

Product datasheet for SC205830

EGFL7 (NM_016215) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	EGFL7 (NM_016215) Human 3' UTR Clone
Symbol:	EGFL7
Synonyms:	NEU1; VE-STATIN; ZNEU1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_016215
Insert Size:	422 bp
Insert Sequence:	>SC205830 3'UTR clone of NM_016215 The sequence shown below is from the reference sequence of NM_016215. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GGGTCCTGCTCCTGCAAGAAAGACTCGTGACTGCCCAGCGCCCCAGGCTGGACTGAGCCCCTCACGCCG
CCCTGCAGCCCCATGCCCTGCCAACATGCTGGGGTCCAGAAACCACCTCGGGTGACTGAGCGGA
AGGCCAGGCAGGGCCTTCTCCTCTTCTCCTCCCTTCTCCTCGGGAGGCTCCCCAGACCCTGGCATGGG
ATGGGCTGGGATCTTCTGTGAATCCACCCTGGCTACCCCCACCTGGCTACCCAACGGCATCCCA
AGGCCAGGTGGGCCCTCAGCTGAGGGAAGTACGAGCTCCCTGCTGGAGCCTGGGACCCATGGCACAGG
CCAGGCAGCCCGAGGCTGGGTGGGCCTCAGTGGGGGCTGCTGCCTGACCCCCAGCACAAATAAAATG
AAACGTGA
ACGCGTAAGCGGCCGCGGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites:	Sgfl-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.



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RefSeq: [NM_016215.5](#)

Summary: This gene encodes a secreted endothelial cell protein that contains two epidermal growth factor-like domains. The encoded protein may play a role in regulating vasculogenesis. This protein may be involved in the growth and proliferation of tumor cells. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Feb 2012]

Locus ID: 51162

MW: 14.8