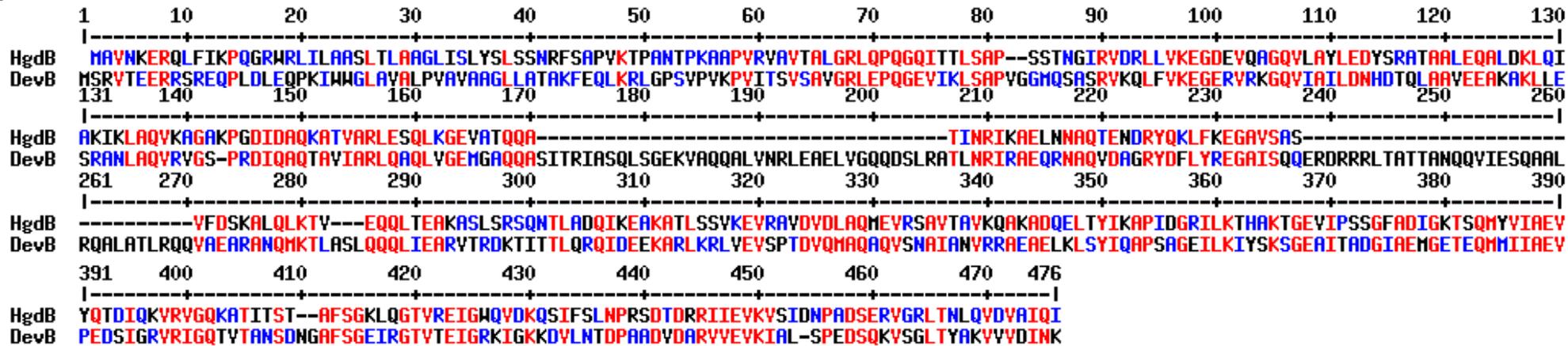


# Supplementary data.

## A



## B

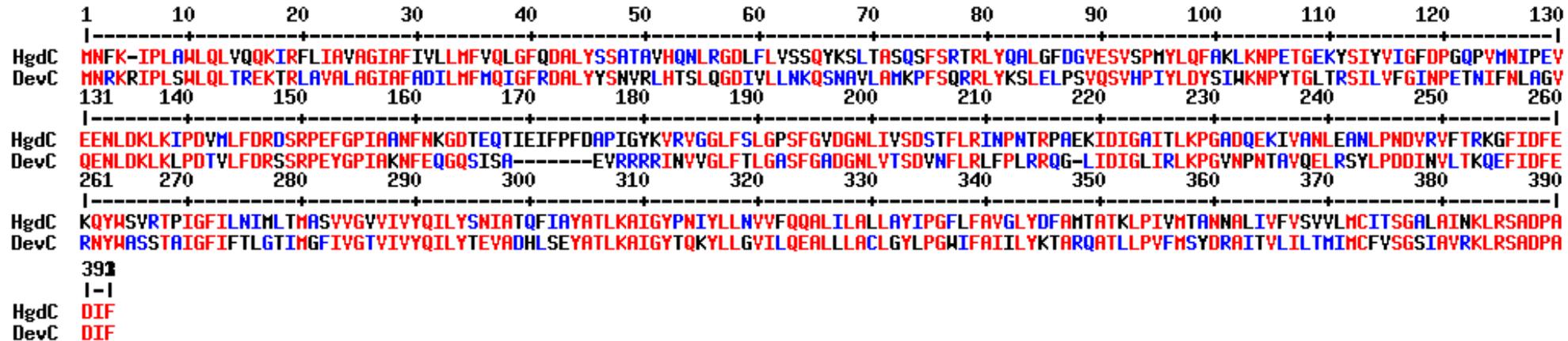


Figure S1. Alignment of amino acid sequences of HgdB with DevB (A) and HgdC with DevC (B).

Alignment was performed using Multalin software [26]. Red color indicates highly similar areas, blue and black colors show low similar areas.

## Supplementary data.

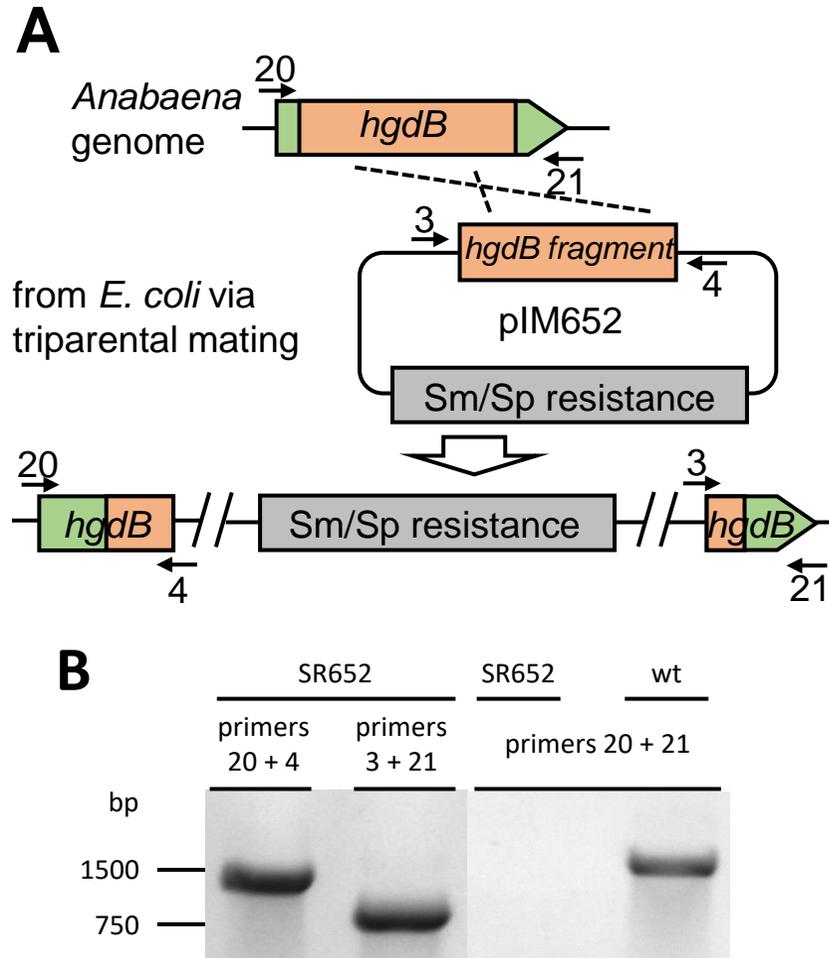


Figure S2. **Segregation of the mutant SR652.**

**A**, Scheme of homologous recombination to obtain the single recombinant SR652 mutant. Arrows with numbers indicate primers (Tab. S1) used for the genotypic analysis of the mutant; **B**, Genotypic analysis of the mutant; SR652 - mutant colony was used as a template for PCR, wt – wild type colony was used as the template. Numbers correspond to those depicted on **A**.

## Supplementary data.

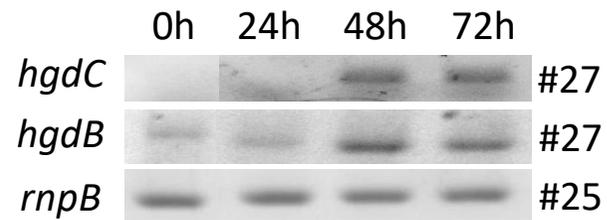
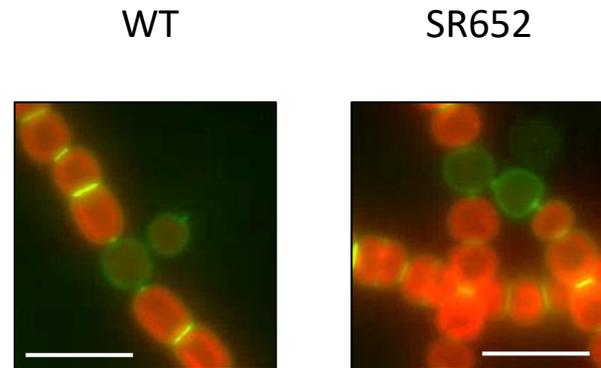


Figure S3. **Time-dependent expression analysis of *hgdC* and *hgdB* genes in the WT culture during nitrogen starvation by RT-PCR.** *rnpB* corresponds to ribonuclease B, which was used as a positive control. Numbers at the right side of the table are numbers of PCR cycles.

## Supplementary data.



**Figure S4. Vancomycin-FL staining of WT and SR652 mutant filaments.**

Vancomycin-FL (green fluorescence) is a fluorescent stain that binds to the peptidoglycan. Red color corresponds to the autofluorescence of cells. Bars are 5  $\mu\text{m}$  long.

## Supplementary data.

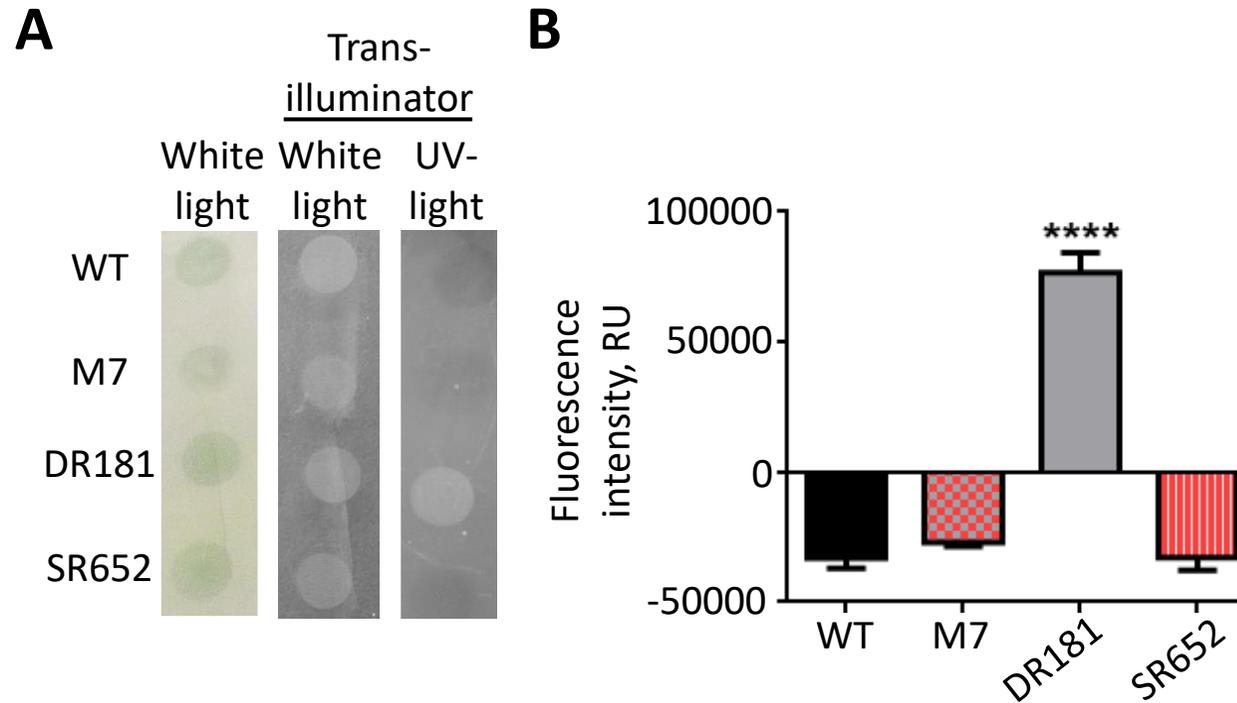


Figure S5. **Retention of ethidium bromide by the SR652 mutant in comparison to the WT, the TolC-deficient mutant DR181, and the DevA-deficient mutant M7.**

**A**, Behavior of *Anabaena* WT and mutant cultures on the solid BG11 medium containing 0.5 µg/ml Ethidium bromide after 16 hours growth starting at 0.1 µg of chlorophyll a; **B**, evaluation of EtBr retention by calculating the relative intensity (RU – relative units) of EtBr fluorescence under UV-light in *Anabaena* colonies. The histogram shows the mean values ± SD of two independent experiments. Student's *t* test *P* value is < 0.0001 (\*\*\*\*).

# Supplementary data.

Table S1. *Anabaena* sp. PCC 7120 strains used in the work.

Number	Name	Reference
1	<i>Anabaena</i> sp. PCC 7120 wild type	[27]
2	SR652	This work
3	SR706	This work
4	M7 (DevA-defficient mutant)	[21]
5	DR181 (ToIC-defficient mutant)	[18]

Table S2. *E. coli* strains used in the work.

Number	Name	Reference
1	NEB10	New England Biolabs
2	Top10	Invitrogen
3	HB101 (pRL528)	[28]
4	J53 (RP4)	[28]
5	Lemo BL21 ( $\lambda$ DE3)	Novagen/Merck

Table S3. Plasmids used in the work.

Number	Name	Reference
1	pIM652	This work
2	pIM706	This work
3	pRL277	[57]
4	pET42a	Novagen/Merck
5	pASK-IBA3	IBA Lifesciences
6	pIM701	This work
7	pIM689	This work

## References:

57. Black, T. A.; Cai, Y.; Wolk, C. P. Spatial expression and autoregulation of *hetR*, a gene involved in the control of heterocyst development in *Anabaena*. *Mol. Microbiol.* **1993**, *9*, 77-84, doi:10.1111/j.1365-2958.1993.tb01670.x.

