

Product datasheet for **SC204106**

IL32 (NM_001012718) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	IL32 (NM_001012718) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	IL32
Synonyms:	IL-32alpha; IL-32beta; IL-32delta; IL-32gamma; NK4; TAIF; TAIFa; TAIFb; TAIFc; TAIFd
ACCN:	NM_001012718
Insert Size:	205 bp
Insert Sequence:	>SC204106 3'UTR clone of NM_001012718 The sequence shown below is from the reference sequence of NM_001012718. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AAGTGCTCTGAACCCCAATCCTCAAAAATGAGATACTGACACCACCTTTGCCCTCCCCGTCACCCGCGCA CCCACCCTGACCCTCCCTCAGCTGTCTGTGCCCGCCCTCTCCCGCACACTCAGTCCCCTGCCTGG CGTTCCTGCCGAGCTCTGACCTGGTGTCTGCGCCCTGGCATCTTAATAAACCTGTTATACTTCC ACGCGTAAGCGGCGCGGCATCTAGATTGAAAGAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
Restriction Sites:	Sgfl-Mlul
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_001012718.4



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Summary: This gene encodes a member of the cytokine family. The protein contains a tyrosine sulfation site, 3 potential N-myristoylation sites, multiple putative phosphorylation sites, and an RGD cell-attachment sequence. Expression of this protein is increased after the activation of T-cells by mitogens or the activation of NK cells by IL-2. This protein induces the production of TNFalpha from macrophage cells. Alternate transcriptional splice variants, encoding different isoforms, have been characterized. [provided by RefSeq, Jul 2008]

Locus ID: 9235

MW: 7.7