

Product datasheet for SC334115

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OriGene Technologies, Inc.

ZASP (NM_001289933) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ZASP (NM_001289933) Human Untagged Clone

Kanamycin (25 ug/mL)

Tag: Tag Free Symbol: ZASP

Mammalian Cell Neomycin

Selection:

E. coli Selection:

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC334115 representing NM_001289933.

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

GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

CAGGTGTCGTGTCAGCCGCAATAA

ACGCGTACGCGCCCCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul

ACCN: NM_001289933

Insert Size: 507 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.

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 001289933.1</u>

RefSeq Size: 737 bp RefSeq ORF: 507 bp

Locus ID: 101927655

Cytogenetics: 7q22.1

MW: 17.1 kDa

Gene Summary: DISCONTINUED: This record has been withdrawn by NCBI. This record was defined by

NM_001289933.2, which has insufficient support for representation. There is insufficient

evidence that it is a distinct gene or represents a conserved coding region.