# Novel Mechanism for Surface Layer Shedding and Regenerating in Bacteria Exposed to Metal-Contaminated Conditions

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11 Lysinibacillus<sub>6</sub>. (Min.5-Max. 8)

### 12 Abstract

- 13 Surface layers (S-layers) are self-assembling, ordered structures composed of repeating protein
- 14 subunits found as components of the cell walls throughout the Bacteria and the Archaea. S-layers act

15 as an interface between prokaryotic cells and their surrounding environment, and provide protection

16 for microorganisms against diverse environmental stresses including heavy metal stress. We have

17 previously characterized the process by which S-layers serve as a nucleation site for metal

18 mineralization in the presence of high concentration of metals. Here, we test the hypothesis originally

19 proposed in cyanobacteria that a "shedding" mechanism exists in prokaryotes for replacing S-layers

- 20 that have become mineral-encrusted. We used a metallotolerant gram-positive bacterium bearing an
- S-layer, *Lysinibacillus* sp. TchIII 20n38, as a model organism. We characterize for the first time a
   mechanism for resistance to metals through S-layer shedding and regeneration. S-layers nucleate the
- formation of Fe-mineral on the cell surface, leading to the encrustation of the S-layer. Using a
- 24 combination of scanning electron microscopy (SEM) and nanoSIMS, we show that mineral-encrusted
- 25 S-layers are shed by the bacterial cells, and the emerging cells regenerate new S-layers as part of
- 26 their cell wall structure. This novel mechanism for the survival of prokaryotes in metal-contaminated

27 environments may also provide elements necessary for the development of renewable systems for

28 metal bioremediation.

# 29 **1** Introduction

30 Environmental contamination by metals and radionuclides from activities such as mining and nuclear

31 power generation pose a serious risk to human health. The sudden, accidental release of high

- 32 concentrations of iron from acid mine drainage from the Gold King Mine polluted the Animas River
- in 2015, mixing downstream with phosphates from agricultural runoff (Rodriguez-Freire et al.,
- 34 2016). Metal contamination affected both water supplies from soluble metals, and sediments after the
- 35 sedimentation of the majority of released metals. Sampling of soils contaminated by metals and
- 36 radionuclides near the former Chernobyl nuclear reactor site (Chapon et al., 2012) and in uranium
- 37 mining waste piles in Germany (Pollmann et al., 2006) have identified bacteria of the genre

- 38 *Lysinibacillus* tolerant to these contaminants. *Lysinibacillus* (formerly classified as part of the
- 39 *Bacillus* genre (Ahmed et al., 2007)) gram-positive bacteria, with a peptidoglycan cell wall enclosed
- 40 by a surface layer ("S-layer") attached non-covalently to the lipopolysaccharides of the outer
- 41 membrane (reviewed in (Sleytr et al., 2014)). These S-layers have proven to be a key mechanism for
- 42 metallotolerance in *Lysinibacillus* as they have been shown to bind U, Pd(II), Cu, Pt(II), and Au(III)
- 43 (Pollmann et al., 2006).
- 44 S-layers, however are not unique to *Lysinibacillus*. They are common components of the cell
- 45 envelopes of both bacteria and archaea. S-layers are formed by self-assembly of repeated protein
- 46 monomers into ordered structures (oblique, square, or hexagonal) depending on the number of
- 47 subunits composing the ordered structure. This self-assembly occurs even in the absence of cells *in*
- 48 *vitro*; a capacity has been exploited in biotechnology in everything from the development of vaccine
- 49 to nanomaterials to filtration technologies (Sleytr et al., 2011).
- 50 S-layers form the interaction interface between prokaryotic cells and their external environment, and
- are therefore in contact with metals and other ions present. Nucleation of mineralization by S-layers
- 52 was first noted in cyanobacteria by Schultze-Lam in the early 1990s (Schultze-Lam et al., 1992).
- 53 Cyanobacterial S-layers were demonstrated to nucleate the formation of carbonates of calcium,
- 54 magnesium, and strontium (Schultze-Lam and Beveridge, 1994). In their 1992 article, Schutlze-Lam
- 55 proposed the hypothesis that mineral-encrusted S-layers are shed from cyanobacteria as part of a
- 56 protective mechanism to ensure that essential cell activities are maintained despite cell wall
- 57 mineralization (Schultze-Lam et al., 1992). This hypothesis was, however, never fully tested. Since
- 58 then, S-layer nucleation of mineralization has been observed in a range of bacteria (Konhauser et al.,
- 59 1994; Phoenix et al., 2000) and archaea (Kish et al., 2016).
- 60 S-layer mineralization is a mechanism for metallotolerance with potential application in
- 61 bioremediation. S-layer shedding and regeneration in metallotolerant bacteria already inhabiting
- 62 contaminated environments provides an added benefit of being a renewable system. Here, we
- 63 describe the shedding and regeneration of mineral-encrusted S-layers in the metallotolerant
- 64 environmental isolate *Lysinibacillus* sp. TchIII 20n38.

# 65 2 Materials and Methods

# 66 **2.1 Culture and growth conditions**

- The bacterial strain used was an environmental strain isolated in 2009 from soils near a radionuclidecontaminated site (Chapon et al., 2012). This strain, referenced as *Lysinibacillus* sp. TchIII 20n38.
- contaminated site (Chapon et al., 2012). This strain, referenced as *Lysinibacillus* sp. TchIII 20n38,
  was cultured at 30 °C in Luria Bertani (LB) medium under aerobic conditions with agitation
- 70 (180 rpm) to mid-exponential, late-exponential, and stationary growth phases ( $OD_{600nm} = 0.3, 0.6, and$
- 71 1.0, respectively). The culture medium was then removed and the cells washed in MilliO-H<sub>2</sub>O by
- 72 gentle centrifugation (2600 x g, 15 min, room temperature). In order to determine the mechanisms of
- 73 resistance of *Lysinibacillus* sp. TchIII 20n38 cells to the presence of heavy metals, the cells were
- resuspended to an equivalent cell density in a Fe-rich solution at a similar pH to that found in the
- 75 Chernobyl isolation (10 mM NaH<sub>2</sub>PO<sub>4</sub>, 10 mM FeSO<sub>4</sub>, pH=4.5), and agitated (150 rpm, 30 °C) with
- for up to 5 days. Cells were filtered and observed by scanning electron microscopy as described
- 77 below.

# 78 **2.2 Mineralization Recovery Time Course**

- 79 In order to test the hypothesis that mineral-encrusted S-layers are shed and regenerated, a time course
- 80 of recovery was followed after Fe-mineralization as follows: *Lysinibacillus* sp. TchIII 20n38 cells
- 81 were grown to mid-exponential growth phase ( $OD_{600nm} = 0.3$ ) in LB (30 °C, 180 rpm). The culture
- 82 medium was then removed and the cells washed in MilliQ-H<sub>2</sub>O by gentle centrifugation ( $2600 \times g$ ,
- 15 min, room temperature). The cells were then resuspended to an equivalent cell density in either a mineralization solution (10 mM NaH<sub>2</sub>PO<sub>4</sub>, 10 mM FeSO<sub>4</sub>, pH=4.5), or a nutrient-free buffered
- 84 mineralization solution (10 mM NaH<sub>2</sub>PO<sub>4</sub>, 10 mM FeSO<sub>4</sub>, pH=4.5), or a nutrient-free buffered 85 solution at the same pH (10 mM NaH<sub>2</sub>PO<sub>4</sub>, pH=4.5), and agitated (30 °C, 150 rpm) for 16 h. The
- mineralization solution was then removed and the cells washed in MilliQ-H<sub>2</sub>O by gentle
- $^{12}$  centrifugation (2600 x g, 15 min, room temperature), and replaced by growth medium (LB, or
- 88 labelled growth medium).
- 89 For NanoSIMS experiments, LB was replaced with a defined growth medium approximating LB but
- 90 containing either <sup>14</sup>N or <sup>15</sup>N (Celtone<sup>®</sup>, Cambridge Isotopes, USA). While specified for use with
- bacteria for isotope analyses, this media was found to contain inhibitory concentrations of trace
- 92 metals, which we were able to remove by precipitation by addition of an excess of buffered
- 93 phosphate (10 mM) over 24 h with agitation at 150rpm, followed by filtration using a 0.22 μm filter.
- 94 This metal-depleted medium containing 5 g/L Celtone<sup>®</sup> was completed with 5 g/L acetate as a C-
- source, and basal salts (15 mM ammonium sulfate, 0.2 mM MgSO<sub>4</sub>,17.6 mM KH<sub>2</sub>PO<sub>4</sub>, 32.7 mM
- 96 NaH<sub>2</sub>PO<sub>4</sub>), as determined by our preliminary optimized experiments with *Lysinibacillus* sp. TchIII
- 20n38. In addition, cultures grown in the presence of <sup>14</sup>N rather than<sup>15</sup>N rapidly ceased vegetative
   growth and sporulated. Therefore instead of a standard labelling medium composed of both <sup>14</sup>N and
- <sup>98</sup> growth and sporulated. Therefore instead of a standard labeling medium composed of both N and <sup>99</sup> <sup>15</sup>N, a 100% <sup>15</sup>N-labeled medium, was used to follow the time course of recovery after Fe-exposure.
- An additional culture was resuspended in a  $100\%^{14}$ N medium and immediately sampled as a
- baseline control for N isotope abundances. Cultures were incubated in the <sup>15</sup>N-labeled medium at
- 102 30 °C with agitation (180 rpm) over a time course of recovery, for both mineralized (M) and non-
- 103 mineralized (NM) cultures. Aliquots were removed immediately after addition of growth medium
- 104 (T0), and then every 24 h (1d, 2d). At each time point, approximately 20mL aliquots were removed
- 105 for optical density measurements, optical microscopy verification of cell morphology, and filtered for
- 106 scanning electron microscopy observations and NanoSIMS analyses as described below. Abiotic
- 107 (non-inoculated) controls were used for comparison to distinguish mineralization due to the presence
- 108 of *Lysinibacillus* sp. TchIII 20n38 cells.

# 109 2.3 Scanning Electron Microscopy

- 110 Aliquots of bacterial cultures as well as abiotic controls (not inoculated with *Lysinibacillus* sp. TchIII
- 111 20n38 cells) were filtered through a 0.2 µm GTTP isopore polycarbonate filters using a Swinnex
- filter holder (Merck Millipore, Darmstadt, Germany). Filters were then air-dried, mounted on
- aluminum supports with carbon tape, and coated with carbon (7-8 nm thickness), gold (7 nm
- thickness), or platinum (5 nm thickness). Scanning electron microscopy (SEM) observations were
- performed using two different instruments; a Hitachi SU 3500 SEM installed at the electron
- microscopy platform of the Muséum National d'Histoire Naturelle (Paris, France), and a Zeiss Ultra
- 117 55 field emission gun SEM equipped with a Brucker EDS QUANTAX detector (Brucker
- 118 Corporation, Houston, TX, USA) installed at IMPMC (Sorbonne Université, Paris, France). For
- 119 observations using the Hitachi SU 3500 instrument observations were made in secondary electron
- 120 mode with an acceleration voltage of 15 kV. SEM-FEG images were acquired in secondary electron
- 121 mode using with the Zeiss Ultra 55 instrument with an in column detector (InLens) at 2 kV to 5 kV
- 122 and a working distance of 3 mm. Energy dispersive X-ray spectroscopy (EDX) analyses were
- 123 performed at 15 kV and a working distance of 7.5 mm after calibration with reference copper.

#### 124 2.4 Nano Secondary Ion Mass Spectrometry (NanoSIMS)

125 NanoSIMS sample preparations followed the protocol of (Miot et al., 2015). Briefly, aliquots of

bacterial cultures sampled from labeled and unlabeled media filtered through 0.2 μm GTTP isopore

- 127 polycarbonate filters previously Au-coated (20 nm thickness) using a Swinnex filter holder (Merck
- 128 Millipore, Darmstadt, Germany). Quantitative ion images were recorded by the NanoSIMS50
- 129 (Cameca, Gennevilliers, France) installed at the National Museum of Natural History of Paris,
- France. All measurements were performed using the same analytical conditions. A Cs + primary beam of 0.8 pA scanned an area of 20  $\mu$ m × 20  $\mu$ m, divided into 256 pixels × 256 pixels, with a
- counting time of 1 ms per pixel. Secondary ion images of  ${}^{31}P^{16}O_{-}$ ,  ${}^{12}C^{14}N_{-}$ , and  ${}^{12}C^{15}N_{-}$  were
- recorded. The mass resolution power was adjusted to 9000 to resolve isobaric interferences at mass
- 134 27 such as  ${}^{13}C^{14}N$  or  ${}^{11}B^{16}O$  from  ${}^{12}C^{15}N$ -. Before any analysis, the surface of each sample was pre-
- sputtered during 5 min with a 80 pA Cs- primary ion beam over 30  $\mu$ m × 30  $\mu$ m to eliminate the
- 136 contamination of the surface, and reached the stable state of sputtering (Thomen et al., 2014).
- 137 Instrument stability was verified throughout the session using a type 3 kerogen standard. NanoSIMS
- data were then processed using the IMAGE software (L. Nittler, Carnegie Institution for Science,
- 139 Washington, DC, USA).

#### 140 2.5 Statistical Analyses

141 The preference of this strain of *Lysinibacillus* for <sup>15</sup>N to maintain vegetative growth eliminated the

- 142 possibility of using a <sup>14</sup>N control throughout the time course of recovery. In order to automatically
- 143 remove the random noise from all the  ${}^{12}C^{14}N$  and  ${}^{12}C^{15}N$  NanoSIMS images, we defined  ${}^{12}C^{14}N$ -

and  ${}^{12}C^{15}N$ - independent thresholds based on their respective distribution for the images of samples

- 145 that were resuspended in the <sup>14</sup>N-labelled medium. Each elemental distribution was fitted by two
- Gaussian components (R-package mixtools, (Benaglia et al., 2009)). We define threshold as the mean
- of the 97.5th percentile of the first Gaussian component (noise) and the 2.5th percentile of the second
- 148 one (signal).
- 149 For each image, the denoised dataset further used for statistical analyses is composed only of pixels
- 150 with both  ${}^{12}C^{14}N$  and  ${}^{12}C^{15}N$  values above the respective thresholds. The isotope abundance  ${}^{12}C^{15}N$ -
- 151  $/({}^{12}C^{14}N + {}^{12}C^{15}N -)$  of this dataset (hereafter named processed  ${}^{15}N/({}^{14}N + {}^{15}N)$  ratio) was used to
- 152 follow the kinetics of incorporation of N (as part of protein production) by non-mineralized and
- 153 mineralized bacteria. For each image, the distribution of processed  ${}^{15}N/({}^{14}N + {}^{15}N)$  ratio was fitted
- using Gaussian mixture modeling (R-package mclust, (Scrucca et al., 2016)) in order to infer
- subpopulations of pixels. The best univariate model, composed of k Gaussian components with either
- equal or unequal variance, was selected based on Bayesian Information Criterion. The mixing
- 157 proportions for the components represent the proportions of the *k* subpopulations of pixels.
- 158 Cluster analysis was performed in order to group the samples according to their subpopulation
- 159 composition. Each image was described by a vector of 10 values, each corresponding to the sum of
- 160 the mixing proportions for the Gaussian components whose mean falls in a given  ${}^{15}N/({}^{14}N+{}^{15}N)$  ratio
- 161 interval (10 intervals of size 0.1 each). An image-to-image distance matrix generated by computing
- 162 Bray-Curtis dissimilarity index between all the pairs of vectors was used for hierarchical
- agglomerative clustering of images (unweighted pair group method with arithmetic mean linkage).
- 164 All analyses were conducted in R version 3.2.3 (R Core team, 2015).

#### 165 **3 Results**

#### 166 **3.1** *Lysinibacillus* sp. S-layers Become Encrusted After Exposure to Iron

- 167 *Lysinibacillus* sp. TchIII 20n38 was isolated from soils contaminated by radionuclides and metals,
- resulting in a moderately acidic pH (5.5) (Chapon et al., 2012). Our work with this strain, like other
- 169 isolates from the same site, has shown that it is resistant to a range of heavy metals and radionuclides
- 170 (article in preparation). To determine the mechanisms of this metallotolerance, we exposed cells to a
- 171 Fe-rich solution at acidic pH in the absence of the preferred carbon source for this strain, acetate,
- while maintaining high levels of phosphate required by *Lysinibacillus* sp.. Non-metabolic metal-
- 173 tolerance mechanisms are favored under these conditions.
- 174 After exposure, Fe-minerals were observed to form on the surface of *Lysinibacillus* sp. cells leading
- 175 to complete encrustation of the cells over time (see Fig. 1). EDX analyses of mineral-encrusted cells
- 176 confirmed the composition as a Fe-phosphate (see Supplementary Figure 1). Some abiotically formed
- 177 Fe-phosphates were also observed, which were easily distinguishable from mineralized S-layers as
- aggregates of larger spherically-shaped minerals not associated with cells, and matching the types of
- 179 minerals observed in non-inoculated controls.
- 180 Cells were fully mineral encrusted after 16 h of exposure to the Fe-rich solution, whether cells were
- 181 exposed in mid-exponential, late-exponential, or stationary growth phase ( $OD_{600nm} = 0.3, 0.6, and$
- 182 1.0, respectively). Longer exposures (20 h and 41 h) did not alter the extent mineralization.
- 183 Attempts to confirm whether the mineralization observed on Lysinibacillus sp. TchIII 20n38 cells
- 184 was due to a completely non-metabolic process, or whether active metabolism by living cells was
- necessary for S-layer mineralization were limited due to an inability to obtain dead cells without
- 186 damaging the S-layer containing cell envelope. This despite multiple trials employing various
- 187 antibiotics targeting non-cell envelope structures, and testing them over a large range of
- 188 concentrations and durations [tetracycline (10-2000  $\mu$ g/mL for 1 h-5 d in LB, buffer, or MilliQ-H<sub>2</sub>O)
- ofloxacin (10-500  $\mu$ g/mL), and heat treatments (up to 55°C)]. The fact that *Lysinibacillus* sp. TchIII
- 190 20n38 cells grow optimally as heterotrophs without added metals suggests that the role of any
- 191 metabolic processes in S-layer mineralization was secondary to non-metabolic processes.

# 192 **3.2** Physiological State of Cells Depends on Times of Mineralization and Recovery

- 193 Replacement of mineral-encrusted cells into a rich growth medium demonstrated that Lysinibacillus
- sp. TchIII 20n38 cells were able to resume proper cell division after complete Fe-mineral
- encrustation. The exposure of mid-exponential growth phase cells ( $OD_{600nm} = 0.3$ ) to the Fe-rich
- solution for 16 h followed by recovery of the cultures in LB showed that cells resumed normal cell
- division (see Supplementary Figure 2). However, the physiological state of the cells during metal
- exposure affected the ability of cells to recover after mineral encrustation, resulting in various
- inhibitions of normal vegetative cell growth and division. For mid-exponential growth phase cells, exposures longer than 16 h to the Fe-rich solution resulted in the death of mineralized cells, and the
- exposures longer than 16 h to the Fe-rich solution resulted in the death of mineralized cells, and the formation of filaments by the small minority of cells observed without mineral-encrustation. In the
- 201 formation of maments by the small minority of cells observed without mineral-encrustation. In the 202 case of both late-exponential growth phase and stationary phase cultures, even 16 hour-long
- 202 case of both late-exponential growth phase and stationary phase cultures, even 16 hour-ic
- 203 exposures to Fe resulted in sporulation and/or cell death.

# 204 3.3 Mineral-Encrusted S-layers Can Be Shed

- 205 SEM observations were made of mid-exponential growth phase *Lysinibacillus* sp. TchIII 20n38 cells
- after a 16 h exposure to the Fe-rich solution. Mineralized S-layers devoid of a cell were observed,
- 207 often with the cells located beside these empty mineralized S-layer shell (see Fig. 2 panel A, left and

right images). After incubation in LB for up to 5 d following Fe-mineralization, cells were seen
 exiting mineralized S-laver cells, with cell division septa visible (see Fig. 2 panel B, top and bottom)

images), in concordance with increases in the optical density of the cultures (see Supplementary

211 Figure 2).

# 212 **3.4** *Lysinibacillus* sp. Cell Morphology Changes During S-layer Shedding

The cell morphology in both mineralized and non-mineralized cultures was altered over the time course of recovery (see Fig. 3). In non-mineralized cultures which were kept at the same pH as the

mineralized cultures but in the absence of iron, cells gradually shrank in size and became ovoid in

shape over two days of incubation in the metal-depleted Celtone® medium. While cell death was

217 minimal for non-mineralized cells, dead cells were easily distinguishable due to both their high

- $^{12}C^{14}N$  counts due to lack of  $^{15}N$  incorporation, and their sustained rod shape (see image of non-
- 219 mineralized cells 1 d in Fig. 4). Non-mineralized cells also formed intracellular polyphosphate
- granules between 1 d and 2 d of incubation, as evidenced by analyses of  ${}^{31}P^{16}O$  counts (see Fig. 4).
- 221 Mineralized cultures showed little change over the first day of incubation. On the second day,

however, cells were observed outside of their mineralized S-layer shells, with biofilm formation

evidenced as a mucoid phenotype (see Fig. 3) together with increases in optical density of the

cultures (see Supplementary Figure 3). SDS-PAGE and mass spectrometry confirmed the presence of

S-layer glycoproteins throughout the time course of recovery (see Supplementary Figures 4 and 5).

Shed, mineralized S-layers maintained the elongated rod shape of non-stressed *Lysinibacillus* sp. TchIII 20n38 cells and restricted  ${}^{31}P^{16}O$ - to their surfaces (see Fig. 4), likely within the Fe-phosphate

228 minerals analyzed by EDX in Figure 2. In comparison, newly emerged cells lacked surface

phosphates, some cells concentrating phosphate as intracellular granules (see Fig. 4). On the third

230 day of incubation, cells in both mineralized and non-mineralized cultures began to sporulate.

# 231 **3.5** Sub-populations of Cells Co-exist During Time course of Recovery

In order to describe the process of S-layer regeneration after mineral encrustation, and to determine if 232 233 the S-layers were indeed regenerated, we followed the recovery of Lysinibacillus sp. TchIII 20n38 234 cells over time after Fe-mineral encrustation. Both SEM observations of cell morphology and NanoSIMS analyses of cell activity using incorporation of nitrogen, needed for the production of new 235 S-layer proteins. <sup>15</sup>N-incorporation is an effective marker, as S-layer proteins are one of the most 236 abundant cellular proteins, and roughly 20% of total protein synthesis can be dedicated to their 237 production (Sleytr et al., 2007). The <sup>15</sup>N incorporation over time was determined using 238  $^{12}C^{15}N/(^{12}C^{15}N+^{12}C^{14}N)$  and statistical analyses of the  $^{15}N/(^{14}N+^{15}N)$  were then performed (see 239 example of sample M T0 rep2 in Fig. 5, Supplementary Figure 6). In order to account for 240 241 differences in cell morphology over time and between mineralized and non-mineralized samples, statistical analyses were performed using all pixels above the established threshold from each image. 242 At least three processed  ${}^{15}N/({}^{14}N + {}^{15}N)$  images were analyzed per sample time point for samples 243 incubated in medium containing <sup>15</sup>N (two images each for natural abundance controls), with 10 to 244 <60 cells per image depending on the physiological state of the cells over the time course. 245 Subpopulations of pixels were identified for each sample according to the distribution of processed 246  $^{15}N/(^{14}N + ^{15}N)$  ratio (see Fig. 5). Figure 5 shows that at the start of the time course (T0), the 247 mineralized cells had  ${}^{15}N/({}^{14}N + {}^{15}N)$  ratios below 0.5 (panel A), with most cells having a ratio near 248 0.3 (panel B). Pixel subpopulations clustered around cells (panel C), with most mineralized cells and 249 abiotic mineralization retaining a small amount of <sup>15</sup>N (panel D) during the brief exposure prior to 250 washing the cells in the first few minutes of the experiment. 251

252 Clustering of the NanoSIMS images according to their distribution of processed  ${}^{15}N/({}^{14}N + {}^{15}N)$  ratio

showed that the samples tended to cluster over the experimental time course into three different groups; "natural abundance" ( $^{14}$ N) T0 controls, low  $^{15}$ N incorporation (NM T0, M T0, M 1d), and

significant <sup>15</sup>N incorporation (NM 1d, NM 2d, M 2d) (see Fig. 6). The fact that controls for the

natural abundance (NM no incubation, M no incubation) grouped separately from the T0 samples

shows that even brief exposure to the <sup>15</sup>N-labelled medium during cell resuspension had an effect on

the isotopic composition of the cells. The T0 samples were therefore used as the baseline of

comparison for all later time-points. The two remaining groups were composed of cells cultivated in

- <sup>15</sup>N-labelled medium, grouped by whether or not cell division had restarted. Low <sup>15</sup>N-incorporation
- samples (M\_T0, NM\_T0, M\_1d) corresponded to cells not yet showing evidence of cell division,
   whereas samples with significant <sup>15</sup>N incorporation (NM 1d, NM 2d, M 2d) showed clear evidence
- whereas samples with significant <sup>15</sup>N incorporation (NM\_1d, NM\_2d, M\_2d) showed clear evidence of active cell division when observed by SEM (see Fig. 3, black arrowheads) and measurements of
- 264 optical density (see Supplementary Figure 3).

265 Variations between replicate images were minimal for all samples with the exception of both

266 mineralized and non-mineralized 2 d samples, as seen in the heat map representation. The weighted

distribution of subpopulations remained low for all samples prior to restart of cell division (see Fig. 6

and Supplementary Figure 6, samples M\_T0, M\_1d, NM\_T0). The increase in pixel subpopulations

269 for samples showing signs of recovery and cell division (samples NM\_1d, NM\_2d, M 2d) reflects

270 the heterogeneity of cell recovery. Heterogeneity in <sup>15</sup>N incorporation was highest for cells

271 immediately after S-layer shedding (M\_2d samples), which is likely a reflection of natural variations

in the capacity of *Lysinibacillus* sp. TchIII 20n38 cells to respond to stress.

# 273 **3.6** S-layers are Regenerated Within Two Days of Fe-Mineral Encrustation

Some cell-free mineralized S-layer shells were observed by SEM immediately after 16 h exposure to the Fe-rich solution (see Fig. 3). However, the morphology of the cells remained generally unchanged through 1 d of incubation in the defined medium. At 2 d of incubation in this medium, SEM observations clearly show a majority of non-mineralized cells alongside the remaining, cellfree, mineralized S-layer shells. Analyses of <sup>15</sup>N uptake using NanoSIMS confirmed this timing, showing no significant <sup>15</sup>N incorporation for mineralized samples prior to 2 d of recovery in labelled medium, compared to T0 control aliquots that were removed immediately after resuspension of cells

in the labelled medium (see Fig. 6). The S-layers of mineralized samples remained at a relatively steady  ${}^{15}N/({}^{14}N + {}^{15}N)$  ratio (see green regions in Supplementary Figure 6, mineralized samples)

compared to the emerging cells, showing that mineralized S-layers did not incorporate <sup>15</sup>N during the

cell division processes giving rise to the emerging cells. This is coherent with S-layer shedding and complete regeneration. S-layer presence before and after mineralization and shedding was confirmed

286 by SDS-PAGE and mass spectrometry (see Supplementary Figure 4 and 5).

# 287 **4 Discussion**

# 4.1 Lysinibacillus sp. TchIII 20n38 Is Highly Adapted for Survival Under Stress Conditions

290 The bacterial isolate used in this study, *Lysinibacillus* sp. TchIII 20n38, is a metallotolerant gram-

291 positive bacterium (*article in preparation*), possessing a cell envelope composed of the plasma

membrane surrounded by a thick layer of peptidoglycan capped by an S-layer forming an ordered

structure at the cell surface. This flexible cage-like structure is in direct contact with the surrounding

294 environment, and thus provides the primary protective element against potentially toxic

environmental concentrations of heavy metals or radionuclides. *Lysinibacillus* sp. TchIII 20n38

demonstrated many adaptive mechanisms to the stresses induced by nutrient limitation and/or the

- 297 presence of iron, as might be expected for a bacterium isolated from a radioactive waste disposal site.
- 298 These included the accumulation and enlargement of intracellular polyphosphate granules and
- sporulation under phosphate-limiting conditions, cell size reduction and morphology alterations from
- 300 rod-shaped to ovoid cells after metal stress, as well as reductions in biofilm after exposure to either
- iron or acidic pH and augmentation in biofilm after a return to neutral pH in the absence of additional
- iron input. Polyphosphate accumulation is a common mechanism used by bacteria, including
- 303 *Lysinibacillus sphaericus*, in response to nutrient stress (depletion of amino acids), and prior to
- 304 sporulation (Shi et al., 2015; Tocheva et al., 2013).

# **4.2** S-layer Shedding Mechanism in *Lysinibacillus* as a Response to Metal Stress

306 S-layers from a variety of prokaryotes are known to induce mineral formation. The S-layers of

- 307 cyanobacteria are able to nucleate selenite and strontium (Schultze-Lam and Beveridge, 1994), while
- 308 the S-layers of thermophilic archaea can form amorphous Fe-phosphate minerals in the quasi-309 periplasmic space between the S-layer and the underlying lipid membrane (Kish et al., 2016). Diverse
- periplasmic space between the S-layer and the underlying lipid membrane (Kish et al., 2016). Divers
- *Lysinibacillus* sp. have been observed to precipitate minerals on their cell surfaces, including Uphosphates (Merroun et al., 2005; Mondani et al., 2011) and calcium carbonate (Kaur and Mukherjee,
- 2013). Here, we show that *Lysinibacillus* sp. TchIII 20n38 cells become encrusted with Fe-minerals
- after exposure to high concentrations of iron under mildly acidic conditions. Mineral precipitation by
- the cell surfaces, including S-layers, prevent damages to cells including oxidative stress, enzyme
- deactivation, protein denaturation, and membrane disruption (Lemire et al., 2013). Iron precipitation
- nucleated by the S-layer proteins prevents an overproduction of free radicals in the cytosol due to
- 317 Fenton chemistry.
- 318 While mineral formation on S-layers is known, the mechanisms for removing such barriers to
- exchange with the surrounding environment are not as well understood. Mechanisms identified to
- date have described partial removal of cell envelope components after metal interactions, particularly
- 321 membrane vesicle formation (Kish et al., 2016; McBroom and Kuehn, 2007; Shao et al., 2014).
- Partial shedding of S-layer fragments has also been observed, both for mineralized cyanobacterial S-
- layers (Schultze-Lam and Beveridge, 1994; Schultze-Lam et al., 1992) and non-mineralized S-layers
- for stationary phase bacteria likely as part of cell wall turnover (Luckevich and Beveridge, 1989).
   During the course of normal cell growth in the closely related *Lysinibacillus sphaericus*, bands of S-
- 325 Juring the course of normal cell growth in the closely related *Lysinibactitus sphaericus*, bands of S 326 laver monomer insertion form on cell surfaces, and in the course of cell division new S-laver
- 327 monomers are only inserted at the newly-formed the poles (Howard et al., 1982).

328 Here we show that *Lysinibacillus* sp. cells were able to recover normal growth after mineral

- 329 encrustation through a shedding of mineralized S-layers, followed by S-layer regeneration. To our
- knowledge, this is the first report of complete S-layer shedding and regeneration. S-layer shedding
- required an additional 24h before cells returned to normal cell division compared to non-mineralized
- cells, as shown by the clustering of <sup>15</sup>N uptake by 2 d mineralized cells with 1 d non-mineralized
- samples as measured by NanoSIMS over a time course after exposure to iron. Uptake of <sup>15</sup>N also
- illustrated that despite extensive mineralization, cells retained active metabolism. The continued
- 335 presence of shed, mineralized S-layers composed of <sup>14</sup>N-based proteins alongside cells bearing newly 336 regenerated <sup>15</sup>N-bearing S-layers resulted in the heterogeneity in <sup>15</sup>N-incorporation for the M 2d
- 337 samples (see Fig. 6). S-layer shedding activity was limited to cells in mid-exponential growth phase,
- 338 providing an advantage over cells in stationary growth phase in Fe-rich conditions.

# **4.3 Potential Applications of S-layer Regeneration in Bioremediation**

- 340 S-layers have a strong potential for use biotechnology, due to the combination their capacities for
- both self-assembly and metal binding. S-layers have been used as templates to nucleate the
- 342 fabrication of metal nanoparticles (Pollmann and Matys, 2007) that can be applied as biocatalysts
- 343 (Creamer et al., 2007) or exploited for their unique magnetic properties (Bartolomé et al., 2012). S-
- 344 layers assembled in vitro have been proposed as bio-filters for bioremediation technologies
- 345 (Pollmann et al., 2006).
- 346 The total S-layer shedding and regeneration shown here after mineral encrustation opens further
- 347 possibilities for a self-renewing bioremediation system, employing soil bacteria already tolerant to
- 348 metals and radionuclides.

### **349 5 Conflict of Interest**

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

### 352 6 Author Contributions

- 353 AK and LR contributed conception and design of the study; AC performed the experiments, LR
- 354 performed NanoSIMS measurements, CL and SZ performed mass spectrometry analyses; ED
- 355 performed the statistical analysis of NanoSIMS data; AK wrote the first draft of the manuscript; AC
- 356 wrote sections of the manuscript. All authors contributed to manuscript revision, read and approved
- the submitted version.

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# 373 9 References

- Ahmed, I., Yokota, A., Yamazoe, A., and Fujiwara, T. (2007). Proposal of Lysinibacillus
- boronitolerans gen. nov. sp. nov., and transfer of *Bacillus fusiformis* to *Lysinibacillus fusiformis*
- 376 comb. nov. and Bacillus sphaericus to Lysinibacillus sphaericus comb. nov. Int J Syst Evol Microbiol
- 377 57, 1117–1125. doi:10.1099/ijs.0.63867-0.

- 378 Bartolomé, J., Bartolomé, F., Bartolomé, F., García, L. M., García, L. M., et al. (2012).
- Strong paramagnetism of gold nanoparticles deposited on a *Sulfolobus acidocaldarius* S layer. *Phys*
- 380 *Rev Lett* 109, 247203. doi:10.1103/PhysRevLett.109.247203.
- Benaglia, T., Chauveau, D., Hunter, D. R., and Young, D. (2009). mixtools: An RPackage for Analyzing Finite Mixture Models. *J Stat Softw* 32, 1–29. doi:10.18637/jss.v032.i06.
- 383 Chapon, V., Piette, L., Vesvres, M.-H., Coppin, F., Marrec, C. L., Christen, R., et al. (2012).
- 384 Microbial diversity in contaminated soils along the T22 trench of the Chernobyl experimental
- 385 platform. *Appl Geochem* 27, 1375–1383. doi:10.1016/j.apgeochem.2011.08.011.
- 386 Creamer, N. J., Mikheenko, I. P., Yong, P., Deplanche, K., Sanyahumbi, D., Wood, J., et al. (2007).
- 387 Novel supported Pd hydrogenation bionanocatalyst for hybrid homogeneous/heterogeneous catalysis.
- 388 *Catal Today* 128, 80–87. doi:10.1016/j.cattod.2007.04.014.
- Howard, L. V., Dalton, D. D., and McCoubrey, W. K. (1982). Expansion of the tetragonally arrayed
  cell wall protein layer during growth of *Bacillus sphaericus*. *J Bacteriol* 149, 748–757.
- Kaur, D. N., and Mukherjee, A. (2013). Biomineralization of calcium carbonate polymorphs by the bacterial strains isolated from calcareous sites. *J Microbil. Biotechnol* 23, 707–714.
- 393 doi:10.4014/jmb.1212.11087.
- Kish, A., Miot, J., Lombard, C., Guigner, J.-M., Bernard, S., Zirah, S., et al. (2016). Preservation of archaeal surface layer structure during mineralization. *Sci Rep* 6, 26152–10. doi:10.1038/srep26152.
- Konhauser, K. O., Schultze-Lam, S., Ferris, F. G., Fyfe, W. S., Longstaffe, F. J., and Beveridge, T. J.
  (1994). Mineral precipitation by epilithic biofilms in the Speed river, Ontario, Canada. *Appl Environ Microbiol* 60, 549–553.
- Lemire, J. A., Harrison, J. J., and Turner, R. J. (2013). Antimicrobial activity of metals: mechanisms,
  molecular targets and applications. *Nat Rev Micro* 11, 371–384. doi:10.1038/nrmicro3028.
- Luckevich, M. D., and Beveridge, T. J. (1989). Characterization of a dynamic S layer on *Bacillus thuringiensis*. *J Bacteriol* 171, 6656–6667.
- McBroom, A. J., and Kuehn, M. J. (2007). Release of outer membrane vesicles by Gram-negative
  bacteria is a novel envelope stress response. *Mol Microbiol* 63, 545–558. doi:10.1111/j.13652958.2006.05522.x.
- 406 Merroun, M. L., Raff, J., Rossberg, A., Hennig, C., Reich, T., and Selenska-Pobell, S. (2005).
- 407 Complexation of uranium by cells and S-layer sheets of *Bacillus sphaericus* JG-A12. *Appl Environ*
- 408 *Microbiol* 71, 5532–5543. doi:10.1128/AEM.71.9.5532-5543.2005.
- 409 Miot, J., Remusat, L., Duprat, E., Gonzalez, A., Pont, S., and Poinsot, M. (2015). Fe
- biomineralization mirrors individual metabolic activity in a nitrate-dependent Fe(II)-oxidizer. *Front Microbiol* 6, 879. doi:10.3389/fmicb.2015.00879.
- 412 Mondani, L., Benzerara, K., Carrière, M., Christen, R., Mamindy-Pajany, Y., Février, L., et al.
- 413 (2011). Influence of uranium on bacterial communities: a comparison of natural uranium-rich soils
- 414 with controls. *PLoS ONE* 6, e25771. doi:10.1371/journal.pone.0025771.

- 415 Phoenix, V. R., Adams, D. G., and Konhauser, K. O. (2000). Cyanobacterial viability during
- 416 hydrothermal biomineralisation. *Chem Geol* 169, 329–338.
- 417 Pollmann, K., and Matys, S. (2007). Construction of an S-layer protein exhibiting modified self-
- 418 assembling properties and enhanced metal binding capacities. *Appl Microbiol Biotechnol* 75, 1079–
  419 1085. doi:10.1007/s00253-007-0937-5.
- Pollmann, K., Raff, J., Merroun, M., Fahmy, K., and Selenska-Pobell, S. (2006). Metal binding by
  bacteria from uranium mining waste piles and its technological applications. *Biotech Adv* 24, 58–68.
  doi:10.1016/j.biotechadv.2005.06.002.
- 423 R Core Team (2015). R: A language and environment for statistical computing. R Foundation for 424 Statistical Computing, Vienna, Austria. Available at https://www.R-project.org/
- 425 Rodriguez-Freire, L., Avasarala, S., Ali, A.-M. S., Agnew, D., Hoover, J. H., Artyushkova, K., et al.
- 426 (2016). Post Gold King Mine Spill Investigation of Metal Stability in Water and Sediments of the
- 427 Animas River Watershed. *Environ Sci Technol.* 50, 11539–11548. doi:10.1021/acs.est.6b03092.
- Schultze-Lam, S., and Beveridge, T. J. (1994). Nucleation of celestite and strontianite on a
  cyanobacterial s-layer. *Appl Environ Microbiol* 60, 447–453.
- Schultze-Lam, S., Harauz, G., and Beveridge, T. J. (1992). Participation of a cyanobacterial S layer
  in fine-grain mineral formation. *J Bacteriol* 174, 7971–7981.
- 432 Scrucca, L., Fop, M., Murphy, T. B., and Raftery, A. E. (2016). mclust 5: Clustering, Classification
  433 and Density Estimation Using Gaussian Finite Mixture Models. *R J* 8, 289–317.
- Shao, P. P., Comolli, L. R., and Bernier-Latmani, R. (2014). Membrane vesicles as a novel strategy
  for shedding encrusted cell surfaces. *Minerals* 4, 74–88. doi:10.3390/min4010074.
- 436 Shi, T., Ge, Y., Zhao, N., Hu, X., and Yuan, Z. (2015). Polyphosphate kinase of *Lysinibacillus*
- 437 *sphaericus* and its effects on accumulation of polyphosphate and bacterial growth. *Microbiol Res*438 172, 41–47. doi:10.1016/j.micres.2014.12.002.
- Sleytr, U. B., Egelseer, E. M., Ilk, N., Pum, D., and Schuster, B. (2007). S-Layers as a basic building
  block in a molecular construction kit. *FEBS J* 274, 323–334. doi:10.1111/j.1742-4658.2006.05606.x.
- Sleytr, U. B., Schuster, B., Egelseer, E. M., and Pum, D. (2014). S-layers: principles and
  applications. *FEMS Microbiol Rev* 38, 823–864. doi:10.1111/1574-6976.12063.
- Sleytr, U. B., Schuster, B., Egelseer, E. M., Pum, D., Horejs, C. M., Tscheliessnig, R., et al. (2011).
  Nanobiotechnology with S-layer proteins as building blocks. *Prog Mol Biol Transl Sci* 103, 277–352.
  doi:10.1016/B978-0-12-415906-8.00003-0.
- Thomen, A., Robert, F., and Remusat, L. (2014). Determination of the nitrogen abundance in organic
   materials by NanoSIMS quantitative imaging. *J Anal At Spectrom* 29, 512–519.
- 448 doi:10.1039/c3ja50313e.

- 449 Tocheva, E. I., Dekas, A. E., McGlynn, S. E., Morris, D., Orphan, V. J., and Jensen, G. J. (2013).
- 450 Polyphosphate Storage during Sporulation in the Gram-Negative Bacterium Acetonema longum. J
- 451 Bacteriol 195, 3940–3946. doi:10.1128/JB.00712-13.

#### 452 **10** Supplementary Material

- 453 The Supplementary Material for this article can be found as a separate file.
- 454 **11 Figures**

A



B

- 455
- 456 Figure 1. Scanning electron microscopy images in secondary electron mode of *Lysinibacillus* sp.
- 457 **TchIII 20n38 cells (A)** grown under optimal conditions, and **(B)** after exposure to an iron-rich
- solution (10 mM NaH<sub>2</sub>PO<sub>4</sub>, 10 mM FeSO<sub>4</sub>, pH=4.5). Scanning electron microscopy images acquired
   in secondary electron mode.







469

470 Figure 3. Scanning electron microscopy images of Lysinibacillus sp. TchIII 20n38 cells

#### 471 (secondary electron mode) for mineralized and non-mineralized cells over a time course of recovery

#### 472 after Fe-exposure.





Figure 4. <sup>12</sup>C<sup>14</sup>N- and <sup>31</sup>P<sup>16</sup>O- NanoSIMS images for both non-mineralized and mineralized cells over a time course of recovery after Fe-exposure. 475

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**Figure 5. Statistical analysis of**  ${}^{15}N/({}^{14}N+{}^{15}N)$  **NanoSIMS image** for mineralized cells after 2 days of recovery (M\_2d\_rep16). (A) Processed  ${}^{15}N/({}^{14}N+{}^{15}N)$  ratio map. (B) Processed  ${}^{15}N/({}^{14}N+{}^{15}N)$ 477 478 ratio histogram (gray bars) and probability density (dark line) estimated by Gaussian mixture 479 modeling. (C) Map of the pixel subpopulations (see (D) for color code). (D) Decomposition of the distribution of processed  ${}^{15}N/({}^{14}N+{}^{15}N)$  ratios (labelled "raw data") into 4 Gaussian components of 480 481 unequal variance. The box boundaries indicate the first and third quartiles. The bold band inside the 482 box corresponds to the median value. The horizontal dashed lines that extend from the box 483 encompass the largest/smallest ratio values that fall within a distance of 1.5 times the box size from 484 485 the nearest box hinge. If any, outliers are shown as individual points. Each component of the Gaussian mixture is represented by a random sample (n=10000) from the corresponding normal 486

- 487 distribution. The box heights represent the proportions of pixel subpopulations (i.e. the mixing
- 488 proportions for the components). A unique color scale is used for all images, the subpopulation color
- 489 depending on the estimated mean of the Gaussian component.





Figure 6. Clustering of <sup>15</sup>N/(<sup>14</sup>N+<sup>15</sup>N) NanoSIMS images according to the distribution of pixel 491 subpopulations. Each row of the heatmap (drawn on the right side of the figure) shows proportion 492 (in %) of the pixel subpopulations for a single NanoSIMS image, and columns show the 493 494  $^{15}N/(^{14}N+^{15}N)$  ratio intervals. Each value corresponds to the sum of the mixing proportions for the 495 Gaussian components whose mean falls in the given interval. Null proportions are hidden. Color scale ranges between white (0 %) and black (100 %). Each image is labelled according to the culture 496 medium (either M or NM for Mineralized or Non-Mineralized, respectively), the <sup>15</sup>N incubation time, 497 and the replicate number (a NanoSIMS image numbering for a given sample). Each image is depicted 498 by a color code according to the <sup>15</sup>N incubation time of the sample (see color legend at bottom left). 499 The heatmap rows are ordered according to the hierarchical clustering of images illustrated on the left 500

- 501 side of the figure. Image labels ending with a star are further illustrated in Figure 6 and
- 502 Supplementary Figure 6.