

pDRIVE-hGAPDH

A plasmid with the native human GlycerAldehyde 3-Phosphate DeHydrogenase promoter

Catalog # pdrive-hgapdh

For research use only

Version # 01E02-MT

PRODUCT INFORMATION

Content:

- 1 disk of lyophilized GT100 *E. coli* bacteria transformed by a pDRIVE plasmid.
- GT100 genotype is: *F*-, *mcrA*, Δ (*mrr-hsdRMS-mcrBC*), \emptyset 80*lacZ* Δ M15, *ΔlacX74*, *recA1*, *endA1*.
- 4 pouches of *E. coli* FastMedia™ Zeo

Shipping and storage:

- Products are shipped at room temperature.
- Transformed bacteria should be stored at -20°C. Bacteria are stable up to one year when properly stored.
- Store *E. coli* FastMedia™ Zeo at room temperature. FastMedia™ is stable 18 months when stored properly.

Quality control:

- Plasmid construct has been confirmed by restriction analysis and sequencing.
- Bacteria have been lyophilized, and their viability upon resuspension has been verified.
- Promoter activity has been confirmed by transient transfection of 293 cells as well as other selected cell lines.

GENERAL PRODUCT USE

pDRIVE is an expression plasmid containing a native or composite promoter of interest. pDRIVE may be used to:

- **Subclone a promoter of interest into another vector.** Unique restriction sites are present at each end of the promoter allowing convenient excision. The 5' sites include *Sda* I, *Pst* I, and *Spe* I. *Sda* I is compatible with *Nsi* I and *Pst* I. *Spe* I is compatible with *Avr* II, *Nhe* I and *Xba* I. The 3' restriction site is *Nco* I which includes the ATG start codon, and is compatible with *Bsp*H I and *Bsp*LU11 I.
- **Compare the activity of different promoters** in transient transfection experiments. Each pDRIVE promoter drives the expression of the *LacZ* reporter gene which allows for testing of the promoter's activity in transient transfection experiments. Furthermore, the *LacZ* gene is flanked by unique restriction sites (*Nco* I and *Eco*R I) for easy replacement with a different gene of interest.

PROMOTER CHARACTERISTICS

Element	Name	Origin	Size bp
Promoter	GAPDH	Human	482
5'UTR	GAPDH	Human	313
Enhancer	-	-	-

GAPDH promoter

Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) is a catalytic enzyme involved in glycolysis. The GAPDH gene is constitutively expressed at high levels in almost all tissues. However, the molecular mechanism which sustains high-level expression of this house-keeping enzyme is still unclear. The expression of GAPDH can be induced by several physiological factors in certain cell types. In adipocytes, insulin increases the levels of GAPDH mRNA through cis-acting sequences located within the GAPDH promoter¹. In endothelial cells, GAPDH is induced by hypoxia at the transcriptional level, this upregulation occurring to a much greater extent in EC than in other cell types².

PLASMID FEATURES

- **LacZ gene** encodes β -galactosidase an enzyme that catalyzes the hydrolysis of X-Gal, producing a blue precipitate that can be easily visualized under a microscope.
 - **SV40 pAn:** The Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA.
 - **Ori pMB1** is a minimal *E. coli* origin of replication with the same activity as the longer Ori.
 - **EM7** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.
 - **Sh ble** gene confers zeocin resistance therefore allowing the selection of transformed *E. coli* carrying a pDRIVE plasmid.
- Note: Stable transfection of clones cannot be performed due to the absence of an eukaryotic promoter upstream of the Sh ble gene.*

METHODS

Growth of pDRIVE-transformed bacteria:

Use sterile conditions to do the following:

- 1- Resuspend the lyophilized *E. coli* by adding 1 ml of LB medium in the tube containing the disk. Let sit for 5 minutes. Mix gently by inverting the tube several times.
- 2- Streak bacteria taken from this suspension on a zeocin LB agar plate prepared with the *E. coli* FastMedia™ Zeo agar provided (see below).
- 3- Place the plate in an incubator at 37°C overnight.
- 4- Isolate a single colony and grow the bacteria in TB supplemented with zeocin using the FastMedia™ Zeo liquid provided (see below).
- 5- Extract the pDRIVE plasmid DNA using the method of your choice.

Selection of bacteria with *E. coli* FastMedia™ Zeo:

E. coli FastMedia™ Zeo is a **new, fast and convenient** way to prepare liquid and solid media for bacterial culture by using only a microwave. *E. coli* FastMedia™ Zeo is a TB (liquid) or LB (solid) based medium with zeocin, and contains stabilizers.

E. coli FastMedia™ Zeo can be ordered separately (catalog code # fas-zn-l, fas-zn-s).

Method:

- 1- Pour the contents of a pouch into a clean borosilicate glass bottle or flask.
- 2- Add 200 ml of distilled water to the flask
- 3- Heat in a microwave on MEDIUM power setting (about 400Watts), until bubbles start appearing (approximately 3 minutes). **Do not heat a closed container. Do not autoclave FastMedia™.**
- 4- Swirl gently to mix the preparation. **Be careful, the bottle and media are hot, use heatproof pads orgloves and care when handling.**
- 5- Reheat the media for 30 seconds and gently swirl again. Repeat as necessary to completely dissolve the powder into solution. But be careful to avoid overboiling and volume loss.
- 6- Let agar medium cool to 45°C before pouring plates. Let liquid media cool to 37°C before seeding bacteria.

Note: Do not reheat solidified FastMedia™ as the antibiotic will be permanently destroyed by the procedure.

References:

- 1- Ercolani Let al. 1988. J Biol Chem. 263(30):15335-41
- 2- Graven KK et al. 1999. Biochim Biophys Acta. 1447(2-3):208-18

TECHNICAL SUPPORT

Toll free (US): 888-457-5873

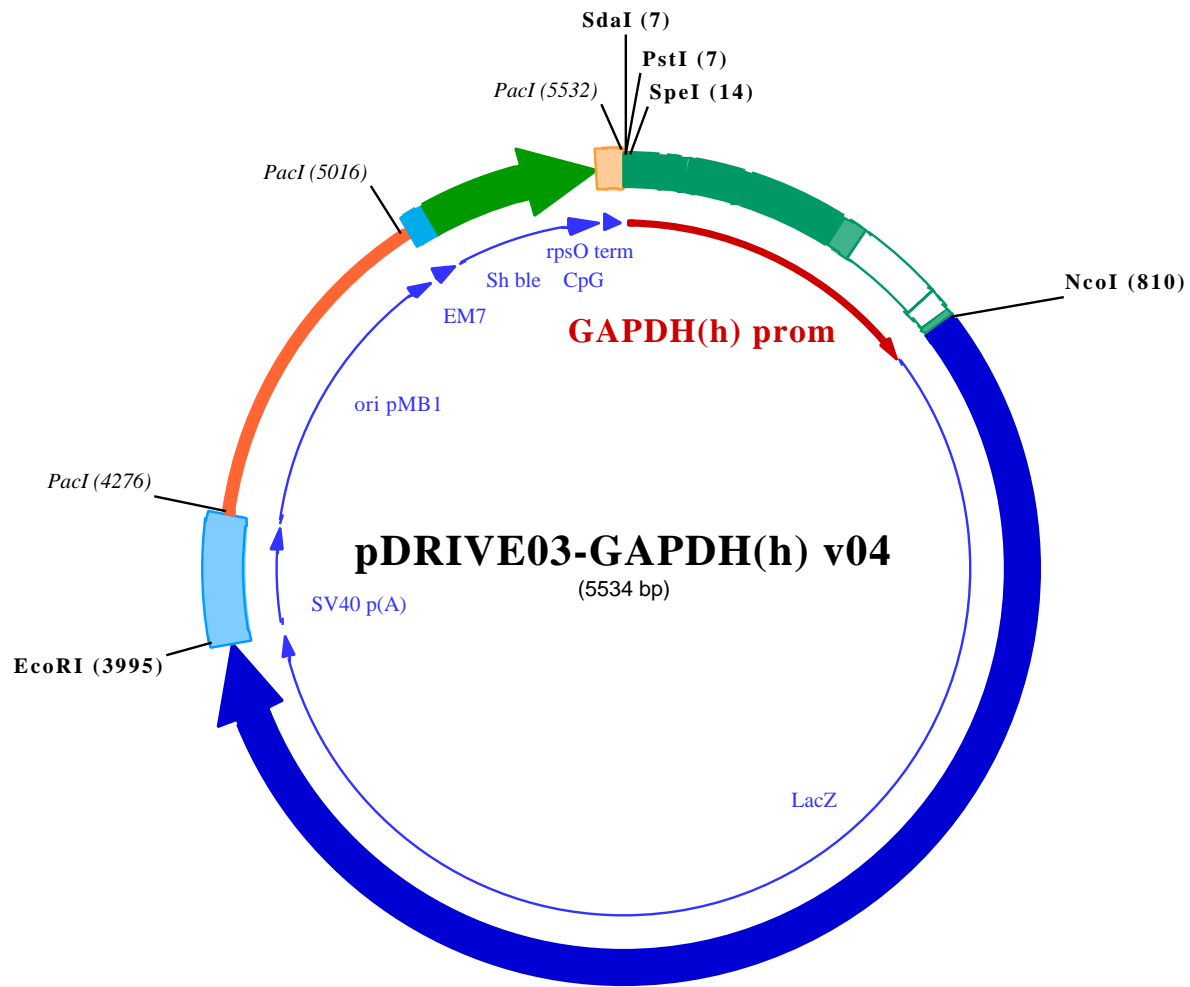
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PstI (7)
SdaI (7) SpeI (14)

1 CCTGCAGGGCCCACTAGTTCCCAACTTTCCCGCCTCTCAGCCTTTGAAAGAAAGAAAGGGGAGGGGCGAGCCGCTGCAGTCGCGAGCGGTGCTGGGC
101 TCCGGCTCCAATTCCCCATCTCAGTCGCTCCCAAAGTCCTTCTGTTTCATCCAAGCGTGAAGGGTCCCCGTCCTTGACTCCCTAGTGTCTGCTGCCA
201 CAGTCCAGTCTCGGGAACCAGCACCGATCACCTCCCATCGGGCCAATCTCAGTCCCTTCCCCCTACGTCGGGGCCACACCGCTCGGTGCGTCCCAGTT
301 GAACCAGGGCGGTGCGGAAAAAAGCGGGGAGAAAGTAGGGCCCGGCTACTAGCGGTTTTACGGGCGCACGTAGCTCAGGCCTCAAGACCTTGGGCT
401 GGGACTGGCTGAGCCTGGCGGGAGCGGGGTCCGAGTACCAGCCTGCCGCGCCCGGTTTCTATAAATGAGCCCGCAGCCTCCCGCTCGCTCTC
501 TGCTCCTCTGTTGACAGTCAGCCGATCTTCTTTTGGCTGCCAGgtgaagacggggcgagagaaacccgggaggctagggaacggcctgaagcgggca
601 ggggcgggcgagcggcgatgtgttcgcgccgctgccccggggcgccctccgcatgacggggcgggcgaggacgtgatcgggcgggcgctg
701 ggcatggaggcctggtgggggaggggaggggagggcgtgggtgtcgccggggccactaggcgctcactgttctctccctccgagCGAGCCACATCG

NcoI (810)

801 CTGAGACACCATGGGGTTCATCATCATCATCATCATGGTATGGTAGCATGACTGGTGGACGCAAAATGGGTCGGGATCTGTACGACGATGACGAT
MetGlyGlySerHisHisHisHisHisHisGlyMetAlaSerMetThrGlyGlyGlnGlnMetGlyArgAspLeuTyrAspAspAspAsp
901 AAGGTACCTAAGGATCAGCTTGGAGTTGATCCCGTCGTTTTACAACGTCGTGACTGGGAAAACCTGGCGTTACCCAACCTAATCGCCTTCAGCACATC
31 LysValProLysAspGlnLeuGlyValAspProValValLeuGlnArgArgAspTrpGluAsnProGlyValThrGlnLeuAsnArgLeuAlaAlaHisP
1001 CCCCTTCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCGCCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAATGGCGCTTTGCCTGGTTTC
64 roProPheAlaSerTrpArgAsnSerGluGluAlaArgThrAspArgProSerGlnGlnLeuArgSerLeuAsnGlyGluTrpArgPheAlaTrpPhePr
1101 GGACCAGAAGCGGTGCGGAAAGCTGGCTGGAGTGGCATCTTCTGAGGCGGATACGTCGCTCGTCCCTCAAACCTGGCAGATGCACGGTTACGATCGG
97 oAlaProGluAlaValProGluSerTrpLeuGluCysAspLeuProGluAlaAspThrValValValProSerAsnTrpGlnMetHisGlyTyrAspAla
1201 CCGATCTACACCAACGTAACCTATCCATTACGGTCAATCCGCGGTTTGTCCACGGAGAATCCGACGGGTTGTTACTCGCTCACATTTAATGTTGATG
131 ProlIeTyrThrAsnValThrTyrProlIeThrValAsnProProPheValProThrGluAsnProThrGlyCysTyrSerLeuThrPheAsnValAspG
1301 AAAGCTGGTACAGGAAGGCCAGACCGCAATATTTTTGATGGCGTTAACTCGGCGTTTCATCTGTGGTGAACGGGCGCTGGGTCGGTACGGCCAGGA
164 luSerTrpLeuGlnGluGlyGlnThrArgIleIePheAspGlyValAsnSerAlaPheHisLeuTrpCysAsnGlyArgTrpValGlyTyrGlyGlnAs
1401 CAGTCGTTTGGCGTCTGAATTTGACCTGAGCGCATTTTTACGCGCGGAGAAAACCGCCTCGCGGTGATGGTGTGCGTGGAGTGACGGCAGTTATCTG
197 pSerArgLeuProSerGluPheAspLeuSerAlaPheLeuArgAlaGlyGluAsnArgLeuAlaValMetValLeuArgTrpSerAspGlySerTyrLeu
1501 GAAGATCAGGATATGGGCGGATGAGCGGCATTTTCCGTCAGCTCTCGTTGCTGCATAAACCGACTACACAAATCAGCGATTTCCATGTTGCCACTCGCT
231 GlyAspGlnAspMetTrpArgMetSerGlyIlePheArgAspValSerLeuGlyGlnLeuAsnGlyGlnLeuSerThrThrGlnIleSerLeuAsnHisPro
1601 TTAATGATGATTTACGCGCGCTGACTGAGGCTGAAGTTACAGATGTGCGCGAGTTGCGTACTACTACGGGTAACAGTTTCTTTATGGCAGGGTGA
264 heAsnAspAspPheSerArgAlaValLeuGluAlaGluValGlnMetCysGlyGluLeuArgAspTyrLeuArgValThrValSerLeuTrpGlnGlyG
1701 AACGCAGTTCGCCAGCGGCACCGCGCCTTTCGCGGTGAAATATTCGATGAGCGGTGGTGGTTATGCCGATCGCGTCACACTAGCTCTGAACGTCGAAAC
297 uThrGlnValAlaSerGlyThrAlaProPheGlyGlyGluIleIeAspGluArgGlyGlyTyrAlaAspArgValThrLeuArgLeuAsnValGluAsn
1801 CCGAACTGTGGAGCGCGAAATCCGAATCTCTATCGTGGCGGTGAGTTGAAGTCACACCGCGCAGCGCTGATTGAAGCAGAAGCTGGCAGTCTGG
331 ProLysLeuTrpSerAlaGluIleProAsnLeuTyrArgAlaValGluLeuHisThrAlaAspGlyThrLeuIleGluAlaGluAlaCysAspValG
1901 GTTTCGCGAGGTGGGATTGAAAATGGTCTGCTGCTGCTGAACGGCAAGCCGTTGCTGATTCGAGGCGTTAACCGTCACGAGCATCATCTCTGCATGG
364 lyPheArgGluValArgIleGluAsnGlyLeuLeuLeuLeuAsnGlyLysProLeuLeuIleArgGlyValAsnArgHisGluHisHisProLeuHisG
2001 TCAGTTCATGGATGAGCAGACGATGGTGCAGGATATCCTGCTGATGAAGCAGAACTTTAACCGCGTGGCGTGTTCGCATTATCCGAACCATCCGCTG
397 yGlnValMetAspGlnGlnThrMetValGlnAspIleLeuLeuMetLysGlnLeuAsnHisPheAsnAlaValArgCysSerHisTyrProAsnHisPro
2101 TGTACAGCTGTGCGACGCTACCGCCTGTATGTGGTGAAGCCAATATTGAAACCCAGCGCATGGTGCATGCAATCGTCTGACCGATGATCCGC
431 TrpTyrThrLeuCysAspArgTyrGlyLeuTyrValValAspGluAlaAsnIleGluThrHisGlyMetValProMetAsnArgLeuThrAspAspProA
2201 GCTGGCTACCGCGATGAGCGAACCGCTAACCGAATGGTGCAGCGCATCGTAATCACCCGAGTGTGATCATCTGGTGGTGGGGAATGAATCAGGCCA
464 rgTrpLeuProAlaMetSerGluArgValThrArgMetValGlnArgAspArgAsnHisProSerValIleIleTrpSerLeuGlyAsnGluSerGlyHi
2301 CGGCGTAATCACGACCGCTGTATCGCTGGATCAAATCTGTCGATCCTTCCCGCCGGTGCAGTATGAAGCGCGGAGCCGACACCCAGCCACCGAT
497 sGlyAlaAsnHisAspAlaLeuTyrArgTrpIleLysSerAlAspProSerArgProValGlnTyrGluGlyGlyGlyAlaAspThrAlaThrAsp
2401 ATTATTTGCCGATGTACGCGCGTGGATGAAGACCAGCCCTTCCCGGCTGTGCCGAAATGGTCCATCAAAAAATGGCTTTCGCTACCTGGAGAGACGC
531 IleIleCysProMetTyrAlaArgValAspGluAspGlnProPheProAlaValProLysTrpSerIleLysLysTrpLeuSerLeuProGlyGluThrA
2501 GCCCGCTGATCTTTGCGAATACGCCACCGGATGGGTAACAGTCTTGGCGGTTTCGCTAAATACTGGCAGGCGTTTCGCTAGTATCCCGCTTACAGGG
564 rgProLeuIleLeuCysGluThrAlaHisAlaMetGlyAsnAlaMetGlyAsnAlaPheAlaLysTyrTrpGlnAlaPheArgGlnTyrProArgLeuGlnG
2601 CGGCTTCTGGGACTGGGTGGATCAGTCGCTGATTAATATGATGAAACGGCAACCCGTTGGTGGCTTACGGCGGTGATTTGGCGATACGCCGAAC
597 yGlyPheValTrpAspTrpValAspGlnSerLeuIleLysTyrAspGluAsnGlyAsnProTrpSerAlaTyrGlyGlyAspPheGlyAspThrProAsn
2701 GATCGCCAGTTCGTATGAACGGTCTGGTCTTTGCCGACCGCACCGCATCCAGCGCTGACGGAAGCAAAACACCAGCAGAGTTTTTCCAGTTCGGTT
631 AspArgGlnPheCysMetAsnGlyLeuValPheAlaAspArgThrProHisProAlaLeuThrGluAlaLysHisGlnGlnPhePheGlnPheArgL
2801 TATCCGGGAAACCATCGAAGTGACCGCAATACCTGTTCCGTCATAGCGATAACGAGTCCCTGCAGTGGTGGCGCTGGATGGTAAGCCGCTGGC
664 euSerGlyGlnThrIleGluValThrSerGluThrLeuPheArgHisSerAspAsnGluLeuLeuLeuLeuLeuLeuLeuLeuLeuLeuLeuLeuLeuLeu
2901 AAGCGTGAAGTGCCTCTGGATGTCGCTCCACAAGGTAACAGTTGATTGAAGTGCCTGAAGTACCGCAGCCGAGAGCGCCGGCAACTCTGGCTACA
697 aSerGlyGluValProLeuAspValAlaProGlnGlyLysGlnLeuIleGluLeuProGluLeuProGlnProGluSerAlaGlyGlnLeuTrpLeuThr
3001 GTACCGTAGTGAACCGAACCAGCGACCGCATGGTTCAGAACCGGGCACATCAGCGCCTGGCAGCAGTGGCGTCTGGCGGAAAACCTCAGTGTGACGCTCC
731 ValArgValValGlnProAsnAlaThrAlaTrpAlaSerGluAlaGlyHisIleSerArgTrpGlnGlnTrpArgLeuAlaLeuAsnLeuSerValThrLeuP
3101 CCGCGGTCACCGCATCCCGCATCTGACCAACCGGAAATGGATTTTTGATCGAGCTGGGTAATAAGCGTTGGCAATTTAACCGCATCAGGCTT
764 roAlaAlaSerHisAlaIleProHisLeuThrThrSerGluMetAspPheCysIleGluLeuGlyAsnLysArgTrpGlnPheAsnArgGlnSerGlyPh
3201 TCTTTCACAGATGGGATGGCGATAAAAAACAACCTGCTGACCGCGTGGCGATCAGTTACCCCGTGCACCCTGGATAACGACATTTGGCGTAAGTGAA
797 eLeuSerGlnMetTrpIleGlyAspLysLysGlnLeuLeuGluThrProLeuArgAspGlnPheThrArgAlaProLeuAspAsnAspIleGlyValSerGlu
3301 CGGACCCGATGACCTAACCGCTGGTCCGAGCGTGAAGCGCGGGCCATTACGAGCCGAAGCAGCGTTGTTGACGTGCACGGCAGATACACTTG
831 AlaThrArgIleAspProAsnAlaTrpValGluArgTrpLysAlaAlaGlyHisTyrGlnAlaGluAlaAlaLeuLeuGlnCysThrAlaAspThrLeuA
3401 CTGATGGGTGCTGATTACGACCGCTACCGGTGGCAGCATCAGGGAAAACCTTATTTATCAGCCGAAAACCTACCGGATGATGGTACTGGTCAAAT
864 laAspAlaValLeuIleThrThrAlaHisAlaTrpGlnHisGlnGlyLysThrLeuPheIleSerArgLysThrTyrArgIleAspGlySerGlyGlnMe
3501 GCGGATTACCGTTGATTTGAAGTGGCAGCGATACCCGCATCCGCGCGGATTGGCTGAACTGCCAGCTGGCGCAGGTAGCAGAGCGGGTAAACTGG
897 tAlaIleThrValAspValGluValAlaSerAspThrProHisProAlaArgIleGlyLeuAsnCysGlnLeuAlaGlnValAlaGluArgValAsnTrp

3601 CTCGGATTAGGGCCGCAAGAAAATATCCCGACCGCCTTACTGCCGCCTGTTTTGACCGCTGGGATCTGCCATTGTGACACATGTATACCCCGTACGTCT
 931▶ LeuGlyLeuGlyProGlnGluAsnTyrProAspArgLeuThrAlaAlaCysPheAspArgTrpAspLeuProLeuSerAspMetTyrThrProTyrValP
 3701 TCCCGAGCGAAAACGGTCTGCGCTGCGGGACGCGGAATTGAATTATGGCCACACCAGTGGCGGGGACTTCCAGTTCAACATCAGCCGCTACAGTCA
 964▶ heProSerGluAsnGlyLeuArgCysGlyThrArgGluLeuAsnTyrGlyProHisGlnTrpArgGlyAspPheGlnPheAsnI leSerArgTyrSerGI
 3801 ACAGCAACTGATGGAACACGCCATCGCCATCTGCTGCACCGGAAGAAGGCACATGGCTGAATATCGACGGTTTTCCATATGGGGATTGGTGGCGACGAC
 997▶ nGlnGlnLeuMetGluThrSerHisArgHisLeuLeuHisAlaGluGluGlyThrTrpLeuAsnI leAspGlyPheHisMetGlyI leGlyGlyAspAsp
EcoRI (3995)

3901 TCCTGGAGCCCCTCAGTATCGGCGGAATTACAGCTGAGCGCCGGTGCCTACCATTACCAGTGGTCTGGTGTCAAAAATAATAATCTAGTCGAGAATTGG
 1031▶ SerTrpSerProSerValSerAlaGluLeuGlnLeuSerAlaGlyArgTyrHisTyrGlnLeuValTrpCysGlnLys•••

4001 CTAGCTCGACATGATAAGATACATTGATGAGTTTGGACAAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCT

 4101 TTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAAACAAGTTAAACAACAACAATTGCATTCATTTTATGTTTCAGGT

 4201 TCAGGGGGAGGTGTGGGAGGTTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTAGATCCATTTAAATGTTAATTAAGTCCATGACCAAAATCCCTT
PacI (4276)

 4301 AACGTGAGTTTTCTGTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTTTTCTGCGCGTAATCTGCTGCTTGC AAAAC

 4401 AAAAAAACACCGCTACCAGCGGTGGTTTGTGTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACCTGGCTTCAGCAGAGCGCAGATACCAAAAT

 4501 ACTGTTCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAAGAACTCTGTAGCACCGCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGCTG

 4601 CCAGTGGCGATAAGTCGTGCTTACCAGGTTGGACTCAAGACGATAGTTACCGGATAAGGCGCAGCGGTCCGGCTGAACGGGGGGTTCGTGCACACAGCC

 4701 CAGCTTGGAGCGAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAAGCCACGCTTCCCGAAGGGAGAAAAGGCGGACAGGTATCCG

 4801 GTAAGCGGCAGGTCGGAACAGGAGAGCGCACGAGGGAGCTTCCAGGGGAAACGCCTGGTATCTTTATAGTCTGTGCGGTTTCGCCACCTCTGACTTG

 4901 AGCGTCGATTTTTGTGATGCTCGTCAGGGGGCGGAGCCTATGGAAAAACGCCAGCAACCGGCCTTTTTACGGTTCCTGGCCTTTTGTGCGCCTTTTGC

PacI (5016)
 5001 TCACATGTTCTTAATTAATTTTTCAAAAGTAGTTGACAATTAATCATCGGCATAGTATATCGGCATAGTATAATACGACTCACTATAAGGAGGGCCATCA
1▶M

 5101 TGGCCAAGTTGACCAGTGTCTCCAGTGTCTCACAGCCAGGGATGTGGCTGGAGCTGTTGAGTTCTGGACTGACAGGTTGGGTTCTCCAGAGATTTTGT
 1▶ etAlaLysLeuThrSerAlaValProValLeuThrAlaArgAspValAlaGlyAlaValGluPheTrpThrAspArgLeuGlyPheSerArgAspPheVa
 5201 GGAGGATGACTTTGCAGGTGTGGTCAGAGATGATGTCACCCTGTTCTCAGCAGTCCAGGACCAGGTGGTGCCTGACAACACCCTGGCTTGGGTGTGG
 34▶ lGluAspAspPheAlaGlyValValArgAspAspValThrLeuPheI leSerAlaValGlnAspGlnValValProAspAsnThrLeuAlaTrpValTrp
 5301 GTGAGAGGACTGGATGAGCTGTATGCTGAGTGGAGTGAGGTGGTCTCCACCAACTTCAGGGATGCCAGTGGCCCTGCCATGACAGAGATTGGAGAGCAGC
 68▶ ValArgGlyLeuAspGluLeuTyrAlaGluTrpSerGluValValSerThrAsnPheArgAspAlaSerGlyProAlaMetThrGluI leGlyGluGlnP
 5401 CCTGGGGGAGAGAGTTTGCCTGAGAGACCCAGCAGGCAACTGTGTCACTTTGTGGCAGAGGAGCAGGACTGAGGATAAGAATTGAGTTTCAGAAAAGG
 101▶ roTrpGlyArgGluPheAlaLeuArgAspProAlaGlyAsnCysValHisPheValAlaGluGluGlnAsp•••

PacI (5532)
 5501 GGGCCTGAGTGGCCCTTTTTTCAACTTAATTAA