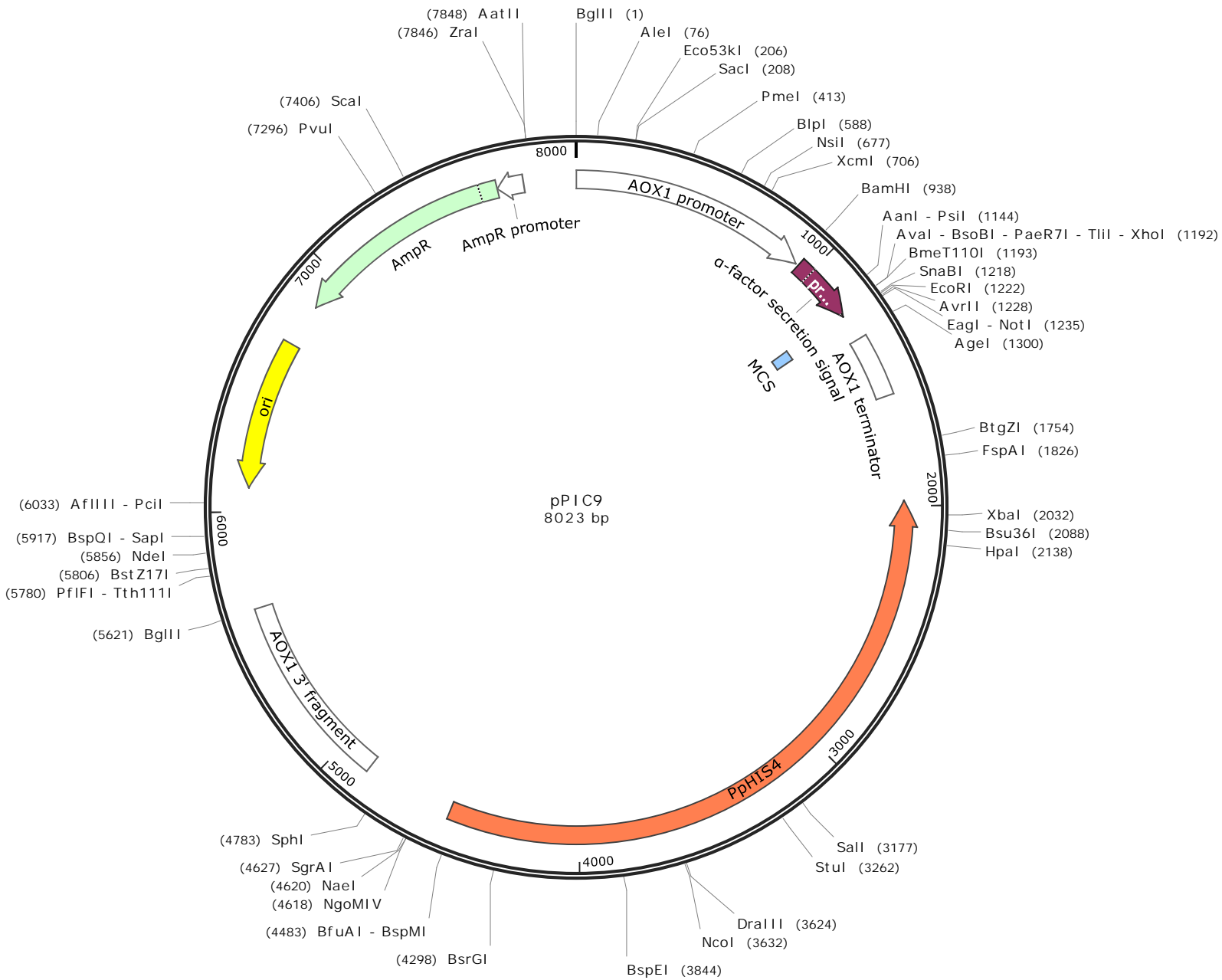


Pichia pastoris HIS4 vector for methanol-inducible expression of a secreted protein.



5' BglI A1eI

...
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AOX1 promoter

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AOX1 promoter

Eco53kI

Sacl

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AOX1 promoter

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AOX1 promoter

PmeI

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AOX1 promoter

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AOX1 promoter

B1pI

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AOX1 promoter

NsiI

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AOX1 promoter

XcmI

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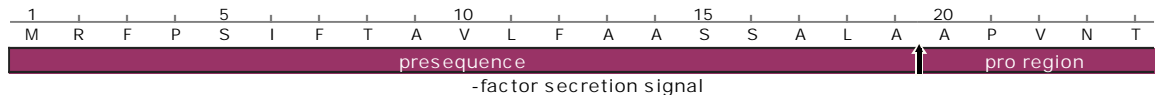
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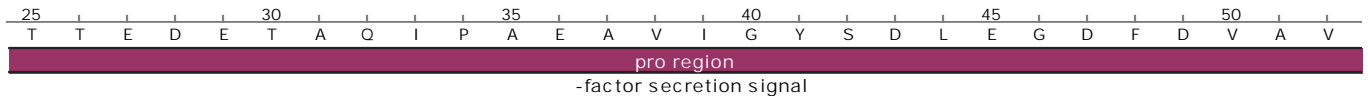
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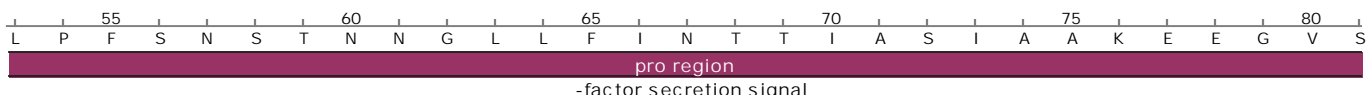
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TiII
XhoI
AvaI
BsoBI
PaeR7I BmeT110I
SnaBI EcoRI AvrII NotI
EagI

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Agel
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AOX1 terminator

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AOX1 terminator

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AOX1 terminator

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1615

AOX1 terminator

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1700

BtgZl

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1785

FspA1

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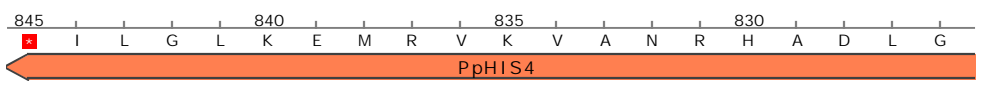
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1955

XbaI

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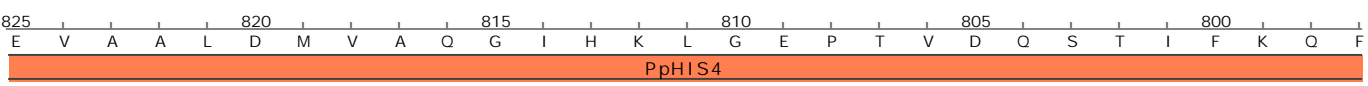
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Bsu36I

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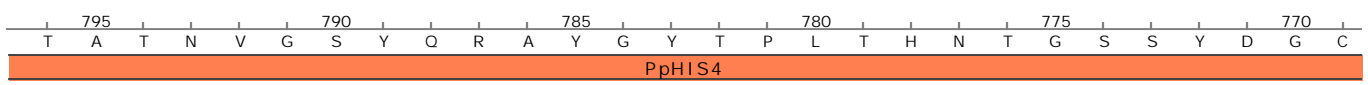
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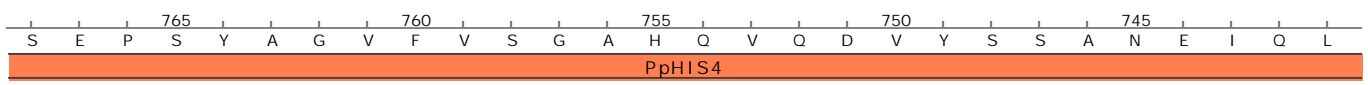
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2210



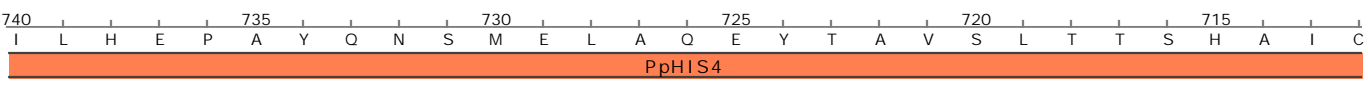
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2295



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2380



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< PpHIS4

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< PpHIS4

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3145

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

3230

PpHIS4

StuI

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

3315

PpHIS4

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3400

PpHIS4

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D L N R E V E D L T V G Y K A C R V L A F Y F L D A C E

3485

PpHIS4

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W A I D E K S K A E A L E D A E E R I K A N L L V E D D

3570

PpHIS4

DraIII

NcoI

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F L R K T Y S G E P A N S K R D W L T A E M A R L G K S Q

3655

PpHIS4

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G F C S T R E L H C F G V G T Q E V V F K L C D G D C D

3740

PpHIS4

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3825

PpHIS4

BspEI

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3910

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PpHIS4

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3995

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PpHIS4

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4080

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PpHIS4

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4165

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4250

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

PpHIS4

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4335

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4420

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PpHIS4

BfuAI

BspMI

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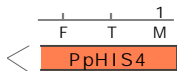
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4590



NgoMIV NaeI SgrAI

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4675

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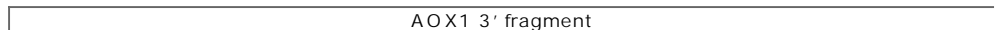
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4845

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4930



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5015

AOX1 3' fragment

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5100

AOX1 3' fragment

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5185

AOX1 3' fragment

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5270

AOX1 3' fragment

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5355

AOX1 3' fragment

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5440

AOX1 3' fragment

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5525

AOX1 3' fragment

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5610

AOX1 3' fragment

BglII

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5695

AOX1 3' fragment

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5780

PfIF1
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5865

SapI
BspQI

TGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGGCGCTCTTCCGCTTCTCGCTCACTGACTCGCTGCGCTCGGTC
ACTTTATGGCGTGTCTACGCATTCTCTTTTATGGCGTAGTCCCGCGAGAAGGCCAAGGAGCGAGTACTGAGCGACGCGAGCCAG

5950

PciI
AflII

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6035

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ACACTCGTTTTCCGGTCTTTTTCCGGTCTTTGGCATTTTTCCGGCGCAACGACCGCAAAAAGGTATCCGAGGCGGGGGGACTGCT

6120

ori

GCATCACAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTTCCCCTGGAAGCTCC
CGTAGTGTTTTTAGCTGCGAGTTCAGTCTCCACCGCTTTGGGCTGTCTGATATTTCTATGGTCCGCAAAGGGGGACCTTCGAGG

6205

ori

CTCGTGCCTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCAAT
GAGCACGCGAGAGGACAAGGCTGGGACGGCGAATGGCCTATGGACAGGCGGAAAGAGGGAAGCCCTTCGCACCGCGAAAGAGTTA

6290

ori

GCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCGTTTCAGCCCGACCG
CGAGTGCACATCCATAGAGTCAAGCCACATCCAGCAAGCGAGGTTTCGACCCGACACACGTGCTTGGGGGGCAAGTCGGGCTGGC

6375

ori

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GACGCGGAATAGGCCATTGATAGCAGAACTCAGGTTGGGCCATTCTGTGCTGAATAGCGGTGACCGTCTGCTGGTGACCATTTGTCC

6460

ori

ATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTG
TAATCGTCTCGCTCCATACATCCGCCACGATGTCTCAAGAACTTCACCACCGGATTGATGCCGATGTGATCTTCTGTCATAAAC

6545

ori

GTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGG
CATAGACGCGAGACGACTTCGGTCAATGGAAGCCTTTTTCTCAACCATCGAGAACTAGGCCGTTTGTGGTGGCGACCATCGCC

6630

ori

TGGTTTTTTTTGTTTGAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGAC
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6715

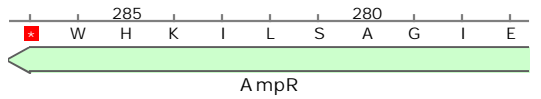
ori

GCTCAGTGGAACGAAAACCTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAGGATCTTCACCTAGATCCTTTTAAATTA
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6800

AATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTC
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6885



AGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGC
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6970

AmpR

CCCAGTGTGCAATGATACCGCGAGACCCACGCTCACCGGCTCCAGATTTATCAGCAATAAACCAGCCAGCCGGAAGGGCCGAGC
GGGTACAGACGTTACTATGGCGCTCTGGGTGCGAGTGGCCGAGGTCTAAATAGTCGTTATTTGGTCCGGTCCGGCCTTCCCGGCTCG

7055

AmpR

GCAGAAGTGGTCTCTGCAACTTTATCCGCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAA
CGTCTTCACCAGGACGTTGAAATAGGCGGAGGTAGGTCAGATAATTAACAACGGCCCTTCGATCTCATTTCATCAAGCGGTCAATT

7140

AmpR

TAGTTTGCGCAACGTTGTTGCCATTGCTGCAGGCATCGTGGTGTACGCTCGTTCGTTTGGTATGGCTTCATTCAGCTCCGGTTCC
ATCAAACGCGTTGCAACAACGGTAACGACGTCCTAGCACCACAGTGCAGAGCAGCAAACCATAACCGAAGTAAGTCGAGGCCAAGG

7225

AmpR

CAACGATCAAGGCGAGTTACATGATCCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTTCGGTCTCCGATCGTTGTGTCAGAAGTA
 GTTGCTAGTTCGCTCAATGTAAGTACTAGGGGGTACAACACGTTTTTTTCGCCAATCGAGGAAGCCAGGAGGCTAGCAACAGTCTTTCAT
 W R D L R T V H D G M N H L F A T L E K P G G I T T L L L
 160 155 150 145 140 135
 AmpR

7310

AGTTGGCCGCAAGTGTATCACTCATGGTTATGGCAGCACTGCATAATTCTTACTGTCATGCCATCCGTAAGATGCTTTTTCTGT
 TCAACCGGCGTCAACAATAGTGAGTACCAATACCGTCTGTGACGTATTAAGAGAATGACAGTACGGTAGGCATTCTACGAAAAGACA
 N A A T N D S M T I A A S C L E R V T M G D T L H K E T
 130 125 120 115 110
 AmpR

7395

GACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACCGAGTTGCTCTTGCCCGGCGTCAACACGGGATAAT
 CTGACCACTCATGAGTTGGTTCAGTAAGACTCTTATCACATACGCCGCTGGCTCAACGAGAACGGGGCCGAGTTGTGCCCTATTA
 V P S Y E V L D N Q S Y H I R R G L O E Q G A D V R S L
 105 100 95 90 85 80
 AmpR

7480

ACCGCGCCACATAGCAGAACTTTAAAAGTGCTCATCATTGGAAAACGTTCTTCGGGGCGAAAACCTCTCAAGGATCTTACCGCTGT
 TGGCGCGGTGTATCGTCTTGAAATTTTACGAGTAGTAACCTTTTGAAGAAGCCCCGCTTTTGAAGATTCTAGAAATGGCGACA
 V A G C L L V K F T S M M P F R E E P R F S E L I K G S N
 75 70 65 60 55 50
 AmpR

7565

TGAGATCCAGTTTCGATGTAACCCACTCGTGCACCCAACTGATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGGGTGAGCAAA
 ACTCTAGGTCAAGCTACATTGGGTGAGCAGTGGGTTGACTAGAAGTCGTAGAAAATGAAAGTGGTCGCAAAGACCCACTCGTTT
 L D L E I Y G V R A G L Q D E A D K V K V L T E P H A F
 45 40 35 30 25
 AmpR signal...

7650

AACAGGAAGGCAAAATGCCGCAAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTTCAATATTAT
 TTGTCCTTCCGTTTTTACGGCGTTTTTTCCCTTATTCGCTGTGCCTTTACAACCTATGAGTATGAGAAGGAAAAAGTTATAATA
 V P L C F A A F F P I L A V R F H Q I S M
 20 15 10 5 1
 signal sequence AmpR AmpR promoter



















7735




TGAAGCATTATCAGGGTATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCA
 ACTTCGTAATAGTCCCAATAACAGAGTACTCGCCTATGTATAAACTTACATAAATCTTTTTATTTGTTTATCCCCAAGGCGCGT
 AmpR promoter

7820

CATTTCCCCGAAAAGTGCCACCTGACGCTAAGAAACCATTATTATCATGACATTAACCTATAAAAAATAGGCGTATCACGAGGCC
 GTAAAGGGGCTTTTACGGTGGACTGCAGATTCTTTGGTAATAATAGTACTGTAATTGGATATTTTTATCCGCATAGTGCTCCGG
 ZraI AatII
 CTTTCGTCTTCAAGAATTAATTCTCATGTTTGACAGCTTATCATCGATAAGCTGACTCATGTTGGTATTGTGAAATAGACGCAGA
 GAAAGCAGAAGTCTTAATTAAGAGTACAACTGTGCAATAGTAGCTATTCGACTGAGTACAACCATAACACTTTATCTGCGTCT
 7905 7990

Enzymes	Sites	
AanI	1	1144
AatII	1	7848
AflIII	1	6033
AgeI	1	1300
AleI	1	76
AvaI	1	1192
AvrII	1	1228
BamHI	1	938
BfuAI	1	4483
BglII	2	1 5621
BipI	1	588
BmeT110I	1	1193
BsoBI	1	1192
BspEI	1	3844
BspMI	1	4483
BspQI	1	5917
BsrGI	1	4298
BstZ17I	1	5806
Bsu36I	1	2088
BtgZI	1	1754
DraIII	1	3624
EagI	1	1235
Eco53kI	1	206
EcoRI	1	1222
FspAI	1	1826
HpaI	1	2138
NaeI	1	4620
NcoI	1	3632
NdeI	1	5856
NgoMIV	1	4618
NotI	1	1235
NsiI	1	677
PaeR7I	1	1192
PciI	1	6033
PfIF1	1	5780
PmeI	1	413
PsiI	1	1144
PvuI	1	7296
SacI	1	208
Sall	1	3177
SapI	1	5917
ScaI	1	7406
SgrAI	1	4627
SnaBI	1	1218
SphI	1	4783
StuI	1	3262
TliI	1	1192
Tth111I	1	5780
XbaI	1	2032
XcmI	1	706
XhoI	1	1192
ZraI	1	7846

Feature	Location	Size (bp)			Type
✓ AOX1 promoter	2 .. 937	936			promoter
/gene	= Pichia pastoris AOX1				
/note	= inducible promoter, regulated by methanol				
✓ -factor secretion signal	949 .. 1215	267			CDS
▶ 3 segments					
/gene	= MF 1				
/product	= N-terminal secretion signal from S. cerevisiae alpha-factor				
/note	= Cleavage by the Kex2 protease occurs after the dibasic KR sequence. The EA dipeptides are then removed by dipeptidyl aminopeptidase A.				
/translation	= MRFPSIFTAVLFAASSALA,APVNTTTEDETAQIPAEAVIGYSDLEGGDFDVAVLFPNSTNNGLLFINTTIIASIAAKEEGVSLEKR,EA FA 89 amino acids = 9.3 kDa				
✓ MCS	1192 .. 1241	50			misc_feature
/note	= multiple cloning site				
✓ AOX1 terminator	1321 .. 1567	247			terminator
/gene	= Pichia pastoris AOX1				
/note	= transcription terminator for AOX1				
✓ PpHIS4	1980 .. 4514	2535			CDS
/gene	= Pichia pastoris HIS4				
/product	= multifunctional enzyme, required for histidine biosynthesis				
/note	= auxotrophic marker for Pichia pastoris				
/translation	= MTFPLLPAYASVAEFDNSLSLVGKAVFPYAADQLHNLIKFTQSTELQVNVQVESSVTEDQFEELIDNLLKLYNNGINEVILDLDLAE RVVQRMIPGARVIYRTLVDKVASLPANASIAVPFSSPLGDLKSFNTGGSRVYAFSETAKLVDVTSTVASGIIPIIDARQLTTEYELS EDVKKFPVSEILLASLTDRPDGLFTTLVADSSNYSLGLVYSSKKSIPAEAIRTQTGVYQSRRHGLWYKGGATSGATQKLLGIELDCDGD CLKFVVEQTVGVGFCHELERTSCFGQSKGLRAMEATLWDRKSNAPESYTKRFLDDEVLLNAKIREEADELAEAKSKEDIAWECADLF YFALVRCAYGVTLDEVERNLDMKSLKVTTRKGDAPGYTKEQPKESKPKVEPSEGRIELCKIDVSKASSQEI EDALRRPIQKTEQ IMELVKPIVDNVRONGDKALLELTAKFDGVALKTPVLEAPFPEELMQLPDNVKRAIDLSIDNVRKFHEAQLTETLQVETCPGVVCSR FARPIEKVGLYIPGGTALPSTSLMLGVPKAVAGCKEIVFASPPKDGTLTPEVIYVAHKVGAKEIVLAGGAQAVAAAMAYGTETVP KCDKIFGPGNQFVTAAKMMVQNDTSALCSIDMPAGPSEVLVIADKYADPDFVASDLLSQAHEHGIDSQVILLAVDMDTKELARIED AVHNQAVQLPRVEIVRKCIAHSTTSLVATYEQALEMSNOYAPEHLILQIENASSYVDQVQHAGSVFVGAYSPESC GDYSSGTH TIPTYGYAROYSGVNTATFQKFITSDVTPFGIKHIGAVMDIAAVFGIDAHRNAVKVRMFKIGI I * 844 amino acids = 92.2 kDa				
✓ AOX1 3' fragment	4869 .. 5625	757			misc_feature
/note	= region downstream of Pichia pastoris AOX1 gene				
✓ ori	6094 .. 6682	589			rep_origin
/direction	= LEFT				
/note	= high-copy-number ColE1/pMB1/pBR322/pUC origin of replication				
✓ AmpR	6853 .. 7713	861			CDS
▶ 2 segments					
/gene	= bla				
/product	= -lactamase				
/note	= confers resistance to ampicillin, carbenicillin, and related antibiotics				
/translation	= MSIQHFRVALIPFFAAFLPVFA,HPETLVKVKDAEDQLGARVGYIELDLNSGKILESFRPEERFPMSTFKVLLCGAVLSRVDAEQE QLGRRIHYSQNDLVEYSPVTEKHLTDGMTVRELCSAAITMSDNTAANLLLTITIGGPKELTAFHNMGDHVRTLDLDRWEPELNEAIPN DERDITMPAAMATTLRKLTLGELLTLASRQLLDWMEADKVAAGPLLRSALPAGWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIY TTGSOATMDFRNROI AFIGASIKHW* 286 amino acids = 31.5 kDa				

Feature	Location	Size (bp)			Type
✓ AmpR promoter	7714 .. 7818	105	<input type="checkbox"/>		promoter
/gene = bla					

Description: Pichia pastoris HIS4 vector for methanol-inducible expression of a secreted protein.

Created: Tuesday, Feb 5, 2013

Last Modified: Tuesday, Feb 5, 2013

Accession Number:

Code Number:

Sequence Author: Invitrogen (Life Technologies)

DNA Type: Synthetic DNA

Laboratory Host Organism: Pichia pastoris

Bacterial Transformation Strain: Unspecified
Dam⁺ Dcm⁺ EcoKI⁺

Comments:

References: