

Product datasheet for SC334556

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CDCA3 (NM_001297603) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: CDCA3 (NM_001297603) Human Untagged Clone

Tag: Tag Free Symbol: CDCA3

Synonyms: GRCC8; TOME-1

Mammalian Cell

Neomycin

Vector:

Selection:

pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF:

>NCBI ORF sequence for NM_001297603, the custom clone sequence may differ by one or

more nucleotides

CCTGTTATCGTGGCCAGAGAGGATGCC<mark>TGA</mark>

Restriction Sites: Sgfl-Mlul

ACCN: NM_001297603

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: <u>NM 001297603.1</u>, <u>NP 001284532.1</u>

 RefSeq Size:
 1625 bp

 RefSeq ORF:
 660 bp

 Locus ID:
 83461

 UniProt ID:
 Q99618

 Cytogenetics:
 12p13.31

Gene Summary: F-box-like protein which is required for entry into mitosis. Acts by participating in E3 ligase

complexes that mediate the ubiquitination and degradation of WEE1 kinase at G2/M phase

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

Transcript Variant: This variant (3) differs in the 3' structure, resulting in a novel 3' coding region and 3' UTR compared to variant 1. The encoded isoform (3) has a distinct C-terminus, and is shorter, compared to 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.