

# Protective Role of Spore Structural Components in Determining *Bacillus subtilis* Spore Resistance to Simulated Mars Surface Conditions

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**Spores of wild-type and mutant *Bacillus subtilis* strains lacking various structural components were exposed to simulated Martian atmospheric and UV irradiation conditions. Spore survival and mutagenesis were strongly dependent on the functionality of all of the structural components, with small acid-soluble spore proteins, coat layers, and dipicolinic acid as key protectants.**

One major aspect of space biological research is the investigation of the responses of bacterial, viral, archaeal, and fungal species, as well as biomolecules, to simulated Martian conditions and their evaluation as potential forward contamination risks in the context of planetary protection (2, 4–7, 11, 12, 24, 30, 37–40, 44–46). For this reason, Mars environmental simulation experiments have been conducted to estimate (i) the survival rates of terrestrial microorganisms and (ii) the persistence of organic molecules on Mars (3, 4, 10, 15, 25, 29, 30, 37, 38, 45). Historically, several studies have explored the resistance of bacterial spores to simulated Martian conditions (8, 10, 12, 14, and references therein); these studies have concentrated mainly on the survival of spores of wild-type strains of various spore-forming species (10, 32, 38–40). More recent experiments have attempted to better understand the molecular factors causing spore resistance to environmental extremes, in which mainly spores of the model organism *Bacillus subtilis* that carry mutations affecting spore protective factors or spore DNA repair systems have been used (9, 18–23; reviewed in references 16, 28, 41, and 42). Spores of *B. subtilis* possess brownish pigmentation, thick layers of highly cross-linked coat proteins, a modified peptidoglycan spore cortex, a low core water content, and abundant intracellular constituents such as the calcium chelate of dipicolinic acid (Ca-DPA) and  $\alpha/\beta$ -type small, acid-soluble spore proteins (SASP), all factors which have been previously found to contribute to spore resistance (reviewed in references 16, 28, 41, and 42). However, the possible roles of these factors in spore resistance to the extreme environmental conditions prevailing on the surface of Mars have not been explored, which is the subject of the present work.

The *B. subtilis* strains used in this study are listed in Table 1, and all of the strains used were congenic to their respective wild-type strains. Spores were obtained by cultivation under vigorous aeration in double-strength liquid Schaeffer sporulation medium (36) under identical conditions for each strain, purified, and stored as described previously (31). When appropriate, chloramphenicol (5  $\mu\text{g}/\text{ml}$ ), erythromycin (1  $\mu\text{g}/\text{ml}$ ), spectinomycin (100  $\mu\text{g}/\text{ml}$ ), or tetracycline (10  $\mu\text{g}/\text{ml}$ ) was added to the medium. Spore preparations consisted of single spores with no detectable clumps and were >99% free of growing cells, germinated spores, and cell debris, as seen in a phase-contrast microscope. Triplicate air-dried spore samples with a thickness of approximately 25 spore layers (for the  $5 \times 10^8$  spore concentration) on presterilized spacecraft-

qualified, chemfilm-treated aluminum 6061 coupons (13 mm in diameter by 1 mm thick) (23) were exposed to simulated Martian conditions in a cylindrical Mars simulation chamber (MSC) as described in detail previously (37). The simulated Mars environmental conditions listed in Table 2 represent a worst-case scenario for high UV flux and thus likely equate to an upper bound for UV effects on *B. subtilis* spore survival under simulated Martian conditions (38). Three types of spore exposure conditions were tested: (i) ambient laboratory conditions (22°C, 54% relative humidity, on the laboratory bench shielded from light), (ii) exposure in the MSC to simulate Mars conditions but shielded from radiation [designated Mars(–)UV], and (iii) exposure to the full suite of simulated Mars conditions, including 8 h of solar radiation [designated Mars(+ )UV]. Further details of the design and geometry of the MSC, radiation dosimetry, and fluence calculations have been described in detail elsewhere by Schuerger et al. (38–40). Spore recovery from coupons was accomplished by stripping off spores with polyvinyl alcohol, and viability assays were performed as described previously (18, 19, 21, 22). The surviving fraction of *B. subtilis* spores was determined from the  $N/N_0$  ratio, with  $N$  being the number of CFU of the exposed sample and  $N_0$  that of the initial spore burden (directly determined after sample preparation). To determine frequencies of mutation to 4-azaleucine resistance (4-azaleu<sup>r</sup>), aliquots of spores of each strain were taken from Mars(+ )UV and Mars(–)UV exposure conditions, as well as laboratory control samples. Aliquots from serial dilutions were spread on Spizizen minimal medium plates supplemented with (final concentrations) tryptophan (50  $\mu\text{g}/\text{ml}$ ), spore germinants (either L-alanine [10 mM] or an equimolar mixture of DPA and  $\text{CaCl}_2$  [60 mM each] [1]), and 4-DL-azaleucine (100  $\mu\text{g}/\text{ml}$ ). Colonies that were 4-azaleu<sup>r</sup> were counted 36 h after incubation at 37°C as described by Rivas-Castillo et al. (35). All data are expressed as averages  $\pm$  standard deviations ( $n = 3$ ). The significance of the differences in the survival rates and 4-azaleu<sup>r</sup> muta-

Received 15 August 2012 Accepted 4 October 2012

Published ahead of print 12 October 2012

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doi:10.1128/AEM.02527-12

TABLE 1 *B. subtilis* strains used in this study

Strain	Genotype and/or phenotype <sup>a</sup>	Missing spore component(s)	Source (reference)
168 (WN131)	Wild-type parent of WN659 and WN661; <i>trpC2</i>	None	Laboratory stock (4)
PS832 (WN552)	Wild-type parent of PS356, PS1899, FB72, and FB108; prototroph	None	P. Setlow (44)
PY79 (WN470)	Wild-type parent of AD17, AD28, and AD142; prototroph	None	A. Driks (34)
AD17 (WN496)	<i>gerE36</i>	Inner coat	A. Driks (34)
AD28 (WN495)	<i>cotE::cat</i> Cm <sup>r</sup>	Outer coat	A. Driks (34)
AD142 (WN469)	<i>gerE36 cotE::cat</i> Cm <sup>r</sup>	Inner and outer coats	A. Driks (34)
PS356 (WN1273)	$\Delta$ <i>sspA</i> $\Delta$ <i>sspB</i>	SASP- $\alpha$ and SASP- $\beta$	P. Setlow (17)
PS1899 (WN1274)	<i>dacB::cat</i> Cm <sup>r</sup>	Core dehydration	P. Setlow (33)
FB72 (WN552)	$\Delta$ <i>gerA::spc</i> $\Delta$ <i>gerB::cat</i> $\Delta$ <i>gerK::erm</i> Spc <sup>r</sup> Cm <sup>r</sup> Erm <sup>r</sup> (referred to as $\Delta$ <i>gerABK</i> DPA <sup>+</sup> spores)	Germination receptors	P. Setlow (44)
FB108 (WN553)	$\Delta$ <i>gerA::spc</i> $\Delta$ <i>gerB::cat</i> $\Delta$ <i>gerK::erm</i> $\Delta$ <i>spoVF::tet</i> Spc <sup>r</sup> Cm <sup>r</sup> Erm <sup>r</sup> Tet <sup>r</sup> (referred to as $\Delta$ <i>gerABK</i> DPA <sup>-</sup> spores)	Germination receptors, DPA	P. Setlow (44)
WN661	<i>trpC2</i> , presumed mutation in <i>cotA</i> <sup>b</sup>	Pigmentation	W. L. Nicholson (unpublished data)

<sup>a</sup> Cm<sup>r</sup>, resistant to chloramphenicol (5  $\mu$ g/ml); Erm<sup>r</sup>, resistant to erythromycin (1  $\mu$ g/ml); Spc<sup>r</sup>, resistant to spectinomycin (100  $\mu$ g/ml); Tet<sup>r</sup>, resistant to tetracycline (10  $\mu$ g/ml).

<sup>b</sup> Reference 13.

tion induction were determined by analysis of variance (ANOVA) using Analyze-it software (Analyze-it Software, Ltd., Leeds, United Kingdom). Values were evaluated in multigroup pairwise combinations, and differences with *P* values of  $\leq 0.05$  were considered statistically significant (18, 19, 22, 34, 43, 44).

Wild-type and structural-component-deficient spores (Table 1) were exposed to simulated Mars environmental conditions for a 24-h period in the MSC with or without 8 h of exposure to simulated Martian solar radiation, including UV (200 to 400 nm) (Fig. 1). It was noted that a 24-h exposure to ambient lab conditions resulted in a 19%  $\pm$  6% average reduction in the survival of wild-type and all mutant spores, with the exception of spores lacking both SASP- $\alpha$  and SASP- $\beta$ , which exhibited a statistically significant 72%  $\pm$  13% loss of spore viability (Fig. 1A). Exposure of spores for 24 h to simulated Mars(-)UV conditions in the MSC did not significantly reduce the viability of wild-type spores but caused a significant reduction in the viability of all of the mutant spores tested, most dramatically, *sspA sspB* mutant spores lacking

the major  $\alpha/\beta$ -type SASP and *gerE cotE* mutant spores lacking inner and outer spore coats (Fig. 1B). Exposure to the full suite of simulated Mars(+)-UV conditions reduced the viability of wild-type spores by  $\sim 1$  order of magnitude, whereas spores of all of the mutant strains tested were significantly more sensitive than wild-type spores, again with spores lacking either  $\alpha/\beta$ -type SASP or spore coats being the most sensitive (Fig. 1C).

To directly compare the contributions of spore structural components to the resistance of spores to simulated Mars conditions, we calculated the relative increase in the sensitivity of each mutant strain with respect to that of the corresponding wild-type strain (Fig. 2). Ranked in order from most to least important with respect to Mars(-)UV treatment were the inner and outer spore coat (*cotE gerE*),  $\alpha/\beta$ -type SASP (*sspA sspB*)  $\gg$  the coat pigment (*cotA*), the inner coat (*gerE*)  $>$  the outer coat (*cotE*), core water content (*dacB*), DPA (*dpaAB*), and germination receptors (*gerA gerB gerK*) (Fig. 2A). Mars(-)UV conditions are dominated by low temperature ( $-10^{\circ}\text{C}$ ) and low pressure ( $\sim 7$  mbar), resulting in an extremely desiccating environment. An intact spore coat and  $\alpha/\beta$ -type SASP are the most important factors for dormant-spore survival under Mars(-)UV conditions, whereas DPA, pigmentation, and spore water content play less important roles. Under Mars(+)-UV conditions, the spore structural components contributing to resistance, from most to least important, were  $\alpha/\beta$ -type SASP (*sspA sspB*)  $>$  the inner and outer spore coats (*cotE gerE*), DPA (*gerA gerB gerK spoVF*)  $\gg$  pigmentation (*cotA*), the outer spore coat (*cotE*)  $>$  core water content (*dacB*)  $>$  the inner coat (*gerE*)  $\cong$  germination receptors (*gerA gerB gerK*) (Fig. 2B). Under Mars(+)-UV conditions, UV radiation is a dominant factor in determining spore survival. In this context, it is not surprising that  $\alpha/\beta$ -type SASP, the spore coat, and DPA make major contributions to spore resistance, as all of these factors have previously been shown to protect air-dried spores from polychromatic UV (21, 34, 43, 44). An intact outer coat, including the outer-coat-associated *cotA*-encoded pigment, was apparently more important to spore survival under Mars(+)-UV conditions than was an intact inner coat (Fig. 2B).

We also measured the frequency of mutation to 4-azaleu<sup>r</sup> in spores exposed to Mars(-)UV or Mars(+)-UV conditions (Fig. 3). The relative importance of spore factors for survival of Mars(-)UV exposure was as follows: intact spore coats,  $\alpha/\beta$ -type

TABLE 2 Environmental conditions used for Mars simulation experiments

Parameter	Value
Avg pressure (kPa) $\pm$ SD	0.69 $\pm$ 0.01
Avg temp ( $^{\circ}\text{C}$ ) $\pm$ SD	$-10 \pm 2$
Avg relative humidity (%) $\pm$ SD	$8 \pm 2$
UV-Vis-NIR radiation <sup>a</sup> fluence rate, $\text{kJ m}^{-2} \text{h}^{-1}$ (total applied fluence) <sup>a</sup>	
Total UV (200–400 nm)	142.9 (1.14 $\text{MJ m}^{-2}$ )
UV-C (200–280 nm)	14.4 kJ (115 $\text{kJ m}^{-2}$ )
UV-B (280–320 nm)	24.8 kJ (199 $\text{kJ m}^{-2}$ )
UV-A (320–400 nm)	103.7 kJ (829 $\text{kJ m}^{-2}$ )
Vis (400–700 nm)	864.0 kJ (6.91 $\text{MJ m}^{-2}$ )
NIR (700–1,100 nm)	882.0 kJ (7.05 $\text{MJ m}^{-2}$ )
Total irradiance (200–1,100 nm)	1,888.9 (15.1 $\text{MJ m}^{-2}$ )
Time (h)	24 <sup>b</sup>
% $\text{CO}_2$	99.99

<sup>a</sup> Details of the radiation dosimetry, fluence rates, and simulation of the UV-Vis-NIR radiation environment of the surface conditions on Mars have been described by Schuergel et al. (38–40). Vis, visible; NIR, near IR.

<sup>b</sup> With or without 8 h of irradiance.

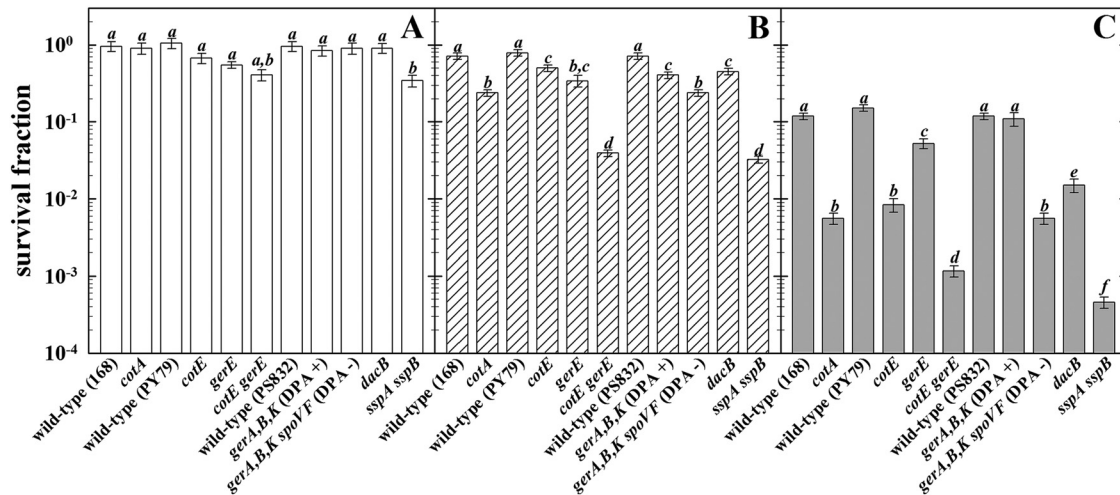


FIG 1 Survival of *B. subtilis* spores in response to simulated Martian environmental conditions, as determined by the ability to form macroscopic visible colonies. The strains used are indicated below the bars; the wild-type strain is listed to the left of the corresponding mutant strain. Spores were exposed as air-dried spore multilayers for 24 h to ambient laboratory bars (white bars; panel A), Mars(-)UV conditions (hatched bars; panel B), or Mars(+)-UV conditions (shaded bars; panel C). Data are expressed as averages and standard deviations ( $n = 3$ ). The lowercase letters above the bars denote groups significantly different by ANOVA ( $P < 0.05$ ).

SASP > an outer coat, DPA >> pigment, reduced core water >>> an inner coat  $\geq$  germination receptors (Fig. 3). Under Mars(+)-UV conditions, spore components exhibited importance in mutagenesis to 4-azaleu<sup>r</sup>, in order from highest to lowest frequency, as follows:  $\alpha/\beta$ -type SASP > intact spore coats > pigment, outer coat, DPA, reduced spore core water, and germination receptors (Fig. 3). Mutagenesis to 4-azaleu<sup>r</sup> was strongly UV dependent, which is due mainly to the two ways for UV to exert its lethal effects, i.e., direct interaction with spore DNA and indirectly

via the generation of reactive oxygen species, as shown previously (22, 35; reviewed in reference 12).

On the basis of the above data, it thus appears that  $\alpha/\beta$ -type SASP and intact spore coat layers are the most important factors in determining spore survival and protection from mutagenic damage under simulated Mars conditions either with or without UV exposure. Spore pigmentation, the outer or inner spore coat layers by themselves, reduced spore water content, and DPA play less important roles in spore survival of Mars(-)-UV exposure but do

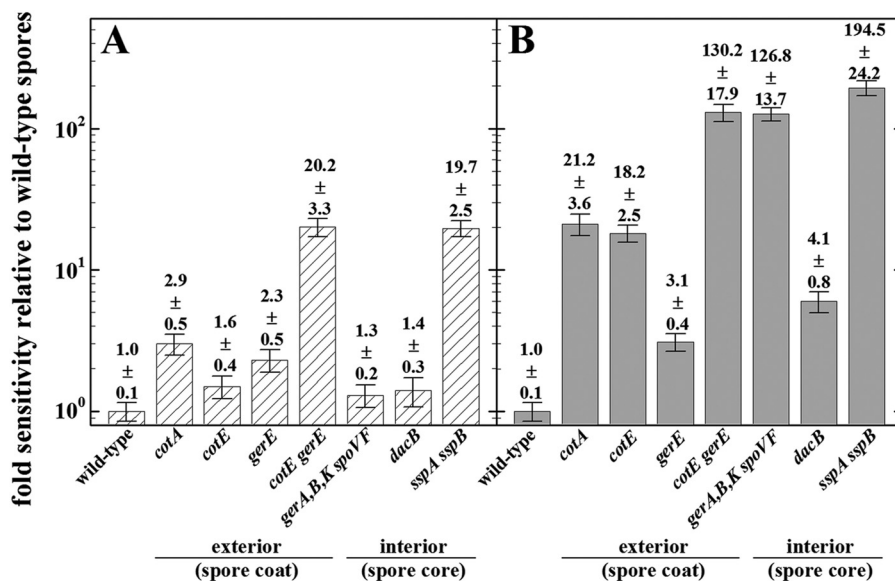
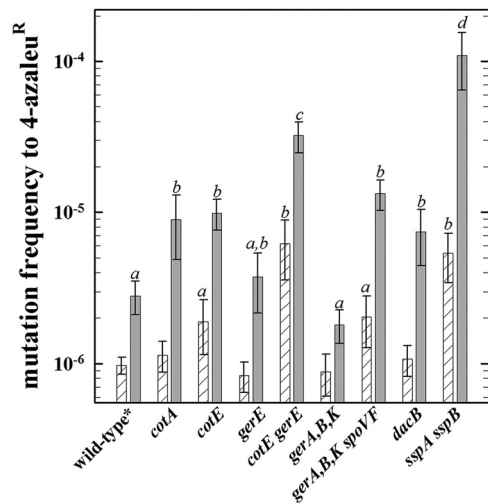


FIG 2 Impacts of the spore-specific structural attributes (exterior and interior) are displayed as relative sensitivities of spores lacking pigmentation, spore coat assembly, dipicolinic acid (DPA) formation, core dehydration, and  $\alpha/\beta$ -type small, acid-soluble spore protein (SASP) formation to Mars(-)-UV conditions (hatched bars; panel A) or Mars(+)-UV conditions (shaded bars; panel B). Relative spore sensitivity was expressed as the ratio of the survival of each mutant strain with respect to the survival of the corresponding wild-type strain from the respective exposure to simulated Martian conditions (from Fig. 1B and C) to that of the ambient control (Fig. 1A). Data are averages and standard deviations ( $n = 3$ ). The actual data values are shown above the corresponding columns (Fig. 1).



**FIG 3** Frequencies of mutation to 4-azaleu<sup>R</sup> of spores of different strains exposed to the full suite of simulated Martian conditions shielded (white hatched bars) and including UV radiation (gray bars). Data are averages and standard deviations ( $n = 3$ ). The lowercase letters above the bars denote groups significantly different by ANOVA ( $P < 0.05$ ). The spontaneous frequency of mutation to 4-azaleu<sup>R</sup> of the wild-type strain (i.e., the mean spontaneous 4-azaleu<sup>R</sup> mutation frequencies of strains 168, PY79, and PS832  $\pm$  the standard deviation) was  $(9.7 \pm 3.6) \times 10^{-7}$ , in good agreement with previous data (35).

play significant roles in spore resistance to Mars(+)UV exposure. Clearly, the data indicate that spore DNA is a major target of both lethal and mutagenic damage, on the basis of the observation that all spore factors contributed significantly to both spore survival (Fig. 2) and the frequency of mutation to 4-azaleu<sup>R</sup> (Fig. 3) in response to the full suite of simulated Mars environmental conditions, including UV. The findings in this communication complement the results of microarray experiments demonstrating the upregulation of DNA repair pathways during the germination of spores exposed to simulated Mars surface conditions (27). Future goals of our work are to (i) compile a complete catalog of all types of cellular damage incurred by spores exposed to simulated Martian conditions; (ii) elucidate the potential additive effects of the sensitive spore phenotypes, such as SASP, core water content, and DPA, with a special focus on the complex interaction of all known spore coat proteins (as reviewed in reference 16 and references therein); and (iii) elucidate the roles of various DNA repair systems in the survival of Mars-exposed spores in order to gain a deeper understanding of the physiological responses used by spore-forming bacteria to cope with the environmental challenges of simulated extraterrestrial environments.

## ACKNOWLEDGMENTS

We thank Krystal Kerney and Andrea Rivas-Castillo for their technical assistance during parts of this work and Adam Driks and Peter Setlow for their generous donations of *B. subtilis* strains. We thank the three anonymous reviewers for insightful comments.

This work was supported in part by grants from the NASA Planetary Protection (NNA06CB58G) and Astrobiology: Exobiology and Evolutionary Biology (NNX08AO15G) programs to W.L.N. and A.C.S. R.M. and G.R. were supported by DLR grant DLR-FuE-Projekt ISS Nutzung in der Biodiagnostik, Programm RF-FuW, Teilprogramm 475.

## REFERENCES

1. Atluri S, Ragkousi K, Cortezzo DE, Setlow P. 2006. Cooperativity between different nutrient receptors in germination of spores of *Bacillus*

2. *subtilis* and reduction of this cooperativity by alterations in the GerB receptor. *J. Bacteriol.* **188**:28–36.
3. Berry BJ, Jenkins DG, Schuerger AC. 2010. Effects of simulated Mars conditions on the survival and growth of *Escherichia coli* and *Serratia liquefaciens*. *Appl. Environ. Microbiol.* **76**:2377–2386.
4. Bruckner JC, Osman S, Conley C, Venkateswaran K. 2008. Space microbiology: planetary protection, burden, diversity and significance of spacecraft associated microbes, p 52–65. In Schaechter M (ed), *Encyclopedia of microbiology*. Elsevier, Oxford, United Kingdom.
5. Fajardo-Cavazos P, Schuerger AC, Nicholson WL. 2008. Persistence of biomarker ATP and ATP-generating capability in bacterial cells and spores contaminating spacecraft materials under Earth conditions and in a simulated Martian environment. *Appl. Environ. Microbiol.* **74**:5159–5167.
6. Fekete A, Kovács G, Hegedüs M, Módos K, Lammer H. 2008. Biological responses to the simulated Martian UV radiation of bacteriophages and isolated DNA. *J. Photochem. Photobiol. B* **92**:110–116.
7. Fendrihan S, et al. 2009. Investigating the effects of simulated Martian ultraviolet radiation on *Halococcus dombrowskii* and other extremely halophilic archaeobacteria. *Astrobiology* **9**:104–112.
8. Foster TL, Winans L, Jr, Casey RC, Kirschner LE. 1978. Response of terrestrial microorganisms to a simulated Martian environment. *Appl. Environ. Microbiol.* **35**:730–737.
9. Fulton JD. 1958. Survival of terrestrial microorganisms under simulated Martian conditions, p 606–613. In Benson OO, Jr, Strughold H (ed), *Proceedings of Physics and Medicine of the Atmosphere and Space*, San Antonio, Texas. John Wiley & Sons, New York, NY.
10. Granger AC, Gaidamakova EK, Matrosova VY, Daly MJ, Setlow P. 2011. Effects of Mn and Fe Levels on *Bacillus subtilis* spore resistance and effects of Mn<sup>2+</sup>, other divalent cations, orthophosphate, and dipicolinic acid on protein resistance to ionizing radiation. *Appl. Environ. Microbiol.* **77**:32–40.
11. Hagen CA, Ehrlich R, Hawrylewicz E. 1964. Survival of microorganisms in simulated Martian environment. I. *Bacillus subtilis* var. *globigii*. *Appl. Microbiol.* **12**:215–218.
12. Hawrylewicz E, Gowdy B, Ehrlich R. 1962. Microorganisms under a simulated Martian environment. *Nature* **193**:497.
13. Horneck G, Klaus DM, Mancinelli RL. 2010. Space microbiology. *Microbiol. Mol. Biol. Rev.* **74**:121–156.
14. Hullo MF, Moszer I, Danchin A, Martin-Verstraete I. 2001. CotA of *Bacillus subtilis* is a copper-dependent laccase. *J. Bacteriol.* **183**:5426–5430.
15. Jensen LL, et al. 2008. A facility for long-term Mars simulation experiments: the Mars environmental simulation chamber (MESCH). *Astrobiology* **8**:537–548.
16. Kerney KR, Schuerger AC. 2011. Survival of *Bacillus subtilis* endospores on ultraviolet-irradiated rover wheels and Mars regolith under simulated Martian conditions. *Astrobiology* **11**:477–485.
17. Leggett MJ, McDonnell G, Denyer SP, Setlow P, Maillard JY. 2012. Bacterial spore structures and their protective role in biocide resistance. *J. Appl. Microbiol.* **113**:485–498.
18. Mason JM, Setlow P. 1986. Essential role of small, acid-soluble spore proteins in resistance of *Bacillus subtilis* spores to UV light. *J. Bacteriol.* **167**:174–178.
19. Moeller R, et al. 2007. Role of DNA repair by nonhomologous-end joining in *Bacillus subtilis* spore resistance to extreme dryness, mono- and polychromatic UV, and ionizing radiation. *J. Bacteriol.* **189**:3306–3311.
20. Moeller R, et al. 2008. Roles of the major, small, acid-soluble spore proteins and spore-specific and universal DNA repair mechanisms in resistance of *Bacillus subtilis* spores to ionizing radiation from X rays and high-energy charged-particle bombardment. *J. Bacteriol.* **190**:1134–1140.
21. Moeller R, Rohde M, Reitz G. 2010. Effects of ionizing radiation on the survival of bacterial spores in artificial Martian regolith. *Icarus* **206**:783–786.
22. Moeller R, Setlow P, Reitz G, Nicholson WL. 2009. Roles of small, acid-soluble spore proteins and core water content in survival of *Bacillus subtilis* spores exposed to environmental solar UV radiation. *Appl. Environ. Microbiol.* **75**:5202–5208.
23. Moeller R, et al. 2011. Role of the Nfo and ExoA apurinic/apyrimidinic endonucleases in radiation resistance and radiation-induced mutagenesis of *Bacillus subtilis* spores. *J. Bacteriol.* **193**:2875–2879.
24. Moeller R, Schuerger AC, Reitz G, Nicholson WL. 2011. Impact of two DNA repair pathways, homologous recombination and non-homologous

- end joining, on bacterial spore inactivation under simulated Martian environmental conditions. *Icarus* 215:204–210.
24. Morozova D, Möhlmann D, Wagner D. 2007. Survival of methanogenic archaea from Siberian permafrost under simulated Martian thermal conditions. *Orig. Life Evol. Biosph.* 37:189–200.
  25. Newcombe DA, et al. 2005. Survival of spacecraft-associated microorganisms under simulated Martian UV irradiation. *Appl. Environ. Microbiol.* 71:8147–8156.
  26. Nicholson WL, et al. 2002. Bacterial endospores and their significance in stress resistance. *Antonie Van Leeuwenhoek* 81:27–32.
  27. Nicholson WL, Moeller R, the PROTECT Team, Horneck G. 2012. Transcriptomic responses of germinating *Bacillus subtilis* spores exposed to 1.5 years of space and simulated Martian conditions on the EXPOSE-E experiment PROTECT. *Astrobiology* 12:469–486.
  28. Nicholson WL, Munakata N, Horneck G, Melosh HJ, Setlow P. 2000. Resistance of bacterial endospores to extreme terrestrial and extraterrestrial environments. *Microbiol. Mol. Biol. Rev.* 64:548–572.
  29. Nicholson WL, Schuerger AC. 2005. *Bacillus subtilis* spore survival and expression of germination-induced bioluminescence after prolonged incubation under simulated Mars atmospheric pressure and composition: implications for planetary protection and lithopanspermia. *Astrobiology* 5:536–544.
  30. Nicholson WL, Schuerger AC, Race MS. 2009. Migrating microbes and planetary protection. *Trends Microbiol.* 17:389–392.
  31. Nicholson WL, Setlow P. 1990. Sporulation, germination, and outgrowth, p 391–450. In Harwood CR, Cutting SM (ed), *Molecular biological methods for Bacillus*. John Wiley and Sons, Sussex, England.
  32. Osman S, et al. 2008. Effect of shadowing on survival of bacteria under conditions simulating the Martian atmosphere and UV radiation. *Appl. Environ. Microbiol.* 74:959–970.
  33. Popham DL, Sengupta S, Setlow P. 1995. Heat, hydrogen peroxide, and UV resistance of *Bacillus subtilis* spores with increased core water content and with or without major DNA-binding proteins. *Appl. Environ. Microbiol.* 61:3633–3638.
  34. Riesenman PJ, Nicholson WL. 2000. Role of the spore coat layers in *Bacillus subtilis* spore resistance to hydrogen peroxide, artificial UV-C, UV-B, and solar UV radiation. *Appl. Environ. Microbiol.* 66:620–626.
  35. Rivas-Castillo AM, Yasbin RE, Robleto E, Nicholson WL, Pedraza-Reyes M. 2010. Role of the Y-family DNA polymerases YqjH and YqjW in protecting sporulating *Bacillus subtilis* cells from DNA damage. *Curr. Microbiol.* 60:263–267.
  36. Schaeffer P, Millet J, Aubert J-P. 1965. Catabolic repression of bacterial sporulation. *Proc. Natl. Acad. Sci. U. S. A.* 54:704–711.
  37. Schuerger AC, et al. 2008. Slow degradation of ATP in simulated Martian environments suggests long residence times for the biosignature molecule on spacecraft surfaces on Mars. *Icarus* 194:86–100.
  38. Schuerger AC, Mancinelli RL, Kern RG, Rothschild LJ, McKay CP. 2003. Survival of endospores of *Bacillus subtilis* on spacecraft surfaces under simulated Martian environments: implications for the forward contamination of Mars. *Icarus* 165:253–276.
  39. Schuerger AC, Nicholson WL. 2006. Interactive effects of hypobaric, low temperature, and CO<sub>2</sub> atmospheres inhibit the growth of mesophilic *Bacillus* spp. under simulated Martian conditions. *Icarus* 185:143–152.
  40. Schuerger AC, Richards JT, Hintze PE, Kern RG. 2005. Surface characteristics of spacecraft components affect the aggregation of microorganisms and may lead to different survival rates of bacteria on Mars landers. *Astrobiology* 5:545–559.
  41. Setlow P. 2006. Spores of *Bacillus subtilis*: their resistance to and killing by radiation, heat and chemicals. *J. Appl. Microbiol.* 101:514–525.
  42. Setlow P. 2007. I will survive: DNA protection in bacterial spores. *Trends Microbiol.* 15:172–180.
  43. Slieman TA, Nicholson WL. 2000. Artificial and solar UV radiation induces strand breaks and cyclobutane dimers in *Bacillus subtilis* spore DNA. *Appl. Environ. Microbiol.* 66:199–205.
  44. Slieman TA, Nicholson WL. 2001. Role of dipicolinic acid in survival of *Bacillus subtilis* spores exposed to artificial and solar UV radiation. *Appl. Environ. Microbiol.* 67:1274–1279.
  45. Stoker CR, Bullock MA. 1997. Organic degradation under simulated Martian conditions. *J. Geophys. Res.* 102:10881–10888.
  46. Tauscher C, Schuerger AC, Nicholson WL. 2006. Survival and germinability of *Bacillus subtilis* spores exposed to simulated Mars solar radiation: implications for life detection and planetary protection. *Astrobiology* 6:592–605.
  47. ten Kate IL, et al. 2003. Investigating complex organic compounds in a simulated Mars environment. *Int. J. Astrobiol.* 1:387–399.