

Product datasheet for **SC205423**

LRRC48 (DRC3) (NM_001130092) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	LRRC48 (DRC3) (NM_001130092) Human 3' UTR Clone
Symbol:	LRRC48
Synonyms:	CFAP134; LRRC48
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001130092
Insert Size:	519 bp
Insert Sequence:	>SC205423 3'UTR clone of NM_001130092 The sequence shown below is from the reference sequence of NM_001130092. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAACGCATCGCC AATTATGTGGAGGCTTCTGGCACTTTCTTAAGAGAAGCAGTTACAGGTAGAGGTTGAAGTCTGAGATTTTCT CCCTGGAGGTCTTCTTAACCTTAGAGTCTGTATAATGAGTACTGTGGGGTGTGGAAACCACCATTTCT GATCCTGCACAATGCCATCCGGTTTGACCAGCACATCCACTGATGAATGGTCACAAAATCAGCAGGGT CGGCCTGAGGAACCTGGCTCCATCCCTGGGGTCTGCACGCTAAGAAGGCTCTCCCGGATGTCTGCGCTG GGTCCCGCGGCAGCATGTCTATGTCTCTGGAAGGAGTGTGGGCATCAAGGGCCAGACAGAGATACTAT GTTGTACAGGGCCAGACAGAGATACTATGTTGTACAGGGCCTGCTTTAGTGCATGCATCATTTCACTCCAC TGCCAGTTTCTCAGCTTCTCTGTGCGGACATCACCAACCAAGGCAAGCCACCATCGTCCTTGGCCTGG ACTGCTTGCCTCTTTACTGTCCTCTTACAGTCCGTC ACGCGTAAGCGGCCGCGCATCTAGATTCTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG </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).


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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:	NM_001130092.2
Summary:	Component of the nexin-dynein regulatory complex (N-DRC) a key regulator of ciliary/flagellar motility which maintains the alignment and integrity of the distal axoneme and regulates microtubule sliding in motile axonemes.[UniProtKB/Swiss-Prot Function]
Locus ID:	83450
MW:	19.3