Bacteriophages for Biofilm Eradication


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Bacteriophages for Biofilm Eradication

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Introduction

Bacteriophages or phages are a unique type of virus that recognize a specific type of bacteria and then infect, replicate and kill the host via cell lysis. The application of phages and their enzymes for treating bacterial biofilms has recently gained significant interest due to a number of significant advantages compared to traditional antibiotics, including high specificity and efficacy, low immunogenicity and production costs. Proteus mirabilis, Stenotrophomonas maltophilia and Pseudomonas aeruginosa are all clinically relevant microorganisms associated with biofilm formation and thus there is great difficulty treating such infections. With the increasing number of multi-drug resistant strains being reported the time for novel antimicrobial agents is now and thus we propose the use of bacteriophages to treat the said bacterial species’ biofilms.

Aims

• To isolate novel bacteriophages which act against Proteus spp., P. aeruginosa and S. maltophilia
• Obtain high and pure phage titre of isolated phages
• Characterize each phage to determine their novelty
• Assess the ability of phages to degrade bacterial biofilms

Methodology

• Environmental enrichment of phages
• Top agar overlay spot test assay
• Top agar plaque assay
• Phenol:chloroform DNA extraction
• Genomic restriction analysis
• MBEC assay

Results

To date we have isolated 6 phages against 6 different Proteus bacteria (Fig. 1), with 2 phages lysing all strains. Furthermore, we have 9 phages which act against P. aeruginosa. Phage qNFS (Fig. 2) has shown great preliminary results when attempting to degrade PA01 biofilms and resulted in a clinically significant two log reduction of the biofilm. The Proteus phages are yet to be tested against biofilms, but are expected to have a similar outcome to that of qNFS. These results so far show the extreme potential of phages to act as biocontrol agents against biofilms in an era of increasing multi-drug-resistance.

<table>
<thead>
<tr>
<th>Pathogen</th>
<th>Phage #</th>
<th>Isolation</th>
<th>Host range</th>
<th>Sequencing</th>
<th>Anti-biofilm assay (MBEC)</th>
</tr>
</thead>
<tbody>
<tr>
<td>P. aeruginosa</td>
<td>9</td>
<td>Completed</td>
<td>Planned</td>
<td>Completed</td>
<td>In Progress</td>
</tr>
<tr>
<td>Proteus spp.</td>
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<td>In progress</td>
<td>In Progress</td>
<td>TBC</td>
<td>In Progress</td>
</tr>
<tr>
<td>S. maltophilia</td>
<td>0</td>
<td>In progress</td>
<td>Planned</td>
<td>TBC</td>
<td>Planned</td>
</tr>
</tbody>
</table>

Fig. 1. (Left) Spot tests of phages isolated from various animal faecal sources and waste water effluent. (Right) Plaque assay confirming the presence and activity of one of the newly isolated phages (BB2000) against P. mirabilis.

Fig. 2. (Left) Plaque assay results at a 10^-4 dilution for qNFS phage active against P. aeruginosa PA01. (Right) Total lysis of PA01 with 10^-1 diluted qNFS phage demonstrates the potential of this phage as a biofilm biocontrol agent.

Future work

• Isolate S. maltophilia phages from various environmental samples
• Optimize our current phage biofilm assay techniques
• Assess the potential for a phage-cold plasma synergistic technique to facilitate total biofilm eradication
• Assess the effect of a cocktail of phages in a biofilm model to see if they enhance the degradation of the biofilm
• Sequence novel phages using MinION nanopore sequencing and identify genes of potential biofilm degradation enzymes
• Obtain phage enzymes and assess their potential as biofilm biocontrol agents

Anti-biofilm assays (MBEC)

P. aeruginosa biofilms grown via the MBEC device (Fig. 3) were exposed to different phage titres of qNFS phage and incubated for 18-24 h before assessing the ability of the phages to degrade the biofilms. A phage titre of 1x10^7 showed a 100x reduction of the PA01 biofilm (Fig. 4).

Fig. 3. Shows the CV stained pegs of the MBEC device used for the qNFS. As can be seen from left to right an increase in staining and hence biomass as the phage titre decreases in value resulting in less biofilm degradation

Fig. 4. Shows the effect of various phage titres and the consequent 100x reduction of the PA01 biofilm when exposed to qNFS bacteriophage

Planned MinION nanopore sequencing

• We plan to sequence all our phages using MinION nanopore sequencer
• Test runs with Escherichia coli phage λ produced 130k reads in 6 h!