

Product datasheet for SC334191

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BORCS5 (NM_001300742) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: BORCS5 (NM_001300742) Human Untagged Clone

Tag: Tag Free Symbol: BORCS5

Synonyms: LOH1CR12; LOH12CR1

Mammalian Cell

Neomycin

Selection:

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

Fully Sequenced ORF: >NCBI ORF sequence for NM_001300742, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

ACCN: NM 001300742

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 001300742.1</u>, <u>NP 001287671.1</u>

 RefSeq Size:
 1701 bp

 RefSeq ORF:
 534 bp

 Locus ID:
 118426

 UniProt ID:
 Q969J3

 Cytogenetics:
 12p13.2

Gene Summary: As part of the BORC complex may play a role in lysosomes movement and localization at the

cell periphery. Associated with the cytosolic face of lysosomes, the BORC complex may recruit ARL8B and couple lysosomes to microtubule plus-end-directed kinesin motor. Thereby, it may

indirectly play a role in cell spreading and motility.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) contains an alternate 5' terminal exon, resulting in the use of an alternate start codon compared to variant 1. The encoded isoform (2) has a distinct N-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.