

# pDRIVE-hCCKAR

A plasmid with the native human CCKAR promoter

Catalog # pdrive-heckar

For research use only

Version # 10D26-MM

## PRODUCT INFORMATION

### Content:

- 1 disk of lyophilized GT115 *E. coli* bacteria transformed by pDRIVE-hCCKAR.
- GT115 genotype is: *F mcrA Δ(mrr-hsdRMS-mcrBC) φ80lacZΔM15 ΔlacX74 recA1 rspL (StrA) endA1 Δdcm uidA(ΔMlu1)::pir-116 ΔsbcC-sbcD*.
- 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® 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

**hCCKAR promoter-** Complete promoter size: 724 bp

Cholecystokinin (CCK) is a gut peptide hormone known to stimulate post-prandial gallbladder contraction and pancreatic enzyme secretion. CCK also plays an important role in pancreatic carcinogenesis. The CCK-A receptor (CCKAR) is predominantly expressed in human pancreatic cancer<sup>1</sup>. A 0.7 kb fragment of the CCKAR promoter was reported to drive the expression of the luciferase gene selectively in pancreatic cancer cells and not in normal cells<sup>2</sup>. This fragment lacks a typical TATA box but contains two GC-Box motifs<sup>3</sup>.

1. Moonka R. *et al.* 1999. Cholecystokinin-A receptor messenger RNA expression in human pancreatic cancer. *J Gastrointest Surg.* 3(2): 134-40. 2. Li Z. *et al.* 2005. Suppression of pancreatic tumor progression by systemic delivery of a pancreatic-cancer-specific promoter driven Bik mutant. *Cancer Lett.* 236(1):58-63. 3. Takata Y. *et al.* 2002. Promoter analysis of human cholecystokinin type-A receptor gene. *J Gastroenterol.* 37(10):815-20.

## 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-1, 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

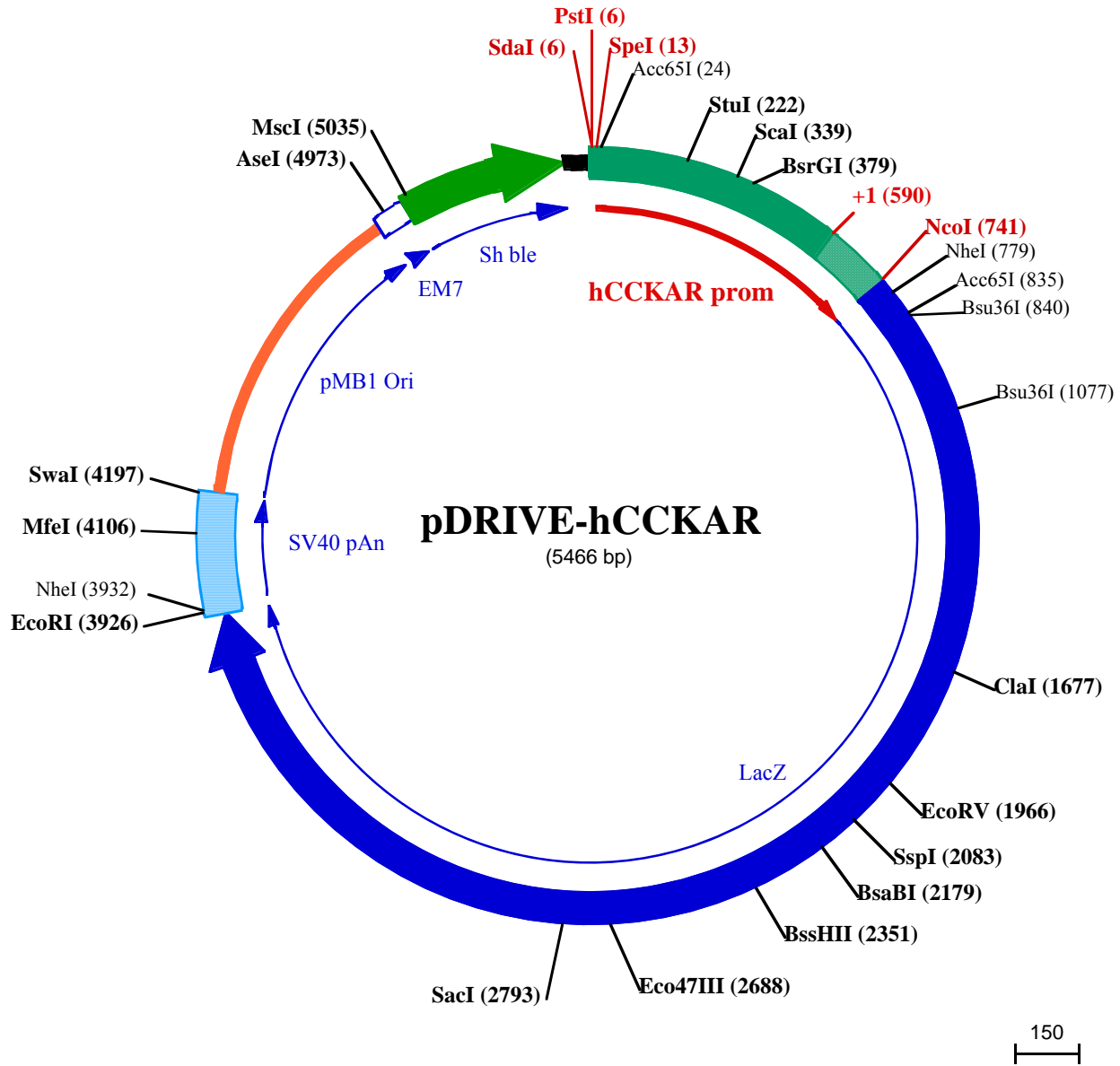
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Eco47III (2688)

2601 CTTACGGCGGTGATTTTGGCGATACGCCGAACGATCGCCAGTCTGTATGAACGGTCTGGTCTTTGCCGACCGCACGCCGCATCCAGCGCTGACGGAAGC

620 ▶ I aTy rGl yGl yAspPheGl yAspThrP roAsnAspA rGlnPheCysMetAsnGl yLeuValPheAl aAspArgThrProHi sProAl aLeuThrGl uAl

SacI (2793)

2701 AAAACACCAGCAGCAGTTTTTCCAGTTCGGTTATCCGGGCAAACCTCGAAGTGACCAGCGAATACCTGTTCCGTCATAGCGATAACGAGCTCCTGCAC

653 ▶ aLysHi sGl nGl nPhePheGl nPheA rGLeuSerGl yGl nThr l l eGl uVal ThrSerGl uTy rLeuPheArgHi sSerAspAsnGl uLeuLeuHi s

2801 TGGATGGTGGCGCTGGATGGTAAGCCGCTGGCAAGCGGTGAAGTGCCTCTGGATGTCGCTCCACAAGGTAACAGTTGATTGAACCTGCCTGAACCTACCGC

687 ▶ TrpMetValAl aLeuAspGl yLysProLeuAl aSerGl yGl uValProLeuAspValAl aProGl nGl yLysGl nLeu l eGl uLeuProGl uLeuProG

2901 AGCCGGAGAGCGCCGGGCAACTCTGGCTCACAGTACCGGTAGTGAACCGAACCGGACCGCATGGTCAGAAGCCGGGCACATCAGCGCTGGCAGCAGTG

720 ▶ InProGl uSerAl aGl yGl nLeuTrpLeuThrValArgValValGl nProAsnAl aThrAl aTrpSerGl uAl aGl yHi s l l eSerAl aTrpGl nGl nTr

3001 GCGTCTGGCGAAAACCTCAGTGTGACGCTCCCGCGCGTCCCACGCCATCCCGCATCTGACCACCAGCGAAATGGATTTTGCATCGAGCTGGGTAAT

753 ▶ pArgLeuAl aGl uAsnLeuSerValThrLeuProAl aAl aSerHi sAl a l l eProHi sLeuThrThrSerGl uMetAspPheCys l eGl uLeuGl yAsn

3101 AAGCGTGGCAATTTAACCGCCAGTACGGCTTCTTTCACAGATGTGGATTGGCGATAAAAAACAACCTGCTGACGCCGTGGCCGATCAGTTACCCCGTG

787 ▶ LysArgTrpGl nPheAsnA rGl nSerGl yPheLeuSerGl nMetTrp l l eGl yAspLysLysGl nLeuLeuThrProLeuArgAspGl nPheThrArgA

3201 CACCGCTGGATAACGACATTTGGCGTAAGTGAAGCGACCCGATTGACCTAACGCCTGGTGAACGCTGGAAGGCGGGGCCATTACCGCCGAAGC

820 ▶ I aProLeuAspAsnAsp l l eGl yValSerGl uAl aThrArg l l eAspProAsnAl aTrpValGl uArgTrpLysAl aAl aGl yHi sTy rGl nAl aGl uAl

3301 AGCGTGTTCAGTGCACGGCAGATACACTTGCTGATGCGGTGCTGATTACGACCCTCACCGTGGCAGCATCAGGGGAAAACCTATTTATCAGCCGG

853 ▶ aAl aLeuLeuGl nCysThrAl aAspThrLeuAl aAspAl aValLeu l l eThrThrAl aHi sAl aTrpGl nHi sGl nGl yLysThrLeuPhe l l eSerArg

3401 AAAACCTACCGGATTTGATGGTAGTGGTCAAAATGGCGATTACCGTTGATGTGAAGTGCCGAGCGATACCCGCATCCGGCGGGATTGGCTGAACCTGCC

887 ▶ LysThrTy rArg l l eAspGl ySerGl yGl nMetAl a l l eThrValAspValGl uValAl aSerAspThrProHi sProAl aArg l l eGl yLeuAsnCysG

3501 AGCTGGCCAGGTAGCAGAGCGGTAACCTGGCTCGGATTAGGCGCCGAAAGAACTATCCGACCGCTTACTGCCGCTGTTTTCAGCCGCTGGGATCT

920 ▶ InLeuAl aGl nValAl aGl uArgValAsnTrpLeuGl yLeuGl yProGl nGl uAsnTy rProAspArgLeuThrAl aAl aCysPheAspArgTrpAspLe

3601 GCCATTGTACAGACATGTATACCCCGTACGCTTCCCGAGCGAAAACGGTCTGCGCTGCGGGACGCGCAATTGAATTATGGCCACACCGAGTGGCGCGGC

953 ▶ uProLeuSerAspMetTy rThrProTy rValPheProSerGl uAsnGl yLeuArgCysGl yThrArgGl uLeuAsnTy rGl yProHi sGl nTrpA rGl y

3701 GACTTCCAGTTCACATCAGCCGCTACAGTCAACAGCAACTGATGGAACAGCCATCGCCATCTGCTGCACGCGGAAGGACACATGGCTGAATATCG

987 ▶ AspPheGl nPheAsn l l eSerArgTy rSerGl nGl nLeuMetGl uThrSerHi sArgHi sLeuLeuHi sAl aGl uGl yThrTrpLeuAsn l l eA

3801 ACGGTTTCCATATGGGATTGGTGGCGACGACTCTGGAGCCGCTCAGTATCGCGGAATTACAGCTGAGCGCCGGTCCGTACCATTACAGTTGGTCTG

1020 ▶ spGl yPheHi sMetGl y l l eGl yGl yAspAspSerTrpSerProSerValSerAl aGl uLeuGl nLeuSerAl aGl yArgTy rHi sTy rGl nLeuValTr

NheI (3932)

3901 GTGTCAAAAAATAAATCTAGTCGAGAAATTCGCTAGCTCGACATGATAAGATACATTGATGAGTTTGGCAAAACCACAACCTAGAATGCAGTGAAAAAAT

1053 ▶ pCysGl nLys•••

4001 GCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACATTATAAGCTGCAATAAACAAGTTAAC

MfeI (4106)

SwaI (4197)

4101 AACACAATTCGATTCATTTTATGTTTCAGGTTTCAGGGGAGGTGGGGAGGTTTTTAAAGCAAGTAAAACCTCTACAAATGGTAGATCCATTTAAA

4201 TGTTAATTAAGTACGATGACCAAAATCCCTTAACGTGAGTTTTTCGTTCCACTGAGCGTCAGACCCGTCAGAAAAGATCAAAGGATCTTCTTGAGATCCT

4301 TTTTTCTGCGGTAATCTGCTGCTTGCACAAACAAAAACCCGCTACCAGCGGTGGTTTGGTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGG

4401 TAACTGGCTTCAGCAGAGCCGAGATACCAATACGTTCTTCTAGTGTAGCCGTAGTTAGGCCACCACTCAAGAACTCTGTAGCACCCTACATACCT

4501 CGCTCTGCTAATCCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCTGTCTTACCAGGTTGGACTCAAGACGATAGTTACCGGATAAGGGCCAGCGG

4601 TCGGGCTGAACGGGGGTTCTGTGCACACAGCCGCTGGAGCGAACGACCTACCCGAACCTGAGATACCTACAGCGTGAGCTATGAGAAAGCCGACGC

4701 TTCCCGAAGGGGAGAAAGGCGACAGGTATCCGGTAAGCGCGAGGTCGGAACAGGAGCGCACGAGGGAGCTTCCAGGGGAAACGCTGGTATCTTTA

4801 TAGTCTGTGCGGTTTCGACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGCGGAGCCTATGAAAAACGCCAGCAACCGCCGCTTT

AseI (4973)

4901 TTACGGTTCCTGGCCTTTGCTGGCCTTTGCTCACATGTTCTTAATTAATTTTTCAAAGTAGTTGACAATTAATCATCGGCATAGTATATCGGCATA

MseI (5035)

5001 GTATAATACGACTCACTATAGGAGGGCCATGAGCCAGTTGACCAGTGTGTCCAGTGTCCAGTGTCCAGCCAGGATGTGGCTGGAGCTGTTGAGTTCTGG

5101 ACTGACAGGTTGGGTTCTCCAGAGATTTTGTGAGGATGACTTTGCAGGTTGGTCAGAGATGATGTCACCTGTTCTATCTCAGCAGTCCAGGACGAGG

24 ▶ ThrAspA rGLeuGl yPheSerArgAspPheValGl uAspAspPheAl aGl yValValA rGAspAspValThrLeuPhe l l eSerAl aValGl nAspGl nV

5201 TGGTGCCTGACAACACCTGGCTTGGTGTGGTGAGAGGACTGGATGAGCTGTGCTGAGTGGAGTGGTGGTCTCCACCAACTCAGGGATGCCAG

57 ▶ a lValProAspAsnThrLeuAl aTrpValTrpValA rGl yLeuAspGl uLeuTy rAl aGl uTrpSerGl uValValSerThrAsnPheA rGAspAl aSe

5301 TGCCCTGCCATGACAGAGATTGGAGAGCAGCCCTGGGGGAGAGAGTTGCCCTGAGAGACCCAGCAGGCAACTGTGTGACCTTTGTGGCAGAGAGCAG

90 ▶ rGl yProAl aMetThrGl u l l eGl yGl uGl nProTrpGl yArgGl uPheAl aLeuArgAspProAl aGl yAsnCysValHi sPheValAl aGl uGl uGl n

5401 GACTGAGGATAAGAATTGAGTTTCAGAAAAGGGGCGCTGAGTGGCCCTTTTTCAACTTAATTA

124 ▶ Asp•••