

Figure S1

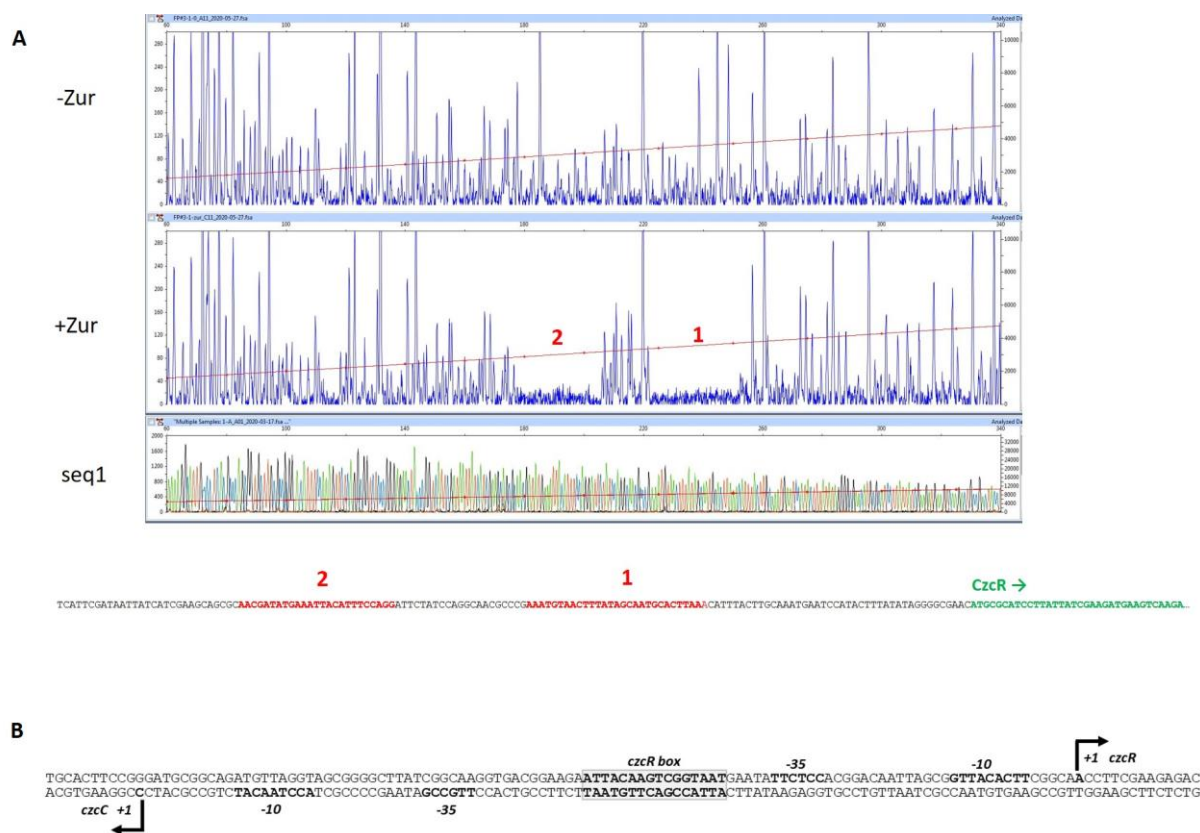


Figure S1. A) DNase I footprinting of *czcR* promoter (template strand) in the absence or presence or of 1 μ M Zur protein. DNA fragments were analyzed by capillary electrophoresis. The two Zur boxes are indicated by the numbers 1 and 2 in red. The sequencing reaction is visible below the figure (seq1) and the sequence with the two boxes indicated in red is shown below the figure. **B)** Transcription start site (+1) of *czcR* and *czcC* determined by 5'RACE. The CzcR box, according to (1) is indicated in the box and the -35, -10 sequences were determined *in silico* using BPROM program (2).

1. Fan, K., Cao, Q. and Lan, L. (2021) Genome-Wide Mapping Reveals Complex Regulatory Activities of BfmR in *Pseudomonas aeruginosa*. *Microorganisms*, **9**.
2. Solovyev, V. and Salamov, A. (2011) In Li, R. W. (ed.), *Metagenomics and its Applications in Agriculture, Biomedicine and Environmental Studies*. Nova Science Publishers, pp. 61-78.