

SUPPLEMENTARY INFORMATION

A platform for rapid prototyping of synthetic gene networks in mammalian cells

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Supplementary Figure 14. Microscopy images of isolated colonies representing the 18 different phenotypes of the circuit library.

Supplementary Figure 15. Annotated maps of vectors used in this study.

Supplementary Table

Supplementary Table 1. Comparison of integration efficiencies of different recombinases.

Supplementary Text

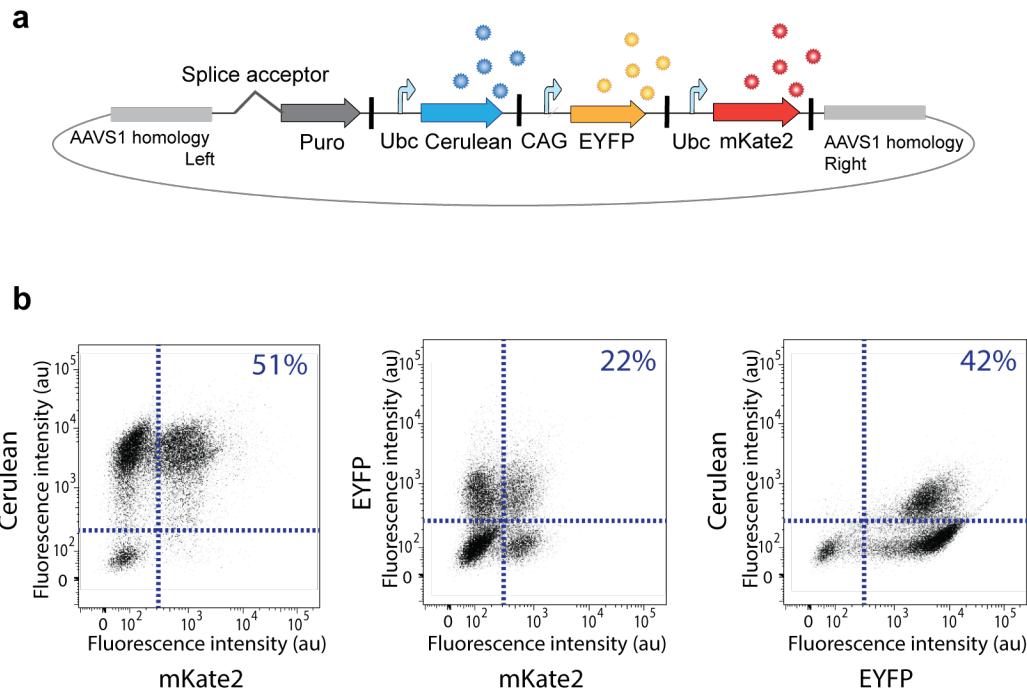
Supplementary Text 1. DNA sequences of the probes used for the Southern Blots.

Supplementary Text 2. DNA sequences of the primers used for genomic PCRs.

Supplementary Text 3. DNA sequences of the genetic components from our mammalian parts library used in this study.

Supplementary Text 4. DNA sequences of all plasmids from the mMoclo library.

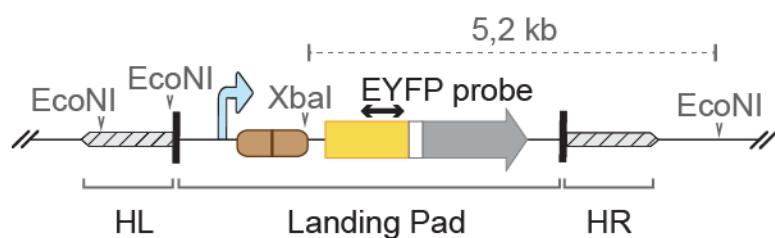
Supplementary Figure 1. Non-homogeneity of transgene expression after zinc-finger nuclease mediated integration of a 4-gene circuit.



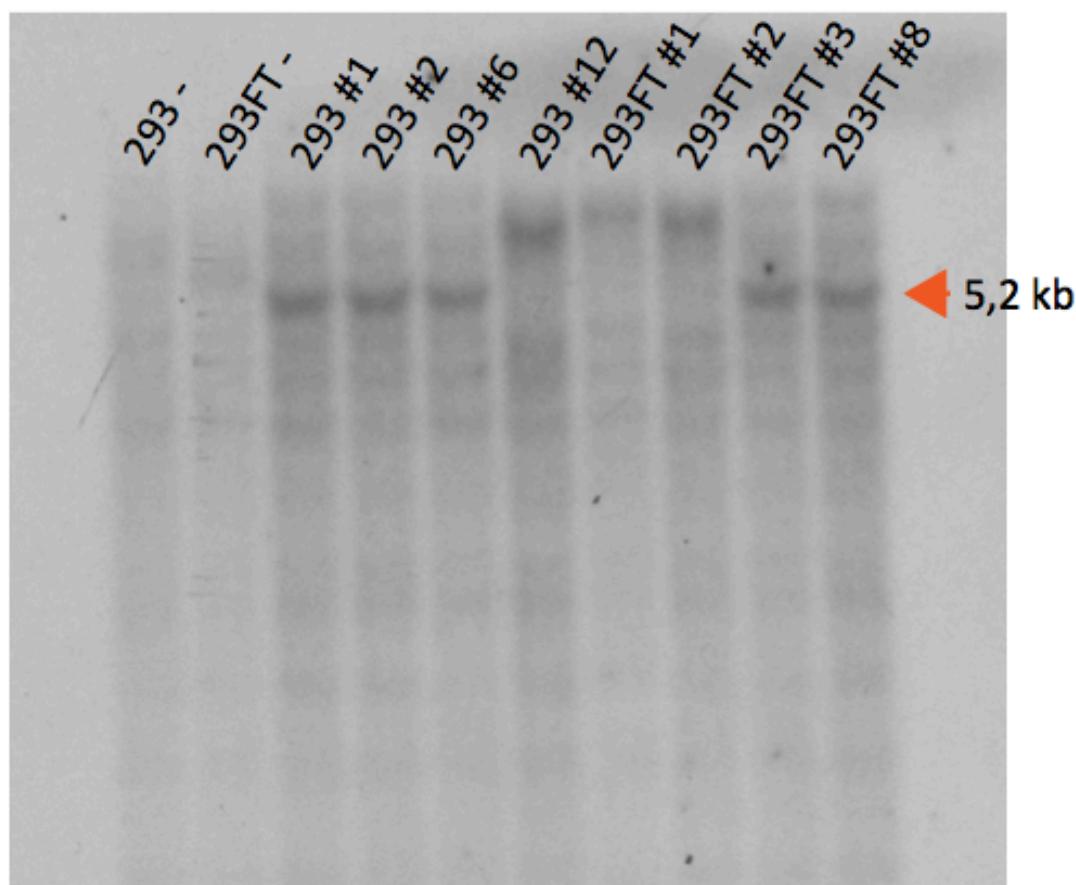
(a) Schematic representation of the 4-gene circuit used for integration; the circuit comprises (1) a promoterless puromycin resistance gene activated only after integration in the AAVS1 locus¹, (2) Ubc promoter constitutively expressing a Cerulean reporter, (3) CAG promoter constitutively expressing an EYFP reporter and (4) Ubc promoter constitutively expressing an mKate2 reporter. AAVS1 homology regions flank the circuit to target integrations into the AAVS1 locus. **(b)** Representative two-dimensional fluorescence density plots demonstrating significant heterogeneity in reporter expression levels within the polyclonal population of cells after integration of the circuit and selection with Puromycin. Selection was started 3 days post transfection and cells were assayed 14 days post transfection.

Supplementary Figure 2. Landing pad chassis cell line construction: Southern blot genomic analysis of landing pad integration with zinc-finger nuclease in wild-type cells.

A



B

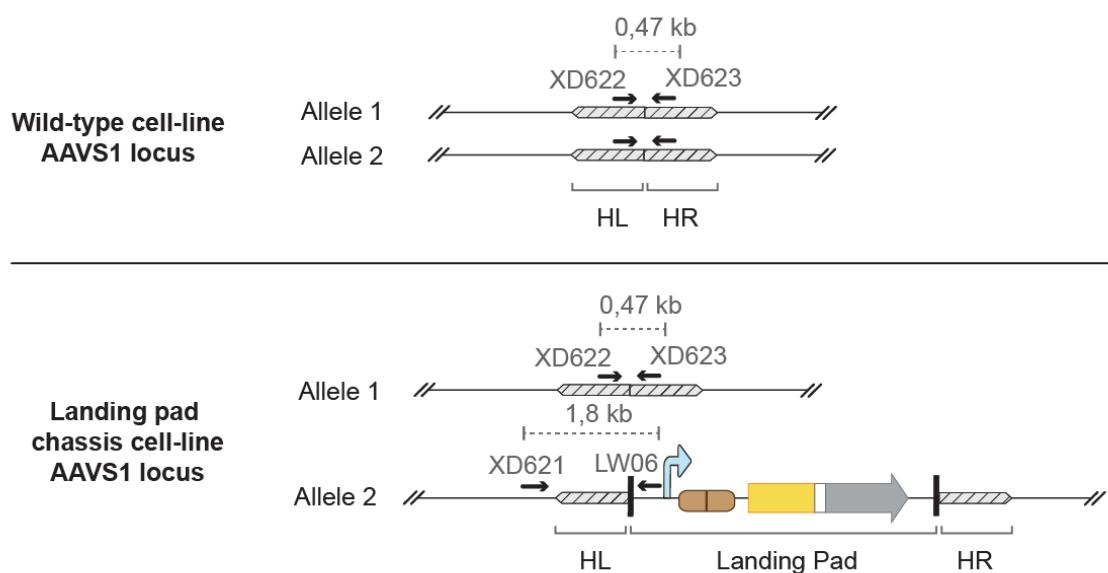


(A) Schematic of the Southern probe and the restriction enzyme cut sites. The landing pad is inserted between the left homology and right homology sequences (HL & HR). The 300 bp probe used for the Southern Blots corresponds to an internal sequence of EYFP. Digestion of chromosomal DNA with EcoNI and XbaI results in a 5.2kb band containing the EYFP gene for the correct clones.

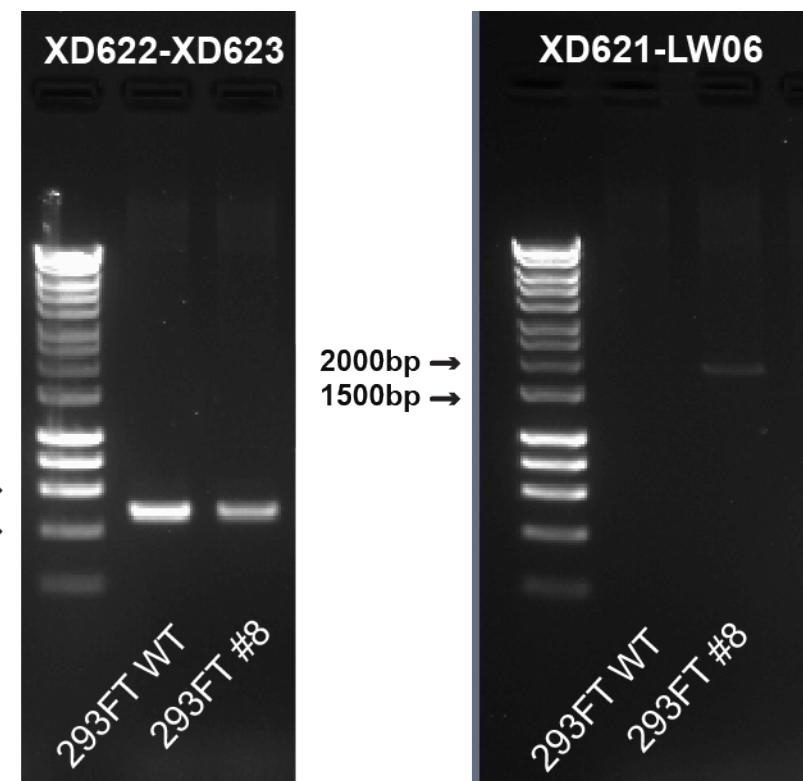
(B) Wild-type cell lines of HEK293FT and HEK293 with no landing pad were used as negative controls (first two lanes on the left). Integration in the correct locus was confirmed for 3 out of 4 clones for HEK293 cell lines, and for 2 out of 4 clones for HEK293FT. Clone HEK293FT#8 (denoted in the text as HEK293FT-LP) was chosen as the monoclonal chassis cell line for all downstream circuit integrations.

Supplementary Figure 3. Landing pad chassis cell line validation: PCR confirmations of mono-allelic landing pad integration.

A



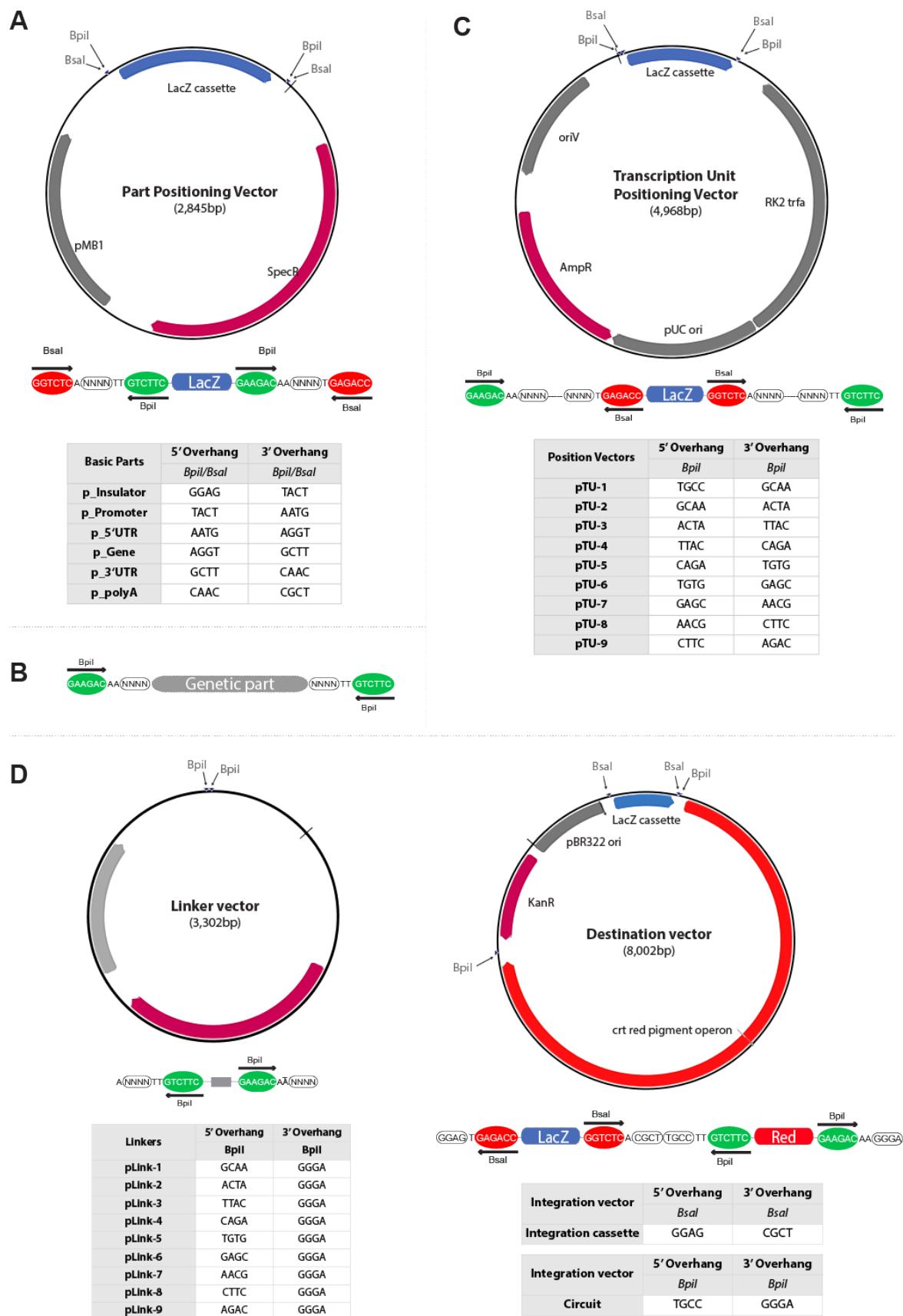
B



(A) Schematic of the PCRs and primers used. Primers XD622 and XD623 allow amplification of a 0.47kb sequence that overlaps the ZFN cleavage site within the AAVS1 locus. XD622 and XD623 are respectively located within the Left Homology arm (HL) and the Right Homology arm (HR) used for integration of the landing pad. The primers XD621 and LW06 allow amplification of a 1.8kb sequence when the landing pad is specifically inserted in the AAVS1 locus in the right orientation.

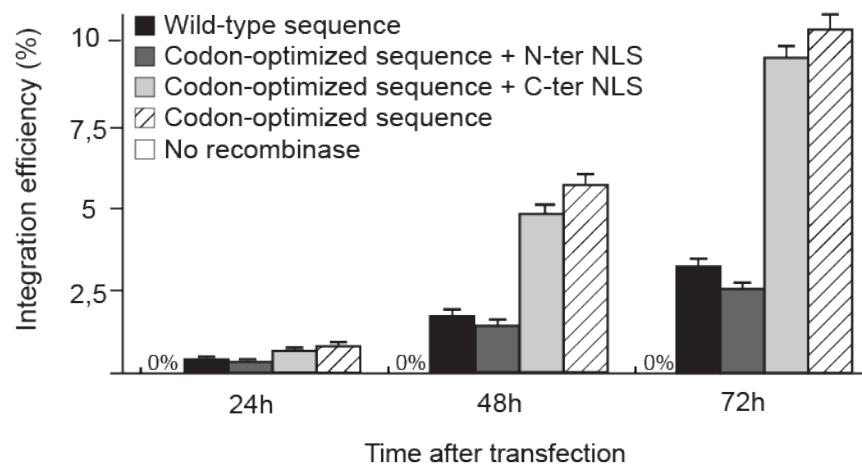
(B) PCR gels on WT and Chassis cell lines. Control PCRs on the wild type cell line with the two primer pairs resulted in a single band at 0.47kb and no band at 1.8kb. PCRs on the chassis cell line resulted in both a band at 0.47kb and a band at 1.8kb, confirming the presence of one non-modified allele and one allele carrying the landing pad.

Supplementary Figure 4. Details of the mMoClo backbones used in the assembly workflow depicted in Figure 2.



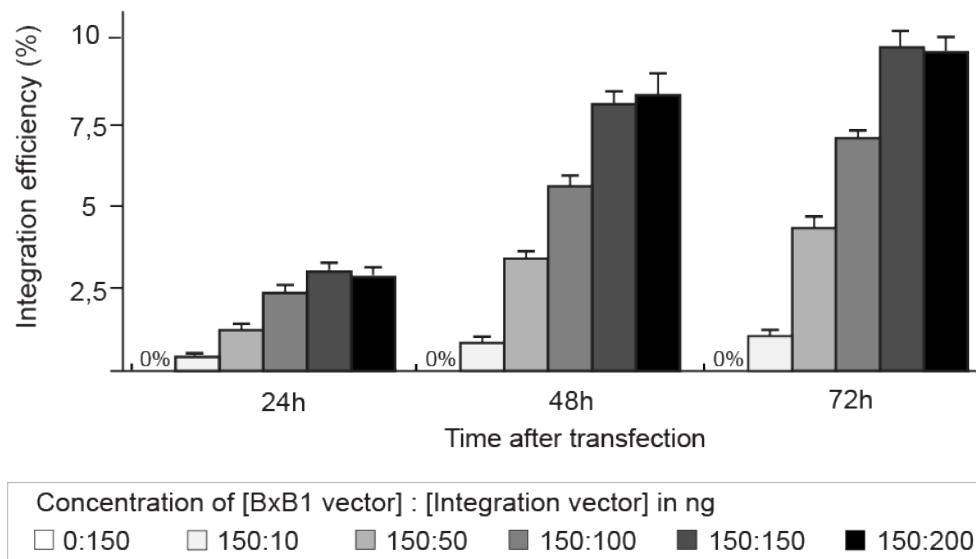
The mMoClo library includes a library of empty backbones that are used to assemble (multi)gene circuits. **(A)** The Part Positioning Vectors are used to create a library of single genetic parts (required later to build functional transcription units): Insulator, Promoter, 5'UTR, Gene coding sequence, 3'UTR and polyA signal. **(B)** To clone a genetic part into the spectinomycin resistant Part Positioning Vector of choice, the part has to be flanked by a BpiI restriction site on each end, together with 4bp overhangs matching the ones from the desired Part Positioning Vector. The BpiI mediated Golden Gate reaction between the genetic part and the Part Positioning Vector replaces the lacZ cassette with the desired genetic part into the Part Positioning Vector and thus allows a rapid and easy visual blue/white screening of the transformed bacteria. The library Part vector created this way does not contain BpiI restriction sites anymore. **(C)** To create a functional transcription unit, the BsaI mediated Golden Gate reaction requires the presence of 6 plasmids: 1 Insulator plasmid, 1 Promoter plasmid, 1 5'UTR plasmid, 1 Gene plasmid, 1 3'UTR plasmid, 1 polyA plasmid (all from the library of parts) and 1 ampicillin resistant Transcription Unit Positioning Vector. Assembly of the transcription unit into the latter one replaces the lacZ cassette and thus allows a rapid and easy visual blue/white screening of the transformed bacteria. The choice of the pTU (Transcription Unit positioning Vector) is directly linked to the position of transcription unit in the final circuit. The transcription unit vector created this way does not contain BsaI restriction sites anymore. **(D)** To create an integration vector (containing a BxB1 recombination site) that is used to assemble a complete circuit, an integration cassette is assembled first into the kanamycin resistant Destination vector, using a mock promoter part together with the BxB1 attB site as the 5'UTR part and a fluorescence/resistance marker of choice for the Gene part. Assembly of this promoterless transcription unit replaces the lacZ gene from the Destination Vector, allowing for a red/green selection of the transformed bacteria. Before this step, the destination vector expresses both lacZ gene and the *crt* red pigment operon, resulting in green pigmentation of the bacteria carrying an intact Destination Vector. The integration vector assembled this way can then be repeatedly used to assemble multi-gene circuits with a BpiI based Golden Gate reaction by combining a set of assembled transcription unit vectors (following each other) together with the linker vector corresponding to the position of the last transcription unit vector. The assembled circuit replaces the *crt* red pigment producing operon and thus allows for easy read/white screening to pick the correct clones.

Supplementary Figure 5. Effect of nucleotide sequence optimization and nuclear localization signal on the integration efficiency of BxB1 recombinase.



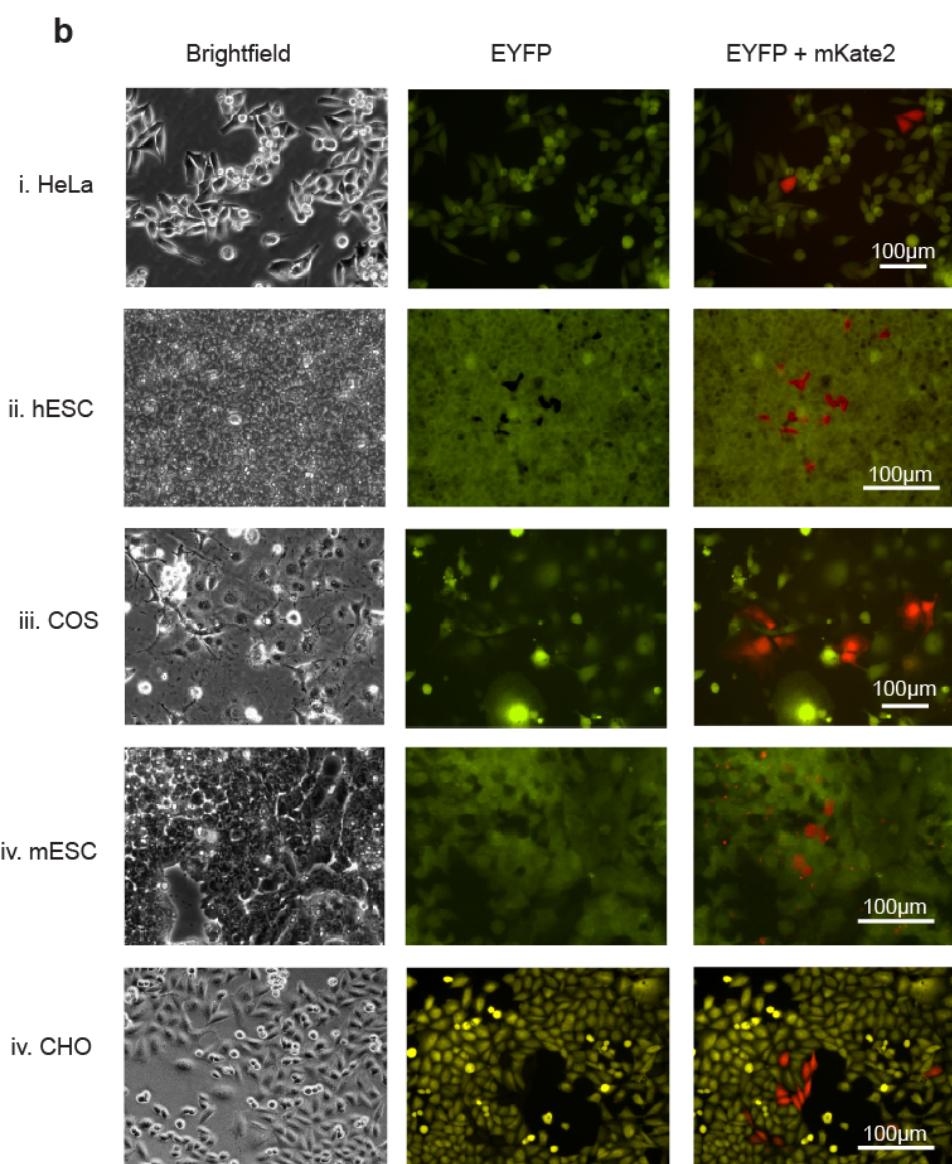
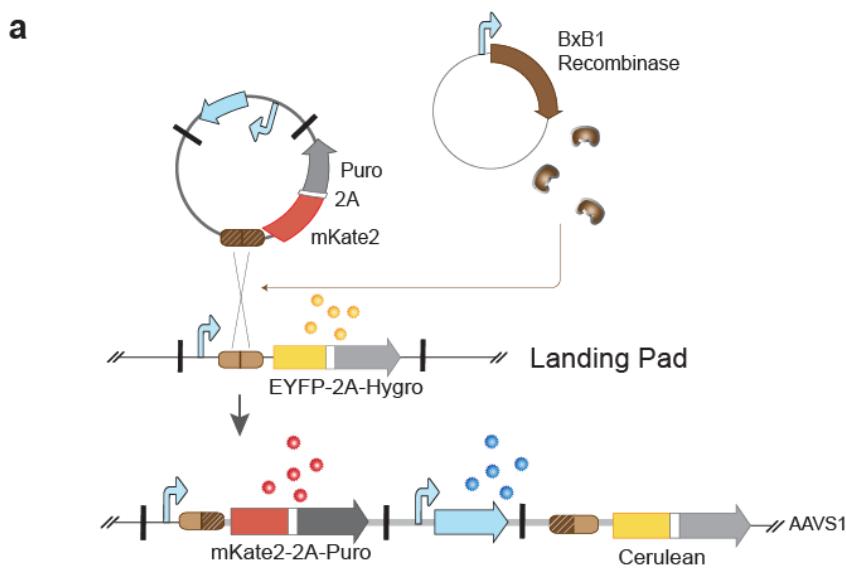
A 12kb integrative plasmid comprising a promoterless Puromycin-2A-mKate2 followed by constitutively expressed Cerulean was co-transfected with BxB1 expressing plasmid into the HEK293FT-LP chassis cell line. Integration efficiency was measured as percentage of mKate2 positive cells within the transfected cell population. The number of cells expressing mKate2 increased with time and reached a maximum 3 days after transfection (we monitored the expression of mKate2 up to 10 days post-transfection, data not shown). Nuclear Localization Signal (NLS) fusion to BxB1 did not improve the integration efficiency. When NLS was added at the N-terminus of the recombinase, significant negative impact on integration efficiency was observed. Our hypothesis is that in this case the NLS may sterically interfere with the N-terminal domain involved in the catalytic cleavage and ligation of DNA. Error bars represent standard deviation from 3 independently repeated experiments.

Supplementary Figure 6. Influence of vector ratios (BxB1 expression vector vs integration vector) on integration efficiency.



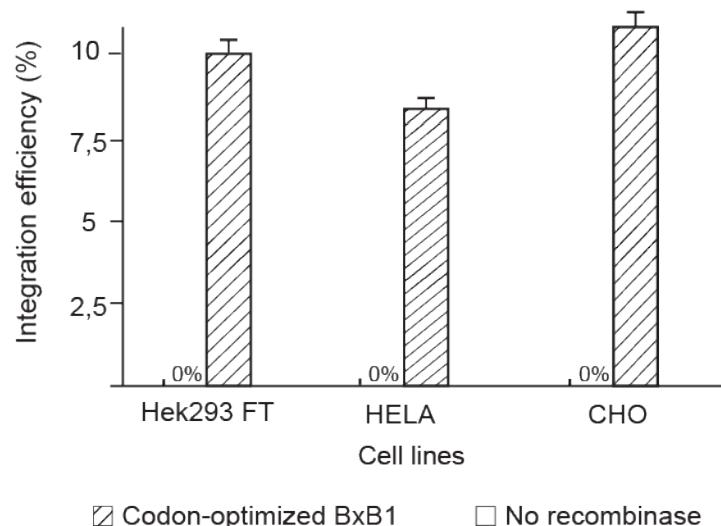
We co-transfected, in different ratios, a 12kb integrative plasmid comprising a promoterless Puromycin-2A-mKate2 construct followed by constitutively expressed Cerulean together with a codon-optimized BxB1 expressing plasmid into the HEK293FT-LP chassis cell line. The transfection mix was supplemented with mock plasmid DNA to always transfet the cells with a total of 400ng of DNA. Integration efficiency was measured as percentage of mKate2 positive cells within the transfected cell population. Error bars represent the standard deviation from 3 independently repeated experiments.

Supplementary Figure 7. Microscopy images of different model chassis cell lines after integration of a simple circuit.



(A) The integrative plasmid comprising a promoterless mKate-2A-puro gene cassette and a constitutively expressed Cerulean fluorescent reporter was co-transfected with the BxB1 recombinase expression plasmid in various monoclonal chassis cell lines. Integration events in the landing pad can be monitored in the transfected cells by both expression of mKate2 and reduction in expression of EYFP. **(B)** Fluorescent microscopy images of monoclonal chassis cell lines four days after co-transfection with BxB1 expression plasmid and the integrative plasmid: (i) Hela cells, (ii) human Embryonic Stem Cells, (iii) COS cells, (iv) mouse Embryonic Stem Cells and (v) CHO cells.

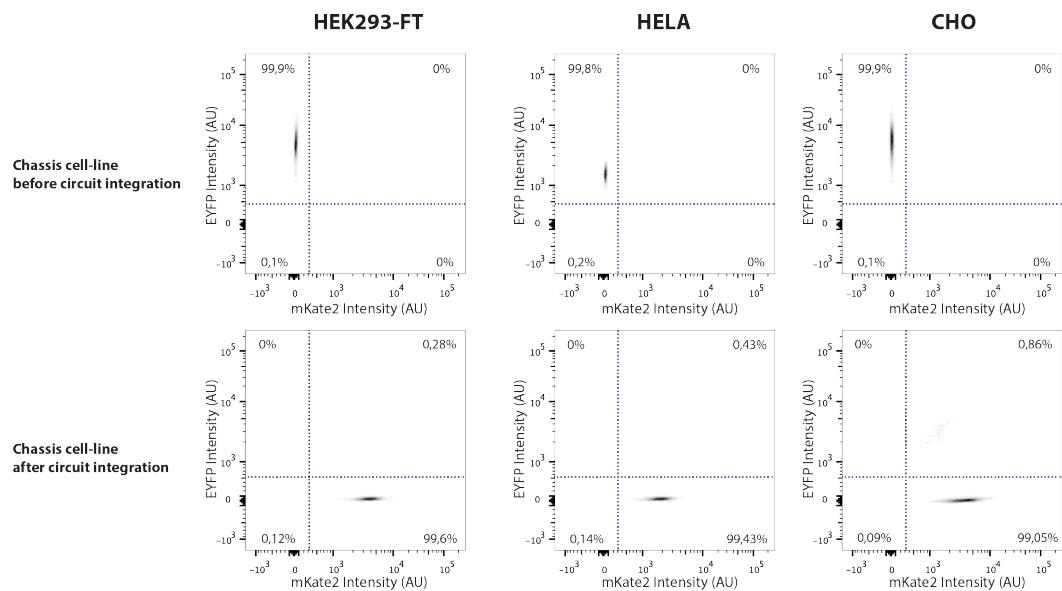
Supplementary Figure 8. Comparison of circuit integration efficiency in three different chassis cell lines.



We co-transfected a 12kb integrative plasmid comprising a promoterless Puromycin-2A-mKate2 followed by constitutively expressed Cerulean together with a codon-optimized BxB1 expressing plasmid into our HEK293FT, HeLa and CHO chassis cell-lines. Integration efficiency was measured as percentage of mKate2 positive cells within the transfected cell population 3 days after transfection.

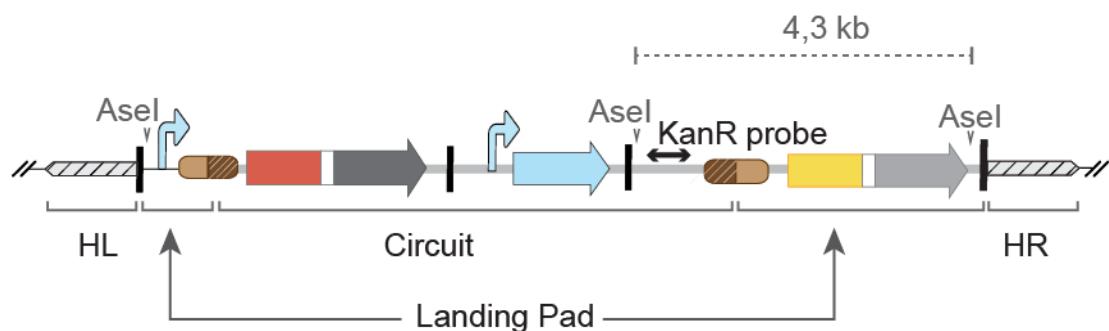
Supplementary Figure 9. Flow cytometry analysis of chassis cell line fluorescence before and after integration of circuits and selection with puromycin.

FACS comparative analysis of EYFP and mKate2 fluorescence levels in different chassis cell-lines before integration of the integration vector carrying the promoterless mKate2-2A-puro cassette, and 14 days after integration and selection with puromycin. After selection, more than 99% of cells that survive selection are mKate2 positive and EYFP negative, demonstrating the specificity of the integration events into the landing pad.

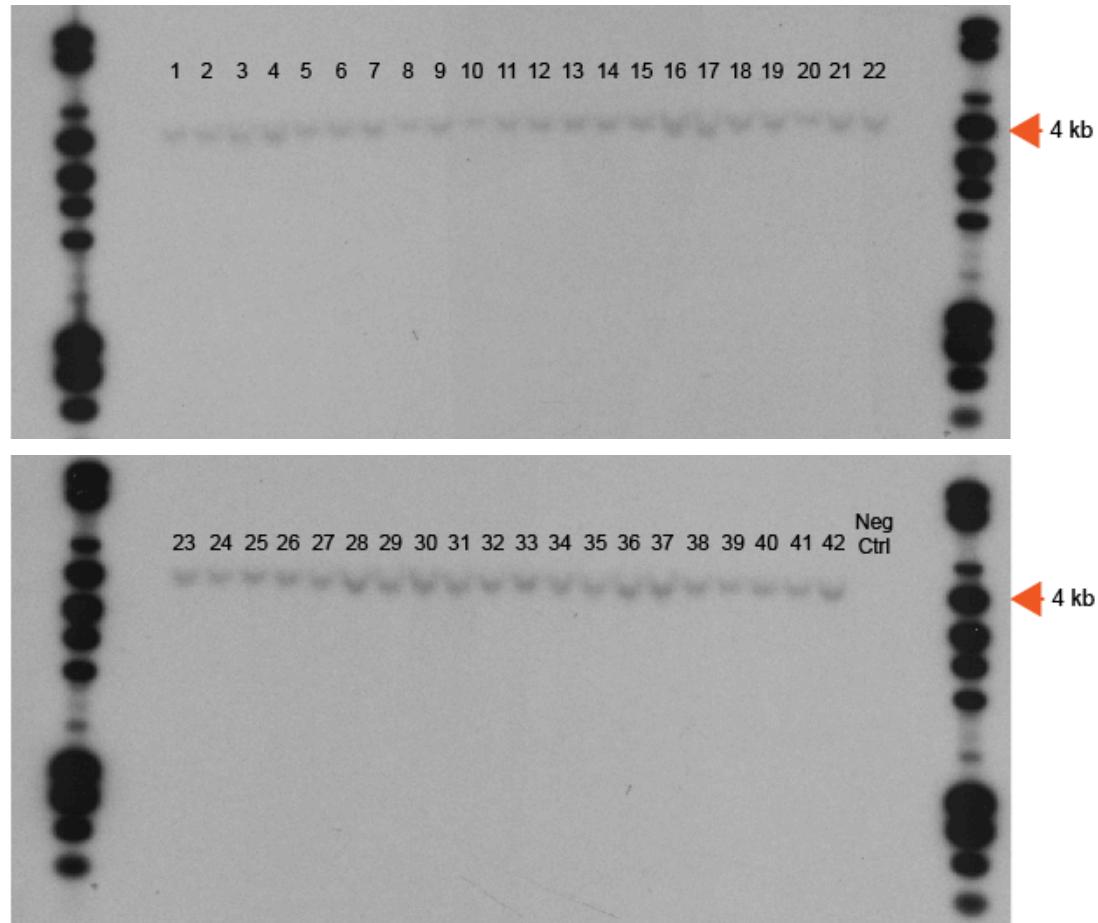


Supplementary Figure 10. Specificity of integration into the landing pad: Southern Blot genomic analysis assessing the specificity of BxB1 mediated integration.

A



B

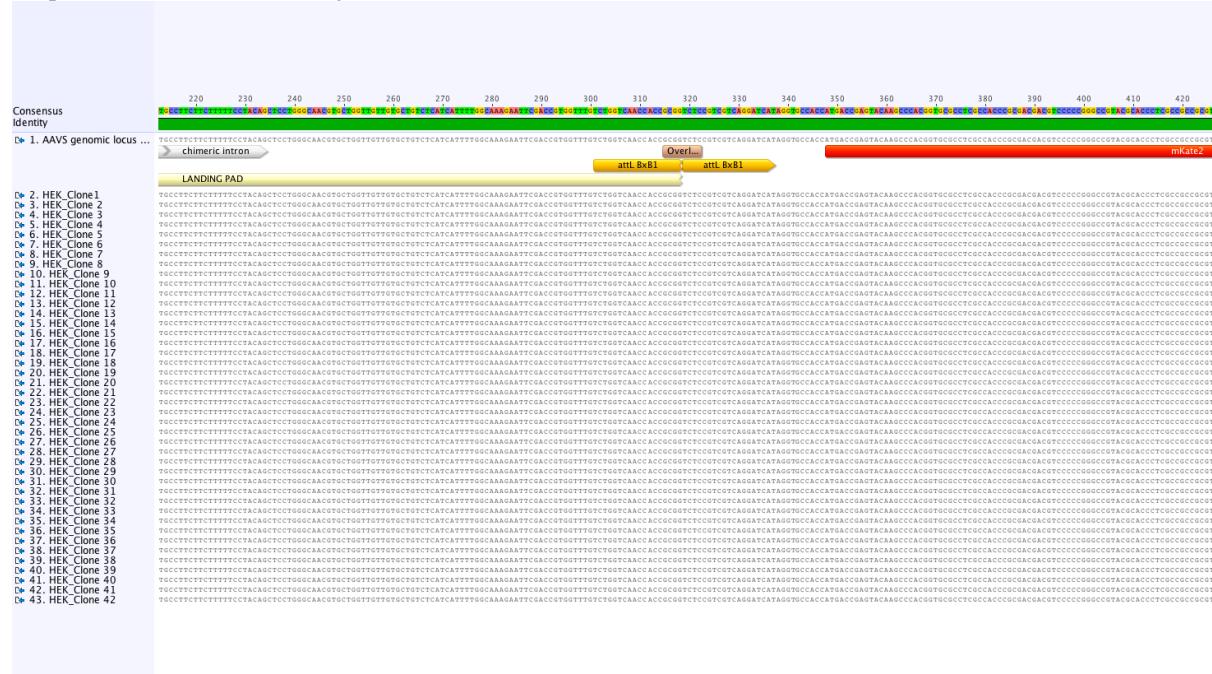


(A) Schematic of the Southern probe and the restriction enzyme cut sites. The 300 bp probe used for the Southern Blots corresponds to an internal sequence of the KanR gene from the integrated plasmid. Digestion of the chromosomal DNA with AseI results in a 4.3kb band containing the KanR gene if the circuit has been correctly integrated into the landing pad.

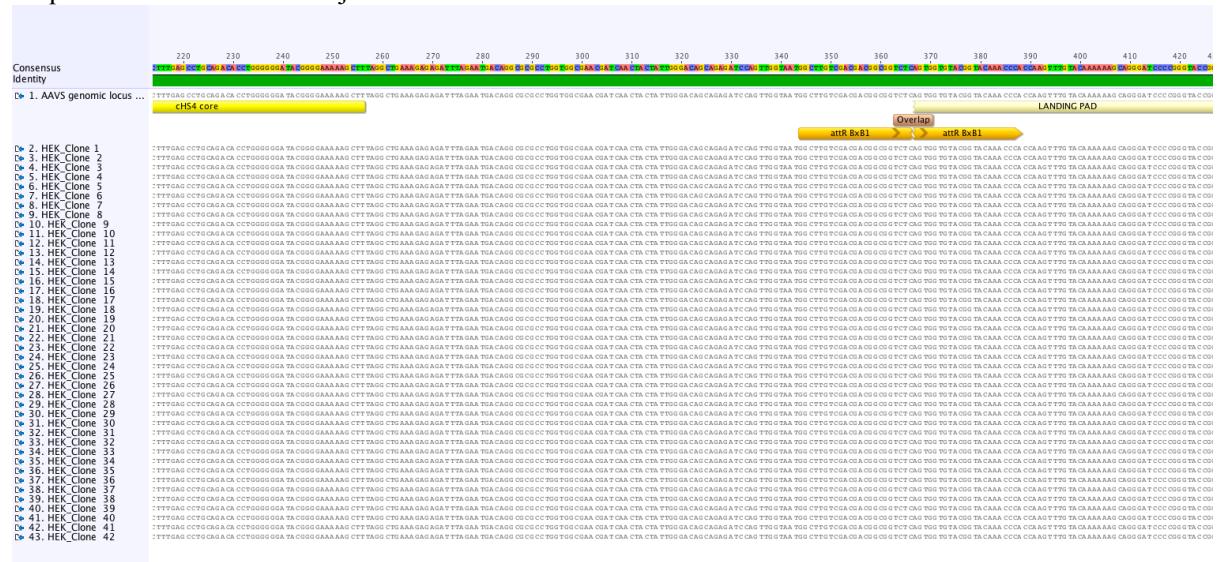
(B) Targeted integration of the integrative plasmid in the landing pad was confirmed for 42 randomly sorted clones that were expanded after selection with puromycin. For all the 42 clones, a single band of the expected size is revealed on the SB. Genomic DNA of the HEK293FT#8 monoclonal chassis cell line before payload integration is used as a negative control (last lane on the right).

Supplementary Figure 11. Sequence alignment of insert flanking sequences of the 42 isolated clones.

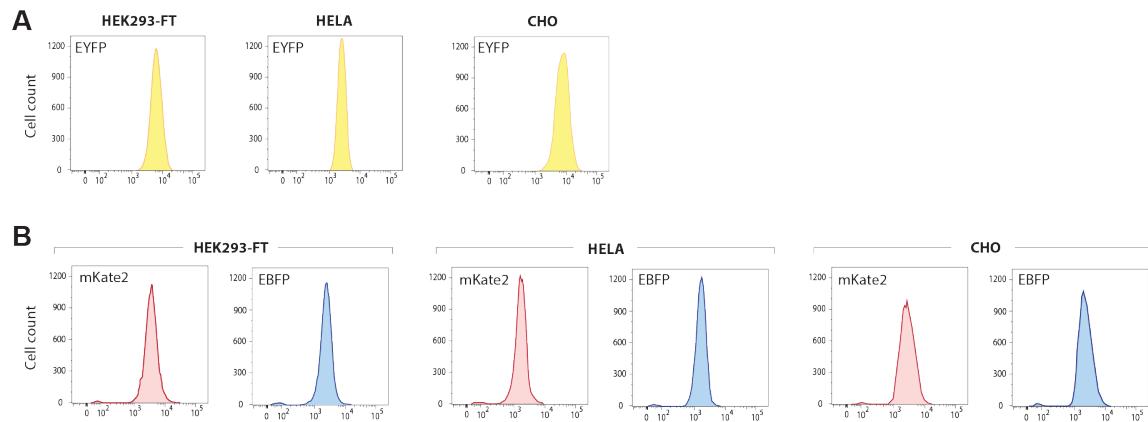
Sequences close to the attL junction



Sequences close to the attR junction

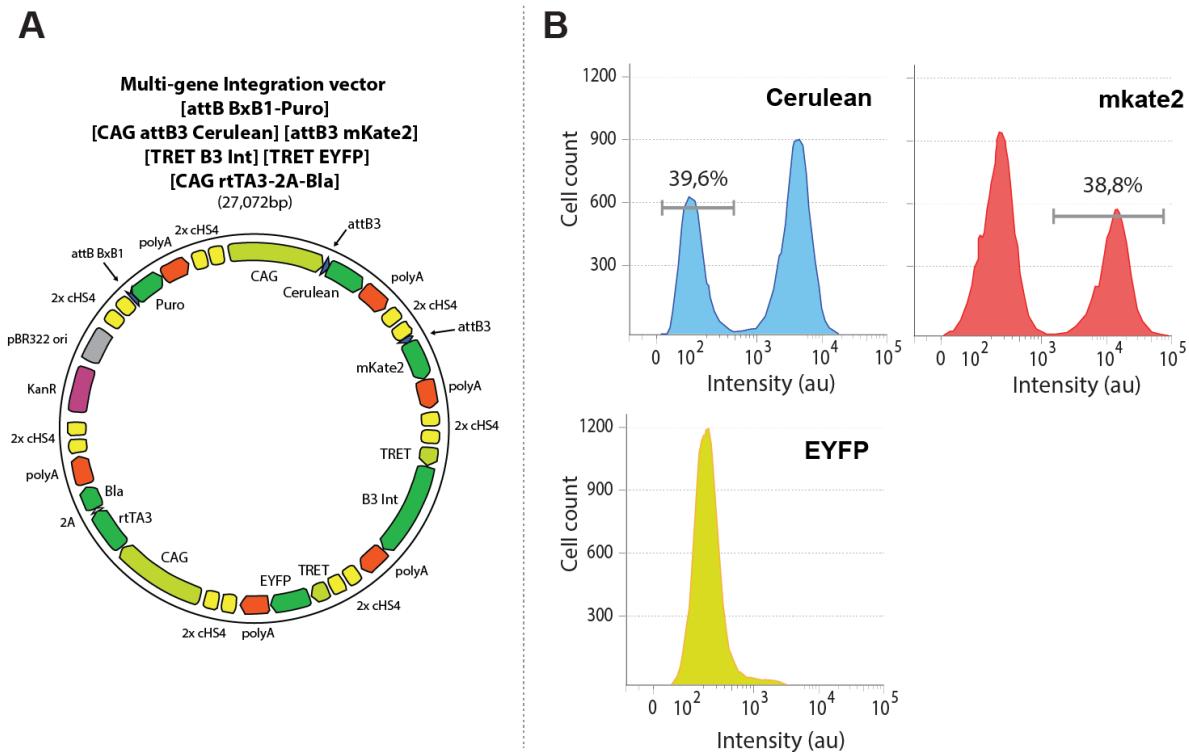


Supplementary Figure 12. Circuit performance in Hek293FT, HeLa and CHO chassis cell-lines.



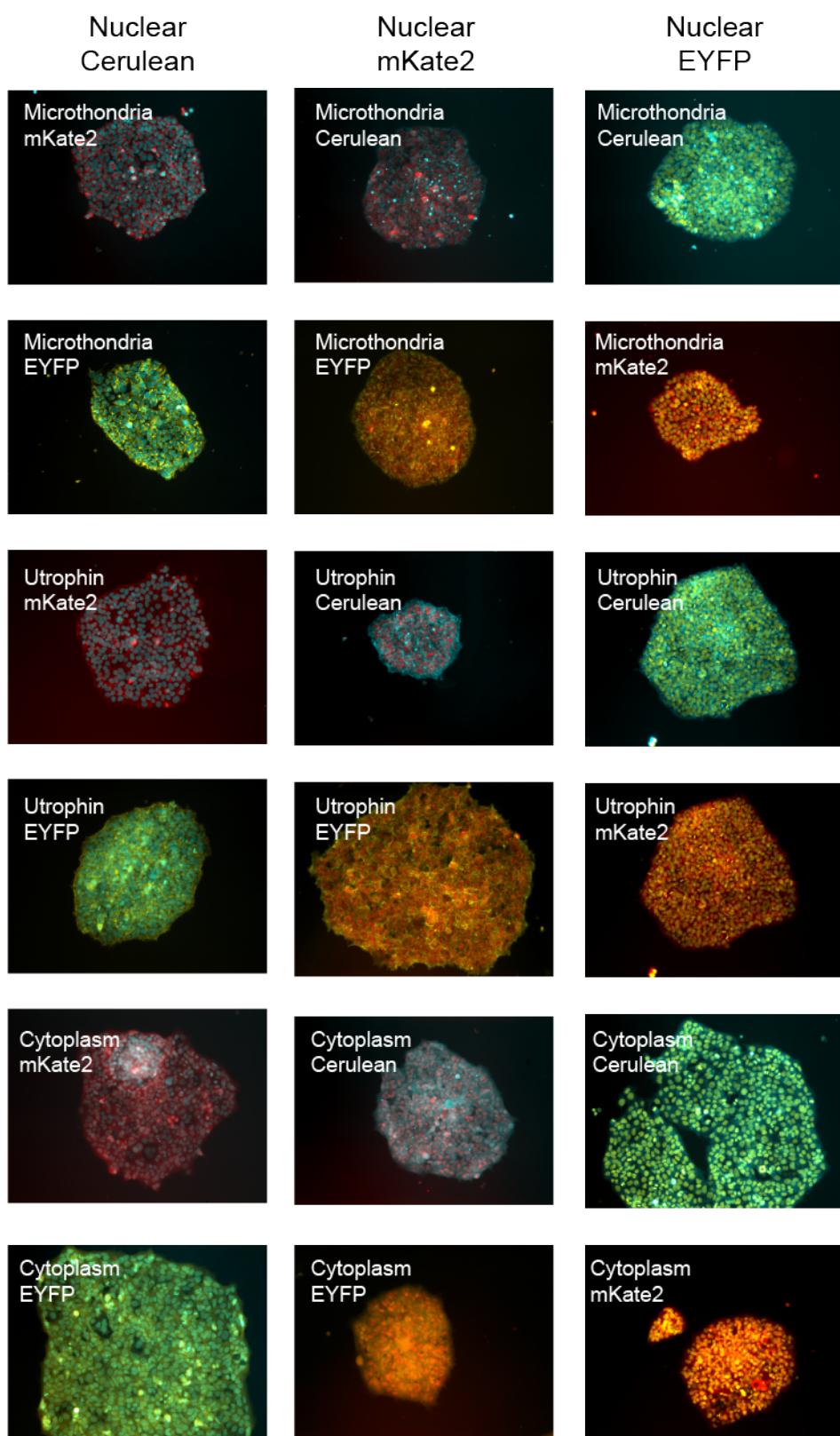
(A) Homogeneity of transgene (EYFP) expression in different landing-pad chassis cell lines. **(B)** FACS histograms of cells selected with puromycin after BxB1 mediated integration in different landing-pad cell lines.

Supplementary Figure 13. Early genomic rearrangement of the 7-gene payload following transfection.

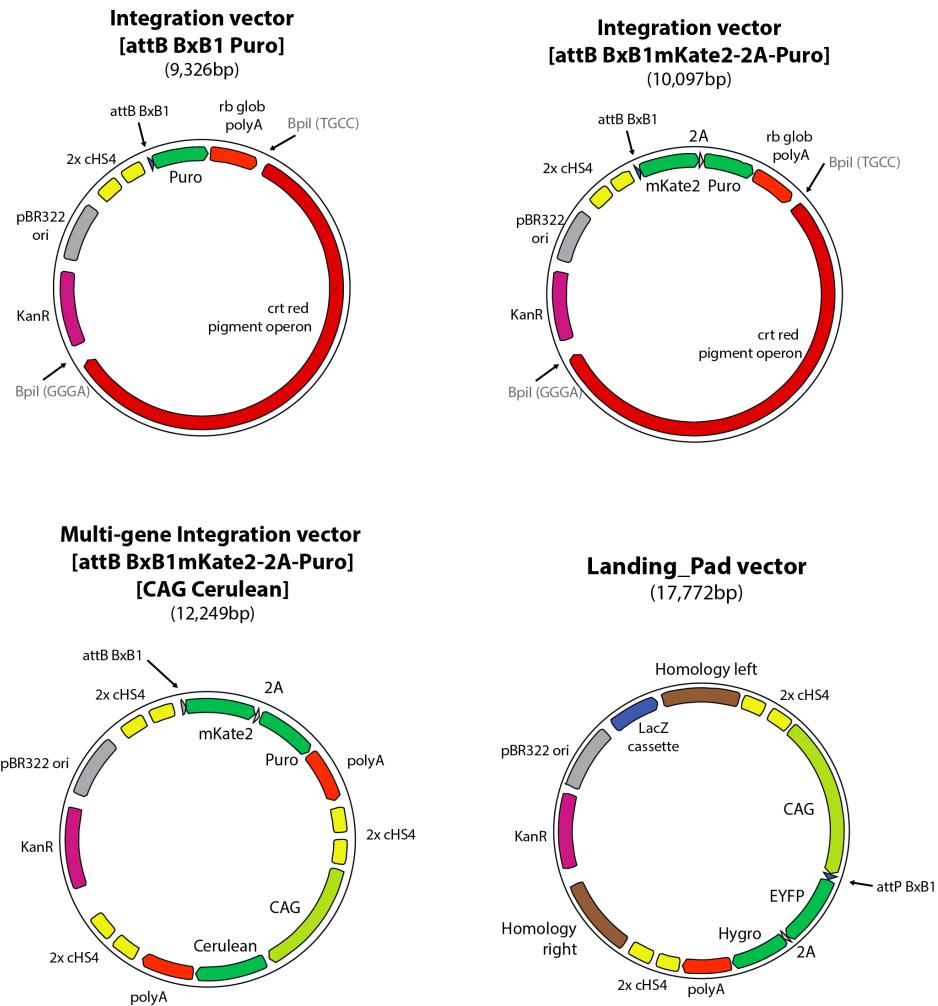


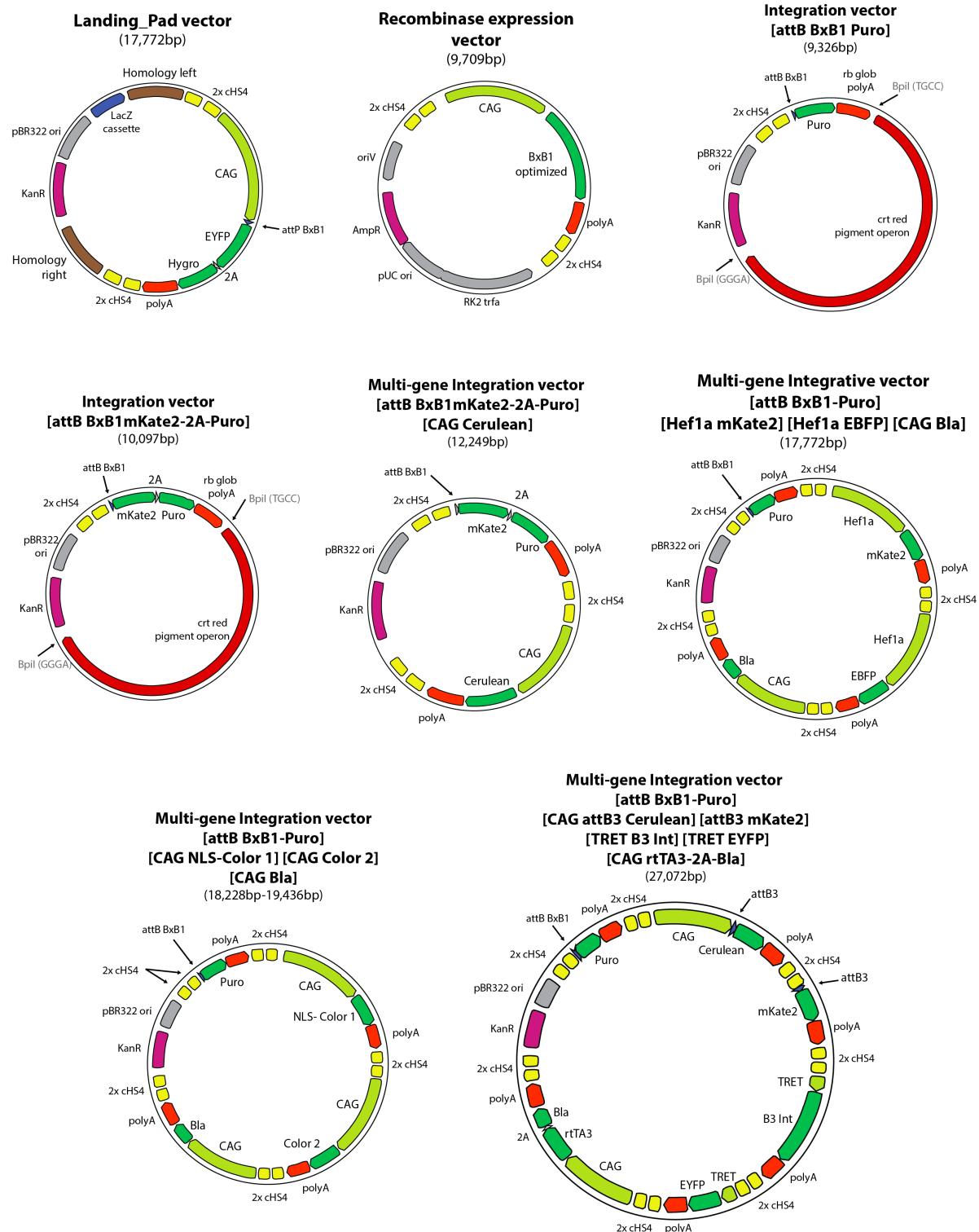
(A) Annotated map of the fully assembled circuit. **(B)** Representative FACS analysis histograms of resistant polyclonal population after integration of the 7-gene payload and selection with Puromycin for 7 days. About 40% of the cells have already excised the Cerulean transcription unit and switched to the final mKate2 expressing state. Given that essentially no cells are in the intermediate state, we hypothesized the switch occurred soon after transfection (complete degradation of the cerulean fluorescent protein takes up to a week). Leakiness of the TRE promoter from the transfected plasmids in the presence of high levels of rtTA3 (constitutively expressed) was sufficient to trigger expression of B3 integrase at low levels. Even in absence of dox, this resulted in moderate excision of the Cerulean expression cassette in the pool of plasmids. The experiment in Figure 3 is therefore performed with cells harboring a non-rearranged payload (red negative sorted cells).

Supplementary Figure 14. Microscopy images of isolated colonies corresponding to the 18 different phenotypes of the circuit library.



Supplementary Figure 15. Annotated maps of vectors used in this study.





Supplementary Table 1. Comparison of integration efficiencies of site-specific recombinases.

Recombinase	Integration efficiency	Cell type	Reference
BxB1	~1x10 ⁻¹	HEK293	This study
PhiC31	~2x10 ⁻³	HEK293	2
R4	~1x10 ⁻⁴	HEK293	3
Cre	~1x10 ⁻⁵	HEK293	4
Flp	~1x10 ⁻⁶	HEK293	5

Supplementary Text 1. DNA sequences of the probes used for the Southern Blots.

YFP (Landing Pad integration test) :

caaagaattcgaccgtggttgtctggtaaccaccgcggctcagtgggtacggtacaaaccaccaagttgtacaaaaaaaggcaggatc
cccggttacgggtgccaccatggtagcaaggcgaggagctgttcacccgggtggcccatcctgggtcgagctggacggcagctaa
acggccacaagttcagcgtgtccggcgagggcgagggcgatgcacccatcggcaagctgaccctgaagtcatctgcaccaccggcaagc
tgcccgtgcccggccaccctcgta

KanR (circuit integration test):

gggatcgcagtggtagtaaccatgcacatcaggagtagggataaaatgcgtatggcggaaaggcataaatccgtcagccagtttgt
ctgaccatctcatctgttaacatcatggcaacgcgtacccgtccatgttcagaaacaactctggccatcggtttccatataaagcgtataggatt
gtcgcacccgtattgtccgcacattatcgcgagccattataccatataatcagcatccatgttggaaatttaatcgcggcctcgacgttcccggt
aatatggctcat

Supplementary Text 2. DNA sequences of the primers used for genomic PCRs.

XD 621 Forward AAVS1 Primer: 5- GCCCTGGCCATTGTCACTT -3

XD 622 Forward AAVS1 CEL-I Primer: 5- TTCGGGTACCTCTCACTCC -3

XD 623 Reverse AAVS1 CEL-I Primer: 5- GGCTCCATCGTAAGCAAACC -3

LW06-LP-RV: 5- ACTCGAGAAATTGGAGCCAAC-3

Supplementary Text 3. DNA sequences of the genetic components from our mammalian part library used in this study.

Pars int p_Insulator:

p_Insulator [2xHS4 core fw]

CCTCGAGACAATTGATTAACATCGATAACGGTACCGAGTTGGCGCGCTGGGAGCTACGGGGACAGCCCCCCCCAAAGCCC
CCAGGGATGTAATTACGTCCCTCCCCGCTAGGGGGCAGCAGCGAGGCCCGGGCTCCGCTCCGGCGCTCCCCCCC
GCATCCCCGAGCCGGCAGCGTGGGGACAGCCCCGGCACGGGGAGGTGGCAGGGATCGCTTTCTGAACGCTCTC
GCTGCTTTGAGCCTGAGACACCTGGGGGATACGGGGAAAAGCTTAGGCTGAAAGAGAGATTAGAATGACAGGG
CGCCTGGCCATACATCGATAACGGTACCGAGTTGGCGCGCTGGGAGCTACGGGGACAGCCCCCCCCAAAGCCCCCAGGG
ATGTAATTACGTCCCTCCCCGCTAGGGGGCAGCGAGGCCGGGGCTCCGCTCCGGCGCTCCCCCGCATCC
CCGAGCCGGCAGCGTGGGGACAGCCCCGGCACGGGGAGGTGGCAGGGATCGCTTTCTGAACGCTCTCGCTGCTC
TTGAGCCTGAGACACCTGGGGGATACGGGGAAAAGCTTAGGCTGAAAGAGAGATTAGAATGACAGGGCGCCTGG
TGGCGAACGATCAAC

p_Insulator [2xHS4 core rev]

CCTCGAGACAATTGATTAACATCGATAACGGTACCGAGTTGGCGCGCTGGGAGCTACGGGGACAGCCCCCCCCAAAGCCC
CCAGGGATGTAATTACGTCCCTCCCCGCTAGGGGGCAGCAGCGAGGCCCGGGCTCCGCTCCGGCGCTCCCCCCC
GCATCCCCGAGCCGGCAGCGTGGGGACAGCCCCGGCACGGGGAGGTGGCAGGGATCGCTTTCTGAACGCTCTC
GCTGCTTTGAGCCTGAGACACCTGGGGGATACGGGGAAAAGCTTAGGCTGAAAGAGAGATTAGAATGACAGGG
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ATGTAATTACGTCCCCCCCCCGTAGGGGCAGCAGCGAGCCGGGGCTCGCTCCGGCCTCCCCCGCATCC
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TTGAGGCTGCAGACACCTGGGGGATAACGGGAAAAAGCTTGGCTAAAGAGAGATTAGAATGACAGGCGCGCTGG
TGGCGAACGATCAAC

p_Insulator [inert sequence]

CCTCGAGACAATTGATTAACATCGATACGGTACCGAGTTGGCGCCTGGGAGCTCACGGGACAGCCCCCCCCAAAGCCC
CCAGGGATGTAATTACGTCCCTCCCCGCTAGGGGGCAGCAGCGAGCCGCCGGCTCCGCTCCGGTCCGGCCTCCCC
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TGGCGAACGATCAAC

Promoters:

Ubc

TTGCCCTTGTAGCTTAAGTGTACTGGCTCCCGCGCCGGTTTGCGCCTCCCGCGGGGCCCTCCCTACGGCGAGCG
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GTAATCATTTGGGTCATATGTAATTTCAGTGTAGACTGTAATTGTCCGCTAAATTCTGGCGTTTGGCTTTGGTT
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Hef1a

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CAAAGTTTTCTTCAATTCAAGTGGATCCAAGGG

CAG

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TGGGGGGGGTGAAGCAGGGGGTGTGGCGCGGGCTGGCTGTAACCCCCCCCCCTGCACCCCCCTCCCCAGTTGTCAGCACG
GCCCGGCTTCGGGTGCCGGCTCCGTCGGGGCGGGCTCGCGCCGCCGCCGGGGTGCCTGCGGGGGGGGGGGGGGG
TGCGGGGGGGGGGGGGGGGCCCTCGGGGGGGGGAGGGCTCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
GAGGCCGCGCGAGCCGAGCCATTGCTTTATGGTAATCGTGCAGAGGGCGCAGGGACTTCCCTTGTCCCAAATCTGGCG
GAGCCGAAATCTGGAGGGCGCCGCACCCCTCTAGCGGGCGGGCGAGCGGTGCGGCCGGCAGGAAGGAAATG
GGCGGGGGAGGGCCTCGTGCCTGCCGCCGCCGTCCCCCTCTCATCTCCAGCCTGGGCTGCCGAGGGGGACGGCTG

CCTCGGGGGGGACGGGGCAGGGCGGGGTTGGCTTCTGGCGTGTGACCGGCGGCTAGAGCCTCTGCTAACCATGTTCAT
GCCTCTTCTTCCCTACAGCTCCTGGCAACGTGCTGGTTGCTCATCATTTGGAAAGAATTCTGTAC

TRE-tight:

ACTAATTGCCCTTCAGGTCCGCTAGACGAGTTACTCCCTATCAGTGATAGAGAACGATGTCGAGTTACTCCCTATCAGTG
ATAGAGAACGATGTCGAGTTACTCCCTATCAGTGATAGAGAACGATGTCGAGTTACTCCCTATCAGTGATAGAGAACG
TATGTCGAGTTACTCCCTATCAGTGATAGAGAACGATGTCGAGTTACTCCCTATCAGTGATAGAGAACGATGTCGAGGT
GGCGTGTACGGTGGGAGGCATAATAAGCAGAGCTCGTTAGTGAACCGTCAGATCGAAAGGGCAATTGACCCAAGTT
GTACAATCACTCTCCTGGTAC

Inert 3' UTR

CCACTGGATTGTACAATTAC

Genes (with Kozak sequence):

Bx_{B1} wild-type recombinase

GCGGCCACCATGAGAGCCTGGTAGTCATCCGCTGTCCCGCGTCACCGATGCTACGACTTCACCGGAGCGTCAGCTGGAGT
CTTGCAGCAGCTCGCAGCAGCGGGCTGGACGTCGTCGGGAGCGGAGGATCTGGACGTCGTCGGGCGGTGATCC
GTTGACCGGAAGCGCAGACCGAACCTGGCCGGTGGCTAGCGTTGAGGAGCACCGTTCGACGTATCGTGGCGTACCG
GGTAGACCGTTGACCGATCGATCCGATCTGCAGCAGCTGGTCCACTGGGCCAGGACCACAAGAACGCTGGTGTCTCC
GCGACCGAACGCACTTCGATACGACGACGCCCTTGCGGCGTGTGAGGAGCACCGTTCGACGTATGGAACGGTGGCGCAGATGGAAT
TAGAAGCGATCAAAGAGCGAACCGTTGGCTGCAATTCAATATCCGCGCCGGAAATACCGAGGATCCCTGCCGCGT
GGGATACCTGCCCTACGCGCTGGACGGGAGTGGCGCTGTGCGGACCCCTGTGAGCAGCAGAGCGCATCTCGAGGTGTA
TCACCGCTGTCGACAACCAGCGGCCCTGACCTGGTGGCCACGACTGGGAGTGGCTGCGGCTGAGCAGAGCGCATCTCGAGGTGTA
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GCGATGCTGGGACTCGCAAGGTAAGACCGTCCGAGACGACGGAGCCCCGCTGGTGCAGGCGATC
CTGACCCGTGAGCAGCTGGAGGCGCTGCGCGTGTGAGCTGCAAGACCTCCGGCGAAGCCCCGCGTGTACCCCGTGC
TGCTGTCGCGGGTTGTTCTGCGCGGTGTGCGGGAGCCCGTGTACAAGTTCGCGGGGGAGGACGTAAGCACCGCGCTA
CCGCTGCCGCTGATGGGTTCCGAAAGCACTGCGGAACGGCACGGTGGCAGTGGCAGTGGGAGCGCTTCGAGGA
GCAGGTGCTGGATCTGCTGGGAGCGGGAGCTGGTGGACCTGACGTCGCTGATCGCTCCCGCCCTACGGGCCGCTCCGAGCAGGAG
GGAGGTGAACCGGGAGCTGGTGGACCTGACGTCGCTGATCGCTCCCGCCCTACGGGCCGCTCCGAGCAGGAG
ACTGGATGCCGTATTGCGCGCTGGCGCGCAGAGGAGACTGGAGGGTCTAGAGGCTGCCGCTGGGAG
GGCGAGACTGGCAGGGTTGGGAGCTGGTGGCGGAGCAGGACACCGCGAAAGAACACCTGGCTCGGTGATGAA
CGTTGGCTGACGTCACGTCACGTCACACACCGGGATGTC
AGGCTCGGCAGCGTGGTCGAACGGTACACACCGGGATGTC
AG

Bx_{B1} mammalian codon optimized recombinase

GCGGCCACCATGAGAGCACTGGTAGTCATCCGACTGAGTAGGGTCACAGACGCAACAACAAGCCCCGAGAGGGCAGCTGGAA
TCATGTCAGCAGCTGTGCGCACAGCGAGGATGGACGTCGAGTGGCAGAGGATCTGGACGTCGAGCGCGCTGTCGAT
CCATTGACAGAAAGCGGAGGCCAACCTGGCAAGGTGGCTTCTGAGGAACAGCCCTTGATGTGATCGTCGCTACA
GAGTGGACAGGCTGACACGCTTATTGACATCTGCACTGGTGCATTGGGCCAGGACCAAGAACACTGGTGTCA
GTGCAACTGAAAGCCACTTCGATACCAACTCCCTTGCGCTGTGTCAGTCAGTGTGGCACCCTGGCCAGATGG
GCTGGAAAGCTATCAAGGAGCGAAACGGAGCTGGCAGCCCATTCAATATTGGGGGGGGAAATACAGAGGATCACTGGCCCC
TTGGGGCTATTCGCTGGGAGCTGGGAGTGGAGACTGGTGGCAGACCCCTCCAGAGGAGAGGAGGATCTGGAAAGT
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AGATTATTGCTCAGCTGAGGGAGGCCACAGGGACGAGAACCTCCAGAGCAACCTGGCTGAGCAGGAGCATGTCAG
TGAGGCTATGCTGGCTACGCAACTCTGAATGGAAAACCGTCCGGAGCAGTGGCAGGAGCACCAACTGGTGGCTGAGGCC
TATTCTGACACCGCAGCTGGAGCTCTGCGGGCAGAACACTGGTGGAAAACCTCCAGAGCAACCTGGCTGAGCACCC
AAGCCTGCTGAGGGTGTGTTCTGCGCCCTGTGGGGAGCCAGCATAAGTTCGCGGGGGAAAGAAAACATCCC
CGCTATCGATGCCGTATGGGATCCCTAAGGACTGTGGAAACGGCACTGTGGCATGGCCAGTGGGAGCCTTGTG
AGGAACAGGTGCTGGATCTGGGAGCGGGAGAGGCTGGAAAAAGTGTGGCTGCTGGCAGCAGCTCGCTGGAG
TGGCAGAAGCTACGTCAGCTGGGAGCTGGGAGACGCGAGGGCTGGGAGTGGCTCAGACAGCAG
AGCTCTGGACGACGCAATTGCTGCACTGGCAGCTGACAGGAGGAACCTGGAGGGGCTGGAGACGACCTAGCGGATGGG
GTGGCGAGAACACAGGCCAGCGGTTGGGAGTGGAGAGGAGCAGGACACAGCAGCAAGAACACTTGCTGAGAAGTA
TGAATGTCAGGCTGACTTCGATGTCGCGGGGGCTGACCCGAACAATCGATTGGCGACCTGCACTGGGAGTATGAACAGCA
CCTGAGACTGGGAGCGTGGTCGAAGACTGCACACTGGGATGTCAGTAA

Bx_{B1} mammalian codon optimized recombinase with N-terminal NLS

GCGGCCACCATGCGAAAGAACGCAAAGTTAGAGCACTGGTAGTCATCCGACTGAGTAGGGTCACAGACGCAACAACA
AGCCCCGAGAGGCAGCTGGAAATCATGTCAGCAGCTGTGCGCACAGCGAGGATGGACGTCGAGTGGCAGAGGATCTG
GACGTGAGCGCGCTGTCGATCCATTGACAGAAAAGCGGAGGCCAACCTGGCAAGGTGGCTTCTGAGGAACAGCCC
TTGATGTCATGTCGCTACAGAGTGGACAGGCTGACACGCTCTATTGACATCTGCACTGGCAGCTGGTGCATTGGGCCAGG
ACCACAAGAACACTGGTGGCTGCAACTGAAAGCCACTTCGATACCAACTCCCTTGCGCTGTGGTCACTGCACTGAT
GGGACCGTGGCCAGATGGAGCTGGAGCTATCAAGGAGCGAAACGGAGCTGGCAGCCATTTCGATATTGGGCCAG
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CCACTGGTGGAGGGCTGAGCCTATTCTGACACCGCAGCTGGAGCTGGCAGAAGCTCTGCGGGCAGAACTGGTGGAAAACCTCCAGAGGCC
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CAGCGACTCCGCTGAGCTGGCAGAAGTCATGCCAGCTGGTGGATCTGACCTCCCTGATCGGATCTCTGATATAGG
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GCACGACCTAGCGGATGGGAGTGGCGAGAACACAGGCCAGCGGTTGGGAGTGGAGAGGAGCAGGACACAGCAGCCAA

GAACACTTGGCTGAGAAGTATGTCAGGCTGACTTTGATGTGCGCGCGGGCTGACCCGAACAATCGATTTGGCGAC
CTGCAGGAGTATGAACAGCACCTGAGAGACTGGGAGCGTGTGAAAGACTGACACTGGGATGTCATAG

BxB1 mammalian codon optimized recombinase with C-terminal NLS

GCCGCCACCATGAGAGCACTGGTGGTCATCCGACTGAGTAGGGTCACAGACGCAACAACAAGCCCCGAGAGGCAGCTGGAA
TCATGTCAGCAGCTGCGCACAGCGAGGATGGGACGTGGCAGAGGATCTGGACGTGAGCGGCCGTGCGAT
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GAGTGGACAGCGTGTGACACGCTATTGACATCGCAGCAGCTGGTCATTGGGCCAGGACCACAGAAACTGGTGGTCA
GTGCAACTGAAGGCCACTCGATACCAACTCTTGGCCGTGTGTCATCGCAGCTGATGGGACCGTGGCCAGATGG
GCTGGAAGCTATCAAGGAGCGAAACCGGAGTGCAAGCCCATTCAATATTGGGCCGGAAATACAGAGGATCACTGCC
TTGGGCTATCTGCTACCCGGGTGGATGGGGAGTGGAGACTGGTGCAGACCCCGTCCAGAGAGAGAGGATTCTGGAAGT
GTACACAGGGTGGTCGATAACCACGAACCACACTGCATCTGGTCGCCACGACCTGAATAGGCGCGCTGCTGAGGCCAAA
AGATTATTTGCTCAGCTGAGGGAGGCCACAGGGACGAGAATGGTCCGTAACCGCCTGAAGCGGAGCATGTCAG
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TATTCTGACACCGCAGCAGCTGGAAAGCTCTGCGGGCAGAACACTGGTGAAGAACCTCCAGAGGCCAACCTGCCGTGAGCACCC
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AGCTCTGGACGACGAATTGCTGCACTGCCAGCTGACAGGAGGAACGGAGGGCTGGAAGCACGACCTAGCGGATGGG
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TGAATGTCAGGCTACTTCGATGTGCGCGCCGGCTGACCCGAACAATCGATTTGGCGACCTGCAAGGAGTATGAACAGCA
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B3 Recombinase

GCCGCCACCATGAGCTGTATATGGATCTTGTGATGATGAACCGCAGCTTGTACCAATAAGTCGTGGAGTGTGAAAG
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TGTGTATCACTTGAAGGACTGTGATGAGCTGTCAGGGGCTTGAGCGATGCTTCGAGCCCTACAAATTCAAATTAGTCG
AATAAGAGGCAACCTCGTTAACACTCTCTTGTGCCCCCTGTTGGCAGCCAGAAGAACACTGGGAAAGAAAGAGGTGG
ACCGCAGGTGGATAACTTGTTCATAGCACCGAGACAGACGAATCGATTTCAAATTCATCTGAACACGTTGGATAGTAT
TGAGACACAAACGAACACGGATGCCAAAGACGGTGTGACTTTCATCTGTTGATGACATTTCACACTGCTGTAGGAAC
AATGACCTGATGAACGTTGATCCCTCACATTAAAGATTGTAAAAACAAATTCTGCGGATACCTGCTGCAAGGCTGAGGTCA
AACAGACTAACAGCAGCAAGTCGAGGAACATTCTCTTCCCACATCCCGAGAATCGATTGCGATCTGTTCTGGCCTTGACG
ATTCTCCCGCACATGCCAGCCATACCCAAAGTCGCGCTTCCGGATCAAGTATCGGAGCAGAAGTGGCAGCTTCCGAGAT
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TCGGGCCGCATCTGATGAAACAGCTTCTGACAAGAATGAACTGGATTCTGGCCAACCTCCCTGGCACTGGGACTGTC
TCCCAGAATCAACCGCAGTCCGGTGCAGCTGGCTACACCCACGGTGGTCGAGATTGCAAGGCCACTGTCGGATTCC
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TGGTCTGATATTCTGAGCGATATCTGAAAGAACGCATGCCGCATATCAGAACACTCCACATTGACGTACGTTCC
TTAACAGACAGTCCGTACCTTGAGGGCTCAAGGGAGCGAACAGGCCAGCAGTCCGCTCAGGATTCCAATACTTAGTAT
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CGTAAACGTGGCGTGGGTGGTCGCCGACCGTTGAGTCGAAGCGCGCGTCTGGCACCAATTACCGGTGA

EYFP

GCCGCCACCATGGTGAGCAAGGGCGAGGAGCTGTTCACGGGGGTGGTGCCCACCTGGTCAGCTGGACGGCGACGTAAC
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TACAAGGCCGCCGCCAGGTGAAGTTGAGGGCGACACCCCTGGTAACCGCATCGAGCTGAAGGGCATCGACTCAAGGAG
GACGGCAACATCCTGGGCCAACAGCTGGAGTACAACACAGCCACACGCTATCATGCCGACAAGCAGAAC
GGCATCAAGGTGAACCTCAAGATCCGCCAACACATCGAGGAGCGCAGCGTGCAGCTGCCGACCACTACCGAGAACACC
CCCATCGGCCAGGGCCCCGTGCTGCTGCCGACAAACACTACCTGAGCTACGCTCAAGCTGAGCAAAGACCCCAACGAG
AAGCGCGATCACATGGTCTGCTGGAGTTGTCGACCGCCGGGATCACTCTGGCATGGACGAGCTGACAAGTGA

mKate2

GCCGCCACCATGGTGAGCAAGGGCGAGGAGCTGTTCACGGGGGTGGTGCCCACCTGGTCAGCTGGACGGCGACGTAAC
TTCAAGTGCACATCCGAGGGCGAAGGCAAGCCCTACGAGGGCACCCAGACCATGAGAAATCAAGGCCGTGAGGGCGGCC
CTCCCTTCGCCCTCGACATCTGGCTACCAGCTTGTACGGCAGAAAACCTTGTACCAACCACACCAGGGCATCCCCGA
CTTCTTAAGCAGTCCCTCCAGGGCTCAGGAGCTGCTCATACACGCTCAAGGAGTACAGAGGGGCTGACCTGACCGCTACC
CAGGACACCGCCTCAGGAGCGCTGCCATCACACGCTCAAGGAGTACAGAGGGGCTGACCTCCATCCAACGCCCTGTGA
TGCAGAAGAAAACACTCGGCTGGAGGCCACCTGAGAGACTGTACCTGGGCTGACGGCGGCCCTGGAAGGCAGAGGCC
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CAAGATGCCGCGTCAACTATGTGAGAGGAGACTGAGAAAGAATCAAGGAGGCCACAAAGAGACATACGTCGAGCAG
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EBFP

GCCGCCACCATGGTGAGCAAGGGCGAGGAGCTGTTCACGGGGGTGGTGCCCACCTGGTCAGCTGGACGGCGACGTAAC
GGCCACAAGTCAGCGTGGGGCGAGGGCGAGGGCGATGCCACCAACGGCAAGCTGACCTGAAGTTCATCTGCACCA
GGCAAGCTGCCGTGCCCTGGCCACCCCTCGTACCGACCCCTGAGCCACGGCGTGCAGTGCCTGCCGCTACCCGACCA
TGAAGCAGCACGACTCTCAAGTCCGCATGCCGAAGGCTACGTCAGGAGCGCACCCTCTCAAGGACGACGACGCC

CTACAAGACCCGCGCCGAGGTGAAGTTCGAGGGCGACACCCTAGTGAACCGCATCGAGCTGAAGGGCGTCGACTTCAGGA
GGACGGCAACATCCTGGGCACAAGCTGGAGTACAACCTCACAGCCACAACATCTATATCATGGCCGTCAAGCAGAAGAA
CGGCATCAAGGTGAACTCAAGATCCGCCACAACGTTGGAGGACGGCAGCGTCACTGCCCCGACACCAC
CCCCATCGGCACGGCCCGTGTGCTGCCGACAGCCACTACCTGAGCACCCAGTCCGTGAGCAAAGACCCCAACGAG
AAGCGCGATCACATGGTCTGTGGAGTTCCGACCGCCGGGATCACTCTCGCATGGACGAGCTGTACAAGTGA

Cerulean

GCGGCCACCATGGTGAGCAAGGGCAGGAGCTGTTCACCGGGGTGGTCCCACCTGGTCAGCTGGACGGCAGCTAAAC
GGCCACAAGTTCAGCGTGTCCGGCAGGGCGAGGGCGATGCCACCTACGGCAAGCTGACCTGAAGTTCATCTGCACCACC
GGCAAGCTGCCGTGCCCTGGCCACCCCTGTGACCACCCGTACCTGGCGTCACTGCTTCGCCCCGTACCCGACCA
TGAAGCAGCACGACTTCAAGTCCGCCATGCCGAAGGCTACGCCAGGGCAGCGCACCATCTTCAGGACGACGGCAA
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GGACGGCAACATCCTGGGCACAAGCTGGAGTACAACGCCATCGCGACAACGCTATATCACCGCCGACAAGCAGAAGAA
CGGCATCAAGGCCAACTCAAGATCCGCCACAACATCGAGGACGGCAGCGTCACTGCCCCGACCACTACCGAGAAC
CCCCATCGGCACGGCCCGTGTGCCGACAACCAACTACCTGAGCACCCAGTCAAGCTGAGCAAAGACCCCAACGA
GAAGCGCGATCACATGGTCTGTGGAGTTCTGTGACCGCCGGGATCACTCTCGCATGGACGAGCTGTACAAGTGA

mKate-2A-Puro

GCGGCCACCATGGTGAGCGAGCTGATTAGGAGAACATGCACATGAAGCTGTACATGGAGGGACCGTGAACAACCAC
TTCAAGTGCACATCCGAGGGCGAAGGCAAGGCCCTACGGGACCCAGACCATGAGAAATCAAGGCGGTGAGGGCGGCC
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TGCAGAAGAAAACACTCGGCTGGAGGCCACCGAGACACTGTACCCGTCAGGGCGCTGGAGGGCAGAGCCGACA
TGGCCCTGAAGCTGTGGCGGGGCCACCTGATCTGCAACCTTAAGGACACATACAGATCCAAGAAACCCGCTAAGAAC
CAAGATGCCCGCCTACTATGTGGACAGGAGACTGGAAGAAAATCAAGGAGGCCACAAAGAGACATACGTCGAGCAGC
ACGAGGTGGCTGTGGCCAGATACTGCGACCTCCCTAGCAAACGGGGCACAACACTTAAATGGAAGCAGGGAGCTAACTTC
GCCTGTCAGCAGGCTGGCAGCTGGAGGAGAACCTGGACCTACCATGACCGAGTACAAGCCCACGGTGCCTCGCCA
CCCGCGACGACGTCCCCGGCGTACGCACCCCGCCGTCGGCGACTACCCGCCACCGCGCACCCGTCGACCC
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CGCATGGCCGAGTTGAGCGTTCCCGTGGCGTCTGCCGACCAACAGATGGAAGGCCCTCTGCCGCGCACCCGCCAAGGAG
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EYFP-2A-Hygro

GCGGCCACCATGGTGAGCAAGGGCAGGAGCTGTTCACCGGGGTGGTCCCACCTGGTCAGCTGGACGGCAGCTAAAC
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GGCAAGCTGCCGTGCCCTGGCCACCCCTGTGACCCACCTCGGCTACGGCTGAGTGTCTGCCGCTACCCGACCA
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AAGCGCGATCACATGGTCTGTGGAGTTCTGTGACCGCCGGGGATCACTCGCATGGACGAGCTGTACAAGGGAGC
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GCATCCGGAGCTTGCGAGGATGCCGCGCTCCGGCGTATATGCTGCCATTGGTCTGACCAACTCTACAGAGCTGG
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CACTCGCCGATGGGAGATGGGGAGGCTACTGA

N-ter H2B localization tag

GCCACCATGCCAGGCCAGCGAAGTCTGCTCCGCCCGAAAAAGGGCTCAAGAACGGCGTGACTAAGGCCAGAAC
GGCGGCAAGAACGCCAGGCCAGCGCAAGGAGAGCTATCCATCTATGTGACAAGGTTCTGAAGCAGGTCCACCC
ACCGGCATTCCTGCAAGGCCATGGCATCATGAATTGTTGTGAACGACATTTGAGCGCATCGCAGGTGAGGCT
GCCCTGGCGCATACAAACAGCGCTGCCGACCCATCTGCCAGGGAGATCCAGACGCCGTCGCCCTGCTGCC
GGCAAGCACGCCGTGCCAGGGTACTAAGGCCATACCAAGTACCCAGCGCTAAG + linker:
GATCCCCGGGTACCGCGCCACC

N-ter Smac mitochondria localization tag

GCGGCCACCATGGCGGCTCTGAAAGAGTTGGCTGCGCAGCGTAACCTCATTCTCAGGTACAGACAGTGTGTTGTT
TGGTGTGGCTAACCTTAAGAAGCGGTGTTCTCAGAATTGATAAGACCATGGCACAAAAGTGTGACGATTGGCT
ACCCCTGTGT + linker: GGATCAGGAGGATCAGGA

C-ter Utrophin localization tag

Linker: GGATCAGGAGGATCAGGA +

TCCGGAAACCATGGCCAAGTATGGAGAACATGAAGCCAGTCCTGACAATGGGCAGAACGAATTCACTGATATCATTAAAGTCC AGATCTGATGAACACAATGACGTACAGAAGAAAACCTTACCAAATGGATAATGCTGATTTCAAAGAGTGGAAACCA CCCATCAATGATATGTTCACAGACCTCAAAGATGGAAGGAAGCTATTGGATCTCTAGAAGGCCTCACAGGAACATCACTGC CAAAGGAACCGTGGTCCACAAGGGTACATGCCCTAAATAACGTCAACAGACTGCTGCAGGTTTACATCAGAACAAATGTGGA ATTAGTGAATATAGGGGAAACGGACATTGTGGATGGAAATCACAAACTGACTTTGGGGTACTTGGAGGCATCATTTGCAC TGCGAGGTGAAAGATGTCATGAAGGATGTCATGTCGGACCTGCAGCACAGAACAGTGAGAACATGCTGCTCAGCTGGGTG CGTCAGACCACCAAGGCCCTACAGCAAGTCAACGTCCCTCAACTTCACCAACAGCTGGACAGATGGACTCGCCCTTAATGCTG TCCTCCACCGACATAAACCTGATCTCTCAGCTGGATAAAGTTGTCAAAATGTCACCAATTGAGAGACTTGAACATGCCCTC AGCAAGGCTCAAACCTATTGGGAAATTGAAAAGCTGTTAGATCCTGAAGATGTTGCCCTCGGCTCCTGACAAGAAATCCA TAATTATGTATTAACATCTTGTGAGGTGCTACCTCAGCAAGTCACCATAGACTGA

Recombination sites:

attB BxB1

GGCTTGTGACGACGGCGGTCTCCGTCGTAGGATCAT

attP BxB1

GTGGTTGTGGTCAACCACCGCGGTCTCAGTGGTGTACGGTACAAACCCA

Inert 5' UTR

TAAGTTGTACAAAAAAGAG

Polyadenylation signal:

rb glob polyA

TGAATTCACTCCTCAGGTGCAGGCTGCCATCAGAAGGTGGTGGCTGGTGGCCAATGCCCTGGCTACAAATACCACTGA GATCTTTTCCCTCTGCCAAAAATTATGGGACATCATGAAGCCCCCTGAGCATCTGACTCTGGCTAATAAAGGAATTAT TTTCATTGCAATAGTGTGTTGAATTTTGTGTCCTCACTCGGAAGGACATATGGGAGGGCAATCATTAAAACATCAGA ATGAGTATTGGTTAGAGTTGGCAACATATGCCCATATGCTGGCTGCCATGAACAAAGGGTGGCTATAAAGAGGTATCA GTATATGAAACAGCCCCCTGTCCTCATTCCATAGAAAAGCTTGACTIONGAGGTAGATTTTTATATTGTTT TGTTGTTATTTCATACATCCAAAATTTCCTTACATGTTTACTAGCCAGATTTCCTCCTCCTGACTACTCCC GTCATAGCTGCCCTCTCTTATGGAGATCCCTCGAC

Supplementary Text 4. DNA sequences of all plasmids from the mMoclo library.

p_Insulator	GTGGCTCTTCAGTGGACGAAAGGGCCTCGTGATACGCCATTTCAGGTTAATGTCATGATAAT AATGGTTTCTTAGACGTCAAGTGGCACTTTCGGGAAATGTCGGCGAACCCCTATTGTTATT TTCTAAATACATTCAAATATGTATCCGCTCATGAGAACATAACCCCTGATAAATGCTTCATAAATAT TGAAAAGGAAGAGTATGCGCTCACGCAACTGGCCAGAACCTTGACCGAACGCGCGTGGTAA CGGCCAGTGGCGTTTCATGGCTTATGACTGTTTTGGGGTACAGTCTATGCCCTGGC ATCCAAGCAGCAAGCGCTTACGCCGTGGTCATGTTGATGTTATGGAGCAGAACGATGTTA CGCAGCAGGGCAGTCGCCCTAAACAAAGTTAACATCATGAGGGAAGCGGTGATGCCGAAGTA TCGACTCAACTATCAGAGGTAGTTGGCGTCATCGAGCGCCATCTGAAACCGACGTTGCTGCCGT ACATTGTACGGCTCCGCACTGGATGGCGGCTGAAGGCCACACAGCGATATTGATTGCTGGTTA CGGTGACCGTAAGGGCTTGATGAAACAACCGGGGAGCTTGACTIONGAGGTAGATTGCTGGAAACTTCG GCTTCCCTGGAGAGGGAGATTCTCGCGCTGAGAAGTCACCATTTGTCACGACCAT CATTCCGTGGCGTTATCCAGCTAACGCGAACGATCGACATTGATCTGGCTATCTGCTGACAAAAGCAAGAG TTGCAGGTATCTCGAGCCACGATCGACATTGATCTGGCTATCTGCTGACAAAAGCAAGAG AACATAGCGTTCCCTGGTAGGTCCAGCGGGAGGAACCTTTGATCCGGTCTGAACAGGAT CTATTGAGGCCTAAATGAAACCTTAAACGCTATGGAACCTGCCGCCCCACTGGCTGGCGATGA GCGAAATGTAGTGTACGTTGCTCCGCATTGGTACAGCGCAGTAACCGGAAATCGCGCCGA AGGATGTCGCTCCGACTGGCAATGGAGGCGCTGCCGGCCAGTATCAGCCCGTCATACTTGAA GCTAGACAGGCTTATCTGGACAAGAAGAATCGCTGGCTGCCGCGAGATCAGTTGGAAGA ATTGTCAGGCTTACGTAAGGGAGATCACCAAGGTAGTCCGCAATAACTGTCAGACCAAGTTA CTCATATATACTTGTAGTTAAAACCTTACATTAAATTAAAAGGATCTAGGTGAAGATCCTT TTTGATAATCTCATGACAAAATCCCTAACGTGAGTTTCGTTCACTGAGCGTCAGACCCCGTA GAAAAGATCAAAGGATCTCTTGAGATCCTTTCTGCGCGTAATCTGCTGCTGCAAACAAAA AAACCAACGCTTACAGCGGTGGTTGGCCGATCAAGAGCTACCAACTTTCCGAAGGTA ACTGGCTTCAGCAGAGCGCAGATAACAAACTGTCCTCTAGTGTAGCGTAGTTAGGCCACCAC TTCAAGAACTCTGTAGCACCGCTACATCACCTCGCTCTGCTAATCCTGTTACCGAGTGGCTGTC AGTGGCGATAACTCGTGTCTTACGGGGTGTGGACTCAAGACGATAGTTACCGGATAAGGGCGACGC GTCGGGCTGAACGGGGGGTCTGCGACACAGCCAGCTGGAGCGAACGACCTACACCGAACGATGA GATACTACAGCGTGAGCTATGAGAACAGCGCACGCTCCGAAGGGAGAAAGGGCGACAGGTAT CCGGTAAGCGGCAGGGTCTGGAACAGGAGAGCGCAGCAGAGGGAGCTCCAGGGGGAAACGCCCTGGT ATCTTATAGTCCTGCGGGTTGCCCCACCTCTGACTTGAGCGTCGATTTGTGATGCTCGTCAG GGGGCGGGAGCCATGGAAAAACGCCAGCAACCGCCCTTTTACGGTTCTGGCTTTGCTGG CCTTTGCTCACATGTTCTCTGCGTTATCCCTGATTCCTGTTGATAACCGTATACCGCCCTTG AGTGAAGCTGATAACCGCTCCGCAAGCCGAACGACCGAGGCGAGCGAGTCAGTGAAGCGAGGAAGC GGAAGAGGCCCAACGCAAACCGCCCTCCCGCGCTGGCGATTCATTAAACTCACTCTG GTCTCAGGAGTTGTCTCGCAGCTGGCACAGGTTCCCGACTGGAAAGCGGGCAGTGA AACGCAATTAAATGAGTTAGCTCACTCATTAGGCACCCCAAGGCTTACACTTATGCTTCCGGCT CGTATGTTGTGAGGATAACAATTTCACACAGGAAACAGCTATGACCATGATTAC
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	GATACCTACAGCGTGAGCTATGAGAAAGGCCACGCCCTCCGAAGGGAGAAAGGCCGACAGGTATCCGGTAAGCGCAGGGTCGGAACAGGAGAGCGCACGAGGGAGCTCCAGGGGAAACGCCCTGGTATCTTATAGTCCTGCGGGTTTCGCCACCTCTGACTTGCAGCGTCGATTTGTGATGCTCGTCAGGGGGCGGAGCCTATGGAAAAAACGCCAGCAACGCCGGCCTTTACGGGTCCTGGCTGGCCTTGTGAGTGCAGCTGATACCGCTGCCGCAGCGAACCGAGCGCAGCGAGTCAGTGAGCGAGGAAGCAGAGCGCCCAATACGCAAACCGCCCTCTCCCCCGCGCTGGCGATTCAATTAACTCTGTGTCCTAAATGTTGCTTCGCAGCTGGCACGACAGGTTCCGACTGGAAAGCAGGGCAGTGAGCGAACGCAATTAAATGTGAGTTAGCTCACTCATTAGGCACCCCAGGCTTACACTTATGCTTCGGCTCGTGTGAGTGCAGCTGAGCGATAACAATTTCACACAGGAAACAGCAGTGACCATGATTACGCCAAGCTGCACTGCCATGCCAGCTGGCGTACCGCAAGGAGGCCGACCGATGCCCTCCCAACAGTTGCGCCTGAATGGCGACTTCAGTACAATCTGCTCTGATGCCCATAGTTAACCGAGCCCCGACACCCGCAACACCCGCTGACCGCCCTGACGGGCTGTCTGCTCCCGCATCGCCTAACAGACAAGCTGTGACGAAGACAAGGTTGAGACCACGAA
P_Gene	GTGGCTCTCAGTGGACAAAGGCCCTCGTGTACGCCCTATTTTATAGGTTAATGTCATGATAAT AATGGTTCTTAGACGTCAAGGTGGCACTTTCGGGAAATGTGCGCGAACCCCTATTGTTATT TTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCCTGATAAATGCTTCAATAATAT TGAAAAGGAAGAGTATGCGCTCACGCAACTGGTCCAGGAACACCTTGCAGCGACGCCGGTGGTAA CGGCCAGTGGGGTTTCATGGCTTGTATGACTGTTTTTGGGGTACAGTCTATGCCCTGGGC ATCCAAGCAGCAAGCGCGTTACGCCGTGGGTGATGTTGATGTTATGGAGCAGCAACGATGTTA CGCAGCAGGGCAGTCGCCCTAAACAAAGTTAAACATCATGAGGGAAGCGGTGATGCCGAAGTA TCGACTCAACTATCAGAGGTAGTTGGCGTACCGAGCCATCTGAGCGCCATCTGAGCGACGTTGCTGCCGT ACATTGTAACGGCTCCGCACTGGATGGCGCTGAAGCCACACAGCGATATTGATTGCTGGTAA CGGTGACCGTAAGGCTTGTGAAACAACACGCCGGAGCTTGTGATCAACAGCACCTTGGAAACTTCG GCTTCCCTGGAGAGGAGCGAGATCTCCGCGTGTAGAAGTCACCATTTGTGACGACGACAT CATTCCGTGGCGTATTCAGCTAACGCCGAAGCTGCAATTGGAGAATGGCAGCGCAATGACATTC TTGCAAGGTATCTCGAGGCCACGATCGACATTGATCTGGCTATCTGCTGACAAAAGCAAGAG AACATAGCGTTCCCTGGTAGGTCCAGCGCGGAGGAACCTTGTGATCCGGTTCTGAACAGGAT CTATTGAGGCCCTAAATGAAACCTTAACGCTATGGAACCTGCCGCCCAGTGGCTGGCGATGA GCGAAATGTAGTGCTTACGTTGTCCCGATTGGTACAGCCAGTAACCGGAAACATCGCAGCCAGTAC AGGATGTCGCTGCCGACTGGCAATGGAGCGCTGCCGGCCAGTATCAGCCCGTACACTTGAA GCTAGACAGGTTATCTTGACAAGAAGAGTCAGCTGGCTGGCCGAGATCAGTTGGAAAGA ATTGTCCTAACGTTAAAGGCAGAGTACCAAGGTAGTCAGCTGGCAATAACTGTCAGACCAAGTTA CTCATATATACTTGTGTTAAACTCTTGTGAGATCCTTTCTGCGCGTAACTCTGCTGCTGCAAACAAAA AAACCCAGCTTACAGCGGTGGTTGGCCGATCAAGAGCTACCAACTCTTCCGAAGGTA ACTGGCTTCAGCAGAGCGCAGATAACAAACTGTCCTCTGCTAATCCTGTTACAGTGGCTGCTGCC AGTGGCGATAACTCGTGTCTACCGGGTTGGACTCAAGAGCATAGTTACCGGATAAGGCCAGCG GTCGGGCTGAACGGGGGGCTGTGACACAGCCAGCTGGAGCGAACGACCTACACCGAACGTA GATACTACAGCGTGAGCTATGAGAAAGCGCCACGCTCCCGAAGGGAGAAAGGCCAGGTT CCAGCGTAAGCGCAGGGCTCGGAACAGGAGAGCGCACGAGGGAGCTCCAGGGGAAACGCCCTGGT ATCTTATAGTCCTGCGGGTTTCGCCACCTCTGACTTGCAGCGTCGATTGTTGATGCTCGTCAG GGGGGCGGAGCCTATGGAAAAACGCCAGCAACGCCGGCTTTACGGTCTGGCTGGCTTTGTGCG CTTTGTGTCACATGTTCTTCCTGCGTTATCCCTGATTCCTGTTGATAACCGTATTACGCCCTTG AGTGAGCTGATACCGCTGCCGCAGCCGAACGACCGAGCGCAGCGAGTCAGTGAGCGAGGAAGC GGAAGAGGCCCAATACGCAAACCGCCCTCTCCCGCGCTGGCGATTCAATTACTCTGTG GTCTCAAGGTTGTCTCCGAGCTGGCACAGCAGGTTCCGACTGGAAACCCCTGGCGTACCCAACTTAATGCCCTGAGCAC ATCCCCCTTCCGCCAGCTGGCGTAATAGCAAGAGGCCGACCGATGCCCTCCCAACAGTT CGCAGCCTGAATGGCGAATGGCGCTGATGCCGTATTTCTCCTACGCATCTGCGGTATTCA CACCGCATATGGCACTTCAGTACAATCTGCTCTGATGCCCATAGTTAACCGAGCCCCGACAC CCGCAACACCCGCTGACCGCCCTGACGGGCTGTCTGCTCCCGCATCGCCTAACAGACAAGCTGTGACGAAGACAAGGTTGAGACCACGAA
P_3'UTR	GTGGCTCTCAGTGGACAAAGGCCCTCGTGTACGCCCTATTTTATAGGTTAATGTCATGATAAT AATGGTTCTTAGACGTCAAGGTGGCACTTTCGGGAAATGTGCGCGAACCCCTATTGTTATT TTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCCTGATAAATGCTTCAATAATAT TGAAAAGGAAGAGTATGCGCTCACGCAACTGGTCCAGGAACACCTTGCAGCGACGCCGGTGGTAA CGGCCAGTGGGGTTTCATGGCTTGTATGACTGTTTTTGGGGTACAGTCTATGCCCTGGGC ATCCAAGCAGCAAGCGCGTTACGCCGTGGGTGATGTTGATGTTATGGAGCAGCAACGATGTTA CGCAGCAGGGCAGTCGCCCTAAACAAAGTTAAACATCATGAGGGAAGCGGTGATGCCGAAGTA TCGACTCAACTATCAGAGGTAGTTGGCGTACCGAGCCATCTGAGCGCCATCTGAGCGACGTTGCTGCCGT ACATTGTAACGGCTCCGCACTGGATGGCGCCCTGAAGCCACACAGCGATATTGATTGCTGGTAA CGGTGACCGTAAGGCTTGTGAAACAACAGCGCCAGCTTGTACAGGACCTTGGAAACTTCG GCTTCCCTGGAGAGAGCGAGATTCTCCGCGTGTAGAAGTCACCATTTGTGACGACGACAT CATTCCGTGGCGTTATCCAGCTAACGCCGAACGCGCAACTGCAATTGGAGAATGGCAGCGCAATGACATTC TTGCAAGGTATCTCGAGGCCACGATCGACATTGATCTGGCTATCTGCTGACAAAAGCAAGAG AACATAGCGTTCCCTGGTAGGTCCAGCGCGGAGGAACCTTGTGATCCGGTCTGAACAGGAT CTATTGAGGCCCTAAATGAAACCTTAACGCTATGGAACCTGCCGCCAGTGGCTGGCGATGA GCGAAATGTAGTGCTTACGTTGTCCCGATTGGTACAGCCAGTAACCGGAAACATCGCAGCCAGGATGCGCTGACCGCAATGACATTC AGGATGTCGCTGCCGACTGGCAATGGAGCGCCCTGCCGGCCAGTATCAGCCCGTACACTTGAA

	GCTAGACAGGCTTATCTGGACAAGAAGAAGATCGCTGGCCTCGCGCGAGATCAGTTGGAAGA ATTGTCCATTACGTAAAAGGCAGATCACCAAGGTAGTCGGCAAATACTGTAGCACCCAAGTTA CTCATATATACTTTAGATTGATTTAAACTTCATTTAATTAAAGGATCTAGGTGAAGGATCCTT TTTGATAATCTCATGACCAAAATCCCTAACGTGAGTTTCGTTCCACTGAGCGTCAGACCCCCGTA GAAAAGATCAAAGGATCTTCTGAGATCCTTTTCTGCGCGTAATCTGCTGTTGCAAACAAAA AAACCACCGCTACCAGCGGTGGTTGTTGCCGATCAAGAGCTACCAACTCTTCCGAAGGTA ACTGGCTTCAGCAGAGCGCAGATACCAAATACTGTCCTCTAGTGTAGCCGTAGTTAGGCCACCAC TTCAAGAACTCTGTAGCACCCTACATACCTCGCTCTGCTAATCCTGTTACCGTAGTGGCTGCTGCC AGTGGCATAAGTCGTCTTACGGGTGGACTCAAGACGGATAGTTACCGGATAAGGCGCAGCG GTCGGCTGAACGGGGGTTCTGTCACACAGGCCAGCTTGAGCGAACGACCTACACCGAAGCTGA GATACTACAGCGTGGACTATGAGAAAGGCCACGCGAGAGCGCACGAGGGAGCTCCAGGGGAAACGCCCTGGT ATCTTATAGTCTGCGGGTTCTGCCACCTCTGACTTGAGCTGCTTCCACTGAGCGTCAGACCCCCGTA GGGGCGGAGCCTATGGAAAAACGCCAGCAACGCCGCTTACGGTTCTGGATAACCGTATTACCGCCTTGT CCTTTGCTCACATGTTCTCCTGCGTTATCCCTGATTCTGTGATAACCGTATTACCGCCTTGT AGTGAGCTGATACCGCTCGCCGAGCGAACGACCGAGCGCAGCGAGTCAGTGGAGCAGGAAGC GGAAGAGGCCAATACGCAAACGCCCTCTCCCGCGCTGGCGATTCTAACACTCTGTG GTCTCACAACCTGCTTGCAGCTGGCACGACAGGTTCCGACTGGAAAGCAGGGCAGTGAGCGC AACGCAATTATGTGAGTTAGCTACTCATTAGGACCCAGGCTTACACTTATGCTCCGGCT CGTATGTTGTTGAGATTGAGCGGATAACAACTTACACAGGAAACAGCTATGACCATGATTAC GCCAAGCTTGCATGCCAGGCTGACTGGAAAACCTGGCTTACCCAACTTACGCCCTGAGCAC ATCCCCCTTCGCCAGCTGGCTAATAGCAAGAGGCCGACCGATGCCCTCCAAACAGTTG CGCAGCCTGAATGGCAATGGCCTGATGGGTATTTCCTTACGCATCTGTGCGGTATTCA CACCGCATATGGTCACTCTAGTACAATCTGCTGATGCCCATAGTTAAGCCAGCCCCGACAC TCGCCAACACCGCTGACGGCCCTGACGGGCTGTGCTGCCATCGCTTACAGACAAGC TGTGACGAAGACAACGCTTGGAGACCAACGAA
P_polyA	GTGGCTCTTCAGTGGACGAAGGGCTCTGATACGCCATTTTTATAGTTAATGTCATGATAAT AATGGTTCTTAGACGTCAAGTGGCCTTTCTGGGAAATGTGCGCGGAACCCCTATTGTTATT TTCTAAATACATTCAAAATATGTATCCGCTCATGAGACAATAACCCCTGATAAATGCTCAATAATAT TGAAAAGGAAGAGTATGGCCTCACGCAACTGGCAGACCCCTGACCGAACGCGAGCGGTGGTAA CGGGCAGTGGGGTTTCTGGCTTATGACTGTTTTGGGGTACAGTCTATGCCCTGGGC ATCCAAGCAGCAAGCGCTTACGCCGTGGTCGATGTTGATGTTATGGAGCAGCAACGATGTTA CGCAGCAGGGCAGTCGCCCTAAACAAAGTTAACATCATGAGGGAAAGCGGTGATGCCGAAGTA TCGACTCAACTATCACAGGGTAGTTGGCTCATCGAGGCCATCTGAAACGACGTTGCTGGCGT ACATTGTCAGGCTCCGCACTGGATGGCCTGAAGGACACAGCGATATTGATTGCTGGTTA CGGTGAGCGTAAGGCTTGTGAAACAAACGCCAGCTTACCGGACCTTGGAAACCTTGGAAACTTGG GCTTCCCTGGAGAGAGCGAGATTCTCGCGTGTAGAAGTCACCATTTGTTGACGACGACAT CATTCCGTGGCTTATCCAGCTAACGCCGAACGCAACTGCAATTGGAGAATGGAGCAGCAATGACATT TTGCAAGGTATCTCGAGCCAGCAGATCGACATTGATCTGGCTATCTGCTGACAAAAGCAAGAG AACATAGCGTTGCCCTGGTAGGTCCAGCGGGAGGAACCTTGTGATCCGGTCTGAACAGGAT CTATTGAGGGCTAAATGAAACCTTAAACGCTATGGAACCTGCCGACTGGCTGGCGATGA GCGAAATGTAAGTCTACGGTCTCCGCAGTGGCAATTGGAGGCCATGGGGCCGACTATGCCGCTACACTTGAA AGGATGTCGCTCCGACTGGCAATTGGAGGCCATGGGGCCGACTATGCCGCTACACTTGAA GCTAGACAGGCTATCTGGACAAGAAGAAGATGCCGTTGGCTGCCGAGATCAGTTGGAAAGA ATTGTCCATTACGTAAGAGCTACCAAGGTAGTCGGAAATAACTGTCAGACCAAGTTA CTCATATATACTTTAGATTGATTTAAACTTCATTTAATTAAAGGATCTAGGTGAAGGATCCTT TTTGATAATCTCATGACCAAATCCCTAACGTGAGTTTCGTTCCACTGAGCGTCAGACCCCCGTA GAAAAGATCAAAGGATCTTCTGAGATCCTTTTCTGCGCGTAATCTGCTGTTGCAAACAAAA AAACCACCGCTACCAGCGGTGGTTGTTGCCGATCAAGAGCTACCAACTCTTCCGAAGGTA ACTGGCTTACGGCAGAGCGCAGATACCAAATACTGTCCTCTAGTGTAGCCGTAGTTAGGCCACCAC TTCAAGAACTCTGAGCACCCTACATACCTCGCTCTGCTAATCCTGTTACCGAGTGGCTGCTGCC AGTGGCATAAGTCGTCTTACGGGTGGACTCAAGACGGATAGTTACCGGATAAGGCGCAGCG GTCGGCTGAACGGGGGTTCTGTCACACAGGCCAGCTTGAGCGAACGACCTACACCGAAGCTGA GATACTACAGCGTGGACTATGAGAAAGGCCACGCGAGGGAGCTCCAGGGGAAACGCCCTGGT CCGGTAAGCGCAGGGCTGGAACAGGAGAGCGCACGAGGGAGCTCCAGGGGAAACGCCCTGGT ATCTTATAGTCTGCGGGTTCTGCCACCTCTGACTTGAGCGTCGATTGTTGATGCTGCTCAG GGGGCGGAGCCTATGGAAAAACGCCAGCAACGCCGCTTACGGTTCTGGCTTTGCTGG CCTTTGCTCACATGTTCTCCTGCGTTATCCCTGATTCTGTGATAACCGTATTACCGCCTTGT AGTGAGCTGATACCGCTCGCCGAGCGAACGACCGAGCGCAGCGAGTCAGTGGAGCAGGAAGC GGAAGAGGCCAATACGCAAACGCCCTCTCCCGCGCTGGCGATTCTAACACTCTGTG GTCTCACAACCTGCTTGCAGCTGGCACGACAGGTTCCGACTGGAAAGCAGGGCAGTGAGCGC AACGCAATTATGTGAGTTAGCTACTCATTAGGACCCAGGCTTACACTTATGCTCCGGCT CGTATGTTGTTGAGATTGAGCGGATAACAACTTACACAGGAAACAGCTATGACCATGATTAC GCCAAGCTTGCATGCCAGGCTGACTGGAAAACCTGGCTTACCCAACTTACGCCCTGAGCAC ATCCCCCTTCGCCAGCTGGCTAATAGCAAGAGGCCGACCGATGCCCTCCAAACAGTTG CGCAGCCTGAATGGCAATGGCCTGATGGGTATTTCCTTACGCATCTGTGCGGTATTCA CACCGCATATGGTCACTCTAGTACAATCTGCTGATGCCCATAGTTAAGCCAGCCCCGACAC TCGCCAACACCGCTGACGGCCCTGACGGGCTGTGCTGCCATCGCTTACAGACAAGC TGTGACGAAGACAACGCTTGGAGACCAACGAA
pTU-1	TCTGTGAAGACAATGCCGAATCGGATCCGGAGTGAGACCGCAGCTGGCACGACAGGTTGCCGA CTGGAAAGCGGGCAGTGAGCGCAACGCAATTATGTGAGTTAGCTACTCATTAGGACCCCAAG CTTTACACTTTATGCTCCGGCTGTATGTTGTTGGAATTGAGCGGATAACAAATTTCACACAG GAAACAGCTATGACCATGTTACGCCAAGCTTGCATGCCCTGAGCGACTCTAGAGGATCCCCG GGTACCGAGCTGAATTCACTGGCCCTGTTTACAACGTCGTGACTGGAAAACCCCTGGCTTA CCCAACTTAACTGCCCTGCCAGCACATCCCCCTTGCAGCTGGCGTAATAGCGAAGAGGCCG ACCGATGCCCTCCCAACAGTTGCCAGCAGCGCTGAATGGCGAATGGCGCCTGATGCCGTATTTC

	CCTTACGCATCTGTGCGGTATTCACACCGCATATGGTCACTCTCAGTACAATCTGCTCTGATGC CGCATAGTTAACGCCAGCCCCGACACCCGCCAACACCCGCTGACGCGCCCTGACGGGCTTGTCTGC TCCCCGCATCCCCTTACAGACAAGCTGTGACGGTCTCACCGCTGCAATTGTCTCTGACGAAGTG GTTTAAACTATCAGTGTGACAGGATATATTGGCGGGTAAACCTAAGAGAAAAGAGCGTTTATTA GAATAATCGGATATTAAAGGGCGTAAAAGGTTATCCGCTGTCATTGTATGTGATGCC ACCACAGGGTTCCCAGATCAGGCCTGGCTGTGAACCCCCAGCCGAACGTGACCCCACAAGGC CCTAGCGTTGCAATGCACCAAGTCATATTGACCCAGGGCTTCCACCAGGCCGTCCTCGC AACTCTCGCAGGCTTCGCCGACCTGCTCGCCACTTCTCACCGGGTGGAAATCCGATCCGCA CATGAGGCCAGGTTCCAGCTGAGCAGCACCCGATTCCAGGTGCCAACAGCGGTGGACGTGAAG CCCATGCCGTCGCTGTAGGGCAGACAGGCTTCGCCCCCTGTAATACCGGCCATTGAT CGACCAGCCCAGGTCTGGCAAAGCTGTAGAACGTGAAGGTGATGGCTGCCGATAGGGGTGC GCTTCGCGTACTCCAACACCTGCTGCCACACCAGTCGTATCGTCGGCCGCACTCGACGCCG GTGTAGGTGATCTCACGTCCTGTTGACGTGAAAATGACCTTGTGTCAGCGCCTCGCGG GATTTCTTGTGCGCGTGGTGAACAGGGCAGAGCAGGGCCGTCGTTGGCATCGTCATCG TGTCCGGCCACGGCGCAATATCGAACAAAGGAAAGCTGCAATTCTTGATCTGCTGTTGTT TCAGCAACGCGGCCGCTGCTGGCAGCTGACGCTGACTGCTTGGCAGGTCTGCCGGGGGTTTCGCT TCTTGGTCTGTCATAGTCTCTGCTGCTGATGGTCTGACGCTTGGCAGGGCAGGGGAGCCAGTTGAC GCTGCGCCTCGATCTGGCGTAGCTGCTGGACCATCGAGCCGACGGACTGGAAGGTTGCG GGGGCGACGCATGACGGTGCAGCTGCGATGGTTTGGCATTCTCGCCGGAAAACCCCGCGTC GATCAGTTCTGCTGTATGCCCTCCGTCAAACGTCGATTCACTCACCCTCTGCGGGATTGC CCCGACTCACGCCGGGCAATGTGCCCTTATTCTGATTGACCCGCTGGTGCCTGGTCCA GATAATCCACCTTATGCCAATGAAGTCGGTCCCGTAGACCGTCTGGCCCTCTGACTTGG TATTCCGAATCTGCCCTGACGAATACCAGCGACCCCTGCCAAATACTTGGCGTGGCCCTCGG CCTGAGAGCCAAAACACTTGTATCGGAAAGACTGGCGCTCTGCTGCTGCGTTTTCATAGGCT CGCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCTGTCGCTGCGTTTTCATAGGCT CGCCCCCCTGACGAGCATCACAAAATCAGCCTCAAGTCAAGAGTGGCAGACCCGACAGGAAT ATAAAGATAACCAAGCGTTCCCCCTGGAAGCTCCCTGTCGCTCTCTGTTCCGACCCGCTGCT TACCGGATACTGTCCGCCCTTCTCCCTCGGAAGCGTGGCCTTCTCATAGCTCACGCTGTAG GTATCTCAGTTGGTGTAGTCGTTGCTCCAAGCTGGCTGTGCAACGACCCCCGTTCAAGC CCGACCGCTGCCCTATCGGTAACTATCGTCTTGAGTCAACCCGTAAGACACGACTTATCGC CACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCAGGGTATGAGGCGTCTCACAGAGTT TTGAAGTGGTGGCCTAAACTACGGTCAACTAGAAGGACAGTATTGGTATCTGCGCTCTGCTGAA GCCAGTTACCTCGGAAAAGAGTTGGTAGCTCTGATCCGCAACAAACACCCTGGTAGCG GTGGTTTTTTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTGA TCTTCTACGGGTCTGACGCTCAGTGGAACAAACTCACGTTAAGGGATTGGTATGAGAT TATCAAAAGGATCTCACCTAGATCCTTTAAATTAAAATGAAGTTAAATCAATCTAAAGTAT ATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGACACTATCAGCAGATCTG TCTATTCGTTCATCCATAGTCTGCTGACTCCCCGCTGTAGATAACTACGATAACGGGAGGGCTT ACCATCTGGCCCGACTGCTGCAATGATACCGCAGAGAACCGCTACCGGCTCCAGATTTATCAG CAATAAACCAGCCAGCGGAAGGGCGAGCGCAGAAGTGGCTCTGCAACATTATCCGCTCATC CAGTCTATTAAATTGTGCGGGGAAGCTAGAGTAAGTAGTGTGCTGCCAGTTAATAGTTGCGCAACGTT GTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTGTGGTATGGCTTCACTCAGCTCCGGT TCCCAACGATCAAGGCAGTTACATGATCCCCATGTTGCAAAAAGCGGTTAGCTCCTCGGT CCTCCGATCGTGTCAAGAATAAGTGGCCGAGTGTATCACTCATGGTTATGGCAGCACTGCAT AATTCTTAACTGTCATGCCATCCGTAAGATGTTCTGACTGGTAGACTGGTAGACTCAACCAAGTCA TTCTGAGAATAGTGTATGGCGACGGAGTTGCTCTGCCCCGGCTCAATACGGGATAACCGC GCCACATGACGAAACTTTAAAGTGTCTCATTTGGAAAAGCTTCTCGGGGGAACAAACTCTAAG GATCTTACCGCTGTTGAGATCAGTTGATGAACTCAGTCCACTCGTGCACCCACTGATCTCAGCATC TTTACTTCACCGCTGTTGGTAGCAAAACAGGAAAGGCAAATGGCAGAAAAGGAA TAAGGGCGACACGGAATGTTGAATACTCATACTCTTCTTTCAATATTGAAGCATTATCA GGGTTATTGTCTCATGAGCGGATACATATTGAATGTATTAGAAAATAACAAATAGGGTTC GCGCACGAATTGGCCAGCCTGCCATTGGGGTGAAGCGCTGCCGGCGAGGGCGCAGCC CCTGGGGGGATGGGAGGCCGCTAGGGCCAGCTGGCAACCGCAATTATGTGAGTTAGCTCACTCATAGGCAACCCAGG CTTTACACTTTATGCTCCGGCTGTATGTTGTGAGGAAATTGTGAGCTGAGCGATAACAATTTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTTGTGATGCCCTGCAAGGTGACTCTAGAGGATCCCG GGTACCGAGCTGCAATTCACTGGCCGCTGTTTACAACGTCGTTGACTGGAAAACCCCTGGCGTTA CCCACTTAATGCCCTGCAAGCACATCCCCCTTCGCCAGCTGGCTAATAGCGAAGAGGCCGC ACCGATGCCCTCCCAACAGTGTGCGCAGCCTGAATGGCGAATGGCGCTGATGCCGTATTTCG CCTTACGCATCTGTGCGGTATTTCACACCGCATATGGTCACTCTCAGTACAATCTGCTGTGATGC
pTU-2	TCTGTGAAGACAAGCAAGAATTCAAGCTGGAGTGAGACCCGAGCTGGCACGACAGGTTGCCGA CTGGAAAGCGGGCAGTGAGCGCAACCGCAATTATGTGAGTTAGCTCACTCATAGGCAACCCAGG CTTTACACTTTATGCTCCGGCTGTATGTTGTGAGGAAATTGTGAGCTGAGCGATAACAATTTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTTGTGATGCCCTGCAAGGTGACTCTAGAGGATCCCG GGTACCGAGCTGCAATTCACTGGCCGCTGTTTACAACGTCGTTGACTGGAAAACCCCTGGCGTTA CCCACTTAATGCCCTGCAAGCACATCCCCCTTCGCCAGCTGGCTAATAGCGAAGAGGCCGC ACCGATGCCCTCCCAACAGTGTGCGCAGCCTGAATGGCGAATGGCGCTGATGCCGTATTTCG CCTTACGCATCTGTGCGGTATTTCACACCGCATATGGTCACTCTCAGTACAATCTGCTGTGATGC

	CGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCCCTGACGGGTTGTCTGC TCCCCGCATCCGCTTACAGACAAGCTGTGACGGTCTCACGCTACTATTGTCCTCTGCACGAAGTGG TTAAACTATCAGTGTGACAGGATATTGGCGGGTAAACCTAAGGAAAAGAGCGTTATTAG AATAATCGGATATTAAAGGGCGTAAAAGGTTTATCGCTCGTCAACCCCCAGCCGAACGACCCCACAAAGGC CCACAGGGTCCCCAGATCAGCGCTGCTGTAACCCCCAGCCGAACGACCCCACAAAGGC CTAGCGTTGCAATGCACCAGGTACATGACCCAGCGTGTCCACCCAGGCCGCTGCCCGCA ACTCTCGCAGGCTTCGCCACCTGCTCGCCACTTCTCACGCGGTGGAATCCGATCCGCAC ATGAGGCGGAAGGTTCCAGCTTGAGCGGGTACGGCTCCCGTGCAGCTGAATAGTCGAACAT CCGTCGGGCCGTGGCGACAGCTTGCCTACTTCTCCATATGAATTCTGTAGTGGTCGCCAG CAAACAGCACGACGATTTCTCGTCATCAGGACCTGGCAACGGACGTTTCTGCCACGGTCC AGGACGCCGAAGGGTGCAGCACCGACAGCTTCCAGGTGCCCACAGCGGACGCTGAAGC CCATCGCCGTCGCTGTAGGCGCACAGGCATTCTGCCGCTCGTGAATACCGGCCATTGATC GACCAGCCCAGGTCTGGCAAAGCTCTGAGAACGTTGATCGGCTCGCCGATAGGGTGC GCTTCGCGTACTCCAACACTGCTGCCACACCAGTCGTACATCGTGGCCCGCAGCTCGACGCCG GTGTAGGTATCTCACGTCCTGTGACGTGAAAATGACCTGTTGCAAGCCTCGCCGCG GATTTCTTGTGCGCGTGGTGAACAGGGCAGAGCGGGCGTGTGTTGGCATCGCTCGCATCG TGTCCGGCCACGGCGCAATATCGAACAGGAAAGCTGCAATTCTTGATCTGCTGTTGTT TCAGCAACGCGGCCGTGCTGGCTGACCTTGTGCACTGATGGTCATCGACTTCGCCAAACCTGCCGCTCCTGTT TCTTGGTCTGTCATAGTCCCTCGCTGTGATGGTCATCGACTTCGCCAAACCTGCCGCTCCTGTT CAAGACGACGCCAACGCTTCCACGGCGGGCGATGGCGGGGAGGGCAGGGCAGTGCAC GCTTCGCGTCTGATCTGGCGTAGCTGCTGGATGGTCTGGCGATCTGCGTGGCG GGGGCGCACGCACTGACGGTGCCTGCGATGGTTTGGCATCTCGCGGAAAACCCCGCGTC GATCAGTTCTGCTGTATGCCCTCCGGTCAAACGTCGATTCACTCACCCTCTGCCGGATTG CCCACACTACGCCGGGCAATGTGCCCTTATTCTGATTGACCCGCTGGTGCCTTGGTGTCCA GATAATCCACCTTATCGGAATGAAGTGGTCCGTAGACCGTCTGGCCCTCTCGTACTTGG TATTCCGAATCTGCCCTGACGAATACCGACGCCCTTGCCAAATACTTGGCGTGGCGCTCG CTTGAGGACCAAACACTTGTGCGGAAGAACGTTGCTGCGCTCTGCTTGTGCGCGCATCGT CGCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCGTGTGCGTGGCGTTTCCATAGGCTC CGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAAGGAGTGGCAGACCCGACAGGACT ATAAAGATAACCGCGTTCCTGGCTTAACACTACGCGTCAACTAGAAGGACAGTATTGGTATTCGCTCTGCTGAG CCGACCGTCTGCGCTTATCGGTAACTATCGTCTTGTGAGTCCAACCCGTAAGACACGACTTATCGC CACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGGCGGTCTACAGAGTTC TTGAAGTGGTGGCTTAACACTACGCGTCAACTAGAAGGACAGTATTGGTATTCGCTCTGCTGAG GCCAGTTACCTTCGAAAAAGAGTTGGTAGCTCTGATCCGGCAAACAAACACCAGCTGGTAGCG GTGGTTTTTGTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCTTGA TCTTCTACGGGCTGACGCTCAGTGGAAACAAACTCACGTTAAGGGATTGGTATGAGAT TATCAAAAGGATCTCACCTAGATCCTTAAATTAAAATGAAGTTAAATCAATCTAAAGTAT ATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCACTGAGGACCTATCTCAGCGATCTG TCTATTCGTTCATCCATAGTGTGCTGACTCCCCGCTGTAGATAACTACGATAACGGGAGGGCTT ACCACATGGCCCCAGTGTCAATGATACCCGCGAGAACACCGCTACCGGCTCCAGATTATCAG CAATAAACCGAGCCGGAAAGGGCGAGCGCAGAGTGGCTCTGCAACTTATCCGCCCTCATC CAGTCTATTAAATTGTCGGGAAAGCTAGAGTAAGTAGTTGCGCTGGTAACTAGTGGCTT GTGCGCATTGCTACAGGCATCGTGGTGTACGCTCGCTGCTGGTATGGCTTCACTCAGCTCCGGT TCCCAACGATCAAGGCAGGTTACATGATCCCCCATGTTGTCGAAAAAGCGGTTAGCTCTCGGT CCTCCGATCGTGTCAAGTAAGTGGCCGAGTGTATCACTCATGTTATGGCAGCACTGCA AATTCTTACTGTATGCCATCCGTAAAGATGCTTCTGTGACTGGTAGTACTCAACCAAGTCA TCTGAGAATAGTGTATGGCGACCGAGTTGCTCTGCCCCGCTCAATACGGGATAATACCGC GCCACATAGCAGAACTTAAAGTGTCACTATTGAAAACGTTCTCGGGCGAAAACCTCTCAAG GATCTTACCGCTGTGAGGACTCGTGTGAGTAAACCCACTGTGCAACCAACTGATCTCAGCATC TTTACTTACCCAGGTTGTCCACATCATCTGTGGAAAACCTCGCTGAAAATCAGGCTTTC GCCGATTGCGAGGCTGGCAGCTCACGTCGCCGCGAAATCGAGGCTGCCCTCATCTGTC ACGCCCGCCGGGTGAGTCGGCCCTCAAGTGTCAACGTCGCCCCCTCATCTGTCAGTGAGGGCC AAGTTTCCGCGAGGTATCCACAACGCCGGCGGGCGGGTGTCTGCACACGGCTTCGACGGCGT TCTGGCGCTTGCAAGGCCATAGACGGCCGAGCCCAGCGGGCAAGGGCAACAGCCGGT AGCGTCGCAAAGGAGATCTGACTGATGGCTGCCGTATCGAGTGGTATTGGCG AGCTGCCGGTGGGGAGCTGTTGGCTGGCTGGCGAGGATATATTGTTGTAACAAATTGAC GCTTAGACAACTAATAACACATTGCCGACGTTTAAATGTA CTGAGTGGGATGCACTGGGCCCC C
pTU-3	TCTGTGAAGACAAACTAGAATTGAGCTCGGAGTGAGACCGCAGCTGGCACGACAGGTTTGGCGA CTGGAAAGCGGGCAGTGAAGCGCAACCGCAATTAAATGAGTTAGCTCACTCATTAGGCACCC CTTACACTTTATGCTCCGGCTGTATGTTGTGGAATTGAGTCAGCGATAACAATTTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTTGTGATGCCCTGAGGTGACTCTAGAGGATCCCC GGTACCGAGCTGCAATTACTGGCCGTCGTTTACAACGTCGCTGACTGGAAAACCTGGCTTA CCCAACTTAATGCCCTTGCAAGCACATCCCCCTTGCCTGAGCTGGTAAATAGCGAAGAGGCC ACCGATGCCCTTCCAAACAGTTGCGCAGCTGTAATGGCGAATGGCGCTGATGGCTT CCTTACGCTATGTCGGTATTTCACACGCCATAGGTGCACTCTAGTACAATCTGCTCTGATGC CGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCCCTGACGGGCTGTCTGC

	<p>GTAACTATCAGTGGACAGGATATTCGGGGTAAACCTAAGAGAAAAGAGCGTTATT GAATAATCGGATATTAAGGGCGTAAAAGGTTATCCGTCGCCCCACTTGTATGCGATGCCA ACCACAGGGTCCCCAGATCAGGCCTGGCTGTAACCCCCAGCCGAACGACCCACAAGGC CCTAGCGTTGAATGCACCAGGTACATTGACCCAGGCGTGTCCACCAAGGCCGTCCTCGC AACTCTCGCAGGCTCGCCGACTCTCGCGCCACTCTCACGCGGTGGAATCCGATCCGCA CATGAGGGAGGTTCCAGCTGAGCGGGTACGGCTCCCGTGCAGGCTGAATAGTCGAACA TCCGTCGGGCGTCGGCGACAGCTGGTACTCTCCATATGAATTCTGTAGTGGTCGCCA GAAACAGCACGACGATTCTCTCGTCGATCAGGACAGGCAACGGGACGTTCTTGCCACGGTC GGTACCGAGCTCGAATTCACTGGCGTGTGTTACAACGTCGTGACTGGAAAACCCCTGGCGTT CCCAACTTAATGCCCTGCAACGACATCCCCCTTCGCCAGCTGGCGTAATAGCGAAGAGGCC ACCGATGCCCTCCCAACAGTTCGCGACGCCCTGAATGGCGAATGGCGCTGATGCGGTATT CCTTACGCACTGTCGCGGTATTTCACACCGCATATGGTCACTCTCAGTAACTCTGCTCTGATGC CGCATAGTTAACCGACCCCCGACACCCGCCAACACCCCGTACGCGCCCTGACGGGGCTTGTCTGC TCCCGGCGATCGCTTACAGAACAGTGTGACGGTCTACGCTGTTGTCAGCAGCGTAAAGTGG TTAAACTATCAGTGGTGTGAGGATATTGCGGGTAAACCTAAGAGAAAAGAGCGTTATT</p>
pTU-5	<p>TCTGTGAAGACAACAGAGAATTGCGATGCCAGGAGTGAACCGCAGCTGGCACCGACAGGTTGCC CTGGAAAGCGGGCAGTGCAGCGAACCGCAATTATGTAAGTTAGCTCACTCATAGGCACCCAGG CTTACACTTATGCTCCGGCTGCTGATGTTGTTGAAATTGTCAGCGGATAACAAATTCTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTGCTGACGCCCTGAGGTGACTCTAGAGGATCCCCG GGTACCGAGCTCGAATTCACTGGCGTGTGTTACAACGTCGTGACTGGAAAACCCCTGGCGTT CCCAACTTAATGCCCTGCAACGACATCCCCCTTCGCCAGCTGGCGTAATAGCGAAGAGGCC ACCGATGCCCTCCCAACAGTTCGCGACGCCCTGAATGGCGAATGGCGCTGATGCGGTATT CCTTACGCACTGTCGCGGTATTTCACACCGCATATGGTCACTCTCAGTAACTCTGCTCTGATGC CGCATAGTTAACCGACCCCCGACACCCGCCAACACCCCGTACGCGCCCTGACGGGGCTTGTCTGC TCCCGGCGATCGCTTACAGAACAGTGTGACGGTCTACGCTGTTGTCAGCAGCGTAAAGTGG TTAAACTATCAGTGGTGTGAGGATATTGCGGGTAAACCTAAGAGAAAAGAGCGTTATT</p>

	ATAATCGGATATTTAAAGGGCGTAAAAGGTTATCCGTCGCCATTGTATGTGCATGCCA CCACAGGGTCCCCAGATCAGGGCTGGCTGTAACCCAGCGACTGACCCCACAAGGCC CTAGCGTTGCAATGCACCAGGTATCATTGACCCAGGGCTTCCACCAGGGCGTGCCTCGCA ACTCTCGCAGGCTCGGCCACCTGCTCGGCCACTTCTCACGCCGGTGGATCCGATCCGAC ATGAGGCGGAAGGTTCCAGCTGAGCGGTACGGCTCCCGTGCAGGCTGAAGAATAGTCGAACAT CCGTGGGGCGTCGGCGAACAGCTGCGTACTTCCTCCATGAAATTCTGTTGAGTGGTGC CAAACAGCACGACGATTCCTCGCATCAGGACCTGGCAACGGGAGCTTCTGCAAGTCC AGGACGCGGAAGCGGTGCAGCAGCACCCGATTCCAGGTGCCAACCGCGTGGACGTGAAGC CATCGCCGCTCGCTGTAGGCAGCACAGGCACTCCCTGGCCCTCGTGAATACCGGCCATTGATC GACCAGCCCAGGTCTGGCAAAGCTCGTAGAACGTGAAGGTATCGGCTGCCGATAGGGGTGC GCTTCGCGTACTCCAACACCTGCTGCCACACCAGTTCGTCATCGTCGCCCGCAGCTGACGCCG GTGTAGGTGATCTTACGTCCTTGTGACGTGGAAAATGACCTTGTGAGCGCCTCGCAG GATTTCCTGTGCGGTGGTAACAGGGCAGAGCGGGCGTGTGTTGGCATCGCTCGCAG TGTCCGGCACGGGCAATATCGAACAGGAAAGCTGCATTCTCTGATCTGCTGCTGCTGTT TCAGAACCGGGCTGCTTGGCTCGTGCACCTGTTTGCCAGGTTCTGCCGGGAGTTTCGCT TCTTGGTCTGCTAGTTCCTCGCGTGTGATGGTATCGACTTCGCAACCTGCCCTCTGTT CAAGACGACCGAACCGCTCCACGGCGGCCATGGCGGGGAGGGCAGGGGAGGGGAGCCAG GCTGTCGGCTCGATCTGGCGTAGCTTGTGACCGATCGAGCCGACGGACTGGAAGGTTTGC GGGGCGCACCGCATGACGGTGCCTGCGATGGTTGGCATCTCGGCCGAAACCCCGCGTC GATCAGTTCTGCTGTATGCCCTCCGGTAAACAGTCCGATTCTACCCCTCTGCGGGATTGC CCCGACTCACGCCGGGCAATGCGCCCTATTCTGATTGACCCGCTGGTGTGCGCTGGTCCA GATAATCCACCTATCGGAATGAAAGTGGCTGGTAGACCGTCTGGCAGGCTGGCTCTCTGACTCTGG TATTCCGAATCTGCCCTGCAAGAACGACGAGGCTGGCAAAACTACTGCGCTGGGCT CCTGAGAGCCAAAACACTTGTGCGGAAGAAGTGTGCGCTCTGCTTGTGCCGGCATGTTG GCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCGTGTGCGTGGCTTTCATAGGCTC CGCCCCCTGACGAGCATCACAAAATGACGCTCAAGTCAGAGGTGGCAGACAGGACT ATAAAGATACCAGGGCTTCCCCCTGGAAGCTCCCTGCGCTCTGTGCGCTCTCGTGGC TACCGGATACCTGTCGCCCTTCTCCCTCGGAAGCGTGGCGTTCTCATAGCTCACGCTGTAG GTATCTCAGTTCGGTGTAGGTGCTTCGCTCCAAAGCTGGCTGTGCGCAACCCCCCGTTCAGC CCGACCGCTGCCCTATCGGTAACTATCGTCTGAGTCAACCCGTAAGACACGACTATCGC CACTGGCAGCAGCCTGGTAAACAGGATTAGCAGAGCGAGGTATGAGCGGTGCTACAGAGTC TTGAAGTGGTGGCTAACTACGGTACACTAGAACGAGTATTGGTATCTGCGCTCTGCTGAA GCCAGTTACCTCGGAAAAAGAGTTGGTAGCTTGTGATCCGGCAACAAACACCACGCTGGTAGCG GTGGTTTTTGTGCAAGCAGCAGATTACGCGCAGAAAAAGGATCTAAGAAGATCCTTGA TCTTTCTACGGGCTGACGCTCAGTGGAAACGAAACTACGTTAACGGGATTGGTATGAGAT TATCAAAAGGATCTCACCTAGATCCTTTAAATTAAATGAAGTTAAATCAATCTAAAGTAT ATAGAGTAAACTTGGTGTACAGTTACCAATGCTTAAGTCACTGAGGGCACCTATCAGGATCTG TCTATTCTGTCATCCATAGTGGCTGACTCCCCCTGCTGTGAGATAACTACGATACGGAGGGCTT ACCATCTGGCCCCAGTCTGCAATGATACCGCAGAACACCGCTCACCGGCTCCAGATTTCAG CAATAAACAGCCAGCCGAAGGGCGAGCGCAGAACGTTCTGCAACTTATCCGCTCCATC CAGTCTATTAAATTGTCGGGGAGCTAGAGTAAGTAGTTGCTGCAAGTAACTGTTGCGCAG TTGCAATTGCTACAGGCATCGTGGTGTACGCTCGTCTGGTATGGCTTATTCTAGCTCCGGT TCCCAACGATCAAGGGAGTTACATGATCCCCATGTTGCAAAAAAGCGTTAGCTCTTGGT CCTCGATGTTGCAAGTAAGTGGCCGAGTGTATCACTCATGGTTATGGCAGCACTGCAT AATTCTCTAATGTCATGCCATCGTAAGTGTCTTCTGACTGGTGTAGACTCAACAAAGTC TTCTGAGAATAGTGTGCGGCCAGCTGCTTGTGAGTCACTGCAACCGGAAACTCTCAG GATCTTACCGCTGTTGAGATCCAGTTGATGTAACCCACTCGTGCACCAACTGATCTCAGCATC TTTACTTCAACAGCGTTCTGGGTGAGCAAAACAGGAAGGCAAATGCCCAAAAGGAA TAAGGGCGACCGAAATGTTGAATACTCATACTCTCCTTTCAATATTATGAAAGCATTTATCA GGGTTATTGCTCATGAGCGGATACATATTGAATGTTAGAAAAATAACAAATAGGGTIC GCGCACGAATTGGCCAGCGCTGCCATTGGGGTGAGGCCGTTGCGGGCCAGGGGCCAGCC CCTGGGGGATGGGAGGGCCGCTTACGGGGGGAGGGTTCAGAAGGGGGGCCACCCCT TCGGCGTGCCTGTCAGCGCACAGGGCGCAGCCCTGGTAAACAAAGGTTATAATATTGGT TTAAAGCAGGTTAAAGACAGGGTAGCGGTGGCCGAAAACGGCGGAAACCCCTGCAATGCT GGATTTCTGCTGTGGACAGCCCTCAAATGTCATAGGTGCGCCCTCATCTGTCAGCACTCTG CCCCCTCAAGTGTCAAGGATCGCCTTCACTGTCACTGTCAGTAGTCGCGCCCTCAAGTGTCAATACCG CAGGGCACTTATCCCCAGGCTTGTCCACATCATCTGTTGGAAACTCGCGTAAACATGGCGTTTC GCCGATTGCGAGGCTGGCCAGCTCCACGTCGCCGGCGAAATCGAGCTGCCCTCATCTGTC ACGCCGCCGGGTAGTCGCCCTCAAGTGTCAACAGTCCGCCCCCTCATCTGTCAGTAGGGCC AAGTTTCCCGAGGTATCCACAAAGCGGGGGCGGGTGTCTGCACACGGCTTCAGCGCGT TTCTGGCGCGTTGCAAGGGCCATAGACGCCGCCAGCGGCCAGGGCAACCAGCCGGTG AGCGTCGCAAAGGAGATCCTGACTGATGGCTGCTGTGGCAGGATATATTGTTGCA GCTTAGACAACCTAATAACACATTGCGGACGTTTTAATGTAACGTTGAGCTGGGATGCA GCTGGCGTGGCC
pTU-6	TCTGTGAAGACAATGTGGAATTCTCGAGGGAGTGAGACCGCAGCTGGCACGACAGGTTGCC CTGGAAAGCGGGCAGTGTAGCGCAACGCAATTAAATGTGAGTTAGCTCACTCATAGGCACCCAG CTTACACTTATGCTCCGGCTCGTGTGGAATTGTGAGCGGAAACAAATTCTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTGCTGATGCCCTGAGGTGCACTCTAGAGGATCCCCG GGTACCGAGCTCGAATTCACTGGCGTGTGTTACAACAGTGTGACTGGGAAACCCCTGCGTTA CCCAACTTAATGCCCTGCAACGACATCCCCCTTGCACGCCAGCTGGCGTAATAGCGAAGAGGCC ACCGATGCCCTCCAAACAGTTGCGCAGCCTGAATGGCGAATGGCGCCTGATGCGGTATTTCT CCTACGCACTGTGCGGTATTTCACACCGCATATGGTCACTCTAGTACAATCTGCTCTGATGC CGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCCCTGACGGGCTTGTCTG TCCCGCCTACCGCTTACAGAACAGCTGTGAGCGGTCTCACGCTGAGGCTTGTCTGCA GTTAAACATGAGGTTAGCGGGGAGCTGTTGGCTGGCTGGCAGGATATATTGTTGCA GCTTAGACAACCTAATAACACATTGCGGACGTTTTAATGTAACGTTGAGCTGGGATGCA GCTGGCGTGGCC

	<p>ACCAACAGGGTTCCCCAGATCAGGCCTGGCTGCTGAACCCCCAGCCGAAC TGACCCCACAAGGC CCTAGCGTTGCAATGCACCAGGTATCATTCATTGACCCAGGCCGTTCCACCCAGGCCGCTGCCTCGC AACTCTTCGAGGCTTCGCCGACCTGCTCGCCCACTTCTTCACGCCGGTGGAAATCCGATCCGCA CATGAGGGCGAAGGTTTCCAGCTTGAGCGGGTACGGCTCCCGTGCAGCTGAATAGTCGAACA TCCGTGGGCCGCTGGCGACAGCTGCGGTACTTCTCCCATATGAATTTCGTTAGTGGTCGGA GCAAACAGCACGACGATTCTCGTCGATCAGGACCTGGCAACGGGACGTTTCTGCCACGTC CAGGACGCCAGCGGTGCAGCAGCACCGATTCCAGGTGCCAACCGGGTGGACGTGAAG CCCATGCCGTGCGCTGTAGGCGCAGACGGCATTCTCGGCCCTCGTGAATACCGGCCATTGAT CGACCAGCCCAGGTCTGGCAAAGCTGTAAGAACGTGAAGGTGATGGCTCGCCATAGGGTGC GCTTCGCGTACTCCAACACTGCTGCCACACCAGTCTGTCATCGTCGCCCGCAGCTGACGCCG GTGTAAGGTGATCTTCACGTCCTGTTGACGTGAAAATGACCTTGTGTTTCGACGCCCTCGCCGG GATTTTCTTGTGCGCTGGTGAACAGGGCAGAGGCCGGCTGCTGTTGGCATCGCTCGCATCG TGTCCGGCCACGGCGCAATATCGAACAGGAAAGCTGCATTCTGATCTGCTGTTCGTGTGTT TCAGCAACGCGGCCCTGCTTGGCCTGCTGACCTGTTGCAAGGTCTCGCCGGCGGTTTCGCT TCTTGGTCGTATAGTTCTCGCGTGTGATGGTCATCGACTTCGCCAACCTGCCGCTCTGTT CAAGACGACGCCAACGCTCCACGGCCGCGATGGCGGGCAGGGCAGGGGAGGCCAGTTGAC GCTGCGCTCGATCTGGCGTAGCTGCTGGACCATCGAGCCGACGGACTGGAAGGTTTGC GGGCGCACGCATGACGGTGCCTGCGATGGTGGCTTCCGCAACAGTCTGCGATTCACCCCTCTG GATCAGTTCTGCTGTATGCCCTCCGGTCAAAGCTGCTGATTCACCCCTCTGCGGGATG CCCGACTCACGGGGGGGCAATGTCCTTATCTCGTATTGACCCCTGGTCAAAG GATAATCCACCTTATCGCAATGAAGTGGTCCCTGAGACCGTCTGCCGCTCTCGTACTTGG TATTCCGAATCTGCCCTGCAAGAATACAGCGACCCCTGCCAAATACTTGCCTGGGCCCTCG CCTGAGAGCCAAAACACTTGTATGCCGAGAAGAAGTGGTGCCTCTGCTTGTGCCGGCATCGT CGCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCTGCTGCCGTTTICATAGGCTC CGCCCCCCTGACGAGCATCACAAAGACGCTCAAGTCAGAGGTGGCAGAACCCGACAGGACT ATAAAGATAACCAAGCGTTCCCCCTGGAAGCTCCCTCGTGCCTCTCTGTTCCGACCCCTGCC TACCGGATACCTGTCGCCCTTCTCCCTTCTGGGAAGCGTGGCGCTTCTCATAGCTCACGCTG GTATCTCAGTTGGTGTAGGTCCTGCTCCTGGCAACAGTGGCTGTGCAAGAACCCCGTT CCGACCGCTGCCCTTATCCGTAACTATCGTCTGAGTCCAACCCGGTAAGACACGACTATCG CACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGGCGGTGCTACAGAG TTGAAGTGGTGGCTAACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCCCTGCTG GCCAGTTACCTTCGAAAAGAGTTGGTAGCTCTGATCCGCAAACAAACCACCGCTGGTAG GTGGTTTTTTGTTGCAAGCAGCAGATTACGCCAGAAAAAGGATCTCAAGAAGATCCTTGA TCTTCTACGGGCTGACGCTCAGTGGAACGAAACTCACGTTAAGGGATTGGTATGAGAT TATCAAAAAGGATCTTACCTAGATCTTTAAATAAAATGAAGTTTAAATCAATCTAAAGTAT ATATGAGTAACCTGGTCTGACAGTTAACATGCTTCACTAGTGGACCCATATCTCAGCGATCT TCTATTCTCATCCATAGTGGCTGACTCCCCCTGCTGAGATAACTACGATACGGGAGGG ACCATCTGGCCCCAGTGTCAATGATACCGCGAGAACCCACGCTCACCGGCTCCAGTT CAATAAACCAGCCAGCGGAAGGGCGAGCGCAGAGTGGCTCTGCAACTTTATCCGCTCC CAGTCTATTAAATTGTTGCCGGAGCTAGAGTAAGTAGTTGCGCAGTTAATAGTTGCG GTTGCATTGCTACAGGATCTGGTGTACGCTCGTGTGTTAGGTTATGGCAGCACTG TCCCACAGTCAAGGCAGTTACATGATCCCCATGTTGCAAAAAAGCGGTTAGCTCTCG CCTCCGATCGTGTGAGAAGTGGCAGTGGTGTACGCTCGTGTGTTAGGTTATGGCAG AATTCTTACTGTCTGAGATCCAGTTCGATGTAACCCACTCGTGCACCCAACTGATCT CAGCATC TTTTACTTCAACAGCTTCTGGGTGAGCAAAACAGGAAGGCCAAATGCCGAAAAAGGG TAAGGGCGACACGAAATGTTGAATACTCATACTCTCCTTTCAATATTGAAGCATTT GGGTATTGTCATGAGCGGATACATATTGAATGTTAGAAAATAACAAATAGGGT GCGCACGAATTGCCAGCGCTGCCATTGGGTGAGGCCGTTGCGGGGAGGG CCTGGGGGATGGGAGGCCGTTAGCGGGGGGGGGGGTTGAGAAGGGGGGG CCTGGCGTGCCTGACCGCAGAGGGCGCAGGGCTGGTTAAAACAAGGTT TTAAAAGCAGGTTAAAGACAGGTTAGCGGTGGCGAAAAACGGGCG GGAAACCCCTGCAAATGTCATAGGTGCGCCCTCATCTGTCAGTACT CCCCTCAAGTGTCAAGGATCGGCCCTCATCTGTCAGTACTCGGCC CAGGGCACTTATCCCCAGGCTTGTCCACATCATCTGTTGGAAACTCG CGTAAACATCAGGCGTCTGCACTCTGCTG GCCGATTGCGAGGCTGGCCAGCTCACGTCGCCGGCGAAATCG AGGCTGCCCTCATCTGTC ACGCCGCCGGGTGAGTCGCCCTCAAGTGTCAACGTCGCC AAAGTTCGCCGAGGTATCCACAACGCCGGCGGGTGTCTGC ACACGGCTTCGACGCC TTCTGGCGCTTGTGAGGGCATAGACGCCGCCAGCCCAGGCC AGCTGCCGGTGGGAGCTGTTGGCTGGCTGGCAGGATATATT GTGGTCAAACAAATTGAC GCTTAGACAATTAAACACATTGCCAGT TTAAATGTA CTGGGTGGATGCA GACTGGGCCCCA C</p>
pTU-7	<p>TCTGTGAAGACAAGAGCGAATTCCATATGGGAGTGAGACCCGAGCTGGCACGACAGGTTGCC CTGGAAAGCGGGCAGTGAAGCGCAACCGCAATTAAATGTTGAGTTGCTACTCATTAGGCACCC CAGGCCCTTACACTTTATGCTTCCGGCTGTTGAGTTGAGCTCATTAGGCACCC CAGGCCCTGAGCTATGACCATGATTACGCCAACGCTGCTGATGCC CAGCTGACTCGACTGGAAAACCCCTGGCGTT CCCAACTTAATGCCCTTGCAAGCACATCCCCCTTCTGCCAGCTGGCGTAATAGCG AAAGAGGCC ACCGATGCCCTCCCAACAGTTGCCAGCCTGAATGGCG GAATGGCGCTGATGCCGTT CCTTACGCATCTGCGGTATTTCACACC GCATATGGTCACTCTCAGTACAATCTGCTCTG CGCATAGTTAACGCGCCAGGCCAACACCC GCTGACGCC TCCCGCATCCGCTTACAGACAAGCTGAGCGCTCAGCTGA CAGCTGTTGCTCTGCA GAGCTGAGGTT GTTTAAACTATCAGTTGAGCGGATATATTGGCGGG TAAACCTAAGAGAAAAGAGCG GTT GAATAATCGGATATTAAAAGGGCGTAAAAGGG TATCCGCTCCATTG TGTATGT GCATGCC ACCACAGGGTTCCCCAGATCAGGCCTGGCTG TAACCCCCAGCCGA ACTGACCC CAGGC</p>

	ACTCTTCGCGAGGCTTCGCCGACCTGCTCGGCCACTTCTCACGGGGTGGAAATCCGATCCGCAC ATGAGGCGGAAGGTTCCAGCTTGAGCGGGTACGGCTCCCGGTGCGAGCTGAAATAGTCGAACAT CCGTCGGGGCAGCTGGCAGACAGCTGGTACTCTCCCATATGAATTCTGTGAGTGGTCGCCAG CAAACAGCACGACGATTCTCTCGCATCAGGACCTGGCAACGGACGTTCTGCCACGGTCC AGGACGCGAAGCGGTGCAAGCAGCACCGATTCCAGGTGCCAACCGCGGTGCGAGCTGAAGC CCATCGCCGCTGCCTGTAGGCGCAGACAGGCTTCCTCCGGCTTCTGTGAAACCGGCCATTGATC GACCAGCCCAGGTCTGGCAAACAGCTCGTAGAACCTGTGAGGTGATGGCTGCCGATAGGGTGC GCTTCGCGTACTCCAAACACTGTGCCACACCGAGTCTGTGATCGTCGCCGAGCTCGACGCCG GTGTAGGTGATCTTACGTCTTGTGACGTGAAAATGACCTGTTTGCAAGGCCCTCGCG GATTTCCTTGTGCGGTGGTGAACAGGGCAGAGCGGGCGTGTGTTGGCATCGCTCGCATCG TGTCCGGCACGGCGAATATCGAACAGGAAAGCTGCATTCTGATCTGCTGCTCGTGTGTT TCAGCAACCGCGCCTGCTTGGCCTCGTGCACCTGTTGCCAGGTCTCGCCGGTTTCGCT TCTTGGTCTGCTAGTCTCGCGTGTGATGGTCTGACTCGACTTCGCCAAACCTGCCCTCTGTT CAAGACGACGCCAACGCTCCACGGCGGATGGCGGGCAGGGCAGGGGAGGCCATGAC GCTGCGCGCTGATCTGGCGACGATGGCTGGCTGAGTGGTCTGGCATCTCGGCCG GGGGCGCACGATGACGGTGCCTGCGATGGTTGGCATCTCGGCCGAAACCCCGCGT GATCAGTTCTGCTGTATGCCCTCCGGTCAAACGTCCGATTCACTCACCCTCTTGCAGGGATTGC CCGACTCACGCCGGGCAATGTGCCATTCTGTGATTTGACCCGCTGGTGCCTGGTCCA GATAATCCACCTTATCGGCAATGAAGTCGGTCCCAGACCGTCTGGCGTCTTCGCTTCTCGACTTGG TATTCCGAATCTTGCCTGACGAATACCAAGCGACCCCTTGCCTAAACTTGTCCGTTGGCGCTCG CCTGAGAGCCAAAACACTTGTGCGGAAGAAGTCGGTGCCTCTGTGTTGCGCTTCCG CGCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCGTGTGGCGTTTCCATAGGCTC CGCCCCCTGACGACATCACAAAATCGACGCTCAAGTCAAGTCAAGGCTGCCGAAACCCGACAGGACT ATAAGATACCAGGGCTTCCCCCTGGAGCTCCCTGTGCTCAAAGCTGGGCTGTGACGAACCCCCCGT TACCGGATAACCTGTCCGCCCTTCTCCCTGGAGCGCTGGCGTTCTCATAGCTCACCGCTGTAG GTATCTCAGTTGGTGTAGGTCGTTGCTCAAAGCTGGGCTGTGACGAACAGCAGTATCGC CCGACCGCTGCCATTATCCGTAACATCGTCTGAGTCAAACCCGTAAGACAGCACTTATCGC CACTGGCAGCAGCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGTCGGTGTACAGAGTT TTGAAGTGGTGGCTAAACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCGCTCTGCTGAA GCCAGTTACCTCGGAAAAAGAGTTGGTAGCTCTGTGAGTCCGGCAAACAAACCCACCGCTGGTAGCG GTGGTTTTTGTGTTGCAAGCAGCAGATTACCGCAGAAAAAAAGGATCTCAAGAAGATCTTGA TCTTTCTACGGGCTGACGCTCACTGGTAAGGTTTAAATTAAATTAAGGATTTAAATCAATCTAAAGTAT TATCAAAAGGATCTTACCTAGATCTTAAATTAAGGATTTAAATCAATCTAAAGTAT ATATGAGTAAACTTGGTGTACAGTTACCAATGCTTAATCAGTGGGACCTATCTCAGCGATCTG TCTATTCTCATCCATAGTGTGACTCCCCGTGTTAGATAACTACGATACGGGAGGGCTT ACCATCTGCCCACTGCTGCAATGATACCGCAGAACCCACGCTACCGGCTCCAGGATTATCAG CAATAAAACAGCCAGCCGAAGGGCGAGCCAGAAGTGGTCTGCAACTTATCCGCCATC CAGTCTATTAAATTGGTCCGGAGCTAGAGTAAGTACTTCGCAAGTAAACTTGTGAAACCGTT GTGCGATTGCTCACAGGATCGTGTGTCAGCTCGTCTGTTGGTATGGCTTCATTGAGCTCCGGT TCCCAACGATCAAGGGAGTTACATGATCCCCATGTTGCAAAAAGCGGGTAGCTCTTGGT CCTCGATGTTGCAAGTAAGTGGCCAGTGTATCACTCATGGTTATGGCAGCA AATTCTTACTGTCATGCCATCCGTAAGATGTTTCTGTGACTGGTAGTACTCAACCAAGTCA TTCTGAGAATAGTGTATCGCGCACCGAGTTGCTCTGCCGGCTCAATACGGGATAATACCGC GCCACATAGCAGAACTTAAAGTGTCTCATATTGAAACAGTCTTCCGGGGCGAAACCTCTCAAG GATCTTACCGCTGTTGAGATCCAGGCTGATGTAACCCACTCGCAGCACCAACTGATCTCAGCATC TTTACTTACCCAGCGTTCTGGTGAAGCAACTTATGGCAGGAAATGGCAGGAAAGGAA TAAGGGCGACAGGAAATGTTGAACTACTCATACTCTTCTTCTCAATTATTGAAGCATTATCA GGGTTATTGTCATGAGCGGATACATATTGAATGTTAGAAAATAACAAATAGGGGTTCC GCGCACGAATTGGCCAGCGCTGCCATTGGGGTAGGGCGTTCGCGGCCAGGGGGCAGCC CCTGGGGGGATGGGAGGGCCCGCTTAGCGGGCGGGAGGGTTGAGAAGGGGGGCACCCCCCT TCGGCGTGCAGCGGTACCGCAGGGCGAGCCCTGGTAAAACAAGGTTATAAATATTGGT TAAAAGCAGGTTAAAAGACAGGGTAGCGGTGGCCAAAAACGGGCGAAACCCCTGCAATGCT GGATTTCTGCCGTGAGCACGGCCCTCAAATGTCATAGGTGCGCCCTCATGTCAGCACTCTG CCCCCTCAAGTGTCAAGGATCGCCCTCATGTCAGTAGTCAGTGGCCCTCAAGTGTCAATACCG AGGGCACTTATCCCAGGCTGTCACATCATGTTGGAAACTCGCGTAAATCAGGCGTTTC GCCGATTGCGAGGCTGCCAGCTCACGTCGCCCGAAATCGAGCTGCCCTCATCTGTCA ACGCCGCCGGGTGAGTCGGCCCTCAAGTGTCAACGTCAGTCCGCCCTCATCTGTCA AAGTTTCCCGAGGTATCCACAACGCCGGCCGGGTGTCGACACGGCTTCGACAGGCGT TTCTGGCGCGTTGCAAGGGCATAGACGGCCAGGCCAGCGAGGGCAACCAGCCGGT AGCGTCGCAAAGGAGATCTGACTGATGGCTGCTGTATGAGTGGTAGTTGTGCG AGCTGCCGGTGGGGAGCTGTTGGCTGGCTGGCAGGGATATAATTGTTGAGTAAACAAATTGAC GCTTAGACAACCTAATAACACATTGCGGACGTTTTAATGACTGGGGTGGATGCA GTTAGAGCAGGTTTAACTTACGTCGACGGTCTCAGCTAGACTTGTCTCTGCACGA GAGGTTTGGCG
pTU-9	TCTGTGAAGACAACCTCGAATTCCATATGGGAGTGAAGCCGAGCTGGCACGACAGGTTGCCGA CTGGAAAGGGCGAGTCAGCGCAACGCAATTAAATGTGAGTTAGCTCACTCATTAGGCACCCAGG CTTACACTTATGCTCCGGCTGTATGTTGTTGAAATTGTGAGCGGATAACAATTTCACACAG GAAACAGCTATGACCATGATTACGCCAAGCTGCTGATGCCAGGTGCAACTCTAGAGGATCCCCG GGTACCGAGCTGAACTTACTGGCCGTGTTTACAACAGCTGTCAGTGGGAAACCCCTGGCGTTA CCCAACTTAACTGCCCTGCAAGCACCCCCCTTCCGCCAGTGGCGTAATAGCGGAAGAGGCCCG ACCGATGCCCTTCCCAACAGTGTGCGAGCCTGTAATGGCGAATGGCGCTGATGCGGTATTTC CCTTACGCTATGTCAGCGGTATTTCACACCGCATATGGTCAGTCACTCTAGTACAATCTGCTCTGATG CGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCCCTGACGGGCTTGTCTG TCCCGCATCCGTTACAGACAAGCTGACGGTCTCAGCTAGACTTGTCTCTGCACGA GTTAAACTATCAGTGTGTTGACAGGATATAATTGGCGGGTAAACCTAAGAGAAAAGAGCGTTTATT GAATAATCGGATATTAAAAGGGCGTAAAAGGTTTATCCGTTGCTCCATTGTTATGTCAGGCCA ACCACAGGGTTCCCCAGATCAGGCCGTGGCTGAAACCCCGAGCCGAACCTGACCCCAACAGGC CTCTGGCTTGCATGCACTGACCCAGGGCTGTTCCACCCAGGGCGTGGCTGCC AACTCTCCGACCCCTTCCCCACCTCTCCCCCAACTTCTTACCCCCCTCCAACTCC

	CATGAGGCAGGAAGGTTCCAGCTTGAGCGGGTACGGCTCCGGTGCAGCTGAAATAGTCGAACA TCCGTGGGCCGTCGGCAGACAGCTTGTGGTACTTCTCCCATAATGAATTTCGTGTAGTGGTCGCCA GCAAACAGCACGACGATTTCCTCGTGTGATCAGGACCTGGCAACGGGACGTTTCTTGCACGGTC CAGGACGGCGGAAGCGGTGAGCAGCGACACCAGGATTCAGGTGCCCACAGCGGGTGGACGTGAAG CCCCATCGCCGTCGCTGTAGCGCAGAGCATTCTCGGCCCTCGTGTAAATACCGGCCATTGAT CGACCAGCCCAGGTCCCTGGCAAAGCTGTAGAACGTGAAGGTGATCGCTCGCCGATAGGGTGC GCTTCGCGTACTCCAACACTGCTGCCACACCAGTCTGTATCGTGCAGCGCCCGCAGCTGACGCCG GTGTAGGTGATCTCACGTCCTGTGACGTGAAAATGACCTTGTGTCAGCGCCTCGCG GATTTCTTGTGCGCGTGTGAAACAGGGCAGAGCAGGGCGTGTGTCAGCGCATCGCATCG TGTGCGGCCACGGCGAATATCGAACAAAGGAAAGCTGCAATTCTGTGATCTGCTGTTGTGTT TCAGCAACGGCGCTGCTGGCCTCGTGCACCTGTTGCAAGGTCTCGCCGGGGTTTTCGCT TCTTGGTCGTATAGTCTCGTGTGATGGTCATCGACTTCGCAACCTGCCCCTCCCTGTT CAAGACGCGAACGCTCCACGGCGCGATGGCGCGGGCAGGGCAGGGGAGGCCAGTTGAC GCTGCGCTCGATCTGGCGTAGCTGCTGACCATCGAGCCGACGGACTGGAAGGTTGCG GGGGCGACGCATGACGGTGCAGCTGCGATGGTTTCCGACATCCTCGCCGAAAACCCCGCCTG GATCAGTTCTGCTGTATGCCCTCCGTCAAACGTCCGATTCACTCACCCTCCTGCGGGATTG CCCAGACTCACGCCGGGCAATGTGCCCTTATTCTGTATTGACCCGCCGCTGTGCGCTTGTG GATAATCCACCTTATCGGAATGAAGTCGGTCCCGTAGACCGCAGCCCTGCCAAATACTTGC CCTTGGCCCTGACGAATACCGCAGGTTAGCAGAGCAGGGTATGAGGCGTCTACAGGTT TATTCCGAATCTGCCCTGACGAATACCGCAGGTTAGCAGAGCAGGGTATGAGGCGTCT CCTGAGAGGAAAACACTTGTGCGGAAAGAAGCTGGTCCGCTCTGCTGTGCGCTTGTG CGCCACATCTAGGATCTGCCAGGAACCGTAAAAGGCCGCGTTGCTGCGTTTTCATAGGTC CGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAGTGGCGAAACCCGACAGGACT ATAAAGATAACCAAGCGTTCCCCCTGGAAGCTCCCTCGTGCCTCTCTGTTCTCATAGCTCACG TACCGGATAACCTGTCCGCCCTTCTCCCTCGGAAGCGTGGCCTTCTCATAGCTCACGCT GTATCTCAGTTGGTGTAGTCGTTGCTCCAAGCTGGGCTGTGTCAGCAACCCCCGTTG CCGACCGCTGCCCTTATCCGTAACTATCGTCTTGAGTCAACCCGGTAAGACACGACTTATCG CACTGGCAGCAGCCACTGTTGAAACAGGATTAGCAGAGCAGGGTATGAGGCGTCTACAGGTT TTGAAGTGGTGGCCTAAACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCGCTGCT GCCAGTTACCTTGGGAAAGAGTTGGTAGCTTGTGATCCGCAACAAACCCAGGCTGGTAGCG GTGGTTTTTTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCT TCTTTCTACGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTGGTATGAGAT TATCAAAAGGATCTCACCTAGATCCTTTAAATAAAAAATGAAGTTAAATCAATCTAAAGTAT ATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCAACCTATCAG TCTATTTCGTTCATCCATAGTGTGCTGACTCCCCGCTGTAGATAACTACGATA ACCATCTGGCCCTAGTGTGCAATGATACCGCAGAACCGCTCAGGCTCCAGATTTATCG CAATAAACAGCCAGCGGAAGGGCGAGCGCAGAAGTGGCCTGCAACTTATCGCCTCCATC CAGTCTATTAAATTGTTGGGAAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTGCG GTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTGTGGTATGGCTTCAAGCTCG TCCCAACGATCAAGGCAGTTACATGATCCCCATGTTGCAAAAAAGCGTTAGCTCTCG CCTCCGATCGTTGCAAGTAAGTTGGCGCAGTGTATCACTCATGGTTATGGCAGCACTGC AATTCTCTACTGTGTCATGCCATCCGTAAGATGTTCTGTGACTGGTAGACTCAACCAAGTC TTCTGAGAATAGTGTATGGCGGACGGAGTTGCTCTGCCGGCGTCAATACGGGATAACCGC GCCACATAGCAGAACCTTAAAGCTGTCATCATGGAAAAGCTTCTCGGGCGAAAACCTCT GATCTTACCGCTGTTGAGATCCGATGTTGAGCAAAACAGGAAAGGCAAAATGCG TTTACTTCAACAGCGTCTGGGTAGCAACTCTTCTTTCAATATTGAAGCATTATCA GGGTTATTGTCATGAGGGATAACATATTGAATGTTAGAAAAATAACAAATAGGGT GCGCACGAATTGGCCAGCCTGCCATTGGGGTGAAGCGCTCGCGGGCGAGGGCGCAGCC CTTGGGGGATGGGAGGCCGCTTAAGCGGGCGGGAGGGTTCGAGAAGGGGGGCA TCGGCGTGCCTGACGGCAGCCGAGGGCGCAGGGCTGGTAAAACAAGGTTATAATATTGGT TTAAAAGCAGGTTAAAGACAGGTTAGCGGGCGAAAACGGGCGAAAACCTGCAAAATGCT GGATTTCCTGTCGAGCAGCCCTCAATGTCATAGTGTGCGCCCTCATCTGTCAGCACT CCCCCTCAAGTGTCAAGGATGCGCCCCCTCATCTGTCAGTAGTCGCGCCCCCTCAAGTGT CAGGGCACTTATCCCAGGCTTGTCCACATCATCTGTGGAAAACTCGCGTAAAATCAG GCCATTGCGAGGCTGGCAGCTCCACGTCGCCGCCGAAATCAGGCTGCCCTCATCTG ACGCCCGCCGGGTGAGTCGGCCCTCAAGTGTCAACGTCCGCCCTCATCTGTCAGTG AAGTTTCCGCGAGGTATCCACAACGCCGGCGGGCGGGTGTCTGCACACGGCT TTCTGGCGGTTGCAAGGCCATAGACGGCCGCGAGCCAGCAGGGCGAGGGCAACCAAGCCGG AGCCTGCGAAAGGAGATCTGACTGATGGCTGCCGTATCGAGTGGTATTTGCG AGCTCGCGGGTGGGAGCTGTTGGCTGGCAGGATATTGTTGTAACAAATTGAC GCTTAGACAACCTAATAACACATTGCCAGCTTTAATGACTGGGTGGATGCA GTGGCCCA
Destination vector	GGGATGGAACGACGCGTAACTCACGTTAAGGGATTGGTCATGAGCTTGCGCCGCTCCGTCAG TCAGCGTAATGCTCTGCCAGTGTACAACCAATAACCAATTGATTAGAAAAACTCATCGAGCA TCAAATGAAACTGCAATTATTATCATATCAGGATTATCAATACCATATTGAAAAGCCGTTCTG TAATGAAGGAGAAAACCTACCGAGGGCAGTTCCATAGGATGCGCAAGATCTGGTATCGGTCTG TICCGACTCTGTCACACATCAATACAACCTATTAAATTCTCCCTCGTCAAATAAGGTTATCAAGT AGAAATCACCAGTGAAGTGAAGACTGAATCCGGTGAAGAATGGCAAAGTTATGCAATTCT CTTCAAGAGCCAGCCATTACGCTCGTCAATGAAATCACTCGCATCAACCAACCGTTATTCA TTCGTGATTGCGCTGAGCGAGACGAAATACGCGATCGCTGTAAAAGGACAATTACAAACAGGA ATCGAATGCAACCGGGCGCAGGAACACTGCCAGCGCATCAACAAATTGACCATCTG TCTCTAATACCTGGAATGCTGTTTCTGGGGATCGCAGTGGTAGTAACCATGCA GTACGGATAAAATGCTGATGGTCGGAAGAGGGCATAAAATTCCGTCAGGCCAGTTAGTCT TCATCGTAAACATCATTGCAACGCTACCTTGCATGTTGAGAAACAAACTCTGGCGCATCG TICCGCATACAAGCGATAGATGTCGCACTGATTGCCAGATTATCGCGAGGCCATTATACCA TATAAATCAGCATCCATGTTGAAATTAAATGCCGCTCGACGTTCCGGTGAATATGGCT ACACCCCTGTATTACTGTTATGTAAGCAGACAGTTATTGTTCATGATGATATTATTATCTT GTGCAATGTAACATCAGAGATTGAGACACGGCCATGCTAGCTCAGCAAAAGGCCAGGAACC

	ATGATAAAAGCTTTCAAGGATCTGGGTGGCGAAGTCGTGTTAACGCCAGAGTCAGCCATATGGA AACGACAGGAACAAGATTGAAGCCGTGCAATTAGAGGACGGTCGCAGGTTCTGACGCAAGCCG TCGCGTCAAATCAGATGTGGTTCATACCTATCCGCACCTGTTAACGCCAGCACCCCTGCCGCCGTTA AGCAGTCCAACAAACTGCAGACTAACGGCATGAGTAACTCTGTGCTCTATTTGGTTGA ATCACCATCATGATCAGCTCGCATCACACGGTTGTTCGGCCGTTACCGCGAGCTGTTG ACGAAATTAAATCATGATGGCCTCGCAGAGGACTTCTCACITTTATCTGCACGCCCTGTGTC CGGATTGTCACTGGCGCTGAAGGTGCGGAGTTACTATGTGTTGGCGCCGTCGGCATTAA GGCACCGCAACCTCGACTGGACGGTGGAGGGGCAAACACTACCGCACCGTATTTTGCGTACCT TGAGCAGCATTACATGCCTGGCTTACGGAGTCAGCTGGTCAGCACCGGATTTACGCCGTTG ATTTCGCGACCAGCTTAATGCCATGGCTCAGCCTTCTGTGGAGCGCTTACCCAGA GCCGCTGGTTCGGCCGATAACCGGATAAAACCTACTAATCTCACCTGGTCGGCGCAGGC ACGCATCCCGGCCGAGCATTCCCTGGCGTATCGGCTCGGCAAACAGCAGCACAGGTTGATGCT GGAGGATCTGATTTGAATAATCCGTCGTTACTCAATCATGCCGTCGAAACAGATGGCAGTGGCT GAAAAGTTTGCAGCAGCTCAAAGTTATTGATGCAAAACCCGGCGCAGCGTACTGATGCTCTA CGCCTGGTGCCGCCATTGTGACGATGTTATTGACGATCAGACGCTGGGCTTCAAGGCCGGCAGC CTGCCTTACAAACGCCGAACAACGCTGTGATGCAACTTGAGATGAAAACCGGCCAGGCCTATGCA GGATCGCAGATGCACGAACCGCGTTGCGGCTTCAAGGAAGTGGCTATGGCTATGATATGCG CCCGCTTACCGCGTTGATCATCTGGAAAGGCTTGCATGGATGTCAGCGAAGCGCAATACAGCC AACTGGATGATAACGCTGCCATTGTCATACGGTGGACGGCTGTCAGGCTTGTGATGATGGCGCA ATCATGGCGTGGGATAACCGCACGCTGGACCGCGCTGACCTGGCTTGGGCTGGCATTTCAAGT GACCAATATTGCTCGCATATTGAGCAGTCGCGATGCCATGCGGCGCTGTATTCAGCCTGG GGCTGGAGCATGAAGGCTGAAACAAAGAGAATTATGCGGACCTGAAAACCGCTCAGGCGCTGAGC CGTATGCCCGTCTGGTGGCAGGAAGCAGAACCTACTATTGCTGCCCCAGCGCTGG AGGGTTGCCCTGCGTCCGCCTGGCAATCGTACGGCGAACAGGTTACCGAAAATAGGTG TCAAAGTTGAACAGGCCGGTCAGCAAGCCTGGGATCAGCGCAGTCACGACCAACGCCGAAAAA TTAACCGCTGCTGGCCCTCTGGTCAAGGCCCTACTTCCCAGTGGGGCTCATCTCCCCG CCCTGCGCATCTGGCAGCGCCCTAGCGGCATGTCTTCCCAGCGCTGCTGAAGTTT GACAGGGCGGCCATAGAGGAAGCGAAAAGAACACAACCTCTTGGCCCTGACGGCGTATG CATACGCTGGCGATTAGCACGAAACAA
pLink-1	GGGTACCGAGCTCGAATTCACTGGCCGTGTTTACAACGTCGTGACTGGAAAACCCCTGGCGTT ACCCAACCTTAATGCCCTGCGCACATCCCCCTTCCGCAGCTGGCGTAATAGCGAAGAGGCCG CACCGATGCCCTTCCAAACAGTTGCCAGCCTGAATGGCGAATGGCCCTGATGCCGTATTT TCCTTACGCATCTGTGCGGTATTTCACACCGCATATGGCACTCTCAGTACAATCTGCTCTGATG CCGCATAGTTAACGCCAGCCCCGACACCCGCAACACCCGCTGACGCCCTGACGGGCTTGTCTG CTCCCGCATCCGTTACAGACAAGCTGTGACCATCTCGGGAGCTGATGTCAGAGGTTTTC ACCGTCATACCGAAACCGCGAGTCGAAAGGCCCTCGTGAATGCCCTATTTTATAGGTTAATG CATGATAATAATGGTTCTTACAGCTGAGCTGAAAGGCCCTCGTGAATGCCCTATTGTCAGAGGTT TTGTTTATTTCTAAATACATTCAAAATATGTATCCGCTCATGAGACAATAACCGTATAATGCTT CAATAATATTGAAAAGAGAAGTATGCGCTACGCAACTGGTCCAGAACACCTTGACCGAACGCAG CGGTGTAACGGCGAGTGGCGTTTATGGCTTGTATGACTGTTTTGGGGTACAGTCTAT GCCTGGGCATCCAAGCAGCAAGCGCGTTACGCCGTGGGCTGATGTTATGGAGCAGCA ACGATGTTACGCAGCAGGGCAGTCGCCCTAAACAAAGTTAAACATCATGAGGGAAGCGGTGATC GCCGAAGTATCGACTACTAGAGGTAGTGGCTCATCGAGGCCATCTGAACCGAACGCTT GCTGGCGTACATTGACGGCTCCCGAGTGGATGGCGGCCGTAAGGCCACACAGTGTATTGATT TGCTGGTTACGGTACGGTAAAGGCTTGTGATGAAACAACCGCGAGCTTGTATCAACGACCTT GAAACTCGGCTTCCCGTGGAGAGAGCGAGATTCTCGCGCTGTAGAAGTCACCATTTGTTGTC CGACGACATATTCCGTGGCGTTATCCAGCTAACGCGAACCTGCAATTGGAGAATGGCAGCGCA ATGACATTCTGCAGGTATCTCGAGGCCAGCCACGATCGACATTGATCTGGCTATCTGCTGACAA AAGCAAGAGAACATAGCGTTGCTTGTAGGTCCAGCGGCCGAGGAACCTTGTATCCGGTCT GAACAGGATCTATTGAGGCCTAAATGAAACCTTAACGCTATGGAAACTCGCCGCCACTGGC TGGCGATGAGCGAAATGTAGTGGCTTACGTTGCTCCGCATTGGTACACGCCAGTAACCGGCAAAA TCGCGCCGAAGGATGCGTGGCGACTGGCAATGGAGGCCCTGCCAGTATCAGCCCGT ATACTTGAAGCTACAGCGGTTATCTTGGACAAGAAGATCGCTGGCTCGCGCAGATCA GTTGGAAGAATTGTCACACTGAAAGCGAGATCACCAAGGTTAGTCGGCAAAATACTGTCAG ACCAAGTTACTCATATACTTAGATTGATTTAAACCTCATTTTAATTAAAGGATCTAGGT GAAGATCTTTTGATAATCTCATGACCAAAATCCCTAACGTGAGTTTCTGCTTCACTGAGCGTC AGACCCCGTAGAAAGATCAAAGGATCTCTTGAGATCTTTCTGCGCTAATCTGCTGCTT GCAAAACAAAAACCCCGCTACCGAGCGGTGGTTGTTGCCGATCAAGAGCTACCAACTCTT TCCGAAGGTAATGGCTTACAGCAGCGCAGATACCAAAACTGTCTCTAGTGTAGCGTAGTT AGGCCACCACTTCAAGAACACTCTGTAGCACCGCCCTACATACCTCGCTGTCAATCTGTT GGCTGCTGGCAGTGGCGATAAGCTGTCTTACCGGGTTGACTCAAGCGATAGTTACCGG AGGCGCAGCGGTGGCTGAACGGGGGGTCTGTCACACAGCCAGCTGGAGCGAACGACCTA CACCGAACTGAGATAACCTACAGCGTGAGCTATGAGAAAGGCCACGCTCCGAAGGGAGAAAGG CGGACAGGTATCCGTAAGCGCAGGGTGGAACAGGGAGAGCGCACGAGGGAGCTCCAGGGGG AAACGCCCTGGTATCTTATAGCTCTGCGGTTGCGCACCTGACTTGAGCGTCGATTTGTG ATGCTCGTCAAGGGGGCGAGCCTATGGAAAAACGCCAGCAACCGGCCCTTTACGGTTCTG CTTTTGCTGGCTTTGCTCACATGTTCTTCTGCTTACCCCTGATTCTGTGATAACCGTAT TACCGCCTTGTAGTGGCTGAGCTGATACCGCTGCCGCAGCGAACGCCAGCGCAGCGAGTCAGTGA GCGAGGAAGCGAAGAGCGCCCAATCGCAACCCGCCCTCCCGCGCCTGTCAGTCA TGCAGCTGGCACGACAGGTTCCCGACTGGAAAGCGGGAGTGTAGCGCAACGCAATTATGTGAG TTAGCTCACTCATAGGCACCCCAGGTTACACTTTATGCTTCCGGCTCGTATGTTGTTGGAAT TGTGAGCGATAACAATTTCACAGGAAACAGCTATGACCATGATTACGCCAAGCTTGCATGCC GCAGGTGCACTCTAGAGGATCCACCCCTTGAAGACAAGAGGATGACATGTGACCGAGGGATTG TCTTCGTTGAAACTTGGGACTTCAGATTCCACCCCTTGAAGCGTGGAAAGGCCCTCCGACCAAGG GAAGTGTGGATGAGGAAATGGAAGAGTTGCTCAGATGGGAGCTGCTTGTGTTTAC AAGAATGCCAGCTTATGCCGTAACCGCAGAGGAAGATTAGTGTGCAATCAGGAAAACGCCGAGA TGGATGTGGCTAAGAAGTTAAAGAACACTGAGGAGACAGAAATCAACTCAACACCTGCA ACTTCCCTAACAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCCAGGCCGAAAGAAAG

	TAAACGCCTAGCTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCAACCAAATGAAATAA CGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCTGGGCCAAA
pLink-2	GGGGATCCTCTAGAGTCGACCTGCAGGCATGCAAGCTTGGCGTAATCATGGTCATAGCTTTCC TGTGAAATTGTTATCCGCTCACAACTTCCACACAACATACAGGCGGAAGCATAAAGTGTAAAGC CTGGGGTGCCTAATGAGTGA GCTAACTCACATTAAATTGCGTGCCTCACTGCCGCTTCCAGTC GGAAACCTGTCGCCCCAGCTGCAATTATGAATCGGCCAACCGCGGGAGAGGCCGTTGCGTA TTGGCGCTCTCGCTCTCGCTCACTGACTCGCTCGCCTCGCTCGGCTGCGCAGCG GTATCAGCTCACTCAAAGGCCGTAATACGGTTATCCACAGAAATCAGGGATAACGCAGGAAAGAA CATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAAGGCCGTTGCTGGCGTTTCC ATAGGCTCCGCCCTGACGAGCATCACAAAATGACGCTCAAGTCAGAGTGCGAAACCCG ACAGGACTATAAGATAACCCAGCGTTCCCCCTGAAAGCTCCCTGTCGCGCTCTCTGTTCCCACC CTGCCGCTTACCGGATACCTGTCGCCCTTCTCCCTCGGAAAGCGTGCCTTCTCATAGCTCA CGCTTAGGGTATCTAGTCGGTGTAGGCTGCTCCAAAGCTGGGCTGTGCAACGAACCCCG CGTTCAGCCCCGACCGCTGCCCTTATCCGTAACATCGCTTGAGTCCAACCCGTAAGACACG ACTTATGCCACTGGCAGCAGCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGCGGTGCT ACAGAGTTCTGAAGTGGGCTAACTACGGCTACACTAGAAGGACAGTATTGATCTGGCT CTGCTGAAGCCAGTTACCTCGGAAAAAGAGTTGGTAGCTTGTATCCGGCAAACAAACCCG TGGTAGCGGTGGTTTTTGTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAG ATCCTTGTATTTCTACGGGCTGACGCTCACTGGAAACGAAACTACGTTAAGGGATTITGG TCATGAGGATTATAAAAGGATCTCACCTAGACCTTAAATTAAAAAGTAAAGTTAAATCAAT CTAAAGTATATGAGTAAATTGCTGACAGTTATTGCGCAGTACCTGGTGTATCTGCCCTT CACGTAGTGGACAATTCTCAACTGATCTGCGCGAGGCCAAGCGATCTTCTTGTCCAAG ATAAGCCTGTCTAGCTCAAGTATGACGGGCTGATACTGGGCCGGCAGCGCTCATTGCCAGT CGGCAGCGACATCCTCGCGCAGTTGCCGTTACTGCGCTGTACCAAATGCCGAAACGTA AGCACTACATTGCTCATGCCAGCCCAGTGGCGAGTCCATAGCGTTAAGGTTTCA AGCGCCTCAAATAGATCTGTCAGGAAACGGATCAAAGAGTTCCCGCGTGGACCTACCAA GGCAACGCTATGTTCTTGTGTTGCTGAGCAAGATGCCAGATCAATGTCGATCTGGCTGCTC GAAGATCTGCAAGAATGTCATGCCGCTGCCATTCTCAAAATTGCACTGCGCTTAGCTGGATA ACGCCACGGAAATGATCTGCGTGCACAAACATGGTACTTACAGCGCGGAGAATCTGCGCT CTCCAGGGAAAGCGAAGTTCCAAAAGGCTGTTGATCAAAGCTGCCGTTGTTCATCAAGC CTTACGGTACCGTAACCGCAAATCAATATCACTGTTGCGCTCAGGCCGATCCACTGCCGA GCCGTACAAATGTACGGCAGCAACGTCGGTTGAGATGCCGCTGATGACGCCAACTACCTCTG ATAGTTGAGTCGATACTCGCGATCACCGCTCCCTCATGATGTTAACTTGTGTTAGGGCAGC TGCCCTGCTGCCAACATCGTGTCTCCATAACATCAAACATCGACCCACGGCGTAACGCCGCTT GCTGCTGGATGCCGAGGCATAGACTGACCCCAAAAAAACAGTCATAACAAGGCGATGAAAC GCCACTGCGCCITTACCCGCTGCGTCCGGTCAAGGTTCTGGACCAGTGCCTGAGCGCATACT CTTCCCTTTCAATTATTGAGCATTATCAGGTTATGCTCATGAGCGGATACATATTGAA TGTATTAGAAAATAACAAATAGGGTTCCGCGCACATTCCCGAAAAGTGCACCTGACGTC TAAGAAACATTATTATCATGACATTAAACCTATAAAATAGCGTATCACGAGGCCCTTCGACTC GCGCCTCGGTGATGACGGTGAAACCTCTGACACATGCACTCCCGAGATGGTCACAGCTG TCTGTAAGCGGATGCCGGAGCAGACAAGCCGTCAGGGCGCTCAGCGGTGTTGGCGGTG CGGGGCTGGCTTAACTATGCCGATCAGAGCAGATTGACTGAGAGTCGACCATATGCCGTG AATAACCGCACAGATGCCAGGAGAAATACCGCATCAGGCCATTGCCATTAGCGTGC ACTGTTGGGAAGGGCAGCTGGCGGGCTTCTGCTATTACGCCAGTGGCGAAAGGGGATGT GCTGCAAGGCAGTTAACGGTAAACGCCAGGGTTTCCCACTGACGAGTTGTAACACGACGC CAGTAATTGAGCTGGTACCCCTTGAAGACAACACTAGAGGATGACATGTGACCGAGGGATTG TCTTCTGAAACTTGGGACTTCAGATTCCACCCCTTGAAGCGTGGAAAAGCCTTCCGACCAAGG GAAGTGTGGATGTAGAGGAAATGAAAGTTGCTCAGATGGGACTTGCTGATTGCTTCAC AAGAATGCCAGCTTATGCCGAAACGCCAGAGGAAATTAAGTGTCAATCAGGAAACGCCGAGA TGGATGTCGGTCAAGAAGTTAAAGAACCTGCAAGGAGACAGAAATCAAAACTCAACCCCTG ACTTCCCTCAAAAGCTCCACAGGAACACAGTAGGGAGGTGAAACACCAGGCCAAAGAAC TAAACGCCTAGCTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCAACAAATGAAATAA CGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCTGGGCCAAA
pLink-3	GGGTACCGAGCTCGAATTCACTGGCCGTTTACAACGTCGTACTGGAAAACCCCTGGCGTT ACCCAACTTAATGCCCTGCAAGCACATCCCCCTTCCGCAAGCTGGCGTAATAGCGAAGAGGCCG CACCGATGCCCTTCCAAACAGTTGCCAGCCTGAATGCCAATGCCCTGATGCCGTATTTC TCCTTACGCATCTGCGGTATTTCACACCGCATATGGCACTCTCAGTACAATCTGCTCTGATG CCGCACTAGTTAACCCAGCCCCGACACCCGCAACACCCGCTGACGCCCTGACGGGCTTGTCTG CTCCCGCATTCCGTTACAGACAAGCTGTGACCATCTCCGGAGCTGATGTCAAGGGTTTC ACCGTCATCACCGAAACGCCGAGCTGAAAGGCCCTGATGCCCTTATTTTATAGGTTAATGT CATGATAATAATGTTCTTACAGCTCAGGTGGCACTTCCGGGAAATGTCGCGGGAAACCCCTAT TGTGTTATTCTAAATACATTCAAAATATGTTACCGCTCATGAGACAATAACCTGATAATGCTT CAATAATATTGAAAAGGAAAGAGTATGCGCTACGCAACTGGTCAGAACCTTGCACGCAACGC CGGTGTTAACGGCGAGTGGCGTTTACGGCTGTTATGACTGTTTTGGGGTACAGTCTAT GCCTCGGCATCCAAGCAGCAAGCGCTTACGCCGGGATGTTGATGTTATGGAGCAGCA ACGATGTTACGCAGCAGGGCAGTCGCCCTAAACAAAGTTAAACATCATGAGGGAAAGCGGTGATC GCCGAAGTATGCACTATCAGAGGTTAGTGGCGTACGAGCGCCATCTGCAACCGACGTT GCTGCCGTACATTGACCGTAAAGGCTGATGAAACAAACGCCGAGCTTGTATCAACGACCTTGT TGCTGGTTACGGTACCGTAAAGGCTGATGAAACAAACGCCGAGCTTGTATCAACGACCTTGT GAAACTTCCGCTTCCCGTGGAGAGGAGGAGATTCTCCGCGCTGAGAAGTCACCATTTG CGACGACATCATTCCGTTACCGCTAAGCGCAACTGCAATTGGAGAATGGCAGCGCA ATGACATTCTGCAAGGTATCTGAGGCCAGCCACGATGCAATTGATCTGGCTATCTGCTGACA AAGCAAGAGAACATAGCGTTGCCCTGGTAGGTCCAGCGGGGAGGAACCTTGTATCCG GAACAGGATCTATTGAGGCCGCTAAATGAAACCTTAACGCTATGGAAACTCGCCGCCC TGGCGATGAGCGAAATGAGTGTGTTACGTTGCGCATTGGTACAGCGCAGTAACCGGCAAA TCGCCGAAAGGATGTCGCTGCCGACTGGCAATGGAGCGCCCTGCCGCCAGTATGCCG ATACTTGAAGCTAGACAGGCTTATCTGGACAAGAAGAGTCGCTTGGCCTGCCGAGATCA GTTGGAAGAATTGTCACACTACGTGAAAGGCGAGATCACCAAGGTAGTCGCAAATAACTGTCAG

	ACCAAGTTACTCATATATACTTAGATTGATTAAAACCTCATTAAATTAAAAGGATCTAGGT GAAGATCCTTTTGATAATCTCATGACCAAATCCCTAACGTGAGTTTCGTTCACTGAGCGTC AGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTCTGCGCGTAATCTGCTGCTT GCAAACAAAAAAACCACCGCTACCAGCGGGTTGTTGCGCGTAAGAGCTACCAACTCTTT TCCGAAGGTAACTGGCTCAGCAGAGCGCAGATAACCAAAACTGTCTTAGTGTAGCCGTAGTT AGGCCACCACTCAAGAACCTGTAGCAGCACCGCCTACATACCTCGCTGCTAATCCTGTTACCAAGT GGCTGCTGCCAGTGGCATAAGTCGTCTTACCGGGTTGACTCAAGACGATACTTACCGGATA AGGCCAGCGGTGGCTGAACGGGGGGTCGTGCACACAGCCCAGCTGGAGCGAACGACCTA CACCGAAGTACGAGATAACCTACAGCGTGAAGCTATGAGAAAGCGCCACGCTTCCGAAGGGGG CGGACAGGTATCCGTAAGCGCAGGGTCGAACAGGAGAGCGCAGCAGGGAGCTCCAGGGGG AAACCGCTGTATCTTTAGTCCTGCGGTTTGCACCTCTGACTTGAGCGTCGATTITGTG ATGCTCGTCAGGGGGGCGAGCCTATGGAAAACGCCAGAACCGGGCTTTACGGTTCTGG CCTTTGCTGGCCTTTGCTCACATGTTCTCGCTTACCCCTGATTCTGTGATAACCGTAT TACCGCCTTGAGTGAAGCTGATACCGCTCGCCGAGCGAACGACCGAGCGCAGTCAGTGA GCGAGGAAGCGGAAGAGGCCAATACGCAAACCGCTCTCCCCGCGCTTGGCGATTCAATTAA TGCAGCTGGCACGACAGTTCCGACTGGAAAGCGGGAGTGAAGCGAACGCAATTATGTGAG TTAGCTCACTATTAGGCACCCAGGTTTACACTTTATGCTTCCGGCTCGTATGTTGTGGAAT TGTGAGCGATAACAATTTCACACAGGAAACAGCTATGACCATGATTACGCCAAGCTTGATGCC GCAGGTCGACTTAGAGGATCCCTTGAAGACAATTACGAGGTGACATGTGACCGAGGGATT GTCTCGTGGAAACTTGGGACTTCAGATTCCACCCCTTGCAGCGTGGGACTTGTGATTGCTICA GGAAGTGTGGATGTAGAGGAATGAAAGTTGTTCTCATGGGACTTGTGATTGCTICA CAAGAATGCCAGCTTATGCCGAAACGAGAGGAAAGATTAGTGTGCAATCAGGAAAACGCCAG ATGGATGTGGCTAAGAACGTTAAAGAACCTGCAGGAGACAGAAATCAAACTCAAAACCTGAGA AACTTCCTCAACAAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCAGGCCAAAGAAA CTAAACGCCCTAGCTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCGAAACAAATGAAATA ACGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCCGGCCAAA
pLink-4	GGGGATCCTCTAGAGTCGACCTGCAGGCATGCAAGCTTGGCTAATCATGGTCATAGCTGTTCC TGTGAAATTGTTATCCGCTCAACATCCACAAACATACAGCGGGAAAGCATAAAGTGTAAAGC CTGGGGCTCTAATGAGTGAAGCTAACCTACATTAAATTGCGTGCCTCACTGCCCGCTTCCAGTC GGGAAACCTGTCGTGCCAGCTGCATTATGAATCGGCCAACCGCGGGAGAGGCCGTTGCGTA TTGGCGCTCTCCGTTCTCGCTCACTGACTCGCTCGCTCGGCTCGCTCGGCGAGCG GTATCAGCTACTCAAAGGGCTAATACGGTTACCCACAGAAATCAGGGATAACCGCAGGAAAGAA CATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAGGCCGTTGCTGGCTTTTCC ATAGGCTCCGCCCCCTGACGAGCATCACAAAATGACGCTCAAGTCAGAGGTGGCGAACCCG ACAGGACTATAAAGATAACCCAGGGTTTCCCTGGAAGGCTCCCTGTCGCTCTCTGTTCCGAC CTGCCGCTTACCGGATACTGTGCCGTTCTCCCTCGGAAAGCGTGGCGCTTCTCATAGCTCA CGCTAGGGTATCTAGTCCGGTAGGGCTGCTCGCTTACGCTGAGTCAACCCGTAAGACACG ACTTATGCCACTGGCAGCAGCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGCGGTGCT ACAGAGTTCTGAAGTGGGCTTAACACTACGGCTACACTAGAAGGACAGTATTGTTATCTGGCT CTGCTGAAGCCAGTTACCTCGGAAAAAGAGTTGTTAGCTTGTATCCGCAAACAAACCC TGGTAGCGGTGGTTTTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAG ATCCTTGTATCTTCTACGGGCTGACGCTCACTGGAAAGAAAACCTACGTTAAGGGATTITGG TCATGAGATTATCAAAGGATCTTCACTAGATCCTTAAATTAAAAAGTAAAGTTTAAATCAAT CTAAAGTATATGAGTAAACTTGTGCTGACAGTTATTGCGCAGTACCTGGGTGATCTCGCCTT CACGTAGTGGACAAATTCTCAACTGATCTGCGCGAGGCCAGCGATCTTCTTGTCCAAG ATAAGCCTGTCTAGCTCAAGTATGACGGGCTGATACTGGCCGGCAGCGCTCCATTGCCAGT CGGCAGCGACATCCTCGCGCAGTTGCCGGTACTGCCGCTGTACCAAATGCCGGACAACGTA AGCACTACATTGCTCATGCCAGCCCAGTCGGCGCGAGTCCATAGCGTTAAGGTTTCACTT AGCCTCTCAAATAGATCCTGTTGAGGACCGGATCAAAGAGTTCCCTCCGCGTGGACCTACCAA GCCAACGCTATGTTCTTGTGAGCAAGATGCCAGTCAATGTCGATCTGGCTGGCT GAAGACTCTGCAAGAATGTCATTGCCGCTGCCATTCTCAAATTGCGATTCTGCGCTTAGCTGGATA ACGCCACGGAAATGATGTCGCTGCAACAAATGGTACTTACAGCGCGAGAATCTCGCTCT CTCCAGGGAAAGCGGAAGTTCCAAAAGGCTGATCAAAGCTGCCGCTTGTGTTCATCAAGC CTTACGGTACCGTAACCGCAAATCAATATCACTGTTGCGCTCAGGCCGCTCCACTGCCG GCCGTACAAATGTACGGCAGCAACGTCGGTCCAGATGGCGCTGATGACGCCAACTACCTCTG ATAGTTGAGTCGATACTCGGCATACCGCTCCCTCATGATGTTAACTTGTGTTAGGGCGAC TGCCCTGCTGCCAACATCGTTGCTGCTCATAACATCAAACATCGACCCACGGCGTACGCCGCTT GCTGCTTGGATGCCGAGGCATAGACTGTAACCCAAAAAAACAGTCATAACAGCGCATGAAAACC GCCACTGCCGCTTACCGCCTGCGTTGCTGAGGCTGGGACCGACTGGCGTGGCGCATACT CTTCTTTTCAATTATTGAGCATGTTACCGTACAGGTTATTGCTCATGAGCGGATACATATTGAA TGTATTAGAAAATAACAAATAGGGTCCGCGCACATTCCCGAAAAGTGCACCTGACGTC TAAGAAACATTATTATCATGACATTAAACCTATAAAATAGGGTATCAGGAGGCCCTTCGACTC GCGCGTTGGTGTAGACGGTGAAACCTCTGACACATGCAAGCTCCGGAGATGGTCACAGCTG TCTGTAAGCGGATGCCGGAGCAGACAAGCCGTCAGGGCGCTGAGCGGGTTGGCGGGT CGGGCGTGGCTTAACATGCGGCATCAGAGCAGATTGACTGAGAGTCACCATATGCCGTGTA AATACCGCAGACATGCGTAAGGAGAAAATACCGCATCAGGCCGCTTCCGATTCAGGCTGCCA ACTGTTGGGAAGGGCAGTCGGTGCAGGCCATTCTCGCTTACGCCAGCTGGCGAAAGGGGATGT GCTGCAAGGCAGTAAGTGGTAAACGCCAGGGTTTCCCGACTCACGACGTTGAAACAGCGGC CACTGAATTGAGCTGGTACCCCTTGAAGACAACAGAGAGGATGACATGTGACCGAGGGATTG TCTCGTGGAAACATTGGGACTTCAGATTCCACCCCTTGCACAGCGTGGAAAGCCTCCGACCAAGG GAAGTGTGGATGTAGAGGAAATGAAAGTTGTTCTCATGGGACTTGTGATTGCTTCAC AAGAATGCCAGCTTATGCCGAAACGCAAGAGGAAAGATTAGTGTGCAATCAGGAAAACGCCGAGA TGGATGTGGTCAAGAAGTTAAAGAACCTGCAAGGAGACAGAAATCAAACTCAAAACCTGAGA ACTTCCCTCAACAAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCAGGCCAAAGAAGC TAAACGCCCTAGCTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCGAAACAAATGAAATA ACGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCCGGCCAAA

pLink-5	<p>GGGTACCGAGCTGAATTCACTGGCCGTCGTTTACAACGTCGTACTGGAAAACCCTGGCGTT ACCCAACTTAATCGCCTTGAGCACATCCCCCTTCGCCAGCTGGCGTAATAGCGAAGAGGCCCG CACCGATGCCCTTCCCAAACAGTTGCGCAGCCTGAATGGCCAATGGGCCCTGATCGGGTATTTC TCCTTACGCATCTGTGGGTATTTCACACCGCATATGGTGCACTCTCAGTACAATCTGCTGTGATG CGCATAGTTAACCGAGCCGACACCCGCCAACACCCGCTGACGCCCTGACGGGCTTGTCTG CTCCCGCATCCGTTACAGACAAGCTGTGACCATCTCCGGAGCTGCATGTGAGGGTTTC ACCGTACATACCGAAACGCCAGTCGAAAGGCCCTGATGACGCCATTATAAGGTTAATGT CATGATAATAATGGTTCTAGACGTAGGTGGACTTCCGGAAATGTGCGCGGAACCCCTAT TTGTTATTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCTGATAATGCTT CAATAATATTGAAAAGGAAGAGTATGCGCTACGCAACTGGTCCAGAACCTGACCGAAC CGGTGTAACGCCAGTGGGTTTCTATGGCTTGTATGACTTTGGGGTACAGTCTAT GCCTGGCATCCAAAGCACAGGCCCTAACACAAAGTTAACATCATGAGGGAGATGGCAGCA ACGATGTTACGCAGGCCAGTCGCCCTAACACAAAGTTAACATCATGAGGGAGATGGCAG GCCGAAGTATCGACTCAACTATCAGAGTAGTGGCGTACATGAGGCCATCTGAACCGAC GCTGCCGTACATTGACGGCTCCGACTGGATGGCGGCTGAAGCCACACAGTGTATATTGATT TGCTGGTACGGTACCGTAAGGCTTGTGAAACAACGCCAGGCTTGTGATCAACGACCTTTG GAAACTTCGGCTTCCCTGGAGAGAGGAGATTCTCCGCGTGTAGAAGTCACCATTTGTGCA CGACGACATCTCCGTGGCGGTTACTGCCGCTACTGCGCTGACGGCTTGTGATGGAGC CGCTGTTGATGGGAGGTTTGTGAGGAGCTGCGCTGACGGCTTGTGATGGAGGAGGATT GTCTCGTGGAAACTTGGGACTTCAGATTCCACCCCTTGCAAGCGTGGAAAGCCTCCGACCAAG GGAAGTGTGGATGTAGAGGAATGGAAGTTGTTCTCAGATGGGACTTGTGATTGCTICA CAAGAATGCCAGCTTATGGTAAACGAGGAAAGATTAGTGTGCAATCAGGAAAACGCCGAG ATGGATGTGGCAAGAAGTAAAGAACCTGAGGAGACAGAAATCAAAACTCAAACCTGCA AACTTCCCTCAACAAGCTCACAGGAAACACAGTAGGGAGGTGAAACACCCAGGCCAAGAAAAG CTAACGCCTAGCTGAAATCCAGGAGTCAATGAGAGCTGAAGGTGATGCCGAACCAAATGA AGCGGGGAGGGCAATACCGAACGCCAACCTCTGGGCCCCAA</p>
pLink-6	<p>GGGGATCCTCTAGAGTCGACCTGAGGCATGCAAGCTTGGCTAATCATGGTATAGCTGTTTCC TGTGAAATTGTTACCGCTCAAACTCCACACAAACATACGAGGCCAGCAAGCTAAAGTGTAAAGC CTGGGGTGCCTAATGAGTGAAGCTAACATTAATTGCGTTGCGCTCACTGCCGCTTCCAGTC GGGAAACCTGTCGTGCCAGCTGCATTAATGAATGCCAACGCCGGGGAGAGGGGGTTGGCTA TTGGCGCTCTCCGCTTCCGCTACTGACTCGCTGCCGCTGGCTGTTGGCTGCCGAGCG GTATCAGCTCACTCAAAGGGCTAATACGGTTATCCACAGAATCAGGGATAACGCAGGAAAGAA CATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAGGCCGGTTGGCTGGCTTTCC ATAGGCTCCGCCCTCTGACGAGCATCACAAAATGACGCCAAGCTCAGAGGCTGGCGAAACCCG ACAGGACTATAAGGATACCCAGGGCTTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCT CTGCCGCTTACCGGATACCTGTCGCCCTTCTCCCTGGAGCGTGGCTGGCTGGCTGGCT CGCTGTAGGTATCTCAGTTGGTGTAGGTGCTGCTCCAGCTGGCTGTGAGTCCAACCCGTAAGAC ACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGCGGTGCT ACAGAGTCTTGTGAGCTGGCTAACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCGCT CTGCTGAAGCCAGTTACCTCGGAAAAGAGTTGGTAGCTTGTATCCGGCAAACAAACCCAGC TGGTAGGGTGGTTTGTGAGGAGCTGAGCAGCAGATTACGCCAGGAAAGGGAGGAGGAGG ATCCCTGATCTTCTACGGGGCTGACGCTGAGTGGAAAGAAAACCTACGTTAAGGGATTGG TCATGAGATTATCAAAGGATCTCAGCTTGTGAGCTTAAATTAAAGGATCTCAAGAAG CTAAGTATATGAGTAAACTTGGTGTGAGCTTATTGCGACTACCTTGGTGTGATCTGCC CACGTAGTGGACAAATTCTCCAACGATCTGCGCGAGGCCAAGCGATCTCTTGTG ATAAGCCTGTCTAGCTCAAGTATGACGGGCTGATACTGGGCCGGAGCGCTCCATTGCCAGT CGGCAGCGACATCTCCGCGCAGTTGCCGTTACTGCCGCTGACCAAATGCCGGACACGTA AGCACTACATTGCGCTCATGCCAGGCCAGTCGGCGGGAGTTCCATAGCGTTAAGGTTTCA AGCCGCTCAAATAGATCTCTGGAGGCCAGGAGTCAAAAGGTTCTCCGCCGCTGGAC GGCAACGCTATGTTCTTGTGTTGAGGAGCTGAGGAGCTGAGTGTGACGCC GAAGATACCTGCAAGAATGTCATTGCGCTGCCATTCTCAAATTGCAAGTGTG ACGCCACGGAAATGTCGCGTGCACAACAAATGGTACTTACAGCGCGGAGAATCTCGCT</p>

	CTCCAGGGAAAGCGAAGTTCCAAAAGGTCGTGATCAAAGCTGCCGTTTTCATCAAGC CTTACGGTCACCGTAACCAGCAAATCAATATCACTGTGTGGCTTCAGGCCATCCACTGCCGA GCCGTACAAATGTACGGCCAGCAACGTCGGTTGAGATGGCGCTCGATGACGCCAACTACCTCTG ATAGTTGAGTCGATACTTCCGCGATCACCGCTTCCCTCATGATGTTAACCTTGTGTTAGGGCGAC TGCCCTGCTCGTAAACATCGTGTCTCCATAACATCAAACATCGACCCACGGCGTAACGCCCTT GCTGCTGGATGCCGAGGCATAGACTGTACCCAAAAAACAGTCATAACAAGGCATGAAAACC GCCACTGCCGTTACCACCGCTGCGTCGGTCAAGGTTCTGGACCAGTGTGAGCGCATACT CTTCCCTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATATTGAA TGTATTAGAAAATAACAAATAGGGTTCCGCGCACATTCCCAGAAAGTGCACCTGACGTC TAAGAAAACATTATTATCATGACATTAACCTATAAAATAGGCATACGAGGCCCTTCGACTC GCGCGTTTGGTATGACCGTAAACCTCTGACACATGCACTGCCAGTCCCCGAGATGGTCACAGCTG TCTGTAAGGGATGCCGGAGCAGAACAGCCCTCAGGGCGCTCAGGGGGTTGGCGGGTGA AATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCCCATTCGCCATTAGGCTGCCA ACTGTTGGAAAGGGCAGTGGTGCAGGCTCTTCGCTATTACGCCAGTGGCAGGGGGATGT GCTGCAAGGCAGTTAAGTGGTAACGCCAGGGTTTCCAGTCACGACGTTGAAAACGACGCC CAGTGAAATTGAGCTCGGTACCCTTGAAGACAAGAGCAGGGATGCACATGTGACCGAGGGATTG TCTTCGTGAAACTTGGGACTTCAGATTCCACCCCTTGAAGCAGCTGGAAAGCCTTCCGACCAAGG GAAGTGTGGATGTAGAGGAAATGAAAGTGGGACTTGTGCTTACGATGGGACTTGTGCTTAC AAGAATGCCAGTTAGCGGTAAGAAGTAAAGAACCTGCAAGGAGACAGAAATCAACTCAAACCTGCAAGAA ACTTCCCTCAACAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCAGGCCCAAAGAAC TAAACGCCAGTGTAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCGAACCAAATGAAATA CGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCTGGGCCAA
pLink-7	GGGTACCGAGCTCGAATTCACTGCCGTTTACAACGCTGTGACTGGAAAACCCCTGGCGTT ACCCAACTTAATGCCCTTGACGACATCCCCCTTCCGCACTGGCGTAATAGCGAAGAGGCCG CACCGATGCCCTTCCCAAACAGTTGCCGAGCCTGAATGCCGAATGCCCTGATGCCGTATTTC TCCTTACGCATCTGTGGTATTTCACCCGATATGGTGCACTCTCAGTACAATCTGCTGTATG CCGCATAGTTAACCCAGCCCCGACACCCGCAACAGCCCTGACGCCCTGACGGGCTTGTG CTCCCGCATCCGTTACAGACAAGCTGTGACCATCTCCGGAGCTGCATGTGTCAGAGGTTTC ACCGTCACTACCGAAACGCCGAGTCGAAGGGCTCGTGTACGCCATTTTTATAGGTTAATGT CATGATAATAATGGTTCTAGACGTAGGTGGACTTCCGGGAAATGTGCGCGGAACCCSTAT TTGTTTATTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCTGATAAAATGCTT CAATAATATTGAAAAGGAAGAGTATGCCCTACGCACTGGTCCAGAACCTTGACCGAACCGAG CGGTGGAACCGCAGTGGGTTTCATGGCTTGTATGACTTTTTGGGGTACAGTCTAT GCCTGGGATCCAAGCAAGCGCTTACGGCTGGGTGATGTTGATTTGAGCAGCA ACGATGTTACGCAAGGGCAGTCGCCCTAAACAAAGTTAACATCATGAGGAAAGCGGTGATC GCCGAGTATGCACTACAGAGGTAGTGGCGTACGAGGCCATCTGAACCGACGTT GCTGCCGTACATTGACGGCTCCCGAGTGGATGGCGGCTGAAGCCACACAGTGTATATTGATT TGCTGGTACGGTACCGTAAGGCTTGTGAAACAACGCCGAGCTTGTACGACGACCTTTG GAAACTTCGGCTCCCTGGAGAGAGGAGATTCTCCGCGCTGTAGAAGTCACCATTTGTGCA CGACGACATCTCCGTGGCTTACAGCTAAGCGCAGACTGCAATTGGAGAATGGCAGCGCA ATGACATTCTGAGGTATCTGAGGCCAGCAGTCAGTGGCTATCTGCTGACAA AAGCAAGAGAACATAGCGTTGCTTGTAGGTCCAGCGGGAGGAACCTTGTGATCCGGTCT GAACAGGATCTTGTGAGGCGCTAAATGAAACCTTACGCTATGGAACCTGCCGCGACTGGC TGGCGATGAGCGAAATGTAGTGCTTACGTTGTCCGCTTACGGCTGAGTACCGCAGTAACCGGAAAGA TCGGCCGAAGGATGTGCGTGGCAGTGGCAATGGAGCGCTGCCGCCAGTATGCCGTC ATACTTGAAGCTAGACAGGCTTACGGGAGAAGAAGATCGCTGGCCTCGCGCAGATCA GTTGAAGAATTGTCACACTGAAAGGCAGATCAGGAAACTCTGAGTGGCTTACGGGAGATCTGCTAG ACCAAGTTACTCATATAACTTGTGAAAGTAAACTCTTAAATTAAAAGGATCTAGGT GAAGATCCTTTGTATACTCATGACCAAATACTCTGAGTGGCTTACGGGAGCTGAGCTGCTC AGACCCCTGAGAAAGGATCTTGTGAGCTTCTGGATCTTCTGGCTGAGTACAGAGCTACCAACTCTT GCAAAACAAAAAAACCCGCTACCGCGGTTGGTGTGACAGCCAGCTGGAGCGAACGACCTA TCCGAAGGTAACTGGCTCAGCAGAGCGCAGATAACCAACTGTCTCTAGTGTAGCCGTAGTT AGGCCACCACTCAAGAACTCTGTAGCACCGCCTACATACCTCGCTGCTGATACTCTGTTACCAAGT GGCTGCTGCCAGTGGCGATAAGTCGTCTTACGGGTTGGACTCAAGACGATAGTTACCGGATA AGGCCAGCGGTGGCTGAACGGGGGGTCGTGACACAGCCAGCTGGAGCGAACGACCTA CACCGAAGTACCTACAGCGTGAAGCTGAGAAGAGGCCAGCTCCGAAGGGAGAAAGG CGGACAGGTATCCGGTAACCGGCAGGGTCCGAACAGGAGAGCGCACGAGGAGCTCCAGGGGG AAACCCCTGGTATCTTATAGTCGTGCGGTTTCGCCACCTCTGACTTGTGAGCTGCTGATTTTG TGATGCTGTCAGGGGGCGAGCGCTTGGAGCTGGGAGGCTTGTGAGCTGCTGATTTTG TGCTTGTGAGCTTGTGAGCTGAGTGGCTTACCGGCTGCTGAGCTGCTGAGCTGCTGAGCTGAGTGGAGT TGAGCTCACTCATTAGGCACCCAGGCTTACACTTTATGCTTCCGGCTCGTATGTTGTGGAAT TGTGAGCGATAACAAATTTCACACAGGAAACAGCTATGACCATGATTACGCCAAGCTTGCATGCC GCAGGTGCACTCTAGAGGATCCCCCTGAGACACAATGCCGATTCGACATGCACTGTGACCGAG GGATTGTCTCGTGGAAACCTGGGACTTCAGATTCCACCCCTTGTCAAGCGTGGAAAGCCTCCGAC CAAGGAAAGTGTGGATGTAGAGGAAATGGAAAGTGGTCTCAGATGGGACTTGTGATGCTG TTCACAAGAATGCCAGCTTATGCGTAAACGCAAGAGGAAGATTAGTCGCAATCAGGAAAAGGCC CGAGATGGATGTGGTCAAGAAGTTAAAGAACCTGCAAGGAGACAGAAATCAAACTCAAACCTG CAGAAACTTCCCTCAACAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCAGGCCCAAAG AAAGCTAAACGCCAGTGTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCGAACCAAATGA AATAAGGGGGCATGGGGCAATACCCAGCAACGCCAACTTCTGGGCCAA
pLink-8	GGGTACCGAGCTCGAATTCACTGCCGTTTACAACGCTGTGACTGGAAAACCCCTGGCGTT ACCCAACTTAATGCCCTTGACGACATCCCCCTTCCGCACTGGCGTAAAGCGCAAGAGGCCG CACCGATGCCCTTCCCAAACAGTTGCGCAGTGGCGTAAAGCGCAATGGGCCCTGATGCCGTATTTC

	TCCCTACCGCATCTGTGCGGTATTCACACCGCATACTGGTCACTCTCAGTACAATCTGCTCTGATG CCGCATAGTTAACGCCAGCCCCGACACCCGCCAACACCCGCTGACCGCCCTGACGGGCTGTCTG CTCCCAGCATCCGTTACAGACAAGCTGTGACCCTCTCCGGAGCTGATGTGTCAGAGGTTTC ACCGTCATACCGAAACCGCGAGTCGAAGGGCCTCGTATACCCCTATTNTTATAGGTTAATGT CATGATAATAATGGTTCTTAGACGTCAAGGGCACTTTGGGGAAATGTGCGCGAACCCCTAT TGTTTATTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCCTGATAAATGCTT CAATAATATTGAAAAGGAAGAGATGCGCTCACGCCACTGGTCCAGAACCTTGACCGAACGAG CGGTGGTAACGGCGAGTGGCGTTCTAGGCTTGTACTGACTGTTTTGGGTACAGTCTAT GCCCTGGGAGTCCAAGCAGCAAGCGCCTACGCCGTGGTCGATGTTGATGTTATGGAGCAGCA ACGATGTTACGCAGCAGGGCAGTCGGCTTAAACAAAGTTAACACATGAGGGAAGCGGTGATC GCCGAAGTATCGACTCAACTATCAGAGTAGTGGCTCATGAGGCCATCTGAGACCGACGTT GCTGGCGTACATTGACGGCTCCGAGTGGATGGCGGCTGAGGCCACACAGTGTATATTGATT TGCTGGTTACGGTGACCGTAAGGCTGATGAAACACCGCGCAGCTTGATCAACGACCTTTG GAAACTTCCGGCTTCCCGTGGAGAGAGCGAGATTCTCCGCTGAGAAGTCAACCTGTTGTGCA CGACGACATCATTCCGGCTTATCCAGCTAACGCCAGCTGACATTGATCTGGCTATCTTGTGACAA ATGACATTCTGAGGTATCTCAGGCCACGATCGACATTGATCTGGCTATCTTGTGACAA AAGCAAGAGAACATACCGTTGGCTTGGTAGGTCCAGCGGCGAGGAACCTTGTATCCGGTCT GAACAGGATCTATTGAGGGCTAAATGAAACCTTAAAGCTGAGGACTCGCCGCCACTGGGG TGGCGATGAGCAGAAATGTAGTGTACGGCTTACGGTCTTCCGCTTGGTACAGCGCAGTAACCGGAAAA TCGCGCCGAGGGATGCGCTGCCACTGGCAATGGAGCGCCTGCCGGCCAGTATCAGCCGTC ATACTTGAAGCTAGACAGGTTATCTTGACAAAGAAGAGATCGCTTGGCTCGCGCAGATCA GTTGGAAGAATTGTCACACTCGTAAAGGGCAGATACCAAGGAGTCAACGACCTTAACTGTCAG ACCAAGTTACTCATATACTTGTGTTAAACCTTACGTTCTGAGGTTGAGCTGCTTCTGAGCGTC GAAGATCTTTGTGATAATCTCATGACCAAATCCCTAACGTGAGTTTCTGTTCACTGAGCGTC AGACCCCGTAGAAAAGATCAAAGGATCTTGTGAGATCTTGTGAGCTTCTGCGCT GCAAACAAAAAACCCCGTACCCAGCGGTGGTTGTTGCTGGGATCAAGAGCTACCAACTCTT TCCGAAGGTAACTGGCTCAGCAGAGCGCAGATACCAAAACTGTCCTTGTAGTGTAGCGTAGTT
pLink-9	GGGTACCGAGCTGAATTCACTGGCCGTCGTTTACAACCGCTGACTGGGAAACCCCTGGCGTT ACCCAACCTTAATGCCCTTGACGACACATCCCCCTTCGCCAGCTGGCTAATAGCGAAGAGGCCG CACCGATGCCCTTCCAACAGTTGCGCAGCTGAATGGCGAATGGCGCTGATGCCGTATTTC TCCTTACGCATCTGTGCGGTATTCACACCCGATATGGTCACTCTCAGTACAATCTGCTCTGATG CCGCATAGTTAACGCCGCCCCGACACCCGCCAACACCCGCTGACCGCCCTGACGGGCTGTCTG CTCCCAGCATCCGTTACAGACAAGCTGTGACCCTCTCCGGAGCTGATGTGTCAGAGGTTTC ACCGTCATACCGAAACCGCGAGTCGAAGGGCCTCGTATACCCCTATTNTTATAGGTTAATGT CATGATAATAATGGTTCTTAGACGTCAAGGGCACTTTGGGGAAATGTGCGCGAACCCCTAT TGTTTATTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCCTGATAAATGCTT CAATAATATTGAAAAGGAAGAGATGCGCTCACGCCACTGGTCCAGAACCTTGACCGAACGAG CGGTGGTAACGGCGAGTGGCGTTCTAGGCTTGTACTGACTGTTTTGGGTACAGTCTAT GCCCTGGGAGTCCAAGCAGCAAGCGCCTACGCCGTGGTCGATGTTGATGTTATGGAGCAGCA ACGATGTTACGCAGCAGGGCAGTCGGCTTAAACAAAGTTAACACATGAGGGAAGCGGTGATC GCCGAAGTATCGACTCAACTATCAGAGTAGTGGCTCATGAGGCCATCTGAGACCGACGTT GCTGGCGTACATTGACGGCTCCGAGTGGATGGCGGCTGAGGCCACACAGTGTATATTGATT TGCTGGTTACGGTGACCGTAAGGCTGATGAAACACCGCGCAGCTTGATCAACGACCTTTG GAAACTTCCGGCTTCCCGTGGAGAGAGCGAGATTCTCCGCTGAGAAGTCAACCTGTTGTGCA CGACGACATCATTCCGGCTTATCCAGCTAACGCCAGCTGACATTGATCTGGCTATCTTGTGACAA ATGACATTCTGAGGTATCTCAGGCCACGATCGACATTGATCTGGCTATCTTGTGACAA AAGCAAGAGAACATACCGTTGGCTTGGTAGGTCCAGCGGCGAGGAACCTTGTATCCGGTCT GAACAGGATCTATTGAGGGCTAAATGAAACCTTAAAGCTGAGGACTCGCCGCCACTGGGG TGGCGATGAGCAGAAATGTAGTGTACGGCTTACGGTCTTCCGCTTGGTACAGCGCAGTAACCGGAAAA TCGCGCCGAGGGATGCGCTGCCACTGGCAATGGAGCGCCTGCCGGCCAGTATCAGCCGTC ATACTTGAAGCTAGACAGGTTATCTTGACAAAGAAGAGATCGCTTGGCTCGCGCAGATCA GTTGGAAGAATTGTCACACTCGTAAAGGGCAGATACCAAGGAGTCAACGACCTTAACTGTCAG ACCAAGTTACTCATATACTTGTGTTAAACCTTACGTTCTGAGGTTGAGTTTCTGTTCACTGAGCGTC GAAGATCTTTGTGATAATCTCATGACCAAATCCCTAACGTGAGTTTCTGTTCACTGAGCGTC AGACCCCGTAGAAAAGATCAAAGGATCTTGTGAGATCTTGTGAGCTTCTGCGCT GCAAACAAAAAACCCCGTACCCAGCGGTGGTTGTTGCTGGGATCAAGAGCTACCAACTCTT TCCGAAGGTAACTGGCTCAGCAGAGCGCAGATACCAAAACTGTCCTTGTAGTGTAGCGTAGTT

	AGGCCACCACTCAAGAACTCTGTAGCACCGCCTACATACCTCGCTTGCTAATCCGTTACCAAGTGGCTGCTGCCAGTGGCGATAAGTCGTCTTACCGGTTGACTCAAGACGATAGTACCGGATAAGGCCAGCGGTGCGGCTGAACGGGGGGGTCGTGCACACAGCCCAGCTGGAGCGAACGACCTAACCGAAACTGAGATACCTACAGCGTGAAGCTATGAGAAAAGCGCCACGCTTCCCAGGGAGAAAGGCGGACAGGTATCCGTAAGCGGCAGGGTCGGAACAGGGAGAGCGCACGAGGGAGCTCCAGGGGGAAACGCCCTGGTATCTTATAGTCCTGCGGGTTGCCACCTCTGACTTGAGCGTCGATTTTGATGCTCGTCAGGGGGCGAGCCTATGGAAAACGCCAGCAACGCCCTTTTACGGTTCTGGCCTTTGCTGGCCTTTGCTCACATGTTCTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATACCGCCTTGAGTGAAGCTGATACCGCTGCCGAGCCAACGCCCTCTCCCAGCGCTGGCGATTCAATTAA TGCAAGCTGGCACGACAGGTTCCGACTGAAAGCGGGCAGTGAGCGCAACGCCAATTAAATGTGAGTTAGCTCACTCATTAGGCACCCAGGCTTACACTTATGCTTCCGGCTCGTATGTTGTGGAAATTGTGAGCGGATAACAATTTCACACAGGAAACAGCTATGACCATGATTACGCCAAGCTTGCATGCCCTGCAAGGTGACTCTAGAGGATCCCCCTGAAAGACAAAGACGAATTGATATCGCACATGTGACCGAGGGATTGCTTGTGAAACTTGGGACTTCAGATCCACCCCTGCAAGCGTGGAAAGCCCTCCGACCAAGGAAAGTCGGATGTAGAGGAATGAAAGTTCTCAGATGGGACTTGCTGATTGCTCACAAGAATGCCAGCTTATGCGGTAACCGCAGAGGAAGATTAGCTGCAATCAGGAAACGCCCGAGATGGATGTCGGTCAAGAAGTTAAAGAACCTGCAGGAGACAGAAATCAAACTCAAACCCCTGCAGAAACTTCCACAAGCTCCACAGGAAACACAGTAGGGAGGTGAAACACCAGGCCGAAAGAAAGCTAAACGCCCTAGCTGAAATCCAGGAGTCATGAGAGCTGAAGGTGATGCCGAACCAAATGAAATAAGCGGGGCATGGGGCAATACCCAGCAACGCCGAACCTCCTGGGCCAAA
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REFERENCES

1. Hockemeyer, D. *et al.* *Nat Biotechnol* **27**, 851–857 (2009).
2. Thyagarajan, B. *et al.* *Molecular and Cellular Biology* **21**, 3926–3934 (2001).
3. Olivares, E. C. *et al.* *Gene* **278**, 167–176 (2001).
4. Fukushige, S., Fukushige, S., Sauer, B. & Sauer, B. *Proc Natl Acad Sci USA* **89**, 7905–7909 (1992).
5. O'Gorman, S., Fox, D. T. & Wahl, G. M. *Science* **251**, 1351–1355 (1991).