

Product datasheet for SC332378

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

EB2 (MAPRE2) (NM_001256420) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: EB2 (MAPRE2) (NM_001256420) Human Untagged Clone

Tag: Tag Free Symbol: EB2

Synonyms: CSCSC2; EB1; EB2; RP1

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC332378 representing NM_001256420.

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

Restriction Sites: Sgfl-Mlul

ACCN: NM_001256420

Insert Size: 825 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.

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: NM 001256420.1

 RefSeq Size:
 4151 bp

 RefSeq ORF:
 825 bp

 Locus ID:
 10982

 UniProt ID:
 Q15555

Cytogenetics: 18q12.1-q12.2

Protein Families: Druggable Genome

MW: 30.7 kDa

Gene Summary: The protein encoded by this gene shares significant homology to the adenomatous polyposis

coli (APC) protein-binding EB1 gene family. This protein is a microtubule-associated protein that is necessary for spindle symmetry during mitosis. It is thought to play a role in the tumorigenesis of colorectal cancers and the proliferative control of normal cells. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Jan 2012] Transcript Variant: This variant (4) lacks an alternate exon in the 5' coding region and uses an alternate start codon, compared to variant 1. The encoded isoform (4) has a distinct N-terminus and 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.