

Product datasheet for **SC332202**

ZNF638 (NM_001252613) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ZNF638 (NM_001252613) Human Untagged Clone
Tag:	Tag Free
Symbol:	ZNF638
Synonyms:	NP220; ZFML; Zfp638
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC332202 representing NM_001252613. Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGTCGAGACCCAGGTTTAATCCTCGAGGAGACTTCCACTTCAAAGGCCACGAGCACCTAACCTTCT
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GAAGATGACACTTTGTCAGAATGTAACAGGTGTCTGATAAAGCTGTTTCTCTCCAGCGAAAGCTTCGG
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 TCTAGGTGA

Restriction Sites:	Sgfl-MluI
ACCN:	NM_001252613
Insert Size:	5874 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:	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_001252613.1</u>
RefSeq Size:	6463 bp
RefSeq ORF:	5874 bp
Locus ID:	27332
UniProt ID:	<u>Q14966</u>
Cytogenetics:	2p13.3-p13.2
MW:	218.2 kDa
Gene Summary:	The protein encoded by this gene is a nucleoplasmic protein. It binds cytidine-rich sequences in double-stranded DNA. This protein has three types of domains: MH1, MH2 (repeated three times) and MH3. It is associated with packaging, transferring, or processing transcripts. Multiple alternatively spliced transcript variants have been found for this gene, but the biological validity of some variants has not been determined. [provided by RefSeq, Jul 2008] Transcript Variant: This variant (4) uses an alternate splice site in the 3' coding region, but maintains the reading frame, compared to variant 1. The encoded isoform (2) is shorter than isoform 1.