

## **Product datasheet for SC333763**

## NEURL3 (NM 001285486) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** NEURL3 (NM\_001285486) Human Untagged Clone

Tag: Tag Free Symbol: NEURL3

Synonyms: LINCR; RNF132

Mammalian Cell

Selection:

Neomycin

**Vector:** pCMV6-Entry (PS100001) **E. coli Selection:** Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC333763 representing NM\_001285486.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

**ACGCGTACGCGGCCGCTC**GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul

**ACCN:** NM 001285486

Insert Size: 387 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).

**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).



**OriGene Technologies, Inc.** 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



## NEURL3 (NM\_001285486) Human Untagged Clone - SC333763

**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: <u>NM 001285486.1</u>

 RefSeq Size:
 1770 bp

 RefSeq ORF:
 387 bp

 Locus ID:
 93082

 Cytogenetics:
 2q11.2

 MW:
 13.9 kDa

Gene Summary: E3 ubiquitin-protein ligase. Seems to utilize UBE2E1. In vitro, generates polyubiquitin chains

via non-canonical lysine residues suggesting that it is not involved in tagging substrates for

proteasomal degradation.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) contains two alternate exons, one of which results in a frameshift, compared to variant 1. The resulting protein (isoform b) uses a downstream, in-

frame AUG and has a distinct C-terminus, compared to isoform a.