

Product datasheet for **RN216770**

Nrg1 (NM_001271125) Rat Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Nrg1 (NM_001271125) Rat Untagged Clone
Tag:	Tag Free
Symbol:	Nrg1
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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Fully Sequenced ORF: >RN216770 representing NM_001271125
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGCC**

ATGTCTGAGCGCAAAGAAGGCAGAGGCAAGGGGAAGGCAAGAAGAAGGACCGGGGATCCCGCGGGAAGC
 CCGGGCCCGCGAGGGCGACCCGAGCCAGCACTGCCTCCAGATTGAAAGAAATGAAGAGCCAGGAGTC
 AGCTGCAGGCTCCAAGCTAGTGCTCCGGTGCGAAACCAGCTCCGAGTACTCCTCACTCAGATTCAAATGG
 TTCAAGAATGGGAACGAGCTGAACCGAAAAATAAACAGAAAACATCAAGATACAGAAGAAGCCAGGGA
 AGTCAGAGCTTCGAATTAACAAAGCATCCCTGGCTGACTCTGGAGAGTATATGTGCAAAGTGATCAGCAA
 GTTAGGAAATGACAGTGCCTCTGCCAACATCACCATTGTTGAGTCAAACGAGTTCATCACTGGCATGCCA
 GCCTCGACTGAGACAGCCTATGTGCTCAGAGTCTCCATTAGAATCTCAGTTTCAACAGAAGGCGCAA
 ACATTCTTCATCCACATCGACATCCACGACTGGGACCAGCCATCTCATAAAGTGC GCGGAGAAGGAGAA
 AACTTTCTGTGTGAATGGGGCGAGTGCTTACGGTGAAGGACCTGTCAAACCCGTC AAGATACTTGTGC
 AAGTGCCCAAATGAGTTTACTGGTGATCGTTGCCAAAAC TACGTAATGGCCAGCTTCTACATGACTTCTA
 GGAGGAAAAGGCAAGAAACAGAGAAGCCTCTAGAAAGAAAATTGGATCATAGCCTTGTGAAAGAATCGAA
 AGCGGAGGAACCTACCAGAAGAGGGTGCTGACAATTACTGGCATCTGTATCGCCCTGCTGGTGGTGGC
 ATCATGTGTGTGGTGGCTACTGCAAAACCAAGAAGCAGCGGCAGAAAGCTTCATGATCGGCTTCGGCAGA
 GTCTTCGGTCAGAACGGAGCAACCTGGTGAACATAGCGAATGGGCTCACCACCCAAACCCACCGCCAGA
 GAACGTGCAGCTGGTGAATCAATACGTATCTAAAAACGTCATCTCCAGTGAGCATATTGTTGAGAGAGAA
 GTGGAGACTTCCTTTTCCACCAGTCATTACACTCCACAGCCATCACTCCACGACTGTCAACCCAGACTC
 CTAGTCACAGCTGGAGTAATGGGCACAGGAGAGCATCATTTTCAGAAAGCAACTCCGTAATCATGATGTC
 TTCGGTAGAGAACAGCAGGCACAGCAGTCCCGCCGGGGGCCACGAGGAGGTCTTCATGGCCTGGGAGGC
 CCTCGTGATAACAGCTTCTCAGGCATGCCAGAGAAACCCCTGACTCCTACAGAGACTCTCCTCATAGCG
 AAAGACATAACCTTATAGCTGAGCTAAGGAGAAACAAGGCTTACAGATCCAAATGCATGCAGATCCAGCT
 GTCAGCAACTCATCTTAGACCCTTCCATTACCCATTTGGGCTCATTCT**TAA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001271125
- Insert Size:** 1455 bp
- OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001271125.1](#), [NP_001258054.1](#)

RefSeq Size: 3466 bp

RefSeq ORF: 1455 bp

Locus ID: 112400

Cytogenetics: 16q12.3

Gene Summary: ligand for ErbB3 and ErbB4 receptors; gene produces many different alternative splicing isoforms; involved in neural and organ development [RGD, Feb 2006]
Transcript Variant: This variant (8) differs in the 5' UTR and coding sequence, lacks an alternate coding exon, and contains two other alternate coding exons compared to variant 1. The resulting isoform (8) contains an alternate internal segment and has shorter and distinct N- and C-termini compared to isoform 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.