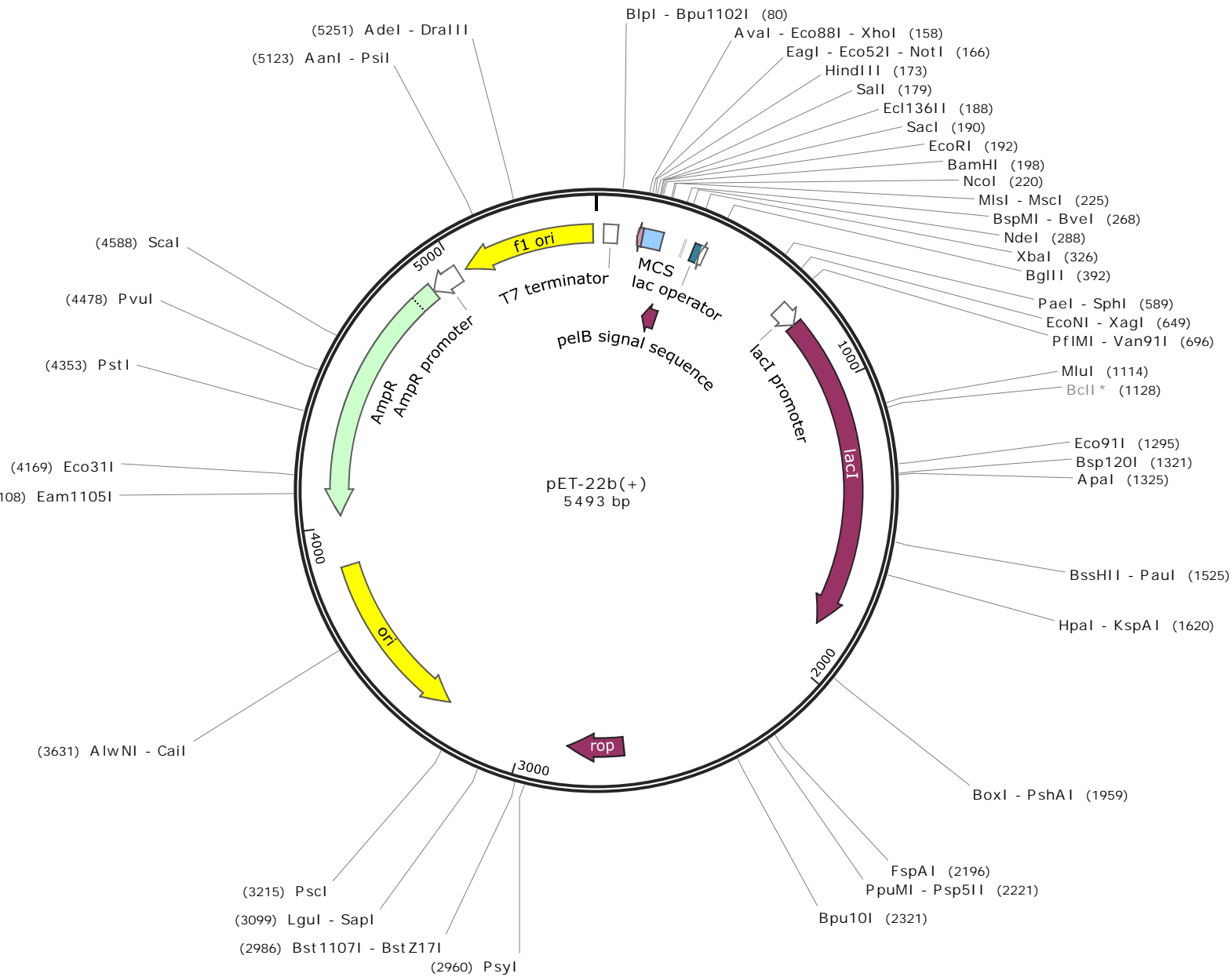


Bacterial vector that encodes a signal sequence for inducible expression of proteins in the periplasm.

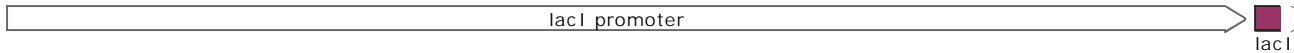




PfIMI  
Van91I

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765



GAAACCAGTAACGTTATACGATGTCGCAGAGTATGCCGGTGTCTCTTATCAGACCGTTTTCCCGCGTGGTGAACCAGGCCAGCCAC  
CTTTGGTCATTGCAATATGCTACAGCGTCTCATACGGCCACAGAGAATAGTCTGGCAAAGGGCGCACCACTTGGTCCGGTCCGGTG

850

K P V T L Y D V A E Y A G V S Y Q T V S R V V N Q A S H  
lacI

GTTTCTGCGAAAACGCGGGAAAAAGTGAAGCGGCGATGGCGGAGCTGAATTACATTCCEAACCGCGTGGCACAACAACCTGGCGG  
CAAAGACGCTTTTTCGCGCCCTTTTTCACCTTCGCGGCTACCGCCTCGACTTAATGTAAGGGTTGGCGCACCGTGTGTTGACCGCC

935

V S A K T R E K V E A A M A E L N Y I P N R V A Q Q L A  
lacI

GCAAACAGTCGTTGCTGATTGGCGTTGCCACCTCCAGTCTGGCCCTGCACGCGCCGTCGCAAATTGTCGCGGGCGATTAAATCTCG  
CGTTTGTGAGCAACGACTAACCAGCAACGGTGGAGGTCAGACCGGGACGTGCGCGGCAGCGTTTAACAGCGCCGCTAATTTAGAGC

1020

G K Q S L L I G V A T S S L A L H A P S Q I V A A I K S R  
lacI

CGCCGATCAACTGGGTGCCAGCGTGGTGGTGTGATGGTAGAACGAAGCGGGCGTGAAGCCTGTAAAGCGGGCGGTGCACAATCTT  
GCGGCTAGTTGACCCACGGTTCGCACCACCACAGCTACCATCTTGCTTCGCCGACGCTTCGGACATTTTCGCCGCCACGTGTTAGAA

1105

A D Q L G A S V V V S M V E R S G V E A C K A A V H N L  
lacI

MluI

BclI \*

CTCGCGCAACGCGTCAGTGGGCTGATCATTAACTATCCGCTGGATGACCAGGATGCCATTGCTGTGGAAGCTGCCTGCACTAATG  
GAGCGCGTTGCGCAGTCACCCGACTAGTAATTGATAGGCGACCTACTGGTCTACGGTAACGACACCTTCGACGGACGTGATTAC

1190

L A Q R V S G L I I N Y P L D D Q D A I A V E A A C T N  
lacI

TTCCGGCGTTATTTCTTGATGTCTCTGACCAGACACCCATCAACAGTATTATTTTCTCCCATGAAGACGGTACGCGACTGGGCGT  
AAGGCCGCAATAAAGAACTACAGAGACTGGTCTGTGGGTAGTTGTGATAATAAAGAGGGTACTTCTGCCATGCGCTGACCCGCA

1275

V P A L F L D V S D Q T P I N S I I F S H E D G T R L G V  
lacI

Eco91I

Bsp120I

ApaI

GGAGCATCTGGTTCGATTGGGTACCCAGCAAATCGCGCTGTTAGCGGGCCCATTAAGTTCTGTCTCGGCGCGTCTGCGTCTGGCT  
CCTCGTAGACCAGCGTAACCCAGTGGTTCGTTAGCGCGACAATCGCCGGGTAATTCAGACAGAGCCGCGCAGACGCAGACCGA

1360

E H L V A L G H Q Q I A L L A G P L S S V S A R L R L A  
lacI

GGCTGGCATAAATATCTCACTCGCAATCAAATTCAGCCGATAGCGGAACGGGAAGGCGACTGGAGTGCCATGTCCGGTTTTCAAC  
 CCGACCGTATTTATAGAGTGAGCGTTAGTTTAAAGTCGGCTATCGCCTTGCCTTCCGCTGACCTCACGGTACAGGCCAAAAGTTG  
 200 205 210 215 220 225  
 G W H K Y L T R N Q I Q P I A E R E G D W S A M S G F Q  
 lacI

AAACCATGCAAATGCTGAATGAGGGCATCGTTCCCACTGCGATGCTGGTTGCCAACGATCAGATGGCGCTGGGCGCAATGCGCGC  
 TTTGGTACGTTTACGACTTACTCCCGTAGCAAGGGTGACGCTACGACCAACGGTTGCTAGTCTACCGCGACCCGCGTTACGCGCG  
 230 235 240 245 250 255  
 Q T M Q M L N E G I V P T A M L V A N D Q M A L G A M R A  
 lacI Paul  
 BssHI I

CATTACCGAGTCCGGGCTGCGCGTTGGTGCGGATATCTCGGTAGTGGGATACGACGATACCGAAGACAGCTCATGTTATATCCCG  
 GTAATGGCTCAGGCCCGACGCGCAACCACGCCTATAGAGCCATCACCCATGCTGCTATGGCTTCTGTGAGTACAATATAGGGC  
 260 265 270 275 280  
 I T E S G L R V G A D I S V V G Y D T E D S S C Y I P  
 lacI

CCGTTAACACCACCATCAAACAGGATTTTCGCCTGCTGGGGCAAACCAGCGTGGACCGCTTGTGCAACTCTCTCAGGGCCAGGCGG  
 GGCAATTGGTGGTAGTTTGTCTAAAGCGGACGACCCCGTTTGGTTCGCACCTGGCGAACGACGTTGAGAGAGTCCCGGTCCGCC  
 285 290 295 300 305 310  
 P L T T I K Q D F R L L G Q T S V D R L L Q L S Q G Q A  
 lacI HpaI  
 KspAI

TGAAGGGCAATCAGCTGTTGCCCGTCTCACTGGTGAAAAGAAAAACCACCCTGGCGCCCAATACGCAAACCAGCCTCTCCCCGCGC  
 ACTTCCCCTTAGTCGACAACGGGCAGAGTGACCACTTTTCTTTTTGGTGGGACCGCGGGTTATGCGTTTGGCGGAGAGGGGCGCG  
 315 320 325 330 335 340  
 V K G N Q L L P V S L V K R K T T L A P N T Q T A S P R A  
 lacI

GTTGGCCGATTCATTAATGCAGCTGGCACGACAGGTTTCCCGACTGGAAAGCGGGCAGTGAGCGCAACGCAATTAATGTAAGTTA  
 CAACCGGCTAAGTAATTACGTCGACCGTGCTGTCCAAAGGGCTGACCTTTCGCCCGTCACTCGCGTTGCGTTAATTACATTCAAT  
 345 350 355 360  
 L A D S L M Q L A R Q V S R L E S G Q  
 lacI

GCTCACTCATTAGGCACCGGGATCTCGACCGATGCCCTTGAGAGCCTTCAACCCAGTCAGCTCCTTCCGGTGGGCGCGGGGCATG  
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CACTGGTCCCGCCACCAAACGTTTCGGCGAGAAGCAGGCCATTATCGCCGGCATGGCGGCCCCACGGGTGCGCATGATCGTGCTC  
 GTGACCAGGGCGGTGGTTTCAAAGCCGCTCTTCGTCGGTAATAGCGGCCGTACCGCCGGGGTGCCACGCGTACTAGCACGAG  
 FspAI

PpuMI  
Psp5I

CTGTCGTTGAGGACCCGGCTAGGCTGGCGGGGTTGCCTTACTGGTTAGCAGAATGAATCACCGATACGCGAGCGAACGTGAAGCG  
GACAGCAACTCCTGGGCCGATCCGACCGCCCCAACGGAATGACCAATCGTCTTACTTAGTGGCTATGCGCTCGCTTGCACCTTCGC

2295

Bpu10I

ACTGCTGCTGCAAAACGTCTGCGACCTGAGCAACAACATGAATGGTCTTCGGTTTTCCGTGTTTTCGTAAAGTCTGGAAACGCGGAA  
TGACGACGACGTTTTGCAGACGCTGGACTCGTTGTTGTACTTACCAGAAGCCAAAGGCACAAAGCATTTCAGACCTTTGCGCCTT

2380

GTCAGCGCCCTGCACCATTATGTTCCGGATCTGCATCGCAGGATGCTGCTGGCTACCCTGTGGAACACCTACATCTGTATTAACG  
CAGTCGCGGGACGTGGTAATAACAAGGCCTAGACGTAGCGTCTACGACGACCGATGGGACACCTTGTGGATGTAGACATAATTGC

2465

AAGCGCTGGCATTGACCCTGAGTGATTTTTCTCTGGTCCC GCCGCATCCATACCGCCAGTTGTTTACCCTCACAACGTTCCAGTA  
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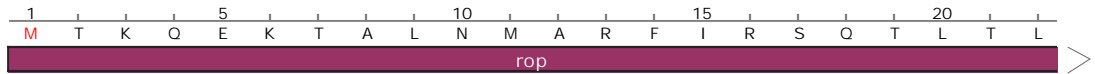
2550

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2635

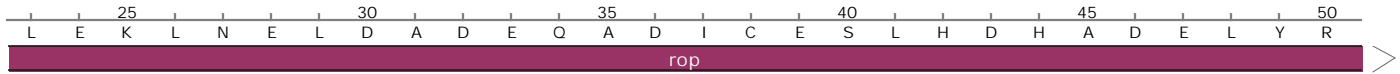
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2720



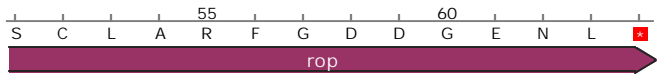
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2805



GCTGCCTCGCGGTTTTGGTGATGACGGTGAAAACCTCTGACACATGCAGCTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGA  
CGACGGAGCGCGCAAAGCCACTACTGCCACTTTTGGAGACTGTGTACGTCGAGGGCCCTCTGCCAGTGTGCAACAGACATTGCGCT

2890



PsuI

TGCCGGGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGTGTGGGGGCGCAGCCATGACCCAGTCACGTAGCGAT  
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2975

BstZ17I  
Bst1107I

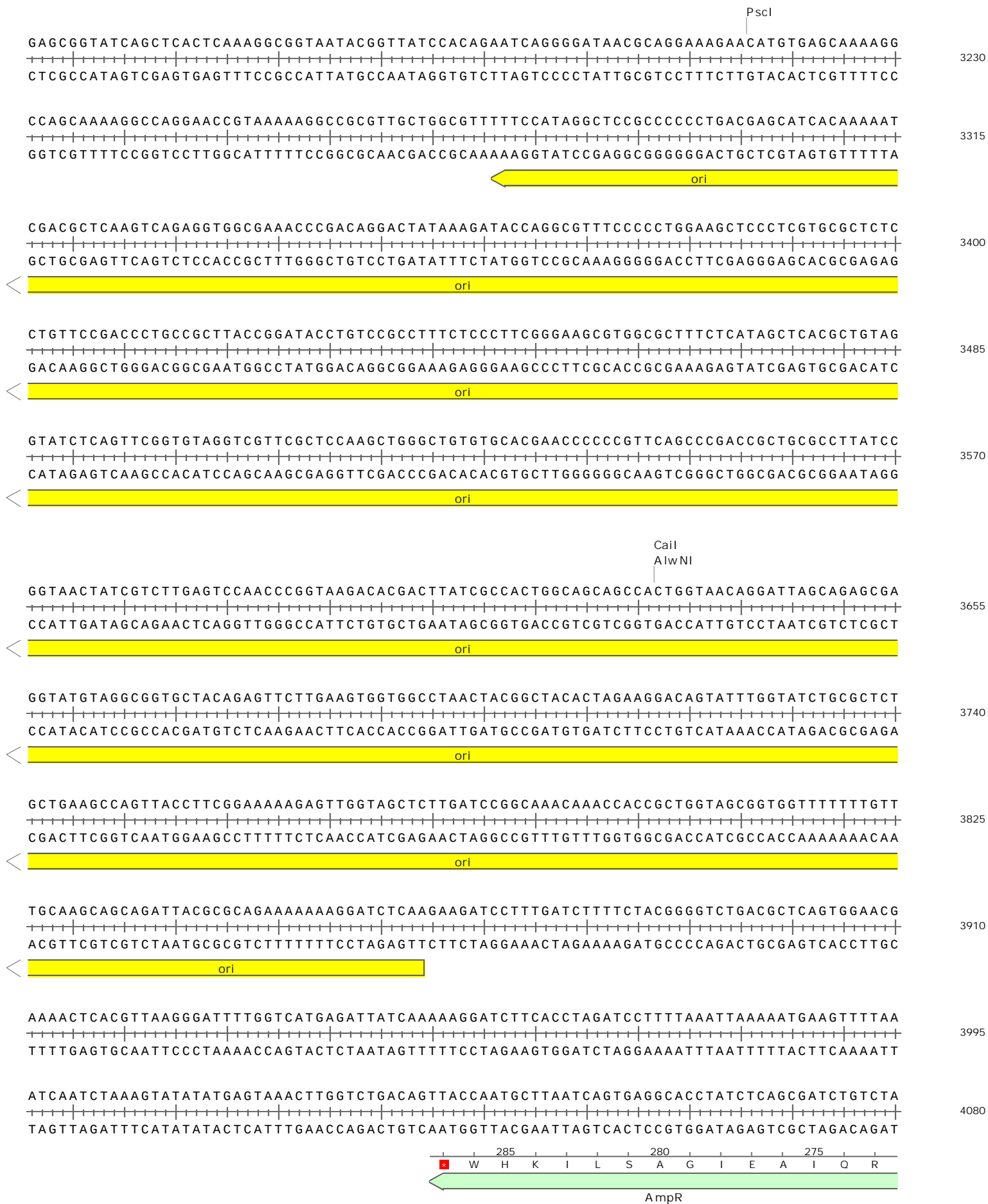
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3060

LguI  
SapI

AGATGCGTAAGGAGAAAATACCGCATCAGGCCTCTTCCGCTTCTCGCTCACTGACTCGCTGCGCTCGGTGCTTCCGCTGCGGC  
TCTACGCATTCTCTTTTATGGCGTAGTCCGCGAGAAGGCGAAGGAGCGAGTGACTGAGCGACGCGAGCCAGCAAGCCGACGCCG

3145



Eam1105I

TTTCGTTTCATCCATAGTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAA  
 AAAGCAAGTAGGTATCAACGGACTGAGGGGCAGCACATCTATTGATGCTATGCCCTCCCGAATGGTAGACCGGGGTCACGACGTT

4165

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

AmpR

Eco31I

TGATACCGCGAGACCCACGCTCACCGGCTCCAGATTTATCAGCAATAAACAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGGTCC  
 ACTATGGCGCTCTGGGTGCGAGTGGCCGAGGTCTAAATAGTCGTTATTTGGTCGGTCGGCCTTCCCGGCTCGCGTCTTACCAGG

4250

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

AmpR

TGCAACTTTATCCGCCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTTGCGCAAC  
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4335

A V K D A E M W D I L Q O R S A L T L L E G T L L K R L

AmpR

PstI

GTTGTTGCCATTGCTGCAGGCATCGTGGTGTACGCTCGTCGTTTGGTATGGCTTCATTCAGCTCCGGTTCCTCAACGATCAAGGC  
 CAACAACGGTAACGACGTCCTGAGCACCACAGTGCAGCAGCAAACCATAACGAAGTAAGTCGAGGGCCAAGGGTTGCTAGTTCCG

4420

T T A M A A P M T T D R E D N P I A E N L E P E W R D L R

AmpR

PvuI

GAGTTACATGATCCCCATGTTGTGCAAAAAGCGGTTAGCTCCTTCGGTCCTCCGATCGTTGTCAGAAGTAAGTTGGCCGCAGT  
 CTCAATGTAAGGGGTACAACACGTTTTTTCGCCAATCGAGGAAGCCAGGAGGCTAGCAACAGTCTTCATTCAACCGGCCTCA

4505

T V H D G M N H L F A T L E K P G G I T T L L L N A A T

AmpR

ScaI

GTTATCACTCATGGTTATGGCAGCACTGCATAATTCTCTTACTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTAC  
 CAATAGTGAGTACCAATACCGTCGTGACGTATTAAGAGAATGACAGTACGGTAGGCATTCTACGAAAAGACACTGACCACTCATG

4590

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

AmpR

TCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGCCACATA  
 AGTTGGTTTACAGTAAGACTCTTATCACATACGCCGCTGGCTCAACGAGAACGGGCGCAGTTATGCCCTATTATGGCGCGGTGTAT

4675

E V L D N Q S Y H I R R G L Q E Q G A D I R S L V A G C L

AmpR

GCAGAACTTTAAAAGTGCTCATCATTGGAAAACGTTCTTCGGGGCGAAAACCTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTC  
CGTCTTGAAATTTTACGAGTAGTAACCTTTTGCAAGAAGCCCCGCTTTTGAGAGTTCCTAGAATGGCGACAACCTCTAGGTCAAG

4760

L V K F T S M M P F R E E P R F S E L I K G S N L D L E

AmpR

GATGTAACCCACTCGTGCACCCAACCTGATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAA  
CTACATTGGGTGAGCACGTGGGTTGACTAGAAAGTCGTAGAAAATGAAAGTGGTCGCAAAGACCCACTCGTTTTTGTCTTCCGTT

4845

I Y G V R A G L Q D E A D K V K V L T E P H A F V P L C

AmpR

signal sequence

AATGCCGCAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTTCAATATTATTGAAGCATTATC  
TTACGGCGTTTTTTCCCTTATTTCCCGCTGTGCCTTTACAACCTATGAGTATGAGAAGGAAAAAGTTATAATAACTTCGTAATAG

4930

F A A F F P I L A V R F H Q I S M

signal sequence

AmpR

AmpR promoter

AGGGTTATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAA  
TCCCAATAACAGAGTACTCGCCTATGTATAAACTTACATAAATCTTTTTATTTGTTTATCCCCAAGGCGCGTGTAAAGGGGCTTT

5015

AmpR promoter

AGTGCCACCTGAAATTGTAACGTTAATATTTTGTAAAATTCGCGTTAAATTTTTGTTAAATCAGCTCATTTTTTAACCAATAG  
TCACGGTGGACTTTAACATTTGCAATTATAAAACAATTTTAAAGCGCAATTTAAAAACAATTTAGTCGAGTAAAAAATTGGTTATC

5100

f1 ori

AanI  
PstI

GCCGAAATCGGCAAAATCCCTTATAAATCAAAGAATAGACCGAGATAGGGTTGAGTGTTGTTCCAGTTTGAACAAGAGTCCAC  
CGGCTTTAGCCGTTTTAGGGAATATTTAGTTTTCTTATCTGGCTCTATCCCAACTCACAACAAGGTCAAACCTTGTCTCAGGTG

5185

f1 ori

Adel  
DraIII

TATTAAGAACGTGGACTCCAACGTCAAAGGGCGAAAAACCGTCTATCAGGGCGATGGCCCACTACGTGAACCATCACCTAATC  
ATAATTTCTTGACCTGAGGTTGCAGTTTTCCCGCTTTTTGGCAGATAGTCCCGCTACCGGGTGATGCACTTGGTAGTGGGATTAG

5270

f1 ori

AAGTTTTTTGGGGTCGAGGTGCCGTAAGCACTAAATCGGAACCTAAAGGGAGCCCCGATTTAGAGCTTGACGGGGAAAGCCG  
TTCAAAAACCCAGCTCCACGGCATTTCGTGATTTAGCCTTGGGATTTCCCTCGGGGGCTAAATCTCGAACTGCCCTTTCCGGC

5355

f1 ori

GCGAACGTGGCGAGAAAGGAAGGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGATGCGGTACGCTGCGCGTAA  
CGCTTGCACCGCTCTTTCCCTTCCCTTCTTTCCGCTTTCCCTCGCCCGGATCCCGCGACCGTTCACATCGCCAGTGCGACGCGCATT

5440

f1 ori

























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GGTGGTGTGGGCGGCGCAATTACGCGGCGATGTCCCGCGCAGGGTAAGCGGT 5'









5493

f1 ori



Enzymes	Sites	
AanI	1	5123
Adel	1	5251
AlwNI	1	3631
ApaI	1	1325
AvaI	1	158
BamHI	1	198
BclI	1*	1128*
BglII	1	392
BipI	1	80
BoxI	1	1959
Bpu10I	1	2321
Bpu1102I	1	80
Bsp120I	1	1321
BspMI	1	268
BssHII	1	1525
Bst1107I	1	2986
BstZ17I	1	2986
BveI	1	268
CaII	1	3631
DraIII	1	5251
EagI	1	166
Eam1105I	1	4108
Ecl136II	1	188
Eco31I	1	4169
Eco52I	1	166
Eco88I	1	158
Eco91I	1	1295
EcoNI	1	649
EcoRI	1	192
FspAI	1	2196
HindIII	1	173
HpaI	1	1620
KspAI	1	1620
LguI	1	3099
MisI	1	225
MluI	1	1114
MscI	1	225
NcoI	1	220
NdeI	1	288
NotI	1	166
PaeI	1	589
Paul	1	1525
PfIMI	1	696
PpuMI	1	2221
PscI	1	3215
PshAI	1	1959
PsiI	1	5123
Psp5I	1	2221
PstI	1	4353
PsyI	1	2960
PvuI	1	4478
SacI	1	190
Sall	1	179
SapI	1	3099
ScaI	1	4588
SphI	1	589
Van91I	1	696
XagI	1	649
XbaI	1	326
XhoI	1	158

Feature	Location	Size (bp)			Type
✓ T7 terminator /note = transcription terminator for bacteriophage T7 RNA polymerase	26 .. 73	48			terminator
✓ 6xHis /product = 6xHis affinity tag /translation = HHHHHH 6 amino acids = 840.9 Da	140 .. 157	18			CDS
✓ MCS /note = multiple cloning site	158 .. 225	68			misc_feature
✓ pelB signal sequence /gene = pelB (fragment) /product = leader peptide for secretion /translation = MKYLLPTAAAGLLLLAAQPAMA 22 amino acids = 2.2 kDa	224 .. 289	66			CDS
✓ RBS /note = ribosome binding site	298 .. 303	6			RBS
✓ lac operator /bound_moiety = lac repressor encoded by lacI /note = The lac repressor binds to the lac operator to inhibit transcription in <i>E. coli</i> . This inhibition can be relieved by adding lactose or isopropyl-β-D-thiogalactopyranoside (IPTG).	334 .. 358	25			protein_bind
✓ T7 promoter /note = promoter for bacteriophage T7 RNA polymerase	359 .. 377	19			promoter
✓ lacI promoter /gene = lacI	686 .. 763	78			promoter
✓ lacI /gene = lacI /product = lac repressor /note = The lac repressor binds to the lac operator to inhibit transcription in <i>E. coli</i> . This inhibition can be relieved by adding lactose or isopropyl-β-D-thiogalactopyranoside (IPTG). /translation = VKPVTLYDVAEYAGVSYQTVSRVFNQASHVSAKTRKVEAAMAELNYIPNRVAQQLAGKQSLIGVATSSLALHAPSQIVAAIKS RADQLGASVVVSMVERSGVEACKAAVHNLLAQRVSGLIINYP LDDQDAIAVEAACTNVPALFLDVS DQTPINSIIFSHEDGTRLG VEHLVALGHQQIALLAGPLSSVSARLRLAGWHKYLTRNQIQPIAERE GDWSAMSGFQQT MQMLNEGIVPTAMLVANDQMALGAM RAITESGLRVGADISVVG YDDTEDSSCYIPPLTTIKQDFRLLGQTSVDRLLQLSQGQAVKGNQLLPVSLVKRKTTLAPNTQTASPR IADSI MOI AROVSRI FSGQ* 360 amino acids = 38.6 kDa	764 .. 1846	1083			CDS
✓ rop /gene = rop /product = Rop protein /translation = VTKQEK TALNMARFIRSQTLTLLEKLNELDADEQADICESLHDHADELYRSC LARFGDDGENL* 63 amino acids = 7.2 kDa	2655 .. 2846	192			CDS
✓ ori /direction = LEFT /note = high-copy-number ColE1/pMB1/pBR322/pUC origin of replication	3276 .. 3864	589			rep_origin

Feature	Location	Size (bp)			Type
✓ AmpR	4035 .. 4895	861			CDS
▶ 2 segments					
/gene	= bla				
/product	= -lactamase				
/note	= confers resistance to ampicillin, carbenicillin, and related antibiotics				
/translation	= MSIQHFRVALIPFFAAFCLPVFA,HPETLVKVKDAEDQLGARVGYIELDLNSGKILESFRPEERFPM MSTFKVLLCGAVLSRIDAGQE QLGRRRIHYSQNDLVEYSPVTEKHLTDGMTVRELC SAAITMSDNTAANLLTTIIGGPKELT AFLHNMGDHSVTRLDRWEP ELNEAIPN DERDTTMPAAMATTLRKLTTGELLTLASRQQLIDWMEADK VAGPLLR SALPAGWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIY TTGSOATMDFERNROI AFIGAS I KHW* 286 amino acids = 31.5 kDa				
✓ AmpR promoter	4896 .. 5000	105			promoter
/gene	= bla				
✓ f1 ori	5027 .. 5482	456			rep_origin
/direction	= LEFT				
/note	= f1 bacteriophage origin of replication; arrow indicates direction of (+) strand synthesis				

Description: Bacterial vector that encodes a signal sequence for inducible expression of proteins in the periplasm.

Created: Saturday, May 12, 2012

Last Modified: Sunday, Dec 1, 2013

Accession Number:

Code Number:

Sequence Author: Novagen (EMD Millipore)

DNA Type: Synthetic DNA

Laboratory Host Organism: Escherichia coli

Bacterial Transformation Strain: Unspecified

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

Comments:

References: