



基本信息

原核抗性:	Amp
平台编号:	bio-109262
克隆菌株:	DH5a
培养条件:	37 度

质粒属性

质粒宿主:	昆虫细胞
质粒用途:	蛋白表达
片段类型:	

片段物种:

原核抗性:	Amp
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真核抗性:

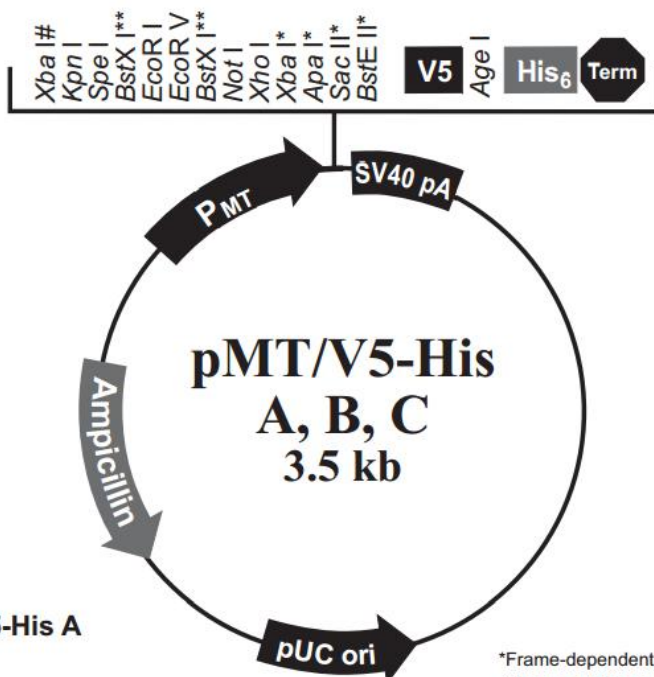
荧光标记:	
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功能简介

质粒简介

pMT/V5-His is a 3.5 kb expression vector designed for use with the Drosophila Inducible Expression System. Upon transfection, the vector allows transient, inducible expression of your protein of interest in Drosophila cells. When cotransfected with the selection vectors, pCoHygro or pCoBlast, included with the appropriate DES[®] Inducible Kit, pMT/V5-His allows selection of stable cell lines exhibiting inducible expression of the protein of interest.

质粒图谱



Comments for pMT/V5-His A

3538 nucleotides

Metallothionein promoter: bases 412-778

Start of transcription: base 778

MT Forward priming site: bases 814-831

Multiple cloning site: bases 854-923

V5 epitope tag: bases 930-971

Polyhistidine region: bases 981-1001

BGH Reverse priming site: bases 1021-1038

SV40 late polyadenylation signal: bases 1163-1168

pUC origin: bases 1601-2334 (complementary strand)

bla promoter: bases 3340-3438 (complementary strand)

Ampicillin (*bla*) resistance gene ORF: bases 2479-3339 (complementary strand)

*Frame-dependent variations

Xba I and *Apa* I are found only in versions A and B.

Sac II is found only in version B.

BstE II is found only in version C.

#This *Xba* I site is *dam* methylated.

**There are two *BstX* I sites in the polylinker.

411 5' end of metallothionein promoter
 CGTTGCAGGA CAGGATGTGG TGCCCGATGT GACTAGCTCT TTGCTGCAGG CCGTCCTATC
 Metal regulatory region
 471 CTCTGGTTCC GATAAGAGAC CCAGAACTCC GGCCCCCACC CGCCCACCGC CACCCCCATA
 Metal regulatory region
 531 CATATGTGGT ACGCAAGTAA GAGTGCCTGC GCATGCCCCA TGTGCCCCAC CAAGAGTTTT
 region
 591 GCATCCCATA CAAGTCCCCA AAGTGGAGAA CCGAACCAAT TCTTCGCGGG CAGAACAAAA
 Metal regulatory regions
 651 GCTTCTGCAC ACGTCTCCAC TCGAATTGAG AGCCGGCCCG CGTGTGCAAA AGAGGTGAAT
 Metal regulatory region
 711 CGAACGAAAG ACCCGTGTGT AAAGCCGCGT TTCCAAAATG TATAAAACCG AGAGCATCTG
 Metal regulatory regions
 771 GCCAATGTGC ATCAGTTGTG GTCAGCAGCA AAATCAAGTG AATCATCTCA GTGCAACTAA
 Start of transcription
 MT Forward priming site
 831 Xba I[#] Kpn I Spe I BstX I* EcoR I EcoR V
 AGGGGGGATC TAGATCGGGGT ACC TAC TAG TCC AGT GTG GTG GAA TTC TGC AGA TAT
 Thr Tyr *** Ser Ser Val Val Glu Phe Cys Arg Tyr
 BstX I* Not I Xho I Xba I Apa I BstB I V5 epitope
 888 CCA GCA CAG TGG CGG CCG CTC GAG TCT AGA GGG CCC TTC GAA GGT AAG CCT ATC
 Pro Ala Gln Trp Arg Pro Leu Glu Ser Arg Gly Pro Phe Glu Gly Lys Pro Ile
 Age I Polyhistidine region
 942 CCT AAC CCT CTC CTC GGT CTC GAT TCT ACG CGT ACC GGT CAT CAT CAC CAT CAC
 Pro Asn Pro Leu Leu Gly Leu Asp Ser Thr Arg Thr Gly His His His His His
 Pme I BGH Reverse priming site
 996 CAT TGA GTTTAAACC CGCTGATCAG CCTCGACTGT GCCTTCTAAG ATCCAGACAT GATAAGATAC
 His ***
 1061 ATTGATGAGT TTGGACAAAC CACAACCTAGA ATGCAGTGAA AAAAAAGCTT TATTTGTGAA
 SV40 late polyadenylation signal
 1121 ATTTGTGATG CTATTGCTTT ATTTGTAACC ATTATAAGCT GCAATAAACA AGTTAACAAC

This Xba I site is *dam* methylated in certain strains of *E. coli*.

* Note that there are two BstX I sites in the polylinker.

质粒序列

LOCUS	Exported	3538 bp ds-DNA	circular SYN
26-NOV-2017			
DEFINITION	synthetic circular DNA		
SOURCE	synthetic DNA construct		
ORGANISM	synthetic DNA construct		
REFERENCE	1 (bases 1 to 3538)		
TITLE	Direct Submission		
FEATURES	Location/Qualifiers		
source	1..3538		
	/organism="synthetic DNA construct"		



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	/label=MT promoter
	/note="Drosophila metallothionein promoter"
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	/label=AmpR
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HNMGDHSVTRLDRW	EPELNEAIPNDERDTTMPVAMATTLRKLLTGELLTLASRQQIDWM
EADKVAGPLLRSA	LPAGWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIYTTGSQATM
DERNRQIAEIGAS	LIKHW"



promoter complement (3340..3444)
/gene="bla"
/label=AmpR promoter

ORIGIN

```
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tcagcgggtg
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caaaaaggat



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