

Product datasheet for **MC219610**

Klc1 (NM_001081959) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Klc1 (NM_001081959) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Klc1
Synonyms:	A1874768; Kn; Kns2
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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Fully Sequenced ORF: >MC219610 representing NM_001081959
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGGATCGCC**

ATGTATGACAACATGTCCACCATGGTGTACATAAAGGAAGAGAAGCTGGAGAAGCTCACGCAGGATGAGA
 TCATCTCTAAGACCAAGCAAGTATCCAGGGGCTGGAAGCCCTGAAGAATGAGCACAACCTCCATCTGCA
 GAGTTTGTCTGGAGACGCTGAAGTCTTGAAGAAGGATGACGAGAGCAACCTGGTGGAAAGAAAATCCAGC
 ATGATCCGCAAGTCCCTGGAGATGCTGGAGCTTGGCTGAGCGAGGCGCAGGTGATGATGGCGCTGTCCA
 ATCACCTGAATGCTGTGGAGTCCGAGAAGCAAAAGCTCCGCGCTCAGTTTCGACGGCTGTGCCAGGAGAA
 CCAGTGGTGGGGATGAGCTGGCAACACGCAGCAGAAGTTGCAGAAGAGCGAGCAGTCGGTGGCGCAG
 CTGGAGGAGGAGAAGAACACCTGGAGTTCATGAACCAGCTGAAGAAGTACGACGACGACATCTCCCCT
 CGGAGGACAAAGACTCTGATTCTTCAAAGAGCCGTTGGATGATCTCTTCCAAAATGACGAGGACGAACC
 AGGACAAGGAATCCAGCAGCAGCAGTGTGCTGCGGCCGCCAGCAGGGCGGCTACGAGATCCCT
 GCAAGGCTGCGCAGCTCCACAACCTGGTATCCAGTATGCTTACAGGGGCGTTACGAGGTGGCGGTGC
 CACTCTGCAAGCAGGCCCTGGAGGATCTGGAGAAGACTTCCGGCCACGACCACCCCGATGTGGCTACCAT
 GCTCAACATCTTGGCCCTGGTGTACAGGGATCAGAAACAAGTATAAAGATGCAGCTAACCTCCTGAACGAC
 GCCCTGGCTATCCGCGAGAAAACCTGGGCAGAGATCACCCCGCGGTGGCAGGACTCTGAACAACCTAG
 CAGTACTGTACGGTAAGCGAGGGAAGTACAAGGAGGCGGAGCCGCTGTGTAACGAGCCCTGGAGATCAG
 GGAGAAGTCTGGGAAAGGATCATCTGATGTTGCCAAACAGTTAAATAACCTGGCCCTGCTGTGCCAG
 AACCAGGGCAAGTACGAGGAGGTGGAGTATTATTACCAGAGGGCCCTGGGCATCTACCAGACGAAGCTGG
 GGCCCGACGATCCCAACGTGGCCAAGACCAAGAACAACCTGGCCCTCTGTTATCTGAAACAAGGGAAGTT
 CAAGCAGGCAGAAAACGCTGTACAAGGAGATTCTACCCGCGCACAGCGGGGAGTTTGGATCTGTGGAC
 GACGAGAACAAGCCATCTGGATGCACGCTGAAGAGAGAGAGGAGTGCAAGGCAAGCAGAAGGACGGGT
 CGGCTTTTGGAGAGTATGGCGGCTGGTATAAAGCCTGCAAAGTGGACAGTCCCACCGTCACAACCACCTT
 GAAAAACCTTGGAGCACTTACCAGCGCAGGGGAAGTTTGAAGCTGCAGAGACATTGGAAGAAGCCGCC
 ATGAGGTCACGTAAGCAGAGAGTGGCTGAAGTGTAAATGACCCTGAGAGCATGGAGAAGCGGAGGAGCC
 GGGAGAGTCTCAATATGGACGTGGTCAAGTACGAGAGTGGCCCTGACGGAGGGGAGGAAGTGAATGAG
 CGTAGAGTGAATGGGGATGGCACTGGATCTTTAAAGCGCAGTGGCTCCTTAGCAAACCTCCGGCTTCC
 ATTAGACGACGAGTGAAGCTGGTTAGGAAGCTGAAGGGAGGAAGCTCACGGGACAGTGAAGCCGAGGA
 ACCCTGGGGCATCCCCGACAGCCTCTTGTGTGAAAAACGACAGCAGCAGCCTAGAAGACGCTAGCAC
 TAACTGA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001081959

Insert Size: 1827 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001081959.1](#), [NP_001075428.1](#)

RefSeq Size: 2480 bp

RefSeq ORF: 1827 bp

Locus ID: 16593

Cytogenetics: 12 61.13 cM

Gene Summary: Conventional kinesin is a tetrameric molecule composed of two heavy chains and two light chains, and transports various cargos along microtubules toward their plus ends. The heavy chains provide the motor activity, while the light chains bind to various cargos. This gene encodes a member of the kinesin light chain family. It associates with kinesin heavy chain through an N-terminal domain, and six tetratricopeptide repeat (TPR) motifs are thought to be involved in binding of cargos such as vesicles, mitochondria, and the Golgi complex. Thus, kinesin light chains function as adapter molecules and not motors per se. Although previously named "kinesin 2", this gene is not a member of the kinesin-2 / kinesin heavy chain subfamily of kinesin motor proteins. Extensive alternative splicing produces isoforms with different C-termini that are proposed to bind to different cargos; however, the full-length nature of some of these variants has not been determined. [provided by RefSeq, Jul 2008]
Transcript Variant: This variant (h) uses an alternate in-frame splice site in the 3' coding region, compared to variant d. The resulting protein (isoform 1H) is shorter than isoform 1D.