

pNiFty3-AN-SEAP

An inducible reporter plasmid selectable with Zeocin™

Catalog code: pnf3-sp6

For research use only

Version 20L03-MM

PRODUCT INFORMATION

Content:

- 20 µg of pNiFty3-AN-SEAP 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).

• NF-κB binding site: Nuclear factor (NF)-κB is a “rapid-acting” primary transcription factor activated by a wide variety of PRRs. NF-κB is a protein complex that belongs to the Rel-homology domain-containing protein family. The prototypical NF-κB is composed of the p65(RelA) and p50 subunits². NF-κB binds specific decameric DNA sequences (GGGRNNYYCC, R-purine Y=pyrimidine) and activates genes involved in the regulation of the innate and adaptative immune response.

• 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 IRF7³.

• SEAP is a secreted form of human embryonic alkaline phosphatase. Unlike endogenous alkaline phosphatases, SEAP is extremely heat stable and resistant to the inhibitor L-homoarginine. It catalyses the hydrolysis of pNitrophenyl phosphate (pNpp) producing a yellow end product. SEAP expression can be readily quantified by collecting samples of culture medium and measuring the hydrolysis of pNpp with a spectrophotometer at 405 nm.

• 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 promoter⁴ and the R segment and part of the U5 sequence (R-U5') of the Human T-Cell Leukemia Virus (HTLV) Type 1 Long Terminal Repeat⁵. 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 *Streptomyces 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 transcription⁶.

References

1. Hess J, et al., 2004. AP-1 subunits: quarrel and harmony among siblings. *J Cell Sci.* 117(Pt 25):5965-73.
2. Kawai T. & Akira S., 2007. Signaling to NF-kappaB by Toll-like receptors. *Trends Mol Med.* 13(11):460-9.
3. 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.
4. Kim D. et al., 1990. Use of the human elongation factor 1α promoter as a versatile and efficient expression system. *Gene* 91(2): 217-23.
5. 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: 466-72.
6. Yu J & Russell J., 2001. Structural and functional analysis of an mRNP complex that mediates the high stability of human β-globin mRNA. *Mol Cell Biol.* 21(17):5879-88.

TECHNICAL SUPPORT

InvivoGen USA (Toll-Free): 888-457-5873

InvivoGen USA (International): +1 (858) 457-5873

InvivoGen Europe: +33 (0) 5-62-71-69-39

InvivoGen Hong Kong : +852 3-622-34-80

E-mail: info@invivogen.com

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.

RELATED PRODUCTS

Product	Catalog Code
ChemiComp GT116	gt116-11
Zeocin™	ant-zn-1

TECHNICAL SUPPORT

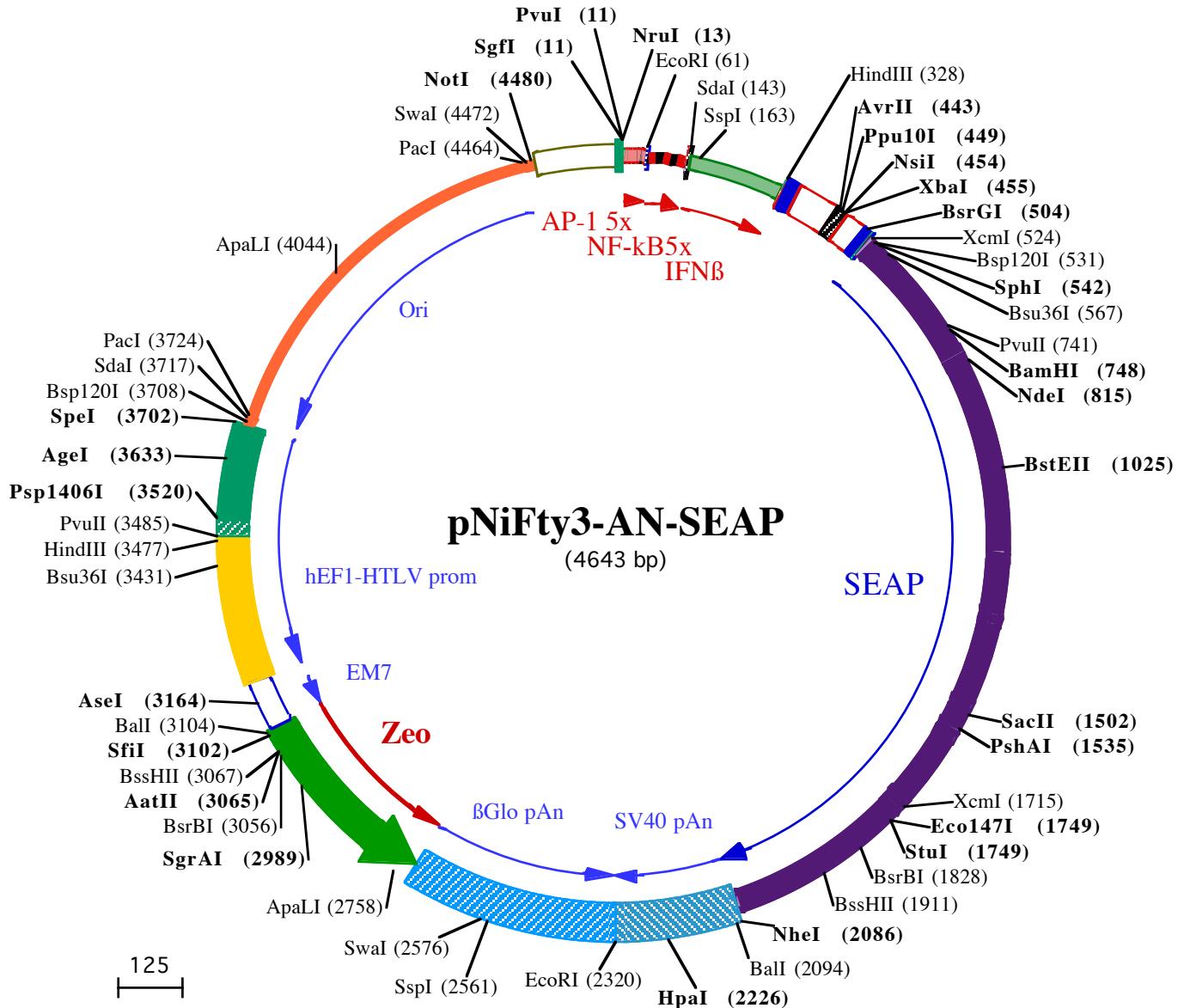
InvivoGen USA (Toll-Free): 888-457-5873

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InvivoGen Europe: +33 (0) 5-62-71-69-39

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E-mail: info@invivogen.com



NruI (13)
PvuI (11)
SgfI (11)

EcoRI (61)

1 GGATTCGATCGCAGCTGAGTCAGTGACTCACTGAGTAAAGAATTCTGGGACTTCCACTGGGACTTTCACTGGGA

101 CTTTCACTGGGACTTTCACTGGGACTTCCACTGGGACTTCCACTcTGAGGagcttgaataaaatgaatattagaagctgttagaataagagaaaatgacagagga

201 AACTGAAAGggAGAACTGAAAGTggaaattccctgaggcagaaaggaccatccctTATAAAtagcacaggccatgaaggaagatcatttcactg

299 cagccttgacagccttgccatcttgAGCTCTGCCTCTCCCTGTGAGTTGtaagtcaactgactgtctatgcctggaaagggtggcag

HindIII (328)

Ppu10I (449)
NsiI (454)
AvrII (443) **XbaI (455)**
SphI (542)

399 gagatggggcagtgcaggaaaagtggcaatatgaacccTGACGCCCTAGGAATGCATCTAGAcattgtactaacctttctcttcctcgtacag

BsrGI (504) **XcmI (524)** **Bsp120I (531)** **Bsu36I (567)**

499 GTTGGTGTACAGTAGCTTCCACCATGATTCTGGGCCCTGCATGCTGCTGCTGCTGCTGCTGGCCCTGAGGCTACAGCTCTCCCTGGGCATCATCCC
1► M I L G P C M L L L L L G L R L Q L S L G I I P
599 AGTTGAGGAGGAGAACCCGACTTCTGAAACCGCGAGGCCGAGGCCGAGGCCCTGGGTGCGCAAGAAGCTGCAGCCTGCACAGACAGCCGCAAGAACCTC
26► V E E E N P D F W N R E A A E A L G A A K K L Q P A Q T A A K N L

PvuII (741) **BamHI (748)**

699 ATCATCTTCTGGCGATGGGATGGGGTGTCTACGGTGACAGCTGCCAGGATCTAAAAGGGCAGAAGAAGGACAAACTGGGCTGAGATAACCCCTGG
60► I I F L G D G M G V S T V T A A R I L K G Q K K D K L G P E I P L

NdeI (815)

799 CTATGGACCGCTTCCATATGGCTCTGTCCAAGACATACAATGTAGACAAACATGTGCCAGACAGTGGAGCCACAGCCACGGCTACCTGTGCGGGT
93► A M D R F P Y V A L S K T Y N V D K H V P D S G A T A T A Y L C G V
899 CAAGGCAACTCCAGACCATTGGCTTGGAGTCAGCCGCCGCTTAACCAAGTCAACACGACAGCGCGAACGAGGTCTCGTGTGAATCGGGCC
126► K G N F Q T I G L S A A A R F N Q C N T T R G N E V I S V M N R A

BstEII (1025)

999 AAGAAAGCAGGGAAGTCAGTGGAGTGGTAACCAACACAGAGTCAGCACGCCCTGCCAGCCGGCACCTACGGTAACCGCAACTGGTACT
160► K K A G K S V G V T T T R V Q H A S P A G T Y A H T V N R N W Y
1099 CGGACGCCGACGTGCCTGCCTCGGCCAGGGGGTGCAGGACATCGCTACGCAGCTCATCTCCACATGGACATTGATGTGATCTGGTGGAGG
193► S D A D V P A S A R Q E G C Q D I A T Q L I S N M D I D V I L G G G
1199 CGCAAAGTACATGTTCGATGGAAACCCAGACCTGAGTACCCAGATGACTACAGCAAGGTGGGACCAGGCTGGACGGGAAGAATCTGGTGCAGGAA
226► R K Y M F R M G T P D P E Y P D D Y S Q G G T R L D G K N L V Q E
1299 TGGCTGGCGAACGCCAGGGCTTGGAGTCAGCCGCCGCTTAACCAAGTCAACACGACAGCGCGAACGAGGTCTCGTGTGACCCATCTCATGGGTCTTTG
260► W L A K R Q G A R Y V W N R T E L M Q A S L D P S V T H L M G L F
1399 AGCCGGAGACATGAAATCAGAGATCCACCGAGACTCCACACTGGACCCCTCCCTGTGGAGATGACAGAGGTGCCCTGCCCTGCTGAGCAGGAACCC
293► P G D M K Y E I H R D S T L D P S L M E M T E A A L R L L S R N P

SacII (1502) **PshAI (1535)**

1499 CCGCGGCTCTCCCTCTCGGGAGGGGGTCATCGACCGTCATCACGAAAGCAGGGCTTACCGGGCACTGACTGAGACGATCATGTTGCAGCAGAC
326► R G F F L F V E G G R I D H G H H E S R A Y R A L T E T I M F D D
1599 GCCATTGAGAGGGCGGGCAGCTACCCAGCGAGGACAGCTGAGCCTCGTCACTGCCGACCACTCCACGTCTCTCCCTGGAGGCTACCCCTGC
360► A I E R A G Q L T S E E D T L S L V T A D H S H V F S F G G Y P L

XcmI (1715)

1699 GAGGGAGCTCCATCTCGGGCTGGCCCCCTGCCAAGGCCCCGGAGAGGAAGGCTACACGGTCTCTATACGGAAACGGTCCAGGCTATGTGCTCAAGGA
393► R G S S I F G L A P G K A R D R K A Y T V L L Y G N G P G Y V L K D

BsrBI (1828)

1799 CGGCGCCCGCCGGATGTTACCGAGAGCGAGAGCGGGAGCCCGAGATCGGCAGCAGCTAGCAGTGGCCCTGGACGAAGAGACCCACGCAGGGCGAGGAC
426► G A R P D V T E S E S G S P E Y R Q Q S A V P L D E E T H A G E D

BssHII (1911)

1899 GTGGCGGTGTTCGCGCGGGCCGAGGCGCACCTGGTCACGGCGCAGGAGCAGACCTTCATAGCGCACGTCACTGGCTTCCCTGGAGGCTACCCCTG
460► V A V F A R G P Q A H L V H G V Q E Q T F I A H V M A F A A C L E

Ball (2094)
NheI (2086)

1999 CCTACACCGCTCGACCTGGCCCCCGCCGGACACCGACGCCGCACCCGGGGCGTCCCGTCAAGCGTCTGGATTGAAAGCTAGCTGGCAG
493► P Y T A C D L A P P A G T T D A A H P G R S R S K R L D •
2099 ACATGATAAGATACATTGATGAGTTGGACAAACCACAATAGAATGCACTGAAATGCTTATTGTGAAATTGATGCTATTGCTTTATTGT

HpaI (2226)

2199 AACCAATTAAAGCTCAATAAACAGTTAACACAATTGCAATTCTATTATGTTAGTCAGGGGAGGTGTGGAGGTTAAAGCAAGTAA

EcoRI (2320)

2299 AACCTCTACAAATGTTGATGAAATTCTAAATACAGCATAGCAAACCTTAACCTCAAATCAAGCCTACTTGAATCTTCTGAGGGATGAATAA

2399 GGCGATAGGCATCAGGGCTTGCACATGTCATTAGCTGTTGCAGCCTCACCTCTTCACTGGAGTTAAGATATAGTGTATTCTTCAAGGTTGAA

SspI (2561) **SwaI (2576)**

2499 CTAGCTTCATTTCTTATGTTAAATGCACTGACCTCCCACATTCCCTTTAGTAAATTCAGAATAATTAAACATCATTGCAATGAA

2599 TAAATGTTTATTAGGCAGATCCAGATGCTCAAGGCCCTCATAATATCCCCAGTTAGTTAGTGGACTAGGAAACAAGAACCTTAATAGAA

ApaLI (2758)

2699 ATTGGACAGCAAGAAAGCGAGCTTCTAGCTTATCCTCAGTCCTGCTCCCTGCCACAAAGTGACGCAGTCAGTTGCCGGCGGGTCGCAGGGCGAACTCCC
1271 • G • D Q E E A V F H V C N G A P D R L A F E R

2799 GCCCCCACGGCTGCTCGCGATCTCGGTATGGCCGGCCGGAGGGTCCCGGAAGTCGTGGACACGACCTCCGACCACCTGGCTACAGCTGTCCAG
103 G W P Q E G I E T M A P G S A D R F N T S V V E S W E A Y L E D L
SgrAI (2989)
2899 GCGCGCACCCACACCCAGGCCAGGGTGTGTCGGCACCCACCTGGCTTGACCGCGCTGATGAACAGGGTACGTGTCGCCGGACACACCGCGAAG
70 G R V W V W A L T N D P V V Q D Q V A S I F L T V D D R V V G A F

BssHII (3067)

BsrBI (3056) AatII (3065)

2999 TCGTCCTCCACGAAGTCCCGGAGAACCCGAGCCGGTCCAGAACTCGACCGCTCCGGCACGTCGCGCGGTGAGCACCGAACGGCACTGGTCA
36 D D E V F D R S F G L R D T W F E V A G A V D R A T L V P V A S T L
SfiI (3102)
Ball (3104)
3099 ACTTGGCCATGATGGCCCTCTATAGTGAAGTCGTATTATACTATGCCGATATACTATGCCGATTAATTGTCAACTACTGTTGTAGGGCGCCGTAC
3 K A M ←
3199 AGCTTGATCTGTAACGGCGAGAACAGAAAAGCAAAGACGTAGAGTTGAGCAAGCAGGGTCAGGCAAAGCGTGGAGAGCCGGCTGAGTCTAGGTA
←
3299 GGCTCCAAGGGAGCGCCGGACAAAGGCCGGTCTGACCTGAGCTTAAACTACCTAGACGGGGACGCAGTCAGGAGGCCACAGGGGGAGGGGG

PvuII (3485)

Bsu36I (3431) HindIII (3477)

3399 CAGAACCGACTCAACCGGTGGATGGGGCTCAGTAGGGCGGGCGGTGAAGGAGAGATGCGAGCCCCTGAAGCTCAGCTGTGTTCTGGCG

Psp1406I (3520)

3499 GCAAACCGTTGCAAAAAGAACGTTACGGCACTACTGCACTTATATACGTTCTCCCCACCCCTGGAAAAAGCGGAGCCAGTACACGACATCAC

AgeI (3633)

3599 TTCCCGAGTTACCCGCGCCACCTCTAGGCACCGGTTCAATTGCCGACCCCTCCCCAACCTCTGGGGACTGTGGCGATGTGCGCTCTGCCA

Bsp120I (3708)

SpeI (3702) SdI (3717) PacI (3724)

3699 CTGACTAGTGGCCCTGCAAGTTAATTAGAACATGTGAGCAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAGGCCGTTGCTGGCTTTCCATA
←
3799 GGCTCCGCCCCCTGACGAGCATCACAAATCGACGCTCAAGTCAGAGGTGGCAAACCCGACAGGACTATAAGATACCGCGTTCCCTGGAG

3899 CTCCCTCGTGCCTCTCTGTTCCGACCTGCGCTTACCGATACTGTCCGCTTCTCCCTGGAAAGCGTGGCGCTTCTCATAGCTACGCTG

ApaLI (4044)

3999 AGGTATCTCAGTCGGTAGGTCGTTCTGCTCCAAGCTGGCTGTGACGAACCCCCCGTTAGCCGACCGCTGCGCTTATCGGTAACTATCGT

4099 TTGAGTCCAACCCGTAAGACACGACTTATGCCACTGGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTTAGGGCTACAGAGTTCTT

4199 GAAGTGGTGGCTAACTACGGCTACACTAGAAGAACAGTATTGGTATCTGCGCTGCTGAAGCCAGTTACCTCGGAAAAAGAGTTGGTAGCTTGA

4299 TCCGGCAAACAAACCACCGTGGTAGCGGTGTTTTGTTGCAAGCAGCAGATTACCGCAGAAAAAAAGGATCTAAGAAGATCCTTGATCTTT

PacI (4464) SwaI (4472) NotI (4480)

4399 CTACGGGTCTGACGCTCAGTGAACGAAACTCACGTTAAGGGATTTGGTCAAGCTGTTAACTAACATACGCTCTCCATCAAACAAACGAAACAAACAAACTAGCAAATAGG

4499 TTATTTCTTACATCTGTGTTGGTTTTGTGTGAATCGTAACATACGCTCTCCATCAAACAAACGAAACAAACAAACTAGCAAATAGG

4599 CTGCCCCAGTGCAGTGCCAGAACATTCTATCGAA