

File S1

Supplementary Information

gRNA sequences

<i>e</i>	GCCACAATTGTCGATCGTCA	(Port <i>et al.</i> 2014)
<i>y</i>	GCGATATAGTTGGAGCCAGC	(Gratz <i>et al.</i> 2014)
<i>w</i>	CCGAGAACCTCACCTATGCC	
<i>MED27</i>	CTCCTCGCCGCTCTCCCCAT	
<i>ham</i>	GGCCATTACAATAAGCTCTT	
<i>dpn</i>	CCTGTGCGGTTGATCTACCA	
<i>mγ</i>	GTGAAGTAATCCTATCTACC	

Sequences of recovered indel alleles

e

ATGGGTT CGCT GCCACAATTGTCGATCGTCA AGGGTCTGCAGCAAGACTT	wt
ATGGGTT CGCTGCCACAATTGTCGATC - TCAAGGGTCTGCAGCAAGACTT	- 1
ATGGGTT CGCTGCCACAATTGTC - - - - - TGCAGCAAGACTT	- 14
ATGGGTT CGCTGCCACAATTGTCGA - - - - CAAGGGTCTGCAGCAAGACTT	- 4
ATGGGTT CGCTGCCACAATTGT - - - - - GTCA AGGGTCTGCAGCAAGACTT	- 5
ATGGGTT CGCTGCCA - - - - - AGGGTCTGCAGCAAGACTT	- 16
ATGGGTT CGCTGCCAC - - - - - A AGGGTCTGCAGCAAGACTT	- 14
ATGGGTT CGCTGCCACAATTGTCGATC - - - - - TGCAGCAAGACTT	- 10
ATGGGTT CGCTGCCACAATTGTCGATC - tgcaaga - CTGCAGCAAGACTT	- 9/+7
ATGGGTT CGCTGCCACAATTGTCGATCGa TCAAGGGTCTGCAGCAAGACTT	+ 1

MED27

GCCGATGGAAC CCGATGGGGAGAGCGGCGAGGAGAGCCG	wt
GCCGATGGAAC CCGATG ctcgaacc GAGCGGCGAGGAGAGCCG	-4/+8
GCCGATGGAAC CCGATG - GGAGAGCGGCGAGGAGAGCCG	-1
GCCGATGGAAC CCGATG ggagagcggagat GGGAGAGCGGCGAGGAGAGCCG	+13

ssODN donor sequences

The homology arms are in upper case, the central V5 sequence in lower case, and the stop codons underlined. The gRNA target sequences are in bold, and are followed by a PAM (NGG). Silent nucleotide changes were introduced in the standard V5 sequence to create a *HindIII* restriction site. Donor sequences were synthesized in the same orientation as the gRNA. For all cases presented here, the insertion of the V5 tag destroyed the gRNA target site so that V5-tagged chromosomes would not be recognized and cleaved again.

ham-V5

GCAACATCAGCAACAACAACAGCAGCAGCGGCAACAACAACAACAGCTCCA**AGGCCATTACA**
ATAAGCTCTggtaagcctatacctaaccctcttcttggtctagatagcacg**TAGGAATCGTA**
AGAGATTAGAGAGCTTGCTTTCCTCGTTGTTAAATGATAAACTTACTTTAACTC

dpn-V5

GCAATGAAATAAATTAACAATGTATAAACGTAACATAGAAGTGG**CCTGTGCGGTTGATCTA**c
gtgctatctagaccaagaagagggttaggtataggcttacc**CCAC**GGCCTCCAAGCGGAGCT
CGACTTTTCGATGGCGCCAGCTAGGAATTTCTTGGACGAGGGC

m γ -V5

CTTTCTATGATCTCTCCTTCGTTTTTTTTTTTTTTCAGTTGTGGT**GTGAAGTAATCCTATCTA**
cgtgctatctagaccaagaagagggttaggtataggcttacc**CCAGGGACGCCAGACGTTCT**
CCTCGTCCTCGCTGTCGTCCTCCATCTTGGTCACATCGATGGTGGTC