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Viable but non-culturable state: a strategy for Staphylococcus aureus survivable in dual-species biofilms with Pseudomonas aeruginosa ?

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3

Polymicrobial Biofilm Infections

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Cystic Fibrosis related-infections

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Cystic Fibrosis Annual Data Report 2017



Cystic Fibrosis related-infections

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Cystic Fibrosis Annual Data Report 2017

Staphylococcus aureus

- Prevalent among people with and without CF
 - \sim 20 % of strains are multidrug-resistant



Pseudomonas aeruginosa

- A leading cause of airway infection
- Associated with a decline in lung function
- ~18 % of strains are multidrug-resistant





AIM:

To investigate the community dynamics of *Pseudomonas aeruginosa* and *Staphylococcus*

aureus, two common co-infecting pathogens in cystic fibrosis infections, growing as dual-

species biofilms.





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- Biofilm structure
- Microbial composition
- Gene expression profile





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Results



Biofilm structure: SEM analysis

Magnification 2750 x











Biofilm structure: SEM analysis



24-h-old







Biofilm structure: SEM analysis

Magnification 2750 x





Mature biofilms producing a thickness of co-aggregated cells surrounded by extracellular biofilm matrix







Effect of secreted compounds produced by P. aeruginosa





Only 1 log of cell reduction (for
 10⁵ and 10⁷ CFU/mL) was observed





Effect of secreted compounds produced by P. aeruginosa







Results



Biofilm bulk fluid quantification







Dual-species biofilms

Single-species biofilm



Results



Biofilm bulk fluid quantification







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()iversidade do Minho

Quantitative assessment of individual populations within dual-species biofilms

Results





23



Results



Quantitative assessment of individual populations within dual-species biofilms





For 24-h biofilms similar counts
 were detected by both methods





Quantitative assessment of individual populations within dual-species biofilms

Results









Quantitative assessment of individual populations within dual-species biofilms



(A) P. aeruginosa 362668 mucoid and S. aureus ATCC 25923 (C) P. aeruginosa 362668 non-mucoid and S. aureus ATCC 25923 (B) P. aeruginosa 362668 mucoid and S. aureus 352845 (D) P. aeruginosa PA14 and S. aureus 352845.



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Quantitative assessment of individual populations within dual-species biofilms



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Virulence expression in dual-species biofilms

P. aeruginosa virulence-related genes:

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Gene	Function
pqsE	НОИО
rhlR	Virulence Regulator (Quorum sensing)
pvdE	Pyoverdine
toxA	Exotoxin A
lasl	Virulence Regulator (Quorum sensing)
algD	Alginate



48 h



Virulence expression in dual-species biofilms



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Virulence expression in dual-species biofilms

S. aureus virulence-related genes:

Gene	Function
sodA	Stress Response
sarA	Virulence Regulator (Quorum sensing)
agrB	Virulence Regulator (Quorum sensing)
hld	Virulence Regulator (Quorum sensing)
icaA	Biofilm formation (PNAG production)
hla	Alfa-hemolysin
uspA	Stress Response



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- ✓ Time-dependent interaction between *P. aeruginosa-S. aureus* in dual-species biofilms.
- ✓ The dual-species consortia dominated by *P. aeruginosa*
- ✓ The presence of S. aureus in high numbers in dual-species biofilms with P. aeruginosa in a VBNC state.

Conclusions





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Conclusions





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Overall, our results underline the importance of select appropriate methodologies to elucidate the

microbial interactions occurring within the dual-species biofilm consortia.





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Nuno Cerca (CEB, University of Minho, Portugal)

MOP Team



NC Team



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Thank you!

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