

Product datasheet for **SC206907**

ZNF415 (NM_001136038) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: ZNF415 (NM_001136038) Human 3' UTR Clone
Symbol: ZNF415
Synonyms: Pact; ZfLp
Mammalian Cell Selection: Neomycin
Vector: pMirTarget (PS100062)
ACCN: NM_001136038
Insert Size: 528 bp
Insert Sequence: >SC206907 3'UTR clone of NM_001136038
The sequence shown below is from the reference sequence of NM_001136038. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG  
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC  
ACTAAGGAGAAACCTTATAAAAGAAATTAAATGGCAAGGTCTTCAGTCAAAGTTTAAATCCTGTGAGT  
CATCAAAGAAATTTATATCAGAGAGAAACCATACAAGTATAATAAATGTGGCAAGGTTTTTCAGTCACAAT  
TCACTCCTACACAGCATCAGAGAATTTCTTCTTGAGAGAATCCTTACGAGTACAGCAAACCTTCATC  
ACAAGTTCAAGCATTCTTGACATCAGAGTCCATGCTAAAGAGAAATCATATACACCTAACTGTGTGGC  
AGAGGCTTCATTTAGGTCTCACAACCTACTAGACATCAAATGTGTAAACATCTTTGTATATTTTGTGC  
ATGTTGAAGCTATTAACCAAGGATCAAACTGTAAACATCCAAGGATTTATGTGAGGAATAATTCAGT  
CTAGTTGTGCTGATAAACTTTTCATATTACACATTGTAGAACAAATGCAAGCCCAAATATGTTAAAACT  
CACACAACATGATATATATTAAGGTTGCAGGATGTTTGAAGTCA  
ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA  
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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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).



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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_001136038.4
Summary:	Involved in transcriptional regulation. Transcriptional activity differed among the various isoforms. All isoforms except isoform 3 seem to suppresses the transcriptional activities of AP-1 and p53/TP53.[UniProtKB/Swiss-Prot Function]
Locus ID:	55786
MW:	20.6