

<b>Fiscal Year:</b>	FY 2018	<b>Task Last Updated:</b>	FY 05/03/2018
<b>PI Name:</b>	Everroad, Craig Ph.D.		
<b>Project Title:</b>	Experimental Evolution of Bacillus subtilis Populations in Space; Mutation, Selection and Population Dynamics		
<b>Division Name:</b>	Space Biology		
<b>Program/Discipline:</b>			
<b>Program/Discipline--Element/Subdiscipline:</b>	SPACE BIOLOGY--Cellular and molecular biology		
<b>Joint Agency Name:</b>		<b>TechPort:</b>	No
<b>Human Research Program Elements:</b>	None		
<b>Human Research Program Risks:</b>	None		
<b>Space Biology Element:</b>	(1) Cell & Molecular Biology (2) Microbiology		
<b>Space Biology Cross-Element Discipline:</b>	(1) Reproductive Biology		
<b>Space Biology Special Category:</b>	None		
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<b>Organization Name:</b>	NASA Ames Research Center		
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<b>Zip Code:</b>	94035-0001	<b>Congressional District:</b>	18
<b>Comments:</b>	NOTE: PI previously at Bay Area Environmental Research Institute until 2018		
<b>Project Type:</b>	FLIGHT	<b>Solicitation:</b>	2014 Space Biology Flight NNH14ZTT001N
<b>Start Date:</b>	07/01/2015	<b>End Date:</b>	09/30/2019
<b>No. of Post Docs:</b>	0	<b>No. of PhD Degrees:</b>	0
<b>No. of PhD Candidates:</b>	0	<b>No. of Master' Degrees:</b>	0
<b>No. of Master's Candidates:</b>	0	<b>No. of Bachelor's Degrees:</b>	0
<b>No. of Bachelor's Candidates:</b>	0	<b>Monitoring Center:</b>	NASA ARC
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<b>Flight Program:</b>	ISS		
<b>Flight Assignment:</b>	NOTE: Extended to 9/30/2019 per F. Hernandez/ARC (Ed., 4/2/19) NOTE: Extended to 6/30/2019 per F. Hernandez/ARC and NSSC information (Ed., 8/8/18) NOTE: Period of performance changed to 7/01/2015-6/30/2018 per NSSC (Ed., 9/14/16) NOTE: End date change to 6/30/2018 per A. Chu/ARC and NSSC; start date to remain at 11/1/2014 per A. Chu/ARC (Ed., 9/23/15)		
<b>Key Personnel Changes/Previous PI:</b>	Ed. Note 8/8/18: PI Craig Everroad is now civil servant at NASA Ames and Robert Bergstrom, Ph.D., Bay Area Environmental Research Institute (BAERI), is CoPI at the BAERI for grant number NNX15AM68A.		
<b>COI Name (Institution):</b>	Bebout, Brad Ph.D. ( NASA Ames Research Center ) Koehne, Jessica Ph.D. ( NASA Ames Research Center ) Ricco, Antonio Ph.D. ( NASA Ames Research Center ) Bergstrom, Robert Ph.D. ( CoPI: Bay Area Environmental Research Institute, grant NNX15AM68A )		
<b>Grant/Contract No.:</b>	Internal Project ; NNX15AM68A		
<b>Performance Goal No.:</b>			

<b>Performance Goal Text:</b>	
<b>Task Description:</b>	<p>The proposed research aims to understand the effects of the space environment on evolutionary processes in the bacterium <i>Bacillus subtilis</i>. Different mutant lines will be 'raced' along solid surfaces to allow continuous selection in the cultures and to maximize the number of generations possible. Deep sequencing of winners will identify evolutionary rates, mechanisms, and targets of selection. We propose printing wax barriers to make paths along a growth surface (agar, membranes) and spotting each starting position of each path with dormant spores of the experimental bacteria to 'race' different mutants. Once on orbit, the material is wetted with growth medium, allowing the individual spots of <i>B. subtilis</i> to grow along their determined paths. This approach provides an opportunity for exponential growth only along the propagating edges, generating continuous bottlenecks thus amplifying selective pressures on the experimental populations. By monitoring the respective growth rate of different mutant lines maintained in each of these experimental conditions, we can estimate relative fitness of the lines. Long-term changes in relative growth rate indicate adaptation. Deep-sequencing of DNA from adapted cells ('winners' at the end of runs) will identify genetic changes within the respective populations. We expect that rates of mutation will differ between microgravity, 1-g, and ground controls, and that the targets of these mutations will differ as the different populations of bacteria adapt to their respective conditions. This research will also utilize the native ability of <i>B. subtilis</i> to uptake foreign DNA. Information-rich environmental DNA is added into the growth medium, and the populations are raced as above. By sampling the winners, and identifying if/what foreign genes are assimilated in each treatment, this experiment will identify potential genes of interest for future studies of genetic adaptation to the space environment. Our approach maximizes the number of generations possible in the 60-day window for this call, and maximizes the potential for evolutionary processes to occur. By performing multi-generational experimental evolution on bacteria on the International Space Station (ISS), the work proposed here aims to advance understanding of the evolutionary processes and challenges facing biological systems in long-term space exploration and habitation.</p>
<b>Rationale for HRP Directed Research:</b>	
<b>Research Impact/Earth Benefits:</b>	Improved understanding of the evolutionary process and in the dynamics of adaptive evolution in a model bacterium.
<b>Task Progress:</b>	<p>The objective of this study is to ascertain how evolutionary processes in bacteria change in response to the spaceflight environment, and specifically to microgravity. We propose to use growth rate as a proxy for fitness, and to 'race' a non-motile mutant of <i>Bacillus subtilis</i> along a membrane wetted with growth media and bounded by impassable printed wax barriers. As cells grow into the fresh media, they will create a front of newly divided cells. These 'racetracks' will be imaged as the cells propagate, and we will be able to observe changes in growth rate over time for treatments in microgravity, 1-g onboard the International Space Station (ISS), and 1-g on the ground. Deep-sequencing of winning lines will identify what genetic changes occurred with respect to the ancestral cells. This year's progress has been minimal due to limited budgets and the unavailability of the flight hardware of the European Modular Cultivation System (EMCS) onboard the ISS. Efforts have focused on transitioning the experiment into new hardware using the Techshot Multi-use Variable-g Platform (MVP). These include several discussions and planning sessions with Techshot representatives and the Space Biology Project Science team as new modules are designed for the MVP. A draft experimental requirements document (ERD) was provided to Techshot, and a Space Biology Compliance Review was conducted in April, 2018.</p> <p>The overall experimental framework and results from our science validation tests were presented as an oral presentation at the 33rd American Society for Gravitational and Space Research (ASGSR) meeting held in Renton, Washington in October, 2017.</p>
<b>Bibliography Type:</b>	Description: (Last Updated: 06/20/2019)
<b>Abstracts for Journals and Proceedings</b>	<p>Everroad RC, Detweiler AM, Koehne J, Ricco A. "Long-term multi-generational evolutionary studies of bacteria in the spaceflight environment." Talk given at the 33rd Annual Meeting of the American Society for Gravitational and Space Research, Seattle, WA, October 25-28, 2017.</p> <p>33rd Annual Meeting of the American Society for Gravitational and Space Research, Seattle, WA, October 25-28, 2017, Oct-2017</p>