

## Product datasheet for SC330769

## ZNF248 (NM\_001267607) Human Untagged Clone

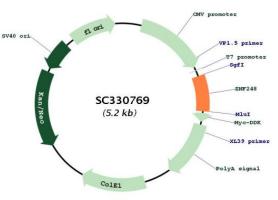
Sgfl-Mlul

## Product data:

## **Product Type: Expression Plasmids Product Name:** ZNF248 (NM\_001267607) Human Untagged Clone Tag: Tag Free Symbol: **ZNF248** bA162G10.3 Synonyms: Vector: pCMV6-Entry (PS100001) >SC330769 representing NM\_001267607. **Fully Sequenced ORF:** Blue=Insert sequence Red=Cloning site Green=Tag(s) CTGCTGGACCCTGCTCAGAAGATTCTATACAGAGATGTGATCCTGGAAAATTATAGCAATCTTGTCTCA GTAGGGTATTGCATTACTAAACCAGAAGTGATCTTTAAGATCGAGCAAGGAGAAGAGCCCTGGATATTA GAAAAAGGATTCCCAAGCCAGTGCCACCCAGAAAGGAAATGGAAAGTTGATGACGTGTTAGAGAGCAGC CAGGAAAATGAAGATGACCATTTTTGGGAGCTTCTATTCCACAACAACAACACTCAGGTTTCAGCTC CTCGGAAGAAAGCAGTTCTTGTGA

**Restriction Sites:** 

Plasmid Map:



ACCN:

NM\_001267607



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<b>ORIGENE</b> ZNF248 (NM_001267607) Human Untagged Clone – SC330769	
Insert Size:	369 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> <li>Centrifuge at 5,000xg for 5min.</li> <li>Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>Close the tube and incubate for 10 minutes at room temperature.</li> <li>Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>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.</li> </ol>
RefSeq:	<u>NM 001267607.1</u>
RefSeq Size:	957 bp
RefSeq ORF:	369 bp
Locus ID:	57209
Cytogenetics:	10p11.21
Protein Families:	Transcription Factors
MW:	14.7 kDa
Gene Summary:	May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (5) differs in the 3' coding region and 3' UTR, compared to variant 1. The resulting isoform (3) is shorter and has a distinct C-terminus, compared to isoform 1. Variants 5 and 12 both encode the same isoform (3). Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.

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