

Product datasheet for **SC206901**

ZNF18 (NM_144680) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	ZNF18
Synonyms:	HDSG1; KOX11; Zfp535; ZKSCAN6; ZNF535; ZSCAN38
Mammalian Cell	Neomycin
Selection:	
Vector:	pMirTarget (PSI00062)
ACCN:	NM_144680
Insert Size:	548 bp
Insert Sequence:	>SC206901 3'UTR clone of NM_144680 The sequence shown below is from the reference sequence of NM_144680. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC TCCCACTTAGGAAAGAAGCCCTTTCAAAGCCAGTAACCAAACTCTCTTTCCCATTTCTATCTCCAG CCCAGTCACAAAAATACTCAGCTCCATCAAGAGGAATTGTGTCTAAGAGGATACCCCTGTTAATCTCCT TTTTCTTGGATTGGAGAGGAGAGAATCTGGACATGGCTTTGGACTTGGAGGATATCTTGGATTGGATT GCACAATGGCTTAAATTCTTGATTCTGCCTCAGGAGAAAGAATAGTCTTCATGTTTCCACTCATCCTTC CTTTGGACCCATCGGGGAAAAAGTCTAAATTGGAGATCCAGTTTTAGAAGTGCTTTCTGGGAAGCATTT AATGGGATTAGCTGTAGTCACTGCTTATGGGAAGAACCTCAGATCAGCCCTTAAATGAGTTCTAGAG CAGGTCTTCTGTTCCAGAAGGGGAGAAGCATAGAGGCCTGTGAGCTCACGTGTGTTCTTTGTCATAGG GGTGAAAACTAATTCAAGTGTCCCTTGTGTTGAAATAAACTAGCAGAGTCACTTTCTATCTTA ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM_144680.4</u>
Summary:	May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]
Locus ID:	7566
MW:	21.1