Supplementary Material

# Supplementary Tables

**Supplementary Table 1**. List of primers used in this study

|  |  |  |
| --- | --- | --- |
| Description | Primer name | Primer sequence |
| *Fn*Cas12a1 system construction  | 2653-1-For | ACTGAGAATTCAATTAAAGGCTCCTTTTGGAGCCTTTTTTTCTGTCAGACCAAGTTTAC |
| 2653-1-Rev | GCAACAACATGAATGGTCATCGGTTTCCGTGTTTCGTAAAGTCTGGAAACGCGGAAGTC |
| 2653-2-For | CGAAACACGGAAACCGATGACCATTCATGTTGTTGCTCAGGTCGCAGACGTTTTGCAGC |
| 2653-2-Rev | GCTCGCGGACCTCATGGACGCCCTCCAGGGCACCCGGAAAACGCCGGACAGCCCCCGGC |
| 2653-3-For | GTCGCTGGTAGGCGCCCGGCCGCCGGGGGCTGTCCGGCGTTTTCCGGGTGCCCTGGAGG |
| 2653-3-Rev | TTATTTTTATAGCACGTGATGAAAAGGACCCAGGTGGCACTTTTCGTACCCGGGGATCC |
| 2653-yeast-For | GATTCTGGTCTAGCTAGAGTCGACTAGAGGATCCCCGGGTACGAAAAGTGCCACCTGGG |
| 2653-yeast-Rev | CGAAAAAAAAACCCCGCCCCTGACAGGGCGGGGTTTTTTTTTCTAGAGTGAGTTTAGTATACATGC |
| *lacZ*-For | TCTAGAAAAAAAAACCCCGCCCTGTCAGGGGCGGGGTTTTTTTTTCGGTCTTCACCGGT |
| LacZ-15-Rev | ACGGGGCCGGGCACCGTCGGGCCGGCGGCGCGGAACTAGTGCTGCTCCTTCGGTCGGAC |
| SA-15-For | TGCCCGTAGACGCACGTCCGACCGAAGGAGCAGCACTAGTTCCGCGCCGCCGGCCCGAC |
| SA-15-Rev | AGTACTTGTTCACAAATTCTTGGTAGATGGACATGCTAGCGGGTTCCTCCTCATGAGTC |
| Cas12a-15-For | TGCAGTGAACAAGTGGACTCATGAGGAGGAACCCGCTAGCATGTCCATCTACCAAGAAT |
| Cas12a-Rev | CTGACAGAAAAAAAGGCTCCAAAAGGAGCCTTTAATTGAATTCTCAGTTATTGCGGTTC |
| *ermE*p\*-For | GACGCACGTCCGACCGAAGGAGCAGCGGTACCAGCCCGACCCGAG |
| *ermE*p\*-Rev | CAAATTCTTGGTAGATGGACATGCTAGCGTGGTGTCCTACCAACCGG |
| *kasO*p\*-For | GACGCACGTCCGACCGAAGGAGCAGCTGTTCACATTCGAACGGTC |
| *kasO*p\*-Rev | CAAATTCTTGGTAGATGGACATGCTAGCAACTCCCCCAGTCCTGCAC |
| *kasO*p\*-1-For | TGTTCACATTCGAACGGTCTCTGCTTTGACAACATGCTGTGCGGTGTTGTAAAGTCGTG |
| *kasO*p\*-1-Rev | AACTCCCCCAGTCCTGCACGCTGTCGTATTCTCCTGGCCACGACTTTACAACACCGCAC |
| Potr\*-For | GACGCACGTCCGACCGAAGGAGCAGCTCAGGCGGACTGCCGCCCG |
| Potr\*-Rev | CAAATTCTTGGTAGATGGACATGCTAGCGGTTCCTCACTCTCCTGGC |
| rpsLp(XC)-For | GACGCACGTCCGACCGAAGGAGCAGCGCCCTGCAGGCGGAAGTCA |
| rpsLp(XC)-BbsI-Rev | CAAATTCTTGGTAGATGGACATGCTAGCTACGTCTCCGTCGTCTACT |
| *Fn*Cas12a2 system construction | 2-*lacZ* For | GTATCTGAAAGGGGATACGCAAGTCTTCCGGTGGAAAGCGGGCAGTG |
| 2-*lacZ* Rev | CGTTCTGAACAAATCCAGATGGAGTATGTCTTCTCAGCCGCTACAGG |
| 2-gapdh For | GTTCGTCCAGAACCGCAATAACTGAGAATTCAGATCTACGCGTTC |
| 2-gapdh Rev | CACTGCCCGCTTTCCACCGGAAGACTTGCGTATCCCCTTTCAGATAC |
| 2-Yes For | CCTGTAGCGGCTGAGAAGACATACTCCATCTGGATTTGTTCAG |
| 2-Yes Rev | CAAAGGATCTTCTTGAGATCC |
| 2-acc For | GCAAGCAGCAGATTACGCGC |
| 2-acc Rev | GAGGAGGAGATCACCGACGAC |
| c-pSG For | GATGACCGCGATGGCGAC |
| 2-pSG Rev | GACTTCCGCCTGCAGGGCAAGCTTCAGCTCGCGGACGTGCTCA |
| 2-rpsl For | TGAGCACGTCCGCGAGCTGAAGCTTGCCCTGCAGGCGGAAGTC |
| 2-rpsl Rev | ATTCTTGGTAGATGGACATGGATCCTACGTCTCCGTCGTCTACTC |
| 2-Cas12a For | GAGTAGACGACGGAGACGTAGGATCCATGTCCATCTACCAAGAAT |
| 2-Cas12a Rev | GAACGCGTAGATCTGAATTCTCAGTTATTGCGGTTCTG |
| 2-*kasO*p\*-pCFc-For | TGCAGGACTGGGGGAGTTGGATCCATGTCCATCTACCAAG |
| 2-*kasO*p\*-pCFc-Rev | TGCAGGACTGGGGGAGTTGGATCCATGTCCATCTACCAAG |
| 2-*ermE*p\*-pCFc-For | CGTCGTGGACTATGAGCACGTCCGCGAGCTGAAGCTTGGTACCAGCCCGACCCGA |
| 2-*ermE*p\*-pCFc-Rev | CTTGGTAGATGGACATGGATCCGTGGTGTCCTACCAACCGGCAC |
| 2-Potr\*-pCFc-For | GCGTCGTGGACTATGAGCACGTCCGCGAGCTGAAGCTTTCAGGCGGACTGCCGCCCGGA |
| 2-Potr\*-pCFc-Rev | CTTGGTAGATGGACATGGATCCGGTTCCTCACTCTCCTG |
| *ActII-orf4* deletion using *Fn*Cas12a1 system with the 23-nt spacer followed by a 19-nt or a 36-nt direct repeat  | 1-*actII*4-19-sp-For | TTTTCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATT |
| 1-*actII*4-19-sp-Rev | ACGCAATTTCTACTGTTGTAGATGAATGCAGCTCCGTATCGACGCG |
| 1-*actII*4-36-sp-For | TTTTCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| 1-*actII*4-36-sp-Rev | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGAATGCAGCTCCGTATCGACGCG |
| 1- *actII*4-KL-For  | CTGTATTATAAGTAAATGCATGTATACTAAACTCACTC  |
| 1- *actII*4-KL-Rev | CTACTGACGCCGGCTGCGCCCCCGTCGAGATTC |
| 1- *actII*4-KR-For | GACGGGGGCGCAGCCGGCGTCAGTAGTTCCCCAG |
| 1- *actII*4-KR-Rev  | ACCCCGCCCCTGACAGGGCGGGGT  |
| 1-Cas12a-SP-check-For | GGCTGGGAAGCATATTTG  |
| 1-Cas12a-SP-check-Rev | CTGCTCCTTCGGTCGGAC  |
| P1 | GTGGGCGAGGAGATCCTGAC |
| P2 | ATGGTGGCGTACCTGATC |
| *ActII-orf4* deletion using *Fn*Cas12a2 system with the 23-nt spacer followed by a 19-nt or a 36-nt direct repeat  | 2- *actII*4-19-sp-For | ACGCAATTTCTACTGTTGTAGATGAATGCAGCTCCGTATCGACGCG |
| 2- *actII*4-19-sp-Rev | GAGTCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATT |
| 2- *actII*4-36-sp-For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGAATGCAGCTCCGTATCGACGCG |
| 2- *actII*4-36-sp-Rev | GAGTCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| 2- *actII*4-KL-For | TGCCGCCGGGCGTTTTTTATCTAGAGGTGAACCCATGGTCGTC |
| 2- *actII*4-KR-Rev | GACCCAGGTGGCACTTTTCGTCTAGAACGACTCTGCGCTTCAATC |
| 2-spacer-2kb check For  | GTGAATGGCCTGTTC |
| 2-spacer-2kb check Rev  | GATCCCCCTAGAGTC |
| ACT deletion using pYL-*kasO*p\*-*Fn*Cas12a2 plasmid | ACT-sp-For 1 | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGAATGCAGCTCCGTATCGACGCG |
| ACT-sp-Rev 1 | GAGTCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| ACT-KL For  | CAGAACGCTCGGTTGCCGCCGGGCGTTTTTTATCTAGACTCTCGAACACGGCGGTCAC |
| ACT-KL Rev  | CACCCACATGGTGGAGGAGACCGGTGAACGGACGCTGACAG |
| ACT-KR For  | CTGTCAGCGTCCGTTCACCGGTCTCCTCCACCATGTGGGTG |
| ACT-KR Rev  | TGATGAAAAGGACCCAGGTGGCACTTTTCGTCTAGAGAACGTCCGCCTGGTCGAGAC |
| P3 | CTGTGCTGCTTTTCGCGCCTG |
| P4 | CGCTTCCCACCGGCCTGTAC |
| P5  | GACGCGGTCACGTTCTGGGAC |
| CDA knockout using pYL-*kasO*p\*-*Fn*Cas12a2 plasmid | CDA-spacer1-For | TTTTTGTTCAGATCGGTGGTCGGACCCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| CDA-spacer1-Rev | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGGGTCCGACCACCGATCTGAACA |
| CDA-spacer2-For | TTTTGCGGGAAGTTCCTGGGCGAGGTCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| CDA-spacer2-Rev | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGACCTCGCCCAGGAACTTCCCGC |
| CDA-spacer3-For | TTTTCTGGAGTTGAGCGGATAGTTCTCATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| CDA-spacer3-Rev | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGAGAACTATCCGCTCAACTCCAG |
| CDA-2KL-For | CATATCATCAATACTTGTCACATGAGAGTACTGATCTACAAGGTCGACG |
| CDA-2KL-Rev | GAGGTGAACGCGGTCGTACTGCCGAAGGTAAG |
| CDA-2KR-For | GCAGTACGACCGCGTTCACCTCTGGGGCTC |
| CDA-2KR-Rev | CTTCAGGCATCAAATTTCAGTGGCCTCGCCCTAGACGACCGCGAG |
| CDA-HR-CM-For | GTGCGAGTATCTGAAAGGGGATACGCGTCTAAGAACTTTAAATAATTTCTACTG |
| CDA-HR-CM-Rev | GAAAAGGACCCAGGTGGCACTTTTCGTCTAGAGCATATCATCAATACTTGT |
| PCM2-CDA-For1 | ACGCTCGGTTGCCGCCGGGCGTTTTTTAGCATATCATCAATACTTGTC |
| PCM2-CDA-Rev1 | ACGTGATGAAAAGGACCCAGGTGGCACTTTTCGTCTAGAGCTGTGCTTTATGAATC |
| P6 | GACCACGGAGTACACGCTGA |
| P7 | CGGCATCCATCTCGAACTCAC |
| P8 | CTCCGACACCACCGAACAG |
| DAP knockout using pYL-*kasO*p\*-*Fn*Cas12a2 and pCRISPomyces-2 plasmids | DAP-spacer-For1 | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATAGTGCGCCGACCGTCCCGTGCTC |
| DAP-spacer-Rev1 | AAACGAGCACGGGACGGTCGGCGCACTATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| DAP spacer For2 | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATaactggggcatcagtgcctggac |
| DAP spacer Rev2 | AAACGTCCAGGCACTGATGCCCCAGTTATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| DAP-KL-For | CAGAACGCTCGGTTGCCGCCGGGCGTTTTTTATCTAGAGTCGATGCCTTCCTGGAGCTG |
| DAP-KL-Rev | GCATTCCCCACGGCTTCACGCGCCGGGGAAGTCCTCTGTC |
| DAP-KR-For | GACAGAGGACTTCCCCGGCGCGTGAAGCCGTGGGGAATGC |
| DAP-KR-Rev | CGTGATGAAAAGGACCCAGGTGGCACTTTTCGTCTAGACACCAGCGCCCACCCGAAC |
| P9 | CAGGAGGGTGGGGAGCAGATC |
| P10 | CATTCGGTCGCGGCACATGC |
| P11 | TCGTGCTCGCCCGAATACAG |
| *RapTH* insertion | Cas12a-*RapTH*-SP-For | TTTTCCGTACTTGGCACATCGGCTGGGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| Cas12a-*RapTH*-SP-Rev | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCCCAGCCGATGTGCCAAGTACGG |
| Cas12a-*RapTH*-SP-Rev | GAGTCCGTACTTGGCACATCGGCTGGGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| TH-KL-For | AAAAACTGTATTATAAGTAAATGCATGTATACTAAACTCACTCTAGACCGAAATGGAGC |
| TH-KL-Rev | AGCTCATCGAGCTCATCGCCAAGAAGCTCGCGGGCTGAGTGATGTTGGAATTGGGTAAC |
| TH-KR-For | TAAATCTGAACTTCAGGCGGTTACCCAATTCCAACATCACTCAGCCCGCGAGCTTCTTG |
| TH-KR-Rev | AAAAAAAAACCCCGCCCCTGACAGGGCGGGGTTTTTTTTTCTAGACAGCGTGATCAAG |
| TH-HR-For | CTGTATTATAAGTAAATGCATGTATACTAAACTCACTCTAGACAGCGTGATCAAGATGG |
| TH-HR-Rev | ACCCCGCCCCTGACAGGGCGGGGTTTTTTTTCCGAAATGGAGCGACTCGAG |
| P18 | GACCGATCAGGGTGAGAC |
| P19 | CAGCTCTCGGCAAAATGGAG |
| Cas12a1-RapTH-SP For | TTTTCCGTACTTGGCACATCGGCTGGGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| Cas12a1-RapTH-SP Rev/ Cas12a2-RapTH-SP For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCCCAGCCGATGTGCCAAGTACGG |
| Cas12a2-RapTH-SP Rev | GAGTCCGTACTTGGCACATCGGCTGGGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| RT-PCR analyses of *Fn*Cas12a in *Streptomyces*  | *hrdB*-T For | GAGCCGCTCCCGGTTCCAC |
| *hrdB*-T Rev | GCATGCGGGCTCCTCACTC |
| RT *Fn*Cas12a For | CAACACGGCCATTCTGTTTATT |
| RT *Fn*Cas12a Rev | CCTTCTTCGGGTTGTCCTTATT |
| RT *hrdB* For | TTGATGACCTCGACCATGTG |
| RT *hrdB* Rev | GCGGTGGAGAAGTTCGACTA |
| Insertion of *kasO*p\* in front of *dptA* gene using pYL-*kasO*p\*-*Fn*Cas12a3 plasmid | Cas9-pYL-*kasO*p\*-For | AGCGTGCAGGACTGGGGGAGTTATGGACAAGAAGTACAGCATCG |
| Cas9-pYL-*kasO*p\*-Rev | CGGGGAACGCGTAGATCTGAATTC |
| Dapt-ccg-sp-For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCGAGAGATGGACATGCAGTCGCA |
| Dapt-ccg-sp-Rev | GAGTTGCGACTGCATGTCCATCTCTCGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| Dapt-atc-sp-For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCGCGAGAGATGGACATGCAGTCG |
| Dapt-atc-sp-Rev | GAGTCGACTGCATGTCCATCTCTCGCGATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| Dapt-cca-sp-For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATTCTCTCGCGGATCCCCTCAGGTG |
| Dapt-cca-sp-Rev | GAGTCACCTGAGGGGATCCGCGAGAGAATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| Dapt-KL-For | AACGCTCGGTTGCCGCCGGGCGTTTTTTATCTAGAAGGACAACTCGTGGACCACG |
| Dapt-KL-Rev | GACTGGGGGAGTTATGGACATGCAGTCGCAGC |
| Dapt- *kasO*p\*-For | CTGCATGTCCATAACTCCCCCAGTCCTGCAC |
| Dapt- *kasO*p\*-Rev | CGACCGCACCTGATGTTCACATTCGAACGGTC |
| Dapt-KR-For | GAATGTGAACATCAGGTGCGGTCGGCCAAC |
| Dapt-KR-Rev | TGATGAAAAGGACCCAGGTGGCACTTTTCGTCTAGAGAGGCCACCCTGTTCGTG |
| Dapt-p-check-F | GAATTCCCCGTCGCGCCG |
| pCFa-YES-Rev | AGCACGTGATGAAAAGGAC |
| Dapt-kas-Check-For | GCCGTAGGAGTTGATGTGG |
| Dapt-kas-check-innner-rev | CGCCGAGTTCGTGAACGAG |
| Dapt-kas-check-Rev | TGCGGTGTTGTAAAGTCGTG |
| *Fn*Cas12a3 system construction | *Fn*Cas12a-N607R-for | GACTTTAGCGcgCGGCTGGGATAAAAACAAG |
| *Fn*Cas12a-N607R-rev | ATCCCAGCCGcgCGCTAAAGTCGAGTTCTC |
| 2653-*Fn*Cas12a-pam-k613v-Fn617r-for | GATAAAAACGTGGAGCCCGACCGCACGGCCATTCTGTTTAT |
| 2653-*Fn*Cas12a-pam-k613v-n617r-rev | GAATGGCCGTGCGGTCGGGCTCCACGTTTTTATCCCAGCCGT |
| 2653-*Fn*Cas12a-Bd616n-for-new- | ACGTGGAGCCCAACCGCACGGCCATTCTGTTTAT |
| 2653-*Fn*Cas12a-Bd616n-rev-new | TGGCCGTGCGGTTGGGCTCCACGTTTTTATC |
| 2653-*Fn*Cas12a-pam-k660r-for | GAAGATTGTGTATAGACTGCTGCCGGGGGC |
| 2653-*Fn*Cas12a-pam-k660r-rev | GCCCCCGGCAGCAGTCTATACACAATCTTCTTGTA |
| 2653-*Fn*Cas12a-pam-k671r-for | CAAAATGCTCCCGCGAGTTTTTTTCTCGGCG |
| 2653-*Fn*Cas12a-pam-k671r-rev | GAGAAAAAAACTCGCGGGAGCATTTTGTTG |
| 2653-*Fn*Cas12a-pam-k180s-for | CGACGTACTTCTCGGGCTTCCATGAGAACCG |
| 2653-*Fn*Cas12a-pam-k180s-rev | CATGGAAGCCCGAGAAGTACGTCGTCCAGC |
| seq-*Fn*Cas12a-1 | CACGAGCACACTGGGGCAC |
| seq-*Fn*Cas12a-2 | AGCGGAAGAGCTGACGTTC |
| seq-*Fn*Cas12a-3 | CAATTTAGCCCAGATCTCCA |
| seq-*Fn*Cas12a-4 | AGGTTATAAGCTCACCTTC |
| *actII-orf4* and *redD* double deletion using pYL-*kasO*p\*-*Fn*Cas12a2 plasmid | AD-P1 | GAGACATCTTTGAAGACTTACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAG  |
| AD-P2 | ACCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAGAAATTATTTAAAG  |
| AD-P3 | GAGACATCTTTGAAGACTTACGCGTC  |
| AD-P4 | ACCGCGTCGATACGGAGCTG  |
| AD-P5 | CAGCTCCGTATCGACGCGGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCTGAG |
| AD-P6 | CGTGAAGAAGACATGAGTGCGCCCTGCGAGACGAGTCTCAGATCTACAACAGTAG  |
| AD-P7 | CAGCTCCGTATCGACGCGGT  |
| AD-P8 | GCCACGTGAAGAAGACATGAGTGC  |
| A-KL-For | GTTCAGAACGCTCGGTTGC  |
| A-KR-Rev | GAGGTCCTGCTCATCACGACTCTGCGCTTCAATC  |
| D-KL-For | GAAGCGCAGAGTCGTGATGAGCAGGACCTCGTG  |
| D-KR-Rev | AGCACGTGATGAAAAGGACCCAGGTGGCACTTTTCGGACAGCGAGTAGCGGAAG |
| AD-P1-1  | TCTTTGAAGACTTACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGAATGC |
| AD-P2-1  | ACCTGACTTCCGCCTGCAGGGCCGCGTCGATACGGAGCTGCATTCATCTACAACAGTAG |
| AD-P5-1  | CGAGTAGACGACGGAGACGTAGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCT |
| AD-P6-1  | CGTGAAGAAGACATGAGTGCGCCCTGCGAGACGAGTCTCAGATCTACAACAGTAGAAAT |
| T7-*actII*-Rev | AGACCCGTTTAGAGGCCCCAAGGGGTTATGCTACGCGTCGATACGGAGCTG |
| P12 | ACGCGTTCGACCGAGATG |
| P13 | TGTGGCTGTGTCGTTGTC |
| *actI-orf1* and *redX* double deletion using pYL-*kasO*p\*-*Fn*Cas12a2 plasmid | AX-P1-1 | TCTTTGAAGACTTACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATGGATTG |
| AX-P2-1 | ACCTGACTTCCGCCTGCAGGGCATGACGACTCTGCGCTTCAATCCATCTACAACAGTAG |
| AX-P4 | ACATGACGACTCTGCGCTTC  |
| AX-P5-1 | CGAGTAGACGACGGAGACGTAGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATCC |
| AX-P6-1 | CGTGAAGAAGACATGAGTTGCCTTCCTCGATCAACAGGTGGATCTACAACAGTAGAAAT |
| AX-P7 | GAAGCGCAGAGTCGTCATGT  |
| AX-P8 | GCCACGTGAAGAAGACATGAGTTG |
| rpsLp(XC)-For | GCCCTGCAGGCGGAAGTCAG  |
| rpsLp(XC)-Rev | TACGTCTCCGTCGTCTACTCG  |
| T7-*actI*-Rev  | AGACCCGTTTAGAGGCCCCAAGGGGTTATGCTAATGACGACTCTGCGCTTC  |
| T7-rpsLp(XC)-For | CTTGGGGCCTCTAAACGGGTCTTGAGGGGTTTTTTGGCCCTGCAGGCGGAAGTCAG |
| *actI*-KL-For | CAGAACGCTCGGTTGCCGCCGGGCGTTTTTTATCTAGAACGGTGAGAAGGTGCTCGTG |
| *actI*-KL-Rev | GAGATCGCACTCGTCCAGCGGCACCCCATCT  |
| *actI*-KR-For | GTGCCGCTGGACGAGTGCGATCTCGACTACGTTC  |
| *actI*-KR-Rev | ACCATGAGCAGTTGCAACGTCCTCGGGACCGGTC  |
| *redX* -KL-For | ACCGGTCCCGAGGACGTTGCAACTGCTCATGGTGGAG  |
| *redX* -KL-Rev | GTCCTCCAGGAGCACGTGGCATAC  |
| *redX* -KR-For | GTATGCCACGTGCTCCTGGAGGACGCAACGCGATGAACTTC  |
| *redX* -KR-Rev | AGCACGTGATGAAAAGGACCCAGGTGGCACTTTTCGCGTCCAGTCCGAGTTGTAC |
| P14 | CTGGAATCGTATCGGAATCTCC |
| P15 | TGAGTACGGACCGCAGCTTC |
| P16 | ACATCGAGGTCGACGTGGCAC  |
| P17 | TCGTTGGTGCCGGAGAAGATC  |
| Site mutations rpsL (nt262 A>G, nt264G>A, nt267C>T) using pYL-*kasO*p\*-*Fn*Cas12a3 plasmid | rpsL-SP-For | ACGCGTCTAAGAACTTTAAATAATTTCTACTGTTGTAGATTGTGAAGGACCTGCCGGGTGTTC |
| rpsL-SP-Rev | GAGTGAACACCCGGCAGGTCCTTCACAATCTACAACAGTAGAAATTATTTAAAGTTCTTAGAC |
| rpsL-KL-For | GTTCAGAACGCTCGGTTGCCGCCGGGCGTTTTTTACAGGTGACGTACGAGCGCGCGCC  |
| rpsL-KL-Rev | AACACCCGGCAGATCTTCCACACGGCCGCCGCGCACGA  |
| rpsL-KR-For | CCGTGTGGAAGATCTGCCGGGTGTTCGCTAC  |
| rpsL-KR-Rev | AGCACGTGATGAAAAGGACCCAGGTGGCACTTTTCGAGCTTGTTCACGAAGCAGATGC  |
| rpsL-check-For | GTCTACTACCGGCTCATG |
| rpsL-check-Rev | CAGTCTTCTCTCGGTCTC |
| Deletion of partial sequences of *traJ* gene from *Fn*Cas12a1 system | 1-Δ*traJ*-For | TGAGCACGTCCGCGAGCTGTCAGCGCTTGTAGTCGATGGCCTC  |
| 1-Δ*traJ*-Rev | AGGCCATCGACTACAAGCGCTGACAGCTCGCGGACGTGCTCATAGTC |
| Replace B1006 terminator of crRNA in *Fn*Cas12a1 with oop terminator | 1*-actII-orf4*-DR36-oop For | CTATGATCGGGGCGTTCCTGC |
| 1-*actII-orf4*-DR36-oop Rev | GCAAAACTAAAAAACTGTATTATAAGTAAATGCATGTATACTAAACTCACACGACTCTGCGCTTCAATC |
| Replace oop terminator of crRNA in *Fn*Cas12a2 system with B1006 terminator | 2-*actII-orf4*-DR36-oop For | CTATGATCGGGGCGTTCCTGC |
| 2-*actII-orf4*-DR36-oop Rev | ATAATTATTTTTATAGCACGTGATGAAAAGGACCCAGGTGGCACTTTTCGGGTGAACCCATGGTCGTC |

**Supplementary Table 2**. Different elements among the *Fn*Cas12a1, *Fn*Cas12a2 and *Fn*Cas12a3 systems

|  |  |  |  |
| --- | --- | --- | --- |
| Descriptions | *Fn*Cas12a1 | *Fn*Cas12a2 | *Fn*Cas12a3 |
| Element for transfer | *oriT*, *traJ* | *oriT,* partial sequences of *traJ* | *oriT*, partial sequences of *traJ* |
| Selection marker  | *aac(3)IV*, *codA(sm)* | *aac(3)IV*, *pSG5* | *aac(3)IV*, *pSG5* |
| Replication origin in *Streptomyces* | *rep* (pIJ101) | *rep* (pSG5) | *rep* (pSG5) |
| Terminator of crRNA cassette | B1006 terminator | oop terminator | oopterminator |
| *Fn*Cas12a | wide-type *Fn*Cas12a | wide-type *Fn*Cas12a | *Fn*Cas12amutationEP16 |

**Supplementary Table 3**. List of plasmids used in this study

|  |  |  |
| --- | --- | --- |
| Plasmids | Description | References |
| pCRISPomyces-2 | CRISPR/Cas system carrying *Sp*Cas9 controlled by the rpsLp(XC) promoter | (Cobb et al., 2015) |
| pWHU2653 | CRISPR/Cas9-*CodA(sm)* combined system carrying *Sp*Cas9 controlled by the *aac(3)IVp* promoter | (Zeng et al., 2015) |
| pYL-rpsLp(XC)-*Fn*Cas12a1 | *Fn*Cas12a1 system carrying Cas12a controlled by the rpsLp(XC) promoter | This study |
| pYL-*kasO*p\*-*Fn*Cas12a1 | *Fn*Cas12a1 system carrying Cas12a controlled by the *kasO*p\* promoter | This study |
| pYL-*ermE*p\*-*Fn*Cas12a1 | *Fn*Cas12a1 system carrying Cas12a controlled by the *ermE*p\* promoter | This study |
| pYL-Potr\*-*Fn*Cas12a1 | *Fn*Cas12a1 system carrying Cas12a controlled by the Potr\* system | This study |
| pYL-rpsLp(XC)-*Fn*Cas12a2 | *Fn*Cas12a2 system carrying Cas12a controlled by the rpsLp(XC) promoter | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2 | *Fn*Cas12a2 system carrying Cas12a controlled by the *kasO*p\* promoter | This study |
| pYL-*ermE*p\*-*Fn*Cas12a2 | *Fn*Cas12a2 system carrying Cas12a controlled by the *ermE*p\* promoter | This study |
| pYL-Potr\*-*Fn*Cas12a2 | *Fn*Cas12a2 system carrying Cas12a controlled by the Potr\* system | This study |
| pYL-rpsLp(XC)-*Fn*Cas12a1-*actII*-*orf4*-DR19 | pYL*-*rpsLp(XC)*-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-rpsLp(XC)-*Fn*Cas12a1-*actII*-*orf4*-DR36 | pYL*-*rpsLp(XC)*-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a1-*actII*-*orf4*-DR19 | pYL*-kasO*p\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a1-*actII*-*orf4*-DR36 | pYL*-kasO*p\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-*ermE*p\*-*Fn*Cas12a1-*actII*-*orf4*-DR19 | pYL*-ermE*p\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-*ermE*p\*-*Fn*Cas12a1-*actII*-*orf4*-DR36 | pYL*-ermE*p\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-Potr\*-*Fn*Cas12a1-*actII*-*orf4*-DR19 | pYL-Potr\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-Potr\*-*Fn*Cas12a1-*actII*-*orf4*-DR36 | pYL-Potr\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-rpsLp(XC)-*Fn*Cas12a2-*actII*-*orf4*-DR19 | pYL*-*rpsLp(XC)*-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-rpsLp(XC)-*Fn*Cas12a2-*actII*-*orf4*-DR36 | pYL*-*rpsLp(XC)*-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-*actII*-*orf4*-DR19 | pYL*-kasO*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-*actII*-*orf4*-DR36 | pYL*-kasO*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-*ermE*p\*-*Fn*Cas12a2-*actII*-*orf4*-DR19 | pYL*-ermE*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-*ermE*p\*-*Fn*Cas12a2-*actII*-*orf4*-DR36 | pYL*-ermE*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-Potr\*-*Fn*Cas12a2-*actII*-*orf4*-DR19 | pYL-Potr\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 19-nt DR sequence | This study |
| pYL-Potr\*-*Fn*Cas12a2-*actII*-*orf4*-DR36 | pYL*-*Potr\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting *actII*-*orf4* and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-ACT | pYL*-kasO*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting ACT gene cluster and a 36-nt DR sequence | This study |
| pCRISPomyces-2-DAP | pCRISPomyces-2 plasmid carrying the 20-nt spacer targeting DAP gene cluster and an 81-nt gRNA sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-DAP | pYL-*kasO*p\**-Fn*Cas12a2 plasmid carrying the 23-nt spacer targeting DAP gene cluster and a 36-nt DR sequence | This study |
| pYL*-ermE*p\*-*Fn*Cas12a1-*rapTH*-DR36 | pYL-*ermE*p*\*-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *rapTH* and a 36-nt DR sequence | This study |
| pYL-Potr\*-*Fn*Cas12a1-*rapTH*-DR36 | pYL-Potr*\*-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *RapTH* gene and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a1-*rapTH*-DR36 | pYL-*kasO*p\**-Fn*Cas12a1 plasmid carrying the 23-nt spacer targeting *RapTH* gene and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a3 | pYL-*kasO*p\*-*Fn*Cas12a3 plasmid carrying *Fn*Cas12a mutant EP16 (N607R/K613V/N617R/K180S/K660R/D616N) | This study |
| pYL-*kasOp\** -*Fn*Cas12a3-DAP (CCG) | pYL-*kasO*p\*-*Fn*Cas12a3 carrying a 36-nt DR with a spacer adjacent to CCG PAM  | This study |
| pYL-*kasOp\**-*Fn*Cas12a3-DAP(ATC) | pYL-*kasO*p\*-*Fn*Cas12a3 carrying a 36-nt DR with a spacer adjacent to ATC PAM | This study |
| pYL-*kasO*p\*-*Fn*Cas12a3-DAP(CCA) | pYL-*kasO*p\*-*Fn*Cas12a3 carrying a 36-nt DR with a spacer adjacent to CCA PAM | This study |
| pYL-*kasO*p\*-*Fn*Cas12a3-rpsL | This plasmid carrying the 23-nt spacer targeting *rpsL* gene and a 36-nt DR sequence | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-*actII*-rpsLp(XC)-T7-*redD*-4kb | This plasmid carrying two 23-nt spacers targeting *actII-orf4* and *redD* respectively and a 36-nt DR sequences | This study |
| pYL-*kasO*p\*-*Fn*Cas12a2-*actI*- rpsLp(XC)-T7-*redX*-4kb | This plasmid carrying two 23-nt spacers targeting *actI-orf1* and *redX* respectively and two 36-nt DR sequences | This study |
| pYL-*kasO*p\*-FnCas12a1-Δ*traJ* | The partial sequences of *traJ* gene were deleted from pYL-rpsLp(XC)-*Fn*Cas12a1 | This study |
| pYL-rpsLp(XC)-FnCas12a1-Δ*traJ* | The partial sequences of *traJ* gene were deleted from pYL-*kasO*p\*-*Fn*Cas12a1 | This study |

**Supplementary Table 4**. Deletion efficiencies in the *actII*-*orf4* gene deletion experiments by using two *Fn*Cas12a systems in the presence of OTC

|  |  |  |
| --- | --- | --- |
| Plasmids | OTC (μM) | Deletion efficiency |
| pYL-Potr\*-*Fn*Cas12a1-*actII*-*orf4*-DR19 | 0 | 9.5%±6.7% |
| 0.6 | 9.5%±6.7% |
| 1.2 | 14.3%±11.7% |
| 3.0 | 28.6%±11.7% |
| pYL-Potr\*-*Fn*Cas12a1-*actII*-*orf4*-DR36 | 0 | 28.6%±14.3% |
| 0.6 | 5.6%±7.9% |
| 1.2 | 23.8%±13.5% |
| 3.0 | 14.3%±11.7% |
| pYL-Potr\*-*Fn*Cas12a2-*actII*-*orf4*-DR19 | 0 | 23.8%±6.7% |
| 0.6 | 9.5%±6.7% |
| 1.2 | 35.7%±7.1% |
| 3.0 | 9.5%±6.7% |
| pYL-Potr\*-*Fn*Cas12a2-*actII*-*orf4*-DR36 | 0 | 21.4%±7.1% |
| 0.6 | 15.1%±11.7% |
| 1.2 | 69.1%±2.4% |
| 3.0 | 22.2%±20.8% |

**Supplementary Table 5**. Efficiencies of multiplex and precise gene editing

|  |  |  |
| --- | --- | --- |
| Targets | Plasmids | Efficiency |
| ACT (deletion) | pYL-*kasO*p\*-*Fn*Cas12a2-ACT | 92.9%±7.2% |
| CDA (deletion) | pYL-*kasO*p\*-*Fn*Cas12a2-CDA-sp1 | 55.6%±7.9% |
| pYL-*kasO*p\*-*Fn*Cas12a2-CDA-sp2 | 18.8%±6.3% |
| pYL-*kasO*p\*-*Fn*Cas12a2-CDA-sp3 | 25.0%±0.0% |
| pYL-*kasO*p\*-*Fn*Cas12a2-CDA-2sp | 25.0%±0.0% |
| DAP(deletion)  | pYL-*kasO*p\*-*Fn*Cas12a2-DAP-sp1 | 25.0%±0.0% |
| pYL-*kasO*p\*-*Fn*Cas12a2-DAP-sp2 | 0.0%±0.0% |
| *RapTH* (insertion） | pYL-*ermE*p\*-*Fn*Cas12a1-*RapTH*-DR36 | 8.3% |
| pYL-Potr\*-*Fn*Cas12a1-*RapTH*-DR36 | 0.0% (0 μM OTC) |
| 0.0% (0.6 μM OTC) |
| 25.0% (1.2 μM OTC) |
| 8.3% (3.0 μM OTC) |
| DAP (insertion) | pYL-*kasO*p\*-*Fn*Cas12a3-DAP(CCG) | 50.0%±12.5% |
| pYL-*kasO*p\*-*Fn*Cas12a3-DAP(CCA) | 40.0%±17.4% |
| pYL-*kasO*p\*-*Fn*Cas12a3-DAP(ATC) | 23.6%±8.6% |
| rpsL (mutation) | pYL-*kasO*p\*-*Fn*Cas12a3-rpsL (CCG) | 12.5%±0.0% |
| *actII-orf4/redD* double deletion | pYL-*kasO*p\*-*Fn*Cas12a2-*actII*-rpsLp(XC)-T7-*redD*-4kb | 14.3%±0.0% |
| *actI-orf1/redX* double deletion | pYL-*kasO*p\*-*Fn*Cas12a2-*actI*-rpsLp(XC)-T7-*redX*-4kb | 14.3%±0.0% |
| *actII-orf4* | pYL-*kasO*p\*-*Fn*Cas12a1-*actII-orf4*-DR36-oop | 95.2%±6.7% |
| *actII-orf4* | pYL-*kasO*p\*-*Fn*Cas12a2-*actII-orf4*-DR36-B1006 | 47.6%±17.3% |

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# References

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