

pDRIVE-hGRP78

A plasmid with the native human Glucose Regulated Protein 78 promoter

Catalog # pdrive-hgrp78

For research use only

Version # 01F07-MT

PRODUCT INFORMATION

Content:

- 1 disk of lyophilized GT100 *E. coli* bacteria transformed by a pDRIVE plasmid.
- GT100 genotype is: *F-*, *mcrA*, $\Delta(mrr-hsdRMS-mcrBC)$, $\emptyset80lacZ\Delta M15$, $\Delta lacX74$, *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 *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 *Eco* R I) for easy replacement with a different gene of interest.

PROMOTER CHARACTERISTICS

Element	Name	Origin	Size bp
Promoter	GRP78	Human	310
5'UTR	GRP78	Human	221
Enhancer	-	-	-

GRP78 promoter

The glucose-regulated protein 78 (GRP78) functions as molecular chaperone. It is expressed constitutively in the endoplasmic reticulum in most cell types under normal growth conditions and are highly induced in stressed cells. Inducing factors are cellular environments of low glucose or oxygen and reagents that disrupt the ER function such as calcium ionophores. The GRP78 promoter is known to retain its strong activity in differentiated and undifferentiated tissues¹. Furthermore it has been demonstrated that *in vivo*, the GRP78 promoter can increase the expression levels of HSV-tk inside tumors resulting in complete eradication of tumor mass, with no recurrence of tumor growth².

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 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 FastMedia™ as the antibiotic will be permanently destroyed by the procedure.

References:

- 1- Kim *et al.* (1990). Differentiation 42:153-9.
2. Gazit G. *et al.* (1999). Cancer Res 59: 3100-6

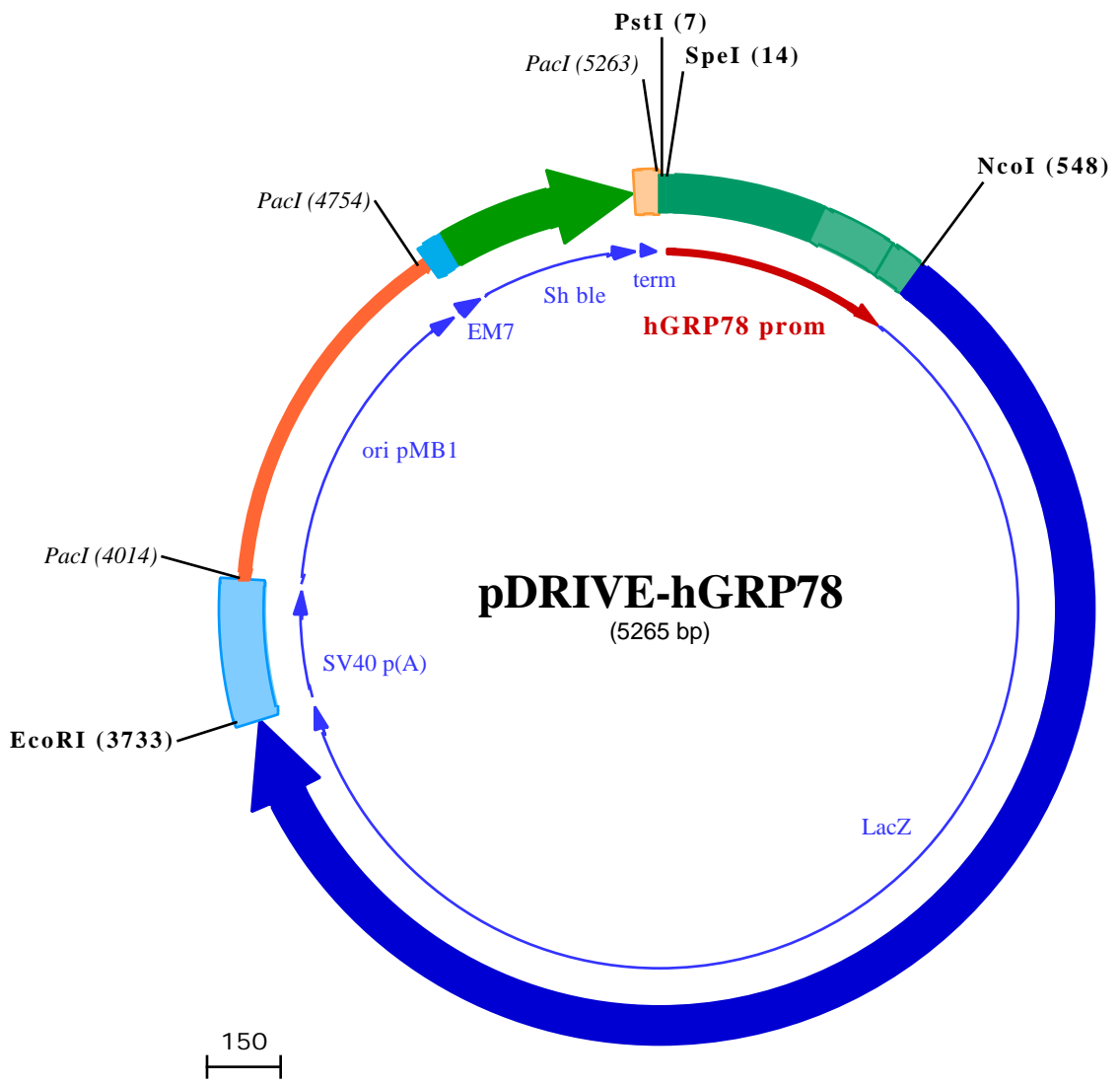
TECHNICAL SUPPORT

Toll free (US): 888-457-5873

Outside US: (+1) 858-457-5873

E-mail: info@invivogen.com

Website: www.invivogen.com



PstI (7) SpeI (14)

1 CTCTGAGGGCCACTAGTGC GGTTACCAGCGGAAATGCCTCGGGGT CAGAAGTCG CAGGAGAGATAGACAGCTGCTGAACCAATGGGACCAGCGGATGGG
101 GCGGATGTTATCTACCATTGGTGAACGTTAGAAACGAATAGCAGCCAATGAATCAGCTGGGGGGCGGAGCAGTGACGTTTATTGCGGAGGGGGCCGCTT
201 CGAATCGGCGGGCGCCAGCTTGGTGGCCTGGGCCAATGAACGGCCTCCAACGAGCAGGGCCTTACCAATCGGGCGCCTCCACGACGGGGCTGGGGAGG
301 GTATATAAGCCGAGTAGGGCAGCGTGAGGTGCGACGCCGGCCAAGACAGCACAGACAGATTGACCTATTGGGTGTTTTCCGGAGTGTGAGAGGGAAGCCGC
401 GCGGCCTGTATTCTAGACCTGCCTTCGCCTGGTTCGTGGCGCCTTGTGACCCCGGGCCCTGCGGCCTGCAAGTCGGAATTGCGCTGTGCTCCTGTG

NeoI (548)

501 CTACGGCCTGTGGCTGGACTGCCTGCTGCTGCCAACTGGCTGGCACCATGGGGGGTCTCATCATCATCATCATCATGGTATGGCTAGCATGACTGGT
601 GACAGCAAATGGGTGGGATCTGTACGACGATGACGATAAGGTAACCTAAGGATCAGCTTGGAGTTGATCCCGTCGTTTTACAACGTCGTGACTGGGAAAA
181 yGlnGlnMetGlyArgAspLeuTyrAspAspAspAspLysValProLysAspGlnLeuGlyValAspProValValLeuHisGlnArgAspTrpGluAs
701 CCCTGGCGTTACCCAACCTAATCGCCTTGACGACATCCCCCTTTCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCAGTCCGCCCTTCCCAACAGTTG
511 nProGlyValThrGlnLeuAsnArgLeuAlaAlaHisProProPheAlaSerTrpArgAsnSerGluGluAlaArgThrAspArgProSerGlnGlnLeu
801 CGCAGCCTGAATGGCGAATGGCGCTTTGCCTGGTTCGCCGCCAGAGCGGTGCGCGAAAGCTGGCTGGAGTGCATCTTCCCTGAGGCCGATACTGTCC
851 ArgSerLeuAsnGlyGluTrpArgPheAlaTrpPheProAlaProGluAlaValProGluSerTrpLeuGluCysAspLeuProGluAlaAspThrValV
901 TCGTCCCCTCAAACCTGGCAGATGCACGGTTACGATGCCCCATCTACACCAACGTAACCTATCCATTACGGTCAATCGCGCGCTTCCACCGGAA
1181 alValProSerAsnTrpGlnMetHisGlyTyrAspAlaProl leTyrThrAsnValThrTyrProl leThrValAsnProProPheValProThrGluAs
1001 TCCGACGGGTTGTTACTCGCTCACATTTAATGTTGATGAAAGCTGGTACAGGAAGGCCAGACGCGAATTATTTTGTATGGCGTTAACTCGCGGTTTCAT
1511 nProThrGlyCysTyrSerLeuThrPheAsnValAspGluSerTrpLeuGlnGluGlyGlnThrArgI leI lePheAspGlyValAsnSerAlaPheHis
1101 CTGTGGTCAACGGCGCTGGTGGTTCAGGCCAGGACAGTCTGTTGCCGCTGAATTTGACCTGAGCGCATTTTTACGCGCGGAGAAAAACCGCTCG
1851 leuTrpCysAsnGlyArgTrpValGlyTrpGlyGlnAspSerLeuProSerGluPheAspLeuSerAlaPheLeuArgAlaGlyGluAlaValProLysLeuA
1201 CGGTGATGGTGTGCTGGTGGAGTGACGGCAGTTATCTGGAAGATCAGGATATGTGGCGGATGAGCGGCATTTTCCGTGACGTCTCGTTGCTGCATAAAC
2181 laValMetValLeuArgTrpSerAspGlySerTyrLeuGluAspGlnAspMetTrpArgMetSerGlyI lePheArgAspValSerLeuLeuHisLysPr
1301 GACTACACAAATCAGCGATTTCCATGTTGCCACTCGCTTAAATGATGATTTACGCCGCGCTGTACTGGAGGCTGAAGTTCAGATGTGCGCGGAGTTGCGT
2511 oThrThrGlnI leSerAspPheHisValAlaThrArgPheAsnAspAspPheSerArgAlaValLeuGluAlaGluValGlnMetCysGlyGluLeuArg
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1701 CGAGGCGTTAACCGTCACGAGCATCATCTCTGCATGGTCAGGTCTAGGTGATGAGCAGACGATGGTGCAGGATATCTGCTGATGAAGCAGAACAACTTTA
3851 ArgGlyValAsnArgHisGluHisHisProLeuHisGlyGlnValMetAspGluGlnThrMetValGlnAspI leLeuLeuMetLysGlnAsnAsnPheA
1801 ACGCCGTGGCTGTTCCGATTTCCGAACCATCCGCTGTTGGTACACGCTGTGCGACCGCTACGGCCTGATGTGGTGGATGAAGCCAATATTGAAACCCA
4181 snAlaValArgCysSerHisTyrProAsnHisValLeuTrpTyrThrLeuGlyCysAspArgTyrGlyLeuTyrValValGluAlaAsnI leGluThrHi
1901 CGCATGGTGCCAAATGAATCGTCTGACCGATGATCCGCGCTGGCTACCGCGCATGAGCGAAGCCGCTAACCGGAATGGTGCAGCGCATCGTAATCACCCG
4511 sGlyMetValProMetAsnArgLeuThrAspAspProArgTrpLeuProAlaMetSerGluArgValThrArgMetValGlnArgAspArgAsnHisPro
2001 AGTGTGATCATCTGGTGGTGGGAATGAATCAGGCCAGGGCCTAATCAGCAGCGCTGTATCGCTGGATCAAATCTGTGATCCTCCCGCCCGTGC
4851 SerValI leI leTrpSerLeuGlyAsnGluSerGlyHisGlyAlaAsnHisAspAlaLeuTyrArgTrpI leLysSerValAspProSerArgProValG
2101 AGTATGAAGGGCGGAGCCGACACCAGGCCACCGATATTTTCCCGCATGTACGCGCGCTGGATGAAGACCAGCCCTTCCCGCTGTGCCGAAATG
5181 InTyrGluGlyGlyGlyAlaAspThrAlaThrAspI leI leCysProMetTyrAlaArgValAspGluAspGlnProGlnGlyAlaValProLysTr
2201 GTCCATCAAAAAATGGCTTTCGCTACCTGGAGAGACGCGCCCGCTGATCCTTTGCGAATACGCCACGCGATGGGTAACAGTCTTGGCGGTTTCGCTAAA
5511 pSerI leLysLysTrpLeuSerLeuProGlyGluThrArgProLeuI leLeuCysGluTyrAlaHisAlaMetGlyAsnSerLeuGlyGlyPheAlaLys
2301 TACTGGCAGCGCTTTCGTCAGTATCCCGCTTACAGGGCGGCTTCTGCTGGGACTGGGTGGATCAGTCGCTGATTAATATGATGAAACCGCAACCCGT
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2401 GGTCCGCTTACCGCGGATTTTGGCGATACGCCGACGATGCCAGTCTGTATGAAACCGCTGGCTTTTCCGACCGCAGCCGATCCAGCGCTGAC
6181 rpSerAlaTyrGlyGlyAspPheGlyAspThrProAsnAspArgGlnPheCysMetAsnGlyLeuValPheAlaAspArgThrProHisProAlaLeuTh
2501 GGAGCAAACACCAGCAGCAGTTTTTCCAGTTCGTTTATCCGGGCAACCATCGAAGTGACCAGCGAATACCTGTCCGTATAGCGATAACGAGCTC
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2601 CTGCACTGGATGGTGGCGCTGGATGGTAAGCCGCTGGCAAGCGGTGAAGTGCCTCTGGATGTCGCTCCACAAGGTAACAGTTGATTGAAGTGCCTGAAC
6851 LeuHisTrpMetValAlaLeuAspGlyLysProLeuAlaSerGlyGluValProLeuAspValAlaProGlnGlyLysGlnLeuI leGluLeuProGluL
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3501 CGCGGCACTTCCAGTTCAACATCAGCCGCTACAGTCAACAGCACTGATGGAACACCGCATCGCCATCTGCTGCACGGGAGAAGGCACATGGCTGA
9851 ArgGlyAspPheGlnPheAsnI leSerArgTyrSerGlnGlnGlnLeuMetGluThrSerHisArgHisLeuLeuHisAlaGluGluGlyThrTrpLeuA

3601 ATATCGACGGTTTCCATATGGGGATTGGTGGCGACGACTCCTGGAGCCCGTCAGTATCGGGGAATTACAGCTGAGCGCCGGTCGCTACCATTACCAGTT
1018▶ snI leAspGlyPheHisMetGlyI leGlyGlyAspAspSerTrpSerProSerValSerAlaGluLeuGlnLeuSerAlaGlyArgTyrHisTyrGlnLe

EcoRI (3733)

3701 GGTCTGGTGTCAAAAATAATAATCTAGTCGAGAATTCGCTAGCTCGACATGATAAGATACATTGATGAGTTTGGACAAAACCACAAGTGAATGCAGTGAA
1051▶ uValTrpCysGlnLys•••

3801 AAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAA

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PacI (4014)

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4301 ATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCGTGCTTACCAGGTTGGACTCAAGACGATGTTACCGGATAAGGGC

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4801 GGCATAGTATAATACGACTCACTATAGGAGGGCCATCATGGCCAAGTTGACCAGTGTGCTCCAGTGTCTCACAGCCAGGGATGTGGCTGGAGCTGTTGA

4900 GTTCTGGACTGACAGGTTGGGGTCTCCAGAGATTTTGTGGAGGATGACTTTGCAGGTGTGGTCAGAGATGATGTCACCCTGTTTATCTCAGCAGTCCAG

5000 GACCAGGTGGTGCCTGACAACACCCTGGCTTGGGTGTGGGTGAGAGGACTGGATGAGCTGTATGCTGAGTGGAGTGAGGTGGTCTCCACCAACTTCAGGG

5100 ATGCCAGTGGCCCTGCCATGACAGAGATTGGAGAGCAGCCCTGGGGGAGAGAGTTTGCCTGAGAGACCCAGCAGGCAACTGTGTGCACTTTGTGGCAGA

PacI (5263)

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