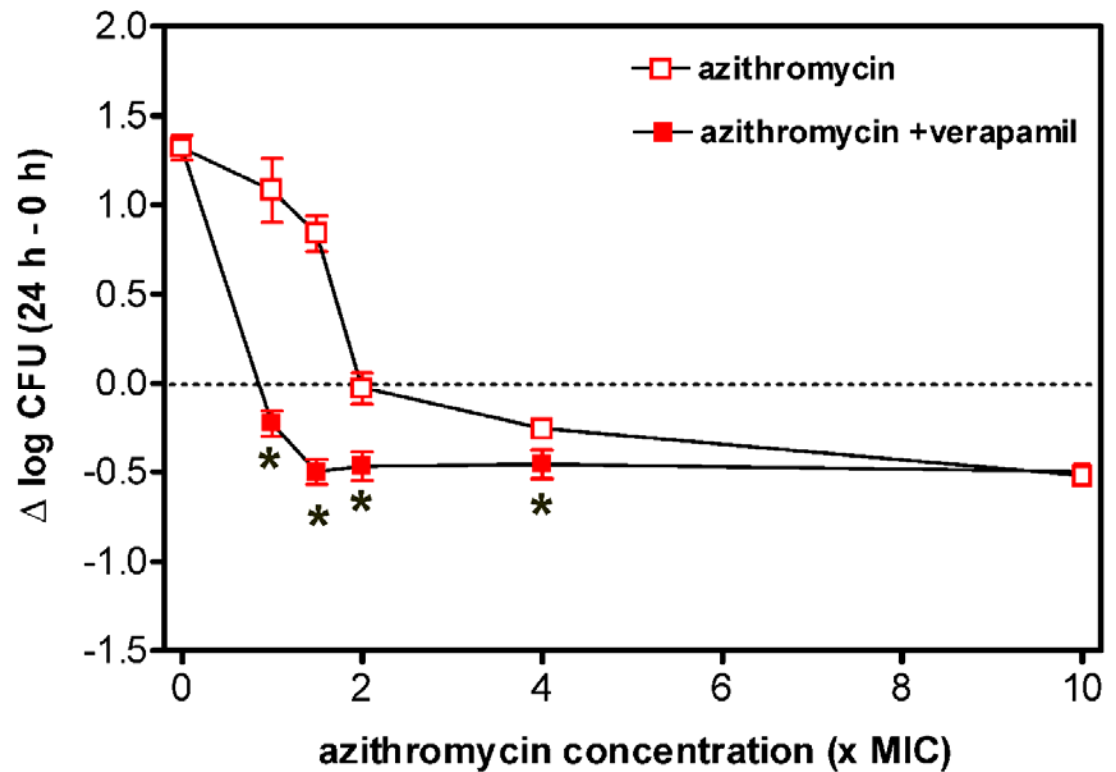
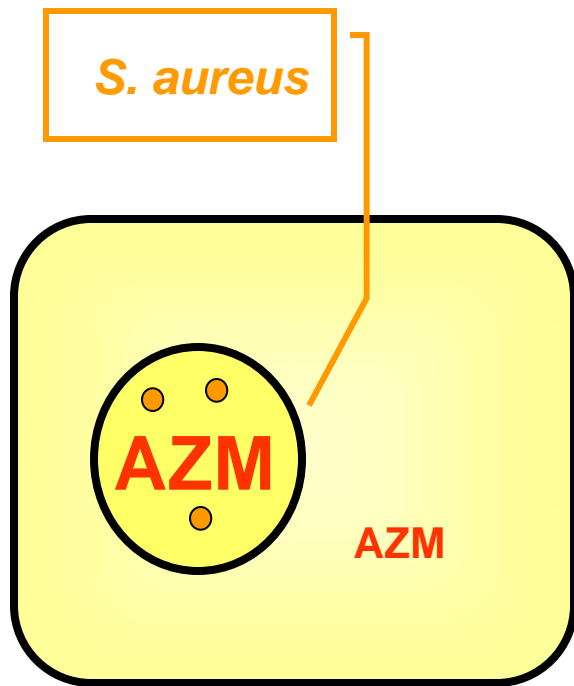


verapamil 20  $\mu\text{M}$ ; 24 h

Seral et al. (2003) JAC 51:1167-73

# Influence of pump inhibitors on intracellular activity

## azithromycin and *S. aureus*



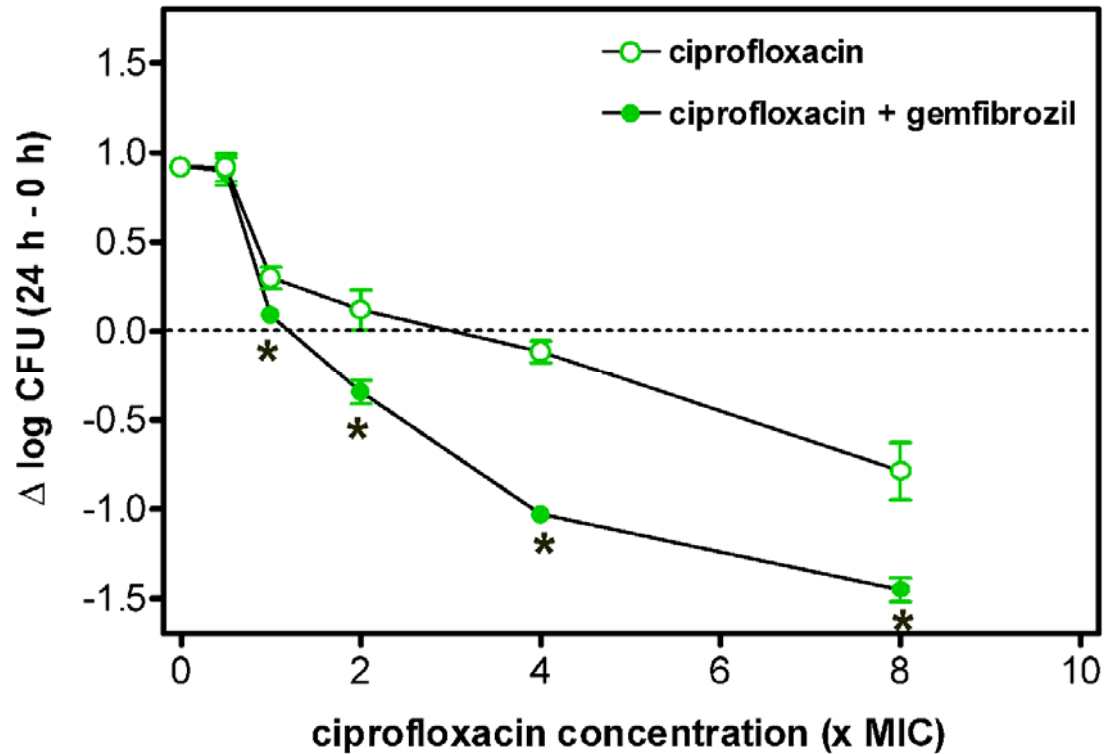


# Influence of pump inhibitors on intracellular activity

## ciprofloxacin and *L. monocytogenes*

*L. monocytogenes*

CIP

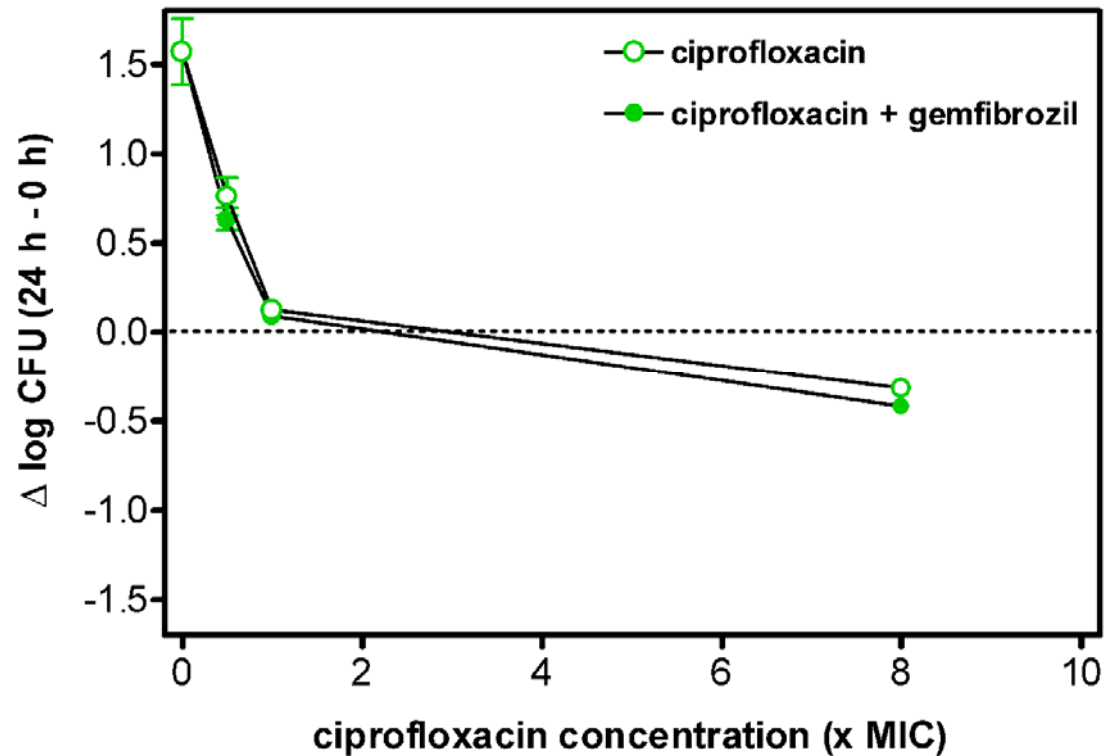
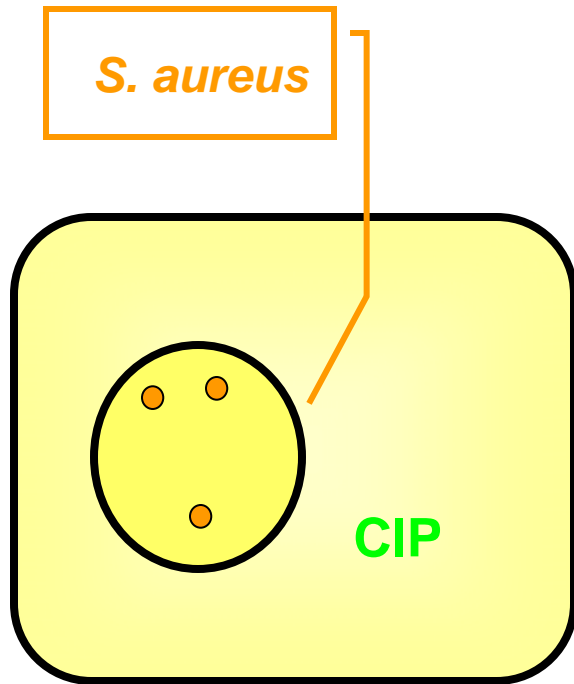


gemfibrozil 250  $\mu\text{M}$ ; 24 h

Seral et al. (2003) JAC 51:1167-73

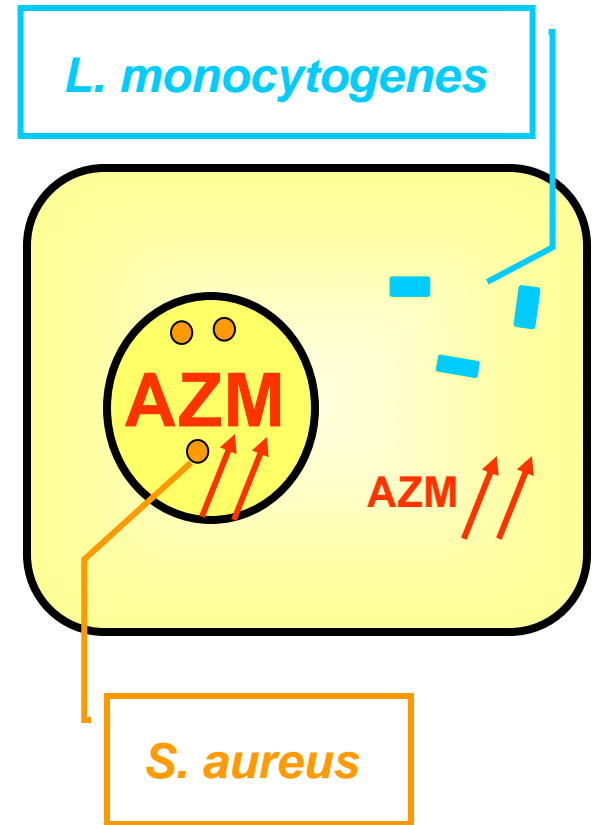
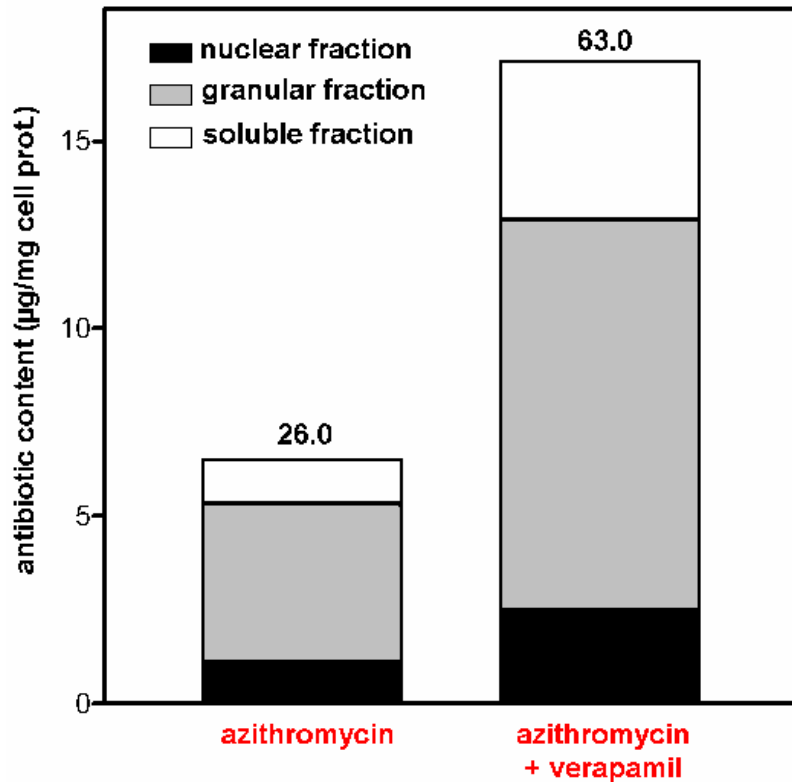
# Influence of pump inhibitors on intracellular activity

## ciprofloxacin and *S. aureus*



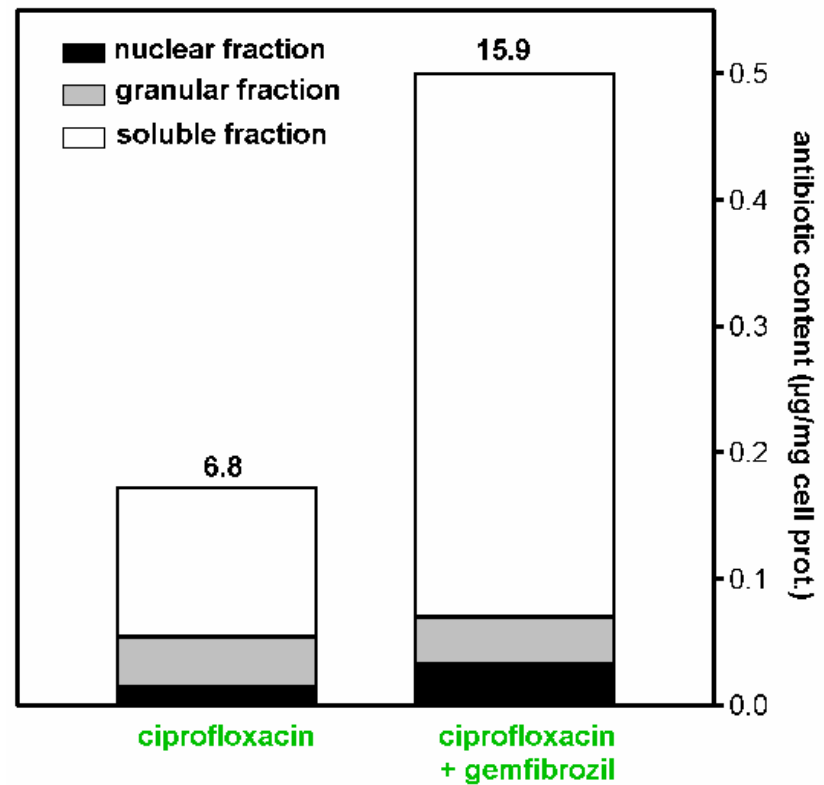
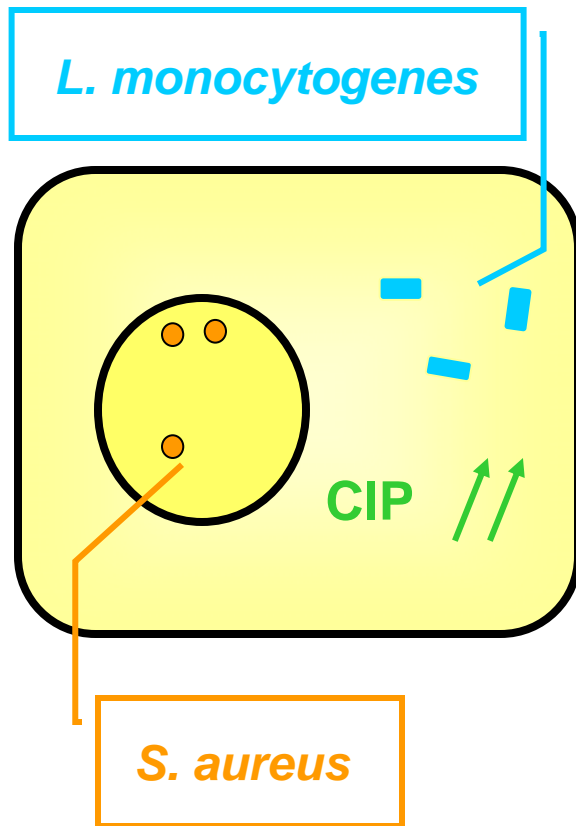
# Influence of pump inhibitors on antibiotic distribution

**verapamil enhances azithromycin concentration  
In cytosol and vacuoles**



# Influence of pump inhibitors on antibiotic distribution

## gemfibrozil enhances ciprofloxacin cytosolic content



# Unexpected antibiotic substrates

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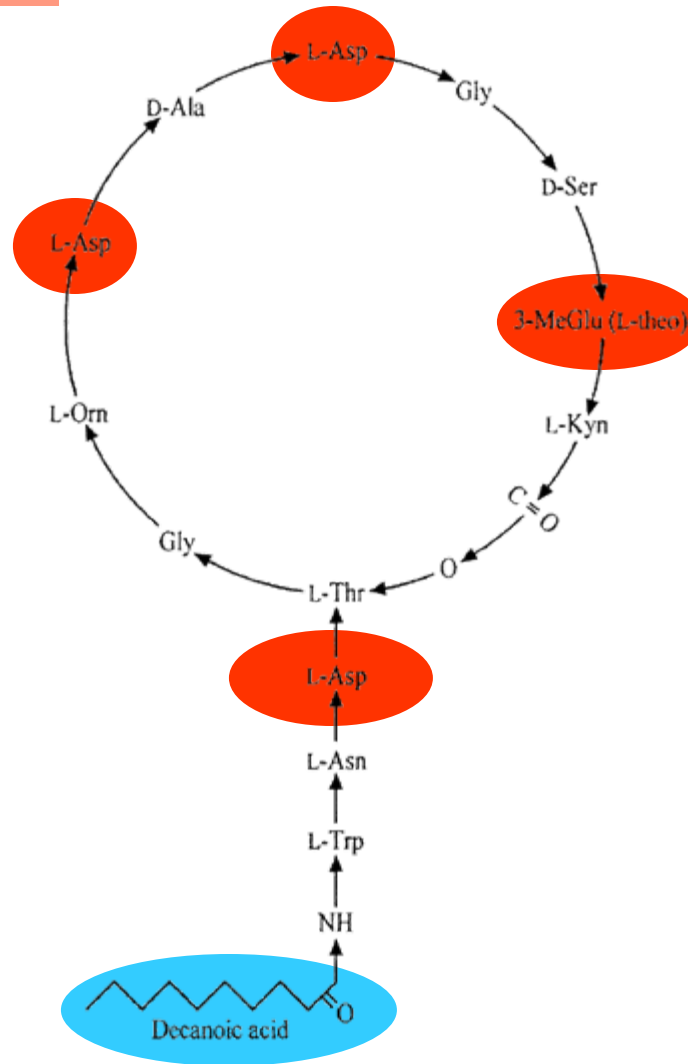
# Unexpected substrate

daptomycin

Log P = - 4  
Log D = -9.5 at pH 7

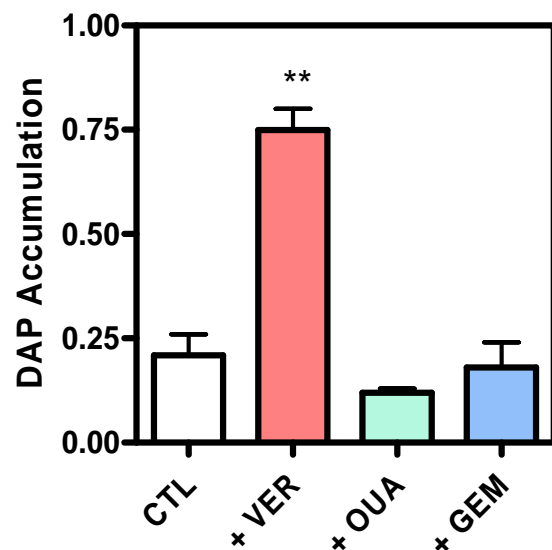
Highly polar molecule !

lipophilic chain

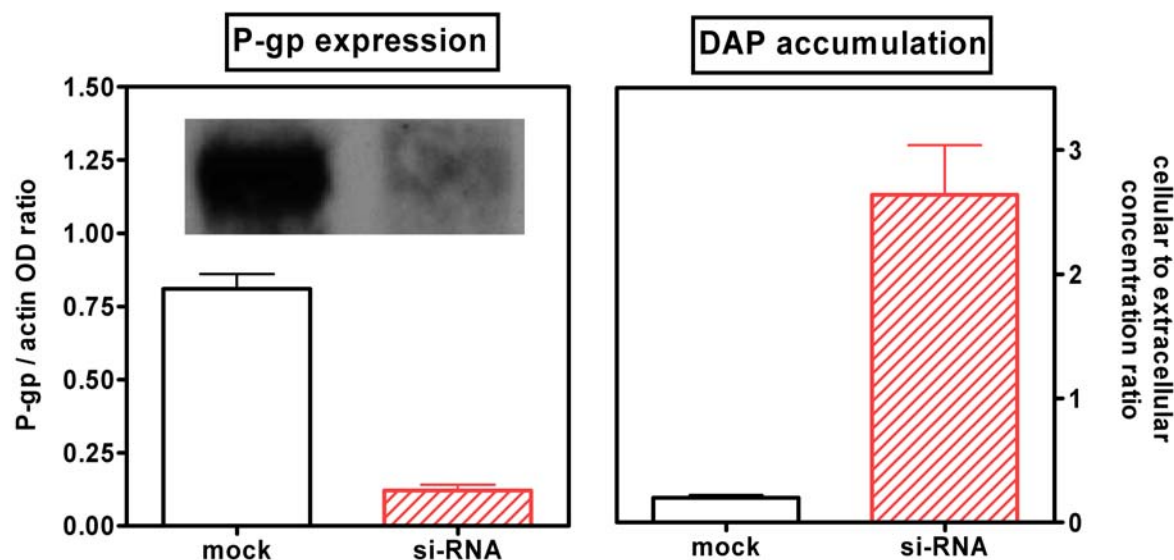


polyanionic  
cyclic peptide

# Daptomycin is substrate of P-gp

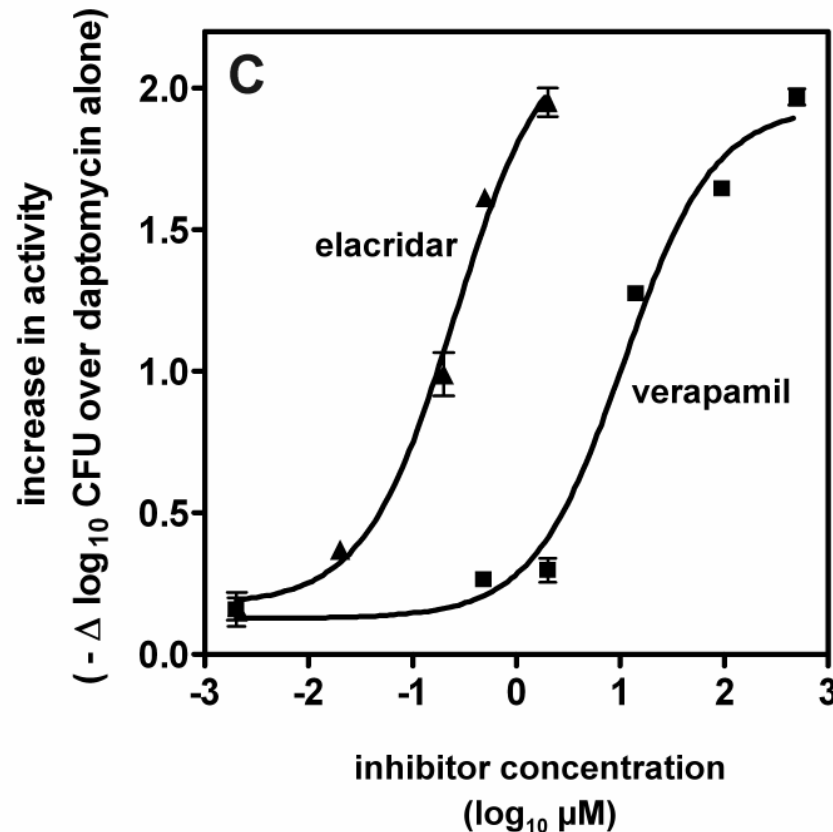


Daptomycin accumulation proportional to P-gp activity and expression level



# Daptomycin is substrate of P-gp

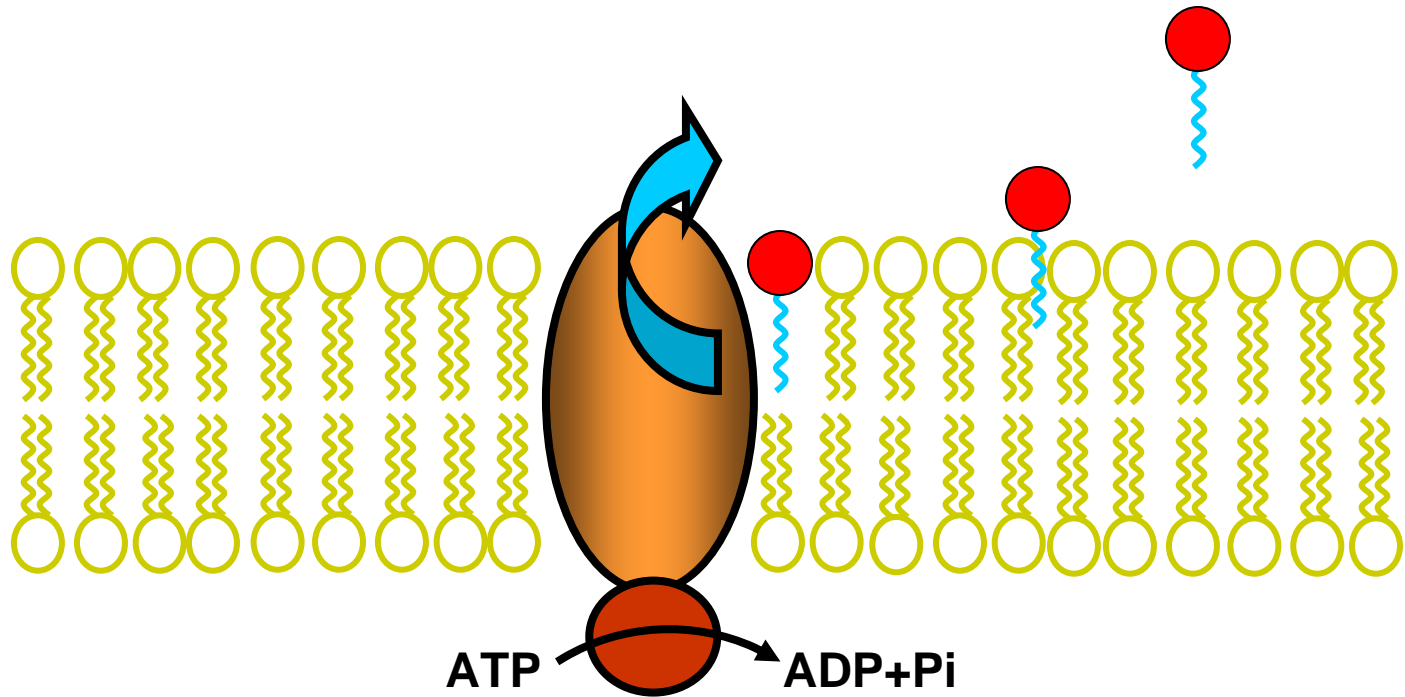
Daptomycin intracellular activity is increased in the presence of P-gp inhibitors





# Putative mechanism

## of daptomycin transport by P-gp

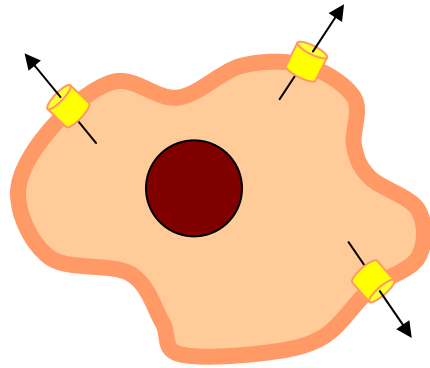


anchoring in the membrane towards the hydrophobic chain  
and extrusion from the membrane

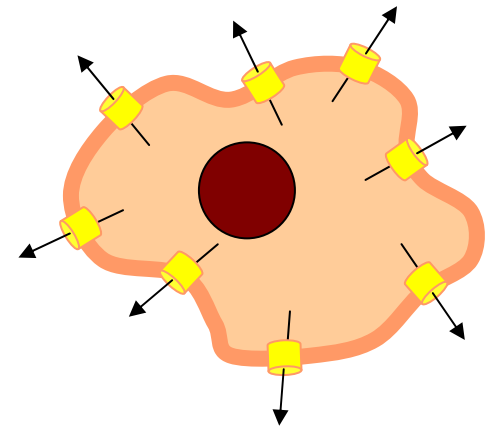
# Can we make eukaryotic cells resistant to antibiotics ?



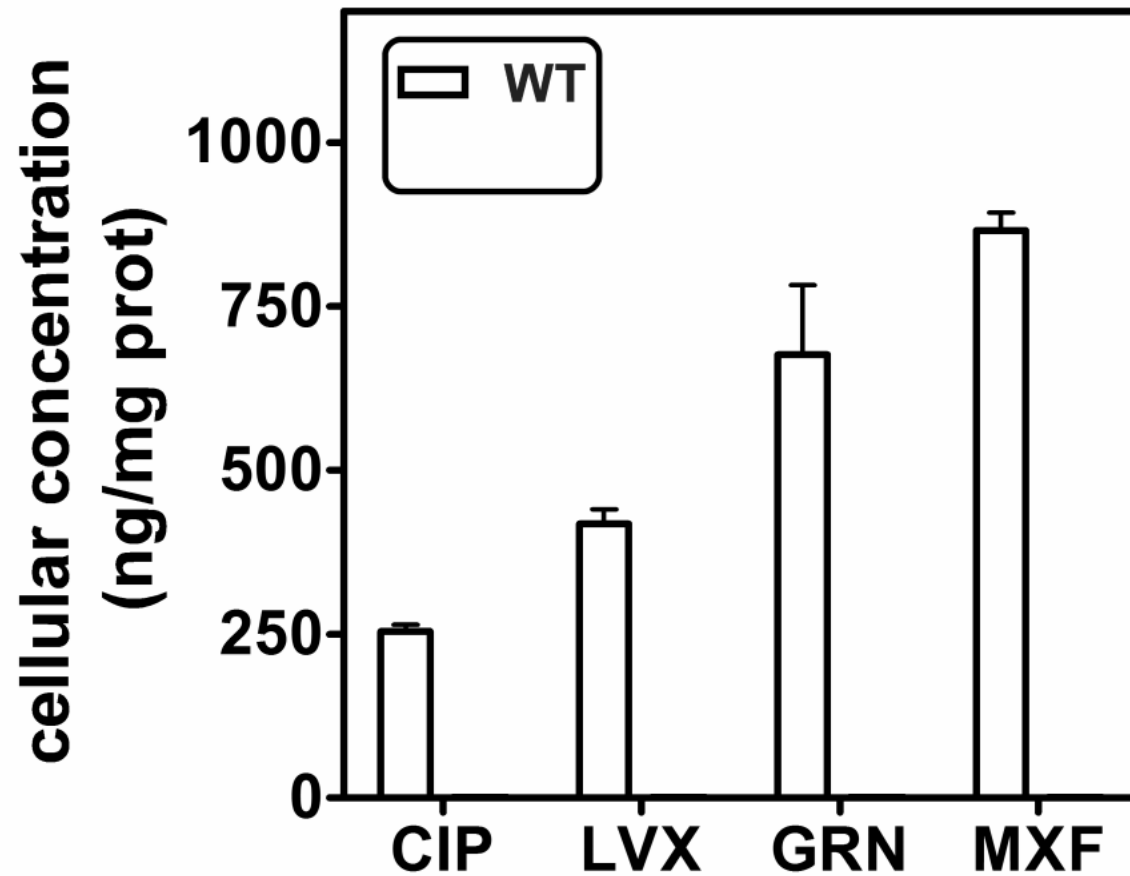
# Can we make eukaryotic cells resistant to antibiotics ?



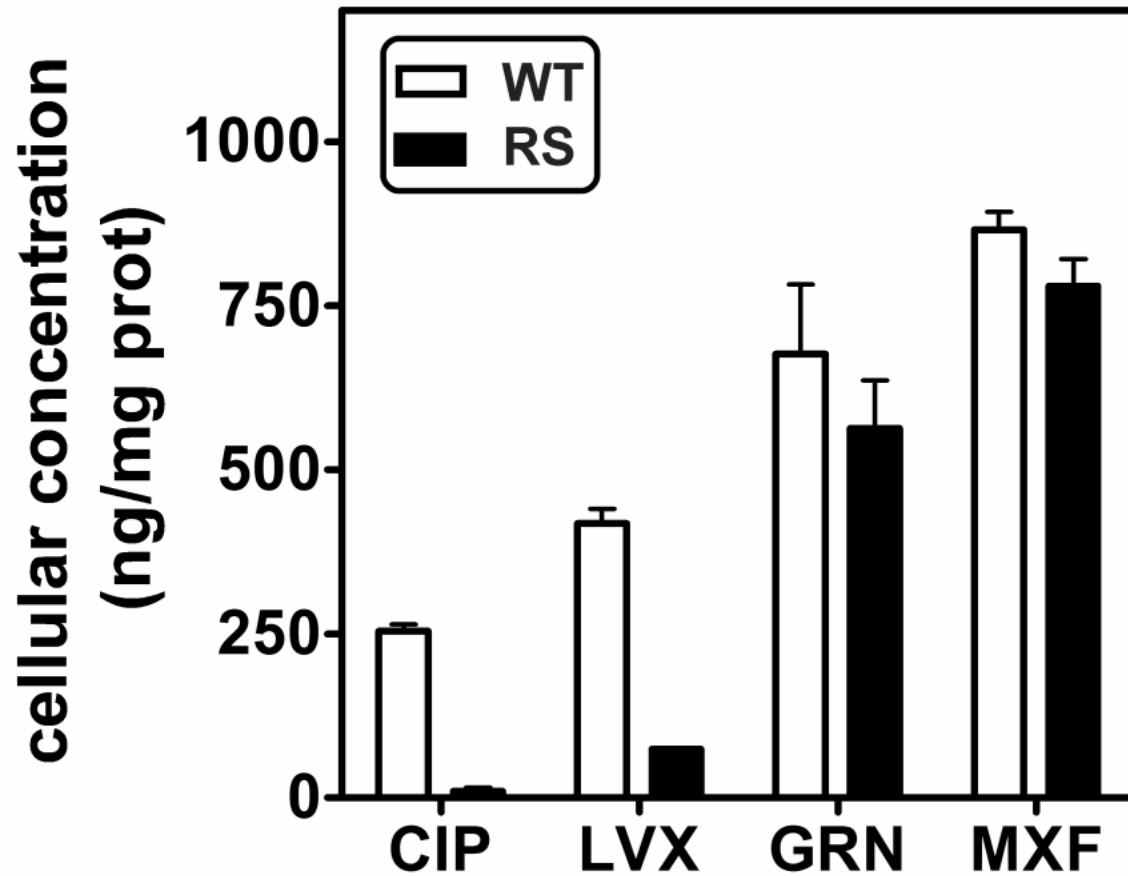
Chronical exposure of J774 macrophages  
to increasing concentrations of ciprofloxacin



# Reduced drug accumulation in resistant macrophages

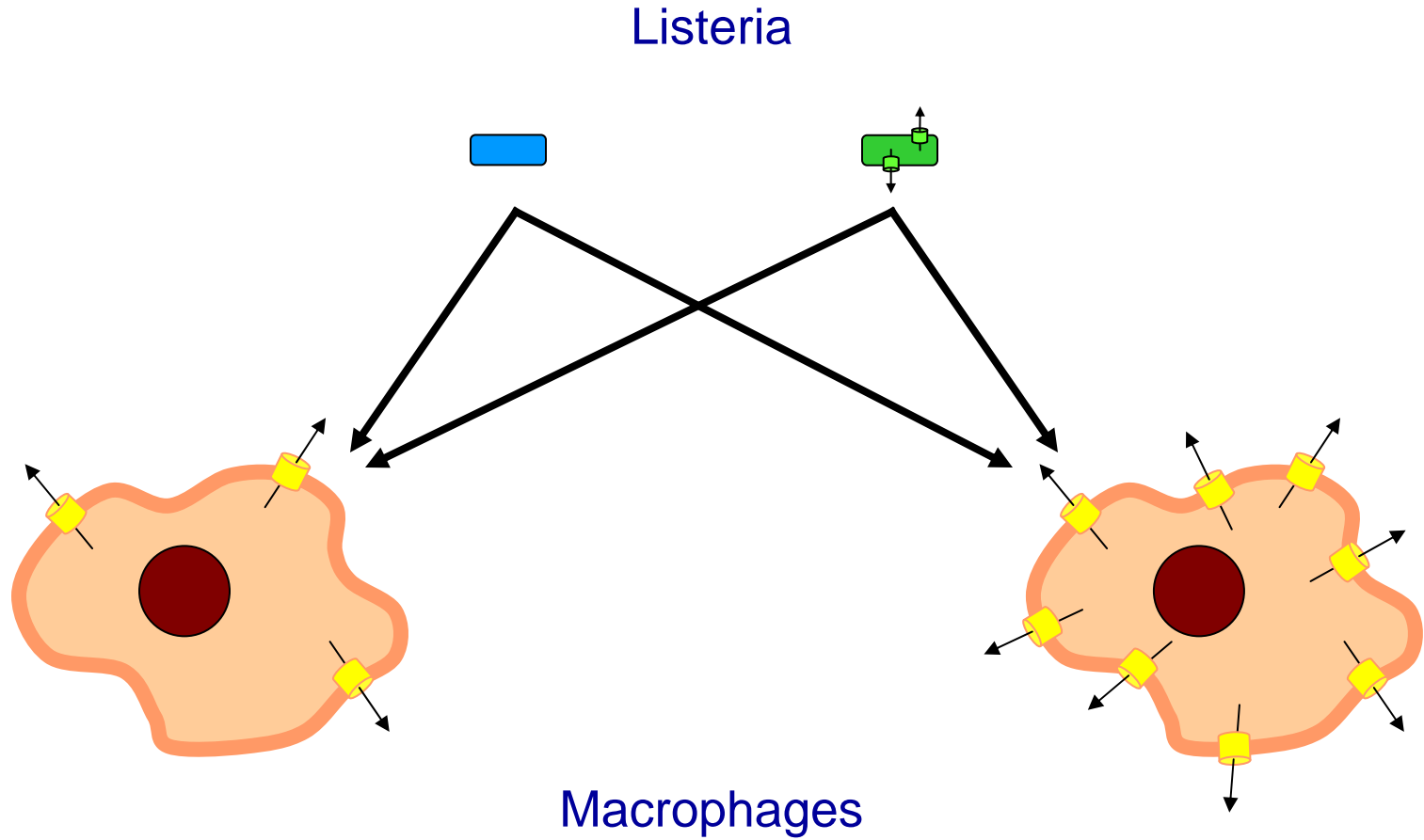


# Reduced drug accumulation in resistant macrophages



# Coworking between bacteria and macrophage pumps to reduce ciprofloxacin activity

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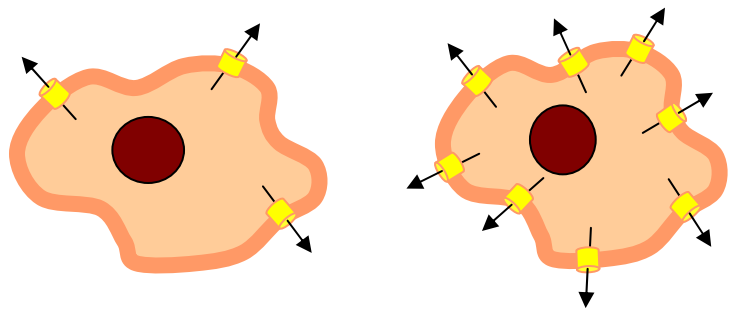


# Coworking between bacteria and macrophage pumps to reduce ciprofloxacin activity

Listeria



Macrophages



FQ	MIC (mg/L)			
	EGD		CLIP	
	Res. (-)	Res. (+)	Res. (-)	Res. (+)
CIP	1.2	1.0	5.0	1.0
MXF	0.6	0.6	0.5	0.25

CIP R; MXF S

FQ	Cellular concentration (ng/mg prot)			
	WT		RS	
	Prob. (-)	Prob. (+)	Prob. (-)	Prob. (+)
CIP	72	263	23	159
MXF	262	208	241	257

CIP R; MXF S

# Coworking between bacteria and macrophage pumps to reduce ciprofloxacin activity

Same substrate specificity of the MFS procaryotic pump and of the ABC eucaryotic pump !

FQ	MIC (mg/L)			
	EGD		CLIP	
	Res. (-)	Res. (+)	Res. (-)	Res. (+)
CIP	1.2	1.0	5.0	1.0
MXF	0.6	0.6	0.5	0.25

CIP R; MXF S

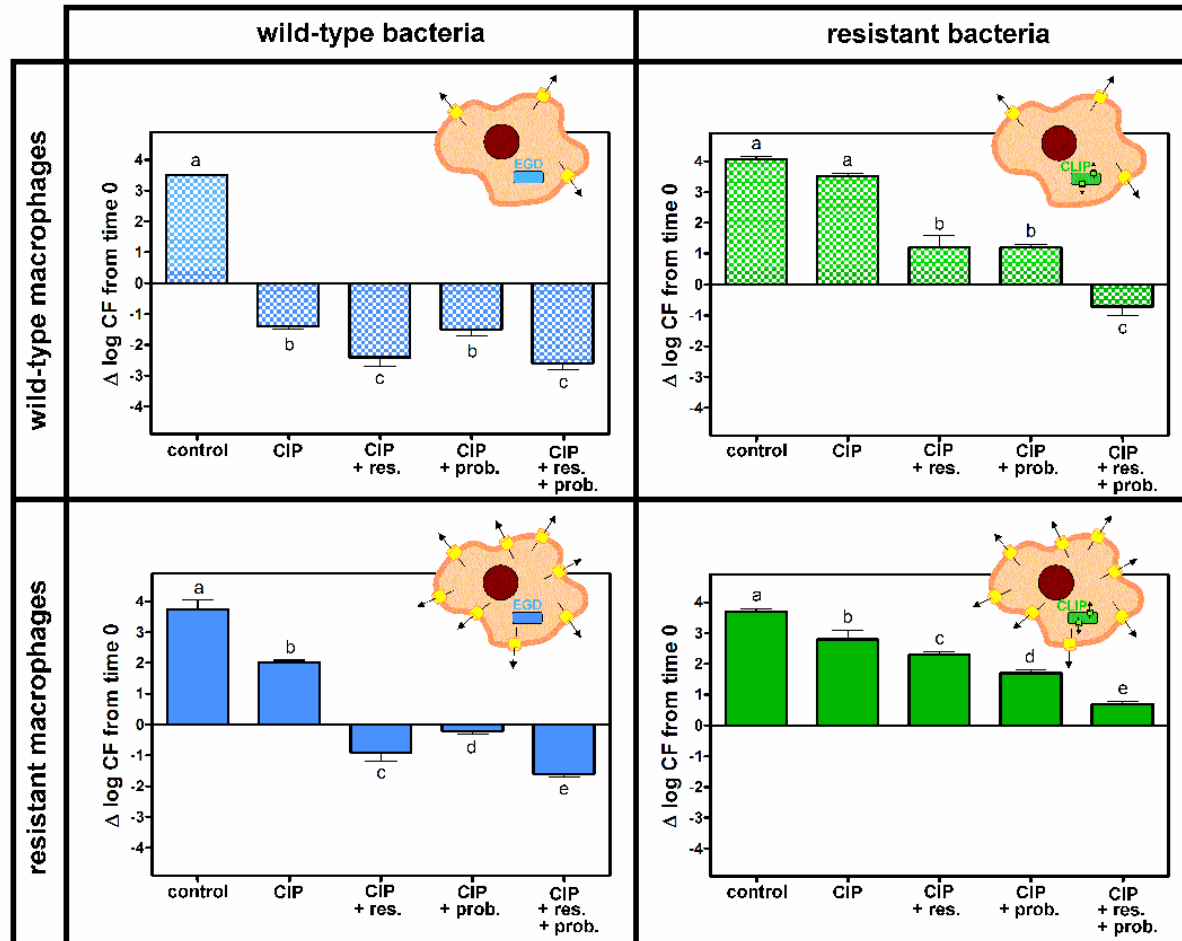
FQ	Cellular concentration (ng/mg prot)			
	WT		RS	
	Prob. (-)	Prob. (+)	Prob. (-)	Prob. (+)
CIP	72	263	23	159
MXF	262	208	241	257

CIP R; MXF S



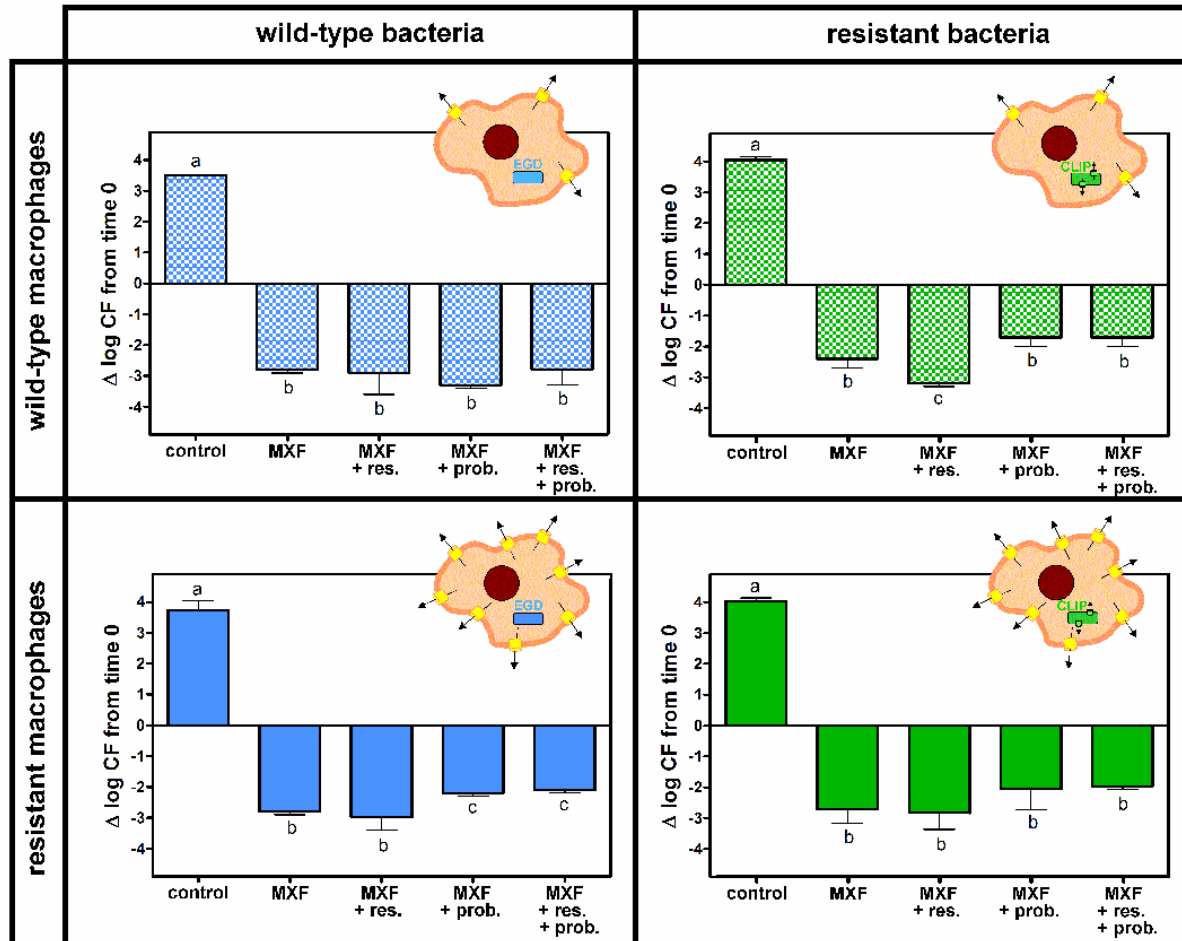
# Coworking between bacteria and macrophage pumps to reduce ciprofloxacin activity

## ciprofloxacin



# No effect of bacteria and macrophage pumps on moxifloxacin activity

## moxifloxacin

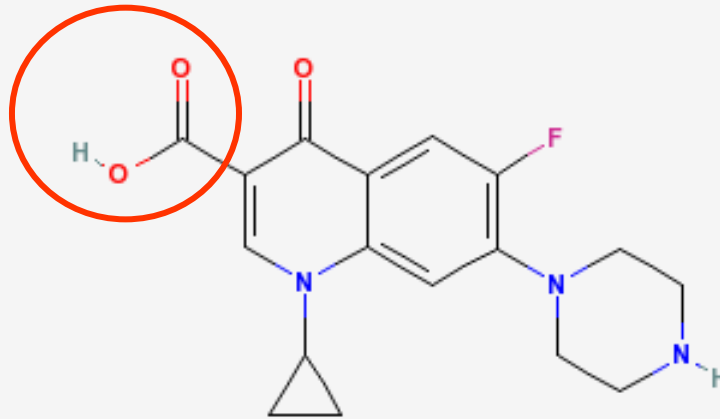


# Competition for transporters as a mechanism of drug interaction

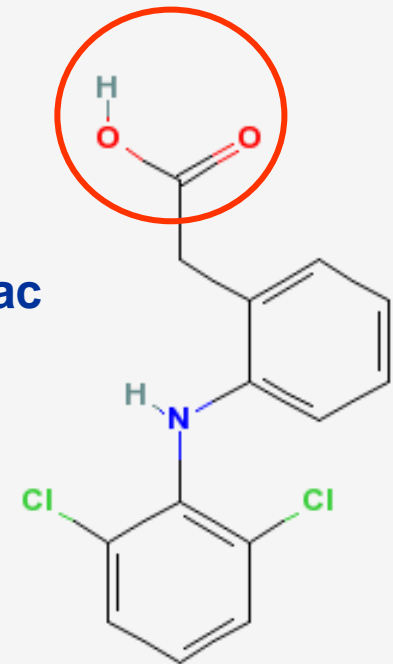


# Ciprofloxacin versus NSAIDs

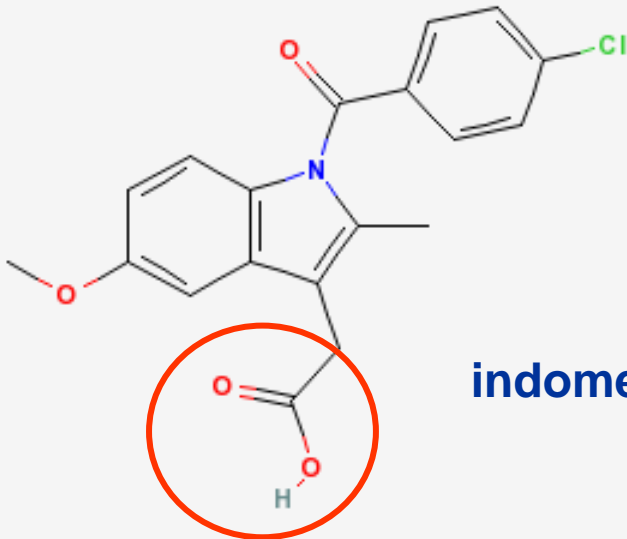
ciprofloxacin



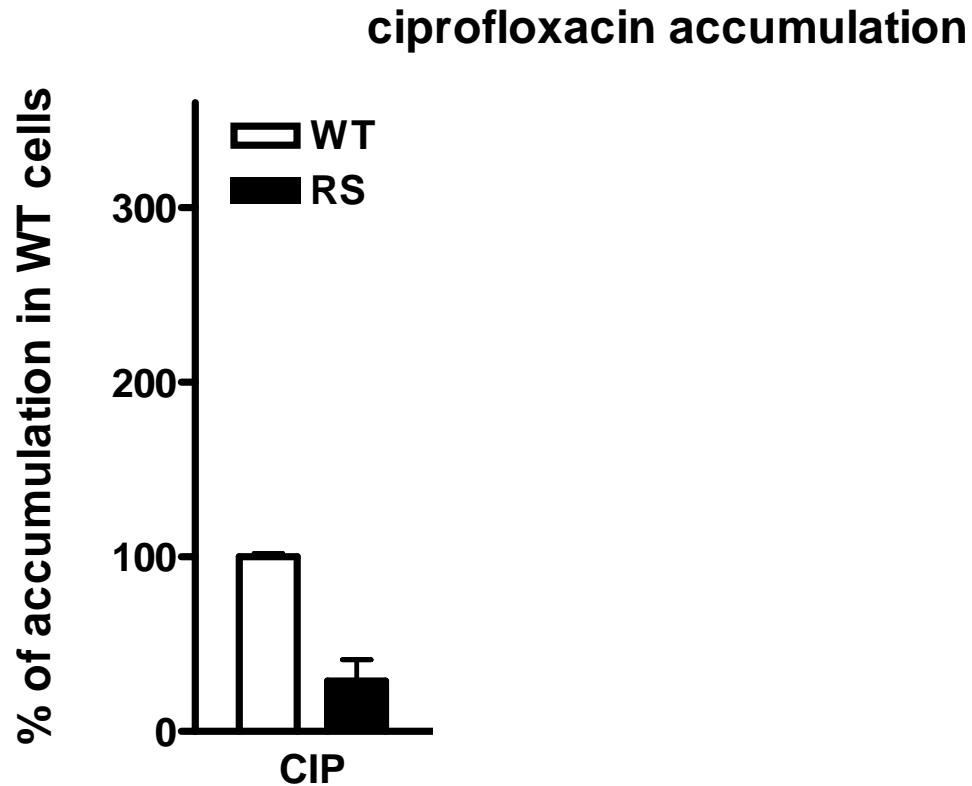
diclofenac



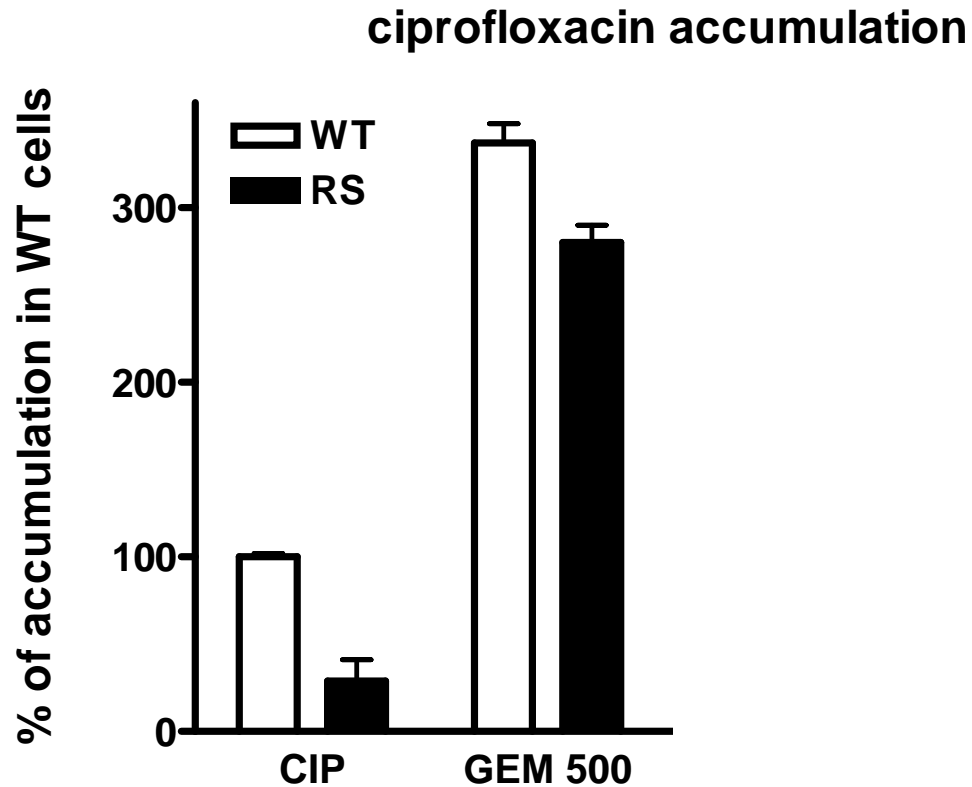
indomethacin



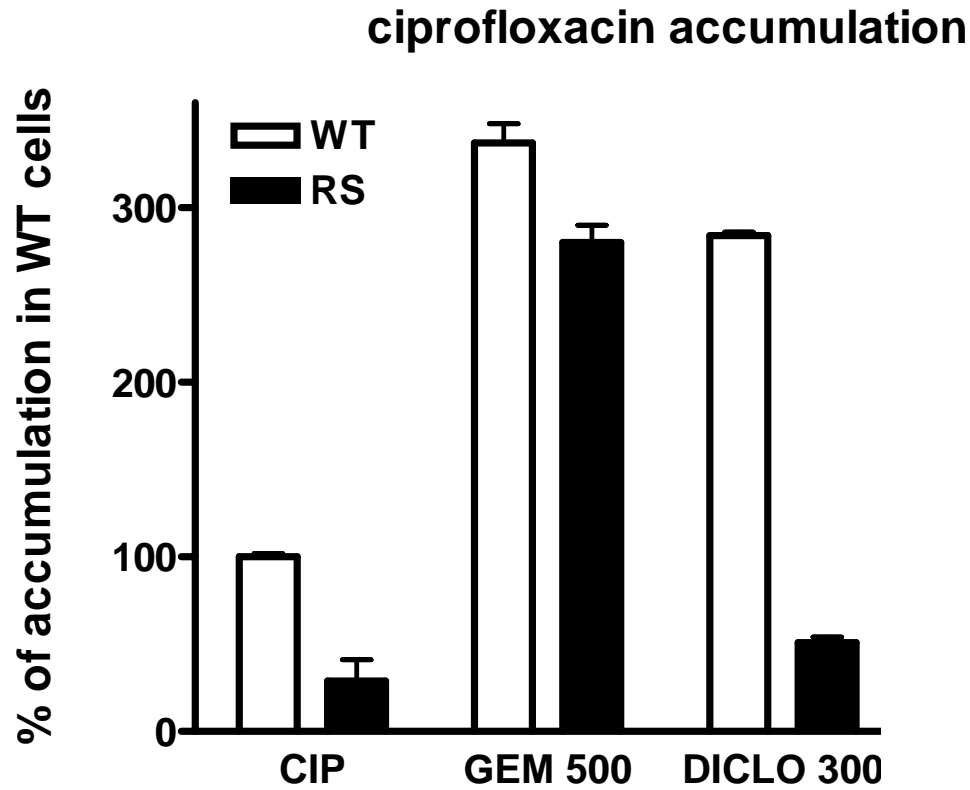
# Ciprofloxacin versus NSAIDs



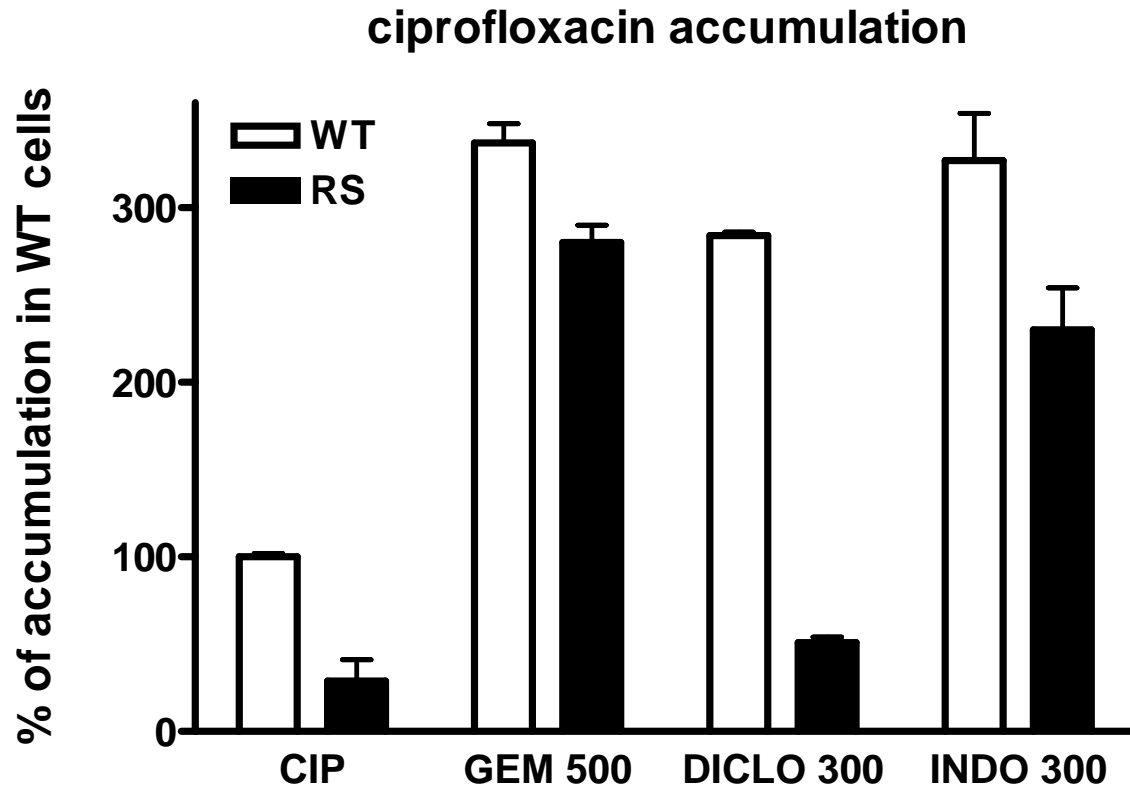
# Ciprofloxacin versus NSAIDs



# Ciprofloxacin versus NSAIDs



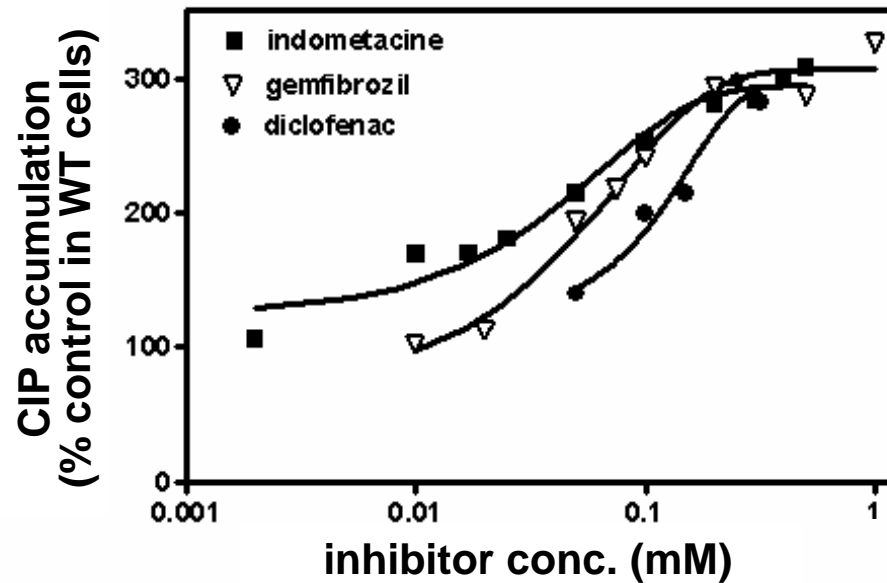
# Ciprofloxacin versus NSAIDs



**indomethacin and, to a lesser extent, diclofenac,  
are inhibitors of ciprofloxacin transport**

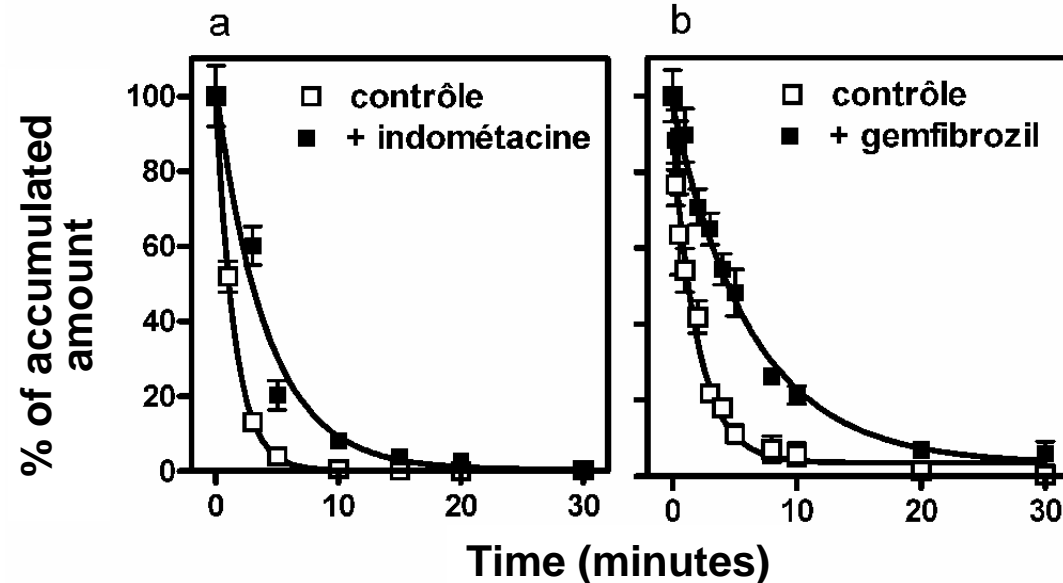


# NSAIDs as inhibitors of ciprofloxacin efflux

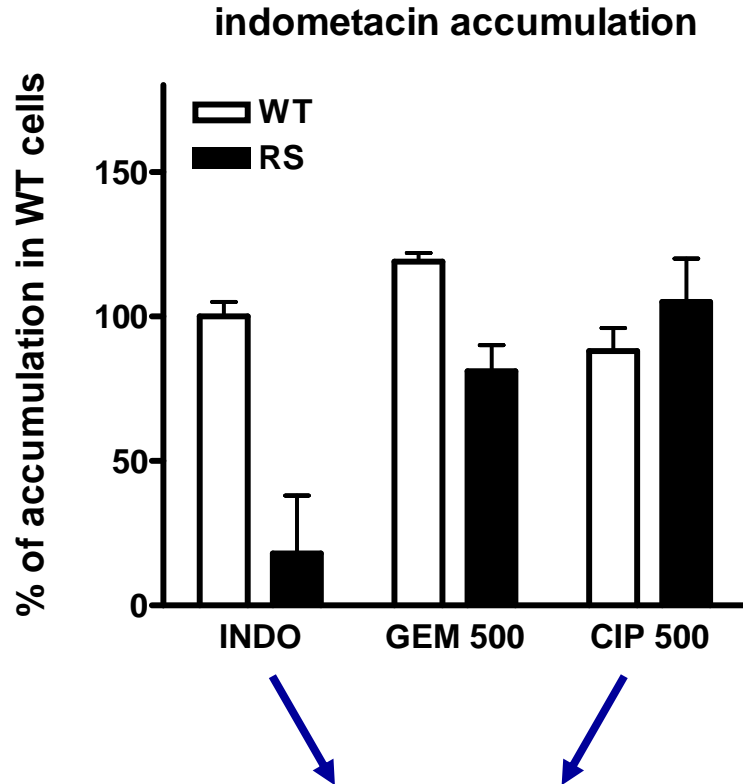


Dose-dependent effect  
on accumulation

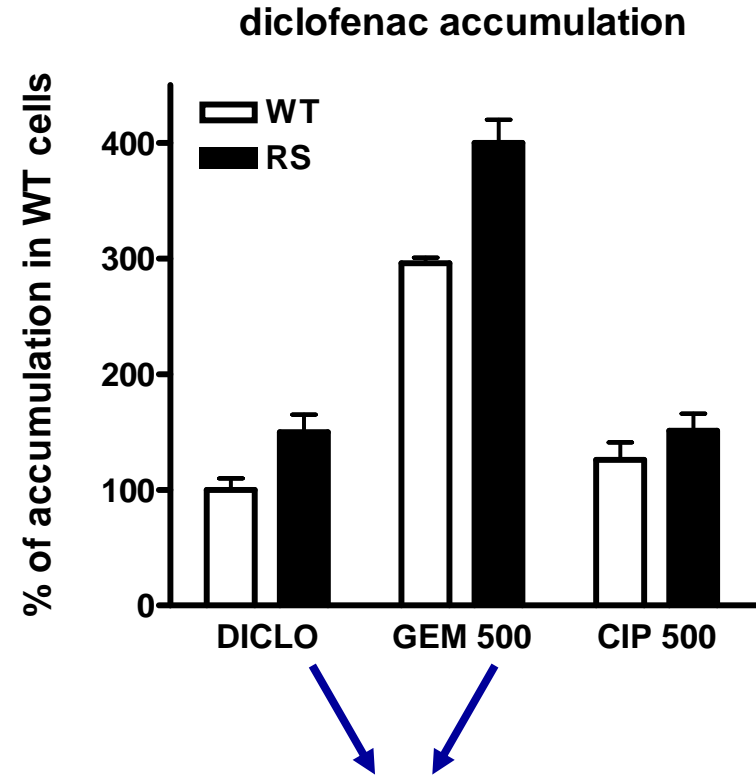
related to slowed-down efflux



# NSAIDs versus Ciprofloxacin



indomethacin shares  
a same transporter  
with ciprofloxacin



diclofenac is substrate  
of another gemfibrozil-inhibitable  
transporter

# Conclusion: avenues for the future

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Do we need to include  
'transport' studies  
with bacteria and eucaryotic cells  
in the early development  
of new antibiotics ?

# The past and present efflux team in Brussels

