

TATION PAGE

Form Approved
OMB No 0704-0188

1a REPC			1b RESTRICTIVE MARKINGS		
AD-A208 975			N/A		
2a. SECU N			3 DISTRIBUTION/AVAILABILITY OF REPORT		
2b. DECLASSIFICATION/DOWNGRADING SCHEDULE N/A			Distribution unlimited		
4. PERFORMING ORGANIZATION REPORT NUMBER(S) N/A			5. MONITORING ORGANIZATION REPORT NUMBER(S) N/A		
6a. NAME OF PERFORMING ORGANIZATION Univ. of British Columbia		6b. OFFICE SYMBOL (If applicable) N/A		7a. NAME OF MONITORING ORGANIZATION Office of Naval Research	
6c. ADDRESS (City, State, and ZIP Code) Department of Microbiology University of British Columbia Vancouver, B.C. V6T 1W5				7b. ADDRESS (City, State, and ZIP Code) 800 N. Quincy St. Arlington, VA 22217-5000	
8a. NAME OF FUNDING/SPONSORING ORGANIZATION Office of Naval Research		8b. OFFICE SYMBOL (If applicable) ONR		9 PROCUREMENT INSTRUMENT IDENTIFICATION NUMBER N00014-89-J-1749	
8c. ADDRESS (City, State, and ZIP Code) 800 N. Quincy St. Arlington, VA 22217-5000				10 SOURCE OF FUNDING NUMBERS	
		PROGRAM ELEMENT NO 61153 N		PROJECT NO RR04106	TASK NO 441h-006
				WORK UNIT ACCESSION NO.	-
11. TITLE (Include Security Classification) Chemical, structural & genetic analysis of the adhesive holdfast of biofouling caulobacters Cal					
12. PERSONAL AUTHOR(S) John Smit					
13a. TYPE OF REPORT Annual		13b. TIME COVERED FROM 3-1-89 TO 6-5-89		14. DATE OF REPORT (Year, Month, Day) 1989, June 5	
15. PAGE COUNT 3					
16. SUPPLEMENTARY NOTATION					
17. COSATI CODES			18. SUBJECT TERMS (Continue on reverse if necessary and identify by block number)		
FIELD	GROUP	SUB-GROUP	Biofouling, holdfast, Caulobacters, Adhesives Biofilms		
19. ABSTRACT (Continue on reverse if necessary and identify by block number)					
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. C. W.					
20. DISTRIBUTION/AVAILABILITY OF ABSTRACT ☒ UNCLASSIFIED/UNLIMITED ☐ SAME AS RPT ☐ DTIC USERS			21. ABSTRACT SECURITY CLASSIFICATION U		
22a. NAME OF RESPONSIBLE INDIVIDUAL M. Marron			22b. TELEPHONE (Include Area Code) (202) 696-4760		22c. OFFICE SYMBOL ONR

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

David White (U. of Tennessee) and Gordon Taylor (U. of Hawaii).

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 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 dimethyldichlorosilane 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:

<u>Cluster</u>	<u>Mutant Group</u>	<u># Detected</u>	<u>Phenotype</u>	<u>Size of Cloned Region</u> (Restriction Enzyme)
A . . .	g5 . . .	6	Reduced level or altered(?) holdfast	9.2 Kb (Eco R1)
B . . .	g7 . . .	2	Holdfast shedder	12 Kb (SstI/EcoR1)
	g8 . . .	2	" " " "	
	g9 . . .	2	" " " "	
C . . .	g1 . . .	6	No Holdfast . . .	14.3 Kb (SstI)
	g2 . . .	16	" " " "	
D . . .	g3 . . .	30	No Holdfast . . .	12.2 Kb (ClaI/KpnI)
	g4 . . .	4	" " " "	
	g10 . . .	1	" " " "	
	g6 . . .	9	Curved or weak stalk, Reduced level or altered(?) holdfast	

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 $2 \times 10^3/\mu\text{g}$. This frequency is low compared to our results with freshwater *Caulobacters* ($5 \times 10^8/\mu\text{g}$) and we are working to improve the efficiency. But it is a very usable efficiency for many experiments.

Publications

Manuscripts currently in preparation

- Merker, R. M., D. Rittschoff and J. Smit. Analysis of the attachment of marine and freshwater *Caulobacters* to surfaces using chemically defined substrates.
- Nivens, D.E. , A. Tunlid, M.J. Franklin, J. Smit, and D. White. Infrared monitoring of interactions between *Caulobacter* species and solid surfaces.
- 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.
- Gilchrist, A., W. Bingle and J. Smit. Introduction of plasmids into freshwater and marine *Caulobacters* by electroporation.
- Mitchell, D. and J. Smit. Identification and cloning of the genes affecting production of the adhesion organelle of *Caulobacter crescentus* CB2.

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.



Accession For	
NTIS GRA&I	☒
DTIC TAB	☐
Unannounced	☐
Justification	
By	
Distribution/	
Availability Codes	
Avail and/or	
Dist	Subanal
A-1	

Distribution List for Annual and Final Reports

1. Put a cover page (Form DD 1473) on your report and attach a copy of the distribution list. Mail one copy of the report to each person on the contractor subset list attached on which your name appears. The other subset list is for your information only. Please don't forget to attach this distribution list to your report - otherwise the folks below think they have mistakenly received the copy meant for the Molecular Biology Program and forward it to us.
2. Mail two copies to (include a DTIC Form 50 with these two copies too)
Administrator
Defense Technical Information Center
Building 5, Cameron Station
Alexandria, VA 22314
3. Mail one copy to each of the following:
 - (a) Dr. Michael Marron
ONR Code 1141
Molecular Biology Program
800 N. Quincy Street
Arlington, VA 22217-5000
 - (b) Administrative Contracting Officer
ONR Resident Representative
(address varies - see copy of your grant)
 - (c) Director,
Applied Research Directorate
ONR Code 12
800 N. Quincy Street
Arlington, VA 22217-5000
 - (d) Director
Office of Naval Technology
Code 22
800 N. Quincy Street
Arlington, VA 22217-5000
 - (e) Director
Chemical and Biological Sci Div
Army Research Office
P. O. Box 12211
Research Triangle Park, NC 27709
 - (f) Life Sciences Directorate
Air Force Office of Scientific Research
Bolling Air Force Base
Washington, DC 20332
 - (g) Director
Naval Research Laboratory
Technical Information Div, Code 2627
Washington, DC 20375