The 1st International Electronic Conference on Antibiotics—The Equal Power of Antibiotics And Antimicrobial Resistance Section: Antimicrobial Resistance Mechanisms and Intrinsic Microbial Factors Contributing to Resistance

Calcium Regulates Resistance of *Pseudomonas aeruginosa* to Polymyxin B

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INTRODUCTION



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Pseudomonas aeruginosa

- Ubiquitous in nature
- A major cause of nosocomial infections (9 -10% of all hospital infections)
- Causes high morbidity and mortality in cystic fibrosis (CF) patients and

immunocompromised individuals

Multi-drug resistant *P. aeruginosa* is a growing global concern

Antibiotic resistance rate over time in pneumonia due to Pseudomonas aeruginosa



Priority 1: CRITICAL[#]

Acinetobacter baumannii, carbapenem-resistant

Pseudomonas aeruginosa, carbapenem-resistant

Enterobacteriaceae*, carbapenem-resistant, 3rd generation cephalosporin-resistant

WHO priority pathogen list for development of new antibiotics

Last hope antibiotics to the rescue

- Nature's peptides
- Small, cationic

Polymyxin B (Pol B) Polymyxin E (Colistin)

Binds to LPS promoting
membrane permeabilization,
results in cell lysis



P. aeruginosa has multifactorial resistance mechanisms

- Restricted uptake and efflux
- Drug inactivation
- Changes in targets



Patik and Kolar, 2019; Czech Repub BIOMED PAP

Ca enhances Polymyxin B resistance in P. aeruginosa



What are the mechanisms of Ca induced resistance to Polymyxin B in *P. aeruginosa?*



Test the involvement of known mechanisms of resistance to Pol B in *P. aeruginosa*

Identify Ca dependent mechanisms of Pol B resistance

Characterize the membrane changes in response to Ca

Known Pol B resistance mechanisms do not play a role in Ca- induced resistance



Known Pol B resistance mechanisms do not play a role in Ca- induced resistance



- Wzz: modulating O-antigen chain length
- RmID: dTDP-4
 - dehydrorhamnose reductase
- CbrB: two-component response regulator

Three novel hypothetical proteins identified to be involved in Ca regulated polymyxin-B resistance



MIC of transposon mutants and in-trans complemented genes compared to WT

MIC of clean deletion mutants compared to WT

Increased transcript abundance of PA2803 and PA3237 in response to Ca



Transcript abundance in WT and *calC*:Tn5 background

Bioinformatics prediction of PA2803, PA3237 and PA5317



Purification of PA2803



Ni-NTA affinity purification L Lys FT W1 W2 1 3 5 7 9 L 11 13 15 17 19 21 23 25 L y y y y y 1 10

- A putative phosphonoacetaldehyde hydrolase- Phosphonatase
- Predicted Ca²⁺/ Mg²⁺ binding
- Cytoplasmic (PSORTb 3.0)

His-tag cleavage and affinity purification



Characterization of PA2803

Determination of oligomeric state of PA2803

Ca binding assay using ANS (8-anilino-1-naphthalene sulfonate)



Phosphatase activity of PA2803

pNPP hydrolysis of Shrimp alkaline phosphatase



pNPP hydrolysis of PA2803





98.04 nM/min/µg of protein

Growth at elevated Ca levels increases the outer membrane permeability of *P. aeruginosa*





- The permeability of the outer membrane increases in a Pol B concentration dependent manner
- No significant change in permeability in mutants tested compared to WT

Short-term exposure to Ca does increase outer membrane permeability of PAO1



Growth at elevated Ca levels increases the inner membrane permeability of *P. aeruginosa*





 No significant change in permeability in mutants tested compared to WT

Inner membrane permeability of PAO1

Differences in LPS and membrane-associated proteins at elevated Ca



Summary



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