

Supplementary Figure S1. The growth curves of different strains. A) The growth curves of E. coli BW25113, BW Δ recA. B) The growth curves of *E. coli* MG Δ hsdR and MG Δ hsdR-recA. C) The effects of RecA complementation to the growth of BW Δ recA. D) The effects of RecA complementation to the growth of XL1-Blue MRF'. Data are averages of three samples with standard deviations (error bars).

Strains and Plasmids	Functions and features	Origin
E. coli strains		
XL1-Blue MRF'	∆(mcrA)183 ∆(mcrCB-hsdSMR-mrr)173 endA1 supE44 thi-1 recA1 gyrA96 relA1 lac [F' proAB lacIªZ⊿M15 Tn10 (Tet ^r)]	Agilent
MG1655	K-12 F ⁻ λ^- ilvG ⁻ rfb-50 rph-1	NBRP E.coli, Japan Coli Genetic
W3110	$F^{-}\lambda^{-}$ rph-1 INV(rrnD, rrnE)	Stock Center (CGSC)
	$LacI^{+} rrnB_{T14} \Delta lacZ_{WJ16} hsdR514$	
BW25113	ΔaraBAD _{AH33} ΔrhaBAD _{LD78} rph-1 Δ(araB– D)567 Δ(rhaD–B)568 ΔlacZ4787(::rrnB-3) hsdP514 rph 1	CGSC
Mach1	Str.W $\Delta recA1398 endA1$ fhuA $\Phi 80\Delta(lac)M15 \Delta(lac)X74 hsdR(r_Km_K^+)$	Invitrogen
Omnimax2	F ⁽ [proAB ⁺ lacI ^q lacZ Δ M15 Tn10(Tet ^K) Δ (ccdAB)] mcrA Δ (mrr-hsdRMS- mcrBC) φ 80(lacZ) Δ M15 Δ (lacZYA- argF)U169 endA1 recA1 glnV44 thi- 1 gyrA96(Nal ^R) relA1 tonA panD	Invitrogen
Stbl2	F- endA1 glnV44 thi-1 recA1 gyrA96 relA1 Δ (lac-proAB) mcrA Δ (mcrBC-hsdRMS-mrr) λ^{-} F ⁻ mcrB mrr hsdS20 (rp ⁻ mp ⁻) recA13	Invitrogen
Stb13	supE44 ara-14 galK2 lacY1 proA2 rpsL20 (Str ^R) xvl-5 λ^{-} leu mtl-1	Invitrogen
Mutant of MG1655		
ΔhsdR	The <i>recA</i> gene deletion mutant of <i>E. coli</i> MG1655	This study
∆hsdR-recA	The genes <i>hsdR</i> and <i>recA</i> were deleted in <i>E</i> . <i>coli</i> MG1655	
Mutant of W3110		
ΔhsdR	The <i>recA</i> gene deletion mutant of <i>E. coli</i> W3110	This study
∆hsdR-recA	The genes <i>hsdR</i> and <i>recA</i> were deleted in <i>E</i> . <i>coli</i> W3110	This study
Mutants of BW25113		

7 Supplementary Table S1. The strains and plasmids used in this study

BW∆recA	The <i>recA</i> gene deletion mutant of <i>E. coli</i> BW25113	
BW3KG	Δ endA, Δ fhuA, Δ galE	This study
BW3KD	Δ endA, Δ fhuA, Δ deoR	This study
BW4K	Δ endA, Δ fhuA, Δ galE, Δ deoR	This study
Strains with		-
Plasmids		
BW25113/vector	BW25113 harboring pCL1920	This study
BW∆recA/vector	BW∆recA harboring pCL1920	This study
BW∆recA/recA	BW∆recA harboring pCL1920-Pnat-recA	This study
XL1-Blue	VI 1 Dive MDE' berbering aCI 1020	This study.
MRF'/vecor	XLI-Blue MRF harboring pCL1920	This study
XL1-Blue	XL1-Blue MRF' harboring pCL1920-Pnat-	This starder
MRF'/recA	recA	This study
Plasmids		
"Dluggerint CV-	Abbreved as pSK-, pUC18 ori, Amp ^R , used	Stratagona
pbluescript SK	to test TE of competent cells	Stratagene
pCL1920	pSC101 ori, Spc ^R	Addgene
pBBR1MCS5	Abbreved as pMCS5, pBBR ori, Gm ^R	Addgene
pBR322	pBR322 ori, Amp ^R and Tet ^R	Addgene
pACYC184	p15A ori, Cmr ^R and Tet ^R	Addgene
pKD4	R6K ori, Kan ^R and Amp ^R , used for gene deletion in <i>E. coli</i>	Addgene
pTKred	pSC101 ori, temperature sensitive, Spc ^R , used for gene deletion in <i>E. coli</i>	Addgene
pCP20	pSC101 ori, temperature sensitive, Amp^{R} , used for gene deletion in <i>E. coli</i>	Addgene
pSK::Pkat-eGFP	derived from pSK-, phbCAB genes under the control of 5 tac promoters	This study
pCL1920::Pnat-	The <i>recA</i> gene of <i>E. coli</i> was cloned in pCL 1920 with its native promoter (Pnat)	This study
pCL1920::NKan- Gm-CKan-300	Plasmid pCL1920 was introduced with 300 bp direct repeat, used to test intramolecular recombination occurring on plasmid DNA	This study
pBR322::NKan- Gm-CKan-300	Plasmid pBR322 was introduced with 300 bp direct repeat, used to test intramolecular recombination occurring on plasmid DNA	This study
pBBR1MCS5::NK an-Gm-CKan-300	Plasmid pMCS5 was introduced with 300 bp direct repeat, used for test intramolecular recombination occurring on plasmid DNA	This study
pSK::NKan-Gm- CKan-300	Plasmid pSK- was introduced with 300 bp direct repeat, used for test intramolecular recombination occurring on plasmid DNA	This study

pCL1920::HR3000	Plasmid pCL1920 cloned with 3000 bp sequence with homology to BW25113 genome, used to test the recombination occurring between plasmid and its	This study
pBR322::HR3000	chromosome Plasmid pBR322 cloned with 3000 bp sequence with homology to BW25113 genome, used to test the recombination occurring between plasmid and its chromosome	This study
pMCS5::HR3000	Plasmid pMCS5 cloned with 3000 bp sequence with homology to BW25113 genome, used to test the recombination occurring between plasmid and its chromosome	This study
pSK::HR3000	Plasmid pMCS5 cloned with 3000 bp sequence with homology to BW25113 genome, used to test the recombination occurring between plasmid and its chromosome	This study
pCL1920::2TR-300	Plasmid pCL1920::Pkat-eGFP was introduced two 300 bp tandem repeats, used for test intramolecular recombination occurring on plasmid DNA	This study
pCL1920::3TR-300	Plasmid pCL1920::Pkat-eGFP was introduced three 300 bp tandem repeats, used for test intramolecular recombination occurring on plasmid DNA	This study
pCL1920::2TR- 1000	Plasmid pCL1920::Pkat-eGFP was introduced two 300 bp tandem repeats, used for test intramolecular recombination occurring on plasmid DNA	This study
pCL1920::3TR- 1000	Plasmid pCL1920::Pkat-eGFP was introduced three 300 bp tandem repeats, used for test intramolecular recombination occurring on plasmid DNA	This study

NO.	Primers	SEQUENCES	Purposes and characteristics
1		GAACATATTGACTA TCCGGTATTACCCG GCATGACAGGAGT	
I. recA-del-fr	recA-del-Ir	AAAAATGGCTCAGC ATTACACGTCTTGA GCGAT	Primers used to amplify DNA fragments from pKD4, and
2.	recA-del-rev	CAGATGCGACCCTT GTGTATCAAACAAG ACGATTAAAAAATCT TCGTTAGTGGAACA CTTAACGGCTGACA TG	used to knock out the <i>recA</i> gene
3.	recA-out-fr	TACCGATATTGCCG GTAGCT	Primers used to check if recA
4.	recA-out-rev	GTCGATGTCACATT	gene was successfully deleted
5.	endA-del-fr	CAGCTTTCGCTACG TTGCTGGCTCGTTT TAACACGGAGTAA GTGATGTACCAGCA TTACACGTCTTGAG CGAT GGGGTTAACAAAA	Primers used to amplify DNA fragments from pKD4, and used to knock out the <i>endA</i>
6.	endA-del-rev	AGAATCCCGCTAGT GTAGGTTAGCTCTT TCGCGCCTGGGAAC ACTTAACGGCTGAC ATG	gene
7.	endA-fr-out	GTGGTACCGCACGA ACTGGCA	Primers used to check if <i>endA</i>
8.	endA-rev-out	GCGACATCACCTGA CCGAGG	gene was successfully deleted
9.	fhuA-del-fr	ATTCTCGTTTACGT TATCATTCACTTTA CATCAGAGATATAC CAATGGCGCAGCAT TACACGTCTTGAGC GAT	Primers used to amplify DNA fragments from pKD4, and used to knock out the <i>fhuA</i> gene
10.	fhuA-del-rev	TGGGCACGGAAATC CGTGCCCCAAAAGA	gene

Supplementary Table S2. The oligoes used in this study

		GAAATTAGAAACG	
		GAAGGTTGCGGAA	
		CACTTAACGGCTGA	
		CATG	
11	flow A cout for	TGTGCCAGCAGAGC	
11.	InuA-out-Ir	GAGATG	Primers used to check if <i>fhuA</i>
10	G A	ACCAGTTCACGCAC	gene was successfully deleted
12.	InuA-out-rev	GGTCA	
		TTATGCTATGGTTA	
		TTTCATACCATAAG	
10		CCTAATGGAGCGAA	
13.	galE-del-fr	TTATGAGACAGCAT	
		TACACGTCTTGAGC	Primers used to amplify DNA
		GAT	fragments from pKD4, and
		TCAACGGGATTAAA	used to knock out the galE
		TTGCGTCATGGTCG	gene
1.4		TTCCTTAATCGGGA	
14.	galE-del-rev	TATCCCTGGGAACA	
		CTTAACGGCTGACA	
		TG	
1 -		GATTAGCGAAGGTG	
15.	galE-out-fr	AAACG	
			Primers used to check if <i>galE</i>
1.0	15	GATTTCCGTCAATG	gene was successfully deleted
16.	galE-out-rev	CTGCA	
		AGTGTAGTATTGAG	
		CGGCTCGCTTCAAT	
17		AACTATTCAGAGGG	
17.	deoR del-Fr	ATTATGGAACAGCA	
		TTACACGTCTTGAG	Primers used to amplify DNA
		CGAT	fragments from pKD4, and
		GATGGCGCGAAAC	used to knock out the <i>deoR</i>
		GTCATCCGGTTATA	gene
10	1	CGTCATTAATACAT	-
18.	deoR del-rev	CAACTTAATGGAAC	
		ACTTAACGGCTGAC	
		ATG	
10	l. D. (C	GCGATCACGGTACG	
19.	deok -out-fr	GTGAT	Primers used to check if <i>deoR</i>
20		CTCAGTGACCATAC	gene was successfully deleted
20.	deoR -out-rev	CTCAGTGACCATAC CGCGT	gene was successfully deleted
20.	deoR -out-rev	CTCAGTGACCATAC CGCGT GATCCTCTAGAGTC	gene was successfully deleted Primers used to amplify

22.	pcl1920-rev- Pnp	GATCGCGTATGCCG CCATG	pCL1920 as template, which was used to construct pCL1920::Pnp-recA
23.	pnp-fr-recA	CATGGCGGCATACG CGATCACGCGGATT TGTCACCTACAG CAGGTCGACTCTAG	Primers used amplify F-recA with its native promoter and use BW25113 genome as
24.	pnp-rev-recA	AGGATCTTAAAAAT CTTCGTTAGTTTCT GCTACGCC	template, which was used to construct pCL1920::Pnp-recA
25.	psk-kan-F	GCGGGACTCTGGGG TTCGAAATG	Primers used amplify linearized vector with pSK-
26.	psk-kan-R	GCGAAACGATCCTC ATCCTGTCTCT	Kan (Xia et al., 2019) as template, which was used for pSK::Pkat-eGFP cloning
27.	Pkat-eGFP-F30	GGCGGCCGCTCTAG AACTAGTGGATCCC CCACTGGGCTATCT GGACAAGGG	Primers used amplify Pkat- eGFP fragment with SmaI treated pSK::Pkat-eGFP (Xia et al. 2019) as template, which
28.	Pkat-eGFP- R30	GATAAGCTTGATAT CGAATTCCTGCAGC CCGCATTCTGCCGA CATGGAA	was used for pSK::Pkat-eGFP cloning with 30 bp homologous ends
29.	pMCS5-fr-spc	CGGCAAATAACAAT TCGTTCAAGCCGAG ATC	Primers used amplify linearized vector with
30.	pMCS5-rev- spc	CTTCCCTCATCGTT GCTGCTCCATAACA TCA AGCAGCAACGATG	pBBR1MCS5 as template, which was used to construct pBBR1MCS5::Spc
31.	spc-fr-pMCS5	AGGGAAGCGGTGA TCGC	Primers used amplify F-spc fragment with pCL1920 as
32.	spc-rev- pMCS5	GAACGAATTGTTAT TTGCCGACTACCTT GGTGATCT	template, which was used to construct pBBR1MCS5::Spc
33.	NKan-fr	GGGAAAACGCAAG CGCAAAGAGAAAG GCTTCCATGTCGCC	Primers used amplify F-NKan fragment with pKD4 as
34.	NKan-rev	AGAATGCTACGTGC TCGCTCGATGCGA	construst NKan-Gm-CKan- 300bp fragment
35.	Gm-fr	AGCATTCTGCCGAC ATGGAAGC	Primers used amplify F-Gm fragment with pBBR1MCS5 as
36.	Gm-rev	GACAAAAAAGAACC	template, which was used to

		GGCGTTGTGACAAT	construst NKan-Gm-CKan-
		TTACCG	300bp fragment
37	CKan fr	GTTCTTTTTGTCAA	Primers used amplify F-CKan
57.	CKall-II	GACCGACCTGTCC	fragment with pKD4 as
			template, which was used to
38.	CKan-rev		construst NKan-Gm-CKan-
		GGIGGAA	300bp fragment
20	11000	TTAAGCCAGCCCCG	Primers used amplify
39.	pc11920-fr	ACACC	linearized vector with
			pCL1920 as template, which
10		CTGTCGTGCCAGCT	was used to construct
40.	pcl1920-rev	GCAT	pCL1920::NKan-Gm-CKan-
			300 and pCL1920::HR3000
		TAATGCAGCTGGCA	Primers used amplify NKan-
	NKan-Gm-	CGACAGGGGAAAA	Gm-CKan-300bp fragment
41.	CKan-300-fr-	CGCAAGCGCAAAG	with F-NKan, F-Gm and F-
	pCL1920	AGAAAGC	CKan as template, which was
	NKan-Gm-	GGGTGTCGGGGCTG	used to construct
42	CKan-300-rev-	GCTTAACATAGAAG	pCL1920NKan-Gm-CKan-
. 2.	pCL1920	GCGGCGGTGGAATC	300
	pellij20	TAATGCAGCTGGCA	200
	HR3000-fr-	CGACAGATGCCTTT	Primers used amplify HR3000
43.	ncl1920	ТАСАСТТССТСААС	fragment with BW25113
	peni)20	GC	genome as template which
		GGGTGTCGGGGCTG	was used to construct
44	HR3000-rev-	GCTTAACTCGGCTT	nCI 1920. HR 3000
	pcl1920	GTTGACCACCAT	peer/20iik3000
		GGCACCTCGCTAAC	Primers used amplify
45.	pBR322-fr	GGATTCA	linearized vector with pBR322
		OUATICA	as template, which was used to
		GTGATACCCCTATT	construct pBR322::NKan-Gm-
46.	pBR322-rev	ТТТАТАССТТА	CKan 300 and
			nBR322HR3000
		ТАТААААТАСССС	p BR 322 IR 3000
	NKan-Gm-	TATCACGGGAAAAC	Primers used amplify NKan-
47.	CKan-300-fr-	GCAAGCGCAAAGA	Gm-CKan-300bp fragment
	pBR322	GAAAGC	with F-NKan, F-Gm and F-
	NKan Gm	GAAAOC	CKan as template, which was
18	$CK_{an}=300 ray$	GGTGCCCATAGAAG	used to construst
4 0.	nRR277	GCGCCCCATAGAAG	pBR322::NKan-Gm-CKan-300
	PDR322 HR3000 fr	GTGACCGCCCCCCTT	
49.	nRR277		Primers used amplify HR3000
	HR3000 rov		fragment with BW25113
50.	nBR300		genome as template, which
	PD1322		

		GGGCTGCAGGAATT	was used to construst pBR322::HR3000 Primers used amplify
51.	pSK-fr	CGATATCAAG	linearized vector with
52.	pSK-rev	GGGGGGATCCACTAG TTCTAGAGC	pBluescript SK ⁻ as template, which was used to construct pSK::NKan-Gm-CKan-300
53.	NKan-Gm- CKan-300-fr- pSK	CTAGAACTAGTGGA TCCCCCGGGAAAAC GCAAGCGCAAAGA GAAAGC	Primers used amplify NKan- Gm-CKan-300bp fragment with F-NKan, F-Gm and F-
54.	NKan-Gm- CKan-300-rev- pSK	ATATCGAATTCCTG CAGCCCCATAGAAG GCGGCGGTGGAATC	CKan as template, which was used to construst pSK::NKan- Gm-CKan-300
55.	pSK-fr- HR3000	GGGCTGCAGGAATT CGATATCAAG	Primers used amplify linearized vector with
56.	pSK-rev- HR3000	CTGGCCGTCGTTTT ACAACG	which was used to construct pSK::HR3000
57.	HR3000-fr- pSK	CGTTGTAAAACGAC GGCCAGATGCCTTT TACACTTGGTCAAC GC	Primers used amplify HR3000 fragment with BW25113 genome as template, which
58.	HR3000-rev- pSK	ATATCGAATTCCTG CAGCCCCTCGGCTT GTTGACCACCAT	was used to construst pSK::HR3000
59.	pMCS5::Spc-fr	GCGTTAATATTTTG TTAAAATTCGCG	Primers used amplify linearized vector with pBBR1MCS5::Spc as
60.	pMCS5::Spc- rev	TAAGCATTCTGCCG ACATGGA	template, which was used to construct pBBR1MCS5- Spc::NKan-Gm-CKan-300 and pBBR1MCS5-Spc::HR3000
61.	NKan-Gm- CKan-300-fr- pMCS5	CTGGTGCTGGGATT ATGATG	Primers used amplify NKan- Gm-CKan-300bp fragment with F-NKan, F-Gm and F-
62.	NKan-Gm- CKan-300-rev- pMCS5	GATATTGTTCAGCG CGGC	used to construst pBBR1MCS5-Spc::NKan-Gm- CKan-300
63.	HR3000-fr- pMCS5	TACCGATGCGATGG CCTAC	Primers used amplify HR3000 fragment with BW25113

64.	HR3000-rev- pMCS5	TCAAGTTCCTCTGC TGTAAGT	genome as template, which was used to construst pBBR1MCS5-Spc::HR3000
65.	pCL-fr-Ntac	TATGGTGCACTCTC AGTACAATCT	Primers used amplify linearized vector with pCL 1920 as template, which
66.	pCL-rev-Ntac	TCCAGCAAAGGTCT AGCAGAA	was used for pCL1920::Ntac- eGFP cloning with 20bp
67.	Ntac-fr-pCL	TCTGCTAGACCTTT GCTGGA	Primers used amplify Ntac fragment with pNTG as
68.	Ntac-fr-pCL	TGTACTGAGAGTGC ACCATATGC	pCL1920::Ntac-eGFP cloning with 20bp homologous ends
69.	pCL-fr-2TR	GAATCGTTTTCCGG GACGC	Primers used amplify linearized vector with pCL1920::pkat-eGFP as
70.	pCL-rev-2TR	CGAAGCCCAACCTT TCATAGAAG	template, which was used to construct pCL1920::2TR- 300bp, pCL1920::2TR-1000bp
71.	2TR-300-fr- pCL	GGGAACTATAAGA CACGTGCTACATCA TGGCAGACAAACA AAAG	Primers used amplify TR300 fragment with pCL1920::pkat- eGFP as template, which was
72.	2TR-300-rev- pCL	TCTATTAACAAGTG TATCACTTGGTCGG TCATTTCGAACCCC AGA	used to construct pCL1920::2TR-300bp
73.	pCL-fr-3TR- 300	GTGATACACTTGTT AATAGAATCGAG	Primers used amplify linearized vector with pCL1920::2TR-300bp as
74.	pCL-rev-3TR- 300	GCACGTGTCTTATA GTTCCCGT	template, which was used to construct pCL1920::3TR- 300bp
75.	3TR-300-fr- pCL	GGGAACTATAAGA CACGTGCTACATCA TGGCAGACAAACA AAAG	Primers used amplify TR300 fragment with pCL1920::pkat- eGFP as template, which was
76.	3TR-300-rev- pCL	TCTATTAACAAGTG TATCACTTGGTCGG TCATTTCGAACCCC AGA	used to construct pCL1920::3TR-300bp

	OTTO 1000 C	CCGTGACAGGTCAT	
77.	2TR-1000-fr-	TCAGACGGGACTGG	Primers used amplify TR1000
	pCL	GCTATCTGGACAAG	fragment with pCL1920::pkat-
		CTTACTGGGTGCAT	eGFP as template, which was
	2TR-1000-rev-	TAGCCATTGGTCGG	used to construct
78.	pCL	TCATTTCGAACCCC	pCL1920::2TR-1000bp
	1	AGA	1 1
	pCL-fr-3TR-	TGGCTAATGCACCC	Primers used amplify
79.	1000	AGTAAGGC	linearized vector with
			pCL1920::2TR-1000bp as
~~	pCL-rev-3TR-	GTCTGAATGACCTG	template, which was used to
80.	1000	TCACGGGATA	construct pCL1920::3TR-
			1000bp
		CCGTGACAGGTCAT	1
81.	3TR-1000-fr-	TCAGACGGGACTGG	Primers used amplify TR1000
	pCL	GCTATCTGGACAAG	fragment with pCL1920::pkat-
		CTTACTGGGTGCAT	eGFP as template, which was
00	3TR-1000-rev-	TAGCCATTGGTCGG	used to construct
82.	pCL	TCATTTCGAACCCC	pCL1920::3TR-1000bp
		AGA	
02	nCL fr NTD	GTCTGCTATGTGGT	Primers used amplify
03.	pCL-II-NIR	GCTATCTG	linearized vector with
			pCL1920 as template, which
		GCCTCTTCGCTATT	was used for pCL1920::NTR-
84.	pCL-rev-NTR		300bp and pCL1920::NTR-
		ACUCCA	1000bp cloning with 20bp
			homologous ends
85	NTR_fr_nCI	TGGCGTAATAGCGA	Primers used amplify NTR
05.	NTR-II-pCL	AGAGGC	fragment with pCL1920::NTR-
			300bp and pCL1920::NTR-
			1000bp as template, which was
86	NTR-rev-nCI	CAGATAGCACCACA	used for pCL1920::NTR-300bp
00.	MIK-IEV-PCL	TAGCAGAC	and pCL1920::NTR-1000bp
			cloning with 20bp homologous
			ends

Supplementary Table S3. Positive rates of homologous inserts into four plasmids

		Positive rates			
	Plasmids	BW25113 (%)	BWΔrecA (%)	XL1-Blue MRF' (%)	
Vector	pSK::NKan-Gm- CKan-300bp	97 ± 1	94 ± 0	97 ± 1	
Vector :: NKan-Gm- CKan- 300bp (Fig. 2A) Vector :: HR3000 (Fig. 2B)	pCL1920::NKan-Gm- CKan-300bp	97 ± 0	97 ± 0	94 ± 2	
	pBR322::NKan-Gm- CKan-300bp	97 ± 0	92 ± 1	93 ± 1	
	pMCS5::NKan-Gm- CKan-300bp	97 ± 0	91 ± 1	92 ± 1	
	pSK::HR3000	95 ± 1	95 ± 1	95 ± 0	
	pCL1920::HR3000	95 ± 1	100 ± 0	100 ± 0	
	pBR322::HR3000	100 ± 0	95 ± 1	95 ± 1	
	pMCS5::HR3000	90 ± 0	90 ± 1	100 ± 0	

with BW25113, BW∆recA and XL1-Blue MRF'

^a The data are averages of clone colonies from three tests (with standard deviations).

Plasmids	Strains	Colonies a	Concentration of plasmids before 3 transfers (ng/ul) ^b	Concentration of plasmids after 3 transfers (ng/ul) ^b
pSK::NKan-	BW25113	R1	514.7	520.67
Gm-CKan-	BW25113	R2	606.4	492.6
300bp	BW∆recA	PC	357.13	312.51
pCL1920::NK	BW25113	R1	26.5	14.85
an-Gm-CKan-	BW25113	R2	18.9	22.829
300bp	BW∆recA	PC	27.74	38.76
pBR322::NKa	BW25113	R1	69.937	67.43
n-Gm-CKan-	BW25113	R2	52.518	50.7
300bp	BW∆recA	PC	73.52	55.56
pMCS5::NKa	BW25113	R1	24.49	29
n-Gm-CKan-	BW25113	R2	43.05	43.015
300bp	BW∆recA	PC	29.06	31.05
	BW25113	R 1	236.7	44.8
pSK::HR3000	BW25113	R2	372.68	227.49
-	BW∆recA	PC	500	416.67
CI 1020.	BW25113	R1	38.222	23.33
HR3000	BW25113	R2	32.523	47.552
	BW∆recA	PC	54.34	36.23
"DD200	BW25113	R1	51.029	51.76
рвк322::	BW25113	R2	31.909	28.8
HK3000	BW∆recA	PC	102.04	89.90
MODE	BW25113	R1	32.562	31.24
pMC55::	BW25113	R2	44.176	31.86
HK3000	BW∆recA	PC	36.76	40.01
-CI 1020-	BW25113	R1	30.864	22.13
pCL1920::	BW25113	R2	17.123	16.39
stac-eGFP	BW∆recA	PC	13.298	17.67
CI 1020	BW25113	R1	34.72	84.75
pCL1920::	BW25113	R2	42.02	44.64
31 R -300	BW∆recA	PC	17.79	29.94
CI 1020	BW25113	R1	22.94	17.18
pCL1920::	BW25113	R2	25	15.52
3TR-1000	BW∆recA	PC	18,31	26.73

Supplementary Table S4. The plasmids extracted in the plasmid stability assay

^a R1 and R2 means the plasmids were extracted from two clonies of BW25113, and PC means the plasmid was extracted from $BW\Delta recA$ used as positive control.

 $^{\rm b}$ The plasmids were extracted from 3 mL of overnight cultures and dissolved in 45 ul of H₂O.

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