

1 Robust orthogonal recombination system for versatile
2 genomic elements rearrangement in yeast *Saccharomyces*

3 *Cerevisiae*

4 Qihui Lin^{a,b*}, Hao Qi^{a,b*}, Yi Wu^{a,b}, Yingjin Yuan^{a,b†}

5 a: Key Laboratory of Systems Bioengineering (Ministry of Education), Tianjin University, Tianjin,
6 300072, PR China

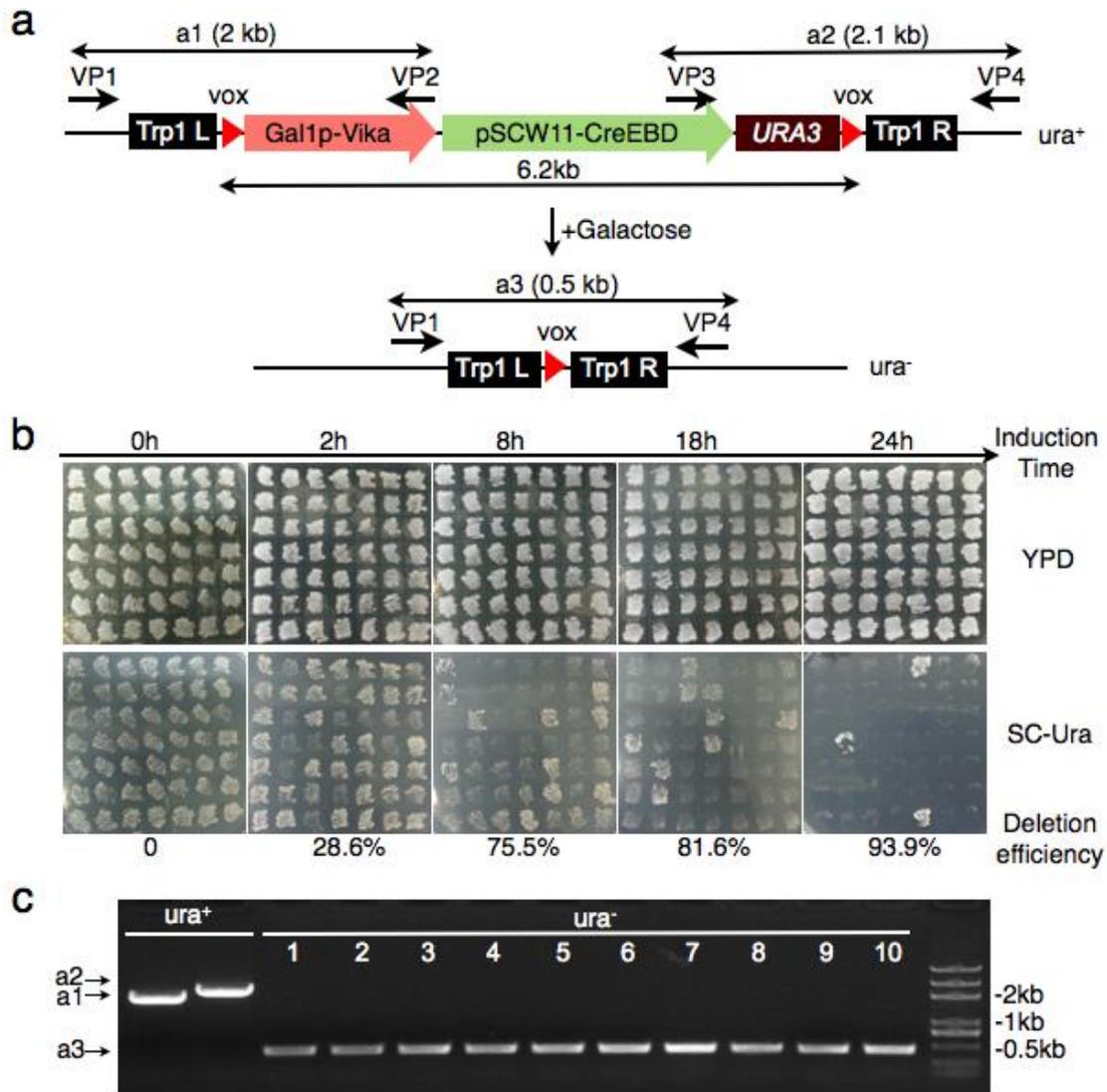
7 b: SynBio Research Platform, Collaborative Innovation Center of Chemical Science and
8 Engineering (Tianjin), School of Chemical Engineering and Technology, Tianjin University, Tianjin,
9 300072, PR China

10

11 * These authors contributed equally to this work.

12 † Correspondence to: yjyuan@tju.edu.cn

13



14

15 **Supplementary Figure 1| Vika mediated DNA cassette self-excision.** (A)
 16 Schematic of DNA cassette encoding recombinase vika and selective marker
 17 URA3 flanked by vox site for self-excision. (B) After inducing expression of
 18 recombinase Vika, cell was collected at various time point and grown on
 19 non-selective YPD and selective SC-Ura plate. (C) Excision was confirmed by
 20 amplified corresponding a3 DNA fragment from cell collected from 10 cell
 21 colonies at time point of 24hr. The most left two lanes are DNA fragement a1 and
 22 a2 amplified from the cell before Vika-mediated excision.

23

24 **Table S1 : Strains used in this study**

Name	Genotype/Description	Source
Strain		
BY4741	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0)	From Prof. Jef Boeke ¹
BY4742	MATα (his3 Δ 1, leu Δ 0, lys2 Δ 0, ura3 Δ 0)	From Prof. Jef Boeke ¹
<i>synIII</i>	MATα (MET15 lys1Δ0 ura3Δ0 his3Δ1leu2Δ0 synIII sup61::HO)	From Porf. Jef Boeke ¹
yLQH201	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0) [pRS415-GAL1p-Cre-loxP-His-loxP]	This study
yLQH202	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0) [pRS415-GAL1p-Cre-vox-Ura3-vox]	This study
yLQH203	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0) [pRS415-GAL1p-Vika-loxP-His3-loxP]	This study
yLQH204	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0) [pRS415-GAL1p-Vika-vox-Ura3-vox]	This study
yLQH205	MATα (MET15 lys1Δ0 ura3Δ0 his3Δ1leu2Δ0 synIII sup61::HO trp1::pSCW11-Cre-pGAL1-Vika-Ura3)	This study
yLQH206	MATα (MET15 lys1Δ0 ura3Δ0 his3Δ1leu2Δ0 synIII sup61::HO trp1::vox-pSCW11-Cre-Ura3-vox)	This study
yLQH207	MATα (MET15 lys1Δ0 ura3Δ0 his3Δ1leu2Δ0 trp1::vox-pSCW11-Cre-pGAL1-Vika-ura3-vox	This study
yLQH208	MATα (MET15 lys1Δ0 ura3Δ0 his3Δ1leu2Δ0 synIII sup61::HO trp1::vox)	This study
yLQH211	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0 HO::vox-RFP-vox-loxP-GFP-loxP-URA3)	This study
yLQH212	MATa (Leu2Δ0 LYS2 met15 his3Δ1 ura3Δ0 HO::vox-RFP-loxP-vox-GFP-loxP-URA3)	This study
Plasmids		
pLM158	pSCW11-CreEBD	This study
pUC57-Vika	Codon-optimized Vika ORF	Genewiz. Inc
pRS vector	pRS413, pRS414, pRS415, pRS416	From Prof.
pYES2	Vector with GAL1p and CYC1t	Gift from Prof. Jef Boeke
pLQH122	pYES2-GAL1p-Cre-CYC1t	This study
pLQH124	pYES2-GAL1p-Vika-CYC1t	This study
pLQH134	pRS415-pGAL1p-Cre-CYC1t	This study
pLQH135	pRS415-pGAL1p-Vika-CYC1t	This study

25

26

27 **Table S2: Primers used in this study**

Primers	Sequence(5'-3')
Expression plasmid construction	
Cre-HindIII-F	TATT <u>AAGCTT</u> TATGTCCAATTTACTGACCGT
Cre-XbaI-R	GCCCTCTAGATTAATCGCCATCTTCCAGCAGG
Vika-HindIII-F	CGCA <u>AAGCTT</u> TATGACTGATTTGACTCCTTTCC
Vika-XbaI-R	CGGATCTAGATTATCTTTGTCTCCTCTTCTGA
Gal1p-EcoRI-Trp1-F	AGACGTATTGTGACCCGGGA <u>ATTCC</u> GGATTAGAAGCCGCCGAGC
CYC1t-EcoRI-Trp1-R	ACAAGTAATTGGGCTGCAGGA <u>ATTTC</u> GCAAATTAAAGCCTTCGAGC
Orthogonal plasmids construction	
pRS415-XhoI-GAL1p-F	ATTGGGTACGGGGCCCCCCTCGAGCGGATTAGAAGCCGCCGAGC
CYC1t-vox-R	AATACGTCTGAGAATGGGCGTTCTCAGACCTATTGCAAATTTAAAGCCTTCGAGC
CYC1t-loxP-R	ATAACTTCGTATAATGTATGCTATACGAAGTTATGCAAATTTAAAGCCTTCGAGC
vox-Ura3-F	AATAGGTCTGAGAACGCCATTCTCAGACGTATTTTCAATTCATCATTTTTTTTT
Ura-vox-R1	AATACGTCTGAGAATGGGCGTTCTCAGACCTATTTTAGTTTTTGCTGGCCGCATC
Ura-vox-pRS415-XhoI-R2	GGAATTCGATATCAAGCTTATCGATACCGTCGACCTCGAGA <u>AATACGAA</u> TACGTCTGAGAATGGGCG
loxP-His3-F	ATAACTTCGTATAGCATACATTATACGAAGTTATCTAGTACACTCTATATTTTTT
His-loxP-R1	ATAACTTCGTATAATGTATGCTATACGAAGTTATCTACATAAGAACA CCTTTGG
His-loxP-pRS415-XhoI-R2	GGAATTCGATATCAAGCTTATCGATACCGTCGACCTCGAGA <u>AATACGAT</u> AACTTCGTATAATGTATG
Sequential and selective deletion experiment	
EcoRI-vox-RFP-F	CATGA <u>ATTCA</u> ATAGGTCTGAGAACGCCATTCTCAGACGTATTAGC TTTTCAATTCAATTCATCA
RFP-vox-KpnI-R	TATGGTACCAATACGTCTGAGAATGGGCGTTCTCAGACCTATTTCATAAAGAGCGACCTCAT
RFP-loxP-KpnI-R	ATTGGTACCATAA <u>ACTTC</u> GTATAATGTATGCTATACGAAGTTATTCAATAAGAGCGACCTCATG
KpnI-vox-GFP-F	TATGGTACCAATAGGTCTGAGAACGCCATTCTCAGACGTATTTGTTTTGCAAAAAGAACAAAAC

KpnI-loxP-GFP-F	TATGGTACCAATACGTCTGAGAATGGGCGTTCTCAGACCTATTTCATAAAGAGCGACCTCAT
GFP-loxP-BamHI-R	TTGGGATCCATAAACTTCGTATAATGTATGCTATACGAAGTTATAAAGCCACGCGTGTGCACC
SCRaMble termination experiment	
trp1L-vox-F	ATGTCTGTTATTAATTTTCAC
trp1L-vox-R	ggcggcttctaaccgAATACGTCTGAGAATGGGCGTTCTCAGACCTATTAGGAACTCTTGGTATTCTT
GAL1p-Vika-CYC1t-F	aataccaagagttcctAATAGGTCTGAGAACGCCCATTTCTCAGACGTATTCGGATTAGAAGCCGCCGAG
GAL1p-Vika-CYC1t-R	aagaatcacctctgggtagtagtatctcattgctcaagacacaagtaatGCAAATTAAAGCCTTCGAG
pSCW11-CreEBD-F	tgaaaacctgcttgagaaggtttgggacgctcgaaggcttaatttgcAATTACTTGTGTCTTGACG
pSCW11-CreEBD-R	caaagaaccgaaatcaaaaaaagaataaaaaaaatgatgaattgaaTCAAGCAAGGTTTTTCAGTA
URA3-vox-F	ctgtacagacgctgtacgcatgtaacattatactgaaaacctgcttgaTTCAATTCATCATTTTTTT
URA3-vox-R	aataactggcaaacggAATACGTCTGAGAATGGGCGTTCTCAGACCTATTTTAGTTTTGCTGGCCGCAT
pSCW11-Cre-F	ATTCTCAGACGTATTGTGACCCGGGAATTCctgcagcccAATTACTTGTGTCTTGACG
pSCW11-Cre-R	AAAGAATAAAAAAAAAAATGATGAATTGAATTGAAATCGATTCAAGCAAGGTTTTTCAGTA
Ura3-Cre-F	TGTAACATTATACTGAAAACCTTGCTTGAATCGATTTCAATTCAATTCATCATTTTTTT
Ura-vox-R	AATAACTGGCAAACCGAATACGTCTGAGAATGGGCGTTCTCAGACCTATTttagtttctgctggccgcat
vox-Trp1R-F	cggccagcaaaactaaAATAGGTCTGAGAACGCCCATTTCTCAGACGTATTGGTTTTGCCAGTTATTTAAA
vox-Trp1R-R	CTATTTCTTAGCATTTTTGACGAAA
VP1	GCACGTGAGTATACGTGATT
VP2	GCAAATTAAAGCCTTCGAGC
VP3	AAACGCTCTAAGGAGAACA
VP4	CTTGCTTTTCAAAGGCCTG

28

29

30

31

32

33 **References:**

34

35

36 1. Annaluru, N. et al. Total synthesis of a functional designer eukaryotic
37 chromosome. *Science* **344**, 55-58 (2014).

38

39