

Product datasheet for **SC206837**

RHEX (NM_001007544) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	RHEX
Synonyms:	Clorf186
Mammalian Cell	Neomycin
Selection:	
Vector:	pMirTarget (PSI00062)
ACCN:	NM_001007544
Insert Size:	527 bp
Insert Sequence:	>SC206837 3'UTR clone of NM_001007544 The sequence shown below is from the reference sequence of NM_001007544. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC CCAGCGGAATATGATCAAGTGCCATGTGAATTCCAAATATTTTAAATGGGGTCCAGTTCTCTATGGAT TCTTACATTTAATTTGTAGGGAATGCCATTTTCCCCCTTAAACAAGGCATGGGGCTCACAAGTCTAT GGAGACAGGCCAAAAAGAATGTGGAGAAGAACTGATAAATACACAGAGGTCCTCAAGACCCATGGAC TCCTGGTCTGTACCCAAAAAGCTGTTCTGTTCTCAAAAACAAAAACAAGGCTTGGCTGGGAAACAGG CCAATGCCCGGCAAGAAAGGTTGAGATCAGATGTTAGGAAGAACTTTCAGGTAAAGTATGAGAACTAT GGAGTCCATCAGCAGAGATAGTAGTGAAGTCTCTCCCAGGGAATTTTAAAAAGGTTGAATCAGCTG TTGTAGAGTTCTATTTGGCAATCTCATGGTTAAATGACTTCCCTTTGAGCTCTTAATTATTGGCAATA AACAACCTCTTTAAAAGTTTTAAATAAAATAGCAACCACCACCA ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
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_001007544.4</u>
Summary:	Acts as a signaling transduction factor of the EPO-EPOR signaling pathway promoting erythroid cell differentiation (PubMed:25092874).[UniProtKB/Swiss-Prot Function]
Locus ID:	440712
MW:	20.6