OF NEW JERSEY

INTRODUCTION

Bacteria rarely exist as single plankotic forms. Several microbial species exist as polymicrobial biofilm communities co-operating the survival of each other. Biofilms are a hallmark feature of various pathogenic and opportunistic pathogenic bacteria that allow them to communicate with each other both *intra-species* and *inter-species* via quorum sensing. We are interested in exploring the quorum sensing phenotypes of the opportunistic bacteria *Pseudomonas aeruginosa* in a polymicrobial setting. *P. aeruginosa* is an opportunistic bacterium that often results in serious infections in health-care settings, especially in immunocompromised patients. This nosocomial opportunistic pathogen produces many virulence factors which are under the regulation of the quorum sensing gene circuit. Two of the very important virulence phenotypes that we are interested in are: production of the blue-green pigment *pyocyanin* and ability to form biofilms.

Our previous work suggested that there is a difference in the pigment production when *Pseudomonas* is co-cultured with other gram positive and gram-negative bacteria. As a continuation of this work, we are currently looking at another quorum sensing phenomenon which is the *biofilm* formation of the bacteria in a polymicrobial setting. We are interested in the polymicrobial biofilm co-culture of *Pseudomonas* aeruginosa with other nosocomial bacterial strains Klebsiella pneumoniae (KP), Escherichia coli (EC), Staphylococcus aureus (SA) and Enterococcus faecalis (EF). Our preliminary results indicate that the polymicrobial biofilms are not only more complex and stronger than the single species biofilms, but also show greater resistance to the antibiotic treatment *in-vitro*. Our studies are aimed at a long-term goal of developing anti-biofilm agents targeting polymicrobial communities rather than the plankotic bacteria. Future experimental efforts will be focused on identifying differential expression of the quorum-sensing genes of *Pseudomonas aeruginosa* in a co-culture model. The research will be an important contribution to the evolving field of antimicrobial peptides targeting bacterial biofilms and bacterial quorum sensing.

Polymicrobial Biofilms

Biofilms are complex, thriving microbial communities attached to biotic and abiotic surfaces via extracellular matrix made of polysaccharides, proteins and DNA. Bacteria living in biofilm are physiologically different from their planktonic counterparts in many ways. The planktonic form is usually characterized by mobility, faster growth and better susceptibility to antibiotics. Biofilm growth can be seen as an adaptation to a lownutrient, stressful environment. The Figure below shows various stages of biofilm formation in bacteria.

The first step of biofilm formation is the reversible attachment to a surface through interactions between the bacterial cell wall and the substrate. The contact triggers the release of extracellular matrix components and the colonies grow creating a differential micro-environment to support intra-species and inter-species communication. The biofilm matures and becomes resistant to the antibiotics which are otherwise effective on various planktonic bacteria.

Standard antibiotic susceptibility testing is done on microbes that are grown planktonically whereas the clinical infections are from polymicrobial biofilms. Therefore, it is possible that interactions between microbes could influence the success of antimicrobial treatment. The current study is aimed at understanding the role of *Pseudomonas aeruginosa* quorum sensing phenotypes in modulating the group behaviors in a mixed microbial community. -Polymicrobial



Investigating the *Biofilm* Formation of the Bacteria *Pseudomonas aeruginosa* in a Polymicrobial Setting

Interactions Quorum Sensing -Antibiotic Resistance

<u>1. Optimization of Biofilm Assay</u>

Staphylococcus aureus and Klebsiella incubated at 37 °C for 24-48 hours the wells were patted dry on a paper towel reproducibility



CONCLUSION

Our results indicate that the biofilm formed by the bacteria *Pseudomonas aeruginosa* in a polymicrobial setting is stronger and has greater resistance to antibiotic treatment than the monoculture biofilm. Our previous results suggested that the virulence factor pyocyanin production by *P. aeruginosa* is highly modulated in a polymicrobial setting. In the current study we observe that the similar regulation is also seen for the biofilm formation of the bacteria in the presence of various other species of bacteria (EC, KP, SA and EF which are also nosocomial bacteria). We hypothesize that there could be a strong inter-connection between the two quorum-sensing phenotypes pyocyanin production and biofilm formation in a polymicrobial setting. To address this, we are currently working on mimicking the polymicrobial environment by working with cell-free supernatant of *Pseudomonas* and other bacteria and assessing their ability to inhibit the biofilm formation. Soon we also would work on generating knock-out/mutant strains of *Pseudomonas* for the pyocyanin gene to establish its role on the polymicrobial-*Pseudomonas* biofilm formation. Eventually the goal is to identify novel drug targets that could serve as quorum quenchers or quorum-sensing inhibitors. This approach has a promising potential for the future in minimizing and delaying the problem of antibiotic resistance.

References & Acknowledgements

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Reilly Goodwin, Stacey Hauk and Kokila Kota School of Theoretical and Applied Science, Ramapo College of New Jersey, Mahwah, NJ, 07430

• Attenuation of *Pseudomonas aeruginosa* biofilm formation by Vitexin: A combinatorial study with azithromycin and gentamicin Manash C Das, Yusuf Akhter et al., Scientific Reports, volume 6,

• The biofilm matrix. Felming HC, Wingender J. Nature Reviews Microbiology. 2010;8:623–633. DOI: 10.1038/nrmicro2415 • Microbiome, biofilms, and pneumonia in the ICU Article in Current Opinion in Infectious Diseases · February 2016 DOI: 10.1097/QCO.000000000000255 • A rapid seamless method for gene knockout in Pseudomonas aeruginosa Weiliang Huang and Angela Wilks; Huang and Wilks BMC Microbiology (2017) 17:199 DOI 10.1186/s12866-017-1112-5

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Future Work

 \rightarrow To test the susceptibility/resistance of the polymicrobial biofilms to various antibiotics at their planktonic MIC values

 \rightarrow To treat the mono-culture biofilms *Pseudomonas* and other nosocomial bacteria with the cell-free supernatants of each bacterial strain \rightarrow To generate the *Pseudomonas* mutant or knock-out strains and identify the role of Pyocyanin in the formation of polymicrobial biofilms