

pDRIVE5SEAP-mCD68

A plasmid with a native tissue-specific mouse CD68 promoter

Catalog # pdrive5s-mcd68

For research use only

Version # 12J25-MM

PRODUCT INFORMATION

Content:

- 1 disk of lyophilized GT116 *E. coli* bacteria transformed by pDRIVE5SEAP-mCD68.
- GT116 genotype is: *F-*, *mcrA*, Δ (*mrr-hsdRMS-mcrBC*), \emptyset 80*lacZ* Δ M15, Δ *lacX74*, *rspL* (*StrA*), *recA1*, *endA1* Δ *dcn* 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 *BspLU* I 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

Mouse CD68 promoter

Complete Promoter Size: 812bp
Plasmid backbone: pDRIVE5-SEAP
Specificity: Macrophages

Murine macrophage and its human homolog CD68 are heavily glycosylated transmembrane proteins expressed specifically in macrophages and macrophage-related cells. The CD68 gene promoter directs macrophage-specific expression. Although the promoter lacks a classical TATA box, it contains other protein binding sites consistent with preferential monocyte/macrophage gene expression¹. CD68 promoter has been used to achieve constitutive expression of IL-10 specifically in macrophages².

1. Jiang Z. *et al.* 1998. Structure, organization, and chromosomal mapping of the gene encoding macrophage-specific protein. *Genomics* 50(2):199-205.
2. Lang R. *et al.* 2002. Autocrine deactivation of macrophages in transgenic mice constitutively overexpressing IL-10 under control of the human CD68 promoter. *J Immunol.* 168(7):3402-11.

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:

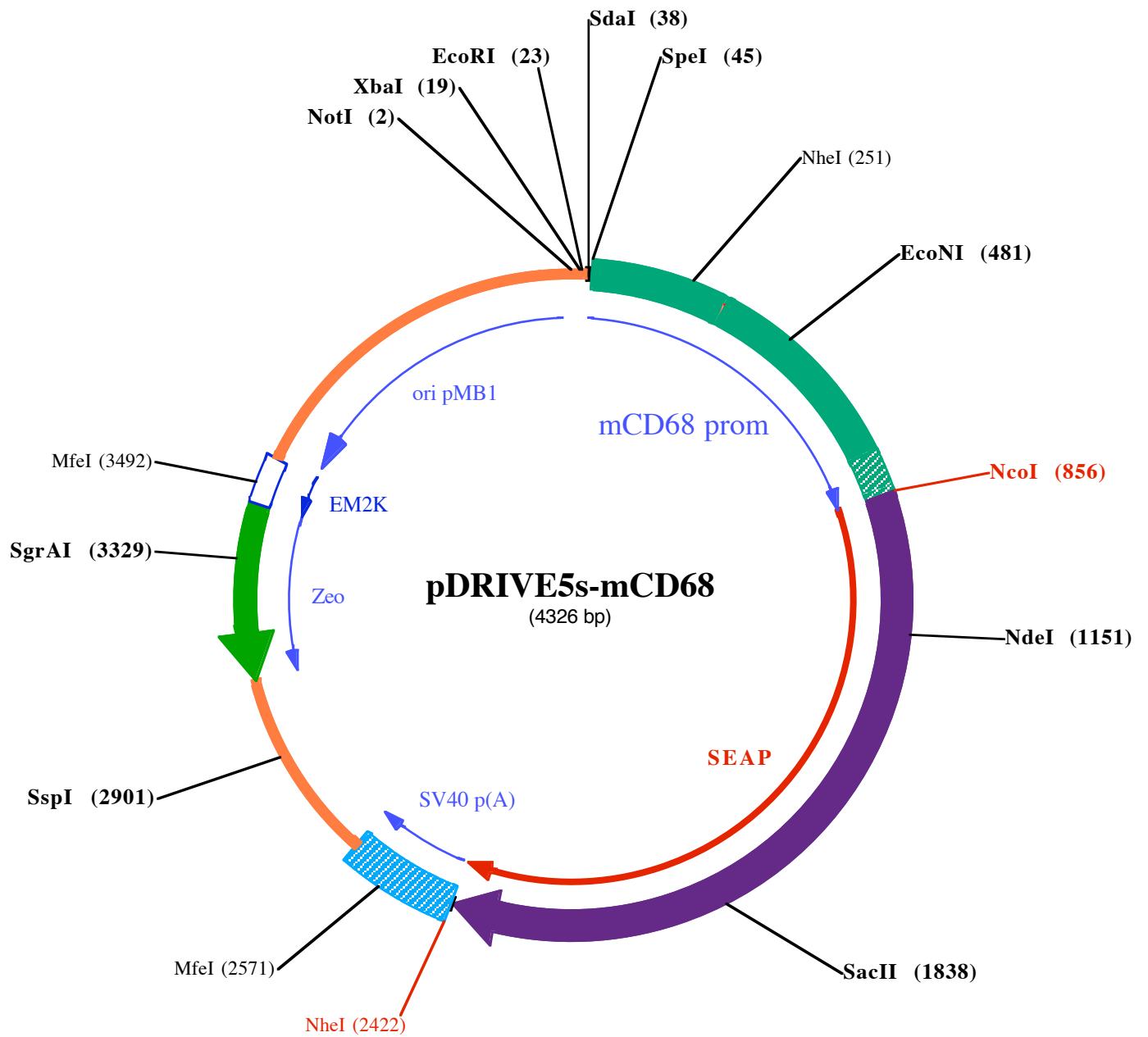
- 1- Pour the contents of a pouch into a clean borosilicate glass bottle or flask.
- 2- Add 200 ml of distilled water to the flask
- 3- Heat in a microwave on MEDIUM power setting (about 400Watts), until bubbles start appearing (approximately 3 minutes). **Do not heat a closed container. Do not autoclave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

Toll free (US): 888-457-5873
Outside US: (+1) 858-457-5873
Europe: +33 562-71-69-39
E-mail: info@invivogen.com
Website: www.invivogen.com


3950 Sorrento Valley Blvd. Suite 100
San Diego, CA 92121 - USA



EcoRI (23)

NotI (2) **XbaI (19)** **SdaI (38)** **SpeI (45)**

1 **GCGGCCGCGTCGACGATATCTAGAATTCGGATCCTGCAGGGCCCACTAGTTGATTACTGAATTTGCCATGTTGC**

75 **TTCCTGCAATACCAAATGACCCACATTACTAACATTTGGTAATTTGCCTCAGTGTTAGGCACAAGACATGAGTG**

149 **AAACTTGCTGCATTAAGACTTAAACTGGCTGGGTATGGTGGCGCAAGCCTTTAATCCCAGCATTCTGCCGGAGG**

NheI (251)

223 **CAGAAGCAGGTGGATTTCTGAGTTCGAGGCTAGCCTGGTCTACAAAAGTGAGTTCAGGACAGCCAAGGCTACA**

297 **CAAATACCTGTCTTGAAAAACCAAAAAAAAAAAAAAATTCAACTGGTTGGTTAGGCCTAGCCAGATGATGTAG**

371 **GTGTCAACTCACCTTGGGGATGTAGGGAGATGGTTGTTCTGTAGCTCTTCTGTCTGCAACTAAATAAATAGG**

EcoNI (481)

445 **CTAAGCTGACCTTTACATCTGCCTTCGGCCTCCTGTGCTAGGATTGGAGTGCATGGGTGGGGTTGCATTTCTT**

519 **ACCAATGAGGAAAGGGCTTCCATTTCTCTGCTGCATTGTAAGCTCCCAAGTAGAGCAAGTTTTGCTTAGGTAA**

593 **GTTCCGTGTGAGTCAGCTGCCTCATTCTCACGAGGTAACCAAGGCTTTGTACCGCCACTGAGAACGTCAGTGC**

667 **CAATCACAGCCTAATTGTGAAAACCAATGGCTTGAGTGGGTGCTAAAGCTGAGGTGTCTGAGTCAGGTTTGGG**

741 **GTGGGATTATTTTAGTTAAGGGAAGTGAGGCTTTTCATTTCTCTTCCAAGAGAAGGCAAAGGGGATTGGATTG**

NcoI (856)

815 **AGGAAGGAAGTGGTGTAGCCTAGCTGGTCTGAGCATCTCTGCCATGGTTCTGGGGCCCTGCATGCTGCTGCTGC**

M V L G P C M L L L

889 **TGCTGCTGCTGGGCCTGAGGCTACAGCTCTCCCTGGGCATCATCCCAGTTGAGGAGGAGAACCCGGACTTCTGG**

110 L L L L G L R L Q L S L G I I P V E E E N P D F W

963 **AACCGCGAGGCAGCCGAGGCCCTGGGTGCCCAAGAAGCTGCAGCCTGCACAGACAGCCGCAAGAACCTCAT**

36 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

1037 **CATCTTCTGGGCGATGGGATGGGGGTGTCTACGGTGACAGCTGCCAGGATCCTAAAAGGGCAGAAGAAGGACA**

60 I F L G D G M G V S T V T A A R I L K G Q K K D

NdeI (1151)

1111 **AACTGGGGCCTGAGATACCCCTGGCTATGGACCGCTTCCCATATGTGGCTCTGTCCAAGACATACAATGTAGAC**

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

1185 **AAACATGTCCAGACAGTGGAGCCACAGCCACGGCCTACCTGTGCGGGTCAAGGGCAACTTCCAGACCATTGG**

110 K H V P D S G A T A T A Y L C G V K G N F Q T I G

1259 **CTTGAGTGCAGCCGCCGCTTTAACAGTGCAACACGACACGCGCAACGAGGTCATCTCCGTGATGAATCGGG**

134 L S A A A R F N Q C N T T R G N E V I S V M N R

1333 **CCAAGAAAGCAGGGAAGTCAGTGGGAGTGGTAACCACCACACGAGTGCAGCACGCTCGCCAGCCGGCACCTAC**

159 A K K A G K S V G V V T T T R V Q H A S P A G T Y

1407 **GCCCACACGGTGAACCGCAACTGGTACTCGGACGCCGACGTGCTGCTCGGCCCGCCAGGAGGGGTGCCAGGA**

184 A H T V N R N W Y S D A D V P A S A R Q E G C Q D

1481 **CATCGCTACGACGCTCATCTCCAACATGGACATTGATGTGATCCTGGGTGGAGGCCGAAAGTACATGTTTCGCA**

208 I A T Q L I S N M D I D V I L G G G R K Y M F R

1555 **TGGGAACCCAGACCCTGAGTACCCAGATGACTACAGCCAAGGTGGGACCAGGCTGGACGGGAAGAATCTGGTG**

233 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

1629 **CAGGAATGGCTGGCGAAGCGCCAGGGTCCCCGGTATGTGTGGAACCGCACTGAGCTCATGCAGGCTTCCCTGGA**

258 Q E W L A K R Q G A R Y V W N R T E L M Q A S L D

1703 **CCCGTCTGTGACCCATCTCATGGGTCTCTTTGAGCCTGGAGACATGAAATACGAGATCCACCGAGACTCCACAC**

282 P S V T H L M G L F E P G D M K Y E I H R D S T

SacII (1838)

1777 **TGGACCCCTCCCTGATGGAGATGACAGAGGCTGCCCTGCGCCTGCTGAGCAGGAACCCCGCGGCTTCTTCTCCT**

307 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

1851 **TTCGTGGAGGGTGGTGCATCGACCACGGTCATCACGAAAGCAGGGCTTACCGGGCACTGACTGAGACGATCAT**

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

1925 **GTTTCGACGACGCCATTGAGAGGGCGGGCCAGCTCACCAGCGAGGAGGACACGCTGAGCCTCGTCACTGCCGACC**

356 F D D A I E R A G Q L T S E E D T L S L V T A D

1999 **ACTCCCACGTCTTCTCCTTCGGAGGCTACCCCTGCGAGGGAGCTCCATCTTCGGGCTGGCCCTGGCAAGGCC**

381 H S H V F S F G G Y P L R G S S I F G L A P G K A

2073 CGGGACAGGAAGGCCTACACGGTCTCTCTATACGGAAACGGTCCAGGCTATGTGCTCAAGGACGGCGCCCGGCC
406 R D R K A Y T V L L Y G N G P G Y V L K D G A R P
2147 GGATGTTACCGAGAGCGAGAGCGGGAGCCCCGAGTATCGGCAGCAGTCAGCAGTGCCCCTGGACGAAGAGACCC
430 D V T E S E S G S P E Y R Q Q S A V P L D E E T
2221 ACGCAGGCGAGGACGTGGCGGTGTTGCGCGCGGCCCGCAGGCGCACCTGGTTCACGGCGTG CAGGAGCAGACC
455 H A G E D V A V F A R G P Q A H L V H G V Q E Q T
2295 TTCATAGCGCACGTCATGGCCTTCGCCGCTGCTGGAGCCCTACACCGCCTGCGACCTGGCGCCCCCGCCGG
480 F I A H V M A F A A C L E P Y T A C D L A P P A G

NheI (2422)

2369 CACCACCGACGCCGCGCACCCGGGGCGGTCCCGGTCCAAGCGTCTGGATTGAAAGCTAGCTGGCCAGACATGATA
504 T T D A A H P G R S R S K R L D •
2443 AGATACATTGATGAGTTTGGACAAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTGAAATTTGTGA

MfeI (2571)

2517 TGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTCATTTTATGT
2591 TTCAGGTT CAGGGGAGGTGTGGGAGGTTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTAATT

2665 CTAAATACAGCATAGCAAACTTTAACCTCCAAATCAAGCCTCTACTTGAATCCTTTTCTGAGGGATGAATAA
2739 GGCATAGGCATCAGGGGCTGTTGCCAATGTGCATTAGCTGTTTGCAGCCTCACCTTCTTTTCATGGAGTTTAAGA
2813 TATAGTGTATTTCCCAAGGTTTGAAGTACTAGCTCTTCATTTCTTTATGTTTTAAATGCACTGACCTCCACATTC

SspI (2901)

2887 CCTTTTTAGTAAATATTCAGAAATAATTTAAATACATCATTGCAATGAAAATAAATGTTTTTTATTAGGCAGA
2961 ATCCAGATGCTCAAGGCCCTTCATAATATCCCCAGTTTAGTAGTTGGACTTAGGGAACAAAGGAACCTTTAAT
3035 AGAAATTGGACAGCAAGAAAGCGAGCTTCTAGCTTATCCTCAGTCTGCTCCTCTGCCACAAAGTGACCGCAGT

125 •

D Q E E A V F H V C N

3109 TGCCGGCCGGGTGCGCGAGGGCGAAGTCCC GCCCCCACGGCTGCTCGCCGATCTCGGTCATGCCC GGCCCGGAG
113 G A P D R L A F E R G W P Q E G I E T M A P G S
3183 GCGTCCCGAAGTTCGTGGACACGACCTCCGACCACTCGGCGTACAGCTCGTCCAGGCCGCGCACCCACACCCA
88 A D R F N T S V V E S W E A Y L E D L G R V W V W

SgrAI (3329)

3257 GGCCAGGGTGTGTCCGGCACCACCTGGTCTGGACCGCGCTGATGAACAGGGTCACGTCGTC CCGGACCACAC
64 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
3331 CGGCGAAGTCGTCTCCACGAAGTCCCGGAGAACCCGAGCCGGTCCGTTCCGGAAGTCCGACCGCTCCGGCGACG
39 A F D D E V F D R S F G L R D T W F E V A G A V
3405 TCGCGCGCGGTGAGCACCGGAACGGCACTGGTCAACTTGGCCATGATGGCTCCTCCTGTCAGGAGAGGAAAGAG
14 D R A T L V P V A S T L K A M

MfeI (3492)

3479 AAGAAGTTAGTACAATTGCTATAGTGAGTTGATTATACTATGCAGATATACTATGCCAATGATTAATTGTCA
3553 AACTAGGGCTGCAGGTTAATTAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGC
3627 GTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGC
3701 GAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCCCTGGAAGTCCCTCGTGCGCTCTCCTGTTCCGACC
3775 CTGCCGTTACCGGATACCTGTCCGCCTTCTCCCTCGGGAAGCGTGGCGCTTTCATAGCTCACGCTGTAG
3849 GTATCTCAGTTCGGTGTAGGTCGTTCCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCGTT CAGCCCGACCGCT
3923 GCGCCTTATCCGTAAC TATCGTCTTGAGTCCAACCCGTAAGACACGACTTATCGCCACTGGCAGCAGCCACT
3997 GGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTA
4071 CACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGAAAAAGAGTTGGTAGCTCTT
4145 GATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAA
4219 GGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAAACGAAAAC TACGTTAAGGGAT
4293 TTTGGTCATGGCTAGTTAATTAACATTTAAATCA