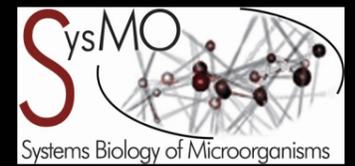


Elucidating Unexpected Reporter Signals for σ^B Activity in *B. subtilis*



Ulf W. Liebal*

Praveen Kumar Sappa §

Thomas Millat*

Leif Steil §

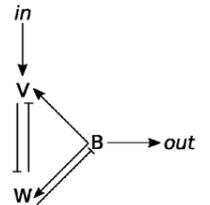
Uwe Völker §

Olaf Wolkenhauer*



Background

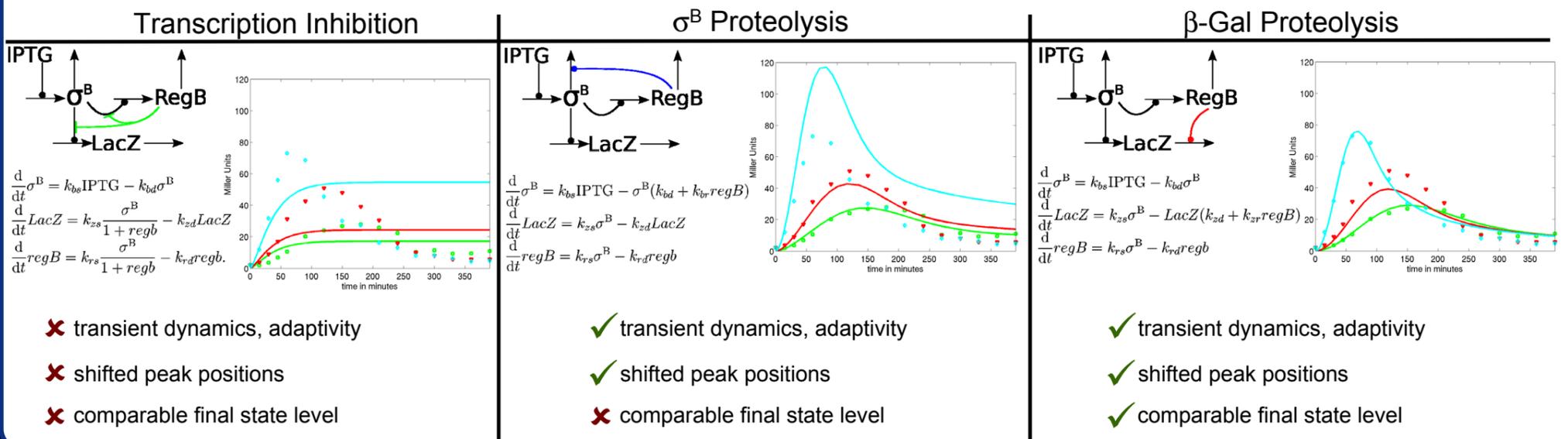
- The σ^B regulon confers *B. subtilis* with the ability to respond to various stress stimuli and adapts it for future stress incidents. Metabolic expenditures are reduced, in turn, expression of about 150 protective genes is activated.
- Starvation activates the anti-anti sigma factor RsbV that inhibits RsbW by complex formation. RsbW itself is the anti sigma factor of σ^B . Hence, an inhibition of RsbW results in release of σ^B and consequently in global expression changes. RsbW can phosphorylate RsbV thus inactivating it. The proteins RsbW and RsbV are transcribed by σ^B and it is assumed that increasing levels of RsbW lead to an inactivation of RsbV and to an adaptive σ^B response.
- To monitor different levels of σ^B -dependent expression, we modulated σ^B expression in a *rsbW* mutant via IPTG induction of a P_{spac} promoter cloned upstream of the σ^B . β -Gal activity of a *ctc::lacZ* fusion was used as reporter for σ^B dependent expression. Direct activation of σ^B in this strain (BSA115) is thought to activate reporter protein synthesis and dysfunctional feedback regulation should result in lasting β -Gal activity.



Surprise and Explanation

- Addition of IPTG induces expression of β -Gal, but activity is transient.
- Maximal β -Gal signal is independent of growth rate
- β -Gal decrease starts before transition to stationary phase for 1mM IPTG.
- All β -Gal signals drop to uninduced signal strength
- High σ^B activation \rightarrow Strong β -Gal decrease
- Low σ^B activation \rightarrow Weak β -Gal decrease

Models of the most simple three explanatory hypotheses are fitted to the experimental data to test how well each is able to reproduce the observations:



Adaptive σ^B behaviour

Hypothesis

Can β -Gal proteolysis contribute to the adaptive σ^B response in a wild type *B. subtilis*?:

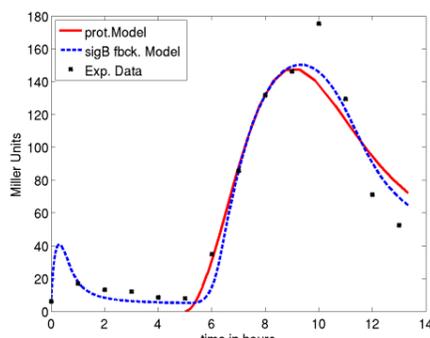
- Taking the β -Gal proteolysis model parameterized with BSA115 experiment as starting point.
- Use of a previous experimental data of β -Gal in wt *B. subtilis*. σ^B response is activated by starvation at the transition to stationary phase.
- Two parameters (IPTG, k_{zs}), that are independent of the putative protease dynamics, are allowed to vary during parameter estimation.

Result

The β -Gal proteolysis model (red) can fit a wt σ^B response with adaptive behaviour as well as a model that relies on negative feedback regulation including RsbW/V proteins (blue).

Conclusion

Therefore, the hypothesis can not be rejected and proteolysis of β -Gal can contribute to adaptivity in the wild type.



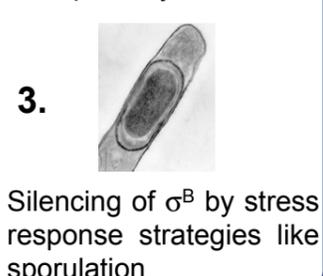
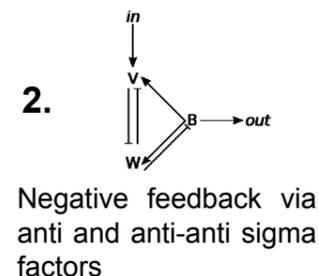
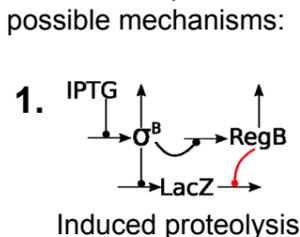
Conclusions and Perspectives

B. subtilis BSA115 shows adaptive, transient dynamics of β -Gal following σ^B activation by IPTG. Modelling and Northern blot experiments suggest σ^B induced β -Gal proteolysis as the origin.

σ^B induced proteolysis of β -Gal in BSA115 was assumed to occur also in *B. subtilis* wild type cells, raising the possibility that β -Gal instabilities might contribute to the transient nature of the σ^B response – testing the ability of the β -Gal proteolysis model to reproduce wild type adaptivity confirmed this assumption.

Therefore, part of the adaptive dynamics of β -Gal reporter signal is due to induced protein proteolysis, questioning the applicability of β -Gal reporters.

Research focus on σ^B has been on its activation and the cause of the transient response is not well known. Our results add proteolysis to the possible mechanisms:



Affiliations, Contact & Acknowledgements

* Systems Biology & Bioinformatics group, University of Rostock, Rostock, Germany

ulf.liebal@uni-rostock.de

www.sbi.uni-rostock.de

§ Institute for Microbiology and Institute for Genetics and Functional Genomics, Ernst-Moritz-Arndt-University Greifswald, Greifswald, Germany

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