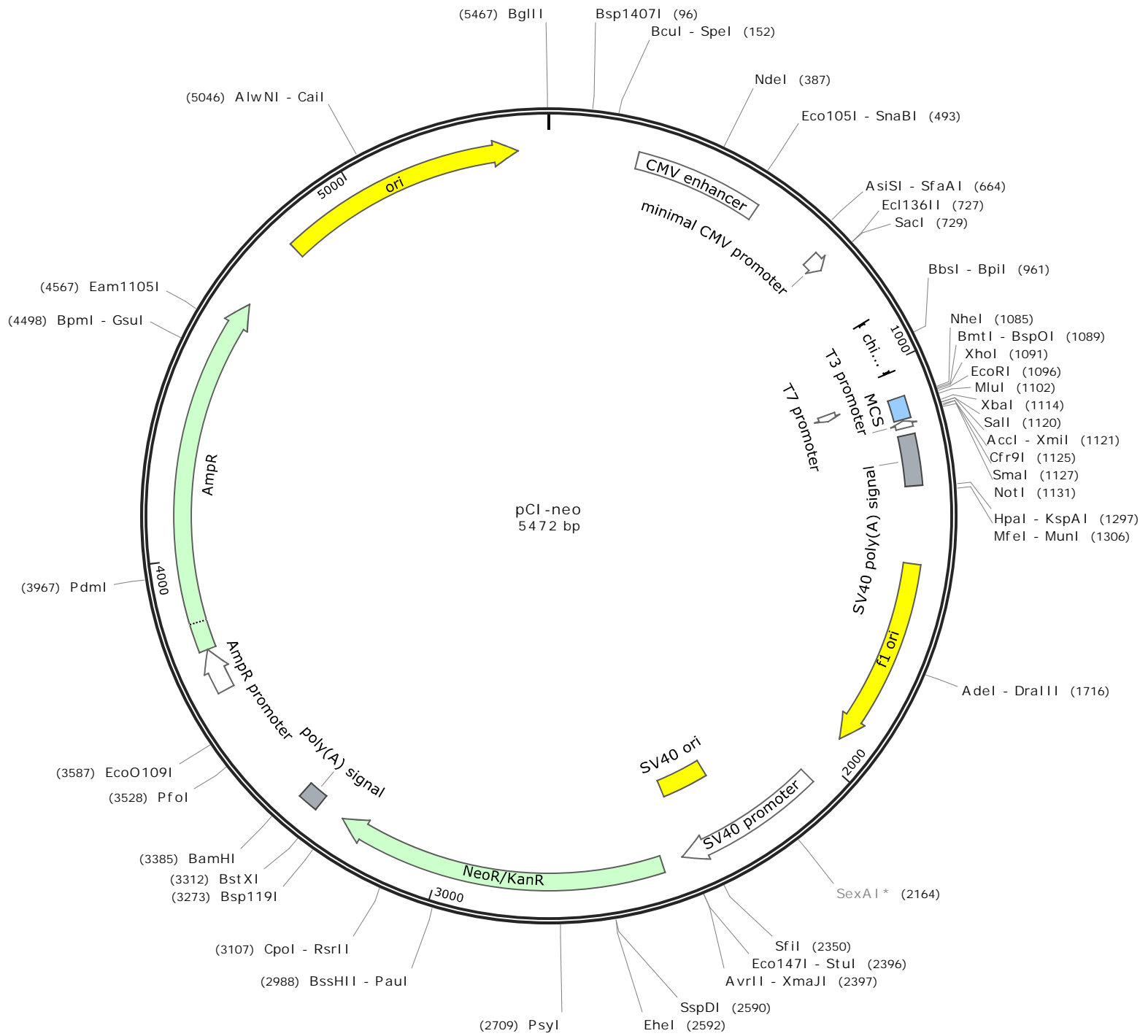


Mammalian cell expression vector with the CMV promoter and a neomycin-resistance marker.





BpiI
BbsI

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 ACCCGAACAGCTCTGTCTCTTCTGAGAACGCAAAGACTATCCGTGGATAACCAGAATGACTGTAGGTGAAACGGAAAGAGAGGTG

1020

chimeric intron

NheI BmtI BspOI XhoI EcoRI MluI

AGGTGTCCACTCCCAGTTCAATTACAGCTCTTAAGGCTAGAGTACTTAATACGACTCACTATAGGCTAGCCTCGAGAATTCACGC
 TCCACAGGTGAGGGTCAAGTTAATGTCGAGAATTCGATCTCATGAATTATGCTGAGTGATATCCGATCGGAGCTCTTAAGTGCG

1105

chimeric intron

MCS

T7 promoter

XbaI SalI XmiI Cfr9I SmaI NotI

GTGGTACCTCTAGAGTCGACCCGGGCGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATGATAAGATAATTGATG
 CACCATGGAGATCTCAGCTGGGCCCGCCGCGAAGGGAAATCACTCCAATTACGAAGCTCGTCTGTACTATTCTATGTAACTAC

1190

MCS

T3 promoter

SV40 poly(A) signal

AGTTTGGACAAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCAT
 TCAAACCTGTTTGGTGTGATCTTACGTCACCTTTTTTACGAAATAAACACTTTAAACACTACGATAACGAAATAAACATTGGTA

1275

SV40 poly(A) signal

HpaI KspAI MfeI MunI

TATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTCATTTTATGTTTCAGGTTTCAGGGGAGATGTGGGAGGTTTTTTAA
 ATATTCGACGTTATTTGTTCAATTGTTGTTGTTAACGTAAGTAAAATACAAAGTCCAAGTCCCCCTCTACACCCTCCAAAAAATT

1360

SV40 poly(A) signal

AGCAAGTAAACCTCTACAAATGTGGTAAATCCGATAAGGATCGATCCGGGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG
 TCGTTCATTTTGGAGATGTTTACACCATTTTAGGCTATTCTAGCTAGGCCCGACCGCATTATCGCTTCTCCGGGCGTGGCTAGC

1445

CCCTTCCAACAGTTGCGCAGCCTGAATGGCGAATGGACGCGCCCTGTAGCGGCGCATTAAAGCGCGGGCGGGTGTGGTGGTTACGC
 GGGAAAGGTTGTCAACGCGTCGGACTTACCGCTTACCTGCGCGGGACATCGCCGCGTAATTCGCGCCGCCACACCACCAATGCG

1530

f1 ori

GCAGCGTGACCGCTACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTTCGCTTTCTTCCCTTCTTCTCGCCACGTTTCGCCGGCTT
 CGTCGCACTGGCGATGTGAACGGTTCGCGGGATCGCGGGCGAGGAAAGCGAAAGAAGGGAAGGAAAGAGCGGTGCAAGCGGCCGAA

1615

f1 ori

TCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGATTTAGTGCTTTACGGCACCTCGACCCCAAAAACTTGATTAG
 AGGGGCGATTCGAGATTTAGCCCCGAGGGAAATCCCAAGGCTAAATCACGAAATGCCGTGGAGCTGGGGTTTTTTGAACTAATC

1700

f1 ori

Adel
DraI I

GGTGATGGTTCACGTAAGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCTTTAATAGTGGAC
CCACTACCAAGTGCATCACCCGGTAGCGGGACTATCTGCCAAAAAGCGGGAAACTGCAACCTCAGGTGCAAGAAATTATCACCTG

1785

f1 ori

TCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGATTTTGCCGATTTTCGGCCTATTG
AGAACAAGGTTTGACCTTGTGTGAGTTGGGATAGAGCCAGATAAGAAAATAAATATTCCTAAAACGGCTAAAGCCGGATAAC

1870

f1 ori

GTTAAAAAATGAGCTGATTTAACAAAAATTTAACGCGAATTTTAAACAAAATATTAACGCTTACAATTTCTGATGCGGTATTTTC
CAATTTTTTACTCGACTAAATTGTTTTTAAATTGCGCTTAAAATTGTTTTATAATTGCGAATGTTAAAGGACTACGCCATAAAAG

1955

f1 ori

TCCTTACGCATCTGTGCGGTATTTACACCCGCATACGCGGATCTGCGCAGCACCATGGCCTGAAATAACCTCTGAAAGAGGAACT
AGGAATGCGTAGACACGCCATAAAGTGTGGCGTATGCGCCTAGACGCGTCTGTGGTACCGGACTTTATTGGAGACTTTCTCCTTGA

2040

TGGTTAGGTACCTTCTGAGGCGGAAAGAACCAGCTGTGGAATGTGTGTCAGTTAGGGTGTGGAAAGTCCCCAGGCTCCCCAGCAG
ACCAATCCATGGAAGACTCCGCCTTTCTTGGTCGACACCTTACACACAGTCAATCCCACACCTTTTCAGGGGTCCGAGGGGTCTGTC

2125

SV40 promoter

SexA1 *

GCAGAAGTATGCAAAGCATGCATCTCAATTAGTCAGCAACCAGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCA
CGTCTTCATACGTTTTCGTACGTAGAGTTAATCAGTCGTTGGTCCACACCTTTTCAGGGGTCCGAGGGGTCTGTCCTTCATACGT

2210

SV40 promoter

AAGCATGCATCTCAATTAGTCAGCAACCATAGTCCCGCCCCTAACTCCGCCCATCCCGCCCCTAACTCCGCCCAGTTCCGCCCAT
TTCGTACGTAGAGTTAATCAGTCGTTGGTATCAGGGCGGGGATTGAGGCGGGTAGGGCGGGGATTGAGGCGGGTCAAGGCGGGTA

2295

SV40 promoter

SV40 ori

SfiI

TCTCCGCCCCATGGCTGACTAATTTTTTTTATTTATGAGAGGCGGAGGCGCCCTCGGCCTCTGAGCTATTCCAGAAGTAGTGAG
AGAGGCGGGGTACCGACTGATTAATAAAAAATAAATACGTCTCCGGCTCCGGCGGAGCCGGAGACTCGATAAGGTCTTCATCACTC

2380

SV40 promoter

SV40 ori

StuI AvrII
Eco147I XmaJI

GAGGCTTTTTTGGAGGCCTAGGCTTTTTGCAAAAAGCTTGATTCTTCTGACACAACAGTCTCGAACTTAAGGCTAGAGCCACCATG
CTCCGAAAAAACCTCCGGATCCGAAAACGTTTTTTCGAACTAAGAAGACTGTGTTGTGAGAGCTTGAATTCGATCTCGGTGGTAC

2465

SV40 promoter

SV40 ori

1
M
NeoR/KanR

ATTGAACAAGATGGATTGCACGCAGGTTCTCCGGCCGCTTGGGTGGAGAGGCTATTCCGGCTATGACTGGGCACAACAGACAATCG
 TAACCTGTTCTACCTAACGTGCGTCCAAGAGGCCGGCGAACCACCTCTCCGATAAGCCGATACTGACCCGTGTTGTCTGTTAGC
 I E Q D G L H A G S P A A W V E R L F G Y D W A Q Q T I
 NeoR/KanR

GCTGCTCTGATGCCGCCGTGTTCCGGCTGTCAGCGCAGGGGCGCCCGTTCTTTTTGTCAAGACCGACCTGTCCGGTGCCCTGAA
 CGACGAGACTACGGCGGCACAAGGCCGACAGTCGCGTCCCCGCGGGCCAAGAAAAACAGTTCTGGCTGGACAGGCCACGGGACTT
 G C S D A A V F R L S A Q G R P V L F V K T D L S G A L N
 NeoR/KanR

TGAACTGCAGGACGAGGCAGCGCGGCTATCGTGGCTGGCCACGACGGGCGTTCTTGGCGAGCTGTGCTCGACGTTGTCACTGAA
 ACTTGACGTCTGCTCCGTGCGCGCCGATAGCACCGACCGGTGCTGCCCGCAAGGAACGCGTGCACACGAGCTGCAACAGTGACTT
 E L Q D E A A R L S W L A T T G V P C A A V L D V V T E
 NeoR/KanR

GCGGGAAGGGACTGGCTGCTATTGGGCGAAGTGCCGGGGCAGGATCTCCTGTCATCTCACCTTGCTCCTGCCGAGAAAGTATCCA
 CGCCCTTCCCTGACCGACGATAACCCGCTTACGGCCCCGTCTAGAGGACAGTAGAGTGAACGAGGACGGCTCTTTCATAGGT
 A G R D W L L L G E V P G Q D L L S S H L A P A E K V S
 NeoR/KanR

TCATGGCTGATGCAATGCGGCGGCTGCATACGCTTGATCCGGCTACCTGCCCATTCGACCACCAAGCGAAACATCGCATCGAGCG
 AGTACCGACTACGTTACGCCGCCGACGTATGCGAACTAGGCCGATGGACGGGTAAGCTGGTGGTTGCTTTGTAGCGTAGCTCGC
 I M A D A M R R L H T L D P A T C P F D H Q A K H R I E R
 NeoR/KanR

AGCACGTA CTCCGATGGAAGCCGGTCTTGTGTCGATCAGGATGATCTGGACGAAGAGCATCAGGGGCTCGCGCCAGCCGAACTGTTC
 TCGTGCATGAGCCTACCTTCGGCCAGAACAGCTAGTCCTACTAGACCTGCTTCTCGTAGTCCCCGAGCGCGGTGCGCTTGACAAG
 A R T R M E A G L V D Q D D L D E E H Q G L A P A E L F
 NeoR/KanR

GCCAGGCTCAAGGCGCGCATGCCCGACGGCGAGGATCTCGTCTGACCCATGGCGATGCCTGCTTGCCGAATATCATGGTGGAAA
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 A R L K A R M P D G E D L V V T H G D A C L P N I M V E
 NeoR/KanR

ATGGCCGCTTTTCTGGATTCATCGACTGTGGCCGGCTGGGTGTGGCGGACCGCTATCAGGACATAGCGTTGGCTACCCGTGATAT
 TACCGGCGAAAAGACCTAAGTAGCTGACACCGGCCGACCCACACCGCCTGGCGATAGTCCTGTATCGCAACCGATGGGCACTATA
 N G R F S G F I D C G R L G V A D R Y Q D I A L A T R D I
 NeoR/KanR

TGCTGAAGAGCTTGGCGGCGAATGGGCTGACCGCTTCCTCGTGCTTTACGGTATCGCCGCTCCCGATTTCGCAGCGCATCGCCTTC
 ACGACTTCTCGAACCGCCGCTTACCCGACTGGCGAAGGAGCACGAAATGCCATAGCGGCGAGGGCTAAGCGTCGCGTAGCGGAAG
 230 235 240 245 250 255
 A E E L G G E W A D R F L V L Y G I A A P D S Q R I A F
 NeoR/KanR

TATCGCCTTCTTGACGAGTTCTTCTGAGCGGGACTCTGGGGTTCGAAATGACCGACCAAGCGACGCCAACCTGCCATCACGATG
 ATAGCGGAAGAACTGCTCAAGAAGACTCGCCCTGAGACCCCAAGCTTTACTGGCTGGTTCGCTGCGGGTTGGACGGTAGTGCTAC
 260 265
 Y R L L D E F F
 NeoR/KanR

GCCGCAATAAAATATCTTTATTTTATTACATCTGTGTGTTGGTTTTTGTGTGAATCGATAGCGATAAGGATCCGCGTATGGTG
 CGGCGTTATTTTATAGAAATAAAAGTAATGTAGACACACAACCAAAAAACACACTTAGCTATCGCTATTCTAGGCGCATACCAC
 poly(A) signal
 BamHI

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 TTGCGCGCTCTGCTTTCCCGGAGCACTATGCGGATAAAAATATCCAATTACAGTACTATTATTACCAAAGAATCTGCAGTCCACC
 EcoO109I

CACTTTTCGGGGAAATGTGCGCGGAACCCCTATTTGTTTATTTTTCTAAATACATTCAAATATGTATCCGCTCATGAGACAATAA
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 AmpR promoter

CCCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTTTCGGG
 GGGACTATTTACGAAGTTATTATAACTTTTTCTTCTCATACTCATAAGTTGTAAGGCACAGCGGGAATAAGGGAAAAAACGCC
 AmpR promoter
 1 5 10 15
 M S I Q H F R V A L I P F F A
 signal sequence
 AmpR

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 GTAAAACGGAAGGACAAAACGAGTGGGTCTTTGCGACCACTTTTCAATTTCTACGACTTCTAGTCAACCCACGTGCTCACCCAAT
 20 25 30 35 40
 A F C L P V F A H P E T L V K V K D A E D Q L G A R V G Y
 signal sequence
 AmpR

PdmI

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 GTAGCTTGACCTAGAGTTGTCGCCATTCTAGGAACTCTCAAAGCGGGGCTTCTTGCAAAGGTTACTACTCGTGAAAATTTCAA
 45 50 55 60 65 70
 I E L D L N S G K I L E S F R P E E R F P M M S T F K V
 AmpR

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 GACGATACACCGCGCCATAATAGGGCATAACTGCGGCCCGTTCTCGTTGAGCCAGCGGCGTATGTGATAAGAGTCTTACTGAACC
 75 80 85 90 95 100
 L L C G A V L S R I D A G Q E Q L G R R I H Y S Q N D L
 AmpR

TTGAGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTGA
 AACTCATGAGTGGTCAGTGTCTTTTCGTAGAATGCCTACCGTACTGTCACTTCTTTAATACGTACAGACGGTATTGGTACTCACT
 105 110 115 120 125
 V E Y S P V T E K H L T D G M T V R E L C S A A I T M S D
 AmpR

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 130 135 140 145 150 155
 N T A A N L L L T T I G G P K E L T A F L H N M G D H V
 AmpR

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 160 165 170 175 180 185
 T R L D R W E P E L N E A I P N D E R D T T M P V A M A
 AmpR

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 AmpR

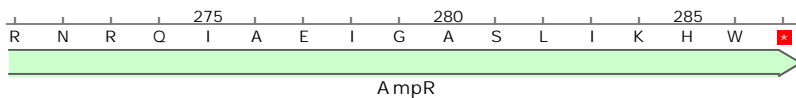
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 A G P L L R S A L P A G W F I A D K S G A G E R G S R G
 AmpR

Eam1105I

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 245 250 255 260 265 270
 I I A A L G P D G K P S R I V V I Y T T G S Q A T M D E
 AmpR

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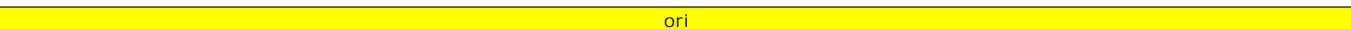


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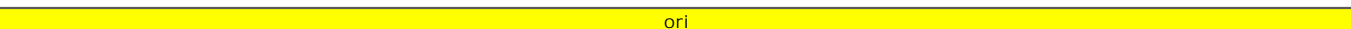
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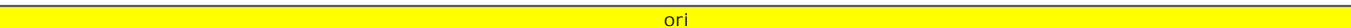


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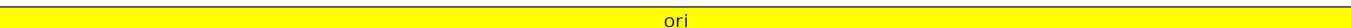
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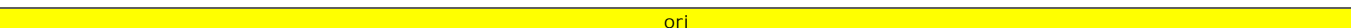
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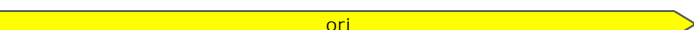
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










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







BglI

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 GACCGGAAAAACGAGTGTACCGAGCTGTCTAGA 5'

Enzymes	Sites	
AccI	1	1121
Adel	1	1716
AlwNI	1	5046
AsiSI	1	664
AvrII	1	2397
BamHI	1	3385
BbsI	1	961
BcuI	1	152
BglII	1	5467
BmtI	1	1089
BpiI	1	961
Bpml	1	4498
Bsp119I	1	3273
Bsp1407I	1	96
BspOI	1	1089
BssHII	1	2988
BstXI	1	3312
CaiI	1	5046
Cfr9I	1	1125
CpoI	1	3107
DraIII	1	1716
Eam1105I	1	4567
Ecl136II	1	727
Eco105I	1	493
Eco147I	1	2396
EcoO109I	1	3587
EcoRI	1	1096
EheI	1	2592
GsuI	1	4498
HpaI	1	1297
KspAI	1	1297
MfeI	1	1306
MluI	1	1102
MunI	1	1306
NdeI	1	387
NheI	1	1085
NotI	1	1131
PaulI	1	2988
Pdml	1	3967
PfoI	1	3528
PsyI	1	2709
RsrII	1	3107
SacI	1	729
Sall	1	1120
SexAI	1*	2164*
SfaAI	1	664
SfiI	1	2350
SmaI	1	1127
SnaBI	1	493
SpeI	1	152
SspDI	1	2590
StuI	1	2396
XbaI	1	1114
XhoI	1	1091
XmaJI	1	2397
XmiI	1	1121

Feature	Location	Size (bp)			Type
✓ CMV enhancer /note = human cytomegalovirus immediate early enhancer	213 .. 517	305			enhancer
✓ minimal CMV promoter /note = human cytomegalovirus (CMV) immediate early promoter	691 .. 729	39			promoter
✓ chimeric intron /note = chimera between introns from human β -globin and immunoglobulin heavy chain genes	890 .. 1022	133			intron
✓ T7 promoter /note = promoter for bacteriophage T7 RNA polymerase	1067 .. 1085	19			promoter
✓ MCS /note = multiple cloning site	1085 .. 1137	53			misc_feature
✓ T3 promoter /note = promoter for bacteriophage T3 RNA polymerase (shorter by one base than the standard T3 promoter)	1141 .. 1158	18			promoter
✓ SV40 poly(A) signal /note = SV40 polyadenylation signal	1176 .. 1297	122			polyA_signal
✓ f1 ori /direction = RIGHT /note = f1 bacteriophage origin of replication; arrow indicates direction of (+) strand synthesis	1483 .. 1938	456			rep_origin
✓ SV40 promoter /note = SV40 enhancer and early promoter	2055 .. 2412	358			promoter
✓ SV40 ori /note = SV40 origin of replication	2263 .. 2398	136			rep_origin
✓ NeoR/ KanR /gene = aph(3')-II (or nptII) /product = aminoglycoside phosphotransferase from Tn5 /note = confers resistance to neomycin, kanamycin, and G418 (Geneticin) /translation = MIEQDGLHAGSPAAWVERLFGYDWAQQTIGCSDAAVFRLSAQGRPVLFVKTDLSGALNELQDEAARLSWLATTGVPCA AVL DVV TEAGRDWLLLGEVPGQDLLSSHLAPA EKVSIMADAMRRLHTLDPATCPF DHQAKHRIERARTRMEAGLVDQDDLDEEHQGLAPAE LFARLKARMPDGEDLVVTHGDA CLPNIMVENGRFSGFIDCGR LGVADRYQDIALATRDIAEELGGEWADRFLVLYGIAAPDSORIA FYRIIDFFF* 264 amino acids = 29.0 kDa	2463 .. 3257	795			CDS
✓ poly(A) signal /note = synthetic polyadenylation signal	3321 .. 3369	49			polyA_signal
✓ AmpR promoter /gene = bla	3675 .. 3779	105			promoter

Feature	Location	Size (bp)			Type
✓ AmpR	3780 .. 4640	861			CDS
▶ 2 segments					
/gene	= bla				
/product	= -lactamase				
/note	= confers resistance to ampicillin, carbenicillin, and related antibiotics				
/translation	= MSIQHFRVALIPFFAAFCLPVFA,HPETLVKVKDAEDQLGARVGYIELDLNSGKILESFRPEERFPMSTFKVLLCGAVLSRIDAGQE QLGRRIHYSQNDLVEYSPVTEKHLTDGMTVRELCSAAITMSDNTAANLLTTIGGPKELTAFLHNMGDHVTRLDRWEPELNEAIPN DERDTTMPVAMATTLRKLTTGELLTASRQQLIDWMEADKVGPLLRSAALPAWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIY TTGSOATMDFRNROIAFIGASIKHW* 286 amino acids = 31.6 kDa				
✓ ori	4811 .. 5399	589			rep_origin
/direction	= RIGHT				
/note	= high-copy-number ColE1/pMB1/pBR322/pUC origin of replication				

Description: Mammalian cell expression vector with the CMV promoter and a neomycin-resistance marker.

Created: Friday, Jun 23, 2000

Last Modified: Saturday, Jan 5, 2013

Accession Number: U47120

Code Number:

Sequence Author: Promega

DNA Type: Synthetic DNA

Laboratory Host Organism: Mammalian Cells

Bacterial Transformation Strain: Unspecified

Dam⁺ Dcm⁺ EcoKI⁺

Comments:

References: