

Product datasheet for SC207005

NEURL4 (NM_001005408) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: NEURL4

Mammalian Cell Selection: Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_001005408

Insert Size: 525 bp

Insert Sequence: >SC207005 3'UTR clone of NM_001005408
The sequence shown below is from the reference sequence of NM_001005408. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon Red=Cloning site

```
GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTCTGTGCCCTGCTGGTACGGGTGGAATGAGGGGTGAGACACCACTACTACAAGCACAGTCGGGCCGCG
GGCCCATGGACTCTGAGTGGCGACTGCCTCCACCTCATTCCCGTGAAGTGGCATGCGCAGGTGCTGG
AGCTTGGCAGCCGCGCAGGAGCATGTAGGCAGGCTCTCAGATGTAGGTGGCAAGTGGCACAGCTCCATG
TCCGGAGGCCGAGCACTCCGTCTGATGGGAGGAGTCGTGGGAGCCAGCTCCAGGCCCTGGTACCCCTC
TTCATGCACTGATTTGGGAACATGACTCCCTTTTACTCCCTACCCACATCACTTAATTTATTTCCG
TTTTTGTCTGTTACTGTGAATCCAGAGGAGTCTCTCCCTGTGCCACATGAAGCTGCTTTTTCCG
GGGCCACCGGGCGGAGTGGGGAAGGTTGGGCGCACGGAAGATGGGGCCTCTGTACAGTTGTTACTGA
CTCTGATTTCTAAGGAGCCAATAAACACCGTCTCAGAGCCTC
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites: SgfI-MluI

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms (SNPs).



Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM_001005408.2</u>
Summary:	The protein encoded by this gene is predicted and it includes two isoforms resulting from two alternatively spliced transcript variants. [provided by RefSeq, Jul 2008]
Locus ID:	84461
MW:	19.2