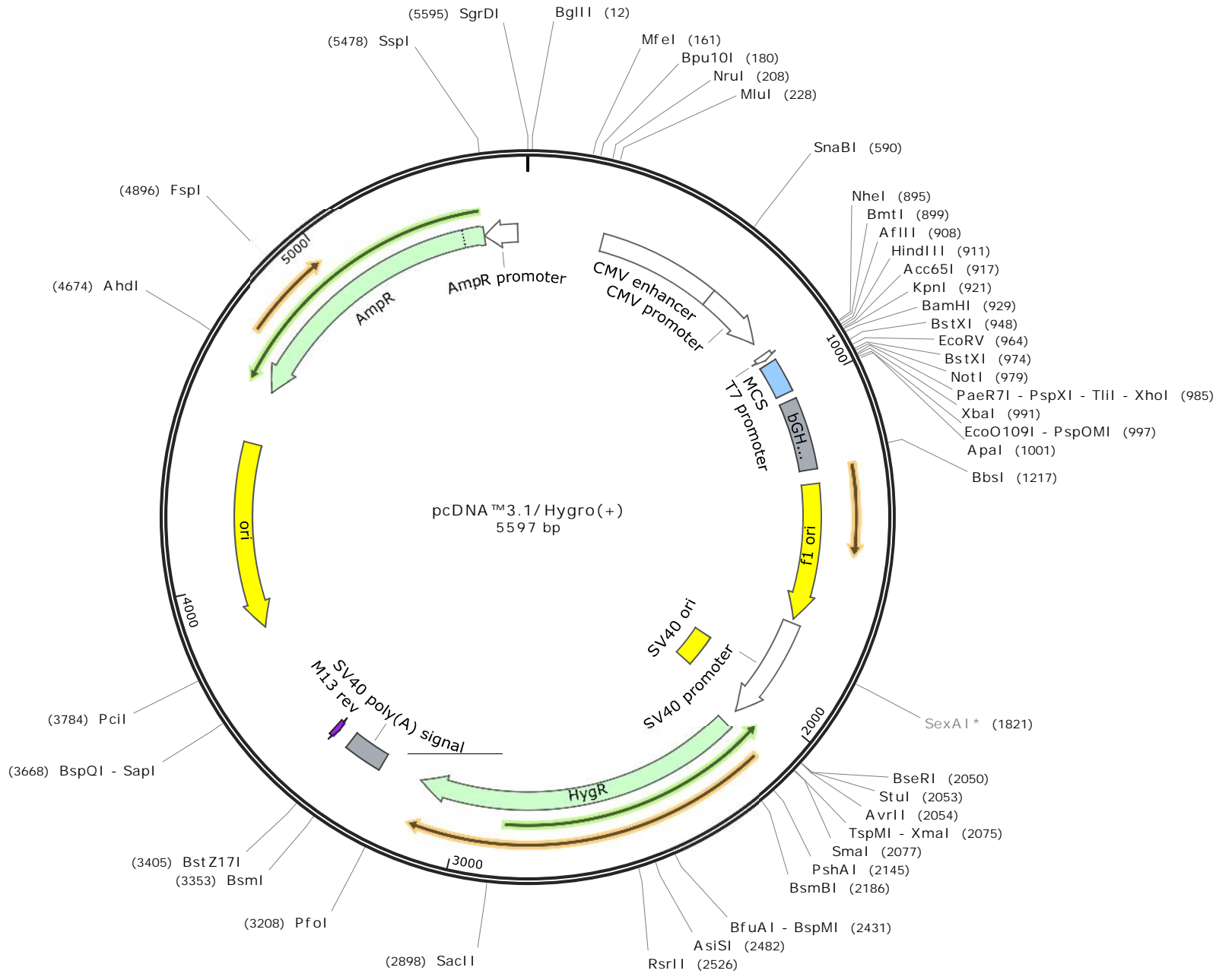


Mammalian expression vector with the CMV promoter. The MCS is in the forward (+) orientation.





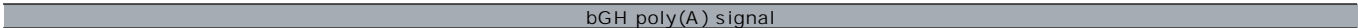
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 GATCAGGTCACACCACCTTAAGACGTCTATAGGTCGTGTCCACCGCCGGCGAGCTCAGATCTCCCGGGCAAATTTGGGCGACTAGT



GCCTCGACTGTGCCTTCTAGTTGCCAGCCATCTGTTGTTTGCCCTCCCGCTGCCTTCCTTGACCCTGGAAGGTGCCACTCCCA  
 CGGAGCTGACACGGAAGATCAACGGTGGTAGACAACAAACGGGGAGGGGGCACGGAAGGAAGTGGGACCTTCCACGGTGAGGGT



CTGTCTTTTCTAATAAAAATGAGGAAATTGCATCGCATTGTCTGAGTAGGTGTCATTCTATTCTGGGGGGTGGGGTGGGGCAGGA  
 GACAGGAAAAGGATTATTTTACTCCTTTAACGTAGCGTAACAGACTCATCCACAGTAAGATAAGACCCCCACCCACCCCGTCTCT



BbsI  
 CAGCAAGGGGGAGGATTGGGAAGACAATAGCAGGCATGCTGGGGATGCGGTGGGCTCTATGGCTTCTGAGGCGGAAAGAACCAGC  
 GTCGTTCCCCCTCCTAACCTTCTGTTATCGTCCGTACGACCCCTACGCCACCCGAGATACCGAAGACTCCGCCTTTCTTGGTCTG



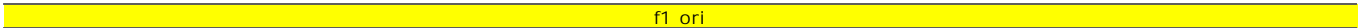
M A S E A E R T S →

TGGGGCTCTAGGGGGTATCCCCACGCGCCCTGTAGCGGCGCATTAAAGCGCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGCTA  
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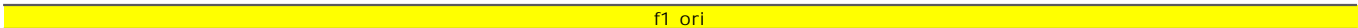
W G S R G Y P H A P C S G A L S A A G V V V T R S V T A →

CACTTGCCAGCGCCCTAGCGCCCGCTCCTTTTCGCTTTCTTCCCTTCTTTCTCGCCACGTTGCGCCGGCTTTCCCGTCAAGCTCT  
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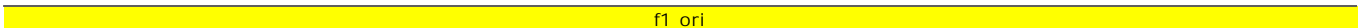
T L A S A L A P A P F A F F P S F L A T F A G F P R Q A L →

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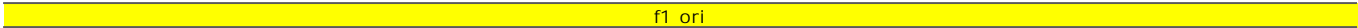


N R G L P L G F R F S A L R H L D P K K L D →

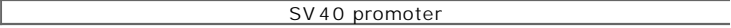
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GAACAACACTCAACCCTATCTCGGTCTATTCTTTGATTTATAAGGGATTTGCCGATTTCCGGCTATTGGTTAAAAAATGAGCT  
 CTTGTTGTGAGTTGGGATAGAGCCAGATAAGAAAATAAATATTCCTAAAACGGCTAAAGCCGGATAACCAATTTTTTACTCGA



GATTTAACAAAAATTTAACGCGAATTAATTCTGTGGAATGTGTGTGTCAGTTAGGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCA  
 CTAATTGTTTTTAAATTGCGCTTAATTAAGACACCTTACACACAGTCAATCCACACCTTTCCAGGGGTCCGAGGGGTCTGTCCTG



SexAI \*

GAAGTATGCAAAGCATGCATCTCAATTAGTCAGCAACCCAGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAAAG  
 CTTTCATACGTTTCGTACGTAGAGTTAATCAGTCGTTGGTCCACACCTTTCAGGGGTCCGAGGGGTCTCGTCCGTCTTCATACGTTTC

SV40 promoter

1870

CATGCATCTCAATTAGTCAGCAACCCATAGTCCC GCCCTAACTCCGCCCATCCCGCCCCTAACTCCGCCCAGTTCGCCCATTTCT  
 GTACGTAGAGTTAATCAGTCGTTGGTATCAGGGCGGGGATTGAGGCGGGTAGGGCGGGGATTGAGGCGGGTCAAGGCGGGTAAGA

SV40 promoter

1955

SV40 ori

CCGCCCCATGGCTGACTAATTTTTTTTATTTATGCAGAGGCCGAGGCCGCTCTGCCTCTGAGCTATTCCAGAAGTAGTGAGGAG  
 GGC GGGGTACCGACTGATTA AAAAAAATAAATACGTCTCCGGCTCCGGCGGAGACGGAGACTCGATAAGGTCTTCATCACTCCTC

SV40 promoter

2040

SV40 ori

BseRI Stul AvrII

XmaI TspMI SmaI

GCTTTTTTGGAGGCCTAGGCTTTTGC AAAAAGCTCCCCGGGAGCTTGTATATCCATTTTCGGATCTGATCAGCACGTGATGAAAAA  
 CGAAAAAACCTCCGGATCCGAAAACGTTTTTCGAGGGCCCTCGAACATATAGGTAAAAGCTAGACTAGTCGTGCACTACTTTTT

2125

SV40 promoter

1  
M K K  
HygR

SV40 ori

M K K

A K A F L E R S S T Y G N E S R I L V H H F L

PshAI

BsmBI

GCCTGAACTCACCGCGACGTCTGTGCGAGAAGTTTCTGATCGAAAAGTTTCGACAGCGTCTCCGACCTGATGCAGCTCTCGGAGGGC  
 CGGACTTGAGTGGCGCTGCAGACAGCTCTTCAAAGACTAGCTTTTCAAGCTGTCTCGCAGAGGCTGGACTACGTGCGAGAGCCTCCCCG

2210

HygR

5 10 15 20 25 30  
 P E L T A T S V E K F L I E K F D S V S D L M Q L S E G  
 R F E G R R R D L L K Q D F L E V A D G V Q H L E R L A

GAAGAATCTCGTGCTTTCAGCTTCGATGTAGGAGGGCGTGGATATGTCCTGCGGGTAAATAGCTGCGCCGATGGTTTCTACAAAG  
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2295

HygR

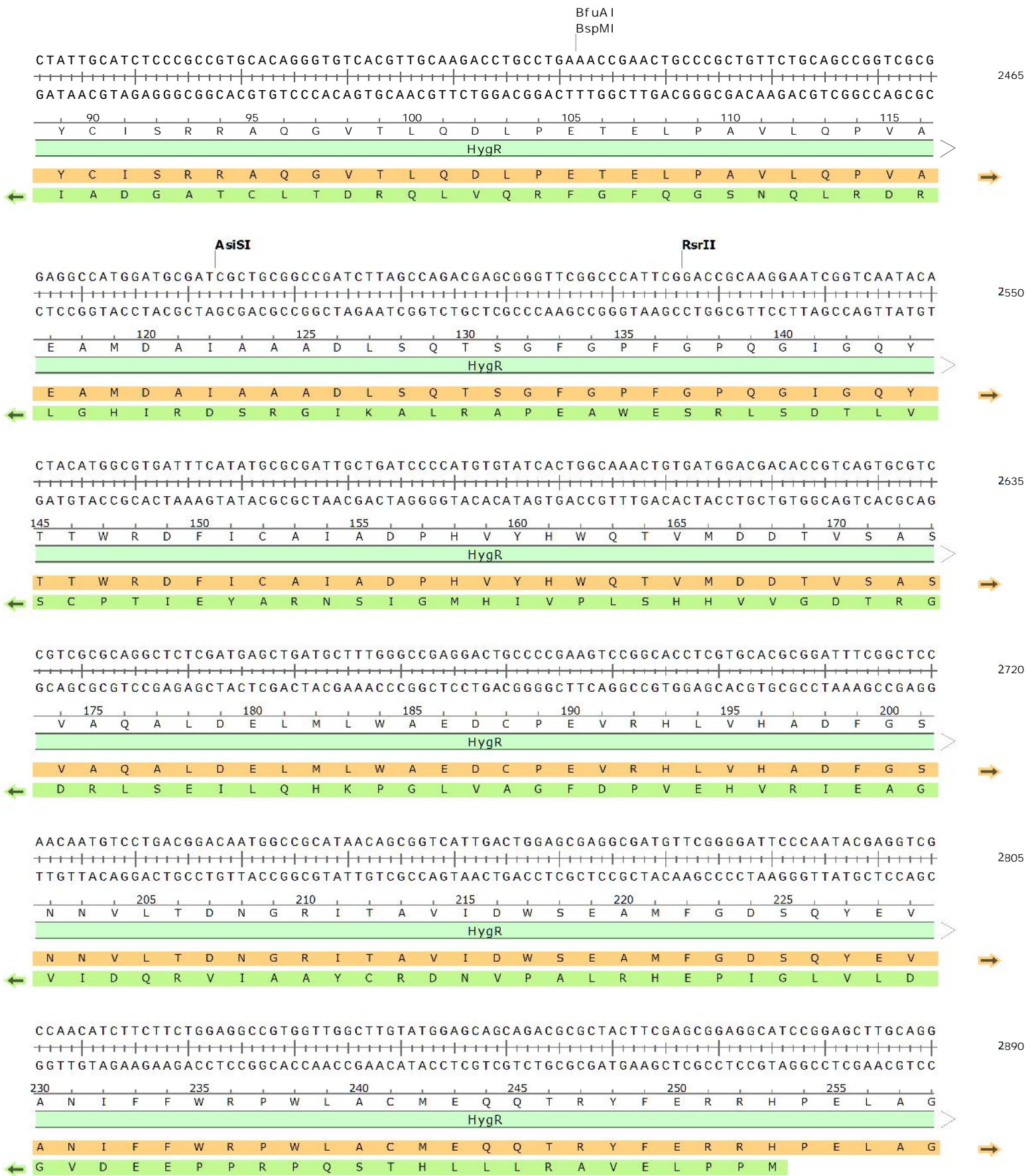
35 40 45 50 55  
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 F F R T S E A E I Y S P T S I D Q P Y I A A G I T E V F

ATCGTTATGTTTATCGGCACTTTGCATCGGCCGCGCTCCCGATTCCGGAAGTGCTTGACATTGGGGGAATTCAGCGAGAGCCTGAC  
 TAGCAATACAAATAGCCGTGAAACGTAGCCGGCGCGAGGGCTAAGGCCTTACGAAGTGTAAACCCCTTAAGTCGCTCTCGGACTG

2380

HygR

60 65 70 75 80 85  
 D R Y V Y R H F A S A A L P I P E V L D I G E F S E S L T  
 I T I N I P V K C R G R E R N R F H K V N P F E A L A Q G



SacI I  
 ATCGCCGCGGCTCCGGGGCGTATATGCTCCGCATTGGTCTTGACCAACTCTATCAGAGCTTGGTTGACGGCAATTTTCGATGATGCA  
 TAGCGGCGCCGAGGCCCGCATATACGAGGCGTAACCAGAAGCTGGTTGAGATAGTCTCGAACCAACTGCCGTTAAAGCTACTACGT  
 S P R L R A Y M L R I G L D Q L Y Q S L V D G N F D D A  
 HygR  
 S P R L R A Y M L R I G L D Q L Y Q S L V D G N F D D A

GCTTGGGCGCAGGGTCGATGCGACGCAATCGTCCGATCCGGAGCCGGGACTGTCTGGGCGTACACAAATCGCCCGCAGAAGCGCGG  
 CGAACCCGCGTCCCAGCTACGCTGCGTTAGCAGGCTAGGCCTCGGCCCTGACAGCCCGCATGTGTTTAGCGGGCGTCTTCGCGCC  
 A W A Q G R C D A I V R S G A G T V G R T Q I A R R S A  
 HygR  
 A W A Q G R C D A I V R S G A G T V G R T Q I A R R S A

CCGTCTGGACCGATGGCTGTGTAGAAGTACTCGCCGATAGTGGAAACCGACGCCCCAGCACTCGTCCGAGGGCAAAGGAATAGCA  
 GGCAGACCTGGCTACCGACACATCTTCATGAGCGGCTATCACCTTTGGCTGCGGGGTCGTGAGCAGGCTCCCGTTTCCTTATCGT  
 A V W T D G C V E V L A D S G N R R P S T R P R A K E  
 HygR  
 A V W T D G C V E V L A D S G N R R P S T R P R A K E

PfoI  
 CGTGCTACGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGGTTGGGCTTCGGAATCGTTTTCCGGGACGCCGGCTGGATGATC  
 GCACGATGCTCTAAAGCTAAGGTGGCGGCGGAAGATACTTTCCAACCCGAAGCCTTAGCAAAGGCCCTGCGGGCCGACCTACTAG  
 CTCCAGCGCGGGGATCTCATGCTGGAGTTCCTCGCCACCCCAACTTGTTTATTGCAGCTTATAATGGTTACAAATAAAGCAATA  
 GAGGTCGCGCCCCCTAGAGTACGACCTCAAGAAGCGGGTGGGGTTGAACAAATAACGTCGAATATTACCAATGTTTATTTTCGTTAT  
 SV 40 poly(A) signal

BsmI  
 GCATCACAAATTTACAAATAAAGCATTTTTTCTACTGCATTCTAGTTGTGGTTTTGTCCAAACTCATCAATGTATCTTATCATGT  
 CGTAGTGTTTAAAGTGTTTATTTTCGTAATAAAGTACGTAAGATCAACACCAACAGGTTTGGAGTAGTTACATAGAATAGTACA  
 SV 40 poly(A) signal

BstZ17I  
 CTGTATACCGTTCGACCTCTAGCTAGAGCTTGGCGTAATCATGGTCATAGCTGTTTCTGTGTGAAATTGTTATCCGCTCACAATT  
 GACATATGGCAGCTGGAGATCGATCTCGAACCGCATTAGTACCAGTATCGACAAAGGACACACTTTAACAATAGGCGAGTGTTAA  
 M13 rev

CCACACAACATACGAGCCGGAAGCATAAAGTGTAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTAATTGCGTTGCGCT  
 GGTGTGTTGTATGCTCGGCCTTCGTATTTACATTTTCGGACCCACGGATTACTCACTCGATTGAGTGTAATTAACGCAACGCGA  
 CACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCCTAT  
 GTGACGGGCGAAAGGTCAGCCCTTTGGACAGCACGGTCGACGTAATTACTTAGCCGGTTGCGCGCCCTCTCCGCCAAACGCATA

SapI  
 BspQI  
 TGGGCGCTCTTCCGCTTCTCGCTCACTGACTCGCTGCGCTCGGTCGTTCCGGCTGCGGCGAGCGGTATCAGCTCACTCAAAGGCG  
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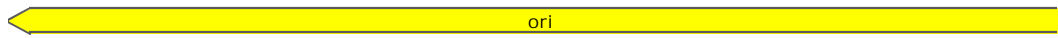
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3825

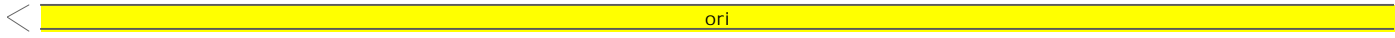
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3910



CCCGACAGGACTATAAAGATACCAGGCGTTTTCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCTGCCGCTTACCGGA  
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3995



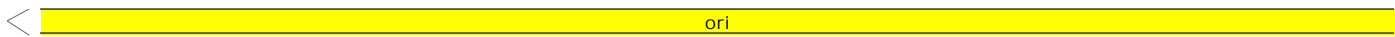
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4080



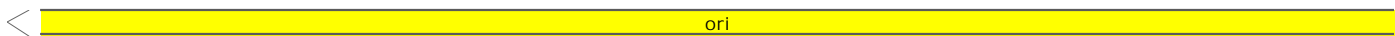
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4165



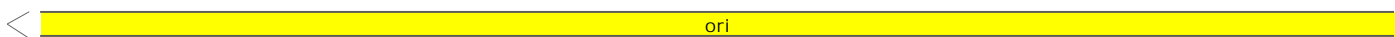
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4250



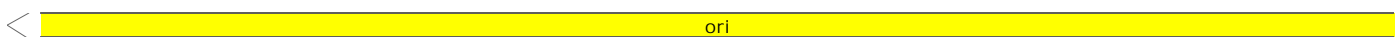
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4335



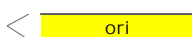
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4420



AGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAAACGAAAACCTCACGTTAAGGGATTTTGGTCATG  
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4505



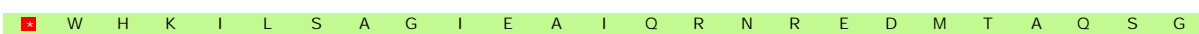
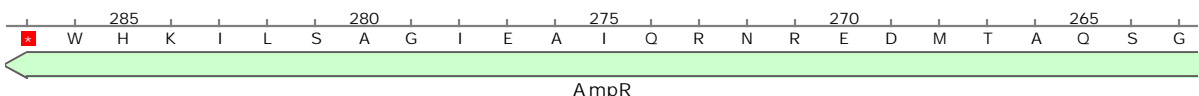
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4590

AhdI

GGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTGACTCCC  
CCAGACTGTCAATGGTTACGAATTAGTCACTCCGTGGATAGAGTTCGCTAGACAGATAAAGCAAGTAGGTATCAACGGACTGAGGG

4675



CGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTCACCGGCT  
 GCAGCACATCTATTGATGCTATGCCCTCCCGAATGGTAGACCGGGGTCACGACGTTACTATGGCGCTCTGGGTGCGAGTGGCCGA

4760

T T Y I V V I R S P K G D P G L A A I I G R S G R E G A

AmpR

M I P R D P R S P A

T T Y I V V I R S P K G D P G L A A I I G R S G R E G A

CCAGATTTATCAGCAATAAACCAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGGTCCTGCAACTTTATCCGCCTCCATCCAGTCTA  
 GGTCTAAATAGTCGTTATTTGGTCGGTCGGCCTTCCCGGCTCGCGTCTTACCAGGACGTTGAAATAGGCGGAGGTAGGTTCAGAT

4845

G S K D A I F W G A P L A S R L L P G A V K D A E M W D I

AmpR

P D L S A I N Q P A G R A E R R S G P A T L S A S I Q S

G S K D A I F W G A P L A S R L L P G A V K D A E M W D I

FspI

TTAATTGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTTTGCGCAACGTTGTTGCCATTGCTACAGGCATCGTGGT  
 AATTAACAACGGCCCTTCGATCTCATTTCATCAAGCGGTCAATTATCAAACGCGTTGCAACAACGGTAACGATGTCCGTAGCACCA

4930

L Q Q R S A L T L L E G T L L K R L T T A M A V P M T T

AmpR

I N C C R E A R V S S S P V N S L R N V V A I A T G I V V

L Q Q R S A L T L L E G T L L K R L T T A M A V P M T T

GTCACGCTCGTCGTTTGGTATGGCTTCATTCAGCTCCGGTCCCAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAA  
 CAGTGCAGCAGCAAACCATACCGAAGTAAGTCGAGGCCAAGGGTTGCTAGTTCGCTCAATGTAAGGGGTACAACACGTTT

5015

D R E D N P I A E N L E P E W R D L R T V H D G M N H L

AmpR

S R S S F G M A S F S S G S Q R S R R V T

D R E D N P I A E N L E P E W R D L R T V H D G M N H L

AAAGCGGTTAGCTCCTTCGGTCTCCGATCGTTGTCAGAAGTAAGTTGGCCGCGAGTGTATCACTCATGGTTATGGCAGCACTGC  
 TTTGCGCAATCGAGGAAGCCAGGAGGCTAGCAACAGTCTTTCATTCAACCGGCGTCACAATAGTGAGTACCAATACCGTCGTGACG

5100

F A T L E K P G G I T T L L L N A A T N D S M T I A A S C

AmpR

F A T L E K P G G I T T L L L N A A T N D S M T I A A S C

ATAATTCTTACTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTAT  
 TATTAAGAGAATGACAGTACGGTAGGCATTCTACGAAAAGACACTGACCACTCATGAGTTGGTTCAGTAAGACTCTTATCACATA

5185

L E R V T M G D T L H K E T V P S Y E V L D N Q S Y H I

AmpR

L E R V T M G D T L H K E T V P S Y E V L D N Q S Y H I



GCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGCTCATCATTGGA  
 CGCCGCTGGCTCAACGAGAACGGGCGCAGTTATGCCCTATTATGGCGCGGTGTATCGTCTTGAAATTTTCACGAGTAGTAACCT

5270

R R G L Q E Q G A D I R S L V A G C L L V K F T S M M P

AmpR

R R G L Q E Q G A D I R S L V A G C L L V K F T S M M P

AAACGTTCTTCGGGGCGAAAACCTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGATGTAACCCACTCGTGACACCAACTGAT  
 TTTGCAAGAAGCCCCGCTTTTGAGAGTTCTTAGAATGGCGACAACCTCTAGGTCAAGCTACATTGGGTGAGCACGTGGGTTGACTA

5355

F R E E P R F S E L I K G S N L D L E I Y G V R A G L Q D

AmpR

F R E E P R F S E L I K G S N L D L E I Y G V R A G L Q D

CTTCAGCATCTTTTACTTTACCCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCCAAAATGCCGCAAAAAAGGGAATAAGGGCGAC  
 GAAAGTCGTAGAAAATGAAAGTGGTCGCAAAGACCCACTCGTTTTTGTCTTCCGTTTTACGGCGTTTTTTCCCTTATTCCCGCTG

5440

E A D K V K V L T E P H A F V P L C F A A F F P I L A V

signal sequence

AmpR

E A D K V K V L T E P H A F V P L C F A A F F P I L A V

SspI

ACGGAAATGTTGAATACTCATACTCTTCCCTTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATA  
 TGCCTTTACAACCTTATGAGTATGAGAAGGAAAAAGTTATAATAACTTCGTAAATAGTCCCAATAACAGAGTACTCGCCTATGTAT

5525

R F H Q I S M

signal sequence

AmpR

AmpR promoter

R F H Q I S M





























SgrDI




TTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAAAGTGCCACCTGACGTC 3'  
 AAACCTTACATAAATCTTTTTATTTGTTTATCCCCAAGGCGCGTGTAAAGGGGCTTTTTACGGTGGACTGCAG 5'

5597

AmpR promoter

Enzymes	Sites	
Acc65I	1	917
AflII	1	908
AhdI	1	4674
Apal	1	1001
AsiSI	1	2482
AvrII	1	2054
BamHI	1	929
BbsI	1	1217
BfuAI	1	2431
BglII	1	12
BmtI	1	899
Bpu10I	1	180
BseRI	1	2050
BsmI	1	3353
BsmBI	1	2186
BspMI	1	2431
BspQI	1	3668
BstXI	2	948 974
BstZ17I	1	3405
EcoO109I	1	997
EcoRV	1	964
FspI	1	4896
HindIII	1	911
KpnI	1	921
MfeI	1	161
MluI	1	228
NheI	1	895
NotI	1	979
NruI	1	208
PaeR7I	1	985
PciI	1	3784
PfoI	1	3208
PshAI	1	2145
PspOMI	1	997
PspXI	1	985
RsrI	1	2526
SacII	1	2898
SapI	1	3668
SexAI	1*	1821*
SgrDI	1	5595
SmaI	1	2077
SnaBI	1	590
SspI	1	5478
StuI	1	2053
TliI	1	985
TspMI	1	2075
XbaI	1	991
XhoI	1	985
XmaI	1	2075

Feature	Location	Size (bp)			Type
✓ CMV enhancer /note = human cytomegalovirus immediate early enhancer	235 .. 614	380			enhancer
✓ CMV promoter /note = human cytomegalovirus (CMV) immediate early promoter	615 .. 818	204			promoter
✓ T7 promoter /note = promoter for bacteriophage T7 RNA polymerase	863 .. 881	19			promoter
✓ MCS /note = multiple cloning site	895 .. 1002	108			misc_feature
✓ bGH poly(A) signal /note = bovine growth hormone polyadenylation signal	1028 .. 1252	225			polyA_signal
✓ f1 ori /direction = RIGHT /note = f1 bacteriophage origin of replication; arrow indicates direction of (+) strand synthesis	1298 .. 1726	429			rep_origin
✓ SV40 promoter /note = SV40 enhancer and early promoter	1740 .. 2069	330			promoter
✓ SV40 ori /note = SV40 origin of replication	1920 .. 2055	136			rep_origin
✓ HygR /gene = aph(4)-Ia /product = aminoglycoside phosphotransferase from E. coli /note = confers resistance to hygromycin /translation = MKKPELTATSVEKFLIEKFDSVSDLMQLSEGEESRAFSFDVGGRGYVLRVNSCADGFYKDRVYRHFASAAALPIPEVLDIGEFSESLT YCSIRRAQGVTLQDLPETELPAVLQPVAAEMDAIAAADLSQTSFGFPGPQGIQQYTTWRDFICAIADPHVYHWQTVMDDTVSA SVAQALDELMLWAEDCPEVRHLVHADFGSNNVLTDNGRITAVIDWSEAMFGDSQYEVANIFFWRPWLCMEQQTRYFERRHPEL AGSPRLRAYMLRIGLDQLYQSLVDGNFDDAAWAQGRCDIVRSGAGTVGRTQIARRSAAVWTDGCVEVLADSGNRRPSTRPRAK F* 341 amino acids = 38.0 kDa	2118 .. 3143	1026			CDS
✓ SV40 poly(A) signal /note = SV40 polyadenylation signal	3273 .. 3394	122			polyA_signal
✓ M13 rev /note = common sequencing primer, one of multiple similar variants	3443 .. 3459	17			primer_bind
✓ ori /direction = LEFT /note = high-copy-number CoIE1/pMB1/pBR322/pUC origin of replication	3845 .. 4430	586			rep_origin
✓ AmpR ▶ 2 segments /gene = bla /product = -lactamase /note = confers resistance to ampicillin, carbenicillin, and related antibiotics /translation = MSIQHFRVALIPFFAAFCPLPVFA,HPETLVKVKDAEDQLGARVGYIELDLNSGKILESFRPEERFPMMSSTFKVLLCGAVLSRIDAGQE QLGRRIHYSQNDLVEYSPVTEKHLTDGMTVRELCSAAITMSDNTAANLLLTITIGGPKELTAFLHNMGDHVTSLDRWEPELNEAIPN DERDRTMPVAMATTLRKLTLGELLTLASRQQLIDWMEADKVGAPLLRSALPAWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIY TTGSOATMDFRNRIAFIGASIKHW* 286 amino acids = 31.6 kDa	4601 .. 5461	861			CDS

Feature	Location	Size (bp)			Type
✓ AmpR promoter	5462 .. 5566	105			promoter
/gene = bla					

Description: Mammalian expression vector with the CMV promoter. The MCS is in the forward (+) orientation.

Created: Tuesday, Jul 31, 2012

Last Modified: Saturday, Jan 5, 2013

Accession Number:

Code Number:

Sequence Author: Invitrogen (Life Technologies)

DNA Type: Synthetic DNA

Laboratory Host Organism: Mammalian Cells

Bacterial Transformation Strain: Unspecified

Dam<sup>+</sup> Dcm<sup>+</sup> EcoKI<sup>+</sup>

Comments: The cloned gene must include a Kozak sequence and start codon.

References: