

# pDRIVE-hFerL/RU5'

A plasmid with a composite promoter comprised of the human Ferritin Light and HTLV 5' UTR

Catalog # pdrive-hferlu5

For research use only

Version # 04L15-SV

## PRODUCT INFORMATION

### Content:

- 1 disk of lyophilized GT100 *E. coli* bacteria transformed by pDRIVE-hFerL/RU5'.
- GT100 genotype is: *F-*, *mcrA*,  $\Delta$ (*mrr-hsdRMS-mcrBC*),  $\Phi$ 80*lacZ* $\Delta$ M15, *ΔlacX74*, *recA1*, *endA1*.
- 4 pouches of *E. coli* Fast-Media® 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* Fast-Media® Zeo at room temperature. Fast-Media® pouches are 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 *BspH* I and *BspLU11* 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 *EcoR* I) for easy replacement with a different gene of interest.

## PROMOTER CHARACTERISTICS

Element	Name	Origin	Size bp
Promoter	FerL	Human	537
5'UTR	HTLV	Viral	267
Intron	-	-	-

### hFerL/HTLV promoter

Ferritin is a ubiquitous iron storage protein. Ferritin is a 24 subunit protein composed of two subunit types termed H (heavy) and L (light) which perform complementary functions in the protein. The synthesis of ferritin is highly regulated by the iron status of the cell. The iron regulation is achieved at the translational level through interaction between a 28-nucleotide iron-responsive element (IRE) located in the 5' UTR of ferritin mRNAs and a cytosolic protein, the iron regulatory protein<sup>1</sup>. To eliminate the iron regulation of the ferritin promoter, the 5' UTR of FerL has been replaced by the 5' UTR of the HTLV. This modification makes the FerL promoter ubiquitous, strong and constitutive.

## 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.
  - **pMB1 Ori** 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* Fast-Media® 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 Fast-Media® Zeo liquid provided (see below).
- 5- Extract the pDRIVE plasmid DNA using the method of your choice.

### Selection of bacteria with *E. coli* Fast-Media Zeo:

*E. coli* Fast-Media® Zeo is a **new, fast and convenient** way to prepare liquid and solid media for bacterial culture by using only a microwave. *E. coli* Fast-Media® Zeo is a TB (liquid) or LB (solid) based medium with zeocin, and contains stabilizers.  
*E. coli* Fast-Media® 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 Fast-Media®.**
  - 4- Swirl gently to mix the preparation. **Be careful, the bottle and media are hot, use heatproof pads or gloves 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 Fast-Media® as the antibiotic will be permanently destroyed by the procedure.*

### References:

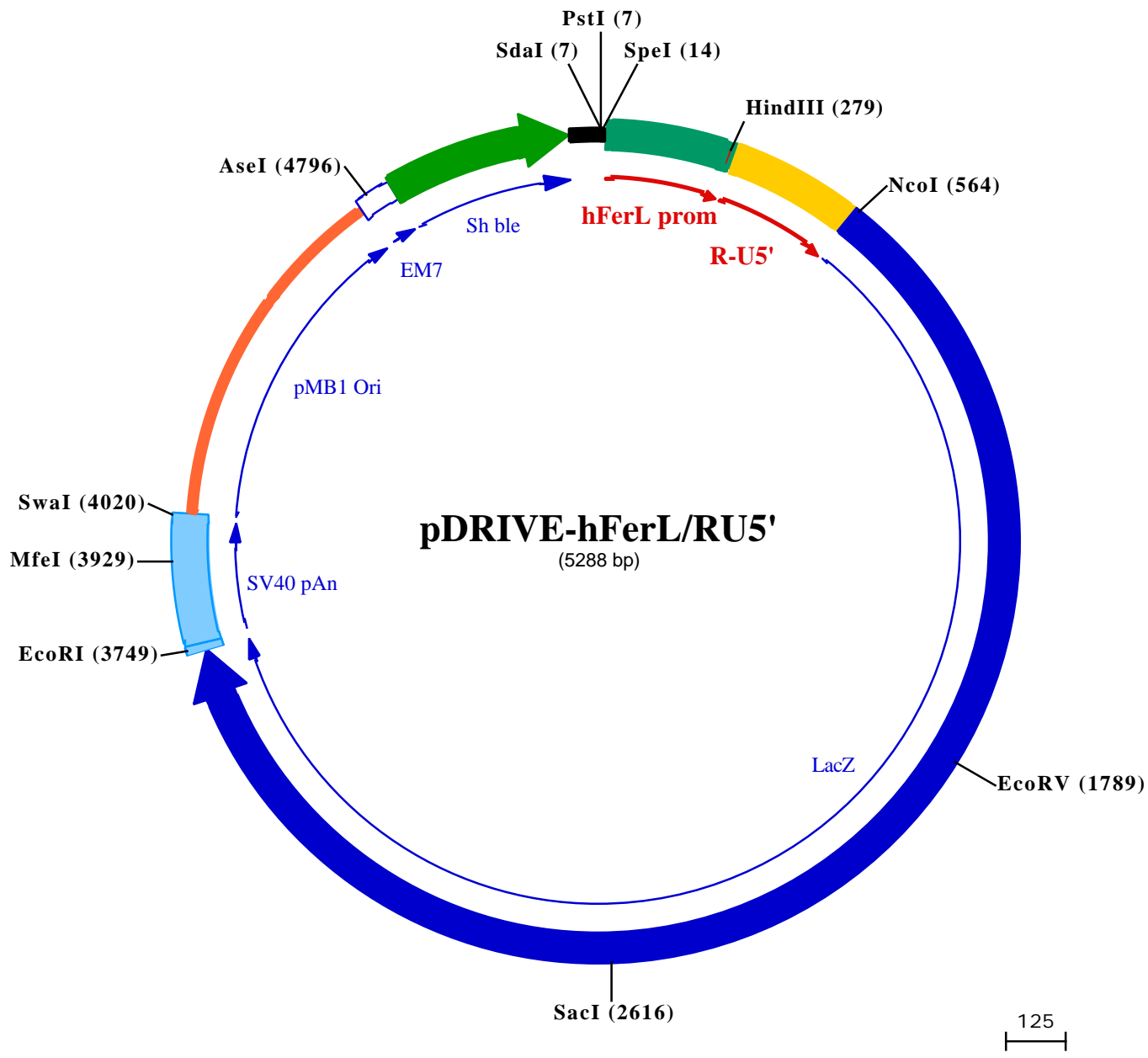
- 1- Eisenstein RS & Munro HN. (1990). *Enzyme* 44(1-4): 42-58.

## TECHNICAL SUPPORT

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

1 CCTGCAGGGCCCACTAGTCAGGGCCCCAACCCCCCAAGCCCCATTTCACAACACGCTGGCGCTACAGGGCGGTGACTTCCCCTTGCTTTGGGGGGGG

101 GGCTGAGACTCCTATGTGTCCGGATTGGTCAGGCACGGCCTTCGGCCCCGCTCCTGCCACCCGAGATTGGCCGCTAGCCCTCCCCGAGCGCCTGCCT

HindIII (279)

201 CCGAGGGCCGGCCACCATAAAAGAAGCCGCCCTAGCCACGTCCCCTCGCAGTTCGGGGTCCCGGGTCTGTCTCAAAGCTTCGAGGGGCTCGCATCTC

301 TCCTTACCGCGCCCGCCCTACCTGAGGCCGCCATCCACGCCGGTTGAGTCCGGTTCTGCCGCTCCCGCTGTGGTGCCTCCTGAACTGCGTCCGCC

401 GTCTAGGTAAGTTTAAAGCTCAGGTCGAGACCGGCCCTTTGTCCGGCGCTCCCTTGAGGCTACCTAGACTCAGCCGGCTCTCCACGCTTTGCCTGACCC

NcoI (564)

501 TGCTTGCTCAACTCTACGTCTTTGTTTTCTGTTTCTGCGCCGTTACAGATCAAGCCACCATGGGGGTTCTCATCATCATCATCATCATGGTATG

601 GCTAGCATGACTGGTGACAGCAAATGGGTCGGGATCTGTACGACGATGACGATAAGGTACCTAAGGATCAGCTTGGAGTTGATCCCGTCTTTTACAAC

701 GTCGTGACTGGGAAACCCCTGGCGTTACCAACTAATCGCCTTGACGACATCCCCCTTCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCG

801 CCCTTCCCAACAGTTGGCAGCCTGAATGGCAATGGCGCTTTCGCTGGTTCCGGCACCAGAAGCGGTGCCGAAAGCTGGCTGGAGTGGCATCTTCTC

901 GAGGCCGATACTGTCGTCGCCCTCAAACCTGGCAGATGACGGTTACGATGCGCCATCTACACCAACGTAACCTATCCCATACGGTCAATCCGCCGT

1001 TTGTTCCACGGAGAATCCGACGGGTTGTTACTCGCTCACATTTAATGTTGATGAAAGCTGGCTACAGGAAGCCAGACCGCAATATTTTTGATGGCGT

1101 TAACTCGGGTTTTCATCTGTGGTCAACGGCGCTGGGTGCGTTACGGCAGGACAGTCTGTTGCCGTCTGAATTTGACCTGAGCGCATTTTTACGGGCC

1201 GGAGAAAACCGCCTCGCGTGATGGTGTGCGTTGGAGTGACGGCAGTTATCTGGAAGATCAGGATATGTGGCGGATGAGCGGCATTTTCCGTGACGTCT

1301 CGTTGCTGCATAAACCGACTACAAAATCAGCGATTTCCATGTTCCACTCGCTTAAATGATGATTTACGCGCGCTGTACTGGAGGCTGAAGTTCAGAT

1401 GTGCGCGGAGTTGCGTGACTACCTACGGTAACAGTTTCTTTATGGCAGGGTGAACCCAGGTCGCCAGCGCCACCGCCCTTTCCGGCGTGAATATC

1501 GATGAGCGTGGTGTATGCCGATCGCGTCACACTACGCTGAAACGTCGAAAACCCGAAACTGTGGAGCGCCGAAATCCCGAATCTCTATCGTCCGGTGG

1601 TTGAATGCACACCGCCGACCGCAGCTGATTGAAGCAGAAGCCGTCGATGTCCGGTTCCCGGAGGTGGGATTGAAAATGCTCTGCTGCTGTAACCG

346 alGluLeuHisThrAlaAspGlyThrLeuIeGluAlaGluAlaCysAspValGlyPheArgGluValArgIeGluAsnGlyLeuLeuLeuLeuAsnGI

EcoRV (1789)

1701 CAAGCCGTTGCTGATTCGAGGCGTTAACCGTCACGAGCATCATCTCTGCATGGTCAAGGTCATGGATGAGCAGACGATGGTGCAGGATATCCTGCTGATG

1801 AAGCAGAACAACCTTAAACCGCGTGCCTGTCGATTATCCGAACCATCCGCTGTGGTACACGCTGTGGCAGCCGCTACGGCCTGATGTTGGATGAAG

1901 CCAATATTGAACCCACGGCATGGTGCCAATGAATCGTCTGACCGATGATCCGCGCTGGCTACCGGCGATGAGCGAACCAGCGAATGGTGCAGCG

2001 CGATCGTAATCACCGAGTGTGATCATCTGGTCCGCTGGGAATGAATCAGGCCACGGCGCTAATCAGCAGCGCTGTATCGCTGGATCAAAATCTGTCGAT

2101 CCTTCCCGCCCGTGCAGTATGAAGGCGCGGAGCCGACACCAGGCCACCGATATTATTTGCCGATGTACCGCGCGTGGATGAAGACCAGCCCTTCC

2201 CGGCTGTGCCGAAATGGTCCATCAAAAAATGGCTTTCGCTACCTGGAGAGACCGCCCGCTGATCCTTTGGCAATACGCCACCGCATGGGTAACAGTCT

2301 TGGCGGTTTTGCTAAATACTGGCAGGCTTTCGTCAGTATCCCGGTTACAGGCGCGTTCGCTGGGACTGGGTGGATCAGTCTGATTAATATGAT

2401 GAAAACGGCAACCGTGGTCCGCTTACCGCGGTGATTTGGCGATACCGGAACGATCGCCAGTCTCTGATGAACGGTCTGGTCTTTCCGACCGCACCG

2501 CGCATCCAGCGTACGGAAGCAAAACACCAGCAGCAGTTTTCCAGTTCGGTTATCCGGGCAACCATCGAAGTGACCAGCGAATACCTGTTCCGCTCA

646 roHisProAlaLeuThrGluAlaLysHisGlnGlnGlnPhePheGlnPheArgLeuSerGlyGlnThrIeGluValThrSerGluTyrLeuPheArgHi

SacI (2616)

2601 TAGCGATAACGAGCTCCTGCACTGGATGGTGGCGCTGGATGGTAAGCCGCTGGCAAGCGGTGAAGTGCCTCTGGATGTCCGCTCCACAAGGTAACAGTTG

2701 ATGAACTGCCTGAACTACCGCAGCCGGAGAGCGCCGCAACTCTGGCTCACAGTACCGTACGCTGCAACCGAACCAGCCGATGGTCCAGAACCCGGC

2801 ACATCAGCGCTGGCAGCAGTGGCGTCTGGCGGAAAACCTCAGTGTGACGCTCCCGCCGCGTCCACGCCATCCCGCATCTGACCACCAGCGAAATGGA

2901 TTTTTCATCGAGCTGGTAATAAGCGTTGGCAATTTAACCCGACGTCAGGCTTTCTTTACAGATGTGGATTGGCGATAAAAAACAACTGCTGACGCCG

3001 CTGCGCGATCAGTTACCCGTCACCCGCTGGATAACGACATTTGGCGTAAGTGAAGCGACCCCGCATTGACCCTAACCGCTGGGTCGAACCGTGGAAAGCGG

3101 CGGGCCATTACCAGCCGAAGCAGCGTGTGTCAGTGCACGGCAGATACACTTGTCTGATGCGGTGCTGATTACGACCGCTCACCGGTGGCAGCATCAGGG

3201 GAAAACCTTATTTATCAGCCGAAAACCTACCGGATTTGATGGTGGTCAAAATGGCGATTACCGTGTGTTGAGTGGCGAGCGCATACACCGCATCCG

3301 GCGCGGATTGGCTGAACTGCCAGCTGGCGCAGGTAGCAGAGCGGGTAAACTGGCTCGGATTAGGGCCGCAAGAAAATATCCCGACCGCCTTACTGCCG

913 AlaArgIeGlyLeuAsnCysGlnLeuAlaGlnValAlaGluArgValAsnTrpLeuGlyLeuGlyProGlnGluAsnTyrProAspArgLeuThrAlaA

3401 CCTGTTTTGACCGCTGGGATCTGCCATTGTGACACATGTATACCCCGTACGTCTTCCCAGCGAAAACGGTCTGCGCTGCGGGACGCGGAATTGAATTA  
946▶ laCysPheAspArgTrpAspLeuProLeuSerAspMetTyrThrProTyrValPheProSerGluAsnGlyLeuArgCysGlyThrArgGluLeuAsnTy  
3501 TGGCCACACCACTGGCGCGGCGACTTCCAGTTC AACATCAGCCGTACAGTCAACAGCAACTGATGAAACCAGCCATCGCCATCTGCTGCACGCGGAA  
979▶ rGlyProHisGlnTrpArgGlyAspPheGlnPheAsnI leSerArgTyrSerGlnGlnGlnLeuMetGluThrSerHisArgHisLeuLeuHisAlaGlu  
3601 GAAGGCACATGGCTGAATATCGACGGTTCCATATGGGGATTGGTGGCGACGACTCCTGGAGCCCGTCAGTATCGGCGGAATTACAGCTGAGCGCCGGTC  
1013▶ GluGlyThrTrpLeuAsnI leAspGlyPheHisMetGlyI leGlyGlyAspAspSerTrpSerProSerValSerAlaGluLeuGlnLeuSerAlaGlyA

**EcoRI (3749)**

3701 GCTACCATTACCAGTTGGTCTGGTGTCAAAAATAATAATCTAGTCGAGAATTCGCTAGCTCGACATGATAAGATACATTGATGAGTTTGGACAAACCACA  
1046▶ rgTyrHisTyrGlnLeuValTrpCysGlnLys•••  
3801 ACTAGAATGCAGTGAAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTAT

**MfeI (3929)**

3901 AAGCTGCAATAAACAAAGTTAACAAACAACAATTGCATTCATTTTATGTTTCAGGTTCAAGGGGAGGTGTGGGAGGTTTTTTAAAGCAAGTAAACCTCTAC

**SwaI (4020)**

4001 AAATGTGGTAGATCCATTTAAATGTTAATTAAGTCCATGACCAAAATCCCTTAACGTGAGTTTTTCGTTCCACTGAGCGTCAGACCCCGTAGAAAAGAT  
4101 CAAAGGATCTTCTTGAGATCCTTTTTTTCTGCGCGTAATCTGCTGCTTGC AAACAAAAAACACCAGCTACCAGCGGTGGTTTGTTCGCCGATCAAGAG  
4201 CTACCAACTCTTTTTCCGAAGGTAAGTGGCTTCAGCAGAGCGCAGATACCAATACTGTTCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAAGAACT  
4301 CTGTAGCACCCGCTACATACCTCGCTCTGCTAATCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCGTGTCTTACCAGGTTGGACTCAAGACGATA  
4401 GTTACCAGGATAAGGCGCAGCGGTTCGGCTGAACGGGGGGTTCGTGCACACAGCCAGCTTGGAGCGAACGACCTACACCGAACTGAGATACCTACAGCGT  
4501 GAGCTATGAGAAAGCGCCACGCTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAGGAGAGCGCACGAGGGAGCTTCCAG  
4601 GGGGAAACCGCTGGTATCTTTATAGTCTGTGCGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGCGGAGCCTATGGAA

**AseI (4796)**

4701 AAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGTGCGCCTTTTGTCTCACATGTTCTTAATTAATTTTTCAAAGTAGTTGACAATTAATC  
4801 ATCGGCATAGTATATCGGCATAGTATAATACGACTCACTATAGGAGGGCCATCATGGCCAAGTTGACCAGTGTGTCCAGTGTCTCACAGCCAGGGATGT  
4901 GGCTGGAGCTGTTGAGTTCTGGACTGACAGGTTGGGGTTCTCCAGAGATTTTGTGGAGGATGACTTTCAGGTTGGTTCAGAGATGATGTCACCCTGTTT  
16▶ lAlaGlyAlaValGluPheTrpThrAspArgLeuGlyPheSerArgAspPheValGluAspAspPheAlaGlyValValArgAspAspValThrLeuPhe  
5001 ATCTCAGCAGTCCAGGACCAGGTGGTGCCTGACAACACCCTGGCTTGGGTGTGGGTGAGAGGACTGGATGAGCTGTATGCTGAGTGGAGTGAGGTGGTCT  
50▶ l leSerAlaValGlnAspGlnValValProAspAsnThrLeuAlaTrpValTrpValArgGlyLeuAspGluLeuTyrAlaGluTrpSerGluValValS  
5101 CCACCAACTTCAGGGATGCCAGTGGCCCTGCCATGACAGAGATTGGAGAGCAGCCCTGGGGGAGAGAGTTTGCCTGAGAGACCCAGCAGGCAACTGTGT  
83▶ erThrAsnPheArgAspAlaSerGlyProAlaMetThrGluI leGlyGluGlnProTrpGlyArgGluPheAlaLeuArgAspProAlaGlyAsnCysVa  
5201 GCAC TTGTGCGAGGAGCAGGACTGAGGATAAGAATTGAGTTTCAGAAAAGGGGGCTGAGTGGCCCTTTTTTCAACTTAATTAA  
116▶ lHisPheValAlaGluGluGlnAsp•••