

Product datasheet for SC335057

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Acidic Calponin (CNN3) (NM_001286055) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Acidic Calponin (CNN3) (NM 001286055) Human Untagged Clone

Tag: Tag Free

Symbol: Acidic Calponin

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001286055, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

ACCN: NM_001286055

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).



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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 001286055.1</u>, <u>NP 001272984.1</u>

RefSeq Size: 1974 bp
RefSeq ORF: 852 bp
Locus ID: 1266
UniProt ID: Q15417
Cytogenetics: 1p21.3

Gene Summary: This gene encodes a protein with a markedly acidic C terminus; the basic N-terminus is highly

homologous to the N-terminus of a related gene, CNN1. Members of the CNN gene family all

contain similar tandemly repeated motifs. This encoded protein is associated with the

cytoskeleton but is not involved in contraction. [provided by RefSeq, Jul 2008]

Transcript Variant: This variant (2) lacks an in-frame exon in the central coding region compared to variant 1. The encoded isoform (2) is shorter than isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for

the transcript record were based on transcript alignments.