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Product datasheet for SC204974

LTK (NM_206961) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	LTK (NM_206961) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	LTK
Synonyms:	TYK1
ACCN:	NM_206961
Insert Size:	389 bp
Insert Sequence:	>SC204974 3'UTR clone of NM_206961 The sequence shown below is from the reference sequence of NM_206961. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AACCTTTGGAATCCCACTTATCGCTCCTGAGCCCCAAGGGGCCCTGAGGGTAAGGACTGAGGCACTGAG GGTCCCTCCCTATACTCCTCAGGCTCCTGGGTGGCCTGTTATGCCAGCGGCCTCTGTTCCCTGCAGTCT GTGCTGTGTGTCTGGGCCTGTCTCGGGGCTGGCCTGGCAGCGCTGCACTTGCCATGCTGGAAACCAGCC CAGGCCTCCCAGGGAAGGGCCCAGCCACTTCCAGCTTTTGATCTTGGGGCCAGAGGCCGCCTTACACAC ACCCCAGGTGTCCATGGGGAGCACTGGATTGCTCTCCCATTATGAGCATCCTTCATCTGGGCAGACCCC CCACCCTGCAGATGCTTCTAATAAAAAGCTCTTCTCATCCTCCA 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.
RefSeq:	NM_206961.4

View online »

Summary:	The protein encoded by this gene is a member of the ros/insulin receptor family of tyrosine kinases. Tyrosine-specific phosphorylation of proteins is a key to the control of diverse pathways leading to cell growth and differentiation. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Oct 2008]
Locus ID:	4058
MW:	13.8