

Table S 1: Bacterial strains used in this study

| Bacterial Strains | Genotype | Reference |
|--|--|------------------------|
| <i>E. coli</i> S17-1 | Conjugation strain; <i>recA pro hsdR</i> ; RP42Tc::Mu-Km::Tn7 integrated into the chromosome | ATCC 47055 |
| <i>P. polymyxa</i> DSM 365 | wild type | DSMZ |
| <i>P. polymyxa</i> DSM 365 Δ <i>ldh1</i> | DSM 365 Δ <i>ldh1</i> | Schilling et al (2020) |
| <i>P. polymyxa</i> DSM 365 Δ <i>pepCQ</i> | DSM 365 Δ <i>pepC</i> | This study |
| <i>P. polymyxa</i> DSM 365 Δ <i>pepQ</i> | DSM 365 Δ <i>pepQ</i> | This study |

In addition to the strains listed above, each plasmid listed in Table S 2 was used for the transformation of *E. coli* S17-1 as well as for the conjugational transformation of *P. polymyxa* DSM 365.

Table S 2: Plasmids used in this study

| Plasmids | Description | Reference |
|-------------------|---|-----------------------|
| pCasPP | <i>P. polymyxa</i> CRISPR-Cas9 genome editing plasmid | Rütering et al (2017) |
| pCasPPH_pegC | <i>pegC</i> targeting knock out plasmid containing repair template | This study |
| pCasPPH_pegQ | <i>pegQ</i> targeting knock out plasmid containing repair template | This study |
| pCRai | <i>PsgsE-AsdCas12a</i> ; <i>Pgapdh</i> -off target gRNA; <i>neo</i> ; <i>oriT</i> , <i>repU</i> | This study |
| pCRaiGFP | <i>PsgsE-AsdCas12a</i> ; <i>Pgapdh</i> -off target gRNA; <i>neo</i> ; <i>oriT</i> , <i>repU</i> ; <i>PsgsE-sfGFP</i> | This study |
| pCRaiGmR_soxS | <i>PsgsE-AsdCas12a-soxS</i> ; <i>Pgapdh</i> -off target gRNA; <i>neo</i> ; <i>oriT</i> , <i>repU</i> ; <i>PsgsE_mRFP-US-PsgsE-sfGFP</i> | This study |
| pCRaiGFP_GltC | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - off target gRNA | This study |
| pCRaiGFP_GltC_a1 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a1 | This study |
| pCRaiGFP_GltC_a2 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a2 | This study |
| pCRaiGFP_GltC_a3 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a3 | This study |
| pCRaiGFP_GltC_a4 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a4 | This study |
| pCRaiGFP_sig70 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - off target gRNA | This study |
| pCRaiGFP_sig70_a1 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a1 | This study |
| pCRaiGFP_sig70_a2 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a2 | This study |
| pCRaiGFP_sig70_a3 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a3 | This study |
| pCRaiGFP_sig70_a4 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-sfGFP</i> - sgRNA_a4 | This study |
| pCRai_GFP_rpoD_a1 | Test plasmid for CRISPRa targeting US region of <i>PsgsE-</i> | This study |

| | | |
|--------------------------------|---|------------|
| | <i>sfGFP</i> - sgRNA_a1 | |
| pCRai_GFP_rpoD_a2 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a2 | This study |
| pCRai_GFP_rpoD_a3 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a3 | This study |
| pCRai_GFP_rpoD_a4 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a4 | This study |
| pCRai_GFP_rpoD | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - off target gRNA | This study |
| pCRai_GFP_soxS_a1 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a1 | This study |
| pCRai_GFP_soxS_a2 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a2 | This study |
| pCRai_GFP_soxS_a3 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a3 | This study |
| pCRai_GFP_soxS_a4 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a4 | This study |
| pCRai_GFP_soxS | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - off target gRNA | This study |
| pCRai_GFP_CRP_a1 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a1 | This study |
| pCRai_GFP_CRP_a2 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a2 | This study |
| pCRai_GFP_CRP_a3 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a3 | This study |
| pCRai_GFP_CRP_a4 | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - sgRNA_a4 | This study |
| pCRai_GFP_CRP | Test plasmid for CRISPRa targeting US region of PsgsE- <i>sfGFP</i> - off target gRNA | This study |
| pCRaiGFP_soxS_sfGFP_T1 | Test plasmid for <i>sfGFP</i> repression sgRNA_T1 | This study |
| pCRaiGFP_soxS_sfGFP_T2 | Test plasmid for <i>sfGFP</i> repression sgRNA_T2 | This study |
| pCRaiGFP_soxS_sfGFP_T3 | Test plasmid for <i>sfGFP</i> repression sgRNA_T3 | This study |
| pCRai_pepQ_T1 | Plasmid for repression of <i>pepQ</i> sgRNA T1 | This study |
| pCRai_pepCQ_T1 | Plasmid for dual repression of <i>pepQ</i> and <i>pepC</i> | This study |
| pCRai_soxS_ldhmultibdh1 | Multiplex knock-down of repression of <i>ldh2</i> , <i>ldh3</i> , <i>ldh4</i> and activation of <i>Pbdh</i> (sgRNA 1) | This study |
| pCRai_soxS_ldhmultibdh2 | Multiplex knock-down of repression of <i>ldh2</i> , <i>ldh3</i> , <i>ldh4</i> and activation of <i>Pbdh</i> (sgRNA 2) | This study |
| pCRai_soxS_ldhmultibdh3 | Multiplex knock-down of repression of <i>ldh2</i> , <i>ldh3</i> , <i>ldh4</i> and activation of <i>Pbdh</i> (sgRNA 3) | This study |
| pCRaiGmR_soxS_mT2 | PsgsE-mRFP-US-PsgsE- <i>sfGFP</i> expression plasmid with gRNA for mRFP repression (sgRNA T2) | This study |
| pCRaiGmR_soxS_dualT2 | PsgsE-mRFP-US-PsgsE- <i>sfGFP</i> expression plasmid with gRNA for mRFP repression (sgRNA T2) and GFP activation (sgRNA a1) | This study |
| pCRaiGmR_soxS_GFPa1 | PsgsE-mRFP-US-PsgsE- <i>sfGFP</i> expression plasmid with gRNA and GFP activation (sgRNA a1) | This study |

Table S 3: Oligonucleotides and primers used in this study. Overhangs used for Golden Gate Assembly or Gibson isothermal assembly are depicted in lower case. Restriction sites are underlined.

| Name | SEQUENCE 5'->3' | Comment |
|---------------------|---|---|
| soxS_fw | ATGTT <u>GGTCTCA</u> ACGCTGATGTCCCATCAGAAAATTATTCAGGATC | Cloning of activator domain |
| soxS_rev | TAAAAG <u>GTCTCGT</u> ACCATTTACAGGCGGTGGCGATAATCG | Cloning of activator domain |
| sig70_fw | ATGTT <u>GGTCTCA</u> ACGCTGATGATAGAGGACACACCCATTCGGT | Cloning of activator domain |
| sig70_rev | TAAAAG <u>GTCTCGT</u> ACCATTTCATTGCCATTCTCCTCCTTCC | Cloning of activator domain |
| cAMP_fw | ATGTT <u>GGTCTCA</u> ACGCTGATGCATCCCTATTATCGTTATTAGAAAAA | Cloning of activator domain |
| cAMP_rev | TAAAAG <u>GTCTCGT</u> ACCATTTAATGCGGATCACTACGAGCAA | Cloning of activator domain |
| GltC_fw | ATGTT <u>GGTCTCA</u> ACGCTGGTGGAATTACGACAGTTACAATACTTCATGAA | Cloning of activator domain |
| GltC_rev | TAAAAG <u>GTCTCGT</u> ACCATTCACTGCCCTAAATCCGCTTTG | Cloning of activator domain |
| RpoD_fw | ATGTT <u>GGTCTCA</u> ACGCTGGTGCTGTATCCTTCCATTGATGAAATG | Cloning of activator domain |
| RpoD_rev | TAAAAG <u>GTCTCGT</u> ACCATTTATTCTTCTTCGCCGCTTTC AAGT | Cloning of activator domain |
| As_CRISPR_fw | GGGGATACGCTAATTTCTACTCTTGATAGATAAGTCTTCTCAGCCG | Construction of pCRai |
| As_CRISPR_rev | AAATCCAGATGGAGTATGTCTTCACCGGTGGAAAGCG | Construction of pCRai |
| AsCRISPR_vec_fw | CACCGGTGAAGACATACTCCATCTGGATTGTTTCAGAACGC | Construction of pCRai |
| AsCRISPR_vec_rev | GTAGAAATTAGCGTATCCCCTTCAGATACTCGCAC | Construction of pCRai |
| enAsCPF1_fw | TTAGGCTTTTACTTAATGACACAGTTTGAAGGCTTTACGAATCTG | Construction of pCRai |
| enAsCPF1_rev | CGTAGATCTGAATTCTTATTACCCGAGACCTACCCAATGCG | Construction of pCRai |
| enAsCPF1_Vector.fw | GGTCTCGGGTAATAAGAATTCAGATCTACGCGTTCCCGC | Construction of pCRai |
| enAsCPF1_Vector.REV | TTCAAAGTGTGTCATTAAGTAAAAGCCTAAAATCCCC TTCGTT | Construction of pCRai |
| GFP_vecint_rev | gcaacgcgcccttttacggTTCCTGGCCATATGACGATCCTCCTTACCTCTCATTG | <i>sfGFP</i> plasmid cloning |
| GFP_vecint_fw | gcccgcaaccggcgcacatcaagcccgcGACTAGCAATATGAAACACGGAAAAAATCAAGC | <i>sfGFP</i> plasmid cloning |
| sgsE-a1_fw | agatTTTATCTCACATAATAGGGCT | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a1_rev | gagtAGCCCTATTATGTGagatAA | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a2_fw | agatAGTCTATATCAATCGGTAAC | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a2_rev | gagtGTTACCGATTGATATAGACT | Test gRNA for activation of PsgsE- <i>sfGFP</i> |

| | | |
|-----------------------|--|---|
| sgsE-a3_fw | agatATCCTCATATTTTCCTAGTA | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a3_rev | gagtTACTAGGAAAATATGAGGAT | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a4_fw | agatCATCGATGGCGACATTGATA | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sgsE-a4_rev | gagtTATCAATGTGCGCCATCGATG | Test gRNA for activation of PsgsE- <i>sfGFP</i> |
| sfGFP1_T1_fw | agatCGTGC GTGGCGAGGGTGAAG | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| sfGFP1_T1_rev | gagtCTTCACCCTCGCCACGCACG | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| sfGFP_T2_fw | agatCCATTAGTTGCGTCACCTTC | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| sfGFP_T2_rev | gagtGAAGGTGACGCAACTAATGG | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| sfGFP_T3_fw | agatAGCTCAATGCGGTTTACCAG | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| sfGFP_T3_rev | gagtCTGGTAAACCGCATTGAGCT | Test gRNA for repression of PsgsE- <i>sfGFP</i> |
| mRFP_T_fw | agatAAAGTTCGTATGGAAGGTTC | gRNA for repression of PsgsE- <i>mRFP</i> |
| mRFP_T_rev | gagtGAACCTCCATACGAACTTT | gRNA for repression of PsgsE- <i>mRFP</i> |
| sfGFP_dual_fw | agatTTATCTCACATAATAGGGCTTAATTTCTACTC | Test gRNA for repression of PsgsE- <i>mRFP</i> and simultaneous activation of PsgsE- <i>sfGFP</i> |
| sfGFP_dual_rev | acaaGAGTAGAAATTAAGCCCTATTATGTGAGATAA | Test gRNA for repression of PsgsE- <i>mRFP</i> and simultaneous activation of PsgsE- <i>sfGFP</i> |
| mRFP_dual_fw | ttgtAGATAAAGTTCGTATGGAAGGTTC | Test gRNA for repression of PsgsE- <i>mRFP</i> and simultaneous activation of PsgsE- <i>sfGFP</i> |
| mRFP_dual_rev | gagtGAACCTCCATACGAACTTTATCT | Test gRNA for repression of PsgsE- <i>mRFP</i> and simultaneous activation of PsgsE- <i>sfGFP</i> |
| GGA_sfGFP_fw | AATTGGTCTCAA ACT CAATATGAAACACGGAAAAAATCAAGCAG | Cloning of PsgsE- <i>mRFP</i> -US-PsgsE- <i>sfGFP</i> expression cassette |
| GGA_sfGFP_rev | AATTGGTCTCA CT ACTAGTATGACGATCCTCCTTACCTCTCATTG | Cloning of PsgsE- <i>mRFP</i> -US-PsgsE- <i>sfGFP</i> expression cassette |
| GGA_mRFP_fw | AATTGGTCTCATCGACCTGCATACTAGCCTGTTACAGGCATATTCATATCAATGTC | Cloning of PsgsE- <i>mRFP</i> -US-PsgsE- <i>sfGFP</i> expression cassette |
| GGA_mRFP_rev | CTTTGGTCTCAAGTTAAGCACCGGTGgagtGACGAC | Cloning of PsgsE- <i>mRFP</i> -US-PsgsE- <i>sfGFP</i> expression cassette |

| | | |
|-----------------------|-------------------------------------|---|
| pepQ_T1_fw | agatCACCAGCAGTGCAGACAATC | gRNA for repression of <i>pepQ</i> |
| pepQ_T1_rev | gagtGATTGTCTGCACTGCTGGTG | gRNA for repression of <i>pepQ</i> |
| pepCQ_T1_fw | agatCCTGACATGACAGCCTCGCCTAATTTCTACT | gRNAs for dual repression of <i>pepCQ</i> |
| pepCQ_T1_rev | caagAGTAGAAATTAGGCGAGGCTGTCATGTCAGG | gRNAs for dual repression of <i>pepCQ</i> |
| pepCQ2_T1_fw | cttgTAGATCACCAGCAGTGCAGACAATC | gRNAs for dual repression of <i>pepCQ</i> |
| pepCQ_T1_rev | gagtGATTGTCTGCACTGCTGGTGATCTA | gRNAs for dual repression of <i>pepCQ</i> |
| bdh1_fw | cttgTAGATGACACTCATTCTGTGGTATA | gRNA cloning for <i>Ppbdh</i> activation |
| bdh1_rev | gagtTATACCACAGAATGAGTGTCATCTA | gRNA cloning for <i>Ppbdh</i> activation |
| bdh2_fw | cttgTAGATTTCTTGTCTTTGCTTCAATT | gRNA cloning for <i>Ppbdh</i> activation |
| bdh2_rev | gagtAATTGAAGCAAAGACAAGAAATCTA | gRNA cloning for <i>Ppbdh</i> activation |
| bdh3_fw | gagtGCTCGTTACTTTTATACAAA | gRNA cloning for <i>Ppbdh</i> activation |
| bdh3_rev | gagtTTTGTATAAAAGTAACGAGCATCTA | gRNA cloning for <i>Ppbdh</i> activation |
| sfGFP_qPCR_fw | CCCTATTCTGGTGGAAGTGGATGG | qPCR primer |
| sfGFP_qPCR_rev | CAGTAGTACAGATGAACTTCAGCGTC | qPCR primer |
| gyrA_qPCR_fw | GAGATATGGCCGCTGCGATG | qPCR primer |
| gyrA_qPCR_rev | GCTCTCTCACCATCGTAGTTCGG | qPCR primer |
| mRFP_qPCR_fw | TACCTGAAACTGTCCTTCCCGG | qPCR primer |
| mRFP_qPCR_rev | GTAGATGAACTCACCGTCTTGCAAG | qPCR primer |
| ldh3_qPCR_fw | GAGTCATTGGATCA GA CGTTGC | qPCR primer |
| ldh3_qPCR_rev | AACTCGGAGTCACCGTGTCTCC | qPCR primer |
| ldh2_qPCR_fw | GATTATCGGGGTTGGACGCATTGG | qPCR primer |
| ldh2_qPCR_rev | CAAGGTTGTGAATGGAATGACTTCCTTG | qPCR primer |
| ldh4_qPCR_fw | GAATGAGCCGGACCTGGATGCG | qPCR primer |
| ldh4_qPCR_rev | CAATCCTCGACTGGATCGGTGTTTC | qPCR primer |
| bdh_qPCR_fw | AGATGCTCAGCATCCATTGACTGG | qPCR primer |
| bdh_qPCR_rev | CAACGACACGGTCGCCAACCTG | qPCR primer |

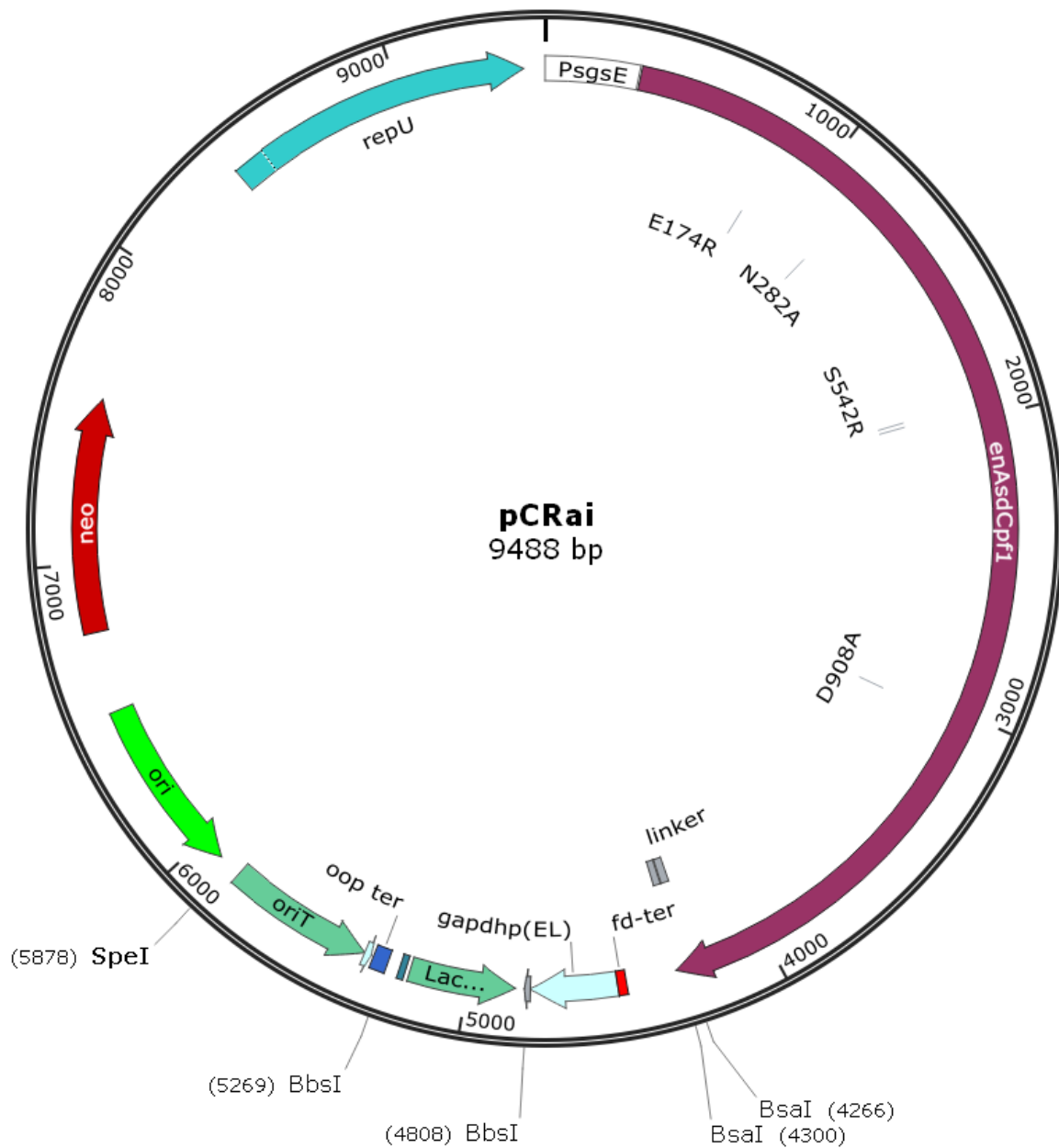


Figure S 1: Plasmid map of pCRai. The conjugational plasmid based on the pUB110 and pCasPP² plasmids was constructed via Gibson Assembly by replacing the Cas9-gRNA expression cassette with an engineered catalytically inactive variant of Cas12a³ (E174R, N282A, S542R, K548R, D908A) and an appropriate CRISPR-array. Variable activator domains can be cloned by replacing a BsaI-cassette linked via a 10 aa flexible linker. The gRNAs can be cloned by Golden Gate Assembly using BbsI by replacing the *lacZ* expression cassette.

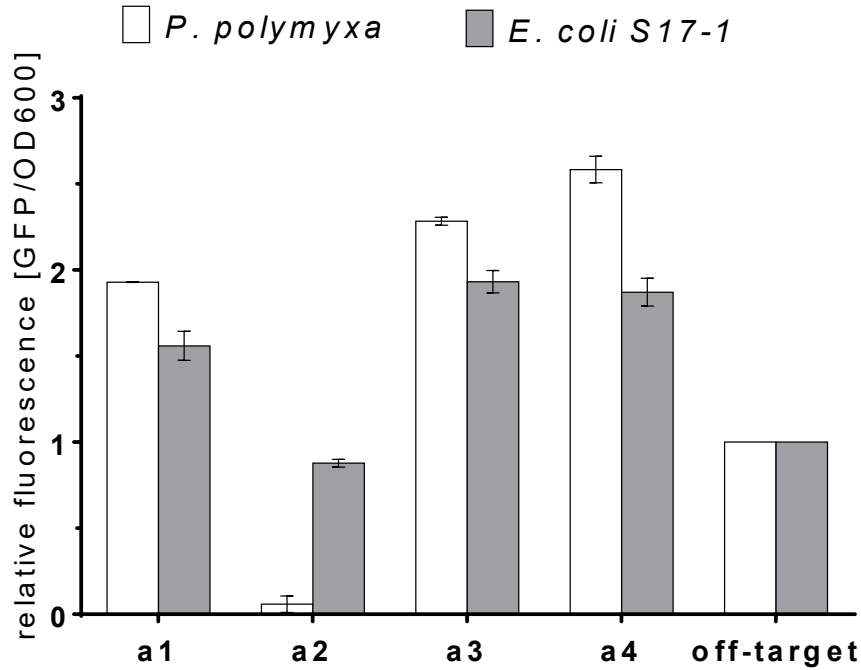


Figure S 2: Relative fluorescence levels of *P. polymyxa* and *E. coli* S17-1 harboring pCRaiGFP_soxS variants. Four different gRNAs (a1-a4) were expressed and normalized fluorescence levels (Ex. 488 nm Em. 515 nm) were compared to a strain expressing an off-target gRNA. The gRNAs a1, a3 and a4 showed increased GFP fluorescence in *P. polymyxa* as well as *E. coli* S17-1 demonstrating similar effects in both, Gram-positive and Gram-negative host organisms.

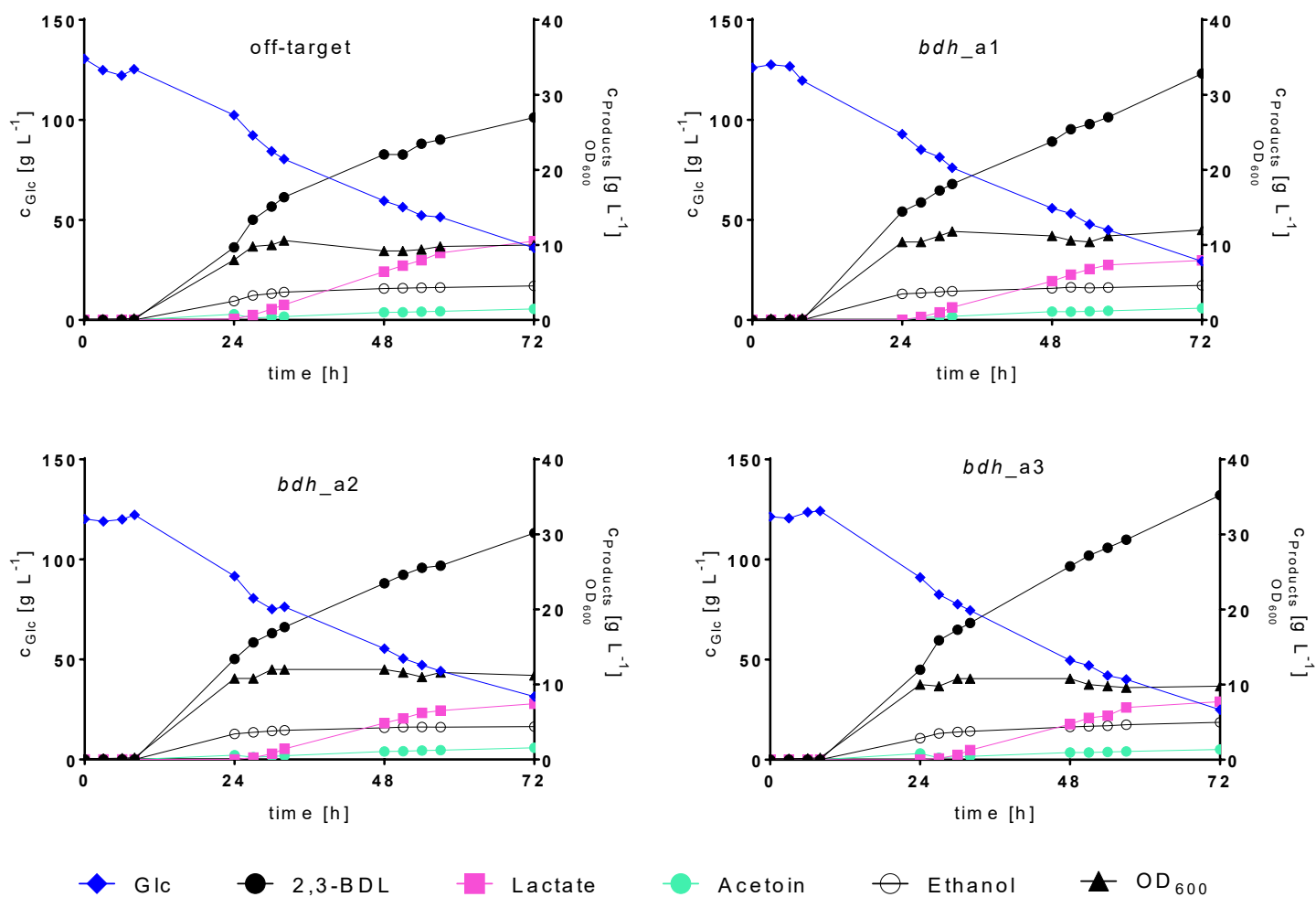


Figure S 3: Profiles of 2,3-BDL fermentations. Overview of fermentation profiles of single batch fermentations. *P. polymyxa* DSM 365 was transformed with pCRai_soxS encoding either off-target gRNAs or a multiplex CRISPR-array targeting the open reading frame of three lactate dehydrogenases and the upstream region of P_{bdh} with three gRNAs (*bdh_a1-a3*) individually. Lactate production was reduced by ~ 20 % in all variants expressing the corresponding gRNAs. Glc: Glucose; 2,3-BDL: 2,3-*R,R*-butantediol

Table S 4 Final product titers and 2,3-BDL yield ($\text{g}_{\text{BDL}} \text{g}_{\text{Glc}}^{-1}$) of 2,3-BDL fermentations. *P. polymyxa* DSM 365 was transformed with pCRai_soxS encoding either off-target gRNAs or a multiplex CRISPR-array targeting the open reading frame of three lactate dehydrogenases and the upstream region of P_{bdh} with three gRNAs (*bdh_a1-a3*) individually. Values represent mean and deviation of biological duplicates after 72 h cultivation at microaerobic conditions.

| | 2,3-BDL | Ethanol | Acetoin | Lactate | Formate | Y_{PS} |
|----------------------|----------------|----------------|----------------|----------------|----------------|-----------------------|
| off-target | 27.46 ± 0.47 | 4.21 ± 0.33 | 1.34 ± 0.14 | 10.30 ± 0.20 | 0.34 ± 0.34 | 0.29 ± 0.01 |
| <i>bdh_a1</i> | 34.30 ± 1.45 | 4.84 ± 0.23 | 1.50 ± 0.08 | 7.69 ± 0.26 | 0.72 ± 0.03 | 0.34 ± 0.00 |
| <i>bdh_a2</i> | 32.44 ± 2.26 | 4.64 ± 0.25 | 1.42 ± 0.15 | 7.74 ± 0.30 | 0.72 ± 0.04 | 0.34 ± 0.00 |
| <i>bdh_a3</i> | 34.74 ± 0.50 | 4.89 ± 0.10 | 1.22 ± 0.08 | 7.81 ± 0.08 | 0.67 ± 0.07 | 0.35 ± 0.02 |

References

- (1) Schilling, C.; Ciccone, R.; Sieber, V.; Schmid, J. Engineering of the 2,3-Butanediol Pathway of *Paenibacillus polymyxa* DSM 365. *Metab. Eng.* **2020**, (in press).
- (2) Rütering, M.; Cress, B. F.; Schilling, M.; Rühmann, B.; Koffas, M. A. G.; Sieber, V.; Schmid, J. Tailor-Made Exopolysaccharides—CRISPR-Cas9 Mediated Genome Editing in *Paenibacillus polymyxa*. *Synth. Biol.* **2017**, 2 (1). <https://doi.org/10.1093/synbio/ysx007>.
- (3) Kleinstiver, B. P.; Sousa, A. A.; Walton, R. T.; Tak, Y. E.; Hsu, J. Y.; Clement, K.; Welch, M. M.; Horng, J. E.; Malagon-Lopez, J.; Scarfò, I.; Maus, M. V.; Pinello, L.; Aryee, M. J.; Joung, J. K. Engineered CRISPR–Cas12a Variants with Increased Activities and Improved Targeting Ranges for Gene, Epigenetic and Base Editing. *Nat. Biotechnol.* **2019**, 37 (3), 276–282. <https://doi.org/10.1038/s41587-018-0011-0>.