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► **To cite this version:**

Thomas Candela, Annette Fagerlund, Christophe Buisson, Nathalie Gilois, Anne-brit Kolstø, et al.. CalY is a major virulence factor and a biofilm matrix protein. *Molecular Microbiology*, 2019, 111 (6), pp.1416-1429. 10.1111/mmi.14184 . hal-02620254v1

HAL Id: hal-02620254

<https://hal.inrae.fr/hal-02620254v1>

Submitted on 30 Nov 2023 (v1), last revised 8 Dec 2023 (v2)

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Journal:	<i>Molecular Microbiology</i>
Manuscript ID	MMI-2018-17131.R1
Manuscript Type:	Research Article
Date Submitted by the Author:	n/a
Complete List of Authors:	<p>Candela, Thomas; Université Paris sud XI, Faculté de pharmacie, EA4043 Fagerlund, Annette; Universitetet i Oslo Farmasoytisk institutt, School of Pharmacy Buisson, Christophe; INRA, Micalis Gilois, Nathalie; INRA, Micalis Kolsto, Anne Brit; Universitetet i Oslo Farmasoytisk institutt, School of Pharmacy Økstad, Ole Andreas; Universitetet i Oslo Farmasoytisk institutt, School of Pharmacy Aymerich, Stéphane; INRA, Micalis Nielsen-LeRoux, Christina; INRA, Micalis Lereclus, Didier; INRA, Micalis Institute GOHAR, Michel; INRA, Micalis</p>
Key Words:	Bacillus, thuringiensis, cereus, anthracis, adhesin

CalY is a major virulence factor and a biofilm matrix protein

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6 Thomas Candela^{a*}, Annette Fagerlund^{b#}, Christophe Buisson^a, Nathalie Gilois^a, Anne-Brit

7 Kolstø^b, Ole-Andreas Økstad^b, Stéphane Aymerich^a, Christina Nielsen-Leroux^a, Didier

8 Lereclus^a, and Michel Gohar^{a#}

9

10

11

12 ^a Micalis Institute, INRA, AgroParisTech, Université Paris-Saclay, 78350 Jouy-en-Josas, France

13 ^b Laboratory for Microbial Dynamics, School of Pharmacy and Centre for Integrative Microbial
14 Evolution, University of Oslo, Oslo, Norway

15 * present address : EA4043, Faculté de Pharmacie, Université Paris Sud, Châtenay-Malabry,
16 France.

17 # present address : Nofima, Norwegian Institute of Food, Fisheries and Aquaculture Research,
18 Ås, Norway

19

20

21

22

23 # Corresponding author: michel.gohar@inra.fr

25 **Abstract**

26 The extracellular biofilm matrix often contains a network of amyloid fibers which, in the human
27 opportunistic pathogen *Bacillus cereus*, includes the two homologous proteins TasA and CalY.
28 We show here, in the closely related entomopathogenic species *B. thuringiensis*, that CalY also
29 displays a second function. In the early stationary phase of planktonic cultures, CalY was located
30 at the bacterial cell-surface, as shown by immunodetection. Deletion of calY revealed that this
31 protein plays a major role in adhesion to HeLa epithelial cells, to the insect *Galleria mellonella*
32 haemocytes, and in the bacterial virulence against larvae of this insect, suggesting that CalY is a
33 cell-surface adhesin. In mid-stationary phase and in biofilms, the location of CalY shifted from
34 the cell surface to the extracellular medium, where it was found as fibers. The transcription study
35 and the deletion of sipW suggested that CalY change of location is due to a delayed activity of
36 the SipW signal peptidase. Using purified CalY, we found that the protein polymerization
37 occurred only in the presence of cell-surface components. CalY is therefore a bifunctional
38 protein, which switches from a cell-surface adhesin activity in early stationary phase, to the
39 production of fibers in mid-stationary phase and in biofilms.

40 Introduction

41 *Bacillus thuringiensis*, an insect pathogen genetically close to the human lethal pathogen *B.*
42 *anthracis* and to the opportunistic food-borne pathogen *B. cereus*, produces biofilm pellicles
43 floating on the culture medium (Auger *et al.*, 2006, Wijman *et al.*, 2007) or sticking to
44 submerged solid surfaces (Houry *et al.*, 2012). The biofilm matrix of these species includes
45 DNA, polysaccharides and proteins (Vilain *et al.*, 2009, Houry *et al.*, 2012). In the closely
46 related species *Bacillus subtilis*, three proteins are required to build the biofilm. TasA forms
47 amyloid fibers (Romero *et al.*, 2011), TapA is required for TasA polymerization and anchoring
48 to cell surfaces (Romero *et al.*, 2014), and BslA coats the biofilm (Hobley *et al.*, 2013). The *B.*
49 *subtilis* *tasA* and *tapA* genes are included in the *tapA-sipW-tasA* operon, in which *sipW* codes for
50 a signal peptidase required for the secretion of TasA and TapA (Stover & Driks, 1999b, Stover
51 & Driks, 1999a). Transcription of *tapA-sipW-tasA* is repressed by SinR and promoted by the
52 anti-SinR protein SinI (Kearns *et al.*, 2005). In *B. thuringiensis*, *B. cereus* and *B. anthracis*, two
53 orthologues of the *B. subtilis* *tasA* are found. One of these orthologues - also named *tasA* - is
54 found just downstream of the signal peptidase gene *sipW*, in the SinR-regulated bicistronic
55 operon *sipW-tasA* (Caro-Astorga *et al.*, 2015, Pflughoeft *et al.*, 2011, Fagerlund *et al.*, 2014).
56 Located downstream from *sipW-tasA* is the second orthologue of *tasA* named *calY*. The *calY*
57 gene is expressed from its own promoter and is also controlled by SinR (Caro-Astorga *et al.*,
58 2015, Pflughoeft *et al.*, 2011, Fagerlund *et al.*, 2014). In *B. cereus*, both CalY and TasA
59 polymerize to form fibers in the biofilm matrix, although TasA was more efficient than CalY for
60 fiber production (Caro-Astorga *et al.*, 2015). Consequently, in this species, the deletion of either
61 *tasA* or *calY* led to a decrease in the biofilm biomass produced in microtiter plate assays (Caro-
62 Astorga *et al.*, 2015).

63 CalY was at first called camelysin, a name which stands for ‘casein-cleaving membrane
64 metalloproteinase’. CalY has indeed previously been described as a cell-surface, membrane-

65 bound zinc metallopeptidase active against casein, plasminogen, actin, collagen or fibrinogen
66 (Fricke *et al.*, 2001, Grass *et al.*, 2004). Because of the limited identity of the protein to other
67 metalloproteases, a new family was created, the M73 peptidase family (MEROPS database
68 accession number MER031615), which so far contains only CalY and TasA, but none of these
69 proteins has identified zinc-binding or catalytic sites. Meanwhile, CalY is often cited in the
70 literature as a protease involved in degradation of host tissues and in toxin activation (Nisnevitch
71 *et al.*, 2006, Nisnevitch *et al.*, 2010, Bai *et al.*, 2002, Irshad *et al.*, 2018) but without
72 experimental evidence for such activity. Furthermore, CalY has also been reported in *B. cereus*
73 to be a cell-surface protein able to bind to fibronectin and to mucin (Sanchez *et al.*, 2009).
74 Both the putative proteolytic and binding activities of CalY suggest that it could take part in
75 pathogenesis, while its ability to form fibers and the consequences of its deletion on biofilm
76 formation shows that it is a component of the biofilm matrix. The *calY* gene is likely to have
77 arisen from *tasA* by gene duplication, a process which can lead to the evolution of moonlighting
78 proteins (Espinosa-Cantu *et al.*, 2015). We therefore hypothesized that CalY could be a
79 bifunctional protein, involved both in the biofilm matrix construction and in adhesion to host
80 tissues. We found that CalY is indeed a major virulence factor and a key component of the
81 biofilm in *B. thuringiensis*, moving from one function to the other according to the subcellular
82 location and to the culture growth state.

83

84

85

87 **Results**

88 **CalY is not a protease**

89 The proteolytic activity previously described for CalY was obtained with a sample purified from
90 *B. cereus* cell-surface extracts, and with azocasein as a substrate (Fricke *et al.*, 1995). Here, we
91 assayed on the same substrate, azocasein, a recombinant CalY (rCalY) purified from *E. coli*. As
92 shown in Table 1, azocasein was not hydrolyzed by rCalY purified from two preparations but
93 was strongly degraded by the two metalloproteases thermolysin and InhA2, used here as positive
94 controls. In addition, a proteolytic activity has never been reported for *B. subtilis* TasA, and there
95 is only one histidine in the CalY amino acids sequence whereas most zinc metalloproteases
96 display two histidines in their catalytic or metal binding site (Cerdeira-Costa & Gomis-Ruth, 2014).
97 Therefore, CalY is most likely not a protease.

98 99 **Transcription analysis reveals that *calY* is the most overexpressed** 100 **gene in biofilms relatively to planktonic cultures in early stationary** 101 **phase**

102 The expression profile of the whole Bt407 genome was compared in 24 h biofilm cultures
103 relatively to early stationary phase planktonic cultures by microarray analysis. Three hundred
104 and five genes displayed a ratio of expression (biofilm/planktonic) greater than 2 (Table S1). By
105 far, the most overexpressed gene in biofilms was *calY*, with an expression ratio of 36, whereas
106 the *tasA* expression ratio was 7. In addition, *sipW-tasA* transcription starts at the onset of
107 stationary phase, one hour earlier than *calY* transcription (Fig. 1). Overall, 12 genes, putatively
108 involved in biofilm formation, were found to be differentially expressed in biofilms. Six of these
109 genes (*sipW*, *tasA*, BTB_c13240, *calY*, *sinI*, *sinR*) are located in the overexpressed *sipW*-

110 *sinR* locus. The six other genes differentially expressed in biofilm include 4 transcriptional or
111 post-transcriptional regulators (*ai2K*, *abrB*, BTB_c16240 and the diguanylate
112 cyclase/phosphodiesterase BTB_c54300), and two genes involved in the
113 biosynthesis and secretion of kurstakin - a lipopeptide shown to be required for biofilm
114 formation (Dubois *et al.*, 2012, Fagerlund *et al.*, 2014, Gelis-Jeanvoine *et al.*, 2016).

115

116 **CalY is a biofilm matrix component**

117 Deletion of *calY* led to a strong and significant decrease in the 48h-aged biofilms biomass, which
118 could not be restored by complementation (Fig. 2A). We followed morphological changes in the
119 formation of the pellicle in a 48 well microtiterplate in HCT medium over a 48h-time period for
120 the wild-type strain, the *calY* mutant strain and the complemented *calY* mutant strain (Fig. 2B).
121 Whereas the pellicle could be seen as soon as 18h in the wild-type strain, it was absent in the
122 mutant strain and was present but incompletely formed in the complemented strain. After 24h of
123 culture, the pellicle was thick in the wild-type strain, but remained very thin and displayed holes
124 and tears in the mutant strain. In 48h-aged cultures, the wild type strain showed dense pellicles in
125 which the structure was hidden by thick biofilm materials, while the mutant strain pellicle was
126 thinner with a clear structure. At both 24h and 48h culture times, the complemented strain
127 pellicle displayed an intermediate phenotype between the wild-type strain and the mutant strain
128 (Fig. 2B), although the complemented strain did produce CalY (Fig. S2).

129 The location of CalY in the biofilm matrix was determined by immunodetection using epi-
130 fluorescence microscopy. The antibody, raised against rCalY, specifically recognizes CalY, but
131 not TasA (see Experimental Procedures and Fig. S2 in supplementary materials). The matrix of
132 the 48h-aged biofilm showed a dense network of CalY fibers in the wild type strain (Fig. 3).
133 Since needle shearing was used to disrupt the biofilm prior to immunodetection, fibers

134 organization might have been disturbed comparatively to the untreated biofilm. In contrast, no
135 CalY fibers could be seen in the biofilm matrix prepared from the *calY* mutant strain. In the *calY*
136 complemented strain, CalY was produced (Fig S2) but aggregated into rare fiber bundles, thicker
137 and longer than the fibers observed in the wild type strain (Fig. 3), which is likely to be the cause
138 of the partial defect in biofilm formation in the complemented strain.

139

140 **CalY overexpression could lead to fiber bundles**

141 We wanted to determine the reason why CalY forms fiber bundles in the complemented strain.
142 Sequencing of pCAL40, the multicopy plasmid used to express *calY* in the complemented strain,
143 revealed no mutation in *calY*, and sequencing of the whole *sipW-sinR* region in the *calY* mutant
144 showed 100% identity with the wild type strain, except for *calY* deletion (Fig S3). Therefore, the
145 formation of fiber bundles by CalY in the complemented strain cannot be a consequence of a
146 mutation in *calY* or in the region targeted by the recombination events. We then hypothesized
147 that an overexpression of *calY* from the multicopy pCAL40 plasmid in the complemented strain
148 could lead to an abnormal protein polymerization leading to the formation of fiber bundles and to
149 a biofilm defect. To verify this hypothesis, we transformed the wild-type strain with pCAL40
150 and determined the effect of this transformation on biofilm formation. The transformed strain
151 produced less biofilm than the wild-type strain, and was not significantly different from the *calY*
152 mutant and the *calY* complemented strains for this phenotype (Fig. 2A), supporting our
153 hypothesis.

154

155 **CalY is a cell-surface adhesin**

156 The adhesion of the wild-type strain, the *calY* mutant and the complemented *calY* mutant grown
157 in planktonic cultures and harvested in early stationary phase was determined for *G. mellonella*

158 (the Greater wax moth) haemocytes and for HeLa cells. Deletion of *calY* reduced drastically and
159 significantly the adhesion to HeLa cells, down to the background level obtained in non-coated
160 plates, where the three strains behave similarly (Fig. 4A). Similar results were obtained for *G.*
161 *mellonella* haemocytes (Fig. 4B), showing that CalY is a major adhesin in *B. thuringiensis*.
162 Complementation restored the adhesion capability of the *calY* strain, in contrast with the absence
163 of complementation in the biofilm phenotype, and CalY overexpression led to an increase,
164 although not significant, in adhesion comparatively to the wild type strain (Fig. 4B). This result
165 suggested that CalY might be located at the cell surface in planktonic cultures. To check for this
166 possibility, the subcellular location of CalY was determined by immunodetection in the wild-
167 type strain, the *calY* strain and the complemented *calY* strain. At the end of the exponential
168 phase, when *calY* is not transcribed (see Fig. 1), CalY could not be detected for any of the three
169 strains (Fig. 4C). In early stationary phase, CalY could be seen on the bacteria cell-surface for
170 the wild-type strain and for the complemented *calY* strain, but not for the *calY* mutant strain.
171 Interestingly, CalY was mainly found at the cell pole when present. In mid-stationary phase and
172 later, CalY was no longer located on the cell-surface in the wild-type strain, but was seen as
173 fibers surrounding the bacteria. CalY fibers were absent from the *calY* mutant strain cultures
174 harvested in mid-stationary phase, and were present as fiber bundles in the complemented *calY*
175 mutant strain cultures harvested at the same time (Fig. 4C).

176

177 **SipW inactivation promotes the cell-surface location of CalY**

178 Because no cell wall binding domain could be found in CalY sequence, we hypothesized that in
179 the early stationary phase of growth, the protein could remain anchored to the bacterium
180 cytoplasmic membrane by the hydrophobic region of its signal peptide. Cleavage of the signal
181 peptide could be delayed if *sipW* transcription occurred later than *calY* transcription, but the
182 reverse situation was observed (Fig. 1). An alternative hypothesis was that SipW could be in an

183 inactive state in early stationary phase, and activated later. To determine if SipW inactivity could
184 lead to a surface location of CalY, we deleted *sipW* and compared CalY location in the wild-type
185 strain, the mutant strain and the complemented strain in cultures harvested in mid-stationary
186 phase. In the wild type strain, CalY was present in the extracellular medium as fibers, but was
187 found only at the cell surface, at the bacterial poles, in the *sipW* mutant strain, and
188 complementation restored the extracellular location of the protein (Fig. 5).

189

190 **CalY polymerization requires an activating factor**

191 While CalY could form fibers in biofilm as well as in planktonic culture, rCalY (produced from
192 *E. coli* without the signal peptide) did not spontaneously polymerize. These results suggested
193 that a factor, either secreted or cell-surface bound, was required for CalY, polymerization. To
194 check this hypothesis, we incubated rCalY with washed bacteria or with a cell-free supernatant
195 obtained from a Bt407 $\Delta calY$ planktonic culture harvested in both early and mid-stationary
196 phases, because CalY fibers were observed in a wild type strain culture harvested in mid- but not
197 in early-stationary phase. The presence of CalY fibers in the two cultures fractions was then
198 determined using the anti-CalY antibody. In cultures harvested in early stationary phase, no
199 CalY fibers could be detected, either in the cell-free supernatant or in the washed bacteria (Fig.
200 6). In contrast, bundles of antibody-reacting fibers were seen in the cell fraction – but not in the
201 cell-free supernatant fraction – obtained from cultures harvested in mid-stationary phase (Fig. 6
202 and Fig. S4). These fibers bundles were present in the Bt407 $\Delta calY$ culture only when rCalY was
203 added, and rCalY alone was unable to form fibers (Fig. 6). Therefore, a cell-associated factor is
204 required for the extracellular CalY polymerization. This result also support the hypothesis that
205 CalY in high concentrations polymerizes as fiber bundles.

206

207 **CalY is a major virulence factor in *B. thuringiensis***

208 CalY involvement in the *B. thuringiensis* pathogenesis against larvae of the lepidoptera *G.*
209 *mellonella*, was assessed by force-feeding (ingestion) or by injection into the hemolymph of
210 various doses of vegetative cells. Dose-response curves were built for the wild-type strain, the
211 *calY* mutant strain and the complemented *calY* mutant strain, and LD50s were calculated from
212 these dose-response curves. Deletion of *calY* resulted in a significant, 12-fold increase in the
213 LD50 obtained by ingestion assays, whereas complementation of *calY* only did not restore the
214 wild-type strain virulence in these assays (Fig. 7). Similarly, injection assays resulted in a 13-
215 fold increase of the LD50 when *calY* was deleted, but here, complementation fully restored the
216 wild-type strain LD50 (Fig. 7).

217

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219 **Discussion**

220 Since its description by Fricke *et al.* in 1995 (Fricke *et al.*, 1995), CalY has been assigned a
221 number of functions. This protein was first reported to be a cell-surface metallopeptidase. We
222 could not detect such an activity on a purified, recombinant sample of the protein. We
223 hypothesize that the metallopeptidase activity reported earlier for CalY is a consequence of the
224 purification process from cell-surface extracts: on the one hand, metallopeptidases could have
225 been co-purified with CalY; and on the other hand, our results show that CalY is present in high
226 quantities as insoluble fibers in mid-stationary phase planktonic cultures, which are likely to be
227 recovered during the cell-envelope preparation process. We therefore conclude that CalY is
228 unlikely to be a metallopeptidase.

229 However, we confirm a recent finding showing that CalY is a component of the biofilm matrix,
230 where it forms fibers (Caro-Astorga *et al.*, 2015). The authors suggested that CalY could play a
231 prominent role in the matrix at the initiation of biofilm formation, and was likely to be replaced
232 by TasA in mature biofilms. Using immunodetection methods and microarray analysis, we could
233 observe that CalY is present as fibers in high quantity in 48h-aged biofilms (Fig. 3), and that
234 *calY* is the most overexpressed gene in mature biofilms compared to planktonic cultures in early
235 stationary phase (Table S1). Deletion of the gene encoding CalY leads to a decrease in the
236 quantity of biofilm produced and in a reduction in the pellicle thickness (Fig. 2).

237 While CalY is produced in high quantities in biofilms, it is also highly expressed in planktonic
238 cultures. Its transcription starts in early stationary phase and increases sharply in mid stationary
239 phase (Fig. 1). Upon entry into stationary phase, when *calY* transcription is still low, CalY is
240 located only at the bacterium cell-surface (Fig. 4). In contrast, later in the stationary phase, CalY
241 is absent from the cell surface and is found as free extracellular insoluble fibers. We have shown
242 here that, when *sipW* is deleted, CalY remains attached to the cell surface (Fig. 5). Therefore, it
243 is likely that CalY is first anchored to the cell membrane by the hydrophobic domain of its signal

244 peptide and then cleaved by SipW to reach the extracellular medium. The transient presence of
245 CalY at the cell surface cannot be a consequence of a delay in *sipW* transcription, because *sipW*
246 transcription starts before *calY* transcription (Fig. 1). Alternatively, SipW activity could require
247 the presence of a cofactor which production would start in mid-stationary phase. This hypothesis
248 is supported by a previous report suggesting the existence of a SipW activator in *B. subtilis*
249 (Tjalsma *et al.*, 2000).

250 The purified CalY used to raise antibodies did not polymerize spontaneously upon incubation.
251 This result suggested that an activating factor, different from SipW (because the purified CalY
252 was without signal peptide), was required for CalY polymerization. We investigated this
253 possibility by mixing purified CalY with planktonic cultures fractions from the Bt407 $\Delta calY$
254 strain (Fig. 6). We found that a cell-associated factor, most likely cell-surface bound, could
255 promote fibers formation by the exogenous CalY. This activating factor was not present in
256 cultures until mid-stationary phase, in agreement with the observation that planktonic cultures of
257 the wild type strain produce CalY fibers only from this time of the stationary phase and later.
258 The need of an activating factor for CalY polymerization is reminiscent of the requirement of
259 TapA for TasA polymerisation in *B. subtilis* (Romero *et al.*, 2011). Yet, since there is no TapA
260 homologue in *B. thuringiensis*, *B. cereus* or *B. anthracis*, the activation mechanism for CalY
261 might be different from the activation of TasA by TapA in *B. subtilis*.

262 The presence of CalY at the cell-surface, and the fact that it can bind mucin or fibronectin
263 (Sanchez *et al.*, 2009), suggested that this protein could interact with host tissues. Our data show
264 that CalY is needed for bacterial adhesion to HeLa cells and to *G. mellonella* haemocytes (Fig.
265 4). To our knowledge, CalY is the first *B. thuringiensis* or *B. cereus* cell-surface adhesin shown
266 to be required for adhesion to HeLa epithelial cells or to insects haemocytes. While the *B. cereus*
267 cell-wall peptidase CwpFM was also reported to impact adhesion on HeLa cells, this effect was
268 likely to be a side-effect of the autolysin activity of this protein (Tran *et al.*, 2010). In *B.*

269 *anthracis*, two cell-wall anchored proteins, displaying LPXTG motifs, were found to bind
270 collagen (Xu *et al.*, 2004), and orthologues of their genes are found on the chromosome of *B.*
271 *cereus* and of *B. thuringiensis*. However, the role of these collagen-binding proteins on the
272 adhesion of bacteria to host cells or to host tissues was not investigated. BslA is a SLH protein
273 (harboring a S-Layer Homology domain), which gene is located on the pathogenicity island of
274 the *B. anthracis* virulence plasmid pXO1, and is therefore not present in *B. cereus* or *B.*
275 *thuringiensis*. BslA is required for adhesion of *B. anthracis* vegetative cells on BJ1 fibroblasts
276 (Kern & Schneewind, 2008). In addition, *bslA* deletion resulted in a dramatic increase in the
277 lethal dose in an anthrax disease guinea pig model (Kern & Schneewind, 2010).

278 Deletion of *calY* also resulted in a strong increase in the lethal dose of *B. thuringiensis* in the
279 insect infection model *G. mellonella*. The dose required to induce 50% mortality upon deletion
280 of *calY* had to be increased in the same order of magnitude, respectively 13 times and 12 times,
281 when bacteria were injected into the insect blood (haemocel) and when bacteria were force-fed
282 (Fig. 7). By this last contamination route, CalY is so far the only known virulence factor,
283 together with the metallopeptidase InhA2 (Fedhila *et al.*, 2003), which plays a major role in the
284 bacterial pathogenesis on its own. CalY activity both by ingestion and by injection possibly
285 reflects the dual function of this protein, acting as a biofilm matrix protein during the
286 colonization of the gut following oral infection, and as an adhesin interacting with haemocytes,
287 fatbody cells or other tissues following injection. CalY is therefore a major virulence factor of *B.*
288 *thuringiensis*, and since it is highly conserved in *B. cereus* and *B. anthracis* (Fig. S5), it is likely
289 that this protein is also involved in the pathogenicity of these species.

290 High concentrations of CalY obtained by an exogenous addition of this protein to a bacterial
291 culture resulted in the formation of thick bundles of fibers in small number instead of the dense
292 network of thin fibers observed in the wild type strain without addition of CalY (Fig. 6). The
293 formation of these fiber bundles is unlikely to stabilize the biofilm. This hypothesis is supported

294 by the fact that an overexpression of CalY, obtained by transformation of the wild type strain by
295 pCAL40, a multicopy plasmid expressing *calY*, led to a decrease in biofilm formation. This can
296 explain why complementation by pCAL40 failed to restore biofilm formation in the $\Delta calY$ strain,
297 but succeeded to restore adhesion to epithelial cells – a phenotype for which we do not expect
298 that CalY polymerization is required. Similarly, virulence in force-feeding assays (not
299 complemented) might require the formation of a biofilm in the insect intestinal tract, while
300 virulence in injection assays (complemented) could need only adhesion on host tissues.

301 Our results show that CalY is a bifunctional protein, expressed at a high level in stationary
302 phase. In *Staphylococcus epidermidis* too, a bifunctional protein has been described, which can
303 either work as an adhesin or promote biofilm formation (Buttner *et al.*, 2015). This protein, Aap,
304 is a cell-surface protein which displays 3 domains. The C-terminal domain C anchors the protein
305 to the bacterial cell surface. The N-terminal domain A is required for the binding of bacteria to
306 epithelial cells. Removal of the A domain by proteolysis activates domain B, which can thereby
307 form dimers and promote intercellular adhesion and biofilm formation. However, unlike CalY,
308 Aap is not released in the extracellular medium and do not form long fibers. Another biofilm
309 bifunctional protein, recently described, is the extracellular protein *B. subtilis* BslA, which plays
310 a role both in the biofilm architecture and in the formation of a hydrophobic layer lining it
311 (Arnaouteli *et al.*, 2017). Unlike these two proteins, CalY subcellular location changes during the
312 culture course. Indeed, CalY can be located at the cell-surface, where it acts as an adhesin
313 thereby promoting the binding of bacterial cells to host tissues. This function provides to CalY a
314 major role in *B. thuringiensis* virulence. It can also be found free in the extracellular medium,
315 where it polymerizes and form insoluble fibers. In this other function, CalY constitutes a major
316 component of the biofilm matrix. The switch between the two functions, summarized in Fig. 8, is
317 likely to be due to a change in the signal peptidase SipW activity, that itself most likely requires
318 a cofactor or an activator. An inactive SipW would keep CalY in the cell membrane, while an

319 active SipW would cleave CalY from the membrane and let it polymerize as fibers in the
320 presence of another, at yet undetermined cell-surface activating factor. The determination of the
321 nature of these activators will certainly be a very promising aspect of future work dedicated to
322 the elucidation of the mechanisms involved in CalY shift from one function to the other.

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324 **Experimental Procedures**

325 **Bacterial strains, plasmids and growth conditions**

326 *Escherichia coli* TG1 (Sambrook *et al.*, 1989) was used as a host for derivatives of
327 pUC19 (New England Laboratories), pQE30 (QIAGEN), pAT113 (Trieu-Cuot *et al.*,
328 1991) pHT304, pHT304-18Z (Arantes & Lereclus, 1991, Agaisse & Lereclus, 1994),
329 pGemT-easy (Promega), and pUC1318Spc (Candela & Fouet, 2005). Strain M15
330 harboring pREP4 (QIAGEN) was used for the production of recombinant CalY. HB101
331 (pRK24) was used for mating experiments (Trieu-Cuot *et al.*, 1987). The *B. thuringiensis*
332 strain (sequenced genome: GenBank accession number CP003889) used here was an
333 acrySTALLIFEROUS derivative of strain 407 (Lereclus *et al.*, 1989), designated hereafter as
334 Bt407. Planktonic cultures were grown in LB medium: the cultures were seeded at OD
335 0.01 and incubated at 30°C, agitation 175 rpm. Antibiotics were used as previously
336 described (Bouillaut *et al.*, 2005).

337

338 **Genetic constructions**

339 Plasmid extraction, endonuclease digestion, ligation and agarose and polyacrylamide gel
340 electrophoresis were carried out as described by Sambrook *et al.* (Sambrook *et al.*, 1989).
341 Polymerase chain reaction (PCR) amplifications were carried out with rTaq according to the
342 manufacturer (GE Healthcare). To produce His-tagged CalY, the *calY* open reading frame,
343 without the leading sequence encoding CalY signal peptide, was amplified by PCR with
344 the *calYs* and *calYa* oligonucleotides (Table 3). The amplified 534bp DNA fragment was
345 inserted into pGemT-easy giving rise to pCAL5. The DNA fragment was then
346 subcloned into pQE30 (QIAGEN) using *Bam*HI/*Hind*III, giving pCAL7.

347 To disrupt *calY*, a 2709bp DNA fragment containing *calY* was amplified by PCR with
348 oligonucleotides *cal1* and *cal4*. The amplified DNA fragment was inserted into pGemT-
349 easy giving rise to pCAL10. pCAL10 was digested with *HpaI*, and a spectinomycin-
350 resistance cassette was inserted giving pCAL20. The *Bam*HI fragment from pCAL20
351 was ligated into pAT113 giving pCAL30 (Trieu-Cuot *et al.*, 1993). This recombinant
352 suicide plasmid was transferred from *E. coli* to Bt407 by heterogamic conjugation
353 (Pezard *et al.*, 1991, Trieu-Cuot *et al.*, 1987) giving rise to 407*calY::spc*. The markerless
354 mutant Bt 407 Δ *sipW* was obtained with the recombinant suicide plasmid pMAD*sipW*.
355 This plasmid was built by inserting the *Bam*HI/*Eco*RI 5'- and *Eco*RI/*Nco*I 3'-regions of
356 *sipW*, amplified by PCR using primers *sipW*AmF/*SipW*AmR or *sipW*AvF/*sipW*AvR, in
357 pMAD (Arnaud *et al.*, 2004) digested by *Bam*HI and *Nco*I, and was transferred in Bt407
358 by electroporation (Lereclus *et al.*, 1989). The *calY* and *sipW* deletions were checked
359 by PCR and by sequencing. To complement 407*calY::spc* with *calY* the PCR fragment
360 containing *calY* and its promoter was amplified with *Pca*I1/*calY*a and inserted into
361 pHT304 (Arantes & Lereclus, 1991) digested by *Hind*III/*Bam*HI to give pCAL40. To
362 complement 407 Δ *sipW* with *sipW* the PCR fragment containing *sipW* and its promoter
363 was amplified with *sipW*cF/ *sipW*cR and inserted into pHT304 (Arantes & Lereclus, 1991)
364 digested by *Hind*III/*Bam*HI to give pHT304*sipW*.

365 To follow *calY* promoter expression, a fusion with this promoter and the *lacZ* reporter
366 was constructed. The *calY* promoter was amplified using *Pca*I1 and *Pca*I2 and cloned
367 into pHT304-18Z (Agaisse & Lereclus, 1994). Likewise, the *sipW* promoter was cloned
368 into pHT304-18Z using *Psip*WF and *Psip*WR primers to follow *sipW* transcription. The
369 resulting plasmids were transferred into Bt407 by electroporation. β -galactosidase specific
370 activity was measured as described previously, and are expressed in units of β -galactosidase per

371 milligram of protein (Perchat *et al.*, 2011). Each assay was carried out at least three times using
372 independent cultures.

373

374 **Peptidase assay**

375 In order to analyze a potential peptidase activity of rCalY, azocasein (A2765, Sigma-Aldrich
376 Inc., USA) was used as substrate. Thermolysin (peptidase family M4) from *Bacillus*
377 *thermoproteolyticus* (P1512, Sigma-Aldrich Inc., USA) and purified InhA2 (peptidase family
378 M6) were used as positive controls. InhA2 was purified from the supernatant of Bt407- Δ plcR
379 [pHT3015Apha3inhA] (Fedhila *et al.*, 2003), a strain overexpressing InhA2. The supernatant
380 from an early stationary LB culture was recovered, centrifuged and 0.22 μ m-filtrated. All proteins
381 were precipitated by 85% ammonium sulphate and resuspended in Bis-Tris HCl 25mM, pH7.2
382 CaCl₂ 4mM. After desalting, InhA2 was purified by ion exchange chromatography
383 (MonoQ5/50GL and HPLC) and eluted with 0-15% NaCl as a single 85 kDa protein. Azocasein
384 was used as a 1% suspension in buffer bis-Tris, HCl pH 7.2, 25 mM, CaCl₂ 4mM, ZnCl₂ 0,1mM.
385 Thermolysin, InhA2 and rCalY were assayed at 10 μ g/ml. Differences between CalY and rCalY
386 folding, because of Bt407 specific post-translational modifications, are unlikely to occur (Fig.
387 S6). The reaction was stopped by the dilution of a 200 μ l sample in 1 ml TCA 10%. The tube
388 was centrifuged to pellet the undigested substrate and the supernatant was mixed with 750 μ l
389 NaOH 1 M before measuring the OD₄₄₀. The assays were repeated three times. The OD₄₄₀ was
390 corrected for blank (without peptidase) values. The peptidase activity was expressed as (nmoles
391 azocasein degraded) / (min x mg peptidase).

392

393 **Microarray analysis**

394 Microarray analysis was performed as described previously (Fagerlund *et al.*, 2014). The Bt407
395 strain was grown in bactopectone medium (10g l⁻¹ bactopectone, 5g l⁻¹ yeast extract, 10g l⁻¹
396 NaCl) at 30°C. For planktonic cultures, an overnight culture was diluted 1:100 in 50ml
397 bactopectone medium, and harvested at the entry point into stationary phase (ie after 3 hours of
398 growth at 250rpm). Biofilm were produced as follows: one gram of glass wool was dry-sterilized
399 in a 500-ml erlenmeyer bottle. Overnight culture was diluted 1:200 in 100ml bactopectone
400 medium, and grown at 50rpm (very slow shaking) in the flask containing glass wool. The glass
401 wool remained semi-submerged in the medium so that the biofilm could grow either on
402 submerged parts of the glass wool or at the liquid/air interface. Cells were harvested after 24
403 hours. To remove unattached cells, the glass wool was gently rinsed twice using 30°C fresh
404 bactopectone medium. The attached cells were subsequently released from the glass wool by
405 shaking in 60% ice-cold methanol. Six biological replicates of planktonic and biofilm RNA,
406 respectively, were compared using the microarrays. Genes showing at least twofold differential
407 expression ($\log_2FC \geq 1.0$ or $\log_2FC \leq -1.0$) between the biofilm and planktonic samples, and with a
408 confidence level (adjusted P value) ≤ 0.05 were selected.

409

410 **Biofilm assays**

411 Biofilms were grown in HCT medium (Lecadet *et al.*, 1980), in glass tubes as described
412 earlier (Fagerlund *et al.*, 2014) or in 48 well microtiterplates seeded at OD 0.01 and
413 incubated at 30°C with no agitation. Biofilm biomass was assessed by measuring the
414 OD₆₀₀ of disrupted biofilms recovered in 1ml PBS. Means were computed from 5 to 10
415 replicates obtained from 3 independent cultures. Each pair of means was compared using
416 the Tukey's range test, with an α -level set at 0.05. Pellicles were observed and
417 photographed with a Leica MZ FLIII binocular microscope and a Sony NEX-5 digital
418 camera.

419

420 Antibody production and immunodetection

421 CalY was overexpressed using the pCal7 plasmid. His-tagged recombinant CalY (rCalY) was
422 purified as described by QIAGEN on Ni-NTA coupled to Superflow resin. rCalY Anti-
423 CalY serum was obtained from rabbits by three injections of 300 μ g of purified protein.
424 Injections and serum preparations were performed by Covalab
425 (<http://www.Covalab.com>). Anti-CalY serum was used at 1:10000 for Western blot
426 detection and at 1:100 for immunodetection on culture samples. Antibody binding in
427 Western blot was revealed with a goat anti-rabbit, peroxidase-conjugated secondary
428 antibody (Pierce antibodies, Thermo Fisher Scientific Inc., USA) and developed using
429 the ECL Western blotting analysis system (Amersham ECL Detection Reagents, GE
430 Healthcare Bio-Sciences Corp., USA). The selected antibodies were quite selective for
431 CalY (Fig. S3). Immunodetection of CalY in culture samples for microscopy purposes
432 was performed as follows. Biofilms recovered from glass tubes assays were
433 homogenized in cold PBS by aspirating/pushing ten times through a 26-gauge needle.
434 Planktonic cultures were centrifuged and the pellets were resuspended in cold PBS.
435 Cultures in PBS were mixed with formaldehyde at a final concentration of 4%, washed
436 twice with cold PBS and resuspended in PBS. Cultures were incubated for 10 minutes
437 at room temperature with anti-CalY antibodies diluted 1:100, washed twice with cold
438 PBS, resuspended in PBS and incubated with an AlexaFluor488-labelled goat anti-
439 rabbit antibody (Molecular Probes, ThermoFisherScientific Inc., USA) diluted 1:100 at
440 room temperature for 10 minutes. Cultures were then washed twice with cold PBS,
441 resuspended in PBS, stained with FM4-64 (Molecular Probes, ThermoFisherScientific
442 Inc., USA) diluted 1:100 and observed on a Zeiss Z1-AxioObserver fluorescence
443 microscope.

444

445 **CalY polymerization**

446 Planktonic cultures of the Bt407 $\Delta calY$ mutant strain were harvested in early- and mid-
447 stationary phases, and 100 μ l of the culture was centrifuged for 2 minutes at 7500 rpm at
448 room temperature. The sedimented bacteria, washed twice with PBS and resuspended
449 in 100 μ l of PBS, and the cell-free supernatant were each incubated with rCalY at a final
450 concentration of 1 μ g/ml for 20 minutes at room temperature, centrifuged for 2 minutes
451 at 7500 rpm at room temperature, resuspended in PBS, and treated for CalY
452 determination as described in 'Antibody production and immunodetection'. Controls
453 (rCalY alone, or sedimented bacteria from a 407 $\Delta calY$ mutant strain planktonic culture
454 harvested in mid-stationary phase and incubated without rCalY) were incubated and
455 treated as described above.

456

457 **Insect virulence assays**

458 The role of CalY in the pathogenicity of the bacterium was determined by comparing the lethal
459 effect of the wild-type strain, the *calY* mutant strain and the complemented *calY* mutant strain in
460 two assays (ingestion and injection) on *G. mellonella* (bred in our laboratory for over 10 years),
461 performed as previously described (Bouillaut *et al.*, 2005). Four (ingestion) or five (injection)
462 concentrations of bacteria in vegetative phase were used, and for each concentration the
463 experiment was repeated three times on a minimum of 20 larvae each time. Phosphate-buffered
464 saline was used for negative controls. Infected larvae were kept at 37°C and mortality was
465 recorded over 24 to 48 hours. The LD₅₀ values were based on mortality data obtained 48 hours
466 post injection and were determined by non-linear regression using JMP9 (SAS Institute Inc.,
467 USA). The model used for the regression was a derivative of the Hill equation $\frac{x^n}{a^n + x^n}$, where x is

468 the dose used and a and n are computed parameters. The parameters, determined with their 95%
469 confidence intervals, represent the LD50 (a) and the steepness of the curve (n).

470

471 **Adhesion assays**

472 For the HeLa cells, adhesion assays were performed as described previously (Ramarao &
473 Lereclus, 2006). Bacteria harvested in early stationary phase were incubated for 15 minutes at
474 37°C in 24-well polystyrene microtiterplates in wells containing confluent Hela cells (supplied
475 by the American Type Culture Collection, reference ATCC CCL-2) (5×10^5 per well) or no HeLa
476 cells. 5×10^6 bacteria were loaded per well. Non-attached bacteria were then removed by washing
477 three times with PBS, and HeLa cells were detached by scraping. Serial dilutions were plated on
478 LB plates to score adherent bacteria *versus* total bacteria. Three replications were performed
479 using three independent cultures. Each pair of means was compared using the Tukey's range test,
480 with an α -level set at 0.05.

481 For *G. mellonella* haemocytes, haemolymph was collected from last instar larvae (similar to
482 those used for infection studies) as follows. Larvae were cleaned by dipping them into sterile
483 water for 15 s, followed by ethanol 70% for 5 s., after what they were dried on sterile filter paper
484 (Whatman grade 5) and let to rest for 15 min. in a sterile Petri dish. The cuticle was then
485 punctured on the third false leg with a 27-gauge sterile needle. Drops of haemolymph were
486 directly collected in chilled anticoagulation buffer (AB) (vanSambeek & Wiesner, 1999) in a 1:4
487 v/v proportion. 500 μ l of this mixture was centrifuged at 3000g at room temperature for 5 min.,
488 and the haemocytes pellet was gently resuspended in 1 ml of a 1:1 v/v mixture of chilled AB
489 buffer and Grace insect medium (Sigma Aldrich G8142). This suspension was kept on ice to be
490 used within one hour. Haemocytes viability was checked using Trypan Blue, and their
491 concentration was assessed using Kova cell counting slides (Kova Intl.com, Garden Grove, CA,
492 USA). Haemocytes final concentration was *ca* 5×10^6 haemocytes /ml.

493 The Bt 407 wt strain, the *calY* strain, the complemented *calY* strain and the wt strain
494 overexpressing *calY* were grown in LB medium at 30°, 175 rpm and harvested in early stationary
495 phase. Bacteria concentrations were assessed by agar plating. To perform the adhesion assay,
496 100 µl from each culture were centrifuged at 6000 g at room temperature and the resulting pellet
497 was gently resuspended with 100 µl of a haemocyte suspension prepared as detailed above,
498 leading to a bacteria/haemocyte ratio (MOI) of 20. The mixture was incubated at room
499 temperature for 20 min. followed first by a treatment with DAPI diluted 1/1000 (to stain
500 haemocytes nuclei), and second by a fixation by formaldehyde 4% (to avoid the development of
501 bacteria cytotoxicity). The preparation was finely washed twice with cold PBS (by
502 centrifugation/resuspension) and immediately observed in phase contrast and epifluorescence
503 microscopy. The microscope tiling procedure was used to produce large images with a high
504 resolution composed of 3 x 3 fields of view, allowing the observation of a high number of
505 haemocytes and bacteria. Bacteria adherent to haemocytes were counted on these images. The
506 counts included at least 200 haemocytes per strain from three independent assays.

507

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509

510 **Acknowledgments**

511 We are grateful to Agnès Fouet for pUC1318spc and to Patrick Trieu-Cuot for pAT113.

512 We thank Sebastien Gelis-Jeanvoine for his help in genomic data handling. Thomas

513 Candela was funded by the region Ile de France under the program DIM Astrea.

514

515

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- 650

651 **Tables**

652

653

Protease :	Thermolysin	InhA2	CalY
	M21663	BTB_c06870	BTB_c13250
replicates	3	3	3
mean	139	99	0.7
sem	25	11	0.3

654

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Table 1: Metallopeptidase activity of CalY

656

The proteolytic activity of CalY was determined using azocasein as a substrate. Pure thermolysin and the metallopeptidase InhA2 were used as positive controls. InhA2 is produced by *B. thuringiensis* in stationary phase. Metallopeptidase activity is expressed in nmoles min⁻¹ mg⁻¹. sem : standard error on the mean.

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primer	sequence	use
calYs	GGATCCattggtggaggaacatttgcattcttagc	CalY overproduction
calYa	AAGCTTtatttttctccccagcttcttggttagc	CalY overproduction & calY complementation
cal1	cgcGGATCCccgaaaacagttaatacgttaaaag	calY deletion
cal4	catgCCATGGgcgcatctgctaaacgttcttccgg	calY deletion
SipWAmF	cgGGATCCgaagcaattagggcgaaagatag	sipW deletion
SipWAmR	gGAATTCgtctctctccctctcctgtg	sipW deletion
SipWAvF	gGAATTCtctggttggtatacttccgttag	sipW deletion
SipWAvR	CATGCCATGGcgttccatactcacgctcaataaac	sipW deletion
sipWcF	gGAATTCgttacgccgtaatacaaaaagg	sipW complementation
sipWcR	aaCTGCAGctaaacggaaagtataaccaaacga	sipW complementation
Pcal1	cccAAGCTTcggaaggacaaaagaaagtagaag	PcalY-lacZ fusion & calY complementation
Pcal2	tgcTCTAGAcacaatcaattccccctagc	PcalY-lacZ fusion
PsipWF	aaCTGCAGgttacgccgtaatacaaaaagg	PsipW-lacZ fusion
PsipWR	gcTCTAGAtcaccaccgctcgtttt	PsipW-lacZ fusion

676

Table 3: Primers used in this study

677

Upper-case letters show restriction sites

678

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679 **Figure captions**

680 **Figure 1: *calY* and *sipW-tasA* transcriptions.**

681 Transcription of *calY* (white circles) or of *sipW-tasA* (black circles) was monitored in planktonic cultures
682 in LB medium through the *lacZ* gene reporter. The inset shows a focus on the transition phase between t_1
683 (OD 1) and t_2 (OD 7). Each circle is the mean of three replications and the error bars are the standard error
684 of the mean. The x-axis time scale is relative to t_0 which is the transition between the exponential phase
685 and the stationary phase.

686 **Figure 2: Role of CalY in biofilm formation.**

687 A: the biomass of biofilms grown in glass tubes in HCT medium was determined for the wild-type strain
688 (wt), the *calY* mutant strain (*calY*), the complemented *calY* mutant strain (*calY_c*), and the wild type strain
689 overexpressing *calY* (*wt_c*). Each bar is the mean of 5 to 10 experiments, and error bars represent the
690 standard error of the mean. Bars with different letters (a or b) represent results that are significantly
691 different ($P < 0.05$), as determined by the Tukey's range test.

692 B: biofilms grown in 48-wells polystyrene microtiter plates (well diameter: 10mm) in HCT medium were
693 photographed at different culture times with a binocular microscope. wt: wild-type strain; *calY*: *calY*
694 mutant strain; *calY_c*: complemented *calY* mutant strain.

695 **Figure 3: CalY location in biofilm.**

696 The presence of CalY in 48h-aged biofilms was determined using an immunodetection method and
697 fluorescence microscopy. Biofilms were grown in glass tubes in HCT medium. wt, wild-type strain;
698 *calY*: *calY* mutant strain; *calY_c*: complemented *calY* mutant strain. Anti-CalY antibody was revealed by a
699 secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained using FM4-64
700 (red).

701 **Figure 4: CalY role in adhesion to epithelial HeLa cells and to *G. mellonella* haemocytes.**

702 A: adhesion to epithelial HeLa cells grown in 24-wells polystyrene microtiterplates (left), or to
703 microtiterplates alone (right), was determined for the wild-type strain (wt), the *calY* mutant strain (*calY*),
704 or the complemented *calY* mutant strain (*calY_c*). Each bar is the mean of 3 experiments, and error bars

705 represent the standard error of the mean. Bars with different letters (a or b) represent significantly
706 different results ($P < 0.05$), as determined by the Tukey's range test.

707 B: adhesion to *G. mellonella* haemocytes prepared from larvae haemolymph was determined for the wild-
708 type strain (wt), the *calY* mutant strain (*calY*), the complemented *calY* mutant strain (*calY_c*) or the strain
709 overexpressing *calY* (*calY₊*). Each bar is the mean of 3 experiments, and error bars represent the standard
710 error of the mean. Bars with different letters (a or b) represent significantly different results ($P < 0.05$), as
711 determined by the Tukey's range test.

712 C: Subcellular location CalY in planktonic cultures. The subcellular location of CalY in planktonic
713 cultures was determined using an immunodetection method and fluorescence microscopy. Planktonic
714 cultures were grown in LB medium and harvested at the end of the exponential phase (exp), in early- and
715 in mid-stationary phase. White arrows point to CalY dots at the cell-surface. wt, wild-type strain; *calY* :
716 *calY* mutant strain; *calY_c* : complemented *calY* mutant strain. Anti-CalY antibody was revealed by a
717 secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained using FM4-64
718 (red).

719 **Figure 5: Role of SipW in CalY subcellular location.**

720 The role of SipW on the subcellular location of CalY was determined using an immunodetection method
721 and fluorescence microscopy. Planktonic cultures were grown in LB medium and harvested in mid-
722 stationary phase. White arrows point to CalY dots at the cell-surface. The lower panel is a 4x –
723 magnification of the upper panel showing a detailed view of CalY presence at the bacterial surface. wt,
724 wild-type strain; *sipW*: *sipW* mutant strain; *sipW_c*: complemented *sipW* mutant strain. Anti-CalY antibody
725 was revealed by a secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained
726 using FM4-64 (red).

727 **Figure 6: Polymerization of CalY.**

728 Cell-free supernatant (S) or pelleted and PBS-washed bacteria (C) were prepared from planktonic cultures
729 grown in LB medium and harvested in mid-stationary phase. rCalY was incubated with the cell-free
730 supernatant (SNY) or with the washed bacteria resuspended in PBS (CY). Controls were rCalY incubated

731 alone (Y), or washed bacteria incubated without rCalY (C). CalY fibers were revealed with a rabbit anti-
732 CalY antibody and a goat anti-rabbit antibody tagged with Alexa488 (green). Phase: phase contrast.

733 **Figure 7: CalY role in the bacterium virulence.**

734 Different concentrations of vegetative cells were injected or force-fed to larvae of the lepidopteran
735 species *G. mellonella*. Mortality was recorded 48h post-treatment, and LD50s values were computed.
736 Bars show LD50s values, and error bars represent the 95% confidence interval. The Y-scale is in log-
737 units. wt: wild-type strain; *calY*: *calY* mutant strain; *calY_c*: complemented *calY* mutant strain. Bars with
738 different letters (a or b) represent significantly different results ($P < 0.05$), as determined by the Tukey's
739 range test.

740 **Figure 8: Schematic representation of CalY functions.**

741 Left panel: in early stationary phase, in planktonic culture, both SipW and CalY are expressed and located
742 on the cell-surface. SipW does not display its peptidase activity and CalY remains anchored in the cell
743 membrane by its signal peptide. CalY works as an adhesin which binds to host cells, and strongly
744 contributes to the bacterial virulence.

745 Right panel: later in the stationary phase or in biofilms, SipW is activated by an unknown, hypothetical
746 factor and cleaves CalY signal peptide. The free, extracellular CalY is activated by an undetermined, cell-
747 surface activating factor (not shown here), and polymerizes to produce amyloid fibers promoting biofilm
748 formation.

749

750

751

752 **Supporting information**

753 **Table S1: Microarray analysis of biofilm vs planktonic cultures.**

754 Planktonic cultures in early stationary phase and 24h-aged biofilms were harvested and compared for
755 genes expression by microarray analysis. Genes showing at least twofold differential expression between
756 the biofilm and planktonic samples, and with a confidence level (adjusted P value) ≤ 0.05 were selected.

757

758 **Fig. S2: CalY Western blot.**

759 Extracts of Bt407 (wt), *calY* mutant strain (calY) and complemented *calY* mutant strain (calYc) were
760 analyzed by Western blot. The anti-CalY antibody was revealed by a HRP-conjugated goat anti-rabbit
761 antibody and by the ECL detection system. CalYr was loaded on the gel as a positive control.

762

763 **Fig. S3: Sequencing of the *sipW-tasA* region in the *calY* strain.**

764 The *sipW-tasA* region from the wild type strain is shown in the mid-panel of the figure (shaded in
765 yellow). This region was sequenced in the *calY* mutant strain, from the end of the BTB_c13210 gene to
766 the beginning of *inhA1* (4428 bp), using 5 PCR fragments amplified from both the direct and the reverse
767 strands. The 10 generated fragments were aligned on the wild type strain sequence (available at
768 <https://www.ncbi.nlm.nih.gov/nucleotide/409171506>) using the LASTZ tool of the Geneious software. In
769 the lower panel, the black color shows the region covered by the assembled sequences. In the upper panel,
770 the green color shows 100% identity between the amplified sequences and the wild type strain sequence.
771 Mismatches were found only in *calY*, which was interrupted by a spectinomycin resistance cassette.

772

773 **Fig. S4: Polymerisation of rCalY.**

774 Examples of CalYr polymerisation in the presence of the cell fraction of different cultures of the 407
775 $\Delta calY$ strain collected in mid-stationary phase. Phase: phase contrast. Alexa488: immunodetection of
776 CalY. Overlay: overlay of the phase and the immunodetection pictures.

777

778 **Fig. S5: CalY conservation in *B. cereus*, *B. anthracis* and *B. thuringiensis*.**

779 A- The CalY predicted sequences from the *Bacillus cereus* ATCC14579 strain, the *Bacillus anthracis*
780 Ames ancestor strain and the *Bacillus thuringiensis* 407 strain were aligned using ClustalW, and
781 displayed 94% sequence identity.

782 B- The TasA and CalY predicted sequences from *B. subtilis*, *B. cereus*, *B. anthracis* and *B. thuringiensis*
783 strains were aligned and the genetic distances were plotted using the tree builder option from Geneious
784 (Biomatters Ltd, New Zealand). CalY sequences are highly conserved and clusterize apart from TasA
785 sequences.

786 **Fig. S6: Search for putative CalY post-translational modifications in the 407wt strain**

787 A: Picture from a 2D-gel electrophoresis experiment showing CalY

788 B: Mapping of the peptides identified by Peptide Mass Fingerprint on the CalY sequence.

789

790

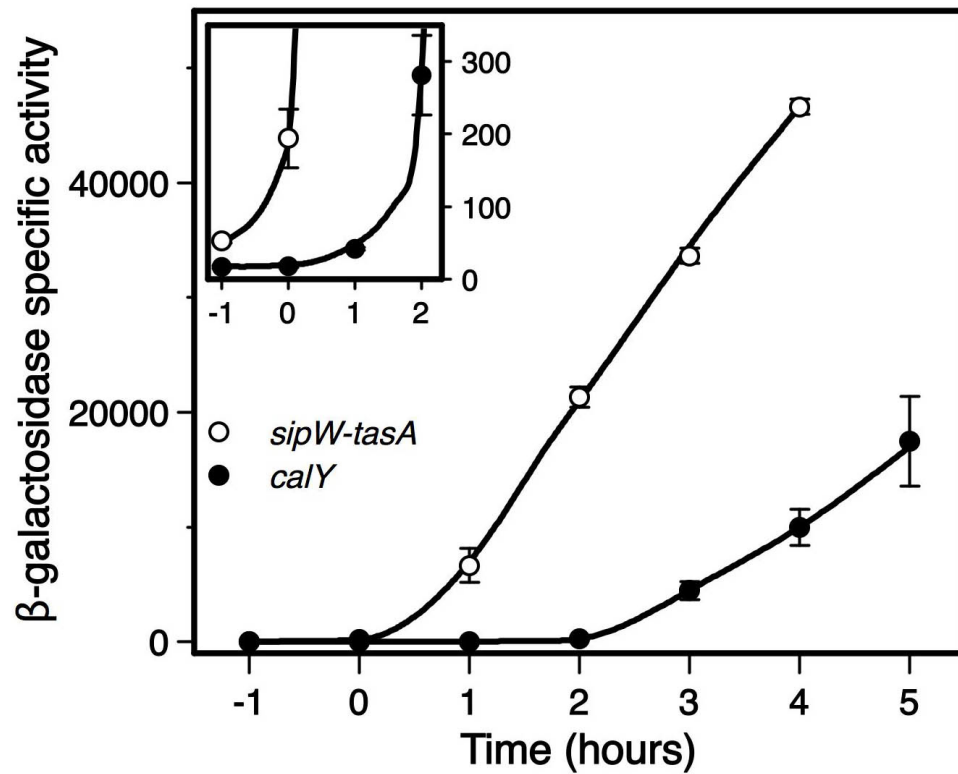


Figure 1: *calY* and *sipW-tasA* transcriptions.

Transcription of *calY* (white circles) or of *sipW-tasA* (black circles) was monitored in planktonic cultures in LB medium through the *lacZ* gene reporter. The inset shows a focus on the transition phase between t-1 (OD 1) and t2 (OD 7). Each circle is the mean of three replications and the error bars are the standard error of the mean. The x-axis time scale is relative to t0 which is the transition between the exponential phase and the stationary phase.

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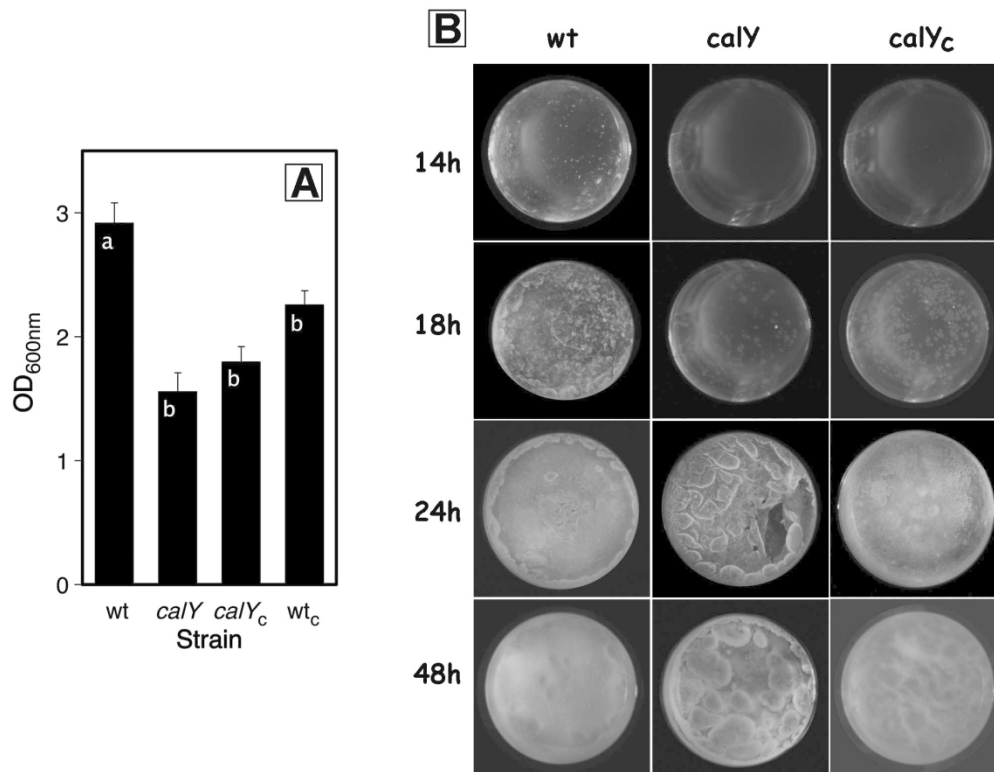
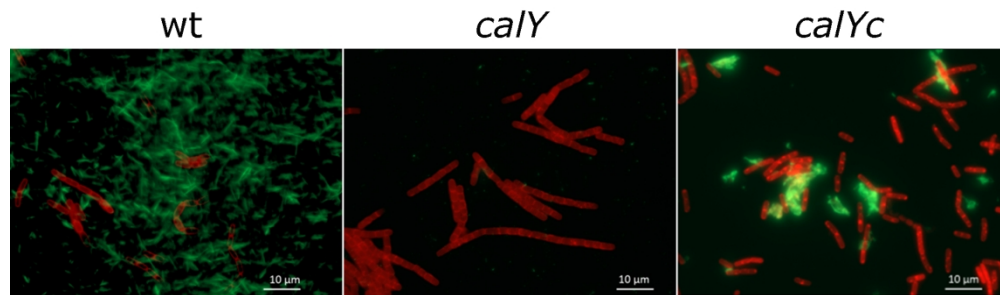


Figure 2: Role of CalY in biofilm formation.

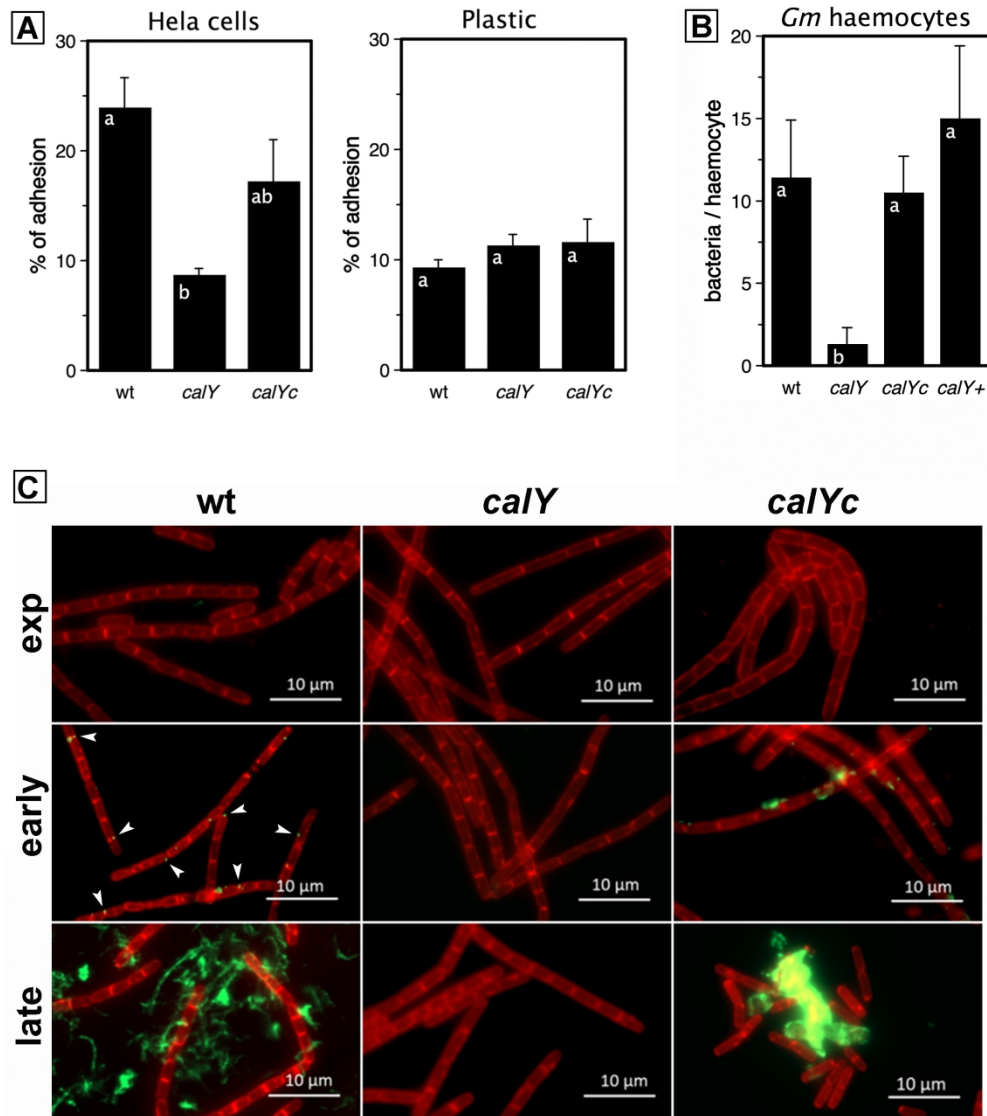
A: the biomass of biofilms grown in glass tubes in HCT medium was determined for the wild-type strain (wt), the calY mutant strain (calY), the complemented calY mutant strain (calY_c), and the wild type strain overexpressing calY (wt_C). Each bar is the mean of 5 to 10 experiments, and error bars represent the standard error of the mean. Bars with different letters (a or b) represent results that are significantly different ($P < 0.05$), as determined by the Tukey's range test.

B: biofilms grown in 48-wells polystyrene microtiter plates (well diameter: 10mm) in HCT medium were photographed at different culture times with a binocular microscope. wt: wild-type strain; calY: calY mutant strain; calY_c: complemented calY mutant strain.

170x134mm (300 x 300 DPI)



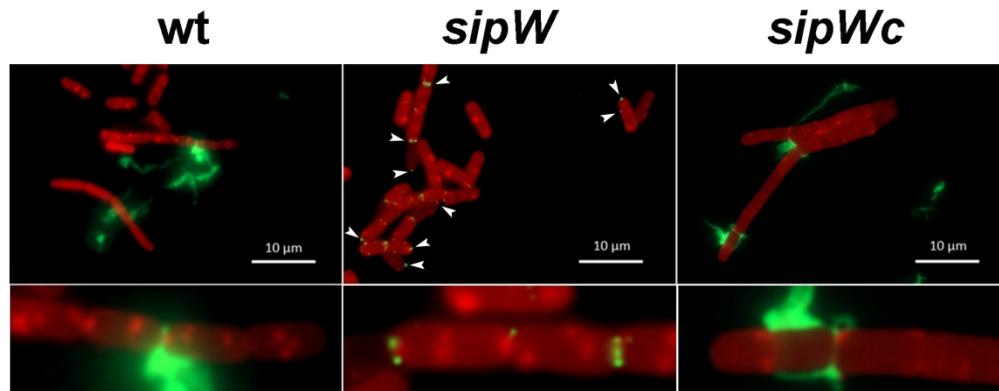
The presence of CaY in 48h-aged biofilms was determined using an immunodetection method and fluorescence microscopy. Biofilms were grown in glass tubes in HCT medium. wt, wild-type strain; calY: calY mutant strain; calYc: complemented calY mutant strain. Anti-CaY antibody was revealed by a secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained using FM4-64 (red).



A: adhesion to epithelial HeLa cells grown in 24-wells polystyrene microtiterplates (left), or to microtiterplates alone (right), was determined for the wild-type strain (wt), the calY mutant strain (calY), or the complemented calY mutant strain (calYc). Each bar is the mean of 3 experiments, and error bars represent the standard error of the mean. Bars with different letters (a or b) represent significantly different results ($P < 0.05$), as determined by the Tukey's range test.

B: adhesion to *G. mellonella* haemocytes prepared from larvae haemolymph was determined for the wild-type strain (wt), the calY mutant strain (calY), the complemented calY mutant strain (calYc) or the strain overexpressing calY (calY+). Each bar is the mean of 3 experiments, and error bars represent the standard error of the mean. Bars with different letters (a or b) represent significantly different results ($P < 0.05$), as determined by the Tukey's range test.

C: Subcellular location CalY in planktonic cultures. The subcellular location of CalY in planktonic cultures was determined using an immunodetection method and fluorescence microscopy. Planktonic cultures were grown in LB medium and harvested at the end of the exponential phase (exp), in early- and in mid-stationary phase. White arrows point to CalY dots at the cell-surface. wt, wild-type strain; calY : calY mutant strain; calYc : complemented calY mutant strain. Anti-CalY antibody was revealed by a secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained using FM4-64 (red).



The role of SipW on the subcellular location of CaY was determined using an immunodetection method and fluorescence microscopy. Planktonic cultures were grown in LB medium and harvested in mid-stationary phase. White arrows point to CaY dots at the cell-surface. The lower panel is a 4x -magnification of the upper panel showing a detailed view of CaY presence at the bacterial surface. wt, wild-type strain; sipW: sipW mutant strain; sipWc: complemented sipW mutant strain. Anti-CaY antibody was revealed by a secondary antibody labelled with Alexa488 (green). Bacterial membranes were stained using FM4-64 (red).

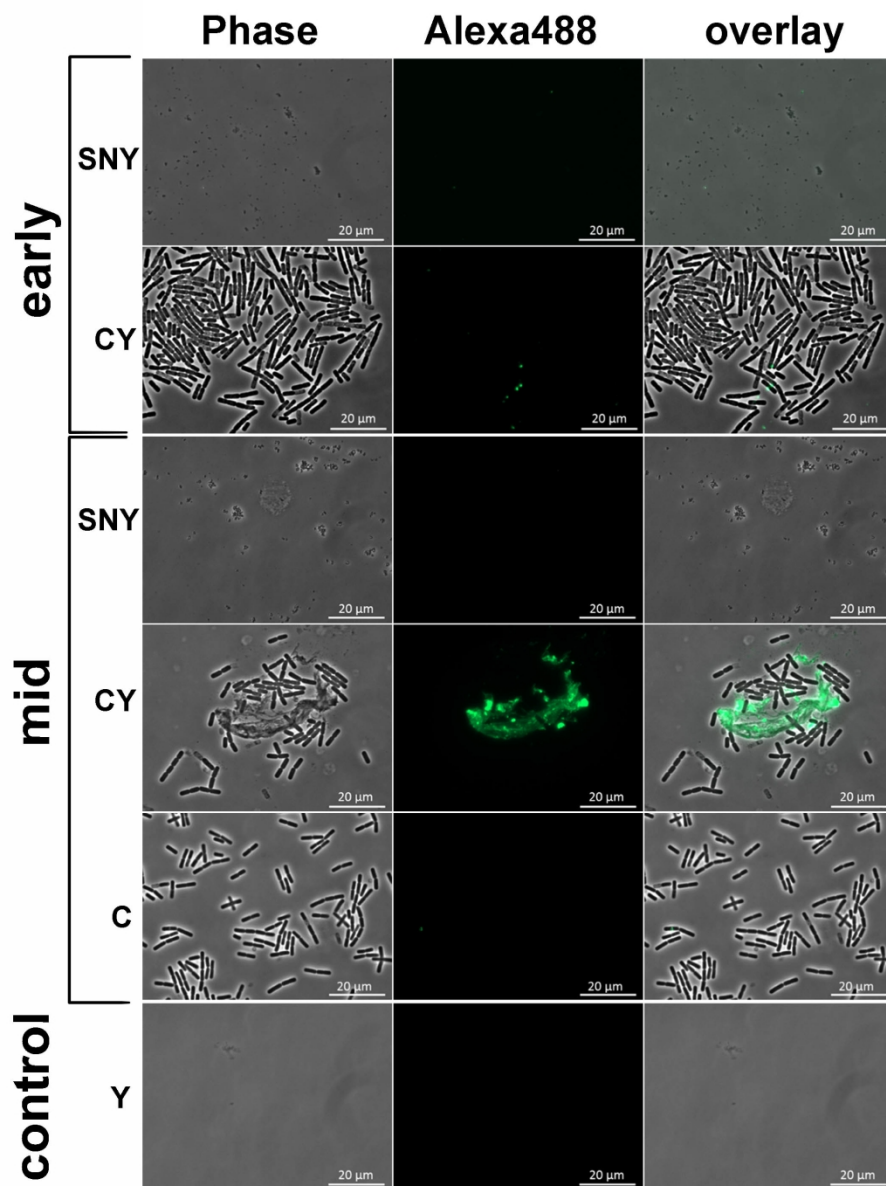


Figure 6: Polymerization of CaY.

Cell-free supernatant (S) or pelleted and PBS-washed bacteria (C) were prepared from planktonic cultures grown in LB medium and harvested in mid-stationary phase. rCaY was incubated with the cell-free supernatant (SNY) or with the washed bacteria resuspended in PBS (CY). Controls were rCaY incubated alone (Y), or washed bacteria incubated without rCaY (C). CaY fibers were revealed with a rabbit anti-CaY antibody and a goat anti-rabbit antibody tagged with Alexa488 (green). Phase: phase contrast.

178x239mm (300 x 300 DPI)

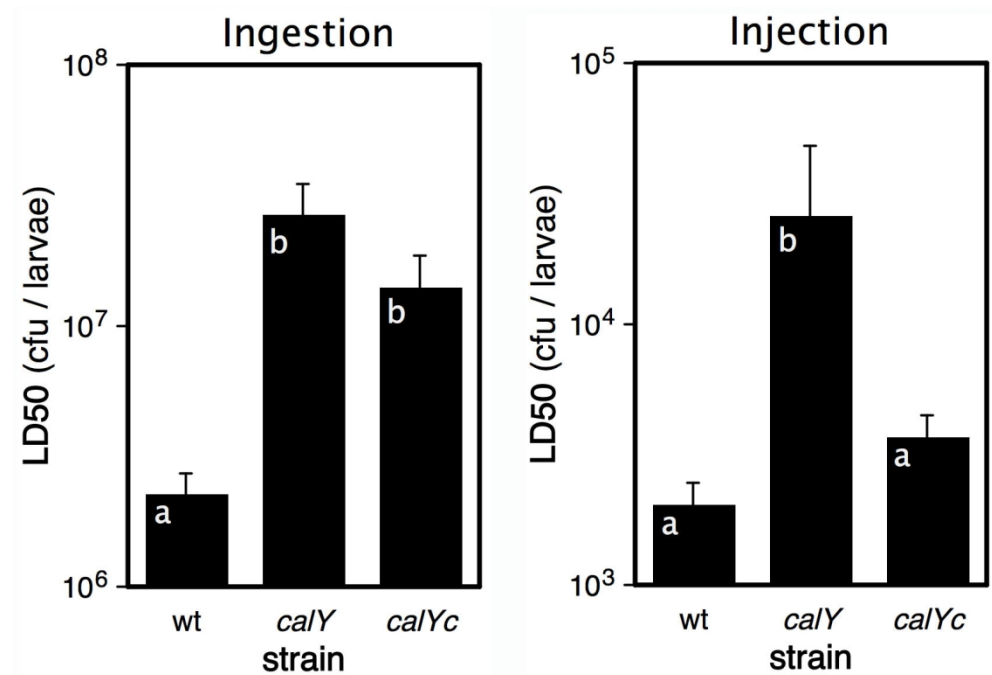
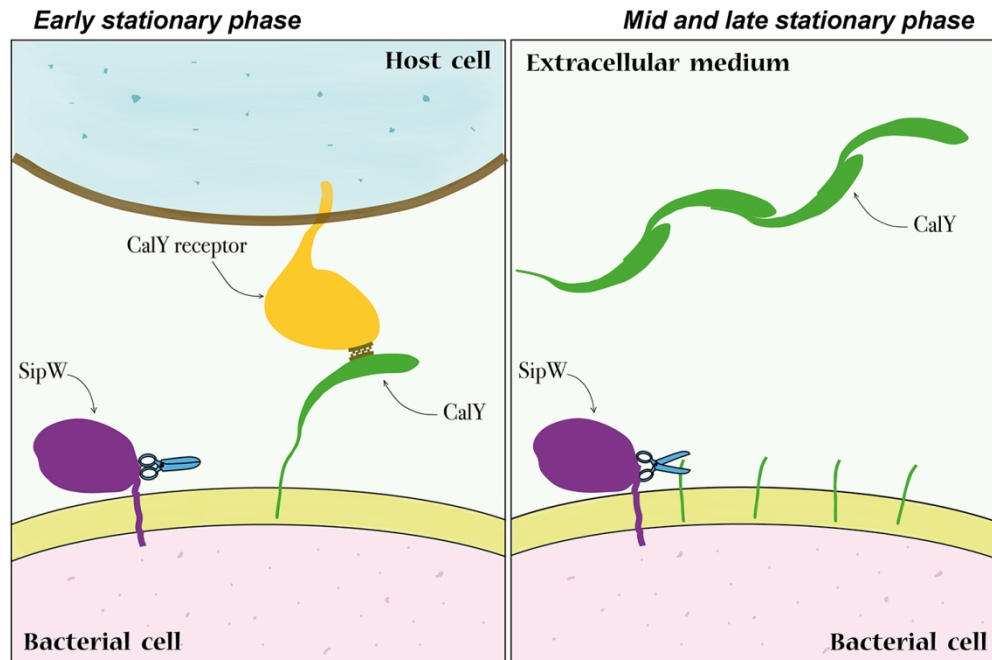


Figure 7: CalY role in the bacterium virulence.

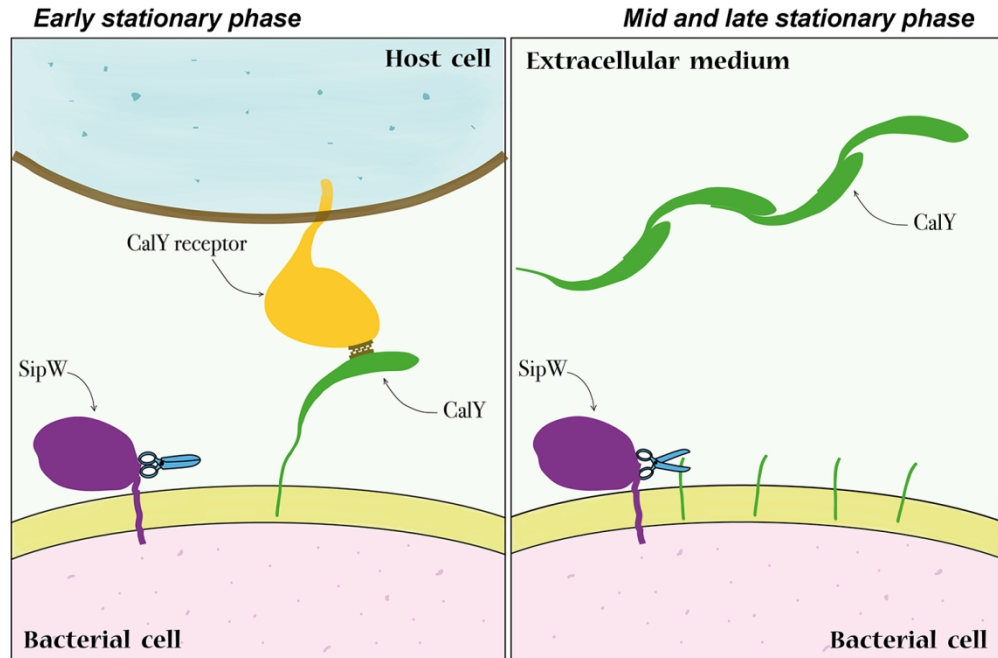
Different concentrations of vegetative cells were injected or force-fed to larvae of the lepidopteran species *Galleria mellonella*. Mortality was recorded 48h post-treatment, and LD50s values were computed. Bars show LD50s values, and error bars represent the 95% confidence interval. The Y-scale is in log-units. wt: wild-type strain; calY: calY mutant strain; calYc: complemented calY mutant strain. Bars with different letters (a or b) represent significantly different results ($P < 0.05$) different, as determined by the Tukey's range test.

144x98mm (300 x 300 DPI)



Left panel: in early stationary phase, in planktonic culture, both SipW and CalY are expressed and located on the cell-surface. SipW does not display its peptidase activity and CalY remains anchored in the cell membrane by its signal peptide. CalY works as an adhesin which binds to host cells, and strongly contributes to the bacterial virulence.

Right panel: later in the stationary phase or in biofilms, SipW is activated by an unknown, hypothetical factor and cleaves CalY signal peptide. The free, extracellular CalY is activated by an undetermined, cell-surface activating factor (not shown here), and polymerizes to produce amyloid fibers promoting biofilm formation.



CalY is a biofilm protein produced in high quantities. We found that in early stationary phase, this protein is located at the cell surface where it promotes the bacterium binding to host cells. Later, in mid- or late-stationary phase, CalY is released in the extracellular medium by the signal peptidase SipW and polymerizes as fibers promoting biofilm formation.