



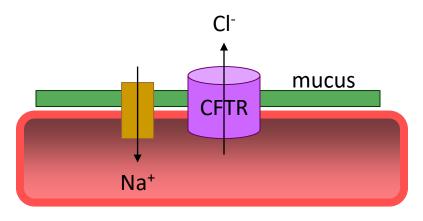
Nuovi approcci terapeutici contro le infezioni da batteri Gram-negativi

Silvia Buroni

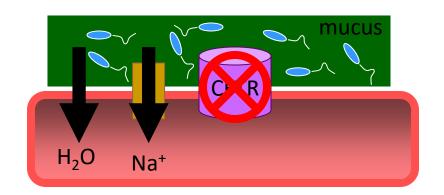
silvia.buroni@unipv.it

Cystic Fibrosis

- Autosomal, recessive disease
- 1 every 2,500 live births
- Caused by a defective <u>Cystic Fibrosis</u> <u>Transmembrane Regulator</u>
- Pulmonary disease
 - Chronic microbial colonization
 - Spectrum of pathogens
 - Pseudomonas aeruginosa
 - Burkholderia cenocepacia

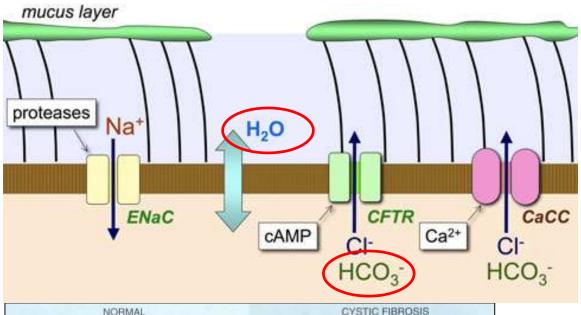


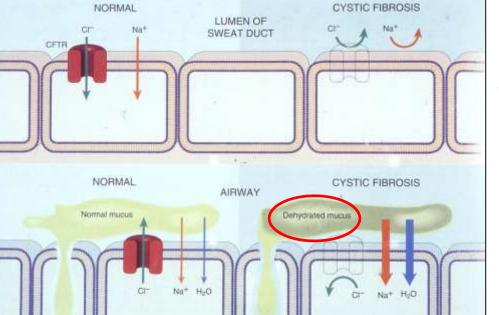
Fully Functioning CFTR



Defective CFTR

Protein Function



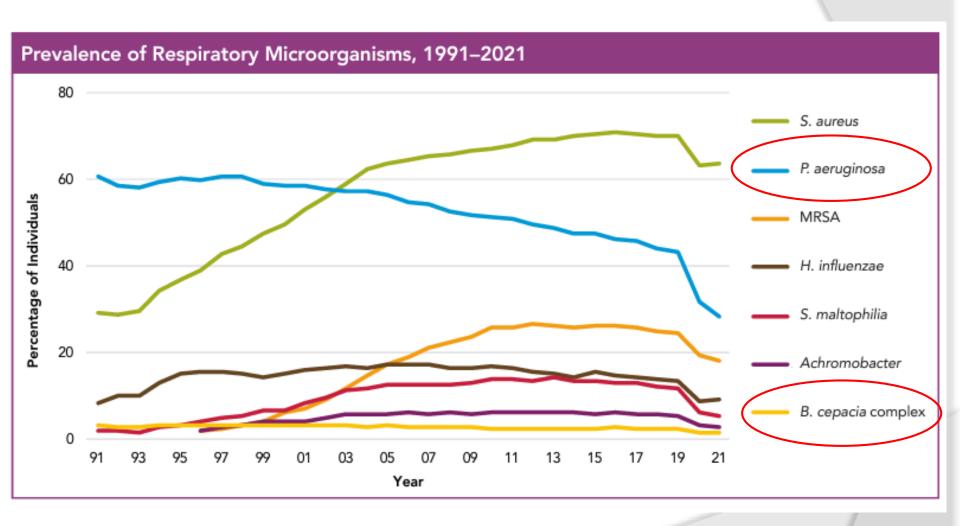


abnormally high chloride concentration in the sweat

most important diagnostic test for CF

airway colonization by bacteria

Infections among cystic fibrosis patients

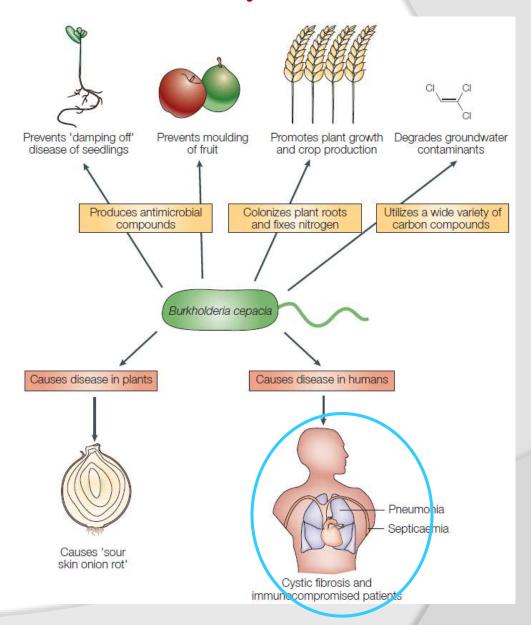


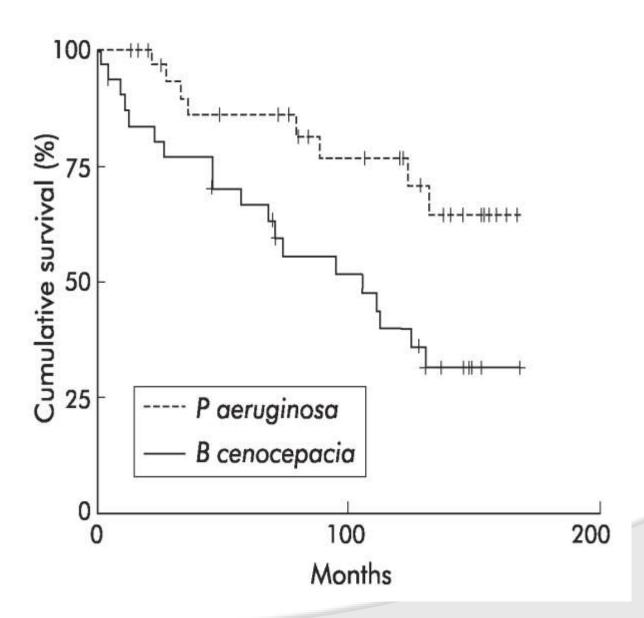
B. cenocepacia: 1-3% patients

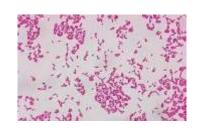
Burkholderia cenocepacia





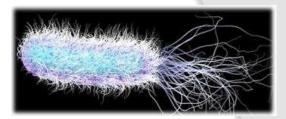






Burkholderia vs Pseudomonas

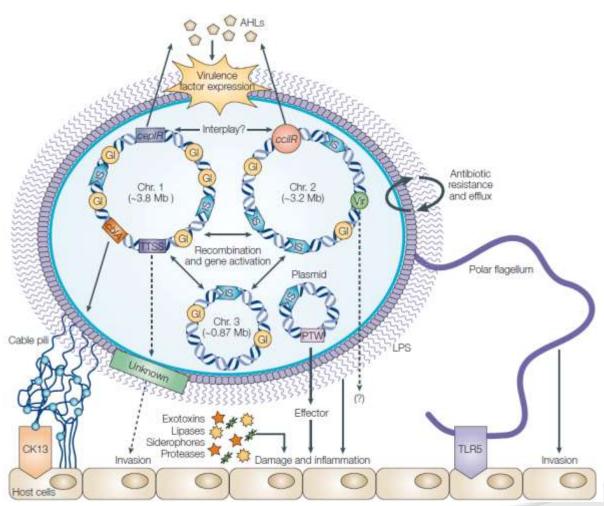






- Opportunistic pathogens
- Ubiquitously distributed in nature (soil, water, plants and industrial settings)
- (i) extraordinary metabolic versatilities
- (ii) abilities to **interact with other bacteria** as well as with their eukaryotic hosts via signal molecules
- (iii) to adapt their genetic repertoire to the needs of a particular niche by **acquiring genetic material** from other bacteria
- ➤ Frequent exchange of genetic material between *P. aeruginosa* an *B. cepacia* → bacteria colonize the same habitats
- B. cepacia is capable of perceiving the QS signals produced by P. aeruginosa but not vice versa

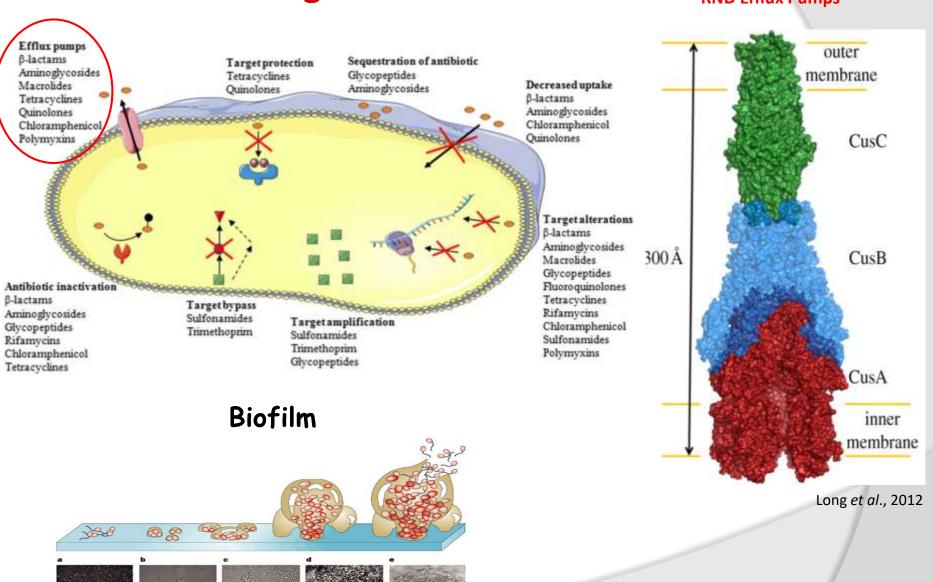
B. cenocepacia Genome Structure and principal Virulence Factors



- -GI= Genomic Islands
- -IS= Insertion Sequence

- Drug resistance;
- 2. Biofilm formation;
- 3. EPS synthesis;
- 4. Cable pili (adherence to mucin and epithelial cells);
- 5. Flagella (invasion);
- 6. Production of catalase and SOD (intracellular survival);
- 6. Secretion of lipases, proteases, hemolysins and siderophores;
- 7. Quorum sensing signals;
- 8. Secretion systems.

Drug Resistance Mechanisms RND Efflux Pumps



Nature Reviews | Drug Discovery

Resistance mechanisms in Burkholderia

Modification of drug target site

- penicillin binding proteins for β-lactams (ΔPBP3 in B. pseudomallei);
- ribosomes for aminoglycosides, macrolides, tetracyclines, and chloramphenicol;
- DNA gyrase for fluoroquinolones (*gyrA* in *B. cepacia*);
- folic acid biosynthesis for trimethoprim and sulphonamides (dhfr in B. cepacia).

Enzymatic drug inactivation

- Overexpression of β-lactamases;
- Aminoglycoside-inactivating enzymes through N-acetylation

O-nucleotidylation

O-phosphorylation.

Reduced membrane permeability

- LPS structure → aminoglycosides, polymyxins and cationic peptides;
- Porines → chloramphenicol, trimethoprim, ciprofloxacin or β-lactams;
- bacterial capsule.

No standard eradication protocol

	Medication	Dosing	Frequency	Route
2 weeks	Tobramycin	10 mg/kg	daily	IV
	Ceftazidime	50 mg/kg	3 times daily	IV
	Temocillin	2 g	Twice daily	IV
3 months	Tobramycin	100-300 mg	Twice daily	Nebulised

Kitt et al. (2016) BMC Pharmacology and Toxicology 17:14

Table 1 Burkholderia Eradication Protocol

	Medication	Dosing	Frequency	Route	Bacteria Targeted
Induction Period (21 Days):	Tobramycin	6 mg/kg (per kinetics)	Daily	IV	PsA, Bcc
	Ceftazidime	2 g	Every 8 h	IV	PsA, Bcc
	Trimethoprim/Sulfamethoxazole	800/160 mg	Twice Daily	Oral	Всс
	Tobramycin inhaled ^a	300 mg	Twice Daily	Nebulized	PsA, Bcc
	Azithromycin	250 mg	Daily	Oral	Anti-inflammatory
Consolidation Period (2 months):	Trimethoprim/Sulfamethoxazole	800/160 mg	Twice daily	Oral	Всс
	Tobramycin inhaled ^a	300 mg	Twice daily	Nebulized	PsA, Bcc
	Azithromycin	250 mg	Daily	Oral	Anti-inflammatory

^aAlternative- TIP (tobramycin dry powder for inhalation, 4 caps (28 mg/cap)) every 12 h

Garcia et al. (2018), BMC Pulmonary Medicine 18:35

Only few antibiotics are effective against B. cenocepacia

A Current approaches

Antibiotics in clinical use

Aztreonam, monobactam (Bosso et al., 1991; Tullis et al., 2012)

Doripenem, carbapenem (Zobell et al., 2014)

Tobramycin, aminoglycoside (Waters *et al.*, 2017)

New compounds in clinical trials

OligoG (inhaled alginate oligosaccharide), perturbation of biofilm formation (www.clinicaltrialsregister.eu/ctr-search/trial/2014-002125-35/DE)







Burkholderia cenocepacia Infections in Cystic Fibrosis Patients: Drug Resistance and Therapeutic Approaches

Viola C. Scoffone¹, Laurent R. Chiarelli¹, Gabriele Trespidi¹, Massimo Mentasti^{2,3}, Giovanna Riccardi¹ and Silvia Buroni¹*

B Alternative approaches

Molecules in clinical use for other diseases Immunosuppressors and corticosteroids, contribution to antibiotic response (Gilchrist et al., 2012; Weidmann et al., 2008)

IFN-γ, increasing of autophagosome formation (Assani *et al.*, 2014)

Thiosulfoacid S-esters, antimicrobial activity (Lubenets *et al.*, 2017)

Cysteamine, mucolytic and antimicrobial properties; restoration of CFTR function (Vu et al., 2017)

Imidazoles, increasing of *in vitro* biofilm eradication (Van den Driessche *et al.* 2017)

Natural products

Plant nanoparticles, antimicrobial activity (Amato et al. 2016)

Fish oils, antimicrobial activity (Mil-Homens *et al.*, 2016; Olveira *et al.*, 2010)

Glycopolymers, antimicrobial properties and enhanced susceptibility of resistant strains (Narayanaswamy et al. 2017)

Other alternatives

Phage therapy and Phage-Antibiotic

Synergy, antimicrobial activity (Semler *et al.*, 2014; Kamal and Dennis, 2015)

Antimicrobial peptides, prevention of biofilm formation (de La Fuente-Nunez *et al.*, 2014)

Quorum sensing inhibitors, decreasing of bacterial virulence (Brackman *et al.* 2012; Scoffone *et al.*, 2016)

RND efflux pumps in B. cenocepacia **55% MexB** BCAS0591-92-93 63% MexB BCAS0764-65-66 71% **MexY** BCAL1674-75-76 **64%** MexB BCAL2820-21-22 50% **MexD BCAL1778 56% MexD** BCAM0925-26-27 **56% MexF** BCAM1945-46-47 **64% MexF** CeoB (10) < BCAM2549-50-51 70% CzrA BCAM0710-11-12 BCAM0433-34-35 Regulatory gene BCAL1811-12-13 Membrane Fusion Protein **Resistance Nodulation Division** BCAS0582-83-84 Outer Membrane Protein BCAL2134-35-36 15 *llpE* gene BCAM1419-20-21 Promoter region

RND Efflux Pumps in *B. cenocepacia*

BMC Microbiology



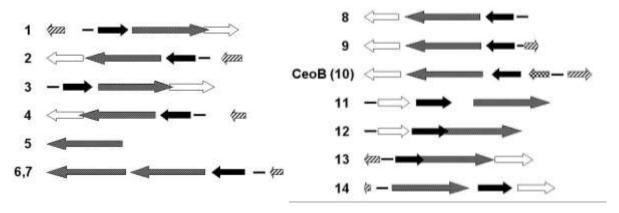
Research article

Open Access

→ RND operons identified in J2315

Efflux pump genes of the resistance-nodulation-division family in Burkholderia cenocepacia genome

Paola Guglierame^{†1}, Maria Rosalia Pasca^{†1}, Edda De Rossi¹, Silvia Buroni¹, Patrizio Arrigo², Giulia Manina¹ and Giovanna Riccardi^{*1}



BMC Microbiology



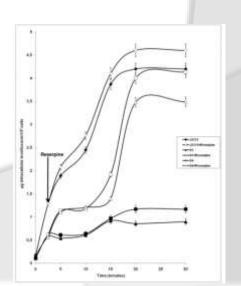
Research article

Open Access

Assessment of three Resistance-Nodulation-Cell Division drug efflux transporters of Burkholderia cenocepacia in intrinsic antibiotic resistance

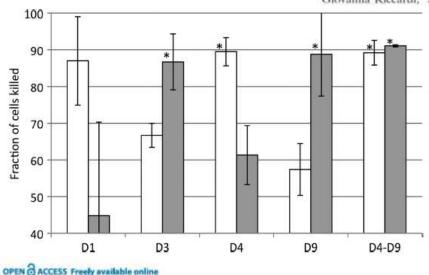
Silvia Buroni^{*1}, Maria R Pasca¹, Ronald S Flannagan², Silvia Bazzini¹, Anna Milano¹, Iris Bertani³, Vittorio Venturi³, Miguel A Valvano² and Giovanna Riccardi¹

- → RND-4 plays a crucial role in the intrinsic resistance of *B. cenocepacia*
 - to Azt, Caf, EtBr, Genta, Tobra, Nal Ac, Cip, Levo, Norflo and Sparflo
- → Transport of **QS** signal molecules



Molecular Mechanisms of Chlorhexidine Tolerance in Burkholderia cenocepacia Biofilms[▽]†

Tom Coenye, 14 Heleen Van Acker, 1 Elke Peeters, 1 Andrea Sass, 2 Silvia Buroni, 3 Giovanna Riccardi, 3 and Eshwar Mahenthiralingam²



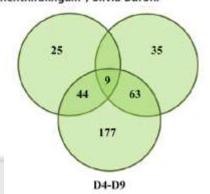
RND-3 and RND-9 → chlorhexidine tolerance for biofilm cells RND-4 → chlorhexidine tolerance for planktonic cells

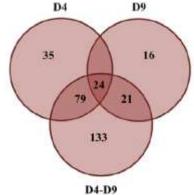
PLOS one

Deciphering the Role of RND Efflux Transporters in Burkholderia cenocepacia

Silvia Bazzini¹⁹, Claudia Udine¹⁹, Andrea Sass²⁹, Maria Rosalia Pasca¹, Francesca Longo³, Giovanni Emiliani⁴, Marco Fondi⁵, Elena Perrin⁵, Francesca Decorosi⁶, Carlo Viti⁶, Luciana Giovannetti⁶, Livia Leoni³, Renato Fani⁵, Giovanna Riccardi¹, Eshwar Mahenthiralingam², Silvia Buroni¹*

transcriptome analysis of ∆RND-4 and ∆RND-9 → wider role than drug resistance





Motility
Chemotaxis
Biofilm production



Differential Roles of RND Efflux Pumps in Antimicrobial Drug Resistance of Sessile and Planktonic *Burkholderia cenocepacia* Cells

Silvia Buroni,^a Nele Matthijs,^b Francesca Spadaro,^a Heleen Van Acker,^b Viola C. Scoffone,^a Maria Rosalia Pasca,^a Giovanna Riccardi,^a Tom Coenye^b

16 RND knock-out mutants

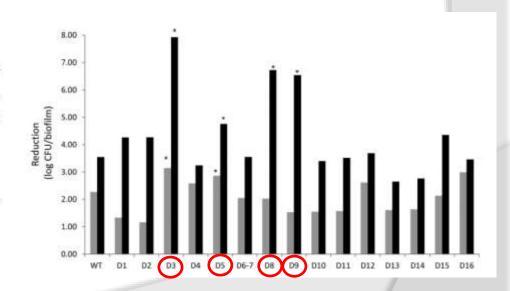
RND-3 and RND-4 involved in resistance of planktonic cells

RND-3, RND-5, RND-8 and RND-9 involved in resistance of sessile cells

TABLE 2 MICs of the various antibiotics for some of the B. cenocepacia strains tested

	MIC (μg/ml) for ^a ;					
Strain	CIP	ТОВ	MIN	MER	CHL	
J2315 (WT)	8	256	16	64	32	
D3	2	2	b	32	16	
D4	2	128	4	64	8	
D16	8	256	4	32	16	

[&]quot;CIP, ciprofloxacin; TOB, tobramycin; MIN, minocycline; MER, meropenem; CHL, chloramphenicol.



b—, no clear minocycline breakpoint was observed for this mutant.

Strategies to fight antibiotic resistance

Essential Proteins

➤ Antibacterial compounds for *B. cenocepacia*: C109

Virulence Factors

➤ 4 antivirulence compounds

Diketopiperazines

Vaccines

From drug to target:

screening of new compounds to find new drugs effective against *Burkholderia*



8000 compounds from AstraZeneca company with unknown chemical structure and mechanism of action. **Resistant**



150 antitubercular compounds from Prof. M. Baltas (CNRs, Toulose). **Resistant**



100 new compounds from Dr. Vadim Makarov

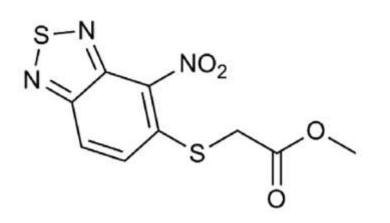
One new compound for *B. cenocepacia* $MIC = 8 \mu g/ml$

10126109

One new compound for *B. cenocepacia* MIC = 16 μg/ml

11026103

C109: a benzothiadiazol derivative active against B. cenocepacia



It is a **bactericidal** compound

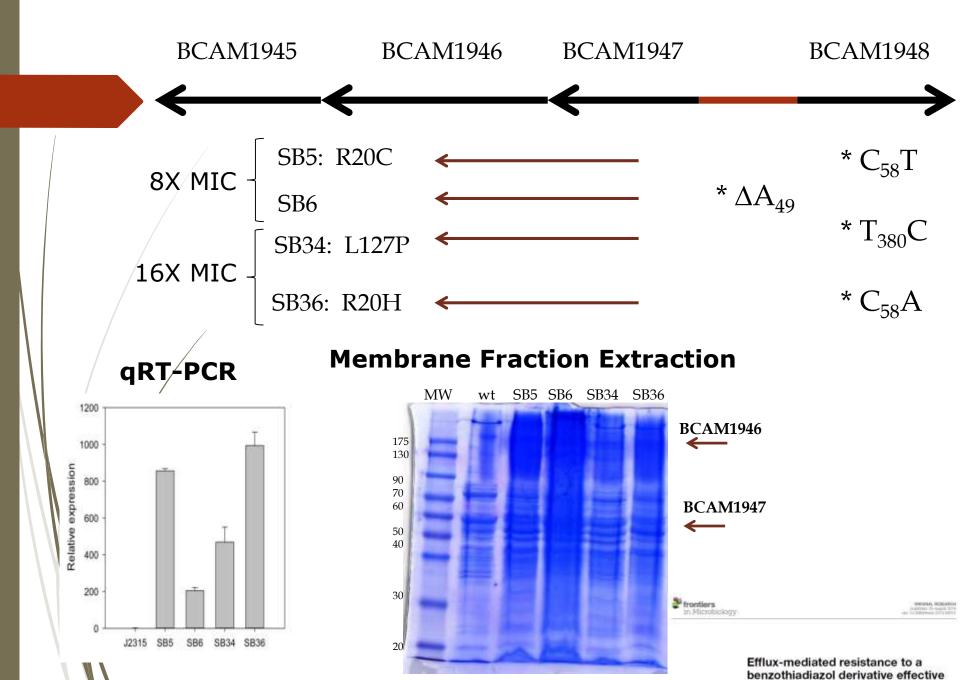
MIC = $8 \mu g/ml$

 TC_{50} on human CF cells = 75 μ M

C109 is effective against the *Burkholderia cepacia* complex strains, more than 50 clinical isolates ...

...other Gram negatives and S. aureus (including MRSA strains)

Strain	MIC 10126109 (μg/ml)
E. coli ATCC 25922	8
A. baumannii ATCC 19606	16
K. pneumoniae ATCC 13883	64
P. aeruginosaPAO1	256
S. aureus ATCC 25923	4



against Burkholderia cenocepacia

Wola C. Scoffonei, Olga Pysicovai, Vadin Malazrovi, Paolo Indennia, Marco Famagali'
Marco Fondii, Renalo Fanir, Edda Da Rosse, Gireama Riccardi and Edvin Burasi'





Competitive Fitness of Essential Gene Knockdowns Reveals a Broad-Spectrum Antibacterial Inhibitor of the Cell Division Protein FtsZ

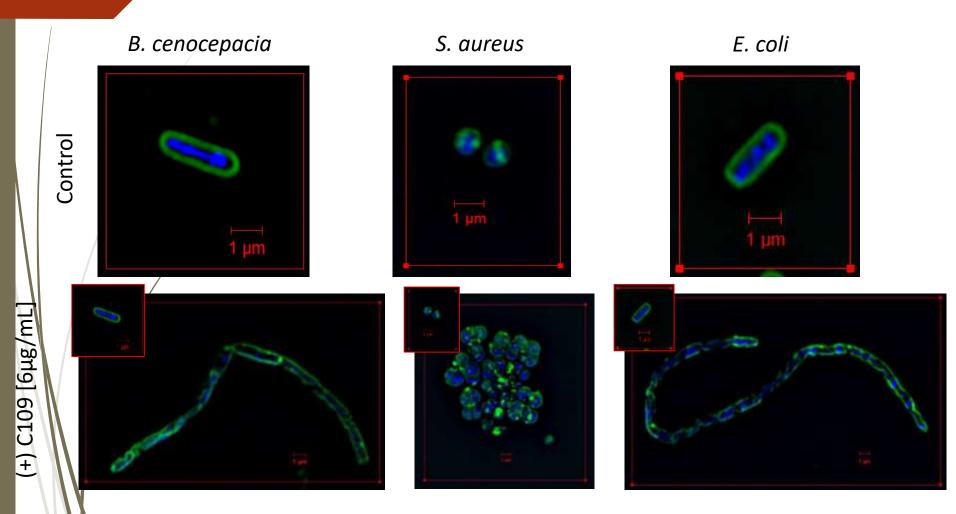
Andrew M. Hogan, *Viola C. Scoffone, *Vadim Makarov, *April S. Gislason, *Haben Tesfu, *Maria S. Stietz, *Ann Karen C. Brassinga, *Michael Domaratzki, *Xuan Li, *Alberto Azzalin, *India Buroni, *Olga Riabova, *Natalia Monakhova, *Olga Riabova, *O

High-density transposon mutant library in B. cenocepacia K56-2 by delivering a transposon element containing an outward tightly regulated rhamnose-inducible promoter (PrhaB)



of PrhaB throughout the bacterial chromosome followed by screening for absence or growth without rhamnose

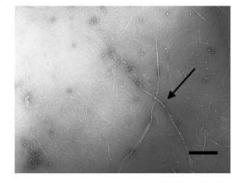
Phenotypic effect of C109 treatment: 3D super resolution microscopy



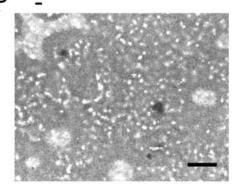
Identification of the cellular target

FtsZ polymers visualized by TEM

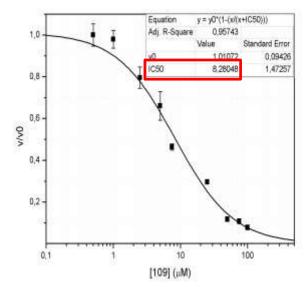
C109 -GTP + GDP -



C109 100 μM GTP + GDP -

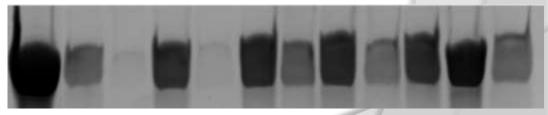


FtsZ GTPase spectrophotometric assay



FtsZ polymerization assay

GTP GDP 100μM 109 50μM 109 25μM 109 10μM 109



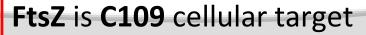
Antimicrobial Agents
occurred
microbiaccor
and Chemotherapy*

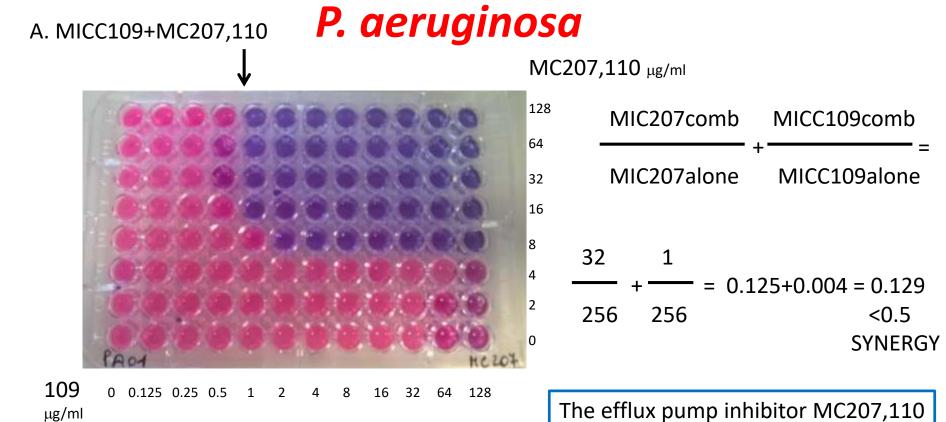
MECHANISMS OF ACTION: PHYSIOLOGICAL EFFECTS

uns Reveals a

Competitive Fitness of Essential Gene Knockdowns Reveals a Broad-Spectrum Antibacterial Inhibitor of the Cell Division Protein FtsZ

Andrew M. Hogan,* Viola C. Scoffone,^b Vadim Makarov,* April S. Gislason,* Haben Tesfu,* Maria S. Stietz,* Ann Karen C. Brassingar,* Michael Domartzkij,* Xuan L.* Alberto Azzalin,* Marco Biggiogera,* Olga Riabova,* Natalia Monakhova,* © Laurent R. Chlarelli,* Colvanna Riczardi,* Silvia Butoni,* © Silvia T. Cardona**





enhances C109 activity against
P. aeruginosa

B. FtsZ purification and enzymatic assay

FtsZ_{PAO} has a 53% as identity with FtsZ_{J2315} \Longrightarrow C109 is effective against the FtsZ_{PAO} purified protein IC50 = 5 μ M

FtsZ is C109 cellular target
In *P. aeruginosa* C109 is extruded by efflux pumps





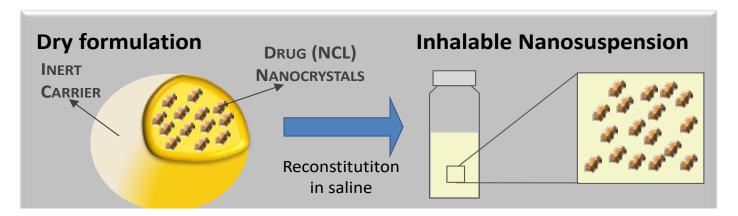




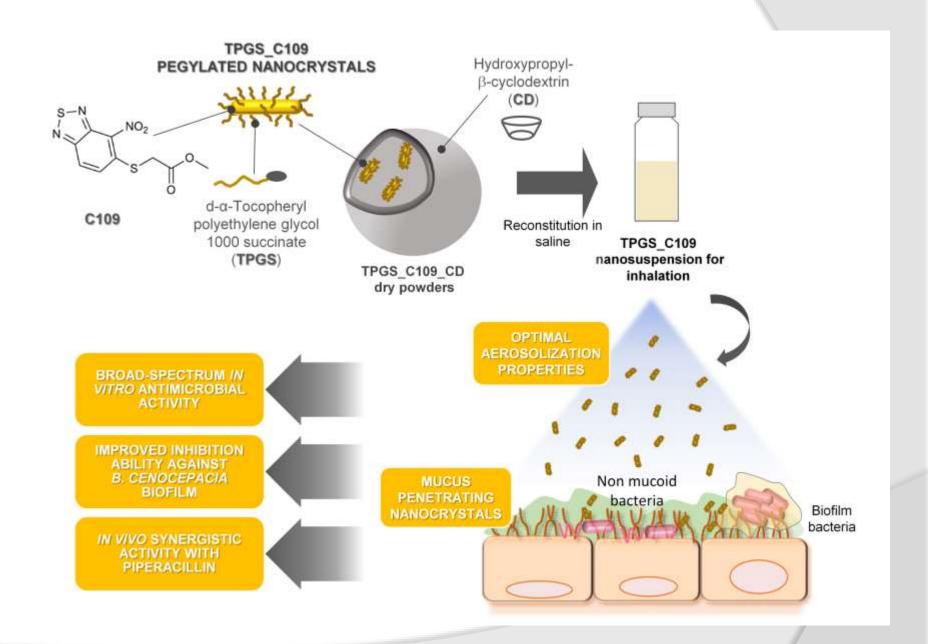




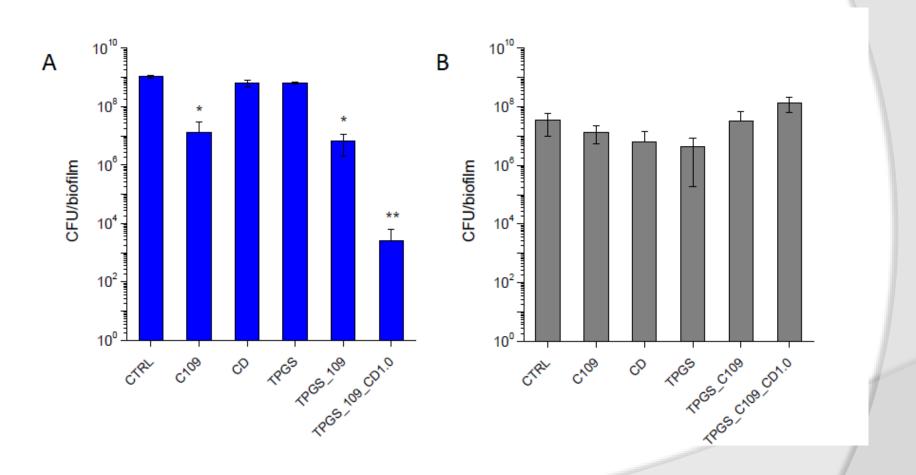




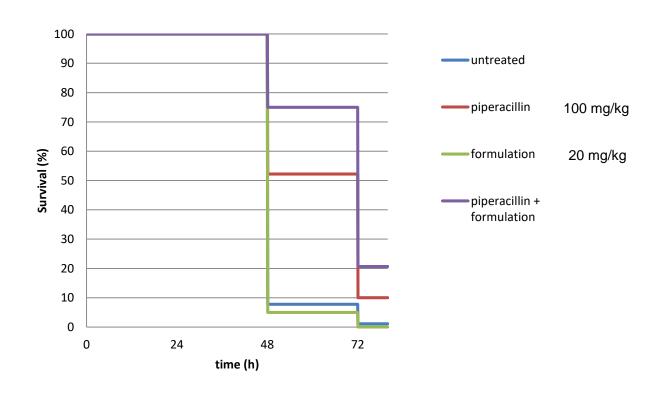
- Biodistribution
- > Efficacy in mouse model of *B. cenocepacia* infection



Inhibition and eradication ability of C109 and of formulations against *B. cenocepacia* J2315 biofilm



Galleria mellonella infection model



Strategies to fight antibiotic resistance

Essential Proteins

➤ Antibacterial compounds for *B. cenocepacia*: C109

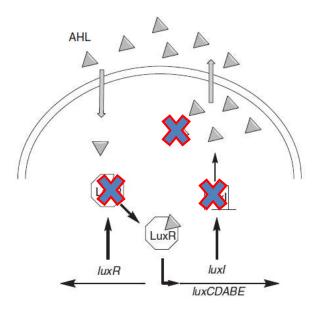
Virulence Factors

➤ 4 antivirulence compounds

Diketopiperazines

Vaccines

Antimicrobial Compounds targeting Virulence Factors



Jiang and Li, 2013

Quorum Sensing Synthases

Cepl (BCAM1870)
Ccil (BCAM0239a)
DfsA (BCAM0581)
Ham cluster

- 1. Bacteria unable to produce virulence factors
- No direct bacteria killing = development of resistance less probable
- Anti-virulence drugs could potentially be used in combination with established or novel antimicrobials to improve the current therapy

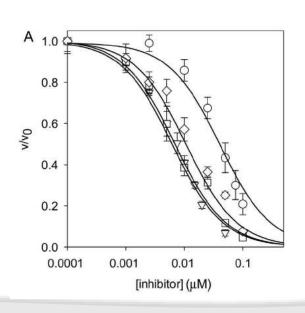
Targeting B. cenocepacia Cepl synthase

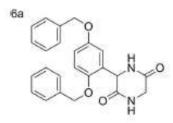
SCIENTIFIC REPORTS

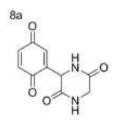
OPEN Discovery of new diketopiperazines inhibiting Burkholderia cenocepacia quorum sensing in vitro and in vivo

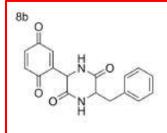
Received: 01 June 2016 Accepted: 08 August 2016 Published: 01 September 2016

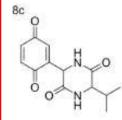
Viola C. Scoffone^{1,*}, Laurent R. Chiarelli^{1,*}, Vadim Makarov^{2,*}, Gilles Brackman³, Aygun Israyilova^{1,4}, Alberto Azzalin^{5,6}, Federico Forneris¹, Olga Riabova², Svetlana Savina², Tom Coenye3, Giovanna Riccardi1 & Silvia Buroni1









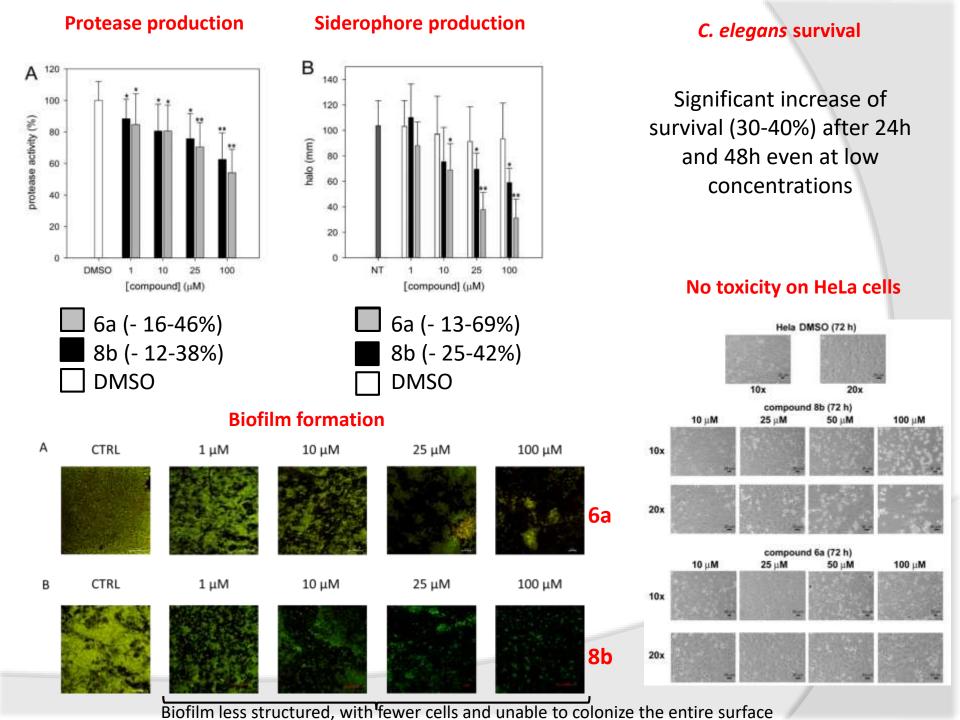


30 μΜ **6a** ○

 $10 \mu M$ 8a ♦

8b ∇ 5 μ M

6 μΜ 8c ⊔



Strategies to fight antibiotic resistance

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➤ Antibacterial compounds for *B. cenocepacia*: C109

Virulence Factors

➤ 4 antivirulence compounds
Diketopiperazines

Vaccines

Vaccines for B. cenocepacia

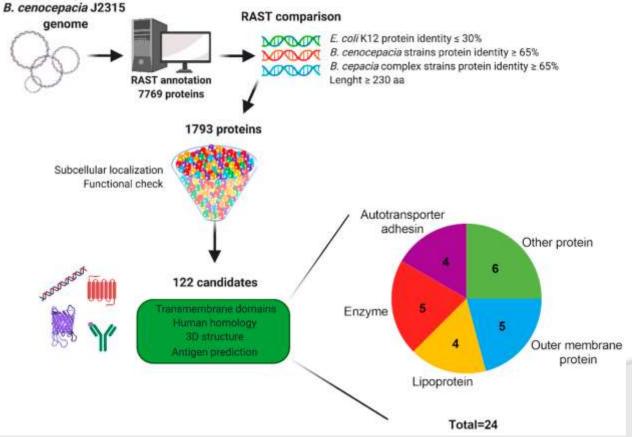




Article

Identification by Reverse Vaccinology of Three Virulence Factors in *Burkholderia cenocepacia* That May Represent Ideal Vaccine Antigens

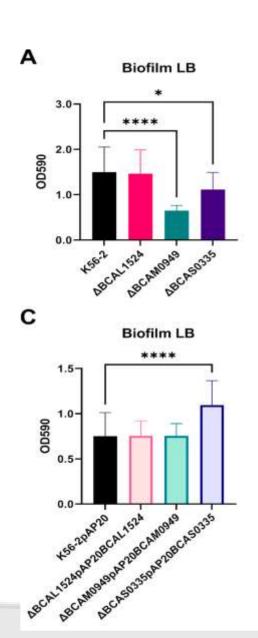
Samuele Irudal ¹©, Viola Camilla Scoffone ¹, Gabriele Trespidi ¹©, Giulia Barbieri ¹©, Maura D'Amato ²©, Simona Viglio ²©, Mariagrazia Pizza ³©, Maria Scarselli ⁴, Giovanna Riccardi ¹ and Silvia Buroni ¹, ⁶©

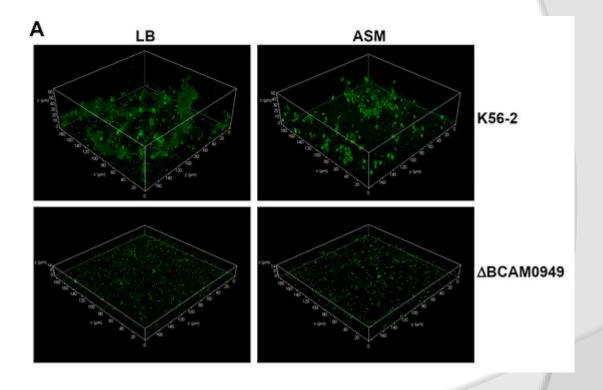


Collage-like protein **BCAL1524** Lipase LipA **BCAM0949** Autotransporter adhesin **BCAS0335**

selected for further analysis

Biofilm formation

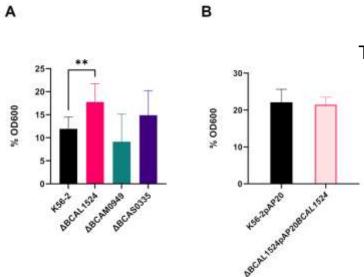




the lipase BCAM0949 and the autotransporter adhesin protein BCAS0335 contribute to biofilm formation

Autoaggregation

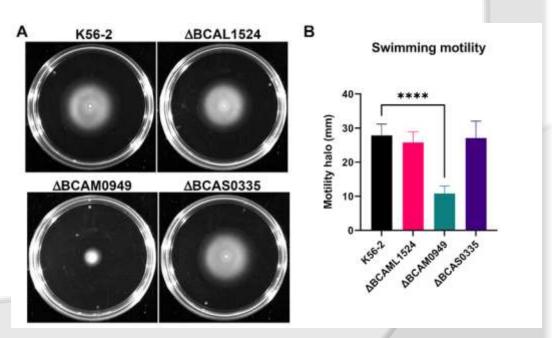
Involved in bacterial colonization and persistence in the host



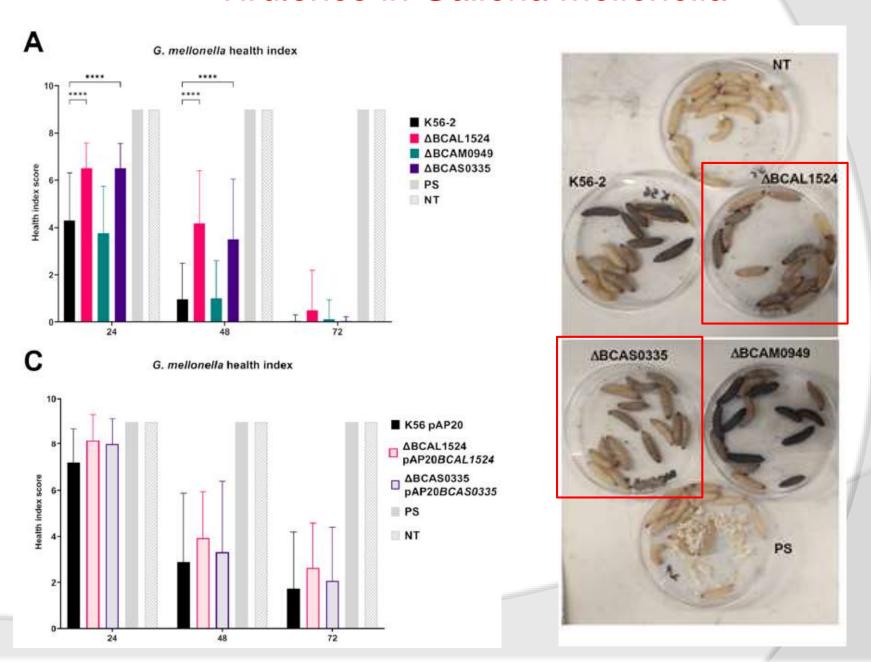
The Collage-like protein BCAL1524 is involved in autoaggregation

Swimming Motility

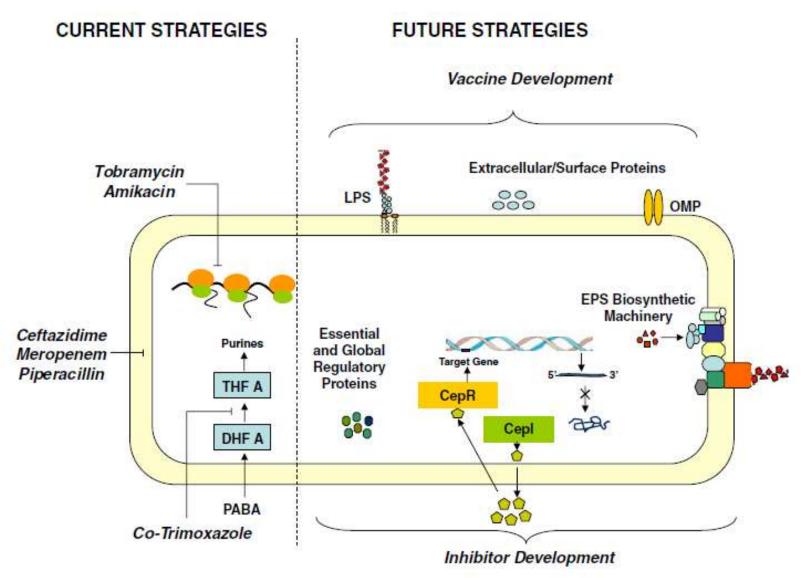
The Lipase LipA BCAM0949 is involved in swimming motility



Virulence in Galleria mellonella



Strategies to fight *B. cenocepacia*





Using a Virtual Screening approach to find new drugs against *Pseudomonas aeruginosa* and Staphylococcus aureus**

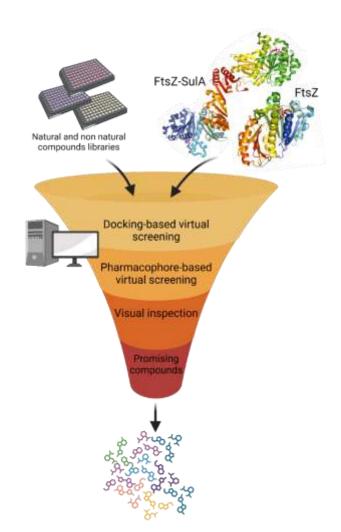


Silvia Buroni and Antonio Coluccia



Background and rationale

- Virtual Screening (VS): fast and cheap method for the selection of small molecules effective in the inhibition of the selected enzymes.
- Filter rules with descriptors to evaluate drug-like and toxicological properties of the compounds → quick identification of new hits to be tested in clinical trials



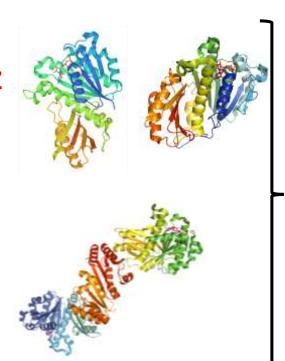
Hypothesis and objectives

VS targeting **cell division** proteins:

S. aureus and P. aeruginosa FtsZ

First protein involved in cell division Assembly of "divisome"

- The complex FtsZ-SulA
- Small protein induced as part of the SOS response to DNA damage
- It binds to FtsZ and blocks cell division until the DNA is repaired
- It regulates FtsZ activity



Essential for bacterial survival

Widely conserved

No antibiotic in clinical use has these two proteins as targets

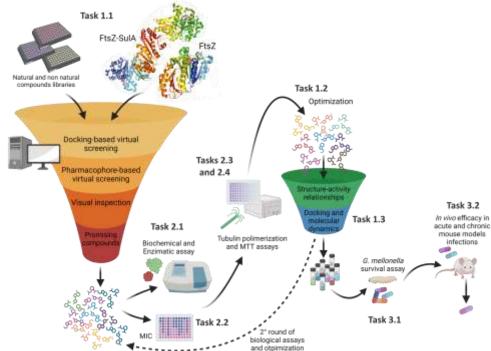
Essential methods

- WP1: chemistry
- Task 1.1 Identification of compounds targeting P. aeruginosa FtsZ or FtsZ-SulA by VS.
- Task 1.2 Compound optimization.
- Task 1.3 Hit to lead optimization.
- WP2: compound characterization
- Task 2.1 Enzymatic assays.
- Task 2.2 MIC evaluation.
- Task 2.3 Evaluation of the activity of the compounds against human tubulin.
- Task 2.4 Evaluation of the toxicity of the compounds by MTT.
- WP3: in vivo assays
- Task 3.1 In vivo assays in Galleria mellonella.
- Task 3.2 *In vivo* assays in mouse models of infection.



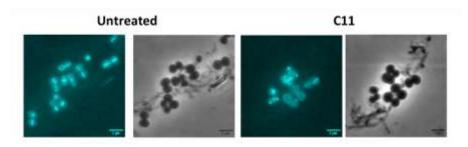
Dr. Alessandra Bragonzi





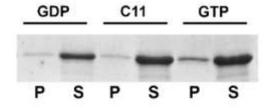
Preliminary results

Compound	% of GTPase residual activity	IC ₅₀	MIC
C109	23 %	1.5 μΜ	2-4 μg/ml
C1	78 %		> 256 μg/ml
C2	35 %		> 256 μg/ml
СЗ	16 %	14.6 μΜ	> 256 μg/ml
C4	71 %		> 256 μg/ml
C5	95 %		> 256 μg/ml
C6	92 %		> 256 μg/ml
C7	100 %		> 256 μg/ml
C8	12 %	8.4 μΜ	> 256 µg/ml
C9	34 %		> 256 μg/ml
C10	87 %		> 256 μg/ml
C11	82 %		2 μg/ml
C12	17 %	60 µM	> 256 μg/ml



In untreated cells FtsZ localizes in the middle of the cell and forms the Z-ring.

In the treated sample, cells are larger and FtsZ is mislocalized.



C11 (100 µM) blocks FtsZ polymerization

Toxicity evaluation

At 25 μM C11: - 20% human tubulin polymerization.
90% of HeLa cells survived the treatment.

Conclusions

- Our final goal is to identify new therapeutic solutions against the two MDR pathogens that mostly affect CF patients, with a new mechanism of action with respect to currently used drugs.
- In the future the same approach could be applied to **other proteins** and bacteria to find new therapeutic solutions for airway pathogens, thus promoting the introduction of innovative treatments for CF patients.

Characterization of the dispirotripiperazine derivative PDSTP as antibiotic adjuvant and antivirulence compound against *Pseudomonas aeruginosa*

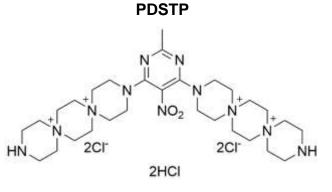
Front. Microbiol. 15:1357708. doi: 10.3389/fmicb.2024.1357708

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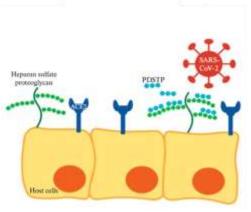
Dispirotripiperazines: broad spectrum antiviral compounds

PDSTP is active against:

- ✓ Herpes simplex virus type-1 and 2
- ✓ Human cytomegalovirus
- ✓ Human immunodeficiency virus
- ✓ Human papilloma virus
- ✓ SARS-CoV-2







Dr. Vadim Makarov

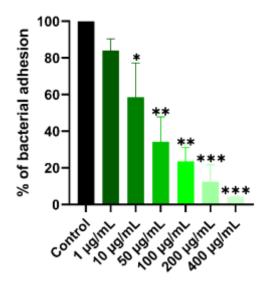
Makarov and Popov, 2022

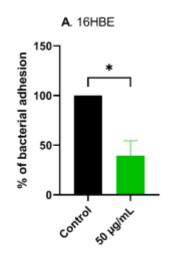
PDSTP impairs P. aeruginosa adhesion

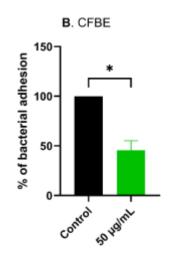
B

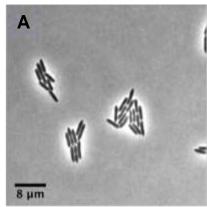
PA01 adhesion on A549 monolayers

PA01 adhesion on 16HBE and CFBE monolayers









3 - 8 μm

No treatment

PDSTP 50 μg/mL

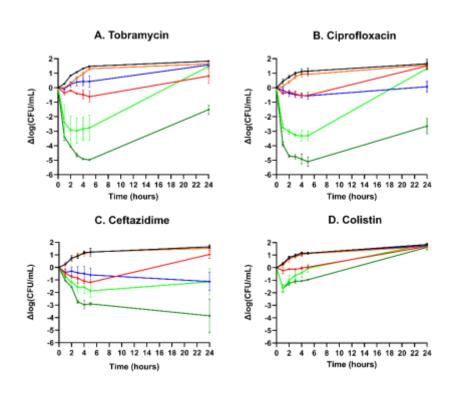
PDSTP enhances efficacy of antibiotics used in clinics

PDSTP-antibiotic combinations tested on PA01 and 9 drug resistant CF clinical isolates

	Antibiotic	MIC reduction
	Tobramycin	0x to 2x
6/7 resistant strains below the CLSI suscept	Ciprofloxacin	2x to 16x
	Ceftazidime	4x to 128x
4/4 resistant strains below the CLSI suscep	lity breakpoint Meropenem	8x to 32x
	Colistin	16x to 32x

MIC of 2 resistant strains decreased from 64-128 μg/mL to 4 μg/mL

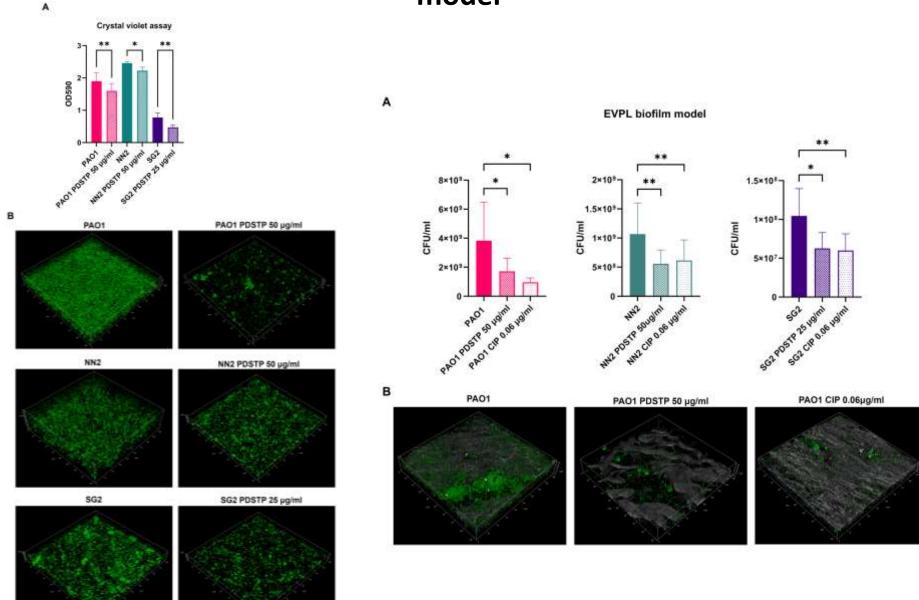
PDSTP-antibiotic combinations tested on PA01 and 9 drug resistant CF clinical isolates



Comparable efficacy of the combinations was observed by time-killing assays on **7/9 clinical isolates**

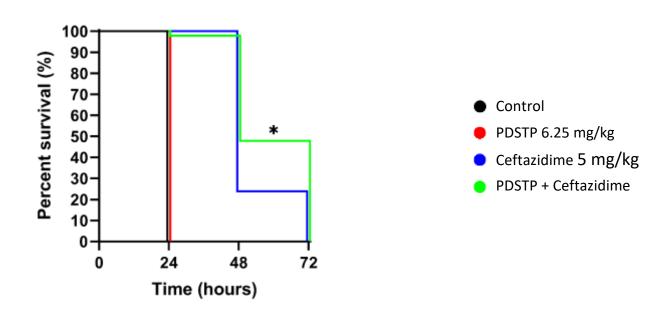
- Control
- PDSTP (50 μg/mL)
- PDSTP (200 μg/mL)
- Antibiotic (1/2 MIC)
- PDSTP (50 μg/mL) + Antibiotic
- PDSTP (200 µg/mL) + Antibiotic

PDSTP inhibits biofilm formation in vitro and in an ex vivo pig lung model



PDSTP - Ceftazidime combination is effective in vivo

Galleria mellonella infection model



Conclusions

Vaccines

Adjuvants





Strategies to fight antibiotic resistance





Essential proteins

Virulence factors

FtsZ

Quorum Sensing



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