

Product datasheet for SC329690

OriGene Technologies, Inc.

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CHURC 1 (CHURC1) (NM_001204064) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: CHURC 1 (CHURC1) (NM_001204064) Human Untagged Clone

Tag: Tag Free Symbol: CHURC 1

Synonyms: C14orf52; chch; My015

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329690 representing NM_001204064.

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

GTTATGCGGCAAAGCCGAAGATACTATCAGTATTCTCCCTGA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001204064

Insert Size: 318 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 001204064.1





CHURC 1 (CHURC1) (NM_001204064) Human Untagged Clone - SC329690

RefSeq Size: 3561 bp

 RefSeq ORF:
 318 bp

 Locus ID:
 91612

 UniProt ID:
 Q8WUH1

Cytogenetics: 14q23.3

Protein Families: Transcription Factors

MW: 12.3 kDa

Gene Summary: Transcriptional activator that mediates FGF signaling during neural development. Plays a role

in the regulation of cell movement (By similarity). Does not bind DNA by itself.

coordinates used for the transcript record were based on transcript alignments.

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) lacks an exon in the coding region, which results in a frameshift, compared to variant 1. The encoded isoform (3) 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