

Plasmid Data Sheet

* **Plasmid Name and Size:** [pcDNA3.1/ZEO\(+\)-lnc-NDRG1-1-4-1; 5561 bp](#)

* developed at the BioMed Resource Core of the 1st Core Facility Lab, NTU-CM

* **General information of lnc-NDRG1-1-4:**

NONCODE TRANSCRIPT ID	NONHSAT129220.2	Strand	-
NONCODE Gene ID	NONHSAG051331.2	Exon Number	2
Chromosome	chr8	CNCI Score	-0.032768
Start Site	133237170	Length	508
End Site	133244503	Assembly	hg38

* **Primers:**

BMRC-2575	ACCGAGCTCGGGATCCATGtaccctatgatgtgcc agattatgccGGACAAGGCAGGGCAGGGCA	pcDNA3.1(+)-NDRG1-OT1_v4 (BamHI) HA tag
BMRC-2492	tctgcagaatgaattcCCGC	pcDNA3.1(+)-NDRG1-OT1_v4 (EcoRI)
BMRC-2494	CTGTGCATCGTCGTCCTTGTAGTCATTC CTCTGACCAAACCTC	pcDNA3.1(+)-NDRG1-OT1_v4-1
BMRC-2577	GACGACGATGACAAGTGAATATTGTAA GTCAGCCAC	pcDNA3.1(+)-NDRG1-OT1_v4-1

* TA cloning vector or Mammalian; Yeast; *E. coli*; expression vector

* plasmid amplify in *E. coli* is high; or low copy number

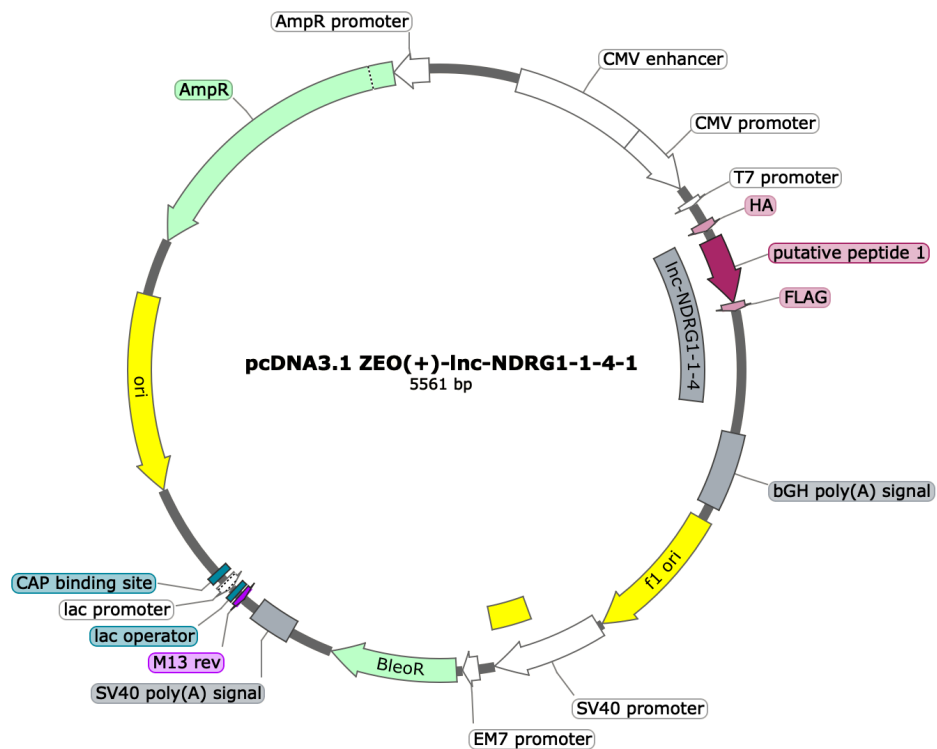
* expression level is low or high

* expression promoter: CMV IE promoter

* antibiotic selection: AmpR; BleoR

BleoR confers resistance to bleomycin, phleomycin, and Zeocin™

* epitope or tag: HA, FLAG



gtctatataagcagagctcctctggctaactagagaaccactgcttactggcttatcgaaattaatacgaactcactatag
cagatatattcgtctcgcagagaccgattgatctcttgggtgacgaatgaccgaatagctttaattatgctgagtgatatc
CMV promoter T7 promoter

ggagaccaagctgGCTAGCGTTAAACTTAAGCTTGGTACCGAGCTCGGGATCCATGtacccttatgatgtgccagatt
cctctgggttcgacCGATCGCAAATTTGAATTCGAACCATGGCTCGAGCCCTAGGTACatgggaatactacacggctcaa
1 1 5
Met Tyr Pro Tyr Asp Val Pro Asp
HA

atgccGGACAAGGCAGGGCAGGGCAGGGCAGGGGCGGATGGGCATCCTTTTCGGAAAAAATTCATAGAGCGCAGAGGGCCTTGGC
tacggCCTGTTCCGTCCCCTCCCCTCGCCCGCCTACCCGTAGGAAAGCCTTTTTTAAGTATCTCGCGTCTCCCGAACC
Inc-NDRG1-1-4
1 5 10 15
Met Gly Ile Leu Ser Glu Lys Ile His Arg Ala Gln Arg Ala Leu Ala
putative peptide 1

Tyr Ala Gly Gln Gly Arg Ala Gly Gln Arg Ala Asp Gly His Pro Phe Gly Lys Asn Ser *
HA (in frame with HA)

TGTGGTGTGGCACTAACGAGCTTCCCTCTCTCCTAGATGGCGGACTGTGGCGGCCTCCCGCAGATCTCCAGGTTCTT
ACACCACAAACCGTGATTGCTCGAAGGGAGAGAGGATCTACCGCTGACACCGCCGGAGGGCGTCTAGAGGGTCCAAGGA
Inc-NDRG1-1-4
20 25 30 35 40
Val Val Phe Gly Thr Asn Glu Leu Pro Ser Leu Leu Asp Gly Gly Leu Trp Arg Pro Pro Ala Asp Leu Pro Gly Ser
putative peptide 1

GTACTIONGCTTGTACTTTTCATTTGGCTCACCGTGGATTTTCTCATAGGAAGTTTGGTCAGAGTGAATGACTACAAG
CATGATGACGGAACATGAAAAGTAAAACCGAGTGGCACCTAAAAGAGTATCCTTCAAACCGTCTCACTTACTGATGTTT
Inc-NDRG1-1-4
45 50 55 60 65 1
Cys Thr Thr Ala Leu Tyr Phe Ser Phe Trp Leu Thr Val Asp Phe Leu Ile Gly Ser Leu Val Arg Val Asn Asp Tyr Lys
putative peptide 1 FLAG

GACGACGATGACAAGTGAATATTGTAAGTCAGCCACTGGgACCCGAGGAtTTCTGGGACCCCGCAGTTGGGAGGAGGAAG
CTGCTGCTACTGTTCACTTATAACATTCAGTCCGTGACCCcTGGGCTCCTaAAGACCCTGGGGCGTCAACCCTCCTCCTTC
Inc-NDRG1-1-4
5
Asp Asp Asp Asp Lys *
FLAG

TAGTCCAGCCTTCCAGGTGGCGTGAGAGGCAATGACTCGTTACCTGCCGCCATCACCTTGGAGGCCTTCCCTGGCCTTG
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Inc-NDRG1-1-4

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TCATCTTTTCAGCCCTAGCCCGTTCTCTCCGACTCATGCCTACCCTTTGATAACAGTGTTCAGAAAGGTCTCCTCAA
Inc-NDRG1-1-4

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AGAATTACTCTATAAACATAAATAAAGGTCTGGTTATTTAAACATTGAAACGTCGCCCTAAGTAAGACGTCTATAGGTC
Inc-NDRG1-1-4

CACAGTGGCGCCGCTCGAGTCTAGAGGGCCCGTTTAAACccgctgatcagcctcgactgtgccttctagttgccagcca
GTGTCACCGCCGGCGAGCTCAGATCTCCCGGGCAAATTTgggcgactagtcggagctgacacggaagatcaacggtcggt
bGH poly(A) signal

DNA Sequence of pcDNA3.1/ZEO(+)-lnc-NDRG1-1-4-1

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