

Product datasheet for **SC336562**

ANKZF1 (NM_001282792) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ANKZF1 (NM_001282792) Human Untagged Clone
Tag:	Tag Free
Symbol:	ANKZF1
Synonyms:	Vms1; ZNF744
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >SC336562 representing NM_001282792.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

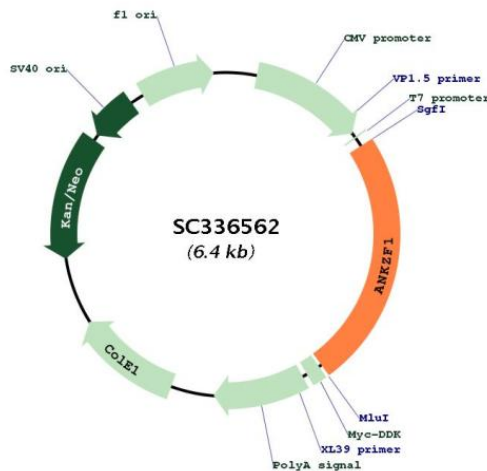
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CACTCTGCTGGAGCCAACCTGAGGCGCTACAATGAAGCCACACTATATAAGGATGTTTCGTGACTGCTG
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Restriction Sites:

Sgfl-MluI

Plasmid Map:



ACCN:

NM_001282792

Insert Size:	1551 bp
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM_001282792.1</u>
RefSeq Size:	2193 bp
RefSeq ORF:	1551 bp
Locus ID:	55139
Cytogenetics:	2q35
MW:	57.6 kDa
Gene Summary:	<p>Plays a role in the cellular response to hydrogen peroxide and in the maintenance of mitochondrial integrity under conditions of cellular stress (PubMed:28302725). Involved in the endoplasmic reticulum (ER)-associated degradation (ERAD) pathway (By similarity). [UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (3) lacks three consecutive exons in the 5' region, and initiates translation at a downstream AUG codon, compared to variant 1. The resulting isoform (2) has a shorter N-terminus, compared to isoform 1.</p>