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# Visualizing Spore Germination Protein Complexes in Action in Bacterial Endospores; Lessons and Challenges Ahead

[Swammerdam Institute for Life Sciences - SILS - University of Amsterdam](#)

Molecular Biology and Microbial Food Safety

Stanley Brul head of laboratory with the input from many PhD students and PETER SETLOW UConn Health

# Bacterial spores; the good are inhabitants of our intestines!

LETTER

OPEN

doi:10.1038/nature17645

## Culturing of ‘unculturable’ human microbiota reveals novel taxa and extensive sporulation

Hilary P. Browne<sup>1\*</sup>, Samuel C. Forster<sup>1,2,3\*</sup>, Blessing O. Anonye<sup>1</sup>, Nitin Kumar<sup>1</sup>, B. Anne Neville<sup>1</sup>, Mark D. Stares<sup>1</sup>, David Goulding<sup>4</sup> & Trevor D. Lawley<sup>1</sup>

See also the Holomicrobiome Initiative

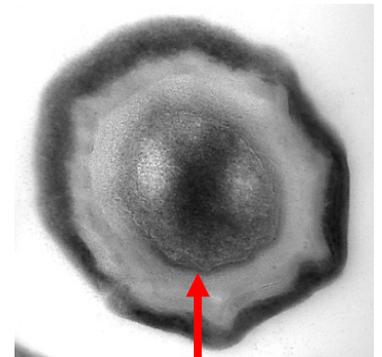
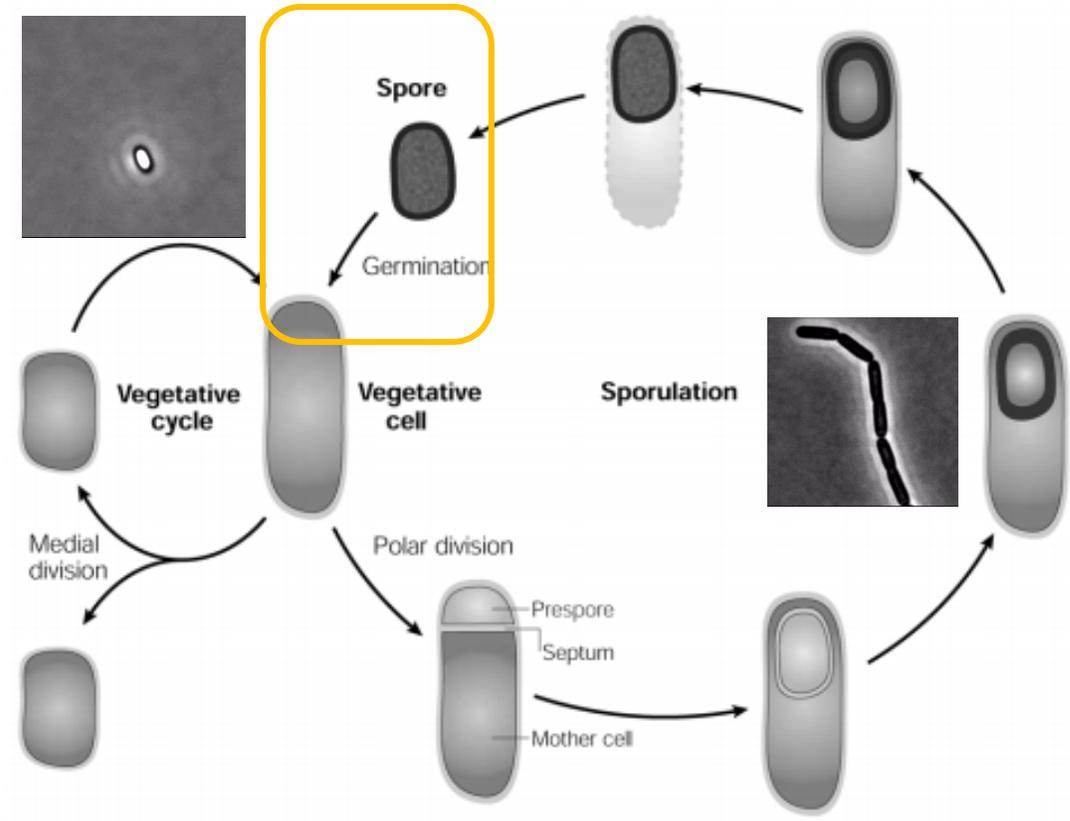
<https://holomicrobiom.nl/en/home-en/>

A Holland-based public-private partnership that aims to map, analyze and model the many contacts between microbiota in Water, Soil-Plant, Animals and Humans

and the Research Priority Area Personal Microbiome Health <https://pmh-uva.nl/>

[Full professor in Microbiome Engineering](#) vacancy open @ SILS-UvA

Main interest in *Bacillus subtilis* and *Bacillus cereus* as a toxigenic spore former that causes food borne disease upon spore germination and outgrowth.



Inner membrane with GRs

the life cycle of spore-forming bacteria

# Germinant Receptor proteins (GRs) were initially visualized @ super-resolution in *B. subtilis*

jove Journal of Visualized Experiments

www.jove.com

Video Article

## Visualization of Germinosomes and the Inner Membrane in *Bacillus subtilis* Spores

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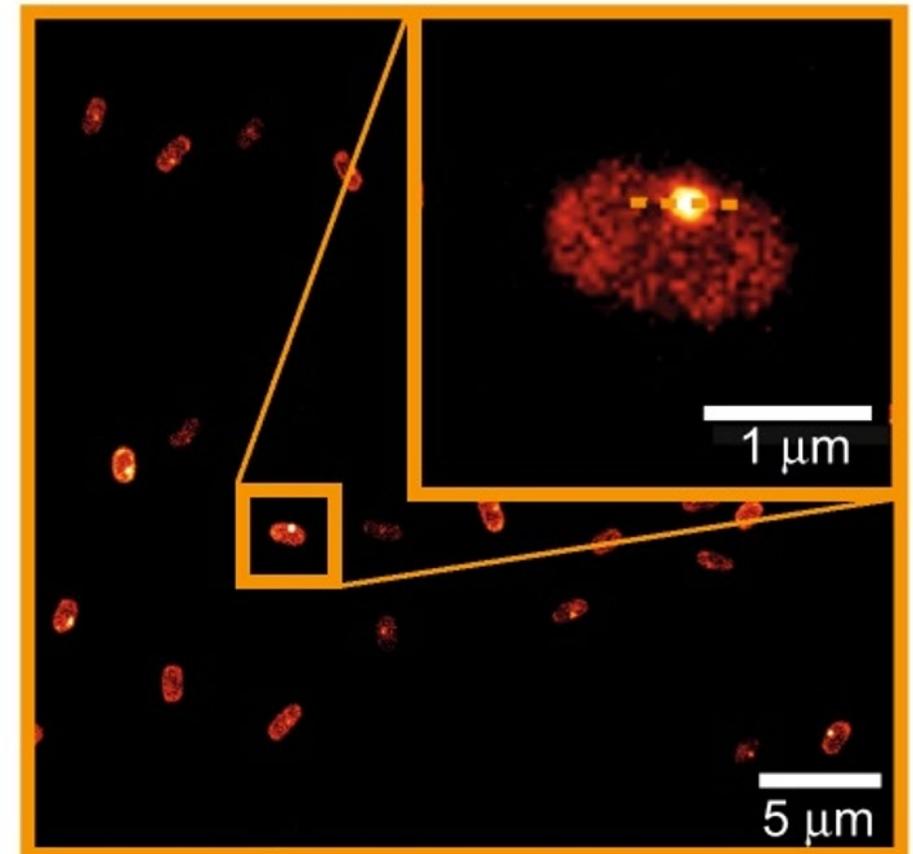
SCIENTIFIC  
REPORTS  
nature research

OPEN

## A live-cell super-resolution technique demonstrated by imaging germinosomes in wild-type bacterial spores

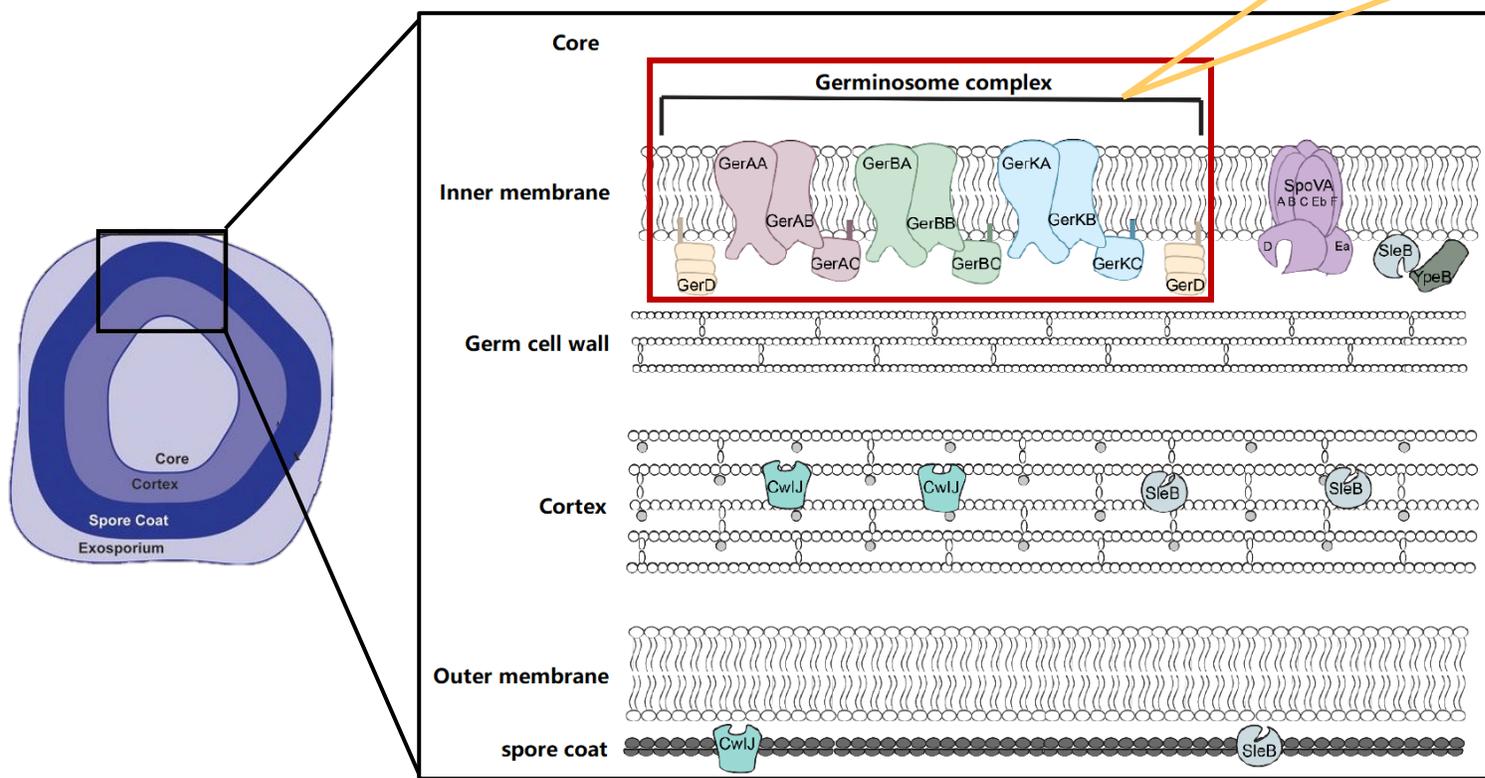
R. M. P. Breedijk<sup>1,6</sup>, J. Wen<sup>2,6</sup>, V. Krishnaswami<sup>1,6</sup>, T. Bernas<sup>3</sup>, E. M. M. Manders<sup>1,4</sup>, P. Setlow<sup>5</sup>, N. O. E. Vischer<sup>2</sup> & S. Brul<sup>2\*</sup>

A germinant receptor protein cluster (germinosome) as shown by annular rescan confocal microscopy of GerKB-sGFP.



# Key germination proteins are clustered in germinosomes in *Bacillus subtilis* spores

How about *B. cereus* spores?



## Germinant receptors (GRs) (sense germinants\*)

- GerA     A / B / C subunit
- GerB     A / B / C subunit
- **GerK**     A / C / B subunit

## Germination protein GerD (as a scaffold)

- **mediates** clustering of germination proteins in germinosomes and
- **promotes** the rapid and cooperative GRs response to germinants!

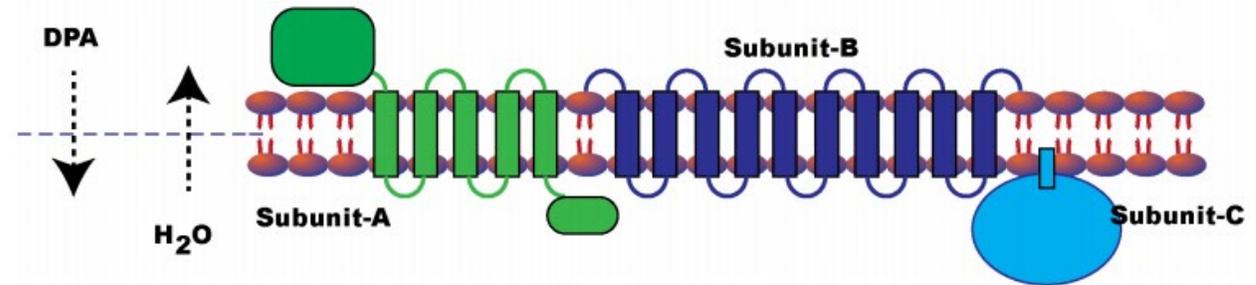
\*GR germinants are generally **amino acids**, sugars or purine nucleosides

*B. cereus* spores have seven different germinant receptors; we focussed on the visualization of the alanine binding tricistronic receptor GerR.

### Germinant receptors

- GerR A / C / B subunit
- GerK A / B / C subunit
- GerI A / B / C subunit
- GerS A / B / C subunit
- GerQ A / B / C subunit
- GerG ? / B / C subunit
- GerL ? / B / C subunit

[can sense L-alanine as a trigger of germination]



Schematic diagram of the topology of germinant receptor subunits in the inner membrane of *Bacillus* spores

### Germination scaffold protein: GerD

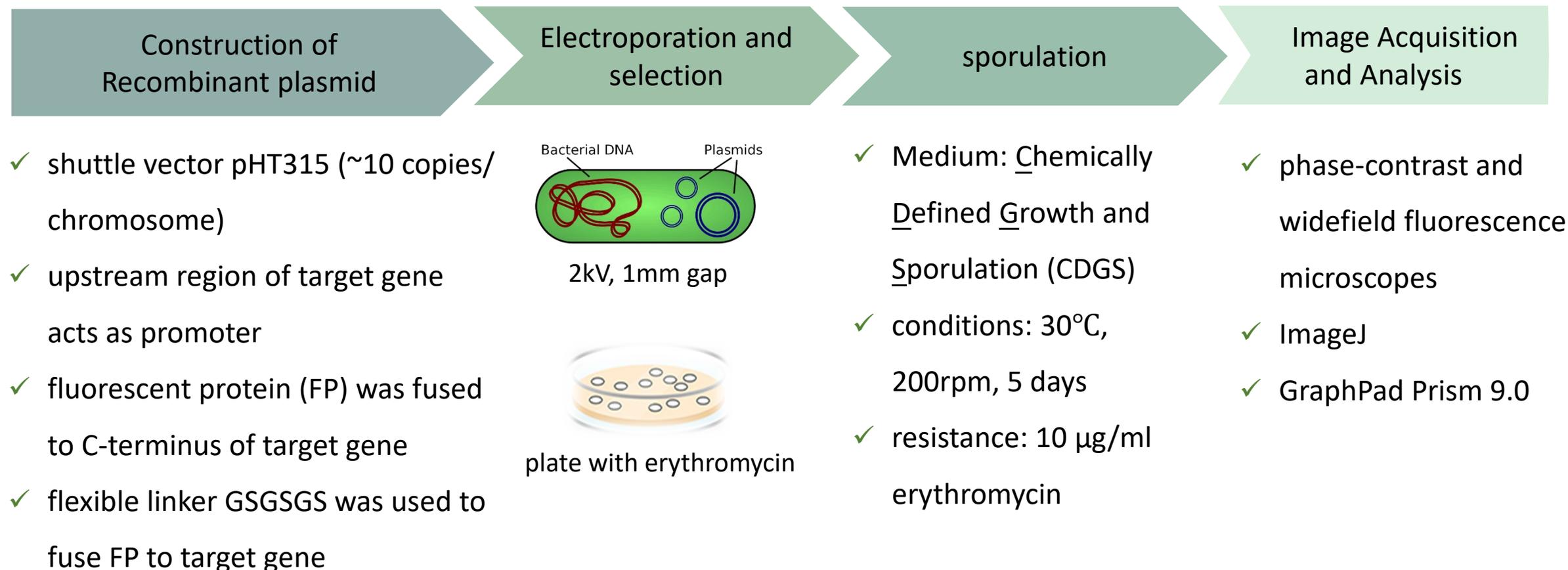
- Compared to GerD of *B. subtilis*: 43% identity in sequence of amino acid

# Research questions (I)

Visualization of germination proteins in putative *Bacillus cereus* germinosomes

Analysis of the interaction of germinant receptor proteins of the gerR operon and GerD in *Bacillus cereus* spores using FRET

# How to visualize GRs and SpoVAEa proteins in *B. cereus* spores ?

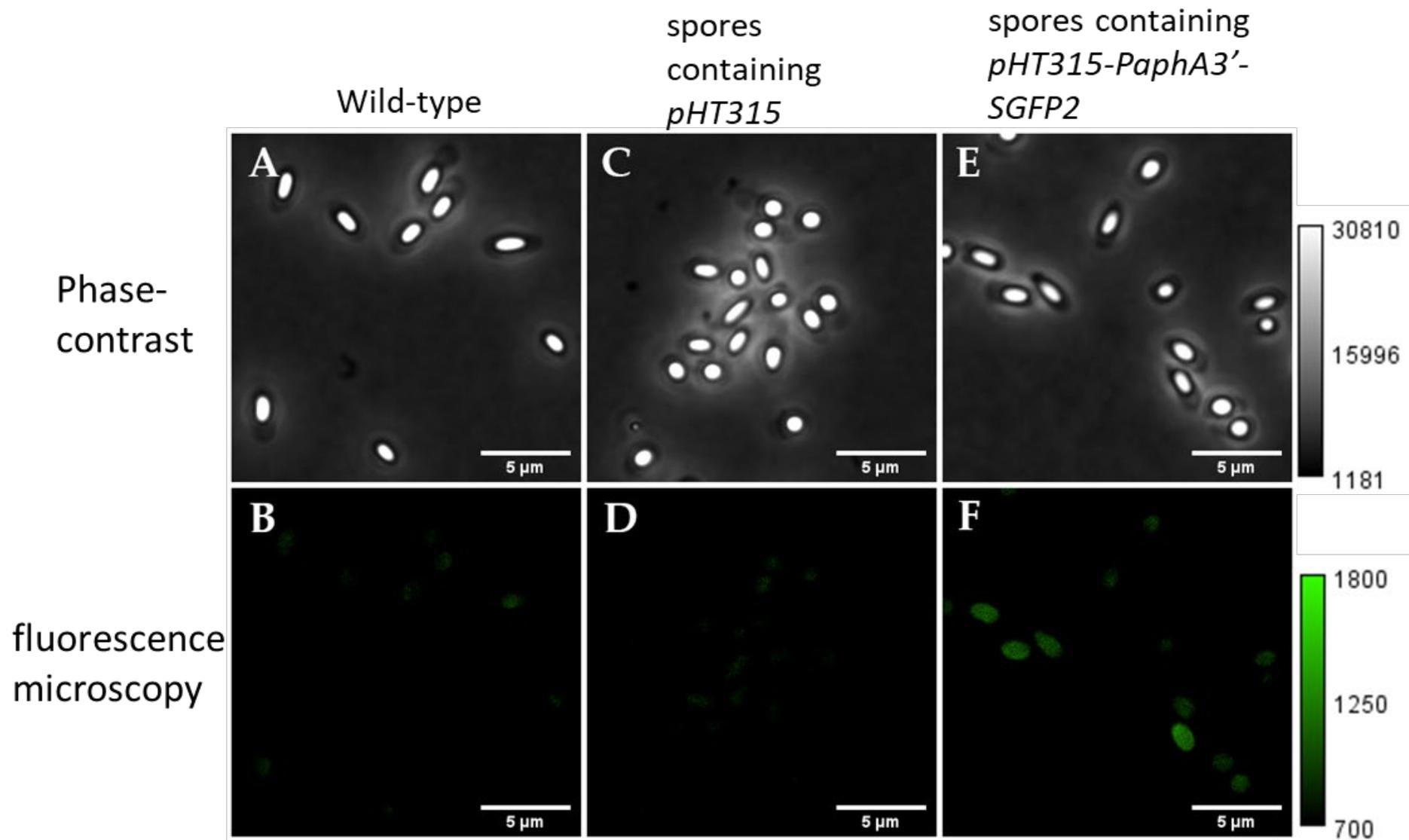


# Research questions (I)

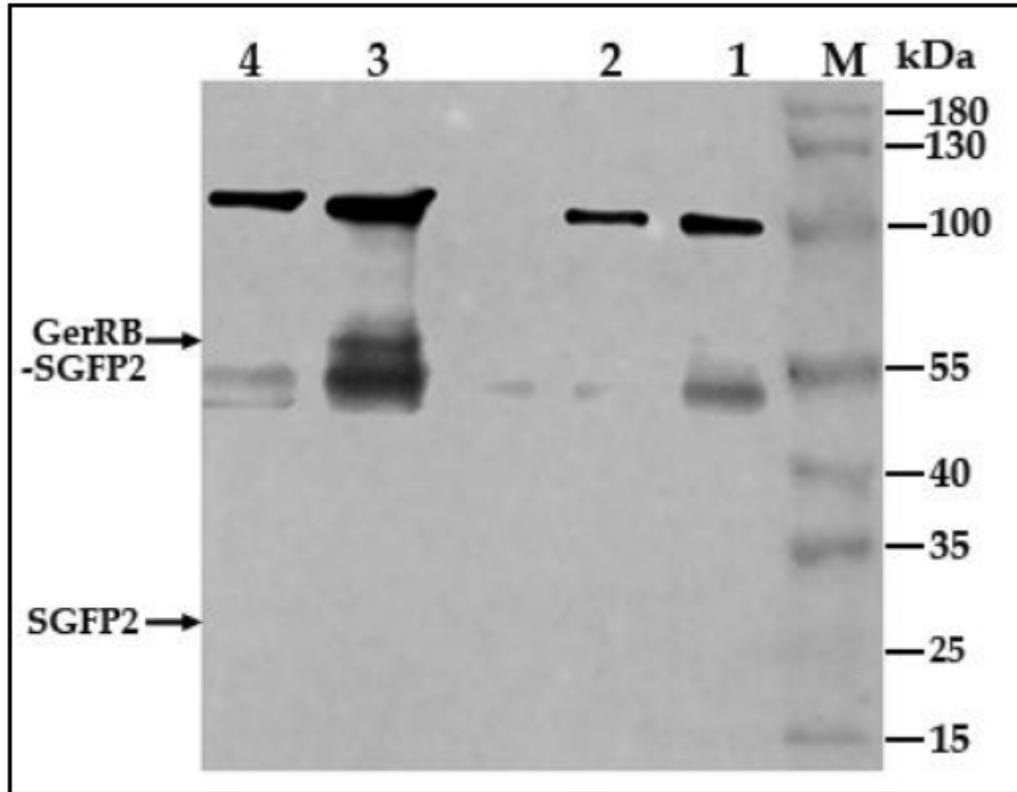
Visualization of germination proteins in putative *Bacillus cereus* germinosomes

Analysis of interaction of germination protein GerR and GerD in *Bacillus cereus* spores using FRET

# Visualization of SGFP2 protein expression from *pHT315* in *B. cereus* spores



Recombinant spores express fusion protein GerRB-SGFP2 encoded from *pHT315* in *B. cereus* spores



lane 1, wild-type spores;

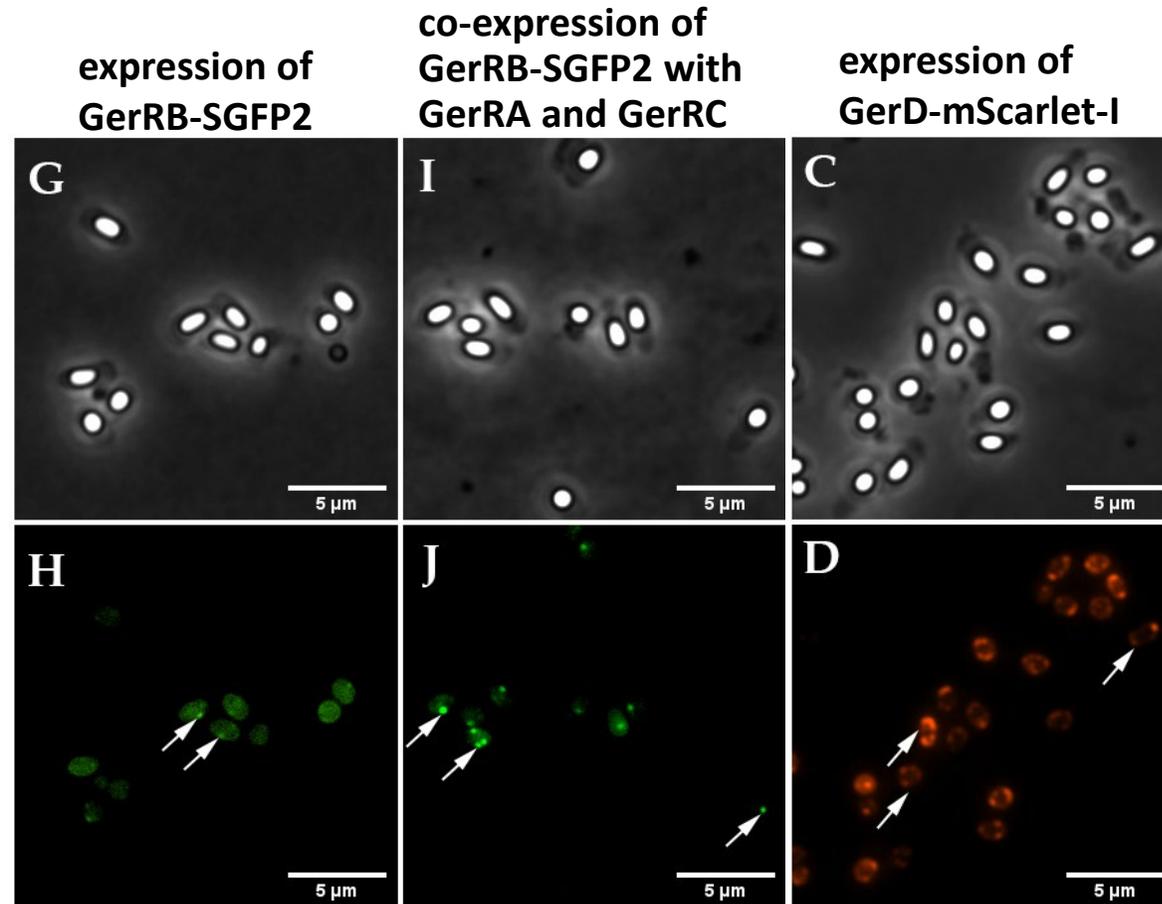
lane 2, spores carrying pHT315;

lane 3, spores carrying pHT315-PgerR-gerRB-SGFP2;

lane 4, spores carrying pHT315-gerRB-SGFP2;

Lane M, prestained protein ladder

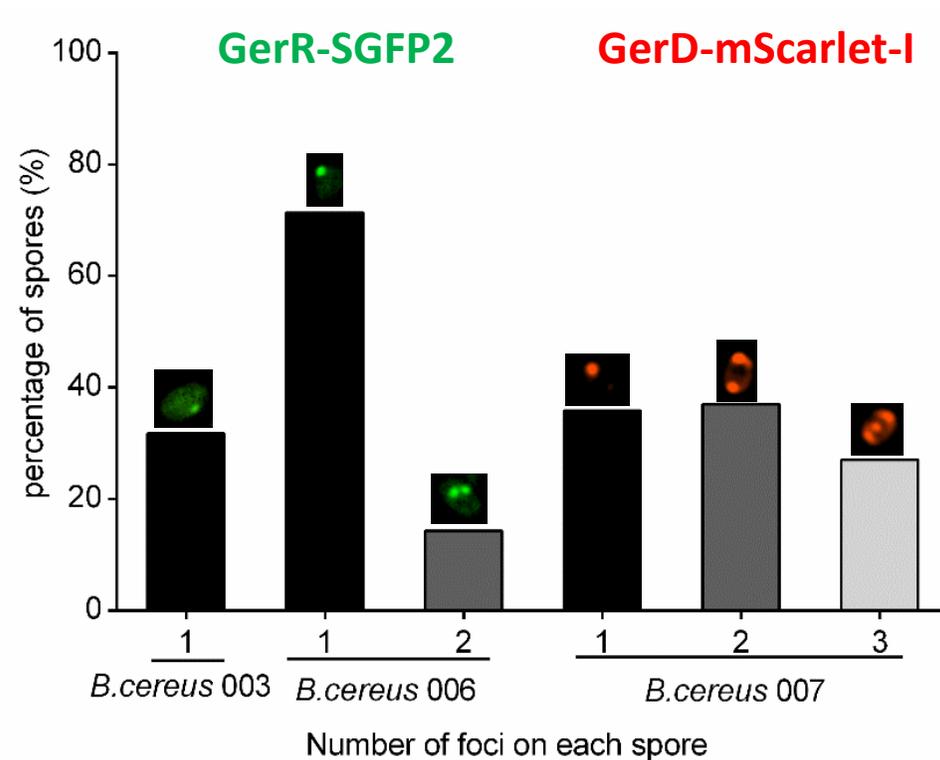
## First time visualization of germination proteins in *B. cereus*



These data of visualization suggested that co-expression of GR subunits improves their stability and are the first evidence for the existence of germinosomes in spores of the pathogen *B. cereus*.

## Percentage of germinosomes in *B. cereus* spores with GerRB-SGFP2 or GerD-mScarlet-I

- **expression of only GerRB-SGFP2 (*B. cereus* 003):**  
one germinosome was observed
- **co-expression of GerRA and GerRC (*B. cereus* 006):**  
one and two germinosomes were observed
- **expression of GerD-mScarlet-I (*B. cereus* 007):**  
one, two and three germinosomes were observed



# CONCLUSIONS

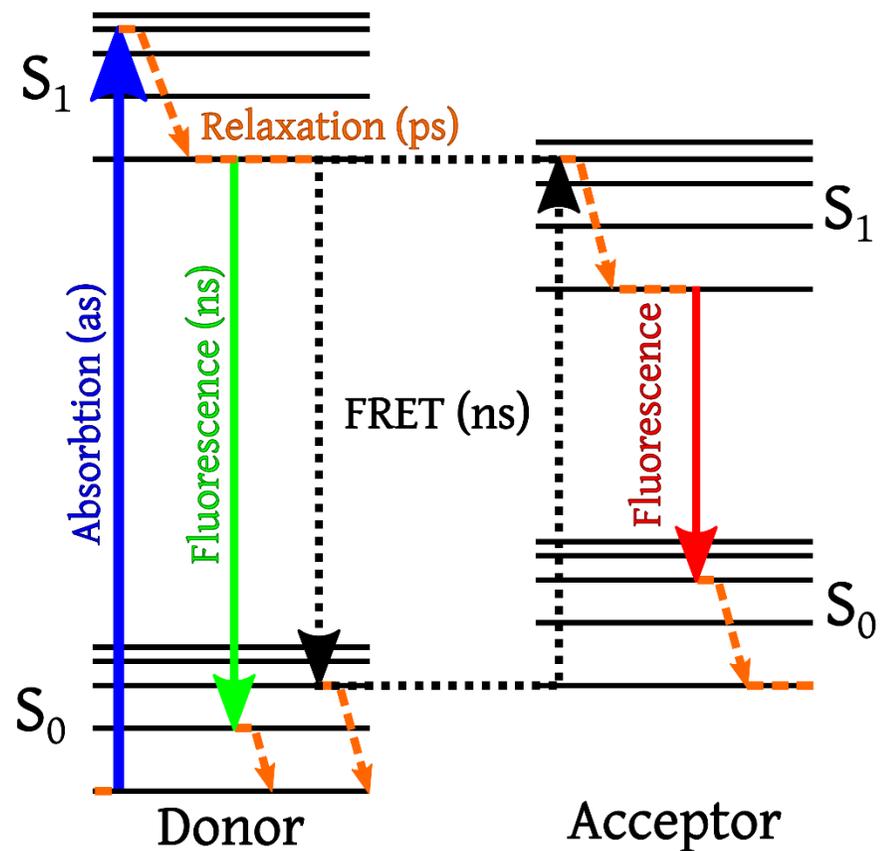
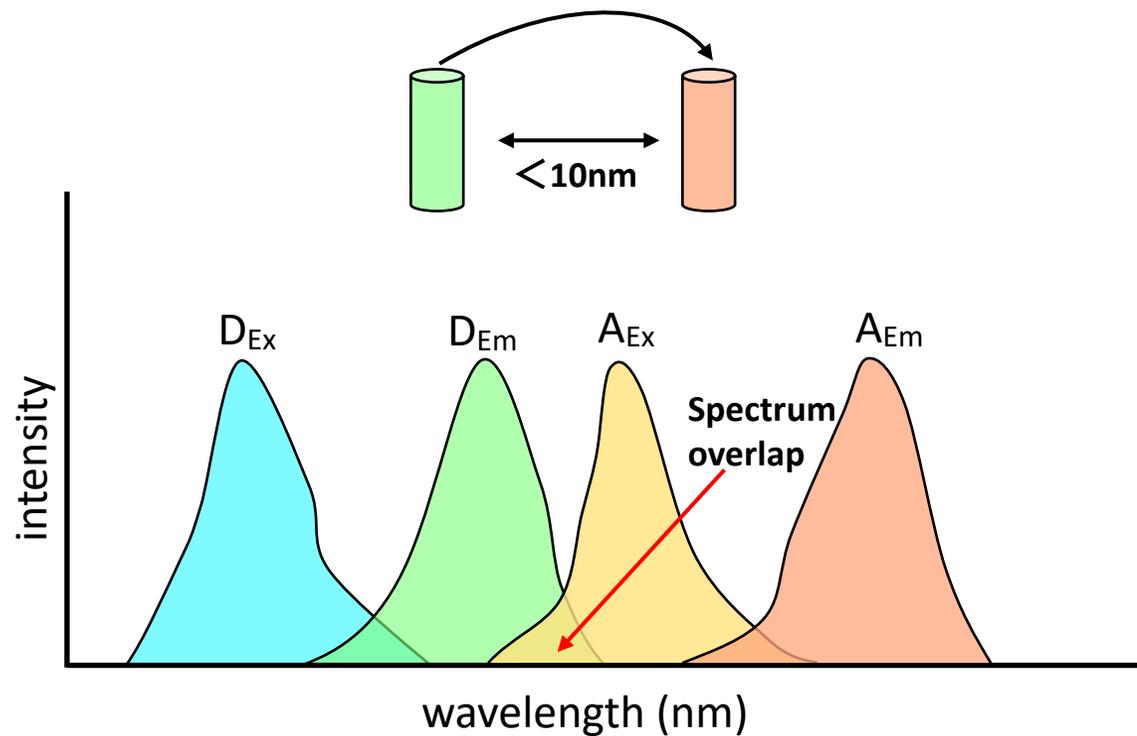
- High resolution wide-field fluorescence microscopy visualized GerRB-SGFP2 specific bright foci (germinosomes) in ~30% of individual dormant spores if only GerRB-SGFP2 was expressed,
- but in ~85% of spores upon co-expression with GerRA and GerRC. Our data corroborates the notion that co-expression of GR subunits improves their stability.
- All spores displayed bright fluorescent foci upon expression of GerD-mScarlet-i under the control of the *gerD* promoter.

# Research questions (I)

Visualization of germination proteins in putative *Bacillus cereus* germinosomes

Analysis of interaction of germination protein GerR and GerD in *Bacillus cereus* spore putative germinosomes using FRET.

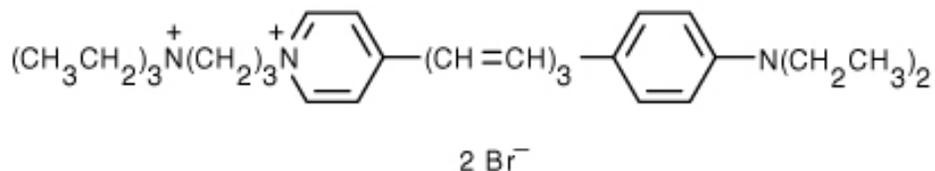
# Fluorescence Resonance Energy Transfer (FRET) analysis principles



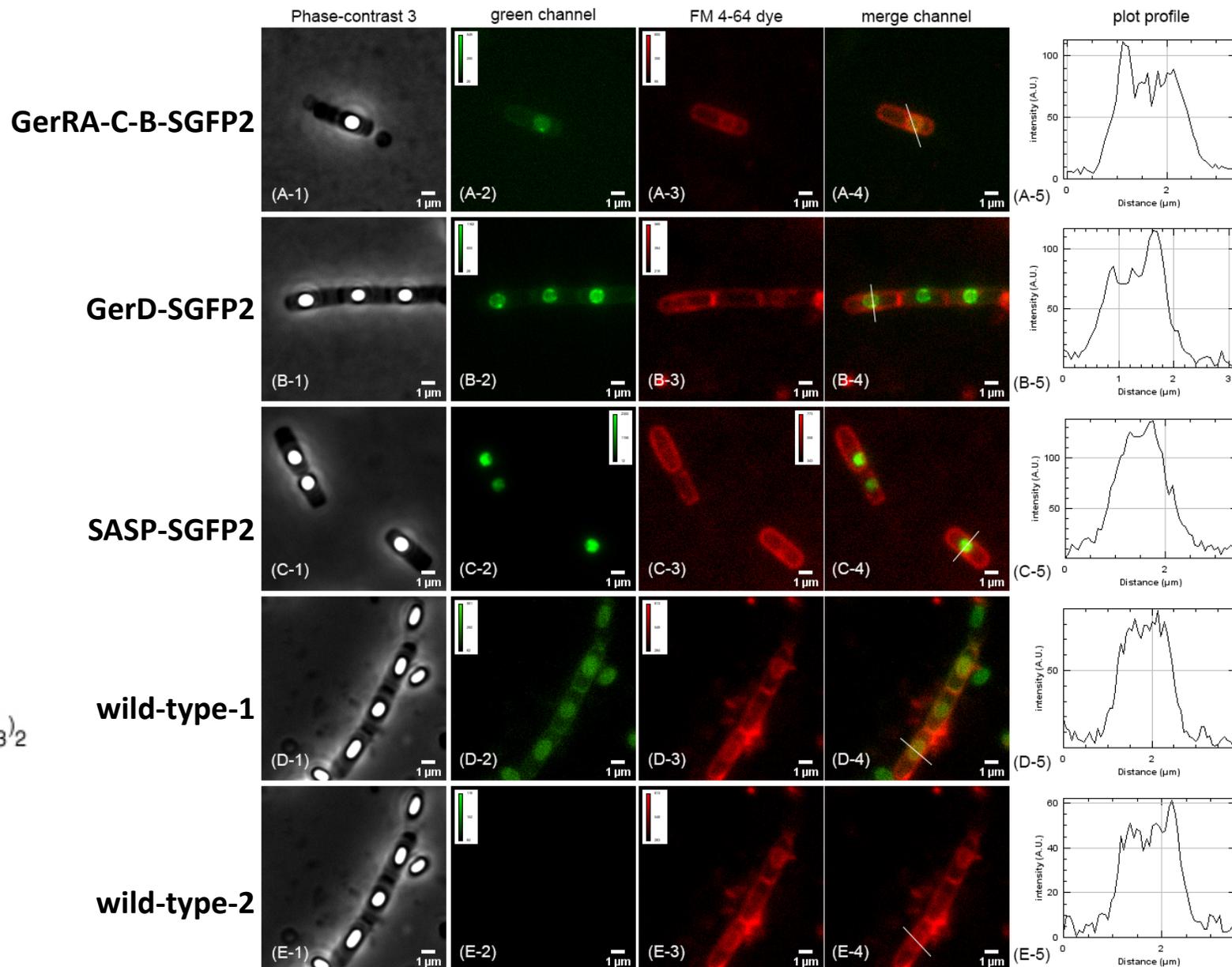
Jablonski diagram illustrating the FRET process. Note that the black dashed line indicates a virtual photon.

## *B. cereus* forespore membrane stained with FM 4-64 indeed harbours GerRB and GerD

- FM-4-64 stains the spore membrane during the sporulating process.
- The germination receptor gerRB and GerD are present in the forespore membrane.
- SASP are, in contrast, as expected localized in the core of forespore.



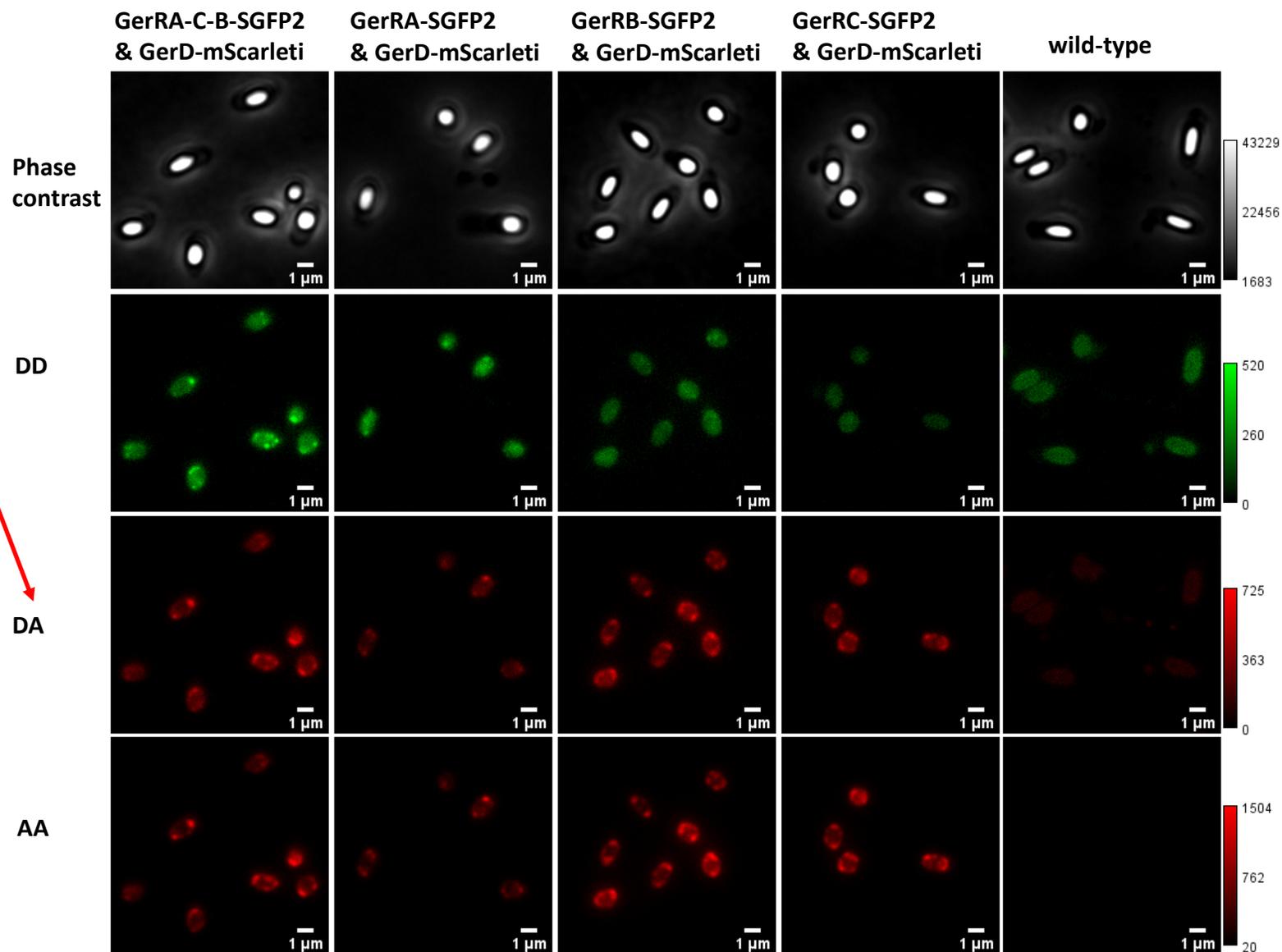
the chemical structure of FM 4-64



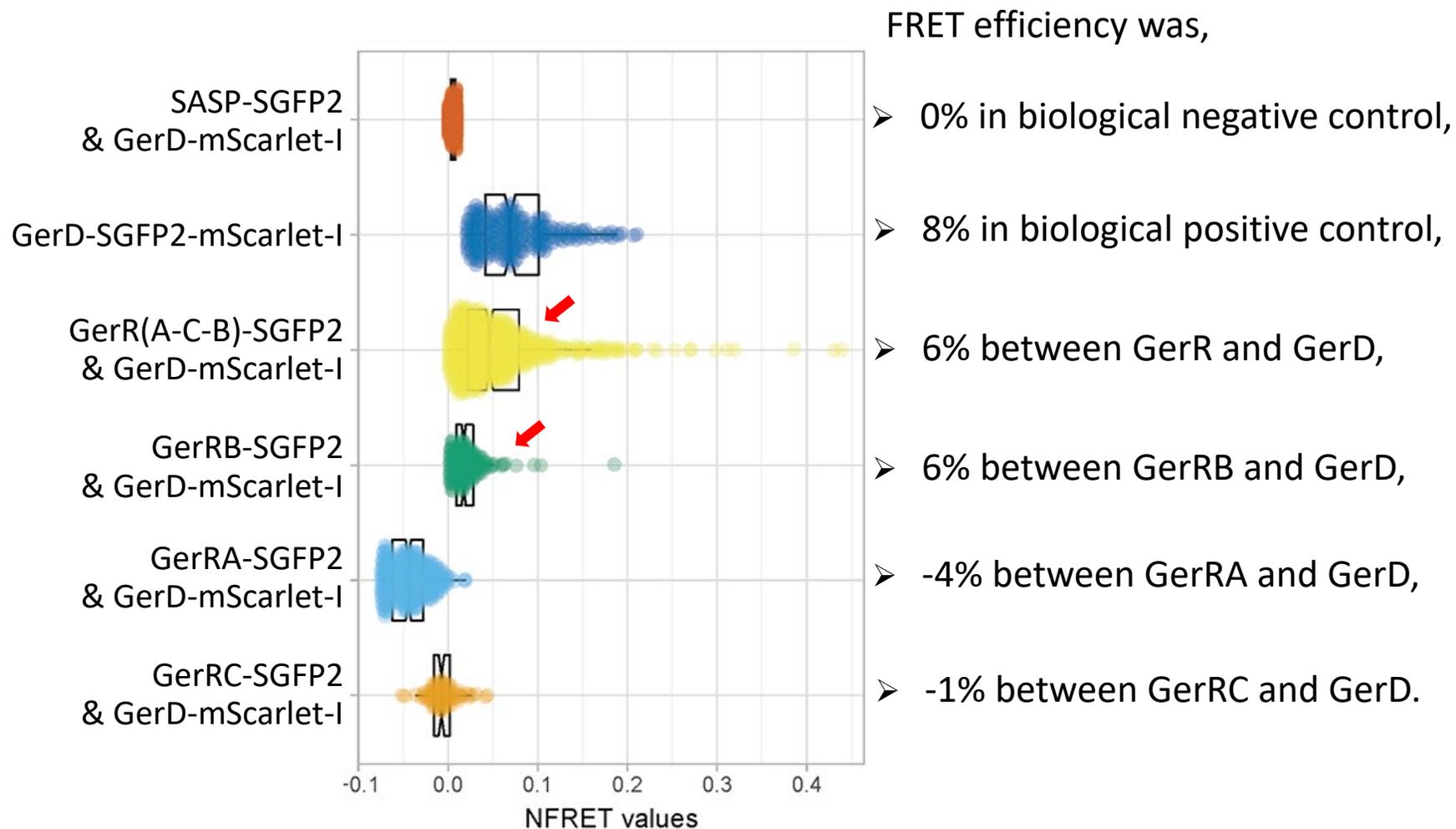
# Protein-protein interactions in the inner spore membrane visualized through FRET

## Definition

Channels	Excitation	Emission
Donor channel (DD)	Donor	Donor
FRET channel (DA)	Donor	Acceptor
Acceptor channel (AA)	Acceptor	Acceptor



# Quantification of the FRET results of interaction between GerR and GerD



**Summary:** GerD mainly interacts with the GerR B subunit, not with A or C- subunits.

## Conclusions:

- There is a close interaction (< 10 nm distance) between GerRB and the GerD protein in *B. cereus* germinosomes.
- The germinant receptor proteins from the GerR operon and GerD were localized in germinosomes in the innermembrane of forespores using reporter proteins and FM 4-64 staining.

# The major *Bacillus* spore germination proteins and their location in the inner membrane

## the inside core of the spore

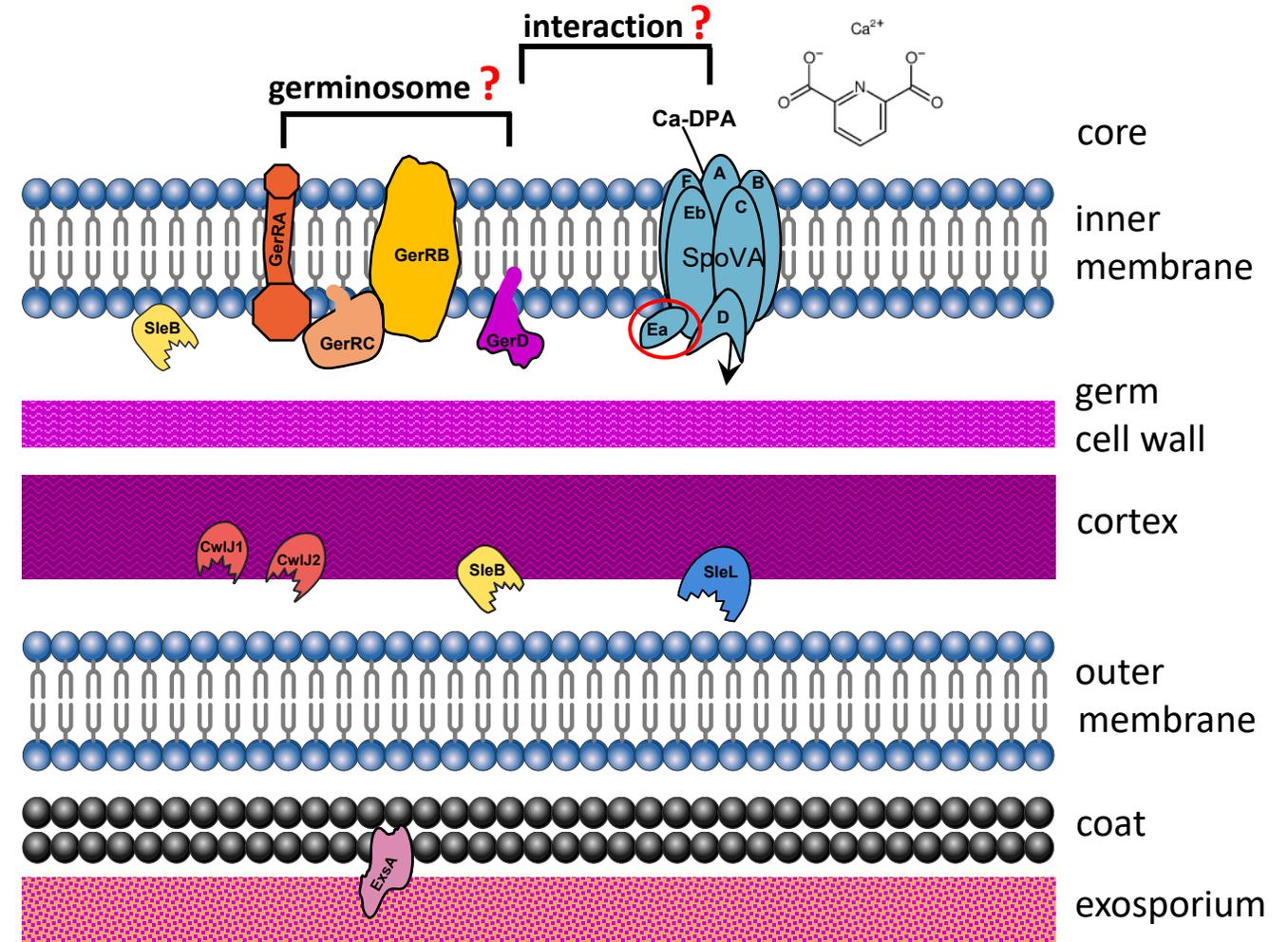
### Germinant receptors

□	<b>GerR</b>	<b>A / C / B subunit</b>
□	GerK	A / B / C subunit
□	GerI	A / B / C subunit
□	GerS	A / B / C subunit
□	GerQ	A / B / C subunit
□	GerG	? / B / C subunit
□	GerL	? / B / C subunit

### Germination scaffold protein: GerD

### SpoVA (stage V sporulation protein A)

A / B / C / D / Eb / **Ea** / F subunit



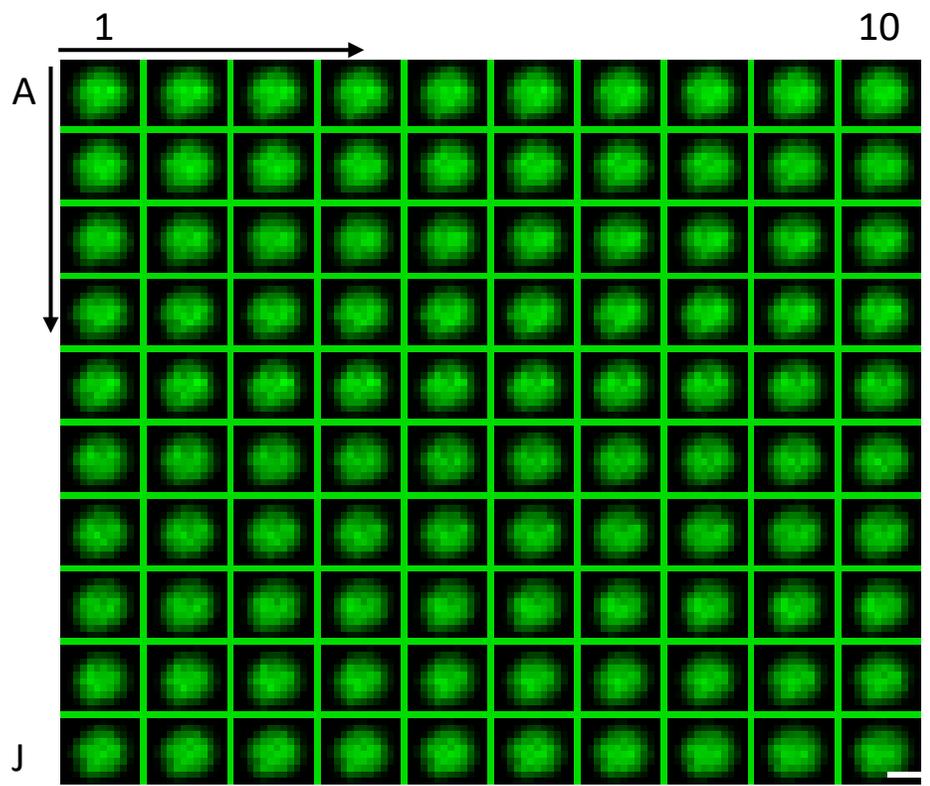
## Research questions (II)

Visualization of SpoVAEa protein dynamics in dormant *B. subtilis* & *B. cereus* spores

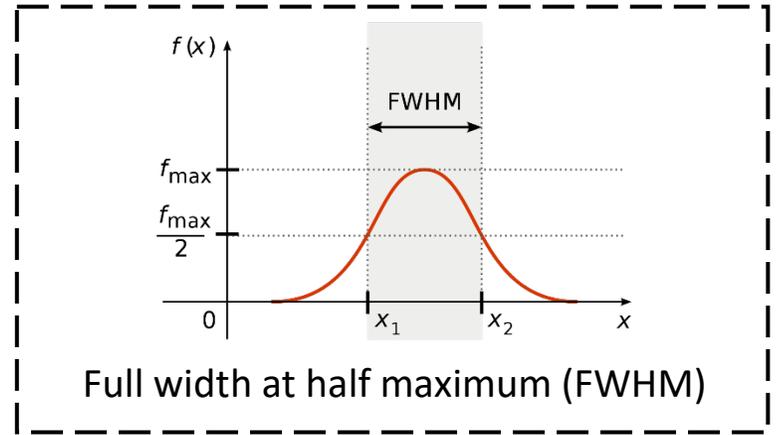
Dynamic changes in the germinosome and SpoVAEa during germination of *B. cereus* spores

# High frequency time-lapse images acquisition and analysis in *B. subtilis* and *B. cereus*

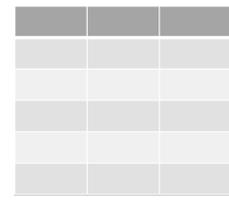
Capture condition: exposure 50ms,  
record 100 frames and no delay between each frame



As an example frames of *B. cereus* spore with montage



Measure the FWHM of each frame  
with plugin Adrian's FWHM in ImageJ



Layout of results

Calculate the percent change of FWHM

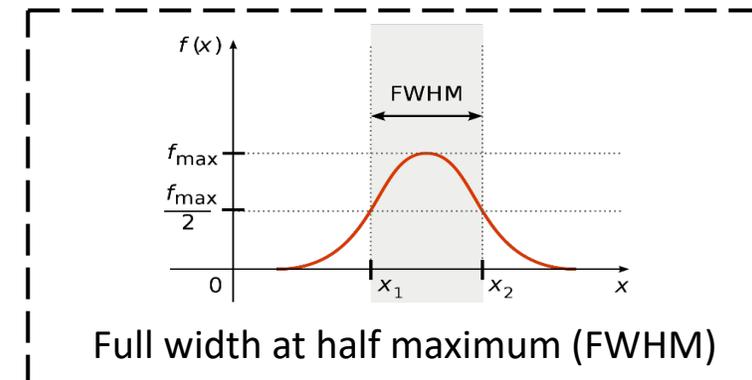
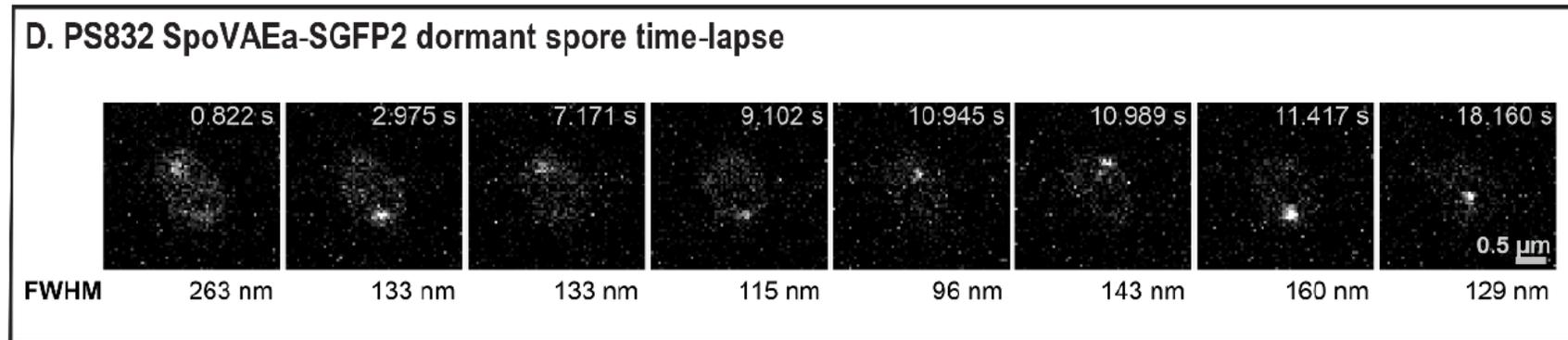
$$\frac{(A2 - A1)}{A1}$$

(A3, A4, A5,.....)

# Super-resolution imaging of high frequency movement of SpoVAEa in *B. subtilis* spores

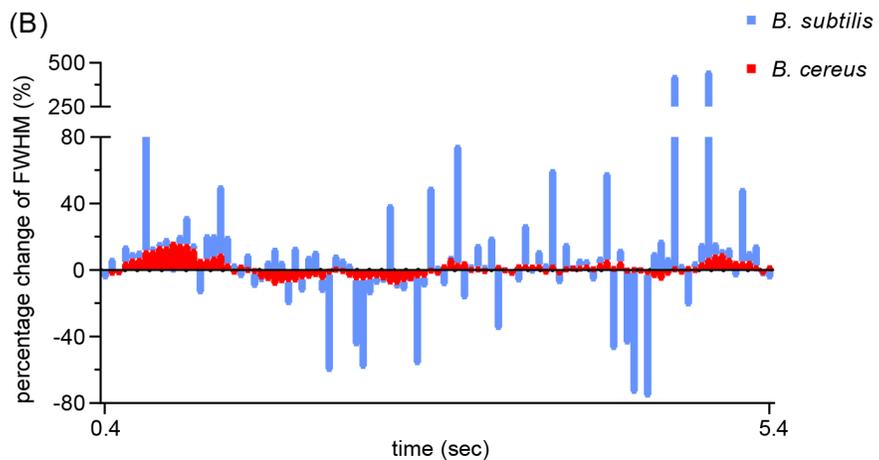
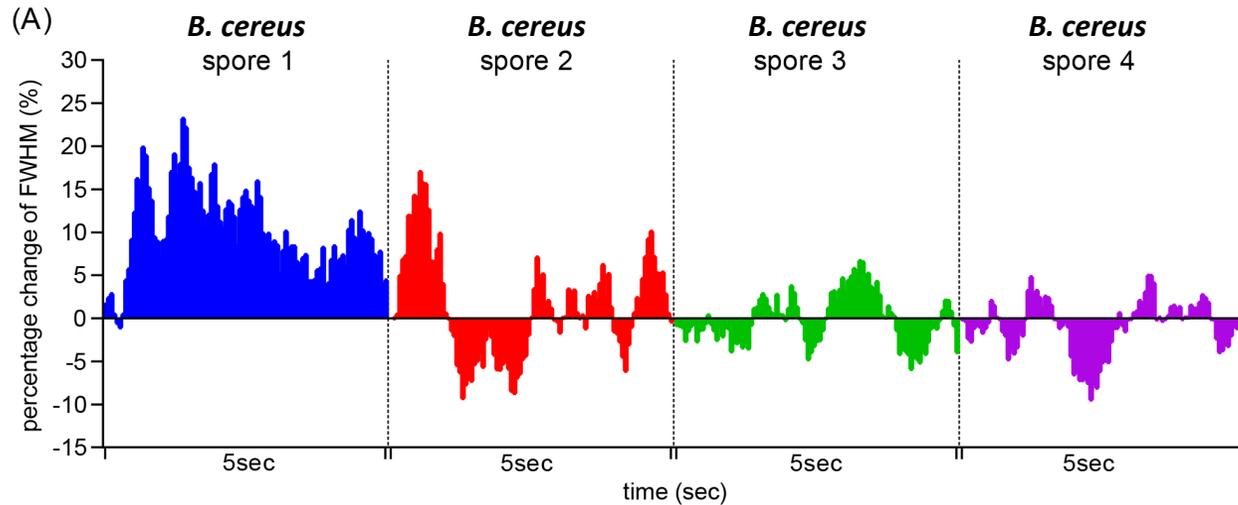
The RCM high frequency time lapse images (22.2 ms/frame) of the SpoVAEa-SGFP2 fusion protein of genomic expression in *B. subtilis* dormant spore.

RCM= Rescan Confocal Microscopy



**Published:** Wen J, Vischer NOE, de Vos AL, Manders EMM, Setlow P and Brul S.  
 Organization and dynamics of the SpoVAEa protein and its surrounding inner membrane lipids, upon germination of *Bacillus subtilis* spores. *Scientific Reports*(2022) 12:4944  
[doi.org/10.1038/s41598-022-09147-3](https://doi.org/10.1038/s41598-022-09147-3)

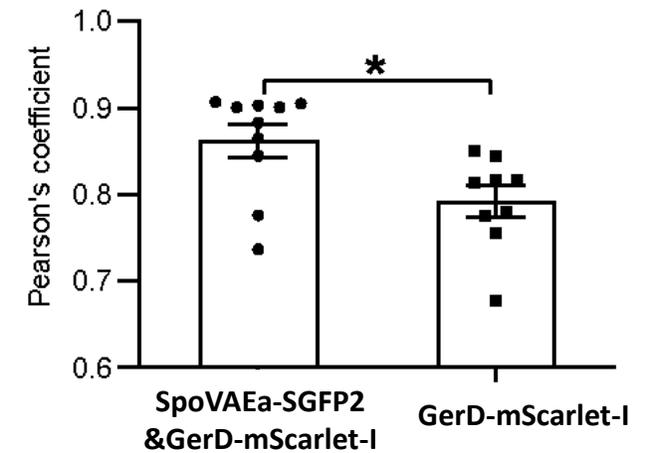
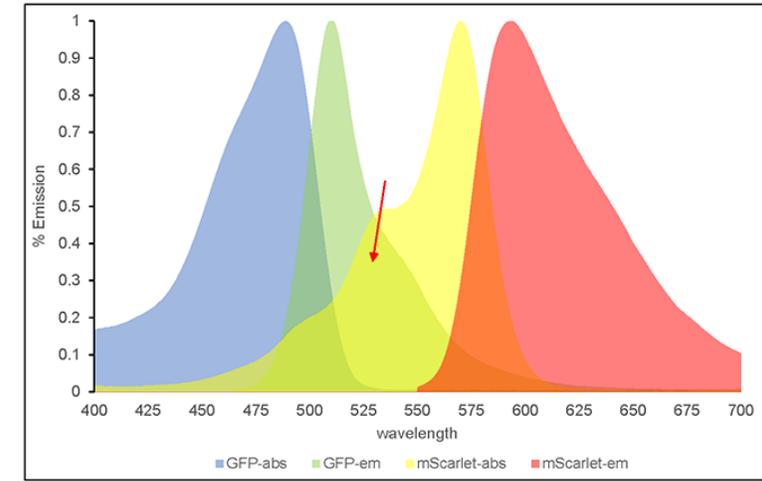
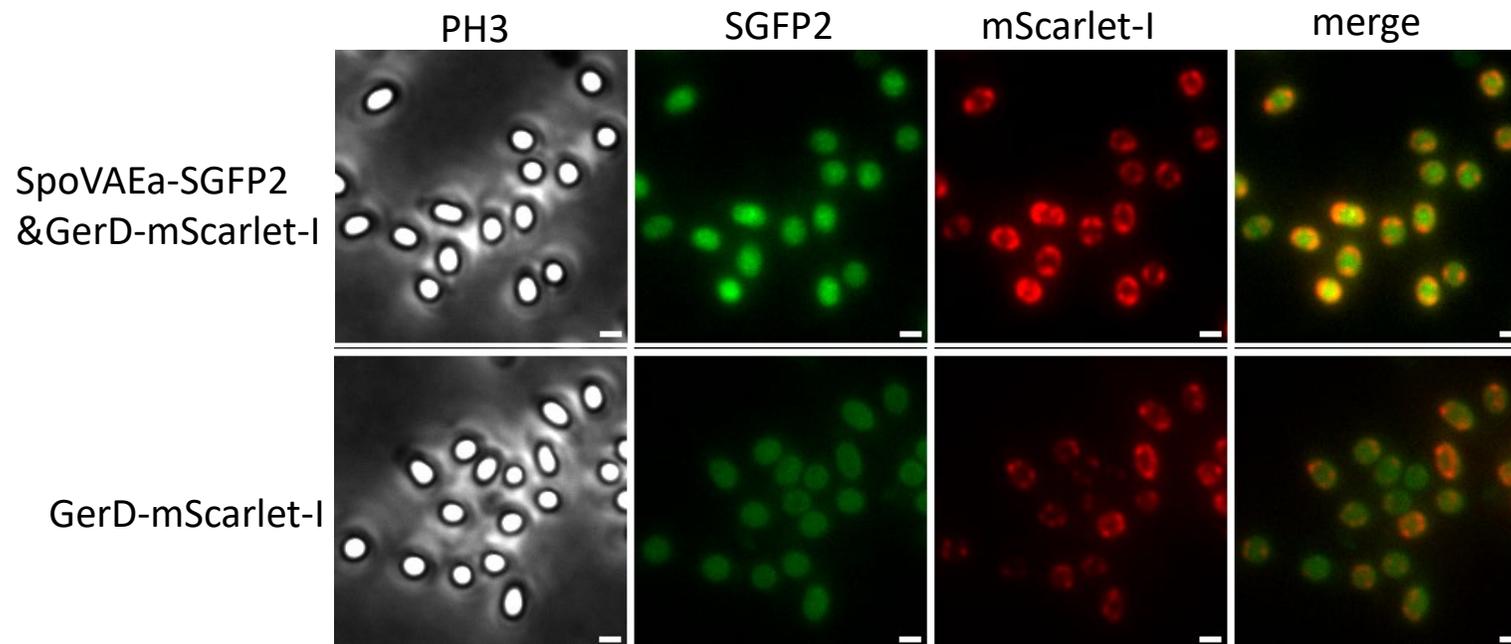
# Comparing the mobility of SpoVAEa foci in dormant spore of *B. cereus* and *B. subtilis*



## Summary:

- In *B. cereus* (panel A,B) and *B. subtilis* (panel B) the SpoVAEa foci of different spores behave differently.
- SpoVAEa fluorescent foci of *B. subtilis* spores\* redistribute at a higher frequency than those of *B. cereus*.
- This difference may be due to the fact that we look at different species with a different protein complement and at genomic expression in *B. subtilis* versus overexpression from a plasmid in *B. cereus*.

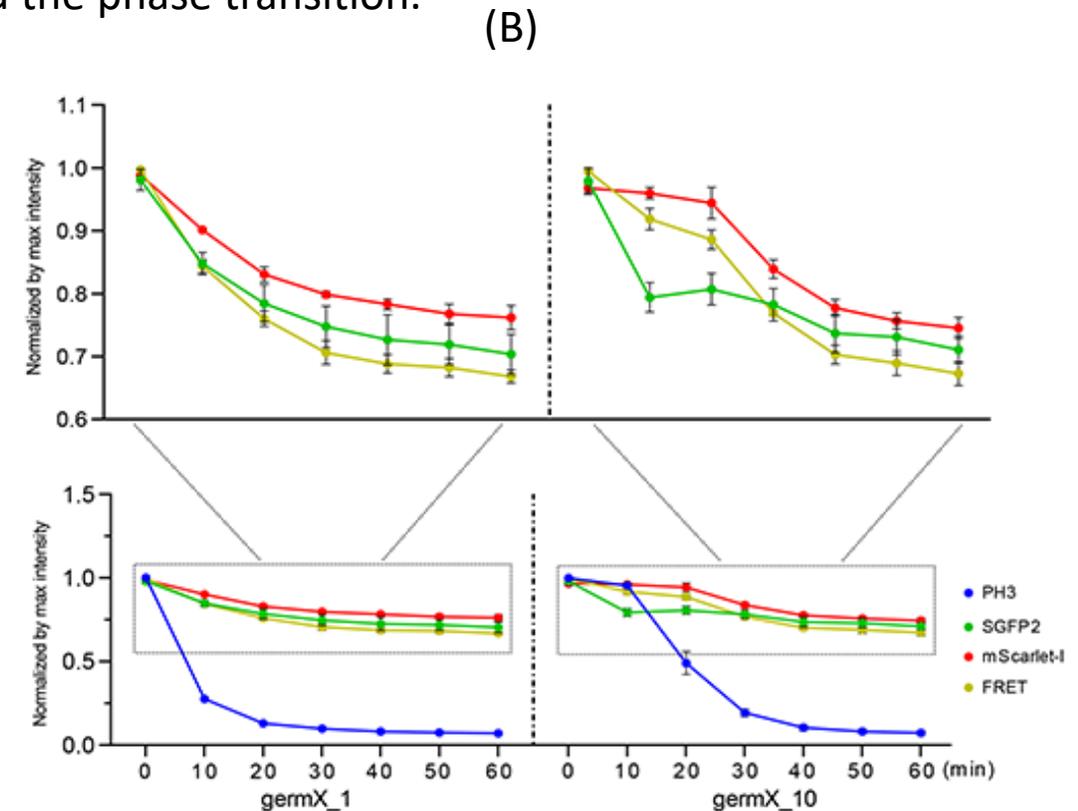
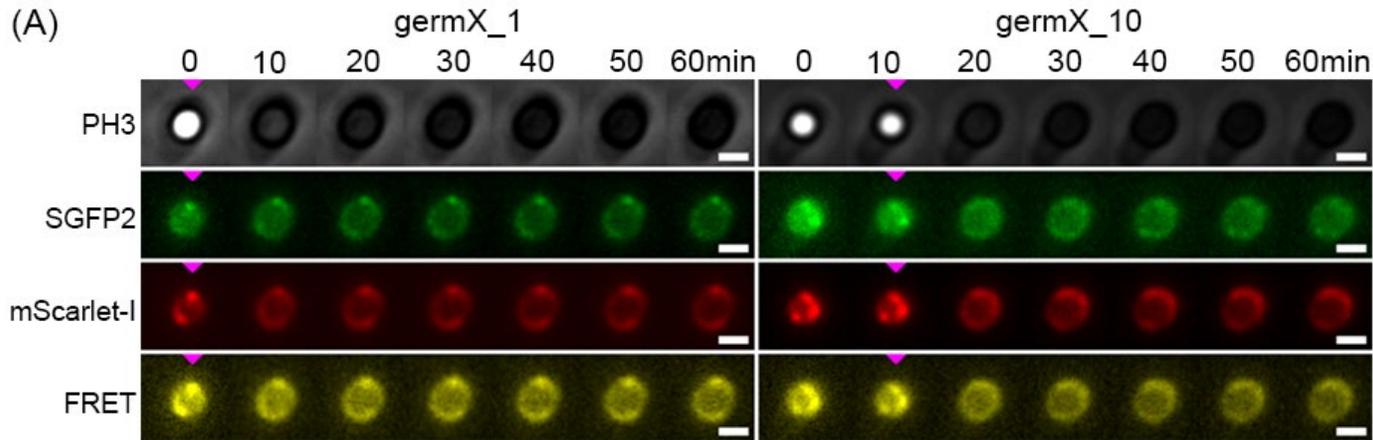
# Co-localization shown between SpoVAEa and germinosome GerD?



# Dynamics of germinant receptor foci FRET and fluorescence upon germination triggering by L-alanine

## Germinosome FRET foci GerRB-SGFP2 GerD-mScarlett *in B. cereus* spores

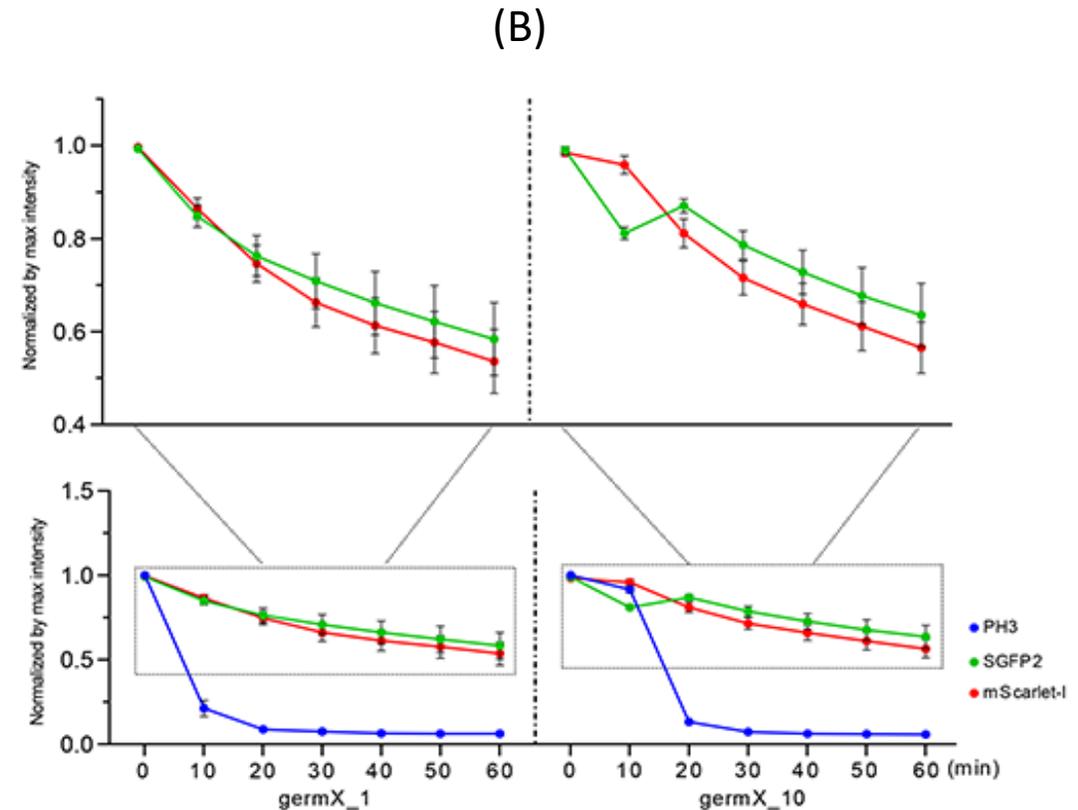
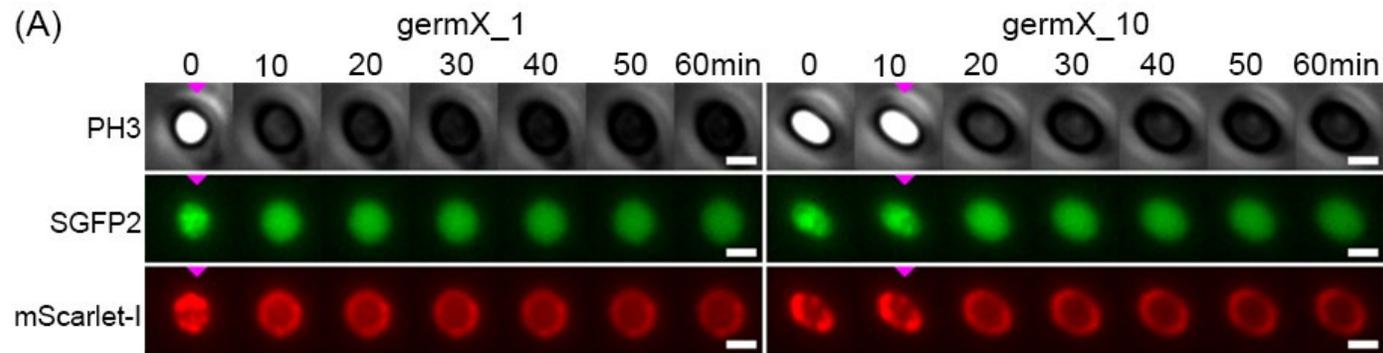
- were lost earlier than loss of SGFP2 foci and mScarlett-I foci,
- were lost to a large extent upon the phase transition,
- but the fluorescent germinosome foci as such persist well beyond the phase transition.



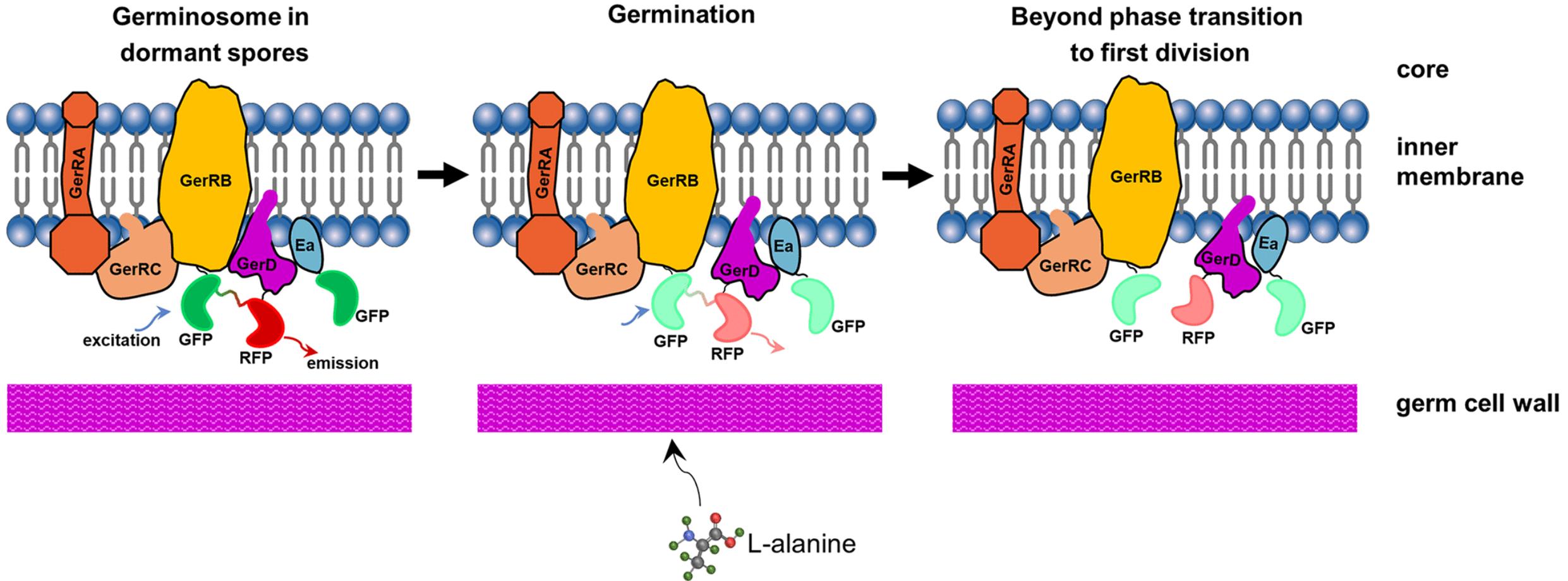
# Dynamics of SpoVAEa and GerD proteins upon germination triggered by L-alanine in *B. cereus* spores

**SpoVAEa-SGFP2 foci were lost upon phase transition.**

**GerD-mScarlet-I foci spread out and continued to exist beyond phase transition.**

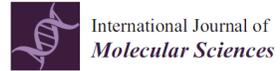


A model for the sequence of events during L-alanine induced spore germination in *Bacillus cereus*.



**Published:** Wang Y, Vischer NOE, Wekking D, Bogian A, Setlow P and Brul S. Visualization of SpoVAEa protein dynamics in dormant spores of *Bacillus cereus* and dynamic changes in their germinosomes and SpoVAEa during germination. (2022) Microbiology Spectrum 10 (3) [doi.org/10.1128/spectrum.00666-22](https://doi.org/10.1128/spectrum.00666-22)

# Back to our model *Bacillus subtilis*:



Article

## Predicting the Structure and Dynamics of Membrane Protein GerAB from *Bacillus subtilis*

Sophie Blinker<sup>1</sup>, Jocelyne Vreede<sup>2,\*</sup>, Peter Setlow<sup>3</sup> and Stanley Brul<sup>1,\*</sup>

# Molecular modelling of the B-subunits' structure & experimental mutant analysis

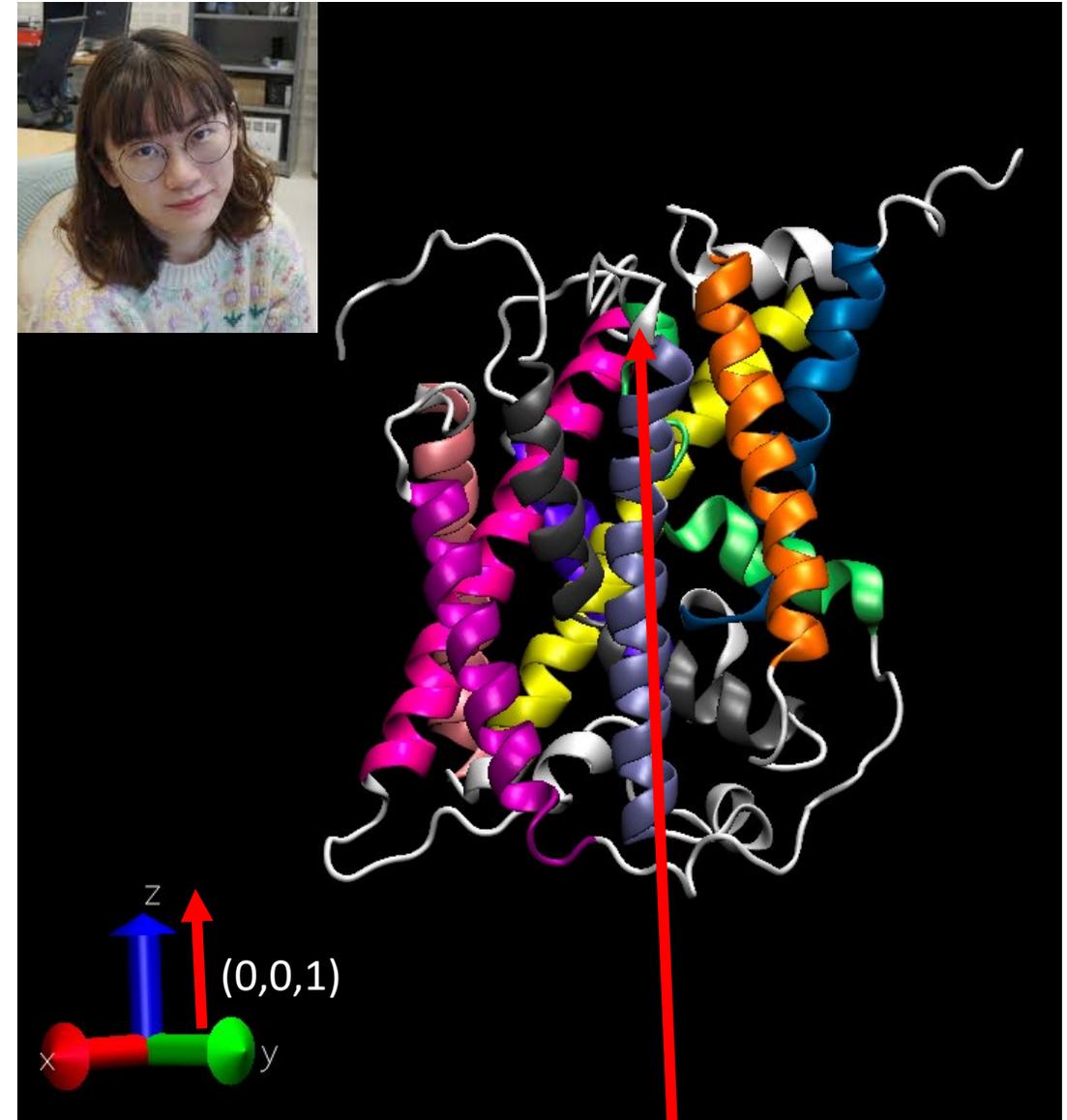
ARTICLE



<https://doi.org/10.1038/s41467-021-27235-2> OPEN

## Dormant spores sense amino acids through the B subunits of their germination receptors

Lior Artzi<sup>1</sup>, Assaf Alon<sup>2</sup>, Kelly P. Brock<sup>3</sup>, Anna G. Green<sup>3</sup>, Amy Tam<sup>3</sup>, Fernando H. Ramírez-Guadiana<sup>1</sup>, Debora Marks<sup>3</sup>, Andrew Kruse<sup>2</sup> & David Z. Rudner<sup>1</sup>



Steered Molecular Dynamics shows a putative water channel



# Take home messages and challenges ahead

- I. First evidence of the existence of germinosomes in spores of the pathogen *B. cereus*.
- II. GerD mainly interacts with GerR B subunit and not with A or C- subunits.
- III. Dynamics data showed that SpoVAEa moves on the surface of the inner membrane and it might play a role in signal transduction between the germinosome, possibly through GerD, and the SpoVA channel.
- IV. There are challenges ahead for bacterial spore germination analysis;
  - exact composition of each germinosome,
  - do GerAB subunits in *B. subtilis* contain a water channel-> Longjiao Chen
  - is such a putative channel conserved in other Bacilli and Clostridia,
  - how is the structure of gerAB, gerRB influenced by L-alanine binding, etc.



# Acknowledgements

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Prof. Peter Setlow

## **Molecular Dynamics @ FNWI UvA**

Dr. Jocelyne Vreede

## **LCAM-FNWI**

Dr. ir. Mark Hink

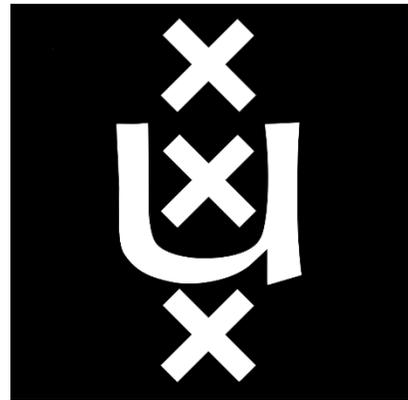
Ing. Ronald Breedijk



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NWO, UvA Research Priority area Systems Biology, Personal Microbiome Health, Urban Mental Health, NWA-ORC, CSC-fellowships, UvA-tenure track funds

Collaborate  
with us!



**Thank you for your attention.**

# Towards the understanding of the structure and function of germination receptor protein GerAB by combining computational and *in vivo* analyses.

Longjiao Chen<sup>1</sup>, Jocelyne Vreede<sup>2</sup>, Peter Setlow<sup>3</sup>, Stanley Brul\*<sup>1</sup>

<sup>1</sup> Molecular Biology and Microbial Food Safety Group, Swammerdam Institute for Life Sciences, University of Amsterdam, the Netherlands.

<sup>2</sup> Computational Chemistry Van 't Hoff Institute for Molecular Sciences, University of Amsterdam, the Netherlands.

<sup>3</sup> Molecular Biology and Biophysics, UCONN Health, USA.

\* s.brul@uva.nl

**Keywords:** germination, germinant receptor, water channel, *Bacillus subtilis*, molecular dynamics.

## **Abstract (<300 words):**

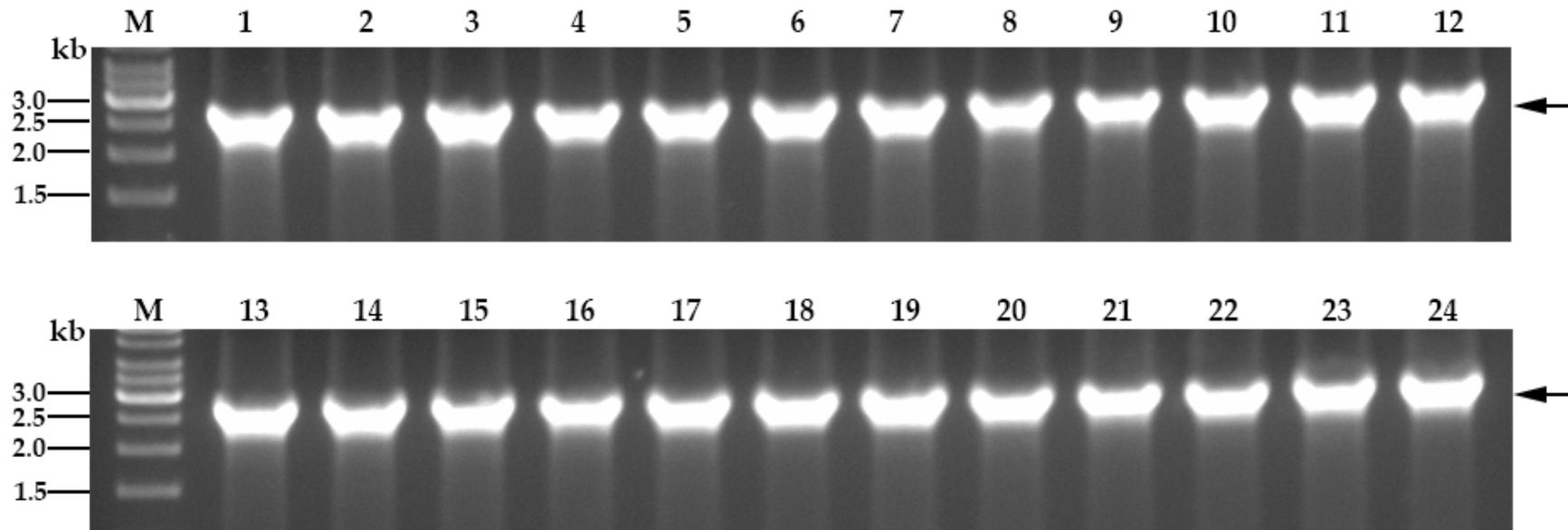
Some species in the *Bacillales* and *Clostridiales* bacterial orders form spores in unfavorable environments. These spores are metabolically dormant, resistant to harsh conditions, and their core, analogous to growing cells' protoplast, has a water content between 25-45% of wet wt compared to the ~80% in vegetative cells. However, upon germination by a variety of germinants, spores take up water relatively rapidly, restoring their water content to that of growing cells.<sup>1</sup> With no known functional water channels in the model spore-forming organism *Bacillus subtilis*, the molecular mechanism of spore water uptake during germination is unknown. Recent work showed that a subunit of the prototypical *Bacillus subtilis* spore germinant receptor GerA, the integral inner membrane protein GerAB, is the sensor initiating spore germination in response to L-alanine.<sup>2</sup> Notably previous work found that GerAB contains what appears to be a water channel.<sup>3</sup> Using Molecular Dynamic (MD) simulation methodology, we found water passing through the GerAB protein *in silico*, indicating the proteins putative water channel may well be functional.<sup>3</sup> At the same time, utilizing Steered MD simulation, we can now also pull single water molecules through the GerAB channel and therefore calculate the free energy of water permeation. These computational methods, as well as the predicted GerAB structure, have provided us with indications of GerAB residues that may be crucial in the water channel's function, and thus suggest mutagenesis experiments to experimentally test *in vivo* the computational modelling predictions. Here we present both the *in silico* and *in vivo* data that reinforce each other and thus jointly help to elucidate the function of GerAB as a water channel.

## **References**

1. Christie, G. & Setlow, P. Bacillus spore germination: Knowns, unknowns and what we need to learn. Cell Signal 74, 109729 (2020).
2. Artzi, L. et al. Dormant spores sense amino acids through the B subunits of their germination receptors. Nat Commun 12, 6842 (2021).
3. Blinker, S., Vreede, J., Setlow, P. & Brul, S. Predicting the structure and dynamics of membrane protein GerAB from *Bacillus subtilis*. Int J Mol Sci 22, 3793 (2021).

## Validation that spores of all transformants carry the plasmid

1. Spores from 24 independent colonies isolated and lysed
2. PCR run on lysate to identify plasmid DNA, and to give an  $\sim 2.6$  kb fragment
3. PCR products run on gel electrophoresis and stained



Spores from all colonies have the plasmid