

Supplementary Information for

Interfering with retrotransposition by two types of CRISPR effectors:

Cas12a and Cas13a

Niubing Zhang^{1,3,#}, Xinyun Jing^{1,#}, Yuanhua Liu², Minjie Chen^{1,3}, Xianfeng Zhu³,

Jing Jiang³, Hongbing Wang⁴, Xuan Li^{1,*}, and Pei Hao^{2,*}

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SUPPLEMENTARY FIGURES

Figure S1

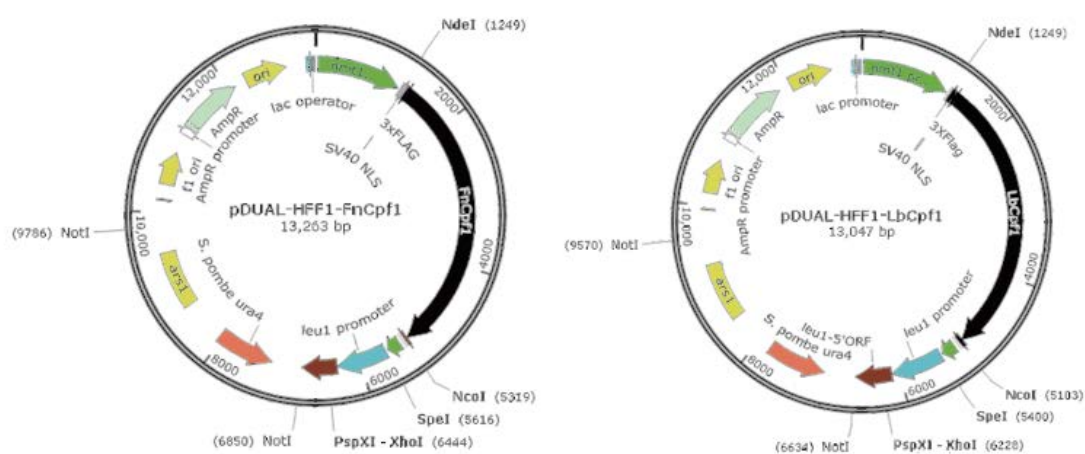


Figure S1. Maps of plasmids of pDUAL-HFF1-FnCpf1 and pDUAL-HFF1-LbCpf1 for chromosome-integrated expression of FnCpf1 and LbCpf1 in *S. pombe*. The FnCpf1 and LbCpf1 expression cassettes contain the *nmt1* promoter and the *ADH1* terminator. The SV40-derived nuclear localization signal (NLS) was fused to both N and C termini of Cpf1; a 3 × Flag was in-frame fused to the N terminus of NLS.

Figure S2

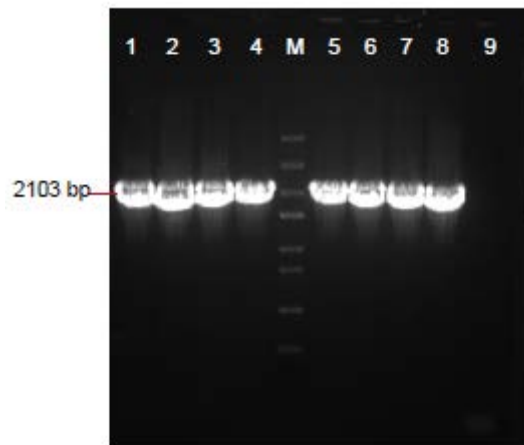


Figure S2. Diagnostic PCR for *S. pombe* transformants with FnCpf1 and LbCpf1 expression cassettes integrated into the chromosome. Primer set ADHterm-F/leu1-R (Supplementary Data 1) was used for diagnostic PCR. Lanes 1-4: *S. pombe* transformants with integrated expression construct of FnCpf1; Lanes 5-8: *S. pombe* transformants with integrated expression construct of LbCpf1; Lane 9: control wild-type *S. pombe* strain (without integrated expression constructs); M: *Trans5K* DNA Marker (Trans Transgen Biotech, Shanghai, China).

Figure S3

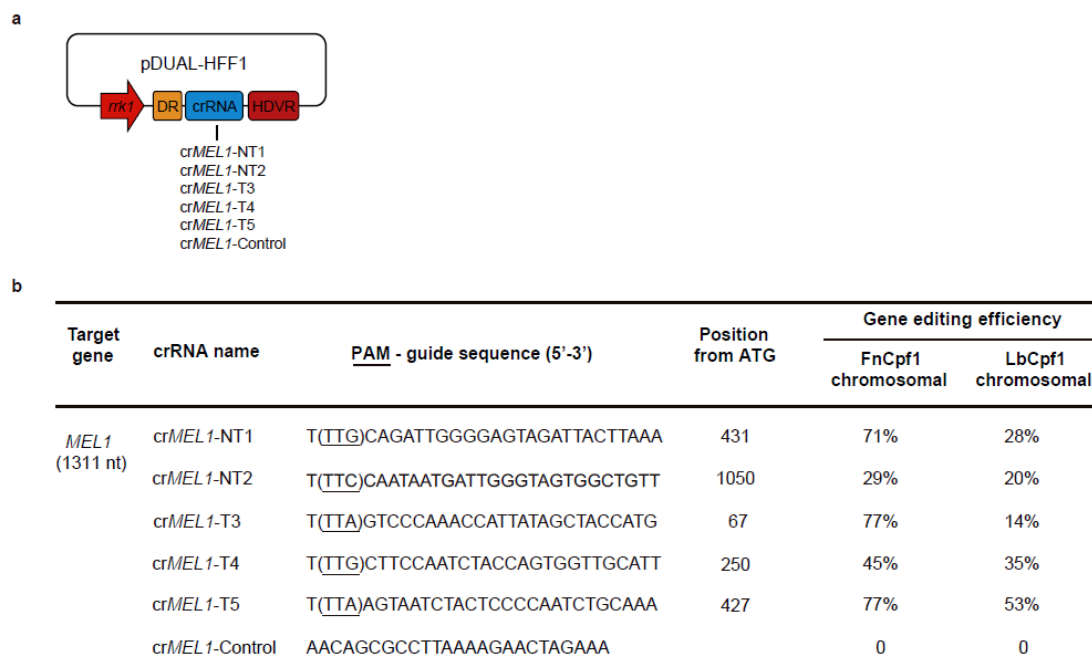
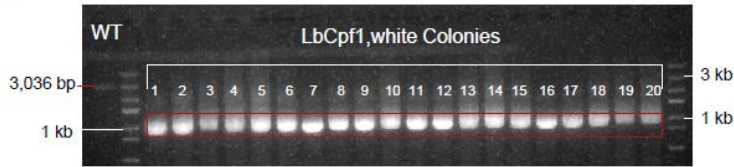


Figure S3. Editing on *MEL1* gene by chromosomal expressing FnCpf1 and LbCpf1 with different crRNAs. (a) Schematic representation the crRNA expression cassette for FnCpf1 and LbCpf1. *rrk1*, *rrk1* promoter; DR, direct repeat of crRNA; HDVR, hepatitis delta virus ribozyme. (b) Targeting sequences and editing efficiency on *MEL1* gene by chromosomal expressing FnCpf1 or LbCpf1 with five different crRNAs in *S.pombe*. The PAM motif (TTN for FnCpf1 and LbCpf1) regions are underlined.

Figure S4

a



b

crMEL1-T5_Col1:

TTCTTATACGGAATACATACAATTATATGCACTGAAAAGTTTTGCCACATTCGCCTCCGTAGTACTTCAAATGCCAATAGTAA
 TTACATTTTACAAGTTGAAAAACAAAAATGGCTGTAGTACTTGTACTCATTAAAGCTTCAAGTTACAATTTTTAGAAATC
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 CGAAAAATTTTTTAAAAAAGTTTATTATTTTTTTCTGTACATCAAATATTCAACGCTAAGAAAACCATAGAAAAATGCG
 CATCAAATTTTAATTAAGATATAGCAGATGCTATAACAAAAGGATTTGTCTAAAACCTAGAGATATTATTAAGCTTTTCTAA
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crMEL1-T5_Col2:

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 CATCAAATTTTAATTAAGATATAGCAGATGCTATAACAAAAGGATTTGTCTAAAACCTAGAGATATTATTAAGCTTTTCTAA
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c

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ACTAACAAATTTTATAAATTCGGGGCTCGTTCTACTATAGC
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Figure S4. Diagnostic PCR products and sequencing of *mel1* mutants generated by LbCpf1 editing. (a) Diagnostic PCR products of white colonies of *mel1* mutants generated by LbCpf1. All colonies show a 998-bp band indicating deletion of *MEL1* from LbCpf1 editing. The wild type strain shows a 3036-bp band. The primer set used for PCR were described in Figure 1c. (b) Sequences of PCR products from two randomly selected *mel1* mutant colonies. The sequence results show the correct deletion of the center portion of *MEL1* sequence. The blue and yellow highlighted sequences represent the upstream and downstream regions of *MEL1* gene, whereas the central portion is deleted. (c) Sequence of the template which was used for editing of *MEL1* gene by FnCpf1 or LbCpf1 through homologous recombination (HR). Blue sequence, the left arm of the HR template, which was amplified using primers of MEL1-P5, mel1-left-p3-new from *S.pombe* genomic DNA. Yellow sequence, the right arm of the HR template, which was amplified using primers MEL1-P3 and mel1-right-P5-new from *S.pombe* genomic DNA. The HR template was obtained by over-lap PCR using primers MEL1-P5, MEL1-P3 from left and right arms.

Figure S5

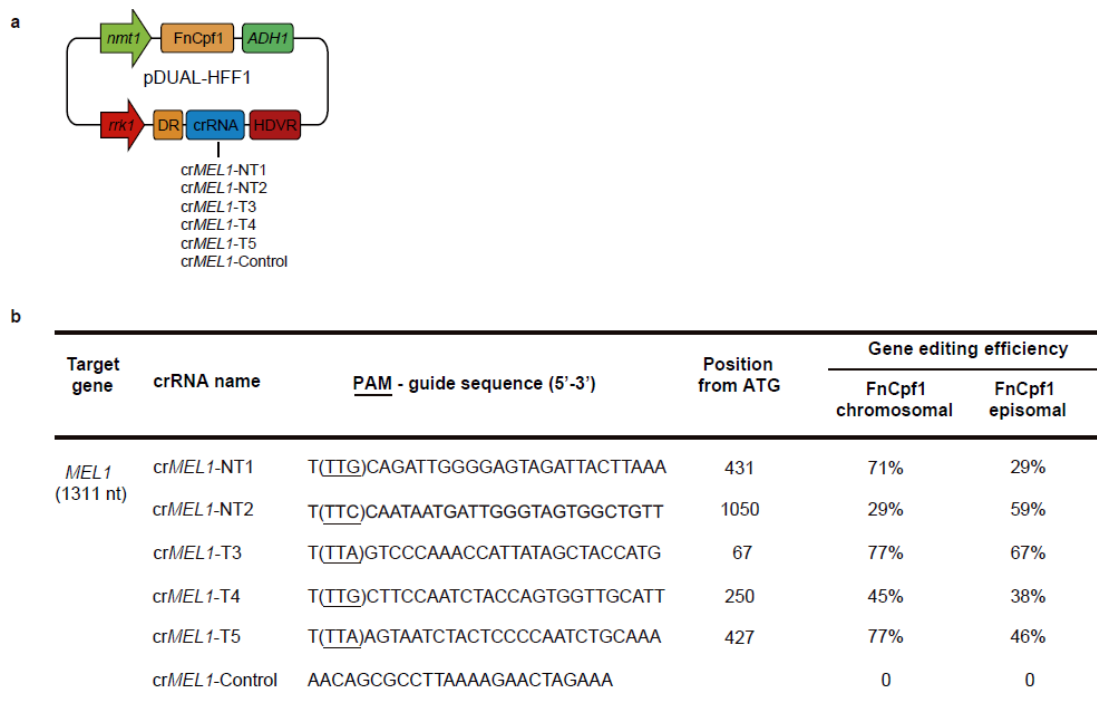


Figure S5. Editing efficiency on *MEL1* gene by chromosomal-expressing FnCpf1 and episomal-expressing FnCpf1 with five different crRNAs. (a) Schematic representation of the FnCpf1 and crRNA expression cassette in multi-copy plasmid pDUAL-HFF1. *rrk1*, *rrk1* promoter; DR, direct repeat of crRNA; HDVR, hepatitis delta virus ribozyme. (b) Targeting sequences and editing efficiency on *MEL1* gene by chromosomal-expressing FnCpf1 and episomal-expressing FnCpf1 with five different crRNAs in *S.pombe*. The PAM motif (TTTN for FnCpf1) regions are underlined.

Figure S6

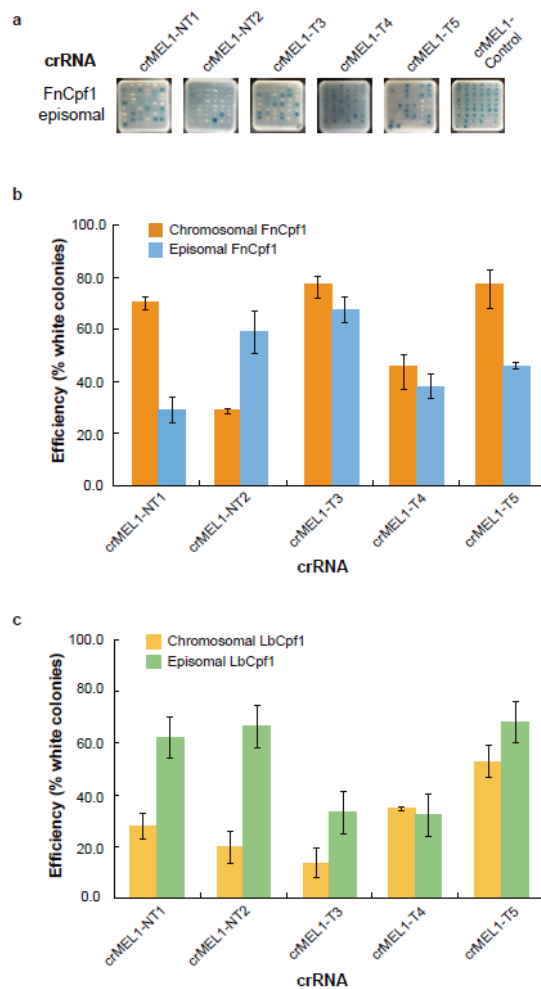


Figure S6. Editing of *MEL1* gene by chromosomal-expressing FnCpf1 and episomal-expressing FnCpf1 with five different crRNAs. (a) The editing on *MEL1* gene by episomal-expressing FnCpf1 with five different crRNAs, estimated by formation of white- (*mell* mutant) and blue-colored (*MEL1* wildtype) colonies, assayed on agar plates containing the X- α -Gal. (b) Efficiency of genome editing on *MEL1* gene by chromosomal expressing FnCpf1 and episomal expressing FnCpf1 with five different crRNAs. Bar represents means \pm s.e.m. with n = 3. (c) Efficiency of genome editing on *MEL1* gene by chromosomal expressing LbCpf1 and episomal expressing LbCpf1 with five different crRNAs. Bar represents means \pm s.e.m. with n = 3.

Figure S7

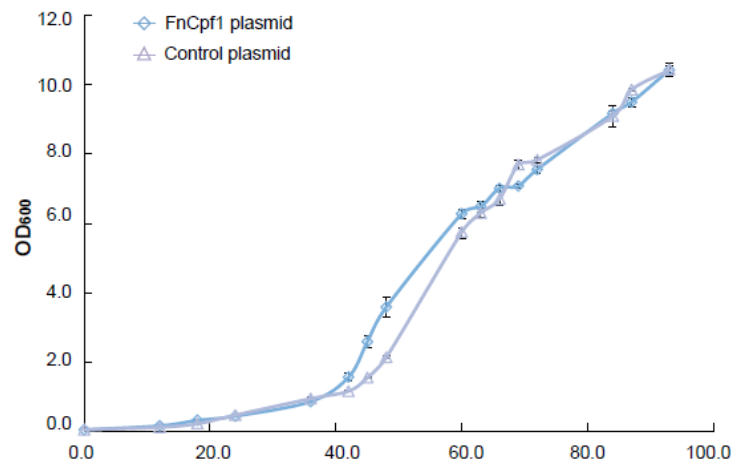


Figure S7. Growth in liquid culture media (EMM + uracil) of *S. pombe* strains with episomal expressing FnCpf1 compared to the control strain with control plasmid pDUAL-HFF1. They had an average double time of 9.2 h and 9.6 h, respectively. Data are means \pm s.e.m. (n = 4 independent experiments)

Figure S8

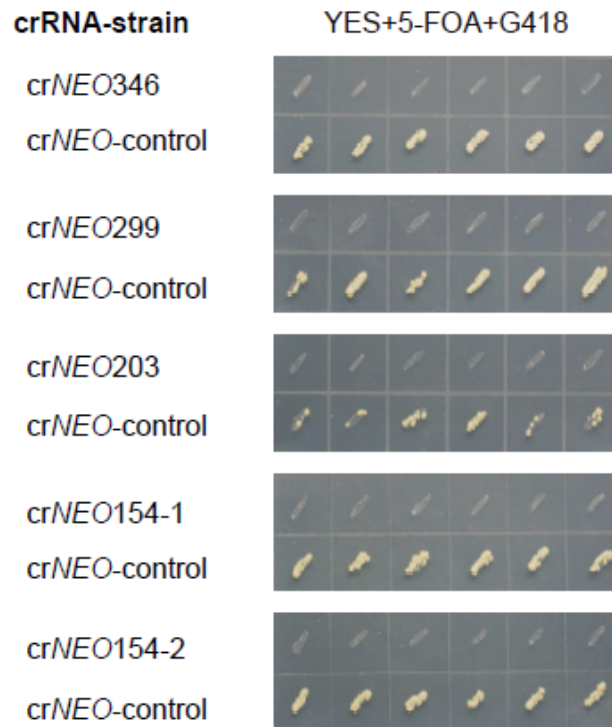


Figure S8. Prolonged crRNA targeting eliminating residual Tf1 retrotransposition by Cas12a.

‘Transfer-and-patch’ assay for detection of Tf1 retrotransposition using plates containing YES media with 5-FOA and G418. The effects of persistent crRNA targeting in their genome-integrated strains were compared to crNEO-control strains side by side.

Figure S9

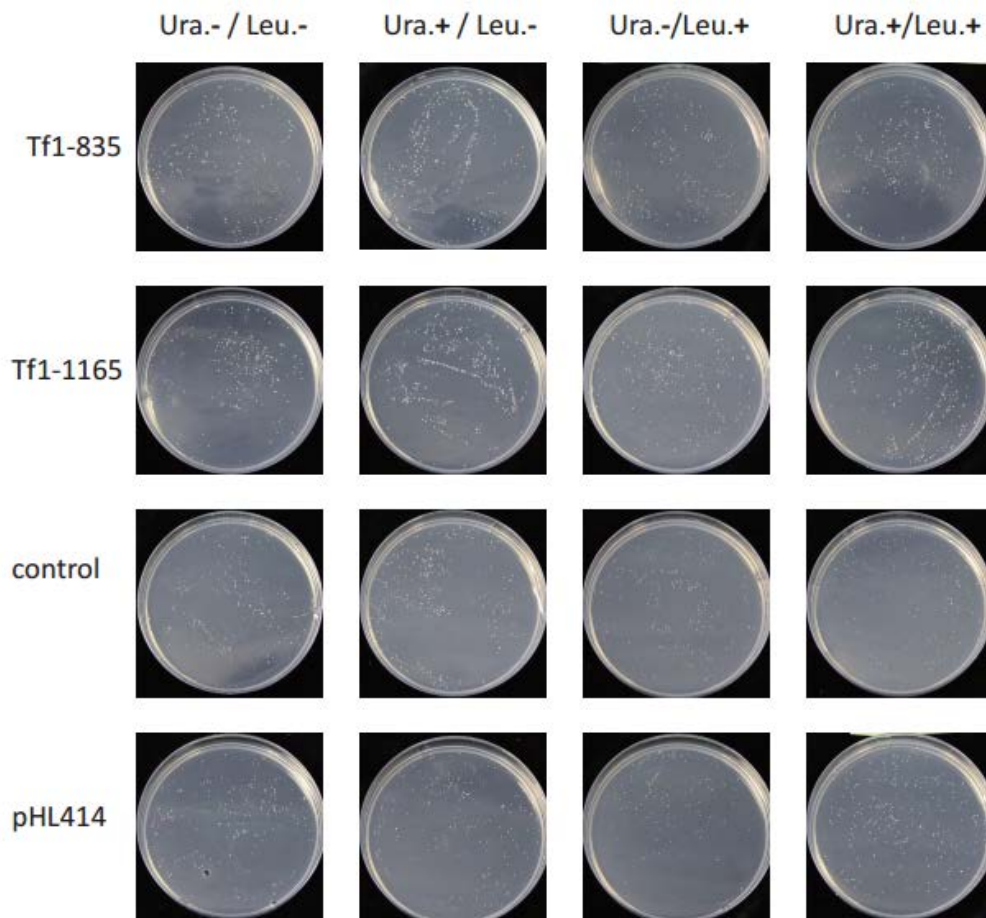


Figure S9. The ending *S. pombe* populations from one batch of liquid cultures (Fig. 5c) were further checked for presence of selection markers for Cas13a gene (Leu+) and Tf1/crRNA-carrying plasmid (Ura+)

100 μ L of 1000-fold diluted cells at late log-phase cultures (Fig. 5c) were plated to four different EMM selective plates (uracil-/leucine-, uracil+/leucine-, uracil-/leucine+, and uracil+/leucine+) respectively. The presence of Cas13a gene and Tf1/crRNA-carrying plasmid were validated by colony-formation on selective media plates.

SUPPLEMENTARY TABLES

Supplementary Table S1. List of strains and plasmids used in this study.

Strains and plasmids	Characteristics	Source/Reference
Strains		
<i>E.coli</i> DH5 α	<i>F- eNDA1 glnV44 thi-1 recA1 relA1 gyrA96 deoRnupG Φ80dlacZΔM15 Δ(lacZYA-argF)U169, hsdR17 (rK-mK), λ-</i>	Takara Biotechnology Co.,Ltd.
<i>S.pombe</i> FY7652	<i>h- leu1-32 ura4-D18</i>	National Bio Resource Project
Plasmids		
FnCpf1-OsU6	<i>E.coli</i> , <i>pBR322 ori</i> , Kan ^r , FnCpf1, crRNA cassette for expression in rice	Gift from Dr. Jiankang Zhu's Lab ¹
LbCpf1-OsU6	<i>E.coli</i> , <i>pBR322 ori</i> , Kan ^r , LbCpf1, crRNA cassette for expression in rice	Gift from Dr. Jiankang Zhu's Lab ¹
pDUAL-HFF1	<i>E.coli</i> - <i>S.pombe</i> shutting vector, <i>ars1 ori</i> , <i>Cole1 ori</i> , <i>fl ori</i> Amp ^r , <i>ura4</i>	RIKEN BRC (RDB:6179)
pDUAL-HFF1-FnCpf1	FY7652, FnCpf1/ <i>nmt1</i>	This study
pDUAL-HFF1-LbCpf1	FY7652, LbCpf1/ <i>nmt1</i>	This study
pDUAL-HFF1-Cas13a	FY7652, LshCas13a/ <i>nmt1</i>	Our Lab ²
pHL414	<i>S. pombe</i> , Tf1 retrotransposon/ <i>nmt1</i>	Gift from Professor Henry L. Levin's lab ³
pBluescript II KS(+)	<i>E.coli</i> , <i>Cole1 ori</i> , <i>fl ori</i> , Amp ^r	
pKS-rrk1-FnCpf1-crRNA-backbone	<i>E.coli</i> , crRNA backbone plasmid for FnCpf1, Amp ^r	This study
pKS-rrk1-LbCpf1-crRNA-backbone	<i>E.coli</i> , crRNA backbone plasmid for LbCpf1, Amp ^r	This study
pKS-rrk1-LshCas13a-crRNA-backbone	<i>E.coli</i> , crRNA backbone plasmid for LshCas13a, Amp ^r	This study

Supplementary Table S2. Intermediate plasmids for crRNA constructs.

Intermediate plasmids	Primers	Cloning sites	Original plasmids	Source
pKS-rrk1-(FnCpf1 crMEL1)-NT1	crMEL1-NT1-P5 crMEL1-NT1-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crMEL1)-NT2	crMEL1-NT2-P5 crMEL1-NT2-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crMEL1)-T3	crMEL1-T3-P5 crMEL1-T3-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crMEL1)-T4	crMEL1-T4-P5 crMEL1-T4-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crMEL1)-T5	crMEL1-T5-P5 crMEL1-T5-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crMEL1)-control	crMEL1-Control-P5 crMEL1-Control-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-NT1	crMEL1-NT1-P5 crMEL1-NT1-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-NT2	crMEL1-NT2-P5 crMEL1-NT2-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-T3	crMEL1-T3-P5 crMEL1-T3-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-T4	crMEL1-T4-P5 crMEL1-T4-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-T5	crMEL1-T5-P5 crMEL1-T5-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(LbCpf1 crMEL1)-control	crMEL1-Control-P5 crMEL1-Control-P3	<i>BspQI</i>	pKS-rrk1-LbCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO346)	crNEO346-P5 crNEO346-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO299)	crNEO299-P5 crNEO299-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO203)	crNEO203-P5 crNEO203-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO154-1)	crNEO154-1-P5 crNEO154-1-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO154-2)	crNEO154-2-P5 crNEO154-2-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(FnCpf1 crNEO-Control)	crNEO-Control-P5 crNEO-Control-P3	<i>BspQI</i>	pKS-rrk1-FnCpf1-crRNA-backbone	This study
pKS-rrk1-(LshCas13a crRNA)-(Tf1-835)	Tf1-835-gRNA-P5 Tf1-835-gRNA-P3	<i>BspQI</i>	pKS-rrk1-LshCas13a-crRNA-backbone	This study
pKS-rrk1-(LshCas13a crRNA)-(Tf1-1165)	Tf1-1165-gRNA-P5 Tf1-1165-gRNA-P3	<i>BspQI</i>	pKS-rrk1-LshCas13a-crRNA-backbone	This study
pKS-rrk1-(LshCas13a crRNA)-Control	Tf1-Control-gRNA-P5 Tf1-Control-gRNA-P3	<i>BspQI</i>	pKS-rrk1-LshCas13a-crRNA-backbone	This study

Supplementary Table S3. crRNA Constructs for targeting the endogenous *MEL1* gene by FnCpf1 and LbCpf1.

Plasmid Name	Genes/Promoters	crRNA primers	Original plasmid	Descriptions	Source
pDUAL-HFF1-(FnCpf1 crMEL1)-NT1	crRNA/ <i>rrk1</i>	crMEL1-NT1-P5 crMEL1-NT1-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(FnCpf1 crMEL1)-NT2	crRNA/ <i>rrk1</i>	crMEL1-NT2-P5 crMEL1-NT2-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(FnCpf1 crMEL1)-T3	crRNA/ <i>rrk1</i>	crMEL1-T3-P5 crMEL1-T3-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(FnCpf1 crMEL1)-T4	crRNA/ <i>rrk1</i>	crMEL1-T4-P5 crMEL1-T4-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(FnCpf1 crMEL1)-T5	crRNA/ <i>rrk1</i>	crMEL1-T5-P5 crMEL1-T5-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(FnCpf1 crMEL1)-control	crRNA/ <i>rrk1</i>	crMEL1-Control-P5 crMEL1-Control-P3	pDUAL-HFF1	for chromosomal expression of FnCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-NT1	crRNA/ <i>rrk1</i>	crMEL1-NT1-P5 crMEL1-NT1-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-NT2	crRNA/ <i>rrk1</i>	crMEL1-NT2-P5 crMEL1-NT2-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-T3	crRNA/ <i>rrk1</i>	crMEL1-T3-P5 crMEL1-T3-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-T4	crRNA/ <i>rrk1</i>	crMEL1-T4-P5 crMEL1-T4-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-T5	crRNA/ <i>rrk1</i>	crMEL1-T5-P5 crMEL1-T5-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-(LbCpf1 crMEL1)-control	crRNA/ <i>rrk1</i>	crMEL1-Control-P5 crMEL1-Control-P3	pDUAL-HFF1	for chromosomal expression of LbCpf1 and episomal expression crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-NT1	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-NT1-P5 crMEL1-NT1-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-NT2	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-NT2-P5 crMEL1-NT2-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-T3	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-T3-P5 crMEL1-T3-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-T4	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-T4-P5 crMEL1-T4-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-T5	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-T5-P5 crMEL1-T5-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study
pDUAL-HFF1-FnCpf1-crMEL1-control	<i>FnCpf1</i> / <i>nmt1</i> crRNA/ <i>rrk1</i>	crMEL1-Control-P5 crMEL1-Control-P3	pDUAL-FnCpf1-HFF1	for episomal expression of FnCpf1 and crRNA	This study

Supplementary Table S4. crRNA constructs for targeting the Tf1 DNA intermediates (generated via splicing of artificial introns and reverse-transcription) by FnCpf1.

Plasmid Name	Promoter	Original plasmid	Descriptions	Source
pHL414-Tf1-neo-intron346	Tf1/ <i>nmt1</i>	pHL414	Tf1 retrotransposition reporters plasmid, artificial intron is 346bp from ATG of the NEO gene	This study
pHL414-Tf1-neo-intron299	Tf1/ <i>nmt1</i>	pHL414	Tf1 retrotransposition reporters plasmid, artificial intron is 299bp from ATG of the NEO gene	This study
pHL414-Tf1-neo-intron203	Tf1/ <i>nmt1</i>	pHL414	Tf1 retrotransposition reporters plasmid, artificial intron is 203bp from ATG of the NEO gene	This study
pHL414-Tf1-neo-intron154	Tf1/ <i>nmt1</i>	pHL414	Tf1 retrotransposition reporters plasmid, artificial intron is 154bp from ATG of the NEO gene	This study
pHL414-Tf1-neo-intron346-(FnCpf1 crNEO346)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron346	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron299-(FnCpf1 crNEO299)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron299	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron203-(FnCpf1 crNEO203)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron203	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron154-(FnCpf1 crNEO154-1)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron154	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron154-(FnCpf1 crNEO154-2)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron154	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron346-(FnCpf1 crNEO-Control)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron346	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron299-(FnCpf1 crNEO-Control)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron299	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron203-(FnCpf1 crNEO-Control)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron203	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pHL414-Tf1-neo-intron154-(FnCpf1 crNEO-Control)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414-Tf1-neo-intron154	for chromosomal expression of FnCpf1, episomal expression crRNA and Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO346	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO299	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO203	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO154-1	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO154-2	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study
pDUAL-HFF1-FnCpf1-crNEO-control	FnCpf1 / <i>nmt1</i> crRNA/ <i>rrk1</i>	pDUAL-FnCpf1-HFF1	for chromosomal expression of FnCpf1 and crRNA; episomal expression Tf1 retrotransposition reporter	This study

Supplementary Table S5. crRNA constructs for targeting the Tf1 RNA intermediates by LshCas13a.

Plasmid Name	Promoter	Original plasmid	target sequence (5' to 3')	Descriptions	Source
pHL414-Tf1-(LshCas13a crRNA)-(Tf1-835)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414	ATCCAAC TAGGTTTAC CATCTTCTTAA	for chromosomal expression of LshCas13a and episomal expression crRNA and Tf1	This study
pHL414-Tf1-(LshCas13a crRNA)-(Tf1-1165)	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414	TGTCGTTCTCCTTTAAA AACTTATTGTT	for chromosomal expression of LshCas13a and episomal expression crRNA and Tf1	This study
pHL414-Tf1-(LshCas13a crRNA)-Control	Tf1/ <i>nmt1</i> crRNA/ <i>rrk1</i>	pHL414	AATGCCTGGCTTGTCG ACGCATAGTCTG	for chromosomal expression of LshCas13a and episomal expression crRNA and Tf1	This study

References

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- 2 Jing, X. *et al.* Implementation of the CRISPR-Cas13a system in fission yeast and its repurposing for precise RNA editing. *Nucleic Acids Res.* **46**, 90 (2018).
- 3 Sangesland, M., Atwood-Moore, A., Rai, S. K. & Levin, H. L. Qualitative and quantitative assays of transposition and homologous recombination of the retrotransposon Tf1 in *Schizosaccharomyces pombe*. *Methods Mol. Biol.* **1400**, 117-130 (2016).

Supplementary Data S1

Primers Name	Sequences (5'-3')	Purpose
Construction of vectors and <i>S. pombe</i> strains for FnCpf1 and LbCpf1 expression		
pHSN-Fn-NdeI-F	GCATCACCACCATCAT CATATG gactataaggaccacg	Amplification of FnCpf1 and LbCpf1
pHSN-Fn-NcoI-R	CATCGTCGTCCTTGTAGT CCATGG ttacttttctttttgctg	Amplification of FnCpf1 and LbCpf1
ADHterm-F	CTCTTATTGACCACACCTCTACC	Confirmation of correct integration FnCpf1 and LbCpf1
leu1-R	GGTCATAAAGTTGAACGGATGTCG	Confirmation of correct integration FnCpf1 and LbCpf1
Construction of Tf1 retrotransposition reporters with artificial introns		
neo-intron299	gaatgctgtttcccggggatcgGTAGGTGCTATTTTACTAGTCTAAGCTAA TCAATAGcagtggtagtaacctgcatcatcaggagtacggataaaatgcttgatggtcggaa gaggcataaattccgtcagccagtttagtctgacctctcatctgtaacatcattggcaacgctaccttg ccatgttcagaaacaactctggcgcacatcgggcttccataacaatgatagattgtcgcacct	To generate the NEO fragments with artificial intron
neo-intron203	gaatgctgtttcccggggatcgagtagtaacctgcatcatcaggagtacggataaaatgctt gatggtcggagaggcataaattccgtcagccagtttagtctgacctctGTAGGTGCTATT TACTAGTCTAAGCTAATCAATAGcatctgtaacatcattggcaacgctaccttg ccatgttcagaaacaactctggcgcacatcgggcttccataacaatgatagattgtcgcacct	To generate the NEO fragments with artificial intron
neo-intron154	gaatgctgtttcccggggatcgagtagtaacctgcatcatcaggagtacggataaaatgctt gatggtcggagaggcataaattccgtcagccagtttagtctgacctctcatctgtaacatcattggca acgctaccttgccatgttcagaaacaGTAGGTGCTATTTTACTAGTCTAAG CTAATCAATAGctctggcgcacatcgggcttccataacaatgatagattgtcgcacct	To generate the NEO fragments with artificial intron
neo-intron346-P5	tcaacaatatttcacctgaGTAGGTGCTATTTTACTAGTCTAAGCTAATC AATAGatcaggatattcttctaatac	To generate the NEO fragments with artificial intron

neo-intron346-P3	gtattagaagaatatcctgatCTATTGATTAGCTTAGACTAGTAAAATAGC ACCTACtcaggtgaaaatattgtga	To generate the NEO fragments with artificial intron
Neo-TYCZ-P5	catcaacaatattttcacctgaatcaggatattcttctaataacctggaatgctgtttcccggggat	To generate the NEO fragments with artificial intron
Neo-TYCZ-P3	atgggtataaatgggctcgcgataatgtcgggcaatcaggtgcgacaatctatcgattg	To generate the NEO fragments with artificial intron

Intermediate plasmids for crRNA constructs

crMEL1-NT1-P5	gat CAGATTGGGGAGTAGATTACTTAAA	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-NT1-P3	gcc TTAAAGTAATCTACTCCCAATCTG	
crMEL1-NT2-P5	gat CAATAATGATTGGGTAGTGGCTGTT	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-NT2-P3	gcc AACAGCCACTACCCAATCATTATTG	
crMEL1-T3-P5	gat GTCCCAAACCATTATAGCTACCATG	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-T3-P3	gcc CATGGTAGCTATAATGGTTTGGGAC	
crMEL1-T4-P5	gat CTTCCAATCTACCAGTGGTTGCATT	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-T4-P3	gcc AATGCAACCACTGGTAGATTGGAAG	
crMEL1-T5-P5	gat AGTAATCTACTCCCAATCTGCAAA	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-T5-P3	gcc TTTGCAGATTGGGGAGTAGATTACT	
crMEL1-Control-P5	gat AACAGCGCCTTAAAAGAACTAGAAA	the oligonucleotide pairs designed for spacer of each crRNA
crMEL1-Control-P3	gcc TTTCTAGTTCTTTTAAGGCGCTGTT	
crNEO346-P5	GAT acctgaatcaggatattcttcta	the oligonucleotide pairs designed for spacer of each crRNA
crNEO346-P3	GCC attagaagaatatcctgattcaggt	
crNEO299-P5	GAT ccggggatcgcagtggtgagtaacc	the oligonucleotide pairs designed for spacer of each crRNA
crNEO299-P3	GCC ggttactcaccactcgcgatccccgg	
crNEO203-P5	GAT gtctgacctctcatctgtaacatc	the oligonucleotide pairs designed for spacer of each crRNA
crNEO203-P3	GCC gatgttacagatgagatggtcagac	

crNEO154-1-P5	GAT ccatgtttcagaaacaactctggcg	the oligonucleotide pairs designed for spacer
crNEO154-1-P3	GCC cgccagagttgtttctgaaacatgg	of each crRNA
crNEO154-2-P5	GAT agaaacaactctggcgcatcgggct	the oligonucleotide pairs designed for spacer
crNEO154-2-P3	GCC agcccgatgcgccagagttgtttct	of each crRNA
crNEO-Control-P5	gat AACAGCGCCTTAAAAGAACTAGAAA	the oligonucleotide pairs designed for spacer
crNEO-Control-P3	gcc TTTCTAGTTCTTTTAAGGCGCTGTT	of each crRNA
Tf1-835-gRNA-P5	aac Atccaactagtttaccattcttcttaa	the oligonucleotide pairs designed for spacer of
Tf1-835-gRNA-P3	gcc ttaagaagaatggtaaacctagtggat	each crRNA
Tf1-1165-gRNA-P5	aac Tgtcgttctccttataaaacttattgtt	the oligonucleotide pairs designed for spacer
Tf1-1165-gRNA-P3	gcc aacaataagttttaaaggagaacgaca	of each crRNA
Tf1-Control-gRNA-P5	acc CAGACTATGCGTCGACAAGCCAGGCATT	the oligonucleotide pairs designed for spacer
Tf1-Control-gRNA-P3	gcc AATGCCTGGCTTGTCGACGCATAGTCTG	of each crRNA

Construction of crRNA vectors to direct targeting by FnCpf1, LbCpf1, and LshCas13a

pDUAL-PspXI-T7	CCTCCAATCTTGTGTTCTTCAA TAATACGACTCACTATAGG	Amplification of crRNA array cassettes inserted into linearized plasmids pDUAL-HFF1
pDUAL-SpeI-T3	CGCTAGGGATAACAGGGTAATAT AATTAACCCTCACTAAAGG	Amplification of crRNA array cassettes inserted into linearized plasmids pDUAL-HFF1
pHL414-NheI-T7	TGCAGCCCGGGGATCCCAGCTGTAATACGACTCACTATAG	Amplification of crRNA array cassettes inserted into linearized plasmids of Tf1 retrotransposition reporters
pHL414-NheI-T3	ATCGCCAGTCACTATGGCGTGCTtctagaAATTAACCCTCACTAA AG	Amplification of crRNA array cassettes inserted into linearized plasmids of Tf1 retrotransposition reporters

pDUAL-SpeI-SK primer	ATCGACGATAAAAGAATCATCTTtctagaactagtgatc	Amplification of crRNA array cassettes inserted into linearized plasmid pDUAL-HFF1-FnCpf1
pDUAL-SpeI-T3	CGCTAGGGATAACAGGGTAATATAATTAACCCTCACTAAAGG	Amplification of crRNA array cassettes inserted into linearized plasmid pDUAL-HFF1-FnCpf1
NheI-TYB-P5	GGGGGATCCCAGCTGGCTAGCAATTAACCCTCACTAAAGG	Amplification of crRNA array cassettes inserted into linearized plasmid pHL414
NheI-TYB-P3	TCACTATGGCGTGCTGCTAGCTAATACGACTCACTATAGG	Amplification of crRNA array cassettes inserted into linearized plasmid pHL414
MEL1-repair fragment		
mel1-left-P5-new	AGGTAAAACCTCCAGAACATTAC	Amplification of repair fragment for <i>MEL1</i> deletion
mel1-left-P3-new	TCATGTGCTAGGTCGATTCTGG GAACAAGAAGTCAATATCGCTTAC	Amplification of repair fragment for <i>MEL1</i> deletion
mel1-right-P5-new	GTAAGCGATATTGACTTCTTGTTCC CCAGAATCGACCTAGCACATGA	Amplification of repair fragment for <i>MEL1</i> deletion
mel1-right-P3-new	GCTATAGTAGAACGAGCCCCGA	Amplification of repair fragment for <i>MEL1</i> deletion
MEL1-P5	AGGTAAAACCTCCAGAACATTAC	Confirmation <i>MEL1</i> deletion
MEL1-P3	GCTATAGTAGAACGAGCCCCGA	Confirmation <i>MEL1</i> deletion

Supplementary Data S2

Interference with Tf1 retrotransposition by CRISPR-Cas12a leaving marginal transposition activities

crRNA-plasmid	# of colonies	YE+Ura.+5-FOA --10 ⁵ cell / mL	YE+Ura.+5-FOA+G418 --10 ⁶ cell / mL	Transposition frequency(%)	Normalized - Transposition frequency(%)
<i>crNEO</i> 346	crNEO346-①	1500	12	0.08	0.1
	crNEO346-②	2500	14	0.06	
	crNEO346-③	2795	19	0.07	
	mean (crNEO346)			0.07	
<i>crNEO</i> control	crNEO-control-1-①	2000	74	0.37	1.0
	crNEO-control-1-②	2000	86	0.43	
	crNEO-control-1-③	2000	160	0.80	
	crNEO-control-2-①	2600	259	1.00	
	crNEO-control-2-②	2200	228	1.04	
	crNEO-control-2-③	2300	210	0.91	
	mean (crNEO-control)			0.76	

Prolonged crRNA targeting eliminating residual Tf1 retrotransposition by CRISPR-Cas12a

crRNA-strain	# of colonies	YE+Ura.+5-FOA --10 ⁵ cell / mL	YE+Ura.+5-FOA+G418 --10 ⁷ cell / mL	Transposition frequency(%)	Normalized - Transposition frequency(%)
<i>crNEO</i> 346	crNEO 346-①	2200	1	0.000	0.00
	crNEO 346-②	2000	1	0.001	
	crNEO 346-③	2500	1	0.000	
	mean (crNEO 346)			0.000	
<i>crNEO</i> control	crNEO control-①	2500	1220	0.488	1.00
	crNEO control-②	2500	1214	0.486	
	crNEO control-③	2500	1257	0.503	
	mean (crNEO control)			0.492	

Interfering with Tf1 retrotransposition by CRISPR-Cas13a via targeting its RNA intermediates

crRNA-plasmid	# of colonies	YE+Ura.+5-FOA+G418 --10⁶ cell / mL	YE+Ura.+5-FOA --10⁵ cell / mL	Transposition frequency(%)	Normalized - Transposition frequency(%)
Tf1-835	Tf1-835-①	820	2100	3.905	0.84
	Tf1-835-②	800	2000	4.000	
	Tf1-835-③	870	2000	4.350	
	mean (Tf1-835-1)			4.085	
Tf1-1165	Tf1-1165-①	340	1700	2.000	0.40
	Tf1-1165-②	400	2150	1.860	
	Tf1-1165-③	400	2020	1.980	
	mean(Tf1-1165-3)			1.947	
Tf1-control	Tf1-control-①	970	1900	5.105	1.00
	Tf1-control-②	1020	2000	5.100	
	Tf1-control-③	910	2100	4.333	
	mean (Tf1-control)			4.846	