Product datasheet for SC204404

ZSCAN2 (NM_017894) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	ZSCAN2 (NM_017894) Human 3' UTR Clone
Symbol:	ZSCAN2
Synonyms:	ZFP29; ZNF854
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_017894
Insert Size:	340 bp
Insert Sequence:	>SC204404 3'UTR clone of NM_017894 The sequence shown below is from the reference sequence of NM_017894. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC TCCCTTCCGGTGGAGGTGACCAGTTTGTAATGAACTTCTGGATCTCCAGTCAGATACTATGGAGGAGAG AGTGATGGAGGGCTGAGGAGAGGCCTCAGGTTGTTATTTGGTTGGGGAGGCACAGAGCAGAGGTTCAGG TCTCAGCAGGTGTCAGAGGTGATTACAGCCCTTCCCATCCACCTTGCAGCAGGGGCTTTGTGCCTTTCA TCTTTAACTGGGCACCTGAGATGGGAACCAATGGGGACCATCCCAGACACATGGGTTTGTCTGTCCCTG ACTTCTGCCTCTGAGATTGTCTTTGTTGGTCTTTGTTTAATAAACGCCCCGACCTTCCTCATGA 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_017894.6

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Summary:	The protein encoded by this gene contains several copies of zinc finger motif, which is commonly found in transcriptional regulatory proteins. Studies in mice show that this gene is expressed during embryonic development, and specifically in the testis in adult mice, suggesting that it may play a role in regulating genes in germ cells. Alternative splicing of this gene results in several transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]
Locus ID:	54993
MW:	12.2