

Product datasheet for SC329645

OriGene Technologies, Inc.

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ZNF 559 (ZNF559) (NM_001202409) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ZNF 559 (ZNF559) (NM_001202409) Human Untagged Clone

Tag: Tag Free
Symbol: ZNF 559
Synonyms: NBLA00121

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329645 representing NM_001202409.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGCTGGCCATTGGTCTTTCCCTTTGTGATCATCTTTCTCAAGATGTTTTCTGTCTTCATGAGTCAAAA
TTTGAAGAGGAAAGGATGGTGGCTGGGTGGTTGACAAATTACTCTCAGGACTCAGTGACCTTTGAGGAT
GTGGCTGTGGACTTCACCCAGGAGGAGTGGACTTTGCTGGATCAAACTCAGAGAAACTTATACAGAGAT
GTGATGCTGGAGAACTATAAGAATCTAGTTGCAGTAGATTGGGAGAGTCATATTAATACCAAATGGTCA
GCACCTCAGCAGAATTTTTTTGCAGGGGAAAACATCCAGTGTGGTGGAAATGAATTCAGAGTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001202409

Insert Size: 339 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: NM 001202409.1





ZNF 559 (ZNF559) (NM_001202409) Human Untagged Clone - SC329645

RefSeq Size: 3309 bp

 RefSeq ORF:
 339 bp

 Locus ID:
 84527

 Cytogenetics:
 19p13.2

Protein Families: Transcription Factors

MW: 13.1 kDa

Gene Summary: May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) includes an additional segment in its 5' UTR, uses a distinct 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 (e) has distinct N- and C-

termini and is shorter than isoform a.