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Product datasheet for SC205962

TNNI3K (FPGT-TNNI3K) (NM_001112808) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	TNNI3K (FPGT-TNNI3K) (NM_001112808) Human 3' UTR Clone
Symbol:	TNNI3K
Synonyms:	CARK; TNNI3K
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_001112808
Insert Size:	476 bp
Insert Sequence:	>SC205962 3'UTR clone of NM_001112808 The sequence shown below is from the reference sequence of NM_001112808. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AATAGTAGCAGCTTTGAGGACAGCAGCTGACAGCATTCGGCGTATACCTAAGGAGAGTTTTTTCCCCGA ACTGACAGCAACGATTCCAACCACGGCAAGCTGGCTTCCAACTATAACATTTTACTCTCAAAGGTCTCC TTAAATTGGGCTTGTTTTTACTTGTCCTATTTAATTCCCCACTATTAGCAGGCTTTGGATTTGTGCCTA AGGAATAATATGCAAAAGAACCAAGACAGAATGTATATGAAGAATTGTTTTTAATTTTGTAAATTAAAA AAAAATTTAGATCGTTACTTGGAAATGGAGCCTAAGTCTGTGGTGGACAGATAATAATTATGTTTTCCT GGGCTGAATTATGTAGACTTGTGTTTGACAGCTATGGGTTTATTTCTTAGAACATTGTTCATTTTCTTT TCTCATTATGTTACTTCTAGTGTTCACCTCTGTGATTAAAGATTCTTTGGTGAAATAGAAAA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
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.

View online »

RefSeq:	NM_001112808.3
Summary:	This locus represents naturally occurring read-through transcription from the neighboring fucose-1-phosphate guanylyltransferase (FPGT) and TNNI3 interacting kinase (TNNI3K) genes. Alternative splicing results in multiple transcript variants that are composed of in-frame exons from each individual gene. [provided by RefSeq, Dec 2010]
Locus ID:	100526835
MW:	18.5