

## **Product datasheet for SC334896**

## OriGene Technologies, Inc.

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## FAM190B (CCSER2) (NM\_001284243) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

**Product Name:** FAM190B (CCSER2) (NM\_001284243) Human Untagged Clone

Tag: Tag Free Symbol: FAM190B

**Synonyms:** bA486O22.1; FAM190B; Gcap14; KIAA1128

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >NCBI ORF sequence for NM\_001284243, the custom clone sequence may differ by one or

more nucleotides

reandacedachare

Sgfl-Mlul

**ACCN:** NM 001284243

**Restriction Sites:** 

**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).





## FAM190B (CCSER2) (NM\_001284243) Human Untagged Clone - SC334896

**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 001284243.1</u>, <u>NP 001271172.1</u>

 RefSeq Size:
 5906 bp

 RefSeq ORF:
 786 bp

 Locus ID:
 54462

 UniProt ID:
 Q9H7U1

 Cytogenetics:
 10q23.1

**Protein Families:** Druggable Genome

**Gene Summary:** Microtubule-binding protein which might play a role in microtubule bundling.

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) differs in the 5' UTR, lacks a portion of the 5' coding region, initiates translation at a downstream start codon, and contains an alternate exon in the 3' coding region, which results in a frameshift, compared to variant 1. The encoded isoform (5) is shorter and has a distinct C-terminus, 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.