

## OriGene Technologies, Inc.

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## Product datasheet for SC329647

## ZNF 559 (ZNF559) (NM\_001202411) Human Untagged Clone

## **Product data:**

Product Type:	Expression Plasmids
Product Name:	ZNF 559 (ZNF559) (NM_001202411) Human Untagged Clone
Tag:	Tag Free
Symbol:	ZNF559
Synonyms:	NBLA00121
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC329647 representing NM_001202411. Blue=Insert sequence <mark>Red</mark> =Cloning site Green=Tag(s)
	ATGGTGGCTGGGTGGTTGACAAATTACTCTCAGGACTCAGTGACCTTTGAGGATGTGGCTGTGGACTTC ACCCAGGAGGAGTGGACTTTGCTGGATCAAACTCAGAGAAACTTATACAGAGATGTGATGCTGGAGAAAC TATAAGAATCTAGTTGCAGTAGATTGGGAGAGTCATATTAATACCAAATGGTCAGCACCTCAGCAGAAT TTTTTGCAGGGGAAAACATCCAGTGTGGTGGAAATGAATTCAGAGTAA
<b>Restriction Sites:</b>	Sgfl-Mlul
Plasmid Map:	
	OW available



ACCN: Insert Size: NM\_001202411 255 bp



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<b>ORIGENE</b> ZNF 55	59 (ZNF559) (NM_001202411) Human Untagged Clone – SC329647
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 001202411.1</u>
RefSeq Size:	3186 bp
RefSeq ORF:	255 bp
Locus ID:	84527
Cytogenetics:	19p13.2
Protein Families:	Transcription Factors
MW:	9.8 kDa
Gene Summary:	May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (7) includes an additional exon in its 5' UTR, uses a downstream start codon, uses an alternate splice site that causes a frameshift in the 3' coding region, and differs in the 3' UTR, compared to variant 1. The encoded isoform (f) has a shorter N-terminus, a distinct C-terminus, and is overall shorter than isoform a. Variants 6, 7 and 8 all encode isoform f.

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