

pDRIVE5SEAP-hCD45

A plasmid with a native tissue-specific human CD45 promoter

Catalog # pdrive5s-hcd45

For research use only

Version # 11A21-MM

PRODUCT INFORMATION

Content:

- 1 disk of lyophilized GT116 *E. coli* bacteria transformed by pDRIVE5SEAP-hCD45.
- GT116 genotype is: *F*-, *mcrA*, Δ (*mrr-hsdRMS-mcrBC*), \emptyset 80*lacZ* Δ M15, Δ *lacX74*, *rspL* (*StrA*), *recA1*, *endA1* Δ *dcm* Δ *sbcC-sbcD*.
- 4 pouches of *E. coli* Fast-Media® Zeo (2 TB and 2 Agar)

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.

GENERAL PRODUCT USE

pDRIVE5-SEAP is an expression plasmid containing a native or composite promoter of interest. pDRIVE5-SEAP 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 are *Sda* 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 pDRIVE5-SEAP promoter drives the expression of the SEAP reporter gene which allows for testing of the promoter's activity in transient transfection experiments. Furthermore, the SEAP gene is flanked by unique restriction sites (*Nco* I and *Nhe* I) for easy replacement with a different gene of interest.

PROMOTER CHARACTERISTICS

Human CD45 promoter

Complete Promoter Size: 856bp
Plasmid backbone: pDRIVE5-SEAP
Specificity: Haematopoietic cells

CD45, also called leukocyte common antigen, is expressed exclusively by all hematopoietic cells except erythrocytes and platelets. CD45 is one of the most abundant leukocyte cell surface component and plays a pivotal role in antigen-stimulated proliferation of T lymphocytes and in thymic development. The CD45 promoter contains two major transcription initiation sites. The nucleotide sequence around the transcription initiation sites does not contain an apparent TATA box or CCAAT box probably due to the existence of multiple transcription initiation sites [1].

1. Hall L. *et al.* 1988. Complete exon-intron organization of the human leukocyte common antigen (CD45) gene. *J. Immunol.* 141(8):2781-87.

PLASMID FEATURES

- **SEAP gene** encodes an engineered secreted embryonic alkaline phosphatase. The levels of SEAP in the culture medium of transfected cells expressing the reporter gene can be assayed with chromogenic or luminescent methods
 - **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.
 - **EM2K** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.
 - **Zeo** gene confers zeocin resistance therefore allowing the selection of transformed *E. coli* carrying a pDRIVE5-SEAP 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 pDRIVE5-SEAP-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 pDRIVE5-SEAP plasmid DNA using the method of your choice.

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

E. coli Fast-Media® Zeo is a **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:

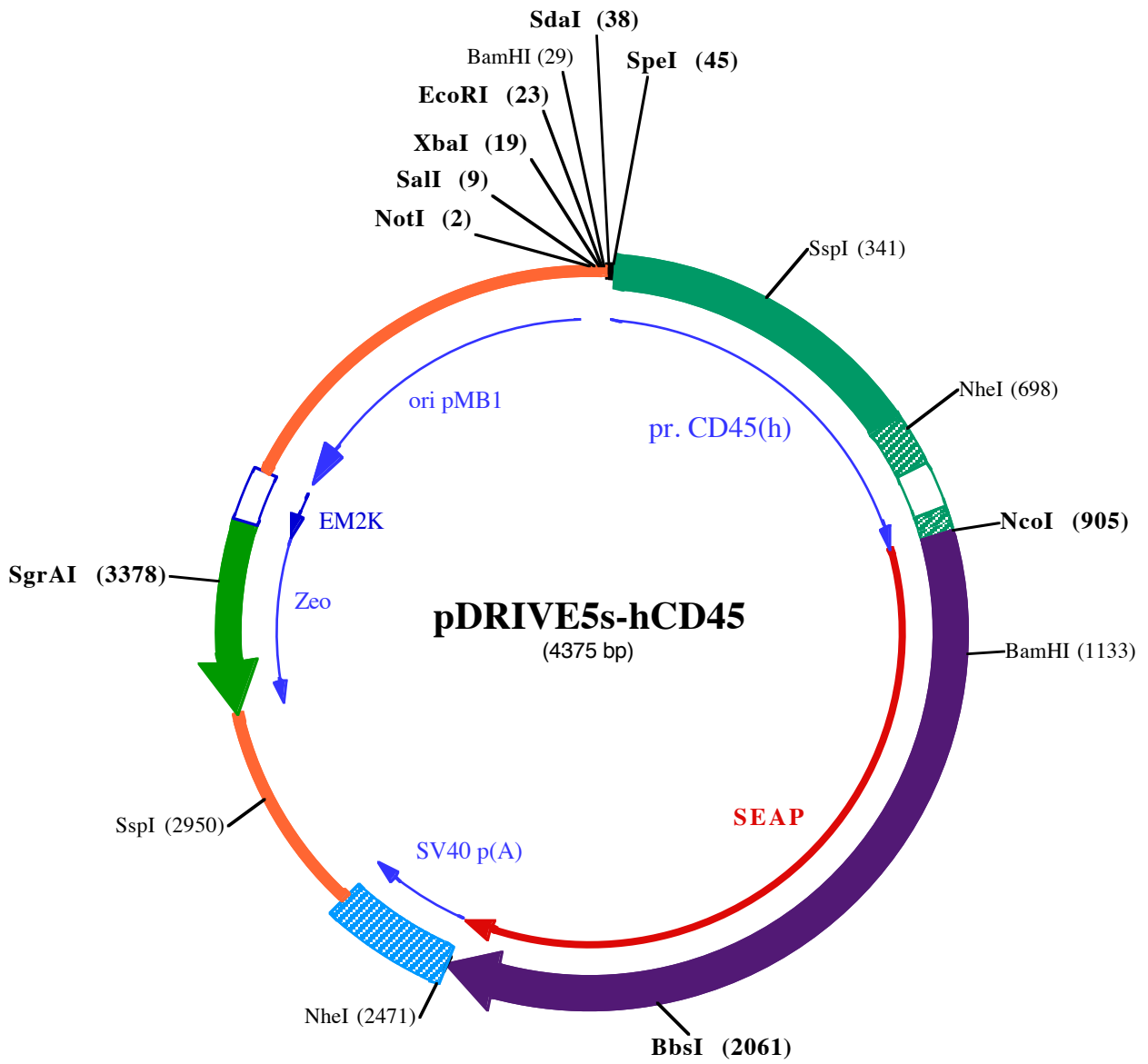
- 1- Pour the contents of a pouch into a clean borosilicate glass bottle or flask.
- 2- Add 200 ml of distilled water to the flask
- 3- Heat in a microwave on MEDIUM power setting (about 400Watts), until bubbles start appearing (approximately 3 minutes). **Do not heat a closed container. Do not autoclave 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.

TECHNICAL SUPPORT

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EcoRI (23) SdaI (38)
NotI (2) SalI (9) XbaI (19) BamHI (29) SpeI (45)

1 CCGGCCGCGTCGACGATATCTAGAATTCGGATCCTGCAGGGCCCACTAGTCCCAAGAACATCTTAAGTACAGAAAACATTAGTTTTGGAAGCAGGGTTT

101 GCTGTAAGTATAGTAGAATGACATTCTGATTCCACTCCTAGCTTCCACAAGGATATCTGTGAAAGATTTGGGCAAACTGTTAAGCTGTCTGAAAGTGC

201 TTTTGCATAAGAAATGGGTTTTACTGCTAAAAGTGCATATTTGCTGAGTTTTGAATGCCTTAATGTAATGATACTGGTGGCAAAAATAACAGATT

SspI (341)

301 AGTAGTTTTTTCATTCATTTGGCCGTCTCAGTAAGTCAAATATTGATACTTTCTACTAAGTCACTTGGCAACCCATTTGTATACCTTATGCTGAAT

401 CTGTTTTGCATCTCTTAAGTAAGAAAATTATTGATTATTTTGTGGGATTAAATTTAAAAAAATGGTAATGGATACTGTAAGGAGCATTATTTGGATG

501 GTTAAAAACATCTTCTCTGATGGGAAAATCTTTTAAAAAGGCTTCTAACTTGGTGTAAATTAAGTAAGGAAGTCAATGCCATTCTACTGACTTA

NheI (698)

601 GAACAACTTTTTACTCTGCAAAAGAGGACCCCTTACAGTATTTTGGAGAAGTTAGTAAACCAGAACTGACATCATCACCTAGCAGTTTCTCATGCAGCT

701 AGCAAGTGGTTTGTCTTAGGTAACAGAGGAGGAAATTTTCTCGTCTGTAAAGAACAGTGAGAgatgcatattttatttatttaccttttacatttt

801 tgattcgttttacagagaaaaactttctacagagatacaattattttgcttttcagAAGGACGCATGCTGTTTCTTAGGACACGGCTGACTTCCAGAT

NcoI (905)

901 ATGACCATGGTTCTGGGGCCCTGCATGCTGCTGCTGCTGCTGCTGGCCTGAGGCTACAGCTCTCCCTGGGCATCATCCAGTTGAGGAGGAGAACC

1001 CGGACTTCTGGAACCGCAGGACCGCAGGCCGCTGGTGGCCCAAGAGCTGCAGCTGCACAGACGCGCAAGAACCTCATCATCTTCTGGGGCA

32▶ P D F W N R E A A E A L G A A K K L Q P A Q T A A K N L I I F L G D

BamHI (1133)

1101 TGGGATGGGGTGTCTACGGTGCAGCTGCCAGGATCCTAAAAGGGCAGAAGAAGGCACAACTGGGGCTGAGATACCCCTGGTATGGACCGTCCCA

65▶ G M G V S T V T A A R I L K G Q K K D K L G P E I P L A M D R F P

1201 TATGTGGCTGTCCAAAGACATAAATGTAGACAAACATGTGCCAGACAGTTGGAGCCACAGCCACGGCTACCTGTGGGGTCAAGGGCACTCCAGA

99▶ Y V A L S K T Y N V D K H V P D S G A T A T A Y L C G V K G N F Q

1301 CCATTGGCTTGTGAGTGCAGCGCCCGCTTTAACCAGTGCACACGACCGGGCAACGAGGTCATCTCCGTGATGAATCGGGCAAGAAAAGCAGGGAAGTC

132▶ T I G L S A A A R F N Q C N T T R G N E V I S V M N R A K K A G K S

1401 AGTGGGAGTGGTAACCCACACAGAGTGACGACGCCTGCCAGCGGCACCTACGCCCACACGGTGAACCAGCACTGGTACTCGGACCGCCAGCTGCC

165▶ V G V V T T T R V Q H A S P A G T Y A H T V N R N W Y S D A D V P

1501 GCCTGGCCCGCAGGAGGGTGCAGGACATCGTACGACGCTCATCTCCAACATGGACATTGATGTGATCCTGGGTGGAGGGCCAAAAGTACATGTTT

199▶ A S A R Q E G C Q D I A T Q L I S N M D I D V I L G G G R K Y M F

1601 GCATGGGAACCCAGACCTGAGTACCAGATGACTACAGCAAGTGGGACAGGCTGGACGGGAAGAACTGGTGCAGGAATGGCTGGGAAGCGCCA

232▶ R M G T P D P E Y P D D Y S Q G G T R L D G K N L V Q E W L A K R Q

1701 GGGTGGCCGGTGTGTGGAACCGACTGAGCTCATGCAGGCTTCCCTGGACCGCTGTGACCCATCTCATGGGTCTCTTTGAGCCTGGAGACATGAAA

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

1801 TACGAGATCCACCGAGATCCACACTGGACCCCTCCCTGATGGAGATGACAGAGGCTGCCCTGCGCTGCTGAGCAGAACCCCGCGGCTTCTTCTCT

299▶ Y E I H R D S T L D P S L M E M T E A A L R L L S R N P R G F F L

1901 TCGTGGAGGGTGGTTCGATCGACCACGGTCATCAGAAAAGCAGGGCTTACCGGCCTGACTGAGACGATCATGTTTCGACGAGCCATTGAGAGGGCCGG

332▶ F V E G G R I D H G H H E S R A Y R A L T E T I M F D D A I E R A G

BbsI (2061)

2001 CCAGCTACCAGCGAGGAGACGCTGAGCTCGTCACTCGCAGCCTCCCACTCTTCTCCTTGGAGGCTACCCCTGCGAGGGAGCTCATCTTC

365▶ Q L T S E E D T L S L V T A D H S H V F S F G G Y P L R G S S I F

2101 GGGCTGGCCCTGGCAAGGCCGGGACAGGAAGGCCCTCACGGTCTCCTATACGGAACCGTCCAGGCTATGTGCTCAAGACGGCGCCGGCGGATG

399▶ G L A P G K A R D R K A Y T V L Y G N G P G Y V L K D G A R P D

2201 TTACCGAGAGCGAGAGCGGAGCCCGAGTATCGGACAGTGCAGCTGCCCCCTGGACGAAGAGACCCACGAGCGGAGCAGCTGGCGGTGTTCGCGCG

432▶ V T E S E S G S P E Y R Q Q S A V P L D E E T H A G E D V A V F A R

2301 CGGCCCGAGGCGCACCTGTTTCAGCGCTGCAGGAGCAGACCTTATAGCGCACGTCATGGCTTCCCGGCTGCTGGAGCCCTACACCGCTGCGAC

465▶ G P Q A H L V H G V Q E Q T F I A H V M A F A A C L E P Y T A C D

NheI (2471)

2401 CTGGCGCCCGCGGCCACCACCGCCGCACCCTGGGGCGGTCCCAGGCTGCCAGGCTGATGAAAGCTAGCTGGCCAGACATGATAAGATACATT

499▶ L A P P A G T T D A A H P G R S R S K R L D •

2501 GATGAGTTGGACAACCAACTAGAAATGCAAGTGAAGAAAATGCTTTATTTGTGAAAATTTGTGATGCTATTGCTTTATTTGTAACATTATAAGCTGCA

2601 ATAACAAGTTAAACAACAATTGCATTCATTTATGTTTTCAGTTTCAGGGGAGGTTGTTGGGAGGTTTTTAAAGCAAGTAAACCTCTACAAATGTGG

2701 TATGGAATTAATCTAAAATACAGCATAGCAAATTTAACCTCAAATCAAGCTCTACTTGAATCCTTTTCTGAGGATGAATAAGGCATAGGCATCA

2801 GGGGCTGTTGCAATGTCATTAGCTGTTTGCAGCCTCACCTTCTTTCATGGAGTTAAGATATAGTGATTTTTCCAAAGGTTTGAAGTACGCTTCATT

SspI (2950)

2901 TCTTTATGTTTAAATGCACTGACCTCCCACATTCCCTTTTAAATGTAATAATTTAGAATAATTTAAATACATCATTGCAATGAAAATAAATGTTTTTA

3001 TTAGGCAGAATCCAGATGCTCAAGGCCCTTCATAATATCCCCAGTGTAGTAGTTGGACTTAGGGAAACAAAGAACCTTTAATAGAAATTTGGACGACAG

3101 AAAGCGAGCTCTAGCTTATCCTCAGTCTGCTCCTCGCCACAAGTGCACGAGTTGCCGGCCGGTCCGCGAGGGCACTCCCGCCCCACGGCTG

125▶ • D Q E E A V F H V C N G A P D R L A F E R G W P Q

3201 CTCGCCGATCTGGTCTATGGCCGGCAGGCGTCCCGAAGTTCGTGGACACGACCTCCGACCCTCGCCGTCAGCTCGTCCAGGCCGCGCACCAC

99▶ E G I E T M A P G S A D R F N T S V V E S W E A Y L E D L G R V W

SgrAI (3378)

3301 ACCCAGGCCAGGGTTGTTCGGCACCACCTGGTCTGGACCGCTGATGAACAGGGTCACGCTGCTCCGGACACACCGCAAGTGTCTCCACGA

65▶ V W A L T N D P V V Q D Q V A S I F L T V D D R V V G A F D D E V F

3401 AGTCCCGGGAACCCGAGCCGCTCGTCCAGAACCTGACCGCTCCGGCGAGTCCGCGCGGGTGGAGCCGGAACGGCACTGGTCAACTTGGCCATGAT

32▶ D R S F G L R D T W F E V A G A V D R A T L V P V A S T L K A M ◀

3501 GGCTCTCTGTCCAGGAGGGAAGAGAAGGTTAGTACAATTTGCTATAGTGAGTTGATTATACTATGCAGATATACTATGCCAATGATTAATTTGTC

3601 AAAGTAGGGCTGCAGGTAATTAAGAACAATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAGGCCCGCTTGGCTGGGTTTTTCCATAGGCTC

3701 CGCCCCCTGACGAGCATCAAAAATCGACGCTCAAGTCCAGAGTGGCGAAACCAGAGGACTATAAAGATACCAGCGTTTTCCCTCGGAAGCTCCC

3801 TCGTGCGCTCTCCTGTTCGACCTGCGCTTACGGGATACCTGTCCGCTTTCTCCTTCCGGAAAGCGTGGCGCTTTTCTCATAGCTCACGCTGTAGGTA

3901 TCTCAGTTCCGTTGAGTGGTTCGCTCAAGCTGGGCTGTGTGCACGAACCCCGTTCAGCCCGACCGCTGCGCTTATCCGGTAACATATCGCTTTGAG

4001 TCCAAACCGGTAAGACAGCTTATCGCCACTGGCAGCAGCACTGGTAACAGGATTAGCAGAGCAGGATGTAGGCGGTGCTACAGAGTCTTGAAGT

4101 GGTGGCCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGATTGGTAGCTCTTGATCCGG

4201 CAAACAACCACCGCTGGTAGCGGTGGTTTTTTGTTTGAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTCTACG

4301 GGGTCTGACGCTCAGTGAACGAAAACACGTTAAGGGATTTGGTCATGGCTAGTTAATTAACATTAAATCA