Plasmid sequences

pDSAP-LRIM1-GFP

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pDSAT-APL1-GFP

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pattB-RfB2 TEP1-GFP,3xP3-RFP (*TEP1-GFP* reporter)

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pattB-RfB2-Vg-TEP1r SV40YFP (line *Vg-TEP1r*)

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1981 ATGCAGCTGG CACGACAGGT TTCCCGACTG GAAAGCGGGC AGTGAGCGCA ACGCAATTAA
2041 TGTGAGTTAG CTCACTCATT AGGCACCCCA GGCTTTACAC TTTATGCTTC CGGCTCGTAT
2101 GTTGTGTGGA ATTGTGAGCG GATAACAATT TCACACAGGA AACAGCTATG ACCATGATTA
2161 CGCCAAGCTC GAAATTAACC CTCACTAAAG GGAACAAAAG CTGGCTAGAA CTAGTGTCGA
2221 CATGCCCGCC GTGACCGTCG AGAACCCGCT GACGCTGCCC CGCGTATCCG CACCCGCCGA
2281 CGCCGTCGCA CGTCCCGTGC TCACCGTGAC CACCGCGCCC AGCGGTTTCG AGGGCGAGGG
2341 CTTCCCGGTG CGCCGCGCGT TCGCCGGGAT CAACTACCGC CACCTCGACC CGTTCATCAT
2401 GATGGACCAG ATGGGTGAGG TGGAGTACGC GCCCGGGGAG CCCAAGGGCA CGCCCTGGCA
2461 CCCGCACCGC GGCTTCGAGA CCGTGACCTA CATCGTCGAC ACTAGTGgat catcaacaag
2521 tttGTACAAA AAAGCAGGCT GGTACCGGGC CCCCCGCTAG CGTCGAGTTC AACTCGACCA
2581 TAATAATTGA TCCGTCAATC CATATTGGTC CGCAATAATG AAAGTTGCAA GAGTACGACG
2641 GTATGAAAAG AGGTTCAGTA AGTTGTAAAC TAATAGTTTC TTCCCAACGT TCAAATGCTG
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3181 TCAGCCCGTA CGACTTAACT ACATTCTCGT CATGGGTTCA AGCCCCAGAT GGACCGTGCC
3241 GCCATACGTA GAGTCAGTCC TATCCTGTTA TGGGGGGTAA TACATAAGAC ACTGAAAGCC
3301 AACCCCACAA GTGGTACAGA CAAGCCTTGa CCGACAATTG TTGTTGAGCC AAACAGAAGA
3361 AGAATCCATT TCGGGAAATG ATTTTATCAT TCAATCAAAC CAGTCAATCA TAAACATCAT
3421 AGTTTTAAAT ACTCAAAACT AGTTGAGATC TTTAAAACAC ATTATTTTAG TTTAATTAAA
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6181 GCAAATGAAA AGACGGGGCG TAATGCACTG CAGTCAGGCA AGCCGATCGG CAAGCTGGTG
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6481 GTGTTGCAGT TTACGCTGTT CAACAACCTT GGAGCGGAGT ATATAGCCGA TGTGACGCTG
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7501 GAGGCACTCA ATAAGCTGAT TGATATGTCT TTCATTGATG CTGATAAAAA CGAACGGTTC
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 9421 AGCCACAACG TCTATATCAT GGCCGACAAG CAGAAGAACG GCATCAAGGT GAACTTCAAG
 9481 ATCCGCCACA ACATCGAGGA CGGCAGCGTG CAGCTCGCCG ACCACTACCA GCAGAACACC
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 9781 CCCTGAACCT GAAACATAAA ATGAATGCAA TTGTTGTTGT TAACTTGTTT ATTGCAGCTT
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 9901 TGCATTCTAG TTGTGGTTTG TCCAAACTCA TCAATGTATC TTAAAGCTTA TCGATACctc
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10141 GAGGCCCGCA CCGATCGCCC TTCCCAACAG TTGCGCAGCC TGAATGGCGA ATGGAAATTG
10201 TAAGCGTTAA TATTTTGTTA AAATTCGCGT TAAATTTTTG TTAAATCAGC TCATTTTTTA
10261 ACCAATAGGC CGAAATCGGC AAAATCCCTT ATAAATCAAA AGAATAGACC GAGATAGGGT
10321 TGAGTGTTGT TCCAGTTTGG AACAAGAGTC CACTATTAAA GAACGTGGAC TCCAACGTCA
10381 AAGGGCGAAA AACCGTCTAT CAGGGCGATG GCCCACTACG TGAACCATCA CCCTAATCAA
10441 GTTTTTTGGG GTCGAGGTGC CGTAAAGCAC TAAATCGGAA CCCTAAAGGG AGCCCCCGAT
10501 TTAGAGCTTG ACGGGGAAAG CCGGCGAACG TGGCGAGAAA GGAAGGGAAG AAAGCGAAAG
10561 GAGCGGGCGC TAGGGCGCTG GCAAGTGTAG CGGTCACGCT GCGCGTAACC ACCACACCCG
10621 CCGCGCTTAA TGCGCCGCTA CAGGGCGCGT CAG
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Promoter sequences:

Lp promoter

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Vg promoter

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TEP1 minimal promoter

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tcgcacaacgtccaatccaacc

TALE sequences:

N terminal domain (start codon is in bold, *Bsa*I overhangs are capitalized):

T**ATG**gccaagaagaagaggaaggtgcaggtggatctacgcacgctcggctacagccagcagcaacaggagaaga
tcaaaccgaaggttcgttcgacagtggcgcagcaccacgaggcactggtcggccatgggtttacacacgcgcacatcgtt
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C terminal domain (*Bsa*I overhangs are capitalized):

AGCAttgttgcccagttatctcgccctgatccggcgttggccgcgttgaccaacgaccacctcgtcgccttggcctgcctcg
gcggacgtcctgcgctggatgcagtgaaaaagggattgccgcacgcgccggccttgatcaaaagaaccaatcgccgtat
tcccgaacgcacatcccatcgcgttgcc

Repeat domain: (Capital letters correspond to *Bsa*I overhangs, nnnnnn correspond to two codons encoding the repeat-variable diresidues (RVD) that determine specificity for DNA bases)

GAACccccggagcaggtggtggccatcgccagcnnnnnnggcggtaagcaggcgctggagacggtgcagcggct
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Vp16 activation domain:

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