

Supplementary Materials

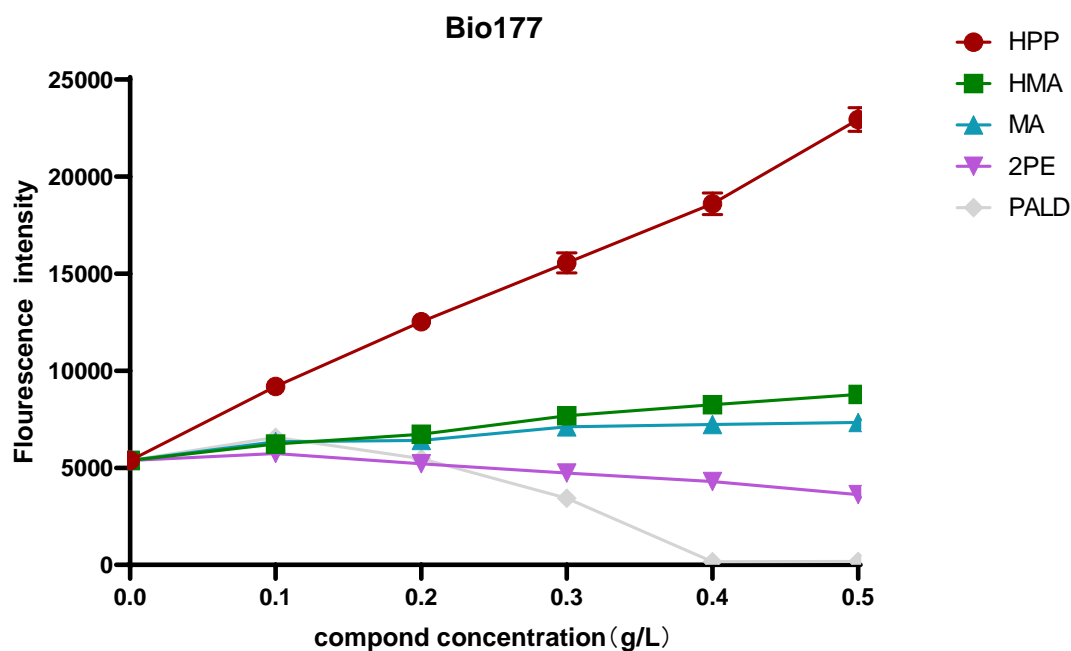
Supplementary Table 1. Strains and plasmids used in this study.

Strains and plasmids	Description	Sources
Strains		
<i>E. coli</i> DH5 α	F- Δ lacU169(Φ 80 lacZ Δ M15) hsdR17 recA1 endA1 supE44 gyrA96 thi-1 relA1	Invitrogen
BW25113	F-, λ -, <i>E. coli</i> K-12 strain BD792 (CGSC6159) lacZ	Invitrogen
BW Δ CD	BW25113 strain knocked out codA	Invitrogen
Plasmids		
gY9s-dual T7-TrrnB-HmaS(ScpA1)-Bio 177	Str ^R , CPA1 promoter, TrrnB, T7 promoter, HmaS Am, CD enzyme, mCherry, pobA promoter, PobR	This study
pHyo094	Cmr ^R , f1 ori, p15A ori, P _{BAD} - UGI	Hyojin, et al. (2020)
PSC101-dCas9(sg-eGFP-1)	Kan ^R , pSC101 ori, Rep 101, lacI, sgRNA1-eGFP, gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-eGFP-1)	Kan ^R , R6K ori, pir, lacI, sgRNA1-eGFP, gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-vec)	Kan ^R , R6K ori, pir, lacI, trc promoter, sgRNA1-eGFP, gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-pykF-1)	Kan ^R , R6K ori, pir, lacI, trc promoter, spacer(pykF), gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-tyrB-2)	Kan ^R , R6K ori, pir, lacI, trc promoter, spacer(tyrB), gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-tyrR-3)	Kan ^R , R6K ori, pir, lacI, trc promoter, spacer(tyrR), gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9(sg-pheA-3)	Kan ^R , R6K ori, pir, lacI, trc promoter, spacer(pheA), gRNA scaffold, dCas9, tracrRNA	This study
R6K-dCas9-gRNA-pykF-tyrB-tyrR-pheA	Kan ^R , R6K ori, pir, lacI, trc promoter, spacer(pheA), trc promoter, spacer(tyrR), trc promoter, spacer(tyrB), trc promoter, spacer(pykF), gRNA scaffold, dCas9, tracrRNA	This study

Supplementary Table 2. Primers used in this study.

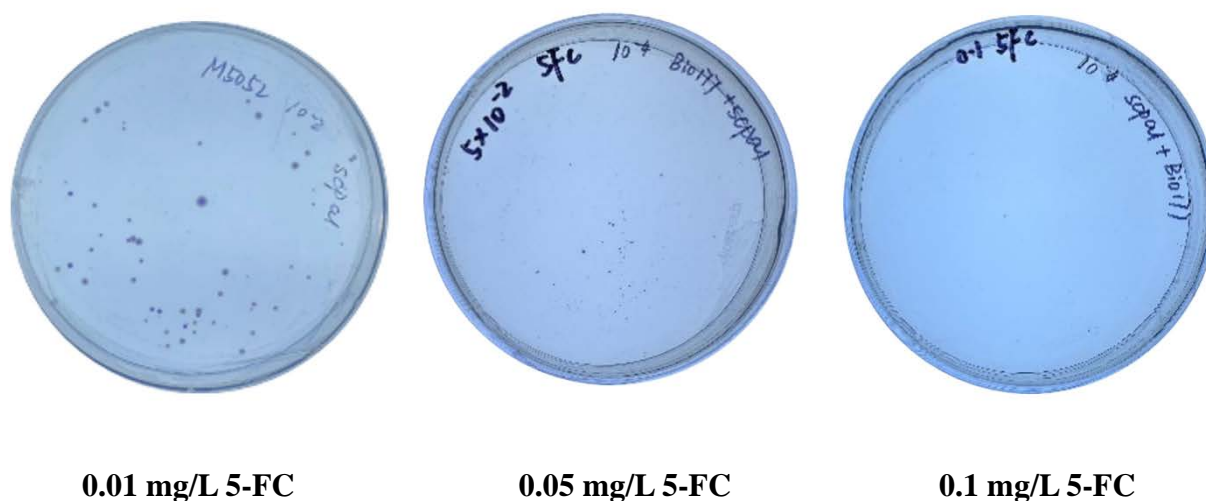
Primers	Sequences
gY9s-dual T7-TrrnB-HmaS(ScpA1)-Bio 177 construction	
(1) Bio 177 cloning	
Bio-F=0811	GTGCTGGGTCTCGCCTTTTACCGTTTGTAATCGATGG
Bio-R=0811	AGCGTGGGTCTCTTATATCTCCTTCTTAAAA GATCTTTTGAA
(2) dual T7-HmaS cloning	
AM-gg-F=0709	GTGCTGGGTCTCGGCAATAATACGACTCAC TATAGGATGGCGGCGCAGGCGGG
AM-R=0811	AGCGTGGGTCTCTTAATACGACTCACTATA GGTTAACG
(3) RE-TrrnB cloning	
RET-F=0811	GTGCTGGGTCTCGTATAAAGGCCAGTCTT TCGACTG
RET-R=0811	AGCGTGGGTCTCTTTGCTGCCTGGCGGCA GTAGCG
(4) TrrnB cloning	
T-F=0811	GTGCTGGGTCTCGATTATGCCTGGCGGCAG TAGCG
T-R=0811	AGCGTGGGTCTCTAAGGCCAGTCTTTTCG ACTG
(5) Bio 177-Str construction	
CD-F==0617	GTGCTGGGTCTCGATTCTTACCGTTTGTAATCGATGGC
Str-R=0617	AGCGTGGGTCTCTTGTGTTATTTGCCGACT ACCTTG
(6) HmaS-scpA1 construction	
cpA1-F=0725	GTGCTGGGTCTCGTAATTATCAAAAAGAGT ATTGACATAAAGTC
Hmas-R=0617	AGCGTGGGTCTCTGAATTCCGACATACAGC AGG

Supplementary Figure 1



Supplementary Figure 1. Evaluating the specificity of biosensor Bio177. HPP: 4-hydroxyphenylpyruvate, HMA: 4-hydroxymandelate, MA: Mandelic acid, 2PE: 2-Phenylethanol, PALD: phenylacetaldehyde.

Supplementary Figure 2



Supplementary Figure 2. Comparison of the functionality between the Bio177 plasmid (low activity) and the gY9s-HmaS-Bio177 plasmid (high activity) in bacteria under stress conditions. Both strains were plated on media with 0.01, 0.05 and 0.1 mg/L of 5-FC and clones containing the hmaS gene were selected. A 100% selection efficiency was achieved on the plate with 0.01 mg/L 5-FC medium. The wild-type strain was impaired in 0.05 mg/L 5-FC medium and eliminated in 0.1 mg/L 5-FC medium.

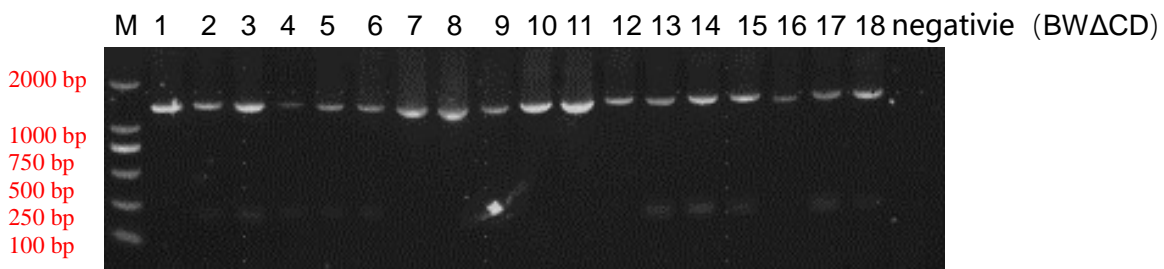
Supplementary Figure 3



gY9s-HmaS^{mut}-Bio177 mutant library plates 1, 2, 3

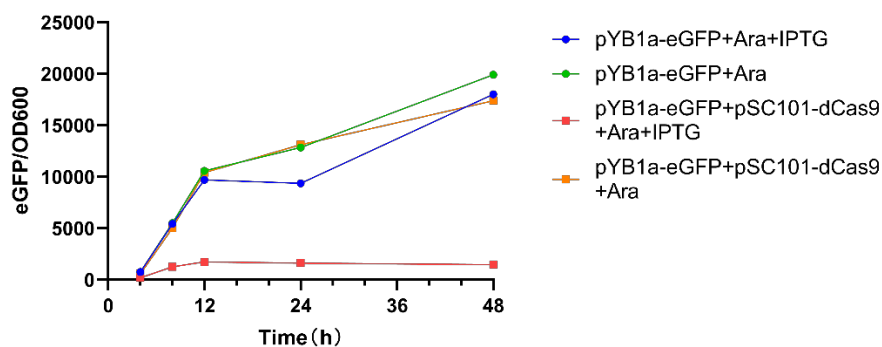
Supplementary Figure 3. By using the error-prone PCR with a low-fidelity DNA polymerase and adjusting Mn^{2+} and Mg^{2+} concentrations and their ratio, the mutation rate was manipulated. The Golden Gate assembly procedure was optimized to increase the insertion probability of mutated fragments into the vector. A mutant library with 10^5 clones was constructed.

Supplementary Figure 4.



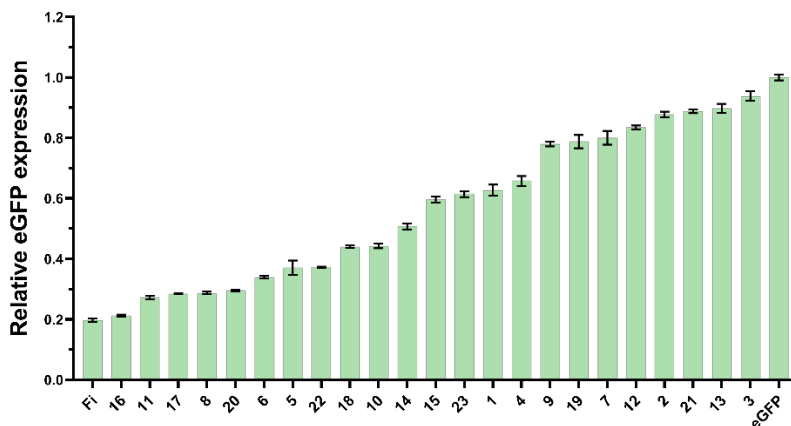
Supplementary Figure 4. Colony PCR to screen for plasmids containing HmaS mutants.

Supplementary Figure 5



Supplementary Figure 5. The inhibitory effect of pSC101-dCas9-eGFP plasmid on CRISPRi. IPTG was used to induce sgRNA expression and Ara was used to induce eGFP expression.

Supplementary Figure 6



Supplementary Figure 6. Evaluation of eGFP inhibition by a sgRNA library using the CRISPRi technology. After co-transformation of R6K-dCas9-eGFP mutant library plasmid and pYB1a-eGFP plasmid, 24 colonies were selected for fluorescence characterization and DNA sequencing of the sgRNA 7-8bp spacer. Fi means Fully Interference, transferred to the strain that containing the non-mutated R6K-dCas9-eGFP plasmid.