

STOP

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[TECHNICAL SUPPORT](#)

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pNiFty3-A-Lucia

An AP-1-inducible secreted luciferase reporter plasmid selectable with Zeocin™

Catalog code: pnf3-lc3

For research use only

Version 20L03-MM

PRODUCT INFORMATION

Content:

- 20 µg of pNiFty3-A-Lucia provided as lyophilized DNA
- 1 ml of Zeocin™ (100 mg/ml)

Storage and stability:

- Products are shipped at room temperature.
- Store lyophilized DNA at -20 °C.
- Store Zeocin™ at 4 °C or at -20 °C. The expiry date is specified on the product label.

Quality control:

- Plasmid construct has been confirmed by restriction analysis and sequencing.

GENERAL PRODUCT USE

Pattern recognition receptor (PRR) activation triggers a complex signaling cascade that leads to the activation of different transcription factors, each playing an important role in the subsequent immune response. To monitor the induction of PRR signaling in response to ligand stimulation in a simple and efficient manner, InvivoGen has designed pNiFty, a family of reporter plasmids expressing a reporter gene under the control of a minimal promoter inducible by these different transcription factors, either individually or in combination. Most pNiFty plasmids are selectable with Zeocin™ in both *E. coli* and mammalian cells, and can be used to generate stable clones. pNiFty plasmids are composed of three key elements: a proximal promoter, repeated transcription factor binding sites (TFBS) and a reporter gene. The proximal promoters are shorter than 500 bp and contain transcription factor binding sites. Upon stimulation in 293 cells, their expression level remains undetectable. With the addition of repeated TFBS, the proximal promoters become inducible by the appropriate stimulus and drive the expression of the reporter gene.

PLASMID FEATURES

- **AP-1 binding site:** Activator protein 1 (AP-1) is a transcription factor activated by most PRRs. AP-1 is a heterodimeric complex composed of members of Fos, Jun and ATF protein families. AP-1 binds to the TPA responsive element (TRE: TGAG/CTCA)¹. AP-1 activation in TLR signaling is mostly mediated by MAP kinases such as c-Jun N-terminal kinase (JNK), p38 and extracellular signal regulated kinase (ERK).
- **IFN-β promoter:** the mouse IFN-β minimal promoter comprises several positive regulatory domains that bind different cooperating transcription factors such as NF-κB, IRF3 and IRF7².
- **Lucia luciferase** is a synthetic CpG-free gene that codes for a secreted coelenterazine-utilizing luciferase. ORF size (from ATG to stop codon): 634 bp. Lucia luciferase activity can be evaluated using QUANTI-Luc™ (cat. code: rep-qlc1), an assay reagent containing all the components required to quantitatively measure the activity of Lucia luciferase and other coelenterazine-utilizing luciferases.
- **SV40 pAn:** The Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA.

• **Ori** is a minimal *E. coli* origin of replication with the same activity as the longer Ori.

• **EF1/HTLV prom** is a composite promoter comprising the Elongation Factor-1α (EF-1α) core promoter³ and the R segment and part of the U5 sequence (R-U5') of the Human T-Cell Leukemia Virus (HTLV) Type 1 Long Terminal Repeat⁴. The EF-1α promoter exhibits a strong activity and yields long lasting expression of a transgene *in vivo*. The R-U5' has been coupled to the EF-1α core promoter to enhance stability of RNA.

• **EM7** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.

• **Zeo:** Resistance to the antibiotic Zeocin™ is conferred by the *Sh ble* gene from *Streptallocteichus hindustanus*. The *Sh ble* gene is driven by the EF1-HTLV promoter in tandem with the bacterial EM7 promoter allowing selection in both mammalian cells and *E. coli*.

• **βGlo pAn:** The human beta-globin 3'UTR and polyadenylation sequence allows efficient arrest of the transgene transcription⁵.

1. Hess J. et al., 2004. AP-1 subunits: quarrel and harmony among siblings. *J Cell Sci.* 117(25):5965-73. **2. Vodjani G. et al., 1988.** Structure and characterization of a murine chromosomal fragment containing the interferon beta gene. *J Mol Biol.* 204(2):221-31. **3. Kim D. et al., 1990.** Use of the human elongation factor 1 alpha promoter as a versatile and efficient expression system. *Gene* 91 (2): 217-223. **4. Takebe Y. et al., 1988.** SR alpha promoter: an efficient and versatile mammalian cDNA expression system composed of the simian virus 40 early promoter and the R-U5 segment of human T-cell leukemia virus type 1 long terminal repeat. *Mol. Cell Biol.* 1: 466-472. **5. Yu J. & Russell J., 2001.** Structural and functional analysis of an mRNP complex that mediates the high stability of human beta-globin mRNA. *Mol Cell Biol.* 21(17):5879-88.

METHODS

Plasmid resuspension

Quickly spin the tube containing the lyophilized plasmid to pellet the DNA. To obtain a plasmid solution at 1 µg/µl, resuspend the DNA in 20 µl of sterile H₂O. Store resuspended plasmid at -20 °C.

Plasmid amplification and cloning

Plasmid amplification and cloning can be performed in *E. coli* GT116 or in other commonly used laboratory *E. coli* strains, such as DH5α.

Zeocin™ usage

This antibiotic can be used for *E. coli* at 25 µg/ml in liquid or solid media and at 50-200 µg/ml to select Zeocin™-resistant mammalian cells.

TECHNICAL SUPPORT

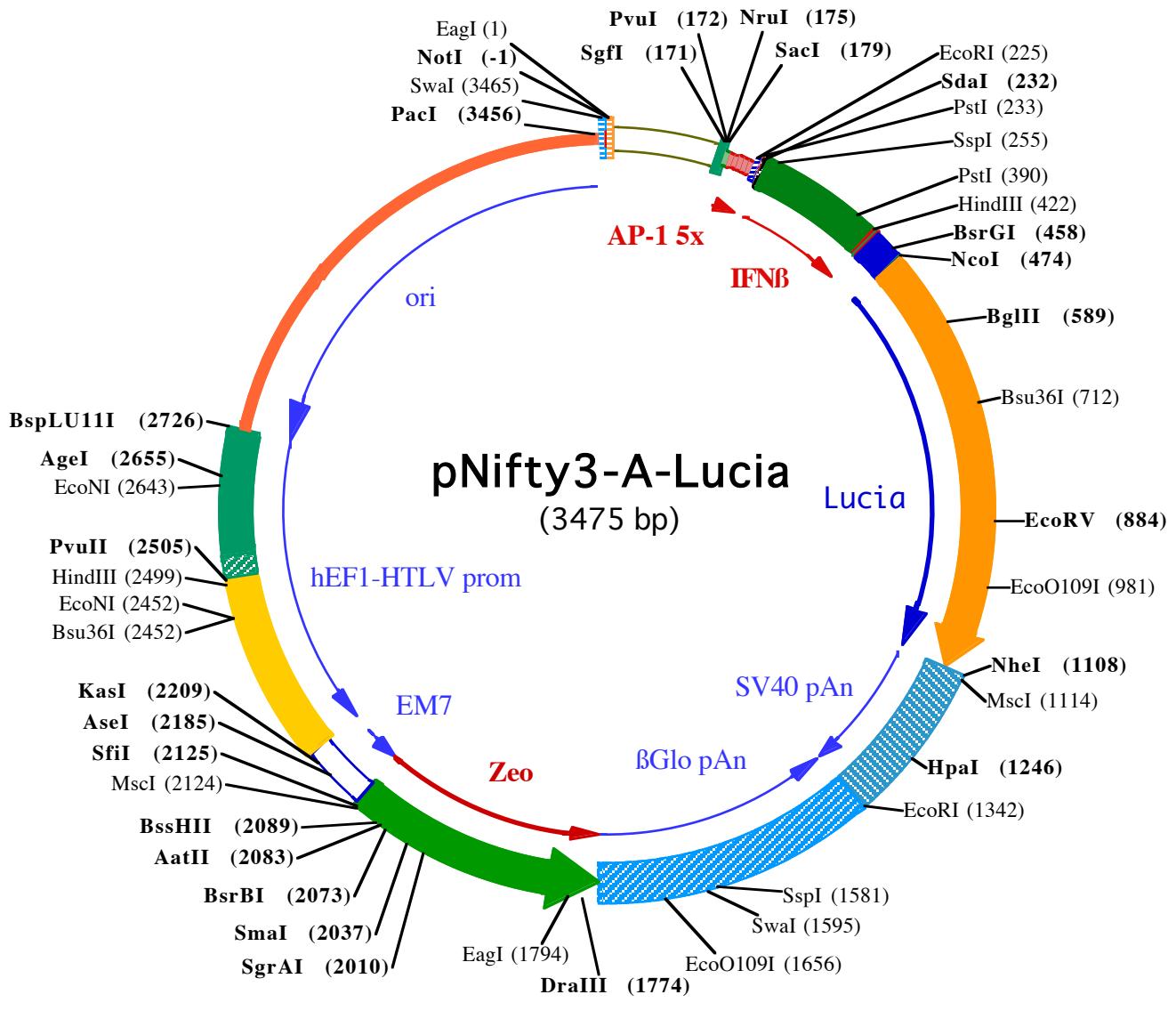
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EagI (1)
NotI (-1)
1 **GCGGCCGAATAAAATATCTTATTTCATTACATCTGTGTTGGTTTGTAATCGTAACATACGCTCTCCATAA**ACAAAACAAAACGAAACA
SacI (179)
NruI (175)
PvuI (172)
SgfII (171)
101 **AAACAAACTAGCAAAATAGGCTGCCAGTCAGGTGCCAGAACATTCTATCGAAGGATCTGCAGCTGAGTC**ACTGACTCA

SdaI (232)
EcoRI (225) PstI (233)
201 **GTGAGTCACTGACTCAGTGAGTAAGGAATTCTCCTGCAGGagcttgaataaaatgaatattagaagctgttagaataagagaaaatgacagaggaAAACT**

PstI (390)
301 **GAAAGGgAGAACTGAAAGTGGaaattccctcgaggcagaaaggaccatccctTATAAAtagcacaggccatgaaggaaagtcattctca**tgcagcc

HindIII (422) BsrGI (458) NcoI (474)
399 **tttgacagccttgccatcttggAACTTCTGCCTTCTCCCTCCTGTGAGTTGGTTGGTACAGTAGCTCCACCATGAAATCAAGGTGCTGTTG**1►MetGl uIleLysValLeuPheA
BglIII (589)
499 **CCCTCATCTGATTGCTGTTGCTGAGGCCAAACCCACTGAAATCAATGAAAGACCTCAATATAGCTGCTGCCCCTCCAAC**TGAAATCAAGGTGCTGTTG
8►laLeuIleCysIleAlaValAlaGl uAl aLysProThrGl uIleAsnGl uAspLeuAsnIleAlaAlaValAlaSerAsnPheAlaThrThrAspLeuGl
599 **GACTGACCTGTTCACCAACTGGGAGACCATGAATGTGATTAGCACTGACACAGAGCAGGTGAAACACAGATGTCGACAGGGCAAGCTGCTGGCAAAAAA**
41►uThrAspLeuPheThrAsnTrpGl uThr MetAsnVal IleSerThrAspThrGl uGl nValAsnThrAspAlaAspArgGl yLysLeuProGl yLysLys
BsU36I (712)
699 **CTCCCCCCAGATGTCCTGAGGGAGCTGGAGGCCATGCCAGAAGGGCTGGTGACAAGAGGCTGCTCATTGCTCTCCACATTAAGTCACCCCTA**
75►LeuProProAspVal LeuArgGl uLeuGl uAl aAsnAl aArgArgAl aGl yCysThr ArgGl yCysLeuIleCysLeuSerHis IleLysCysThrProL
EcoRV (884)
799 **AGATGAAGAAATTATCCTGGCAGGTGCCACACTTATGAAGGTGAAAGGAGCTGCTCAGGGAGGGATTGGAGAGGCAATTGTTGATCCCAGAGAT**
108►ysMetLysLysPhel IleProGl yArgCysHisThr TyrGl uGl yGl uLysGl uSerAl aGl nGl yGl yGl uAl aIleValAspIleProGl uI
EcoO109I (981)
899 **TCCTGGCTCAAGGATAAGGAGCCACTGGACAGTTATTGCTCAAGTGGACCTCTGTGCTGATTGACCACACTGGCTGTGAAGGGCCTGCAATGTC**
141►eProGl yPheLysAspLysGl uProLeuAspGl nPhel IleAl aGl nValAspLeuCysAl aAspCysThrThrGl yCysLeuLysGl yLeuAl aAsnVal
999 **CAGTGCCTGACCTCTGAAGAAGTGGCTTCCCAGAGGTGTACCACTTTGCCAGAACATTGAGGTGGACAAATCAAGGGTCTGGCTGGGG**
175►Gl nCysSerAspLeuLeuLysTrpLeuProGl nArgCysThrThrPheAlaSerLysIleGl nGl yArgValAspLysIleLysGl yLeuAl aGl yA
Mscl (1114)
NheI (1108)
1099 **ACAGATGATACTAGCTGGCCAGACATGATAAGATAACATTGATGAGTTGGACAAACCAACTAGAACATGCACTGAAAAAAATGTTATTGTGAAATT**
208►spArg***

HpaI (1246)
1199 **TGTGATGCTATTGCTTATTGTAACCATTATAAGCTGCAATAAACAAGTTAACACAACATTGATTCAATTGATTTATGTTCAGGTTAGGGGAGGTGT**

EcoRI (1342)
1299 **GGGAGGTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTCTAAACATAGCATA**GGAAACTTAACTCCAAATCAAGCCTACTTGA

1399 **ATCCTTTCTGAGGGATGAATAAGGCATAGGCATCAGGGCTGTCATAGCTGTTGCAGCCTCACCTTTCATGGAGTTAAGATAT**

SspI (1581) SwaI (1595)
1499 **AGTGTATTTCAGGGTAAACTAGCTCTCATTCTTATGTTAAATGCACACTGACCTCCACATTCCCTTTAGTAAATTCAGAAATAATT**

EcoO109I (1656)
1599 **TAAATACATCATTGCAATGAAATAATGTTTTATTAGGCAGATGTCAGGCCCTCATATACTCCCCAGTTAGTAGTTGACTTAGG**

DraIII (1774) EagI" (1794)
1699 **GAACAAAGGAACCTTAATAGAAATTGGACAGCAAGAAAGCGAGCTTAGCTTATCCTCAGTCCTGCTCTGCCACAAAGTCACGCCAGTTGCC**
127►•••Gl y•••AspGl nGl uGl uAl aVal PheHi sVal CysAsnGl yAl a

1799 **CGGGTCGCGAGGGCGAACCTCCGCCACGGCTGCTCGCCGATCGGTCATGGCGCCGGAGGCCTCCGGAA**GATTCGTGGACACGACCTCGAC
111►ProAspArgLeuAl aPheGl uArgGl yTrpRoGl nGl uGl yIleGl uThr MetAlaProGl ySerAl aAspArgPheAsnThrSerValValGl uSerT
1899 **CACTCGCGTACAGCTCGCAGGCCGACCCCACAGCCAGGGTGTGCGACCCACTGGCTGGACCGCGCTGATGAACAGGGTCACGT**
77►rpGl uAl aTyrLeuGl uAspLeuGl yArgVal TrpValItrpAl aLeuThrAspProVal Val Gl nAspGl nValAl aSerIlePheLeuThrValAs

AatII (2083)
SgrAI (2010) SmaI (2037) BsrBI (2073) BssHII (2089)
1999 **CGTCCGGACACCCGGGAAGTCGTCTCCACGAAGTCCGGAGAACCGAGCCGGTCGGTCCAGACTCGACCGCTGGCGACGTGCGCGCGT**
44►pAspArgVal Val Gl yAl aPheAspAspGl uVal PheAspArgSer PheGl yLeuArgAspThr TrpPheGl uValAl aGl yAl aValAspArgAl aThr

SfiI (2125) MscI (2124) AscI (2185)
2099 **GAGCACCGGAACGGCACTGGCACTTGGCCATGATGCCCTCCTATAGTGAGTCGTATTATACTATGCCGATATACTATGCCGATGATTAAATTGCAAC**
11►LeuVal ProValAl aSer Thr LeuLysAl aMet

KasI (2209)
2199 **TACTGTTGTAGGCAGCGGTACAGCTTGATCTGTAACGGCGCAGAACAGAAAAGAACAGTAGAGTTGAGCAAGCAGGGTCAGGCAAAGCGT**

EcoNI (2452) Bsu36I (2452)
2299 **GGAGAGCCGGCTGAGTCTAGGTAGGCTCCAGGGAGCGCCGGACAAAGGCCGGTCTGACCTGAGCTTAACTTACCTAGACGGCGACGCTAG**

2399 **GAGGCACCACAGGCCGGAGCGGGCAGAACGCGACTCAACCGCGTGGATGGCGCCCTCAGGTAGGGCGCCGGCGTGAAGGAGAGATGCGAGCCCCTC**

PvuII (2505)
HindIII (2499)
2499 **GAAGCTTCAGCTGTGTTCTGGCGAAACCGTTGCAGAAAGAACGTTACGGCGACTACTGCACTTATACGGTCTCCCCACCCCTGGAAAAAG**

EcoNI (2643) AgeI (2655)

2599 **GCGGAGCCAGTACACGACATCACTTCCCAGTTACCCGCCACCTCTAGGCACCGTTCAATTGCCACCCCTCCCCCAACTTCGGGACT**

2699 **BspLU11I (2726)**
2699 **GTGGGCGATGTGCGCTCTGCCACTGACACATGTGAGCAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAGGCCGTTGCTGGCTTTCCATAGGC**

2799 **TCCGCCCCCTGACGAGCATCACAAAAATGACGCTCAAGTCAGAGTGGCAGAACCCGACAGGACTATAAGATAACCAGCGTTCCCCCTGGAAGCTC**

2899 **CCTCGTGCCTCTCGTCCGACCCCTGCCGTTACCGGATACCTGTCGCCCTTCTCCCTCGGAAGCGTGGCTTCTCATAGCTCACGCTGTAGG**

2999 **TATCTCAGTTGGTGTAGTCGTCGCTCCAAGCTGGCTGTGACGAACCCCCGTTAGCCGACCGCTGCCCTATCCGTAACATCGTCTTG**

3099 **AGTCCAACCCGTAAGACACGACTTATGCCACTGGCAGCCACTGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTGAA**

3199 **GTGGTGGCTAAGTACGGCTACACTAGAAGAACAGTATTGGTATCTGCCTGCTGAAGCCAGTTACCTCGAAAAAGAGTTGGTAGCTCTGATCC**

3299 **GGCAAACAAACCAACCGCTGGTAGCGGTGGTTTGCAAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTGATCTTCTA**

SwaI (3465)

PacI (3456)

3399 **CGGGGTCTGACGCTCAGTGGAACGAAACTCACGTTAAGGGATTTGGTATGGCTTAATTAA**CATTTAAATCA****
