
SUPPLEMENTARY INFORMATION

A CRISPR-Cas9 Assisted Non-Homologous End-Joining Strategy for One-step Engineering of Bacterial Genome

Tianyuan Su¹, Fapeng Liu¹, Pengfei Gu¹, Haiying Jin¹, Yizhao Chang¹, Qian Wang², Quanfeng Liang¹, Qingsheng Qi^{1,2}

¹State Key Laboratory of Microbial Technology, Shandong University, Jinan 250100, People's Republic of China

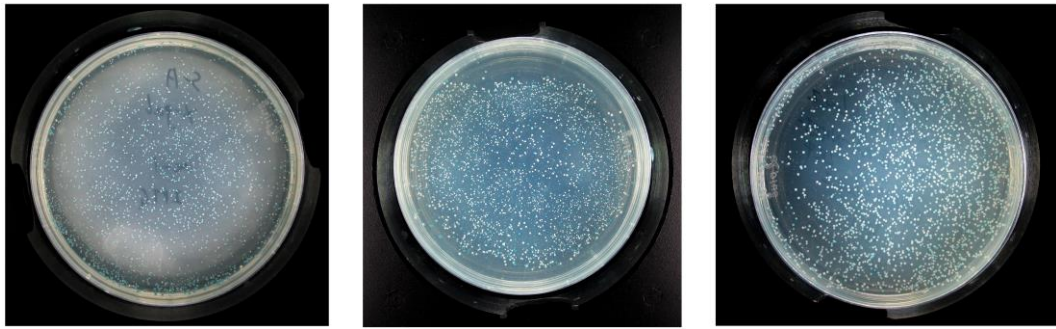
²National Glycoengineering Center, Shandong University, Jinan, 250100, People's Republic of China

* Corresponding author: Qingsheng Qi

E.mail: qiqingsheng@sdu.edu.cn

Tel: +86-531-88365628;

Fax: +86-531-88362897



HindIII-linearized pUC19

SmaI-linearized pUC19

CRISPR-cleaved pUC-lacZ

Supplementary Figure 1. Blue-white screening of the transformants to determine the efficiency and fidelity of the heterogenous NHEJ pathway in *E. coli*. White colonies represent the *lacZ* genotype and blue colonies represent the *lacZ*⁺ genotype.

HindIII-linearized pUC19

WT	TTACGCCAAGCTTGCATGCCTGCAGGTCGACTCTAGAGGATCCCGGGGTACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG
8 random selected white colonies	TTACGCCAAGCTTGCATGCCTGCAGGTCGACTCTAGA GG -----GTACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAGGTCGACTCTAGAGGATCC CGT -----AGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAGGTCGACTCTAGAGGATCC CGT -----AGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAGGTCGACTCTAGA GG -----GTACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAG CGT -----GAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAG CGT -----ACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAG GGT -----ACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG TTACGCCAAGCTTGCATGCCTGCAGG TC -----ACCGAGCTCGAATTCAGTGGCCGTCGTTTTACAACGTCGTG N₈

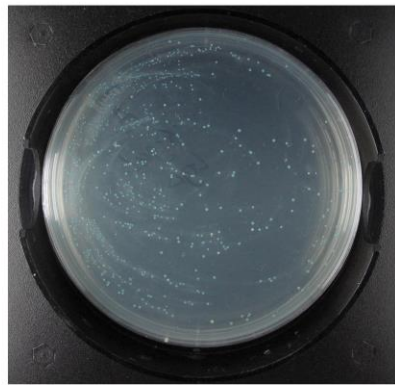
SmaI-linearized pUC19

WT	TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA
8 random selected white colonies	TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA TTGTGTGGAATTGTGAGCGGATAACAATTTACACAGGAAACAGCTATGACCATGATTACGCC AAGCTT GCATGCCTGCAGGTCGACTCTA N₈

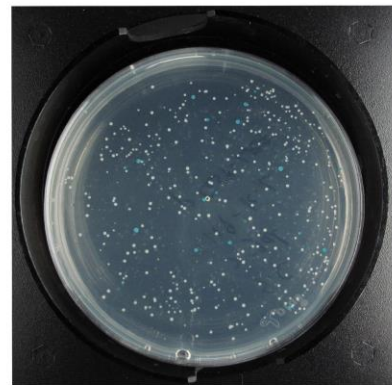
CRISPR-cleaved pUC-lacZ

WT	GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC
8 random selected white colonies	GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC GGGTACCGAGCTCGAATTCAGTGG CCGTCGTTTTACAACGCTGCTGACTGGGAAAACC CTGGCGTTACCCAACCTTAATCGCCTTGCAGCAC N₈

Supplementary Figure 2. Sanger sequencing analysis of 8 random selected white colonies (*lacZ*⁻ genotype) generated by repair of *HindIII*/*SmaI*-linearized pUC19 or Cas9-cleaved pUC-lacZ, respectively. The recognition sequences of *HindIII*/*SmaI* and the LR4-CRISPR target are shown in orange. The PAM sequences are shown in red. The light blue color indicates micro-homology sequence for ending-joining and the dashed line highlighted in yellow represents the deleted nucleotides by NHEJ.



MG1655
crRNA

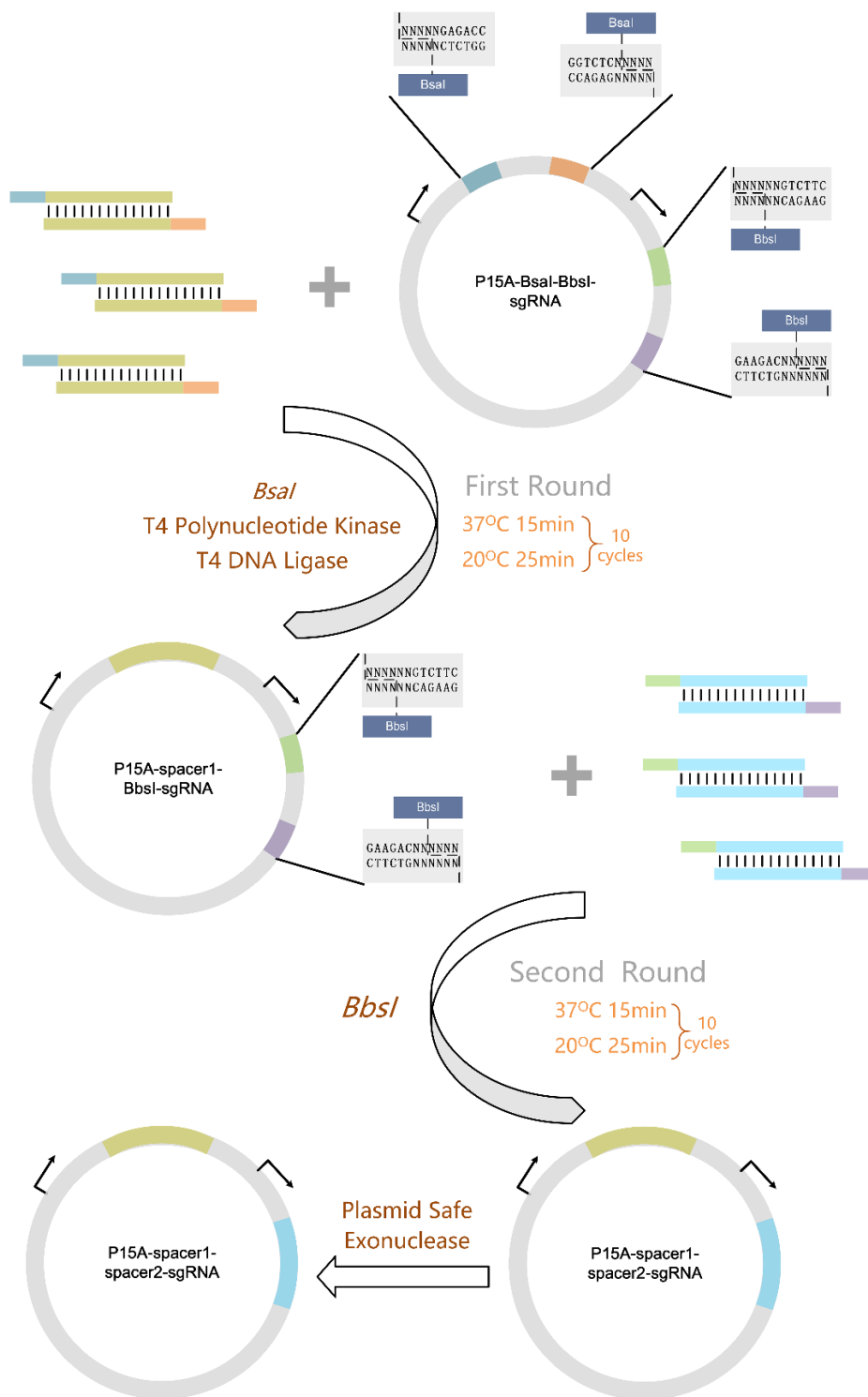


MG1655
sgRNA

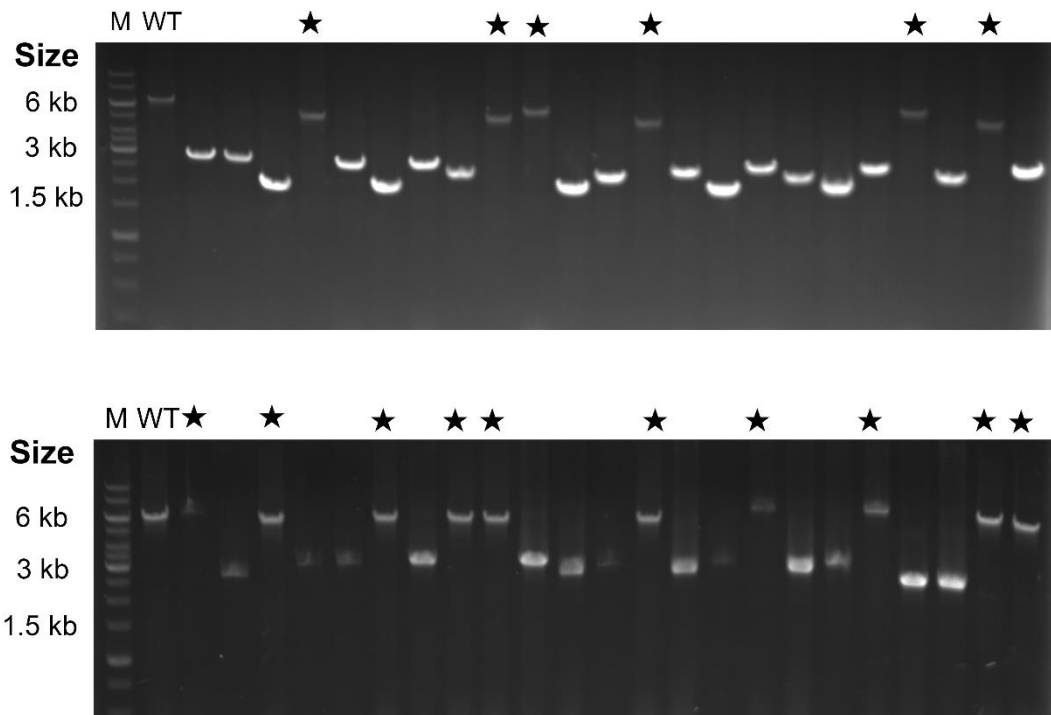
Supplementary Figure 3. Blue-white screening of the transformants to evaluate the efficiency and positivity rate of the improved CA-NHEJ system using L4 target site in *E. coli* MG1655. White colonies represent the *lacZ* genotype and blue colonies represent the *lacZ*⁺ genotype.

WT	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
8 random selected blue colonies	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA
	CGAGGTTTATAGACTATGCTGTTTTGAATGGTCCCAAACGGTTTTCCAGTCACGACGTTGTA AAAACGAGTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC	TTCA

Supplementary Figure 4. Sanger sequencing analysis of the LR4 CRISPR array in 8 random selected blue colonies grown on the X-gal plate. The repeat sequences and the LR4 spacer of CRISPR array are shown in blue and orange, respectively. The dashed line highlighted in yellow indicates the deleted sequences.



Supplementary Figure 5. One-step digestion-ligation method based on golden gate cloning to easy clone of specific sgRNA pair cassettes in plasmid p15A-BsaI-BbsI-sgRNA. The one-step digestion-ligation cloning strategy for ligation of the sgRNA pairs into the p15A-BsaI-BbsI-sgRNA plasmid was accomplished through two rounds iterative golden gate cloning in a 30- μ l reaction system. Tan: spacer 1; Light blue: spacer 2.



Supplementary Figure 6. Gel electrophoresis of the PCR products to distinguish the mutation types of *lacZ* by sgRNA pair L4&LR8. Asterisk indicates the frameshift mutation of *lacZ*.

Supplementary Table 1. Primers used in this study (The homologous sequences used for assembling were in bold, restriction sites were underlined).

Primers	Nucleotide sequence(5'-3')
Primers for plasmids construction	
lacZ-F	CCCAA <u>AAGCTT</u> ACAGCTATGACCATGATTACGGATT
lacZ-R	TCCAGAT <u>CTAGACCTT</u> ACGCGAAATACGGGCAGACAT
Cm-Ori-F	TAGTGTGAGATCTCATCGGGCTTACTCGATGCATGCGCTAACCGTTTTTATCAGGCTCTGGG
Cm-Ori-R	ATCCGTACGCCTGCAGGTCTAGATTAATTAACGCCGGCGGCGGCGCTATGGACAGTTTTCCCTTTGATAT
cas9-F	GCGGCCGCCGCCGGCGTTAATTAATCTAGACCTGCAGGCGTACGGATTACGAAATCATCCTGTGGAGC
cas9-R	CGGTTAGCGCATGCATCGAGTAAGCCCGATGAGATCTCACACTACTCTTCTTTTGCCTATTATAAC
23119-ligd-F	ACCGTA <u>AAGATCT</u> TTTGACAGCTAGCTCAGTCCTAGGTATAATGCTAGCTACTAGAGAAAGAGGAGAAATACTAGATGGGTT CGGCGTCGGAGCA
23119-ligd-R	TTAATTCATTCGCGCACCCACCTCACTGG
mku-F	GCGGCCGGACAAGAAACCCAGTGAGGTGGTGCGCGAATGAATTAAGAGGAGAATACTAGATGCGAGCCATTTGGA CGGG
mku-R	ACAATG <u>ATGCAT</u> TTTAGCGCAAGAAGACAAAAATCACCTTGCGCTAATGCTCTGTTACAGTCACGGAGGCGTTGGGACG
CRISPR-F	TCCAGAT <u>CTAGATGCCT</u> CTAGCACGCGTACCAT
CRISPR-R	TCCAGAT <u>CTAGATGACCGA</u> ATTCAACTCAACAAGT
NHEJ-F	ACCGATA <u>ACTAGT</u> TTTGACAGCTCAGTCCTAGGTATAATGCTAGC
NHEJ-R	ACCGATA <u>ACTAGT</u> TTTAGCGCAAGAAGACAAAAATCACCTTGCGC

Spc-F	TCTGTTGTTTGTCTGGTGAAGTGGATCCCTTAGTAAAGCCCTCGCTAGATT
Spc-R	ACCATCATACTAAATCAGTAAGTTGGCAGCATCACCCGACGGCTCGTTCGCCAGCCAGGACAGA
P15A-F	CGTCGGGTGATGCTGCCAACTTA
P15A-R	ACTTATATCGTATGGGGCTGACTTC
sgRNA-F	AGCACCTGAAGTCAGCCCCATACGATATAAGTCTATAAAAATAGGCGTATCACGAGGC
sgRNA-R	TTACTAAGGGATCCAGTTCACCGACAAACAACAGA

Primers for gene knockout

lac-F	CGATACCGAAGACAGCTCATGT
lac-R	TATCATCGCCGGGCTTGCCCCGT

Primers for fragment deletion analysis

LacZ-JF	ATGTCCGCCGTAGCCCCTCCGATGAT
LacZ-JR	CCAGCAGGAACGGTACTTCAAAC
Lac-JF	AGCGCCTCGTCATCAATACCAAT
Lac-JR	TTCCCACAAGACAACAACCACTCC
LC-JF	CAGAACAATGAGCAGACGGAATA
LC-JR	GGAGGCTAACAGTGTCGAATAAC
MLC-JF	AGCCCTTCCATCAGAGCGACCA
MLC-JR	CACACAACCGGCACAAACCACC

Supplementary Table 2. Plasmids used in this study.

Plasmids	Relevant genotype	Reference
pUC19	Cloning vector, Amp ^R	Lab stock
pCRISPR	pUC19 containing CRISPR array, kan ^R	1
pCas9	pACYC184 containing tracr RNA, <i>cas9</i> and CRISPR array, Cm ^R	1
pwtCas9	pUC19 containing aTc-inducible promoter P _{LtetO-1} expressing <i>cas9</i> , Amp ^R	2
pTKRED	Temperature-conditional replicon containing γ , β , <i>exo</i> (red recombinase), Spc ^R	3
pCP20	Helper plasmid, Cm ^R	4
pUC-lacZ	pUC19 containing <i>lacZ</i> gene, Amp ^R	This study
pCas9 (Ts)	Temperature-conditional replicon containing <i>cas9</i> , Cm ^R	This study
pCas9 (Ts)-NHEJ	pCas9 (Ts) containing p23119- <i>mku-ligd</i> , Cm ^R	This study
pCas9 (Ts)-LR4	pCas9 (Ts) containing LR4-CRISPR array, Cm ^R	This study
pCas9 (Ts)-NHEJ-LR4	pCas9 (Ts)-NHEJ containing LR4-CRISPR array, Cm ^R	This study
pcurCas9 (Ts)-NHEJ	pCas9 (Ts)-NHEJ containing <i>lacI</i> and trc-p15A-sgRNA, Cm ^R	This study
pwtCas9-NHEJ	pwtCas9 containing p23119- <i>mku-ligd</i> , Amp ^R	This study
p15A-gRNA	p15A replicon containing sgRNA, spc ^R	This study
p15A-BsaI-BbsI-gRNA	p15A replicon containing two sgRNA modules for one-step constructing sgRNA pairs, spc ^R	This study
pCRISPR-L4	pCRISPR containing L4 CRISPR array, kan ^R	This study
pCRISPR-LR4	pCRISPR containing LR4 CRISPR array, kan ^R	This study
p15A-L4	p15A-gRNA containing sgRNA L4, spc ^R	This study
p15A-L5	p15A-gRNA containing sgRNA L5, spc ^R	This study
p15A-LR4	p15A-gRNA containing sgRNA LR4, spc ^R	This study
p15A-LR6	p15A-gRNA containing sgRNA LR6, spc ^R	This study

p15A-LR7	p15A-gRNA containing sgRNA LR7, spc ^R	This study
p15A-LR8	p15A-gRNA containing sgRNA LR8, spc ^R	This study
p15A-L4&LR8	p15A-BsaI-BbsI-gRNA containing sgRNA pair L4&LR8, spc ^R	This study
p15A-LI10&LA0	p15A-BsaI-BbsI-gRNA containing sgRNA pair LI10&LA0, spc ^R	This study
p15A-LI10&CR0	p15A-BsaI-BbsI-gRNA containing sgRNA pair LI10&CR0, spc ^R	This study
p15A-ME17&CR0	p15A-BsaI-BbsI-gRNA containing sgRNA pair ME17&CR0, spc ^R	This study

Supplementary Table 3. CRISPR target sequences used in this study.

Spacer name	Nucleotide sequence (5'-3')
L4-CRISPR	GGTTTTCCCAGTCACGACGTTGTAAAACGA
LR4-CRISPR	TCCGCCGTTTGTCCACGGAGAATCCGAC
L4	GTCACGACGTTGTAAAACGA
L5	GTGAGCGAGTAACAACCCGT
LR4	GTTCCCACGGAGAATCCGAC
LR6	CAACGTGACCTATCCCATTA
LR7	CCATCGCCATCTGCTGCACG
LR8	CTCCTGGAGCCCGTCAGTAT
LI10	TACGATGTTCGCAGAGTATGC
LA0	ATTGGCAATAACGTCTGGAT
CR0	CCCGCCACTACTGGAGAGAA
ME17	GTTTCGCGATCGACTCGTAC

Supplementary Note 1

The sequences of core elements in this study

P₁₂₃₁₁₉-ligd-mku

ttgacagctagctcagtcctaggtataatgctagctactagagAAAGAGGAGAAATACTAGATG
GGTTCGGCGTCGGAGCAACGGGTGACGCTGACCAACGCCGACAAGG
TGCTCTATCCCGCCACCGGGACCACAAAGTCCGATATCTTCGACTACT
ACGCCGGTGTTGCCGAAGTCATGCTCGGCCACATCGCGGGACGGCCG
GCGACGCGCAAGCGCTGGCCTAACGGCGTCGACCAACCCGCGTTCTT
CGAAAAGCAGTTGGCGTTGTCGGCGCCGCCTTGGCTGTCACGTGCAA
CGGTGGCGCACCGGTCCGGGACGACGACCTATCCGATCATCGATAGC
GCAACCGGGCTGGCCTGGATCGCCCAACAGGCGGCGCTGGAGGTGC
ACGTGCCGCAGTGGCGGTTTGTTCGCCGAGCCCGGATCAGGTGAGTTA
AATCCGGGCCC GGCAACGCGTTTGGTGTTTCGACCTGGACCCGGGCGA
AGGCGTGATGATGGCCCAGCTGGCCGAGGTGGCGCGCGCGGTTTCGTG
ATCTTCTCGCCGATATCGGGTTGGTCACCTTCCCGGTCACCAGCGGCA
GCAAGGGATTGCATCTGTACACACCGCTGGATGAGCCGGTGAGCAGC
AGGGGAGCCACGGTGTTGGCCAAGCGCGTCGCGCAGCGATTGGAGC
AGGCGATGCCCGCGTTGGTCACCTCGACCATGACCAAAAAGCCTGCGG
GCCGGGAAGGTGTTTGTGGACTGGAGCCAGAACAGCGGCTCGAAGA
CCACCATCGCGCCGTACTCACTACGTGGCCGGACGCATCCGACCGTC
GCGGCGCCACGCACCTGGGCGGAGCTCGACGACCCCGCACTGCGTC
AGCTCTCCTACGACGAGGTGCTGACCCGGATTGCCCGCGACGGCGAT
CTGCTCGAGCGGCTGGATGCCGACGCTCCGGTAGCGGACCGGTTGAC
CCGATAACGCCGCATGCGCGACGCATCGAAAACCTCCCGAGCCGATTCC
CACGGCGAAACCCGTTACCGGAGACGGCAATACGTTTCGTCATCCAGG
AGCATCACGCGCGTCGGCCGCACTACGATTTCCGGCTGGAATGCGAC
GGCGTGCTGGTTTCGTGGGCGGTACCGAAAAACCTGCCCGACAACAC
ATCGGTTAACCATCTAGCGATAACACACCGAGGACCACCCGCTGGAATA
CGCCACGTTTCGAGGGCGCGATTCCCAGCGGGGAGTACGGCGCCGGCA
AGGTGATCATCTGGGACTCCGGCACTTACGACACCGAGAAGTTCCAC
GATGACCCGCACACGGGGGAGGTTCATCGTGAATCTGCACGGCGGCCG

GATCTCTGGGCGTTATGCGCTGATTTCGGACCAACGGCGATCGGTGGCT
GGCGCACCGCCTAAAGAATCAGAAAGACCAGAAGGTGTTTCGAGTTCG
ACAATCTGGCCCCAATGCTTGCCACGCACGGCACGGTGGCCGGTCTA
AAGGCCAGCCAGTGGGCGTTTCGAAGGCAAGTGGGACGGCTACCGGTT
GCTGGTTGAGGCTGACCACGGCGCCGTGCGGCTGCGGTCCCGCAGC
GGGCGCGATGTCACCGCCGAGTATCCGCAATTGCGGGCATTGGCGGA
GGATCTCGCCGATCACCACGTGGTGCTGGACGGCGAGGCCGTTCGTAC
TTGACTCCTCTGGTGTGCCAGCTTCAGCCAGATGCAGAATCGGGGC
CGCGACACCCGTGTCGAGTTCTGGGCGTTCGACCTGCTCTACCTCGA
CGGCCGCGCGCTGCTAGGCACCCGCTACCAAGACCGGCGTAAGCTGC
TCGAAACCCTAGCTAACGCAACCAGTCTCACCGTTCCTCGAGCTGCTGC
CCGGTGACGGCGCCCAAGCGTTTGCGTGCTCGCGCAAGCACGGCTGG
GAGGGCGTGATCGCCAAGAGGCGTGACTCGCGCTATCAGCCGGGCCG
GCGCTGCGCGTCGTGGGTCAAGGACAAGCACTGGAACACCCAGGAA
GTCGTCATTGGTGGCTGGCGCGCCGGGGAAAGGCGGGCGCAGCAGTG
GCGTCCGGTTCGCTGCTCATGGGCATCCCCGGTCCAGGTGGGCTGCAG
TTCGCCGGGCGGGTTCGGTACCGGCCTCAGCGAACGCGAACTGGCCAA
CCTCAAGGAGATGCTGGCGCCGCTGCATACCGACGAGTCCCCCTTCG
ACGTACCACTGCCCGCGCGTGACGCCAAGGGCATCACATATGTCAAGC
CGGCGCTGGTTGCAGAGGTGCGCTACAGCGAGTGGACTCCGGAGGG
CCGGCTGCGTCAATCAAGCTGGCGTGGGCTGCGGCCGGACAAGAAAC
CCAGTGAGGTGGTGCGCGAATGAtactagagATTAAAGAGGAGAATACTA
GATGCGAGCCATTTGGACGGGTTTCGATCGCCTTCGGGCTGGTGAACG
TGCCGGTCAAGGTGTACAGCGCTACCGCAGACCACGACATCAGGTTC
CACCAGGTGCACGCCAAGGACAACGGACGCATCCGGTACAAGCGCGT
CTGCGAGGCGTGTGGCGAGGTGGTTCGACTACCGCGATCTTGCCCGGG
CCTACGAGTCCGGCGACGGCCAAATGGTGGCGATCACCGACGACGAC
ATCGCCAGCTTGCTGAAGAACGCAGCCGGGAGATCGAGGTGTTGGA
GTTTCGTCCCCGCCGCCGACGTGGACCCGATGATGTTTCGACCGCAGCT
ACTTTTTGGAGCCTGATTTCGAAGTCGTCGAAATCGTATGTGCTGCTGG
CTAAGACACTCGCCGAAACCGACCGGATGGCGATCGTGCAATTCACGC
TGCGCAACAAGACCAGGCTGGCGGCGTTGCGCGTCAAGGATTCGGC
AAGCGAGAGGTGATGATGGTGCACACGTTGCTGTGGCCCGATGAGAT
CCGCGACCCCGACTTCCCGGTGCTGGACCAGAAGGTGGAGATCAAAC

CCGCGGAACTCAAGATGGCCGGCCAGGTGGTGGACTCGATGGCCGAC
GACTTCAATCCGGACCGCTACCACGACACCTACCAGGAGCAGTTACAG
GAGCTGATCGACACCAAACCTCGAAGGTGGGCAGGCATTTACCGCCGA
GGACCAACCGAGGTTGCTGGACGAGCCCGAAGACGTCTCCGACCTGC
TCGCCAAGCTGGAGGCCAGCGTGAAGGCCGCGCTCGAAGGCCAACTCA
AACGTCCCAACGCCTCCGTGA

tracRNA-Cas9-CRISPR array

AAAAAAGCACCGACTCGGTGCCACTTTTTCAAGTTGATAACGGACTA
GCCTTATTTTAACTTGCTATGCTGTTTTGAATGGTTCCAACAAGATTAT
TTTATAACTTTTATAACAAATAATCAAGGAGAAATTCAAAGAAATTTATC
AGCCATAAAACAATACTTAATACTATAGAATGATAACAAAATAAACTACT
TTTTAAAAGAATTTTGTGTTATAATCTATTTATTATTAAGTATTGGGTAAT
ATTTTTGAAGAGATATTTTGAAAAAGAAAAATTAAAGCATATTAACCT
AATTCGGAGGTCATTAAACTATTATTGAAATCATCAAACCTCATTATG
GATTTAATTTAACTTTTTATTTAGGAGGCCAAAATGGATAAGAAATA
CTCAATAGGCTTAGATATCGGCACAAATAGCGTCGGATGGGCGGTGAT
CACTGATGAATATAAGGTTCCGTCTAAAAAGTTCAAGGTTCTGGGAAA
TACAGACCGCCACAGTATCAAAAAAATCTTATAGGGGCTCTTTTATTT
GACAGTGGAGAGACAGCGGAAGCGACTCGTCTCAAACGGACAGCTCG
TAGAAGGTATACACGTCGGAAGAATCGTATTTGTTATCTACAGGAGATT
TTTTCAAATGAGATGGCGAAAGTAGATGATAGTTTCTTTCATCGACTTG
AAGAGTCTTTTTTGGTGGAAGAAGACAAGAAGCATGAACGTCATCCTA
TTTTTGGAATATAGTAGATGAAGTTGCTTATCATGAGAAATATCCAAC
TATCTATCATCTGCGAAAAAATTGGTAGATTCTACTGATAAAGCGGAT
TTGCGCTTAATCTATTTGGCCTTAGCGCATATGATTAAGTTTCGTGGTC
ATTTTTTGATTGAGGGAGATTTAAATCCTGATAATAGTGATGTGGACAA
ACTATTTATCCAGTTGGTACAAACCTACAATCAATTATTTGAAGAAAAC
CCTATTAACGCAAGTGGAGTAGATGCTAAAGCGATTCTTTCTGCACGA
TTGAGTAAATCAAGACGATTAGAAAATCTCATTGCTCAGCTCCCCGGT
GAGAAGAAAAATGGCTTATTTGGGAATCTCATTGCTTTGTCATTGGGT
TTGACCCCTAATTTTAAATCAAATTTTGATTTGGCAGAAGATGCTAAAT
TACAGCTTTCAAAGATACTTACGATGATGATTTAGATAATTTATTGGC
GCAAATTGGAGATCAATATGCTGATTTGTTTTTGGCAGCTAAGAATTTA

TCAGATGCTATTTTACTTTTCAGATATCCTAAGAGTAAATACTGAAATAAC
TAAGGCTCCCCTATCAGCTTCAATGATTAACGCTACGATGAACATCAT
CAAGACTTGACTCTTTTAAAAGCTTTAGTTTCGACAACAACCTCCAGAA
AAGTATAAAGAAATCTTTTTTGATCAATCAAAAAACGGATATGCAGGTT
ATATTGATGGGGGAGCTAGCCAAGAAGAATTTTATAAATTTATCAAACC
AATTTTAGAAAAATGGATGGTACTGAGGAATTATTGGTGAAACTAAAT
CGTGAAGATTTGCTGCGCAAGCAACGGACCTTTGACAACGGCTCTATT
CCCCATCAAATTCACTTGGGTGAGCTGCATGCTATTTTGAGAAGACAA
GAAGACTTTTATCCATTTTAAAAGACAATCGTGAGAAGATTGAAAAA
ATCTTGACTTTTCGAATTCCTTATTATGTTGGTCCATTGGCGCGTGGCA
ATAGTCGTTTTGCATGGATGACTCGGAAGTCTGAAGAAACAATTACCC
CATGGAATTTTGAAGAAGTTGTTCGATAAAGGTGCTTCAGCTCAATCAT
TTATTGAACGCATGACAACTTTGATAAAAAATCTTCCAAATGAAAAAGT
ACTACCAAACATAGTTTGCTTTATGAGTATTTTACGGTTTATAACGAAT
TGACAAAGGTCAAATATGTTACTGAAGGAATGCGAAAACCAGCATTTC
TTTCAGGTGAACAGAAGAAAGCCATTGTTGATTTACTCTTCAAAACAA
ATCGAAAAGTAACCGTTAAGCAATTAAGAAGATTATTTCAAAAAAAT
AGAATGTTTTGATAGTGTTGAAATTTACAGGAGTTGAAGATAGATTTAAT
GCTTCATTAGGTACCTACCATGATTTGCTAAAAATTATTAAGATAAAG
ATTTTTTGGATAATGAAGAAAATGAAGATATCTTAGAGGATATTGTTTTA
ACATTGACCTTATTTGAAGATAGGGAGATGATTGAGGAAAGACTTAAA
ACATATGCTCACCTCTTTGATGATAAGGTGATGAAACAGCTTAAACGTC
GCCGTTATACTGGTTGGGGACGTTTGTCTCGAAAATTGATTAATGGTAT
TAGGGATAAGCAATCTGGCAAAACAATATTAGATTTTTTGAATCAGAT
GGTTTTGCCAATCGCAATTTTATGCAGCTGATCCATGATGATAGTTTGA
CATTAAAGAAGACATTCAAAAAGCACAAGTGTCTGGACAAGGCGATA
GTTTACATGAACATATTGCAAATTTAGCTGGTAGCCCTGCTATTA AAAA
AGGTATTTTACAGACTGTAAAAGTTGTTGATGAATTGGTCAAAGTAATG
GGGCGGCATAAGCCAGAAAATATCGTTATTGAAATGGCACGTGAAAAT
CAGACAACCTCAAAGGGCCAGAAAATTCGCGAGAGCGTATGAAACG
AATCGAAGAAGGTATCAAAGAATTAGGAAGTCAGATTCTTAAAGAGCA
TCCTGTTGAAAATACTCAATTGCAAATGAAAAGCTCTATCTCTATTAT
CTCCAAAATGGAAGAGACATGTATGTGGACCAAGAATTAGATATTAATC
GTTTAAGTGATTATGATGTCGATCACATTGTTCCACAAAGTTTCCTTAA

AGACGATTCAATAGACAATAAGGTCTTAACGCGTTCTGATAAAAAATCGT
GGTAAATCGGATAACGTTCCAAGTGAAGAAGTAGTCAAAAAGATGAAA
AACTATTGGAGACAACCTTCTAAACGCCAAGTTAATCACTCAACGTAAG
TTTGATAATTTAACGAAAGCTGAACGTGGAGGTTTGAGTGAACCTTGAT
AAAGCTGGTTTTATCAAACGCCAATTGGTTGAAACTCGCCAAATCACT
AAGCATGTGGCACAAATTTTGGATAGTCGCATGAATACTAAATACGATG
AAAATGATAAACTTATTTCGAGAGGTTAAAGTGATTACCTTAAAATCTAA
ATTAGTTTCTGACTTCCGAAAAGATTTCCAATTCTATAAAGTACGTGAG
ATTAACAATTACCATCATGCCCATGATGCGTATCTAAATGCCGTCGTTG
GAACTGCTTTGATTAAGAAATATCCAAAACCTTGAATCGGAGTTTGTCTA
TGGTGATTATAAAGTTTATGATGTTTCGTAAAATGATTGCTAAGTCTGAG
CAAGAAATAGGCAAAGCAACCGCAAATATTTCTTTTACTCTAATATCA
TGAACCTTCTTCAAAACAGAAATTACACTTGCAAATGGAGAGATTTCGCA
AACGCCCTCTAATCGAAACTAATGGGGAAACTGGAGAAATTGTCTGGG
ATAAAGGGCGAGATTTTGCCACAGTGCACAAAGTATTGTCCATGCCCC
AAGTCAATATTGTCAAGAAAACAGAAGTACAGACAGGCGGATTCTCCA
AGGAGTCAATTTTACCAAAAAGAAATTCGGACAAGCTTATTGCTCGTA
AAAAAGACTGGGATCCAAAAAATATGGTGGTTTTGATAGTCCAACGG
TAGCTTATTCAGTCCTAGTGGTTGCTAAGGTGGAAAAAGGGAAATCGA
AGAAGTTAAAATCCGTTAAAGAGTTACTAGGGATCACAATTATGGAAA
GAAGTTCCTTTGAAAAAATCCGATTGACTTTTTTAGAAGCTAAAGGAT
ATAAGGAAGTTAAAAAAGACTTAATCATTAAACTACCTAAATATAGTCTT
TTTGAGTTAGAAAACGGTTCGTAAACGGATGCTGGCTAGTGCCGGAGA
ATTACAAAAGGAAATGAGCTGGCTCTGCCAAGCAAATATGTGAATTT
TTTATATTTAGCTAGTCATTATGAAAAGTTGAAGGGTAGTCCAGAAGAT
AACGAACAAAAACAATTGTTTGTGGAGCAGCATAAGCATTATTTAGAT
GAGATTATTGAGCAAATCAGTGAATTTTCTAAGCGTGTTATTTTAGCAG
ATGCCAATTTAGATAAAGTTCTTAGTGATATAACAAACATAGAGACAA
ACCAATACGTGAACAAGCAGAAAATATTATTCATTTATTTACGTTGACG
AATCTTGGAGCTCCCGCTGCTTTTAAATATTTTGATACAACAATTGATC
GTAAACGATATACGTCTACAAAAGAAGTTTTAGATGCCACTCTTATCCA
TCAATCCATCACTGGTCTTTATGAAACACGCATTGATTTGAGTCAGCTA
GGAGGTGACTGAAGTATATTTTAGATGAAGATTATTTCTTAATAACTAA
AAATATGGTATAATACTCTTAATAAATGCAGTAATACAGGGGCTTTTCAA

GACTGAAGTCTAGCTGAGACAAATAGTGCGATTACGAAATTTTTTAGA
CAAAAATAGTCTACGAGGTTTTAGAGCTATGCTGTTTTGAATGGTCCCA
AAACTGAGACCAGTCTCGGAAGCTCAAAGGTCTCGTTTTAGAGCTATG
CTGTTTTGAATGGTCCCAAAC

P_{J23119}-*Bsal*-sgRNA

TTGACAGCTAGCTCAGTCCTAGGTATAATACTAGTTGAGACCAGTCTC
GGAAGCTCAAAGGTCTCGTTTTAGAGCTAGAAATAGCAAGTTAAAAT
AAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTGC

- 1 Jiang, W. Y., Bikard, D., Cox, D., Zhang, F. & Marraffini, L. A. RNA-guided editing of bacterial genomes using CRISPR-Cas systems. *Nat Biotechnol* **31**, 233-239, doi:10.1038/nbt.2508 (2013).
- 2 Qi, L. S. *et al.* Repurposing CRISPR as an RNA-guided platform for sequence-specific control of gene expression. *Cell* **152**, 1173-1183, doi:10.1016/j.cell.2013.02.022 (2013).
- 3 Kuhlman, T. E. & Cox, E. C. Site-specific chromosomal integration of large synthetic constructs. *Nucleic Acids Res* **38**, doi:ARTN e9210.1093/nar/gkp1193 (2010).
- 4 Cherepanov, P. P. & Wackernagel, W. Gene disruption in *Escherichia coli*: TcR and KmR cassettes with the option of Flp-catalyzed excision of the antibiotic-resistance determinant. *Gene* **158**, 9-14 (1995).