



IGEbio

广州艾基生物技术有限公司

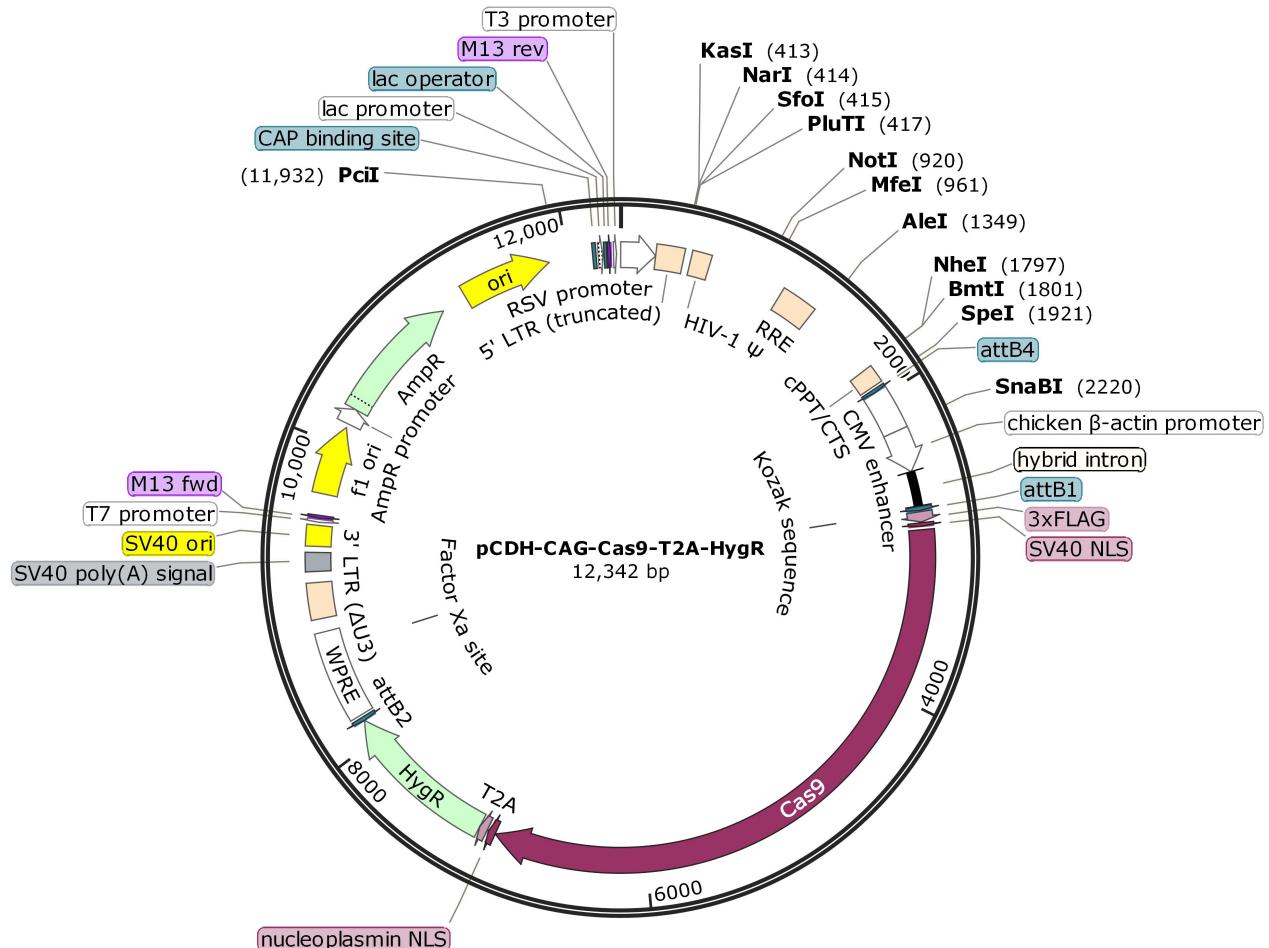
标准产业单元三期四栋 301 室

番禺区，国际生物岛

广州市，广东省

pCDH-CAG-Cas9-T2A-HygR Vector Information

Created with SnapGene®



载体名称: pCDH-CAG-Cas9-T2A-HygR

质粒类型: 慢病毒表达载体

表达水平: 低拷贝

启动子: CAG promoter

克隆方法: 多克隆位点, 限制性内切酶

克隆位点: MCS

载体大小: 12342bp

5' 测序引物及序列: --

3' 测序引物及序列: --

载体标签: 3xFLAG, SV40 NLS

载体抗性: Amp

筛选标记: HygR

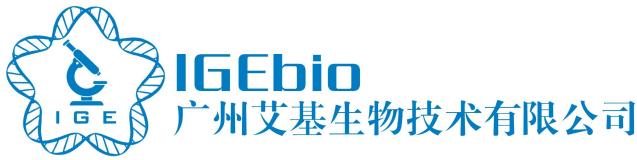
产品目录号: ----

稳定性: 稳定表达 Stable

组成型/诱导型: 组成型

病毒/非病毒: 慢病毒

克隆菌株: Stbl3



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LOCUS Exported 12342bp ds-DNA circular SYN 13-JUL-2018
DEFINITION pCDH-CAG-Cas9-T2A-HygR.
ACCESSION .
VERSION .
KEYWORDS pCDH-CAG-Cas9-T2A-HygR
SOURCE synthetic DNA construct
ORGANISM synthetic DNA construct
REFERENCE 1 (bases 1 to 12342)
AUTHORS .
TITLE Direct Submission
JOURNAL Exported Monday, June 24, 2019 from SnapGene 3.2.1
<http://www.snapgene.com>
FEATURES Location/Qualifiers
source 1..12342
/organism="synthetic DNA construct"
/mol_type="other DNA"
promoter 3..229
/note="RSV promoter"
/note="Rous sarcoma virus enhancer/promoter"
LTR 230..410
/note="5' LTR (truncated)"
/note="truncated 5' long terminal repeat (LTR) from HIV-1"
misc_feature 457..582
/note="HIV-1 Psi"
/note="packaging signal of human immunodeficiency virus type 1"
misc_feature 1075..1308
/note="RRE"
/note="The Rev response element (RRE) of HIV-1 allows for Rev-dependent mRNA export from the nucleus to the cytoplasm."
misc_feature 1803..1920
/note="cPPT/CTS"
/note="central polypyrimidine tract and central termination sequence of HIV-1"
protein_bind 1938..1958
/gene="mutant version of attB"
/bound_moiety="BP Clonase(TM)"
/note="attB4"
/note="core recombination site for the Gateway(R) BP reaction"
enhancer 1959..2244
/note="CMV enhancer"
/note="human cytomegalovirus immediate early enhancer;"

contains an 18-bp deletion relative to the standard CMV
 enhancer"
 promoter 2246..2523
 /note="chicken beta-actin promoter"
 intron 2524..2751
 /note="hybrid intron"
 /note="hybrid between chicken beta-actin (CBA) and minute
 virus of mice (MMV) introns (Gray et al., 2011)"
 protein_bind 2756..2780
 /gene="mutant version of attB"
 /bound_moiety="BP Clonase(TM)"
 /note="attB1"
 /note="recombination site for the Gateway(R) BP reaction"
 regulatory 2781..2790
 /regulatory_class="other"
 /note="Kozak sequence"
 /note="vertebrate consensus sequence for strong initiation
 of translation (Kozak, 1987)"
 CDS 2790..2855
 /codon_start=1
 /product="three tandem FLAG(R) epitope tags, followed by an
 enterokinase cleavage site"
 /note="3xFLAG"
 /translation="DYKDHDGDDYKDHDIDYKDDDK"
 CDS 2862..2882
 /codon_start=1
 /product="nuclear localization signal of SV40 (simian virus
 40) large T antigen"
 /note="SV40 NLS"
 /translation="PKKKRKV"
 CDS 2907..7007
 /codon_start=1
 /product="Cas9 (Csn1) endonuclease from the Streptococcus
 pyogenes Type II CRISPR/Cas system"
 /note="Cas9"
 /note="generates RNA-guided double strand breaks in DNA"
 /translation="DKKYSIGLDIGTNSVGWAVITDEYKVPSSKKFKVLGNTDRHSIKKN
 LIGALLFDSGETAATRLKRTARRRYTRRKNRICYLQEIFSNEAKVDDSFHRL
 ESFLVEEDKKHERHPIFGNIVDEVAYHEKYPTIYHLRKKLVDSTDKADLRLIYL
 LAHMIKFRGHFLIEGDLNPNSDVKLFIQLVQTYNQLFEENPINASGVDA
 KAILSARLSKSRRLENLIAQLPGEKKNGLFGNLIALSGLTPNFNSNFD
 LAEDAKLQLSKDTYDDDLDNLLAQIGDQYADLFLAAKNLSDA
 ILSILRVNTEITKAPLSASMIKRYDEHHQDLTLLKALVRQQLPE
 EKYKEIFFDQSNSKNGYAGYIDGGASQEEFYKFIKPILEKMDGTE
 ELLVKLNREDLLRKQRTFDNGSIPHQIHLGELHAILRRQEDFYPFL
 KDNREKIEKILTFRIPYYVGPLARGN"

SRFAWMTRKSEETITPNFEEVVDKGASAQSFIERNFTNFDKNLPNEKVLPHSLLYEYF
 TVYNELTKVKYVTEGMRKPAFLSGEQKKAIVDLLFKTNRKVTVKQLKEDYFKKIECFDS
 VEISGVEDRFNASLGTYHDLLKI KDKDFLDNEENEDILEDIVLTLTFEDREMIEERL
 KTYAHLFDDKVMQQLKRRRTGWRGRSLRKLINGIRDQKSGKTILDFLKSDGFANRNFMQ
 LIHDDSLTFKEDIQKAQVSGQGDSLHEHIANLAGSPAIIKGILQTVKVVDDELVKVMGRH
 KPENIVIEMARENQTTQKGQKNSRERMKRIEEGIKELGSQLKEHPVENTQLQNEKLYL
 YYLQNQGRDMYVDQELDINRLSDYDWDHVIPQSFLKDDSIDNKVLTRSDKNRGKSDNVPS
 EEVVKMKNYWRQLLNAKLTQRKFNDLTKAERGGLELDKAGFIKRQLVETRQITKHV
 AQILDLSRMNTKYDENDKLIREVKVITLKSCLVSDFRKDFQFYKVREINNYHHADAYLN
 AVVGTALIKKYPKLESEFVYGDYKVDVRKMIAKSEQEIGKATAKYFFYSNIMNFFKTE
 ITLANGEIRKRPLIETNGETGEIVWDKGRDFATVRKVLSMPQVNIVKKTEVQTGGFSKE
 SILPKRNSDKLIAKRDWDPKKGFFDSPTVAYSVLVAKVEKGSKKLKSVKELLGIT
 IMERSSFEKNPIDFLEAKGYKEVKKDLI IKLPKYSLELENGRKRMLASAGELQKGNEL
 ALPSKYVNFLYASHYEKLGSPEDNEQKQLFVEQHKHYDEIIEQISEFSKRVILADA
 NLDKVLSAYNKHRDKPIREQAENI IHLFTLTNLGAPAAFKYFDTTIDRKRYTSTKEVLD
 ATLIHQSQITGLYETRIDSQLGGD"

CDS	7008..7055	/codon_start=1 /product="bipartite nuclear localization signal from nucleoplasmin" /note="nucleoplasmin NLS" /translation="KRPAATKKAGQAKKK"
CDS	7065..7118	/codon_start=1 /product="2A peptide from Thosea asigna virus capsid protein" /note="T2A" /note="Eukaryotic ribosomes fail to insert a peptide bond between the Gly and Pro residues, yielding separate polypeptides." /translation="EGRGSLLTCGDVEENPGP"
CDS	7119..8144	/codon_start=1 /gene="aph(4)-Ia" /product="aminoglycoside phosphotransferase from E. coli" /note="HygR" /note="confers resistance to hygromycin" /translation="MKKPELTATSVEKFLIEKFDSVSDLMLSEGEESRAFSFDVGGRG YVLRVNSCADGFYKDRYVYRHFAAALPIPEVLDIGEFSESLTYCISRRAQGVTLQDLP ETELPAVLQPVAEAMDAIAADLSQTSGFGFPQGIGQYTTWRDFICAIADPHVYHWQ TVMDDTVSASVAQALDEMLWAEDCPEVRHLVADFGSNNVLTDNGRITAVIDWSEAMF GDSQYEVANIFFWRPWLAACMEQQTRYFERRHPELAGSPRLRAYMLRIGLDQLYQSLVDG NFDDAAWAQGRCDAIVRSGAGTVGRTQIARRSAAVTDGCVEVLADSGNRPSTRPRAK E"

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protein_bind complement(8145..8169)
/gene="mutant version of attB"
/bound_moiety="BP Clonase(TM)"
/note="attB2"
/note="recombination site for the Gateway(R) BP reaction"

misc_feature 8187..8775
/note="WPRE"
/note="woodchuck hepatitis virus posttranscriptional
regulatory element"

CDS complement(8658..8669)
/codon_start=1
/product="Factor Xa recognition and cleavage site"
/note="Factor Xa site"
/translation="IEGR"

LTR 8863..9096
/note="3' LTR (Delta-U3)"
/note="self-inactivating 3' long terminal repeat (LTR) from
HIV-1"

polyA_signal 9168..9289
/note="SV40 poly(A) signal"
/note="SV40 polyadenylation signal"

rep_origin 9329..9464
/note="SV40 ori"
/note="SV40 origin of replication"

promoter complement(9485..9503)
/note="T7 promoter"
/note="promoter for bacteriophage T7 RNA polymerase"

primer_bind complement(9513..9529)
/note="M13 fwd"
/note="common sequencing primer, one of multiple similar
variants"

rep_origin 9671..10126
/direction=RIGHT
/note="f1 ori"
/note="f1 bacteriophage origin of replication; arrow
indicates direction of (+) strand synthesis"

promoter 10152..10256
/gene="bla"
/note="AmpR promoter"

CDS 10257..11117
/codon_start=1
/gene="bla"
/product="beta-lactamase"
/note="AmpR"

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/note="confers resistance to ampicillin, carbenicillin, and
related antibiotics"
/translation="MSIQHFRVALIPFFAAFCLPVFAHPETLVKVKDAEDQLGARVGYI
ELDLNSGKILESRPPEERFPMMSTFKVLLCGAVLSRIDAGQEQLGRRIHYSQNDLVEYS
PVTEKHLTDGMTVRELCSAITMSDNTAANLLLTTIGGPKELTAFLHNMGDHVTRLDRW
EPELNEAIPNDERDTTMRVAMATTLRKLLTGELLTLASRQQLIDWMEADKVAGPLLRSA
LPAGWFIADKSGAGERGSRGIIAALGPDGKPSRIVVIYTTGSQATMDERNRQIAEIGAS
LIKHW"

rep_origin    11288..11876
/direction=RIGHT
/note="ori"
/note="high-copy-number ColE1/pMB1/pBR322/pUC origin of
replication"

protein_bind  12164..12185
/bound_moiety="E. coli catabolite activator protein"
/note="CAP binding site"
/note="CAP binding activates transcription in the presence
of cAMP."

promoter      12200..12230
/note="lac promoter"
/note="promoter for the E. coli lac operon"

protein_bind  12238..12254
/bound_moiety="lac repressor encoded by lacI"
/note="lac operator"
/note="The lac repressor binds to the lac operator to
inhibit transcription in E. coli. This inhibition can be
relieved by adding lactose or
isopropyl-beta-D-thiogalactopyranoside (IPTG)."

primer_bind   12262..12278
/note="M13 rev"
/note="common sequencing primer, one of multiple similar
variants"

promoter      12299..12317
/note="T3 promoter"
/note="promoter for bacteriophage T3 RNA polymerase"

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