

Product datasheet for **SC204973**

LTK (NM_001135685) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	LTK (NM_001135685) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	LTK
Synonyms:	TYK1
ACCN:	NM_001135685
Insert Size:	389 bp
Insert Sequence:	>SC204973 3'UTR clone of NM_001135685 The sequence shown below is from the reference sequence of NM_001135685. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AACCTTTGGAATCCACTTATCGCTCCTGAGCCCAAGGGGCCCTGAGGGTAAGGACTGAGGCACTGAG GGTCCCTCCCTATACTCCTCAGGCTCCTGGGTGGCCTGTTATGCCAGCGGCTCTGTTCCCTGCAGTCT GTGCTGTGTCTGGGCCTGTCTCGGGGCTGGCCTGGCAGCGCTGCACTTGCCATGCTGGAACAGCC CAGGCCTCCCAGGAAGGGCCCAGCCACTTCCAGCTTTTATCTTGGGGCCAGAGGCCGCTTACACAC ACCCAGGTGTCATGGGAGCACTGGATTGCTCTCCATTATGAGCATCCTTCATCTGGGCAGACCCC CCACCCTGCAGATGCTTCTAATAAAAAGCTCTTCTCATCCTCCA ACGCGTAAGCGGCCGCGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCCTTCTATGAAAGG 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:	<u>NM_001135685.2</u>



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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