

Product datasheet for SC207628

NCF2 (NM_001127651) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	NCF2 (NM_001127651) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	NCF2
Synonyms:	NCF-2; NOXA2; P67-PHOX; P67PHOX
ACCN:	NM_001127651
Insert Size:	586 bp
Insert Sequence:	>SC207628 3'UTR clone of NM_001127651 The sequence shown below is from the reference sequence of NM_001127651. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site
	GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GATTTGGAAAGCACTCGGAGAGAAGTCTAGGATGTTTCACAAACTACAAAGCTGAAGAAAATGAAGCCC TATTACTTGTTTGTAAGATTTAGCACCCTTCTGCTGTATACTGTACTGAGACATTACAGTTTGGAAGTG TTAACTATTTATTCCCTGTTAAAATTTAACCTACTAGACAATGATGTGAGTACCCAGGATGATTTCCTG GGGCACAGTGGGTGAGGAGATGGGGACAGGTGAATGGAGGAGTTAGGGGAGAGAAAAGTGGATGGA
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.



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	NCF2 (NM_001127651) Human 3' UTR Clone – SC207628
RefSeq:	<u>NM 001127651.3</u>
Summary:	This gene encodes neutrophil cytosolic factor 2, the 67-kilodalton cytosolic subunit of the multi-protein NADPH oxidase complex found in neutrophils. This oxidase produces a burst of superoxide which is delivered to the lumen of the neutrophil phagosome. Mutations in this gene, as well as in other NADPH oxidase subunits, can result in chronic granulomatous disease, a disease that causes recurrent infections by catalase-positive organisms. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jun 2010]
Locus ID:	4688
MW:	22.1

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