

Product datasheet for **SC200412**

ADAM22 (NM_004194) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	ADAM22 (NM_004194) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	ADAM22
Synonyms:	ADAM 22; DEE61; EIEE61; MDC2
ACCN:	NM_004194
Insert Size:	118 bp
Insert Sequence:	>SC200412 3'UTR clone of NM_004194 The sequence shown below is from the reference sequence of NM_004194. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC TTTAGACCTCGGTCTAATTCAACTGAGTAGTCTGAACCTCTAATGGAAAAAGGCACTCCAAGGTCCTG TGCAAATAGTGTCAAAAAGTAAATTAAGCAGATAGTCAAAGAAGTAAA ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
Restriction Sites:	Sgfl-Mlul
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.
RefSeq:	<u>NM_004194.5</u>



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Summary:

This gene encodes a member of the ADAM (a disintegrin and metalloprotease domain) family. Members of this family are membrane-anchored proteins structurally related to snake venom disintegrins, and have been implicated in a variety of biological processes involving cell-cell and cell-matrix interactions, including fertilization, muscle development, and neurogenesis. Unlike other members of the ADAM protein family, the protein encoded by this gene lacks metalloprotease activity since it has no zinc-binding motif. This gene is highly expressed in the brain and may function as an integrin ligand in the brain. In mice, it has been shown to be essential for correct myelination in the peripheral nervous system. Alternative splicing results in several transcript variants.[provided by RefSeq, Dec 2010]

Locus ID:

53616

MW:

4.4