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► To cite this version:

Solène Fastenackels, Michèle Mock, Jean-Nicolas Tournier, Pierre L Goossens. Early expression of capsule during Bacillus anthracis germination. Research in Microbiology, 2023, 174 (6), pp.104054. 10.1016/j.resmic.2023.104054 . hal-04077748

HAL Id: hal-04077748 https://hal.science/hal-04077748v1

Submitted on 21 Apr 2023 $\,$

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PII: S0923-2508(23)00029-3

DOI: https://doi.org/10.1016/j.resmic.2023.104054

Reference: RESMIC 104054

To appear in: Research in Microbiology

Received Date: 28 October 2022

Revised Date: 10 March 2023

Accepted Date: 15 March 2023

Please cite this article as: S. Fastenackels, M. Mock, J.-N. Tournier, P.L. Goossens, Early expression of capsule during *Bacillus anthracis* germination, *Research in Microbiologoy*, https://doi.org/10.1016/j.resmic.2023.104054.

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4	Solène Fastenackels ^{a,} Michèle Mock ^b , Jean-Nicolas Tournier ^c , Pierre L. Goossens ^{d*}
5	
6	
7	Affiliations :
8	^a Laboratory "Immune Microenvironment and Immunotherapy", INSERM U1135, Centre
9	d'Immunologie et des Maladies Infectieuses Paris (CIMI-Paris), Paris, France
10	^b Institut Pasteur, c/o Yersinia Unit, 26 rue du Docteur Roux, 75724 Paris cedex 15
11	^c IRBA, Div.DNRBC/D.2MI, 1, place Général Valérie André, 91223 Brétigny/Orge
12	^d Institut Pasteur, Yersinia Unit, 26 rue du Docteur Roux, 75724 Paris cedex 15
13	
14	
15	
16	
17	
18	
19	
20	
21	
22	
23	
24	^a solene.fastenackels@sorbonne-universite.fr
25	^b <u>michele.mock@pasteur.fr</u>
26	^c jntournier@gmail.com
27	^{d*} pierre.goossens@pasteur.fr Correspondance and reprints
28	

29 Abstract

30 Bacillus anthracis is a spore-forming bacterium that produces two major virulence 31 factors, a tripartite toxin with two enzymatic toxic activities and a pseudo-proteic capsule. 32 One of the main described functions of the poly-gamma-D-glutamate capsule is to enable B. 33 anthracis bacilli to escape phagocytosis. Thus, kinetics of expression of the capsule filaments 34 at the surface of the emerging bacillus during germination is an important step for the 35 protection of the nascent bacilli. In this study, through immunofluorescence and electron 36 microscopic approaches, we show the emergence of the capsule through a significant surface 37 of the exosporium in the vast majority of the germinating spores, with co-detection of BclA and capsular material. This suggests that, due to an early capsule expression, the extracellular 38 39 life of *B. anthracis* might occur earlier than previously thought, once germination is triggered. 40 This raises the prospect that an anti-capsular vaccine may play a protective role at the initial 41 stage of infection by opsonisation of the nascent encapsulated bacilli before their emergence 42 from the exosporium. 43

- 44
- 45 Keywords
- 46 Bacillus anthracis; spore; capsule; germination

47 **1. Introduction**

The virulence of *Bacillus anthracis*, the etiological agent of anthrax, relies predominantly on two main factors, a tripartite toxin with two enzymatic activities (lethal and edema factors, metalloprotease and adenylcyclase activities respectively) and a pseudoproteic capsule made of poly-gamma-D-Glutamic acid (PDGA)[1] The capsule protects the bacteria from phagocytosis, is poorly immunogenic [2] and potentially disruptive for the immune response [3, 4]; it enables the bacteria to bind to the liver endothelium [5] and plays a role in sequestration inside pulmonary capillaries at the terminal phase of the infection [6].

55 B. anthracis is a spore-forming bacterium. The spore is a specialised cell form that remains dormant and survives for many years, formed of concentric shells surrounding the 56 57 core that contains the bacterial genetic material: a membrane: a peptidoglycan layer - the 58 cortex: a protein multilayer -the coat: and the exosporium -the outermost structure of the 59 spore [7-9]. The exosporium is composed of a basal layer and an external hair-like nap 60 formed by a single collagen-like glycoprotein called BclA [10, 11]). Upon germination there 61 is an influx of water into the spore, swelling of the spore core, dismantling of the cortex and 62 coat leading to resumption of metabolic activity and cell growth [7-9].

Early expression of the toxin genes has been shown [12], as soon as 15 min after germination triggering for mRNA detection. This suggests a potential early role for them in subverting the host defense. Temporal expression of the capsule is also critical to enable protection of the nascent bacilli. How the capsule emerges from a germinating spore that is surrounded by the exosporium and the coat layers is thus of importance.

In this study, we addressed the kinetics of capsule expression during *B. anthracis* germination. We observed early detection of capsular material as soon as 15 min after germination triggering. Capsular material was detected in the majority of spores as emerging all around the external layers of the spore.

72 2. Materials and methods

73

74 2.1 Bacterial growth

75 The B. anthracis strains used were the encapsulated non-toxinogenic RPG1 strain 76 derived from the RPLC2 strain after transduction of the pXO2 of the ATCC 4229 strain [13], 77 the parental non-encapsulated RPLC2 strain [14] and the Δ -bclA RPG1-derived strain 78 (constructed as described previously [10]); the RPLC2 strain is derived from the Sterne strain 79 with one point mutation in the *lef* gene and two point mutations in the *cya* gene leading to 80 enzymatically inactive toxins. To synchronise spore germination, spores $(1 \times 10^7 \text{ mL} - 1)$ 81 were shortly incubated in BHI for 3 min at 37° [13]. This incubation step triggered germination in more than 90% of the spores, allowing the analysis of a homogeneous 82 83 germinating spore population, as germination was less efficiently triggered in the minimum 84 culture R-medium. After centrifugation (13,000 rpm, 2 min), the germinating spores were 85 resuspended in R-medium with 0.6% sodium bicarbonate [15] and incubated at 37 °C in a 5% 86 CO₂ atmosphere; this growth condition mimics the *in vivo* environment and ensures 87 coordinated *in vitro* expression of the capsule and the toxins. At various timepoints, bacteria 88 were collected, washed in Dulbecco's Phosphate Buffered Saline with MgCl₂ and CaCl₂ 89 (Sigma, ref D8662) and then analysed by immunofluorescence or ultrastructural studies (see 90 below). For the *in vivo* approach, $1 \ge 10^8$ germinated spores obtained after the BHI incubation step 91 92 were injected intraperiteonally in female outbred Swiss OF1 mice (weight, 22-24 g; Charles 93 River, L'Arbresle, France). One hour later, the animals were euthanasised, a peritoneal lavage 94 was performed with 10 mL cold Dulbecco's Phosphate Buffered Saline without MgCl₂ and 95 $CaCl_2$ (Sigma, ref D8537); the suspension of bacterial and peritoneal cells were then 96 centrifuged in PBS with MgCl₂ and CaCl₂ (1200rpm, 5 min) to pellet the majority of 97 peritoneal cells, the supernatant was then centrifuged (4000rpm, 8mn) to pellet the bacterial 98 suspension. After resuspension in PBS with MgCl₂ and CaCl₂, the bacteria were processed for 99 immuno-staining. Animals were housed in the Institut Pasteur animal facilities, licensed by 100 the French Ministry of Agriculture and in compliance with European regulations. All animal 101 experiments were conducted according to European Union guidelines 102 https://ec.europa.eu/environment/chemicals/lab_animals/3r/ key_resources/portals_en.htm 103 and were approved by the animal care and use committee at the Institut Pasteur. 104

105 2.2 Immunofluorescence and ultrastructural studies

The germinated spores were fixed in 4% paraformaldehyde for 15 min, deposited on poly-L-lysine-coated (PLL) glass coverslips as described previously [16]. In brief, coverslips were sequentially incubated for 1 hour at room temperature with appropriate dilutions of the primary antibodies and of the species-specific secondary fluorochrome conjugates in PBS 0.25% gelatin. Each incubation was followed by three washes of 10 minutes with PBS. In some experiments, the host cell nuclei were stained with 100 nM Topro-3 in the Vectashield mounting medium.

113 The exosporium protein BclA was detected with the mouse monoclonal antibody 114 4B7G12 developed in the laboratory, revealed by an Alexa488-labelled donkey anti-mouse 115 secondary polyclonal antiserum. For the exosporium of the Δ -*bclA* RPG1 strain, devoid of the 116 BclA protein, primary labelling was performed with a murine polyclonal anti-exosporium 117 antiserum raised against Δ -*bclA* spores developed in the laboratory.

118 The PDGA capsular material was detected with a polyclonal rabbit immune serum 119 revealed by an Alexa594-labelled goat anti-rabbit polyclonal antiserum as previously 120 described [17]. No cross reaction between the various antibodies was observed (data not 121 shown). Confocal microscopy was performed on a Leica SP5 confocal microscope using a 122 63x, oil immersion objective. Image deconvolution was performed as already described [18]. 123 Quantification of the spore populations exhibiting distinct patterns of BclA and 124 PGDA-labelling was performed at various times of the germination kinetics and the 125 vegetative cells were excluded (Table 1).

Transmission electron microscopy was performed on dendritic cells (DC) at 1h of infection with RPG1 and RPLC2 spores as previously described [19]. In brief, DC were infected with the spores and analysed after 30 min. Cells were fixed for 3 h in 3% glutaraldehyde in 0.1 M phosphate buffer and postfixed for 1 h with 1% OsO4 in 0.1 M cacodylate buffer. The pellet was embedded in epoxy resin and sectioned for transmission electron microscopy.

132 Scanning electron microscopy was performed on 45 min-germinated spores bound to 133 PLL-coated coverslips; after overnight fixation in 2.5 % glutaraldehyde in 0.1 M sodium 134 cacodylate buffer at 4 °C, and incubation with 1/100 diluted in BHI (brain heart infusion) 135 anti-PDGA specific rabbit polyclonal antibodies and anti-BclA mouse monoclonal antibodies 136 as described above and then incubated with a 15 nm immuno-gold conjugate (British Biocell 137 International) coupled with an anti-rabbit goat antibody, and of 10 nm immuno-gold 138 conjugate (British Biocell International) coupled with an anti-mouse goat antibody. Finally, a 139 negative staining was performed with phosphotungstic acid 2 % in water.

Of note, all image acquisitions were carried out on PLL-coated supports. This originated from the observation that carbon-coated grids preferentially bound spores that do not present capsular material at the surface in contrast to PLL-coated grids. Indeed, only 5 % of the spore population was positive for anti-capsular material labelling on carbon-coated grids versus 61 % for the PLL-coated grids, as assessed through immunofluorescence detection as described above. Such approach is thus critical to avoid misrepresentation of a particular subpopulation.

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147 **3. Results**

148 The presence of PDGA capsular material and the BclA protein - an abundant protein 149 of the spore surface - was detected through immuno-fluorescence labelling with specific 150 antibodies (Fig. 1&2); quantitative analysis of both fluorescence signal topographic 151 distribution was performed on the samples (Table 1 built on kinetics as shown in Fig. 1&2). 152 At 30 and 45 min (Fig. 1), many germinating spores of the encapsulated RPG1 strain 153 were labelled with both anti-capsular material and anti-BclA antibodies; co-detection of 154 capsular material and BclA was observed around the whole, or greater part of, spore surface, 155 in 67 % of the bacterial population at 30 min (forms 5+6 in Table 1). This proportion for the 156 forms 5+6 was still around 57 % at 60 min (Table 1); at this timepoint, 29 % of the bacterial 157 cells (form 8 in Table 1) were labelled as encapsulated bacteria that were devoid of anti-BclA 158 labelling. This population (form 8 in Table 1) increased, as expected, to reach 76 % of the 159 labelled bacteria at 120 min, with a few remnants of anti-BclA labelled exosporium; the 160 remaining bacterial cells were anti-BclA labelled spores that did not undergo the germination 161 step (Fig. 1 and form 1 in Table 1). Interestingly, a particular topographic distribution of fluorescence was observed in 8 % of the germinating spores at 30 min; capsular material was 162 163 observed as dots on some locations of the BclA layer (form 2 in Table 1).

Triple labelling (Topro-3 for DNA, and anti-BclA and anti-capsular material labelling) was performed at 45 min, at a time point when the majority of the germinating spores were labelled for BclA and capsule (Fig. 1, time point 45 min); all germinating spores harboured DNA and thus were not empty exosporium shells. Using mouse anti-exosporium polyclonal antibodies, similar co-detection of capsular material and the exosporium layer at 45 min was also observed with spores from a strain deleted of the *bclA* gene (Fig 1).

Raising the question of the timepoint of initial detection of the anti-capsular material
labelling, we observed that capsular material could be detected as early as 15 min of
incubation in R medium 0.6% sodium bicarbonate in a 5% CO₂ atmosphere (Fig 2).
Quantitatively, this fluorescence signal was detected in 60% of the germinating spores (forms
4+5+6 in Table1). No capsular material labelling was detected immediately after the 3mingermination step in BHI, while all spores were labelled with the anti-BclA antibody (Table 1
and Fig. 2).

To confirm that this phenomenon also occurs *in vivo*, the germinated spores were
inoculated in mice into the peritoneum cavity and bacterial growth allowed to develop *in vivo*;
the bacterial cells were recovered after 1h of growth and immunolabelled as above (Fig. 1,

180 uttermost right panel). Similar co-detection of capsular material and BclA around the bacterial181 surface was observed.

Analysis of the fluorescent images at 30 and 45 min suggested that the capsular material was detected outside the BclA layer (Fig. 1). To analyse this particular aspect in more detail, two complementary approaches were developed, confocal microscopy and image deconvolution (Fig. 3a top and bottom panels respectively). Both approaches strongly suggested the capsular material was localised outside the BclA layer. Image deconvolution also suggested that the capsular material was distributed as dots around the surface (Fig. 3a bottom panel).

189 To confirm this observation, an ultrastructural approach was followed, through 190 electron microscopy (Fig 3b and c). Transmission electron microscopy without labelling 191 showed deposition of electron dense material as dots around the most external surface of the 192 encapsulated germinating RPG1 spores inside the hair-like BclA layer (Fig 3b, left), while 193 this was not observed with the parental non-encapsulated RPLC2 strain (Fig 3b, right). 194 We then used a specific qualitative analysis through scanning electron microscopy 195 with labelling by immunogold co-detection of antibodies directed against capsular material 196 (15 nm beads) and BclA (10 nm beads). We were able to observe spores that presented small 197 packs of labelled capsular material that emerge in various locations of the labelled 198 exosporium (Fig. 3c).

,

199 4. Discussion

In this study, we addressed the kinetics of capsule expression and topographic distribution during germination of *B. anthracis* spores, using immunofluorescence and electron microscopy approaches. We show co-detection of both antigens peaking at 30-45 min after germination triggering.

Topographically, the capsular material was detected in the majority of the germinating spores around the whole surface of the exosporium. This strongly suggests that the external spore layers were fissured and the capsular material synthesised by the nascent bacilli emerged through these fissures and breaks. Indeed, dots of capsular material could be detected around the BclA layer in 8% of the germinating spores at 30 min (form 2 in Table 1). This phenomenon was also observed for *in vivo* infection.

210 Our data thus strongly suggest that *B. anthracis* behaves similarly to *B. atrophaeus* 211 and *B. subtilis*, where fissures and breaks have been observed in the coat during germination. 212 [20, 21]. Interestingly, John Ezzell reported a similar finding many years ago [22]; this 213 observation was exploited as a means for *B. anthracis* spore identification through this 214 phenomenon of 'spore encapsulation'. Of note, we could also observe bacterial cells where 215 the nascent bacillus emerged from the spore coat and exosporium layers only at one end, 216 leaving behind an empty shell. This was in accordance with previously published data 217 reporting that, during spore germination and outgrowth, the outgrowing cell escapes from its 218 exosporium shell by popping through a region of the exosporium that could not be stained by 219 antibodies to the spore specific alanine racemase [23].

Toxins and capsule are expressed in a coordinated manner, at the genetic and molecular levels [24, 25]. Our study shows that capsular material could be detected as soon as 15 min. PA mRNA has been reported as early as 15 min after germination, thus highlighting the precocity of toxin production [26]. Similarly, lethal and edema factors were detected in the plasma of mice as early as one hour after intranasal inoculation [27]. Two consequences of this early capsular material expression and distribution on the surface of the germinating spores may be envisioned.

(i) The lifestyle of *B. anthracis*, an extracellular pathogen, at this early stage is
considered to follow two different initial scenarios at the portal of entry into the infected host
- an extracellular or an intracellular step - the physiological importance of each one depending
on the route of entry. On one hand, during a cutaneous infection, *B. anthracis* spores
germinate very efficiently in the extracellular tissues [28].Furthermore, as reported in
inhalational and gastrointestinal infections, as soon as a lesion is present, efficient spore

233 germination occurs at the lesion site [29]. On the other hand, an intracellular pathway has 234 been described, playing a role in the crossing of the respiratory or intestinal epithelial barriers 235 during inhalational and gastrointestinal infections [30-32]; spores are engulfed and, through 236 trafficking, emerge from the cells on the other side of the epithelium. Our study thus suggests 237 that, in both scenarios, early expression of capsular material around the germinating spore 238 would help protect the emerging nascent bacilli at a crucial timepoint when *B. anthracis* is 239 susceptible to the host innate defenses, as it is converting from the resistant dormant spore to 240 the virulent toxinogenic encapsulated bacillus. Combined with early expression of toxins [26, 241 27] this would help subvert the host defenses.

242 (ii) a second outcome of such early capsular material expression on the germinating spores is

that they may become a target for a pre-existing anti-capsular vaccine immune response.

244 Current veterinary vaccines rely on live attenuated spores of the toxinogenic non-encapsulated

245 Sterne strain, while human vaccines are mainly based on the common toxin component

246 protective antigen (PA) [33]. However these PA-based vaccines for human use are protective

only in some specific animal models [34]. To enhance their efficacy, other components have

248 been added experimentally, especially for protection against inhalational anthrax. Addition of

249 inactivated spores as a source of spore antigens notably increases the efficiency of an anthrax

vaccine in the mouse model [14, 35]. Addition of spore or bacilli components, or more

251 specifically of capsular material, have been considered and tested [17, 36, 37]; in particular,

the group of Arthur Friedlander has further demonstrated the role of anti-capsule immunity in

vaccine development [38, 39]. Our data suggest that the approaches of adjoining capsular

254 material to the current anthrax human vaccines would thus be beneficial earlier than

255 considered; opsonisation of the nascent encapsulated bacilli, even before emergence from the

exosporium, would play a protective role at the initial stage of infection.

257 Acknowledgments

- 258
- 259 We would like to express our thanks to Tam Mignot for the deconvolution analysis, Evelyne
- 260 Tosi-Couture (Pasteur Institute) and Guy Brochier (IRBA) for their expertise in ultrastructural
- analysis, and Jean-Philippe Corre for his contributions and discussions.
- 262

263 **Conflict of interest**

264 All authors have no conflict of interest

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265 **References**

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365 Legends to figures

366

367 Figure 1: Co-detection of capsular material and exosporium during germination of B. 368 anthracis spores of the encapsulated RPG1 strain and the *bclA*-deleted derived strain (Δ -369 *bclA*): germinating spores were immuno-labelled 30 and 45 min after germination triggering 370 for 3 min in BHI and further growth in R-medium 0.6% Bicarbonate in 5% CO2 atmosphere. 371 Representative germinating spores are represented at higher magnification beside groups of 372 bacterial cells for each time points. Red fluorescence for anti-capsular material labelling, 373 green fluorescence for monoclonal anti-BclA or polyclonal anti-exosporium labelling for the 374 Δ -bclA strain. Simultaneous DNA labelling (Topro-3, blue fluorescence) is shown at 45 min. 375 For the *in vivo* germinating spores, 1 x 10⁸ germinated RPG1 spores obtained after the 3min-376 BHI incubation step were inoculated intraperitoneally and recovered one hour later. Images 377 are representative of at least two to three experiments (except for the *in vivo* experiment for 378 ethical reasons).

379

Figure 2: Kinetics, from 15 min to 2h, of co-detection of capsular material and exosporium BclA during germination of *B. anthracis* spores of the encapsulated RPG1 strain; same growth conditions and immunofluorescence labelling as in Fig.1. The 3-min time point represents the spores after germination triggering in BHI, in conditions where the capsule is not expressed.

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Figure 3: Same growth conditions as in Fig.1. (a) Same immunofluorescence labelling as in 386 387 Fig.1; (top) confocal analysis of germinating spores at 45 min, (bottom) image deconvolution 388 in the same experimental conditions. (b and c) spore germination was studied independently 389 by transmission electron microscopy w/o labelling (JNT's lab) and by scanning electron 390 microscopy with specific labelling of BclA and capsular material (PLG's lab) - (b) 391 Transmission electron microscopic analysis of extracellular RPG1 spores (left) and the non-392 encapsulated RPLC2 parental strain (right) 1h after dendritic cell infection (see Materials and 393 methods). (c) scanning electron microscopic analysis of immunolabelled RPG1 germinating 394 spores at 45 min (see Fig. 1); 10 nm beads for the anti-BclA antibodies, 15 nm beads for the 395 anticapsular material antibodies (see Materials and methods); to facilitate reading of the 396 image is shown in the insert a colored representation of the 10 nm anti-BclA beads (colored in 397 green) and 15 nm anti-capsule beads (colored in red) in a representative given area of the 398 spore.

- **Table 1:** Quantitative analysis of the topographical distribution of the immunofluorescence
- 400 during germination of the encapsulated RPG1 strain; a numbered schematic representation of
- 401 the type of observed immunofluorescence signals is shown in the two left columns. After a
- 402 3min-step of germination triggering ("0"min), germinated spores were incubated in R
- 403 medium 0.6% bicarbonate in a 5% CO2 atmosphere, recovered at various time points
- 404 thereafter and immunolabelling performed after binding on PLL-coated glass coverslips (see
- 405 Materials and Methods). Enumeration of the labelled spores is expressed as percentage (n=
- 406 number of spores enumerated on successive fields). red: anticapsular material labelling, green:
- 407 anti-BclA labelling.
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Form type	Immuno- fluorescence distribution	"0" min (n=103)	15 min (n=55)	30 min (n=64)	60 min (n=59)	90 min (n=50)	120 min (n=50)
1	\bigcirc	100%	16 %	9 %	10 %	26 %	24 %
2			24 %	8 %			
3				2 %	•		
4	\bigcirc		31 %	12 %	3 %	4 %	
5	\bigcirc		27 %	31 %	10 %	2 %	
6	\bigcirc		2 %	36 %	47 %	16 %	
7				2 %			
8					29 %	52 %	76 %



Fig 1: Fastenackels et al



Fig 2: Fastenackels et al





Fig 3: Fastenackels et al