We are primarily engaged in an analysis of the adhesives that enable the stalked Caulobacter bacteria to attach and remain adherent to surfaces in the environment for long periods of time. The specific goals are: 1) Determine the chemical composition and structural arrangement of monosaccharides and other substituents within the holdfasts of selected marine and freshwater Caulobacters. 2) Characterize the types of surfaces to which holdfasts will adhere. 3) Clone and analyze the genes specifying the holdfasts of selected marine and freshwater Caulobacters. 4) Develop the capabilities of selected marine Caulobacters for molecular genetic experimentation. 5) To evaluate the occurrence, stability and behavior of Caulobacters on surfaces and in complex biofilms, in collaboration with other ONR-funded researchers.
Annual Report on grant N00014-89-J-1749
Tuesday, June 6, 1989

Principle Investigator: John Smit

Contractor University of British Columbia

Contact Title: Chemical, structural and genetic analysis of the adhesive holdfast of biofouling Caulobacters

Start Date: 1 March 1989

Research Objectives:
1) Determine the chemical composition and structural arrangement of monosaccharides and other substituents within the holdfasts of selected marine and freshwater Caulobacters. This also includes determining the periodicity and regulation of synthesis during the cell cycle.
2) Further characterize the types of surfaces to which holdfasts will adhere.
3) Clone and analyze the genes specifying the holdfasts of selected marine and freshwater Caulobacters.
4) Continue developing and evaluating the capabilities of selected marine Caulobacters for molecular genetic experimentation.
5) To evaluate the occurrence, stability and behavior of Caulobacters on surfaces and in complex biofilms, in collaboration with other ONR-funded researchers.

Progress and Planned Activities

Chemical analysis of the adhesive holdfast produced by Caulobacters

Based on our studies using lectins to determine the composition of holdfasts, we have selected 2 marine Caulobacters (MCS6 and MCS24) and 2 freshwater Caulobacters (Caulobacter crescentus CB2A and Caulobacter subvibrioides) for in-depth composition analysis. This is also done in concert with decisions regarding the strains selected for gene isolation (MCS6 and Caulobacter crescentus CB2A). As we generate mutants with altered holdfasts, the composition of these will be determined as well.

Our first efforts have been directed to CB2A. We have been concentrating on an isolation procedure based on the observation that holdfasts bind very tightly to colloidal gold particles. Once bound, the holdfast is no longer adhesive and the complex can be isolated readily by CsCl density centrifugation, relying on the high density imparted by the gold binding. However, the method did not work adequately well with wild type cells. We attempted to dissociate holdfast from the main cell body by extensive sonication. This proved to be inadequate; antisera raised against the holdfast-colloidal gold complexes produced sizable activities to the lipopolysaccharide coat of Caulobacters.

We are now using a mutant we isolated after UV mutagenesis that sheds its holdfast into the medium. The holdfast can be collected by attachment to the colloidal gold particles and appears to be free of all cell constituents. As discussed below, we believe we can introduce this shedding phenotype into strains with altered holdfast composition, to enable chemical analysis of the alterations.

With holdfast attached to the colloidal gold we shall be doing a composition chemical analysis, raising specific antisera for the development of an immunoassay and acquire surface-enhanced Raman spectroscopy "fingerprints" for normal and altered holdfast, as a collaboration with both
On the adhesive properties of the holdfast.
As part of studies aimed at discerning what types of surfaces to which the Caulobacters will attach, glass surfaces were covalently modified with a variety of chemical substituents (provided by Dan Rittschoff, Duke University Marine Labs), resulting in surfaces ranging from highly charged to very hydrophobic. From a quantitative static flow attachment assay we learned was that Caulobacters will attach to virtually all surfaces at some frequency, but appear to prefer substrates that are moderately hydrophobic. Freshwater Caulobacters attach better to very hydrophobic surfaces than do marine Caulobacters. By growing marine strains that tolerate low ionic strength media in a freshwater medium, we learned that the salts in seawater are apparently responsible for lowered adhesiveness to hydrophobic surfaces. Of practical significance was the finding that dimethylchlorosilane treated glass (ie classical "silanizing") was reasonably effective in discouraging attachment, a convenience for many future experiments, especially holdfast isolation procedures. We are presently using the same assay to determine the preference of the Caulobacter adhesive for various metal surfaces, by evaporating metals onto glass surfaces. We know the holdfast binds tightly to metals such as gold and silver, but wish to learn if there is a preference for these metals over other types of surfaces.

On the genes that specify the holdfast structure.
We prepared a library of 16000 independent transposon Tn5 insertions in Caulobacter crescentus CB2A, isolated 78 holdfast-defective mutants and began genetic analysis of the mutants to learn where the transposon was inserted. We have learned that the mutants cluster in 4 regions throughout the genome and have a number of phenotypes. Among them is the "shedder" phenotype discussed above. We also have cloned a segment of DNA from each of the 4 regions. Our results are summarized in this table:

<table>
<thead>
<tr>
<th>Cluster</th>
<th>Mutant Group</th>
<th># Detected</th>
<th>Phenotype</th>
<th>Size of Cloned Region (Restriction Enzyme)</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>g5</td>
<td>6</td>
<td>Reduced level or altered(?) holdfast</td>
<td>9.2 Kb (Eco RI)</td>
</tr>
<tr>
<td>B</td>
<td>g7</td>
<td>2</td>
<td>Holdfast shedder</td>
<td>12 Kb (SstI/EcoRI)</td>
</tr>
<tr>
<td></td>
<td>g8</td>
<td>2</td>
<td>&quot;&quot;</td>
<td>&quot;&quot;</td>
</tr>
<tr>
<td></td>
<td>g9</td>
<td>2</td>
<td>&quot;&quot;</td>
<td>&quot;&quot;</td>
</tr>
<tr>
<td>C</td>
<td>g1</td>
<td>6</td>
<td>No Holdfast</td>
<td>14.3 Kb (SstI)</td>
</tr>
<tr>
<td></td>
<td>g2</td>
<td>16</td>
<td>&quot;&quot;</td>
<td>&quot;&quot;</td>
</tr>
<tr>
<td>D</td>
<td>g3</td>
<td>30</td>
<td>No Holdfast</td>
<td>12.2 Kb (ClaI/KpnI)</td>
</tr>
<tr>
<td></td>
<td>g4</td>
<td>4</td>
<td>&quot;&quot;</td>
<td>&quot;&quot;</td>
</tr>
<tr>
<td></td>
<td>g10</td>
<td>1</td>
<td>Curved or weak stalk, Reduced level or altered(?) holdfast</td>
<td></td>
</tr>
<tr>
<td></td>
<td>g6</td>
<td>9</td>
<td>&quot;&quot;</td>
<td>&quot;&quot;</td>
</tr>
</tbody>
</table>

We are in the process now of a wide variety of experiments aimed at further characterization of the genetic regions involved with the holdfast production and are still searching for additional genes since we believe there was a degree of non-randomness in the transposon mutagenesis.

We are also repeating the process in the marine Caulobacter MCS6. In addition we will learn whether there is significant homology between the freshwater and marine holdfast genes. If
true, we will be able to use the cloned segments from the freshwater Caulobacter to speed up identification of marine holdfast genes.

**On the development of molecular genetic capabilities of marine Caulobacters.**
A very important development during the last several months was to determine the conditions for electroporation of both marine and freshwater Caulobacters. This is a method for introducing plasmids into bacteria without the necessity of transformation or conjugation techniques. The marine Caulobacters were particularly troublesome, since the technique tolerates very little ionic strength. We were able to adapt strain MCS6 (chosen for genetic studies) for growth in very low ionic strength medium and have successfully electroporated in plasmids at a frequency of $2 \times 10^3/\mu g$. This frequency is low compared to our results with freshwater Caulobacters ($5 \times 10^8/\mu g$) and we are working to improve the efficiency. But it is a very usable efficiency for many experiments.

**Publications**

*Manuscripts currently in preparation*


- Ong, C and J. Smit. The adhesive holdfast of *Caulobacters*: Isolation and analysis of mutants defective in the attachment of the organelle to the cell.


**Training Activities**

One graduate student (Canadian, Caucasian) and one postdoctoral (South African, Caucasian) are currently supported by this contract. This will rise to 2 of each by 1 September 1989.
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