

Product datasheet for SC335037

DACT2 (NM 001286351) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: DACT2 (NM_001286351) Human Untagged Clone

Tag: Tag Free Symbol: DACT2

Synonyms: bA503C24.7; C6orf116; DAPPER2; DPR2

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >NCBI ORF sequence for NM_001286351, the custom clone sequence may differ by one or

more nucleotides

TGA

Restriction Sites: Sgfl-Mlul

ACCN: NM 001286351

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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DACT2 (NM_001286351) Human Untagged Clone - SC335037

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 001286351.1</u>, <u>NP 001273280.1</u>

 RefSeq Size:
 2043 bp

 RefSeq ORF:
 843 bp

 Locus ID:
 168002

 UniProt ID:
 Q5SW24

 Cytogenetics:
 6q27

Gene Summary: Involved in regulation of intracellular signaling pathways during development. Negatively

regulates the Nodal signaling pathway, possibly by promoting the lysosomal degradation of Nodal receptors, such as TGFBR1. May be involved in control of the morphogenetic behavior of kidney ureteric bud cells by keeping cells epithelial and restraining their mesenchymal character. May play an inhibitory role in the re-epithelialization of skin wounds by attenuating

TGF-beta signaling (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) contains alternate 3' exon structure and it thus differs in the 3' coding region and 3' UTR, compared to variant 1. The encoded isoform (c) has a distinct C-terminus and is shorter than isoform a. 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.