

Supporting Information

Synthetic gene circuits combining CRISPR interference and CRISPR activation in *E. coli*: importance of equal guide RNA binding affinities to avoid context-dependent effects

Içvara Barbier¹, Hadiastri Kusumawardhani^{1,+}, Lakshya Chauhan^{1,2,+}, Pradyumna Vinod Harlapur², Mohit Kumar Jolly², and Yolanda Schaerli^{1,*}

¹Department of Fundamental Microbiology, University of Lausanne, Switzerland

²Centre for BioSystems Science and Engineering, Indian Institute of Science, India

⁺These authors contributed equally to the project

^{*}Corresponding author: yolanda.schaerli@unil.ch

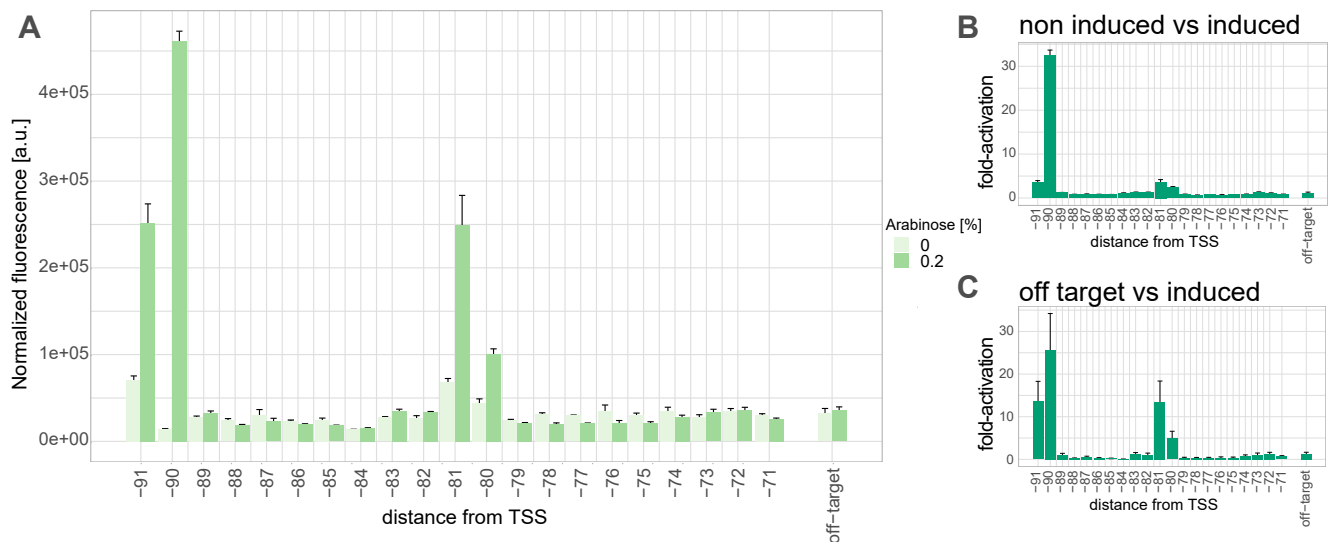


Figure S1: Activation is sensitive to the distance to the transcriptional start site (TSS)

A. Normalized green fluorescence at indicated arabinose concentrations. Different binding site position upstream of the TSS were tested as indicated in the x axis. **B.** Fold-activation at 0.2% arabinose compared to fluorescence level at 0% arabinose. **C.** Fold-activation at 0.2% arabinose compared to off-target construct (scRNA 1 - binding site 4) at 0% arabinose. Mean and s.d. represent three biological replicates.

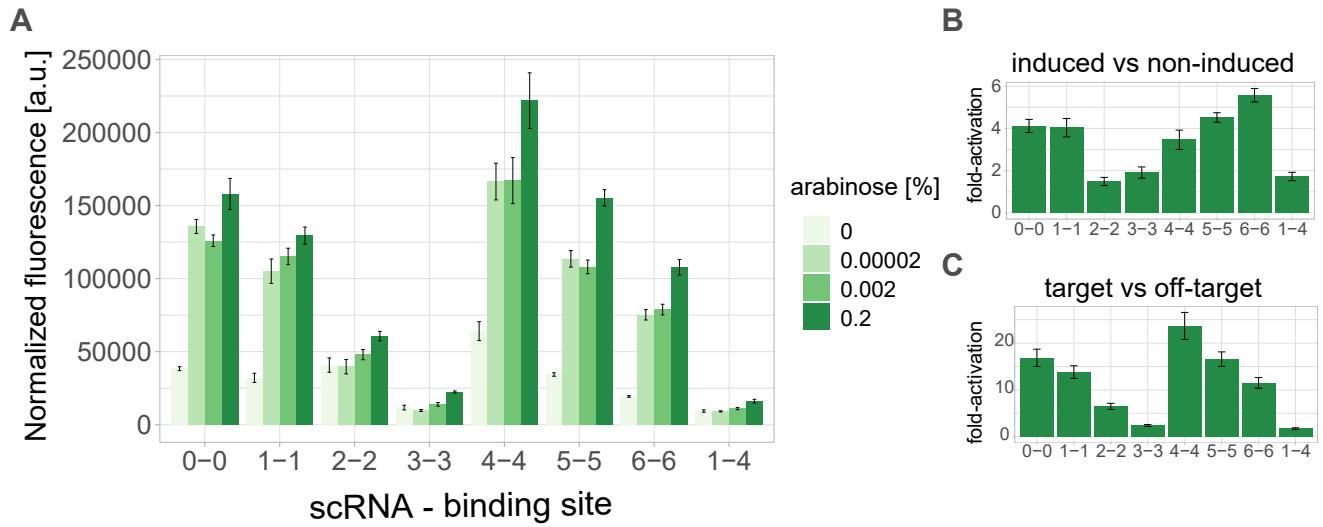


Figure S2: Induction of CRISPRa at different arabinose concentrations

A. Normalized green fluorescence at indicated arabinose concentrations. Different combinations of scRNAs and binding sites were tested as indicated in the x axis. **B.** Fold-activation at 0.2% arabinose compared to fluorescence level at 0% arabinose. **C.** Fold-activation at 0.2% arabinose compared to off-target construct (scRNA 1 - binding site 4) at 0% arabinose. Mean and s.d. of three biological replicates.

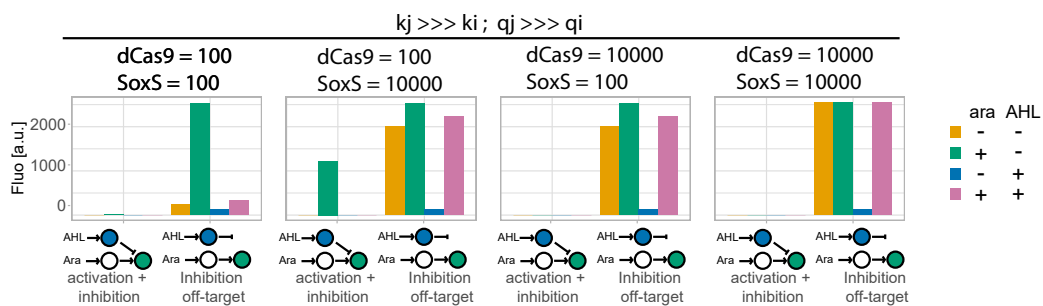


Figure S3: Increasing dCas9 and SoxS quantities does not lead to correct function.

Qualitative model of GFP intensity with or without arabinose and AHL induction (0,10) and different values of MCP-SoxS and dCas9. k_j and q_j are fixed at 100000, k_i and q_i are fixed at 1. For the off-target controls, q_j is set to 0.

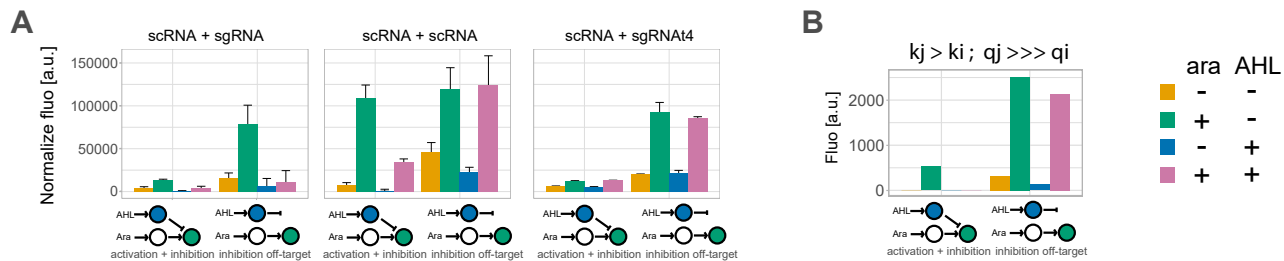


Figure S4: CRISPRa/i induction with different guide RNAs for CRISPRi

A. *CRISPRa* with *scRNA* and *CRISPRi* with *sgRNA*, *scRNA* or *sgRNAt4* in on-target circuits or off-target controls. Bar plots at 0 or maximum concentration of AHL (0.1 μ M) and arabinose (0.2%). Mean and s.d. represent three biological replicates. **B.** Modeling as in figure 2 but with $qj = 100000$ and $kj = 1000$

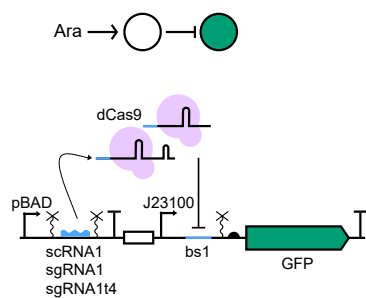
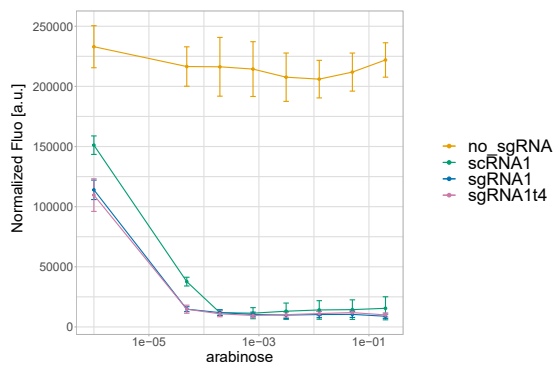
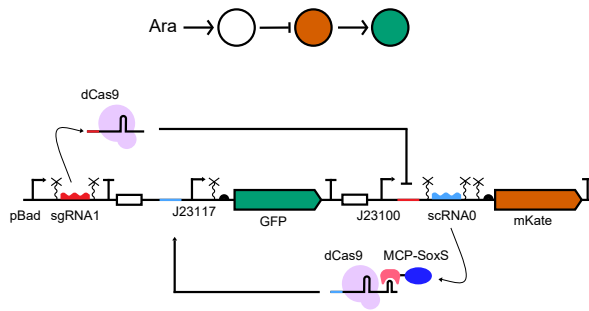
A**B**

Figure S5: Inhibition with sgRNA or scRNA

A. Details of the circuit design and schematic representation of the circuit. **B.** GFP fluorescence in presence of different concentrations of arabinose. Mean and s.d. represent three biological replicates.

A



B

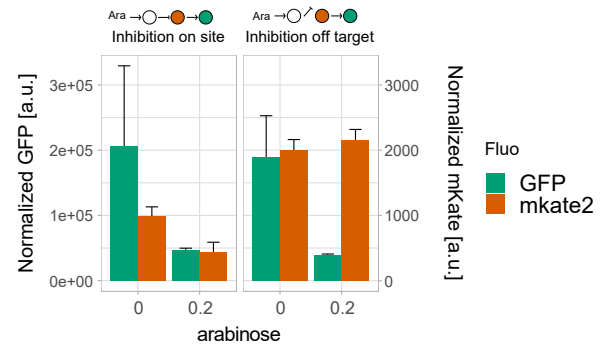


Figure S6: Cascade circuit with sgRNA and scRNA leads to unpredicted behavior of the off-target control
A. Details of the circuit design and schematic representation of the circuit. It is the same as in Figure 4, but instead of using scRNA1, we used here sgRNA1 **B.** Bar plots represent the GFP and mKate fluorescences of the circuit and of an off-target control in absence (0%) or presence of arabinose (0.2%). Mean and s.d. represent three biological replicates.