

Product datasheet for **SC207128**

ZNF350 (NM_021632) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: ZNF350

Synonyms: ZBRK1; ZFQR

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PSI00062)

ACCN: NM_021632

Insert Size: 544 bp

Insert Sequence: >SC207128 3'UTR clone of NM_021632
The sequence shown below is from the reference sequence of NM_021632. 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
GTCTTATTTTATGTTACAGAAAACCAAGGAAGAAAACCTCAGATCTATGTGGAAAAGGGTTGAGCAAA
AATTTGTAGTTTATTATGTGGCCGAAAAGCATACACTGAGAGAACATGTATAAGGCTGAGATAGCCTGA
TAAACTCATCCTATTTAAATGTATGCTGTGATACAGGCAATTTGATGTTAACCTAAGCACATACAC
AGCAATTGCTCGACTGTGTCAATTAATGAGTAAAGGAAGCCCAAGTACTTTTAAGTGGAGGAAATGAA
GTACTGTGAATTTTAAACAGATGAGATAATTGGTGTGGGAGTGTTCAGTGAAGTGAAGCTTCAT
AAGGGGCAGTTGGTGGGAAAGGGCTAATACTGTAATGGTGGATGTGGGACACATATGTGTCAATTCAG
TTTCCTTAGACCAGTGCCAGTGTGGAGGATTAATTGAAACCAGGACCATAAACTTTCAGAACTTTGTA
TTATGCAGAGGGATCTGTGAACTAGTCAATTTGTTTCATTAAACAAATATTTTAAATTA
ACGCGTAAGCGGCCGCGGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTGATTCCACCGCCGCTTCTATGAAAGG
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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).



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_021632.4</u>
Summary:	Transcriptional repressor. Binds to a specific sequence, 5'-GGGxxxCAGxxxTTT-3', within GADD45 intron 3.[UniProtKB/Swiss-Prot Function]
Locus ID:	59348
MW:	20.4