

## Product datasheet for **SC204461**

### DUSP15 (NM\_001012644) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	DUSP15 (NM_001012644) Human 3' UTR Clone
Symbol:	DUSP15
Synonyms:	C20orf57; VHY
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001012644
Insert Size:	356 bp
Insert Sequence:	<p>&gt;SC204461 3'UTR clone of NM_001012644 The sequence shown below is from the reference sequence of NM_001012644. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b>=Stop Codon <b>Red</b>=Cloning site</p> <pre>GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA<b>GCGATCGCC</b> CGGTGTCTGTCCCGAAGGGCGGCAAG<b>TGA</b>GGATGCAGTCCAGCCGTGGCTCCCCACTTCCGACTGGCT CCCTTCGGGGGTGTCTGCGCCTTCCACGCCCCCAGGACGGGCCAGAGGCTGGGGGAGCCCCGCGGC GGCCTGAACCCTGCCTCCCGCGCCCGCCTGCTCGTCCGCGTCTGCAGTCAGCGTCCCCAACCTGTGCG TCTCTGTGTCCGGGCCGGCCTGCTGCAGCCACCTGGTGCCTTAGTCCTTGGGCTGGGGGAGGGGCCCA CCCTTAAAGGCGCGGGAGGGGAGGGAGAGTGGAGGGTTTGACGGGCTGGAGGGTATTAAGAG ACACAGAAGAA <b>ACGCGT</b>AAGCGGCCGCGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG</pre>
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\\_001012644.3](#)

**Summary:** The protein encoded by this gene has both protein-tyrosine phosphatase activity and serine/threonine-specific phosphatase activity, and therefore is known as a dual specificity phosphatase. This protein may function in the differentiation of oligodendrocytes. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Feb 2016]

**Locus ID:** 128853

**MW:** 12