

Product datasheet for SC306825

USH1G (NM_173477) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: USH1G (NM_173477) Human Untagged Clone

Tag: Tag Free
Symbol: USH1G

Synonyms: ANKS4A; SANS

Mammalian Cell Neomycin

Selection:

Vector: pCMV6-Entry (PS100001)

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

OriGene Technologies, Inc.

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Fully Sequenced ORF: >SC306825 representing NM_173477.

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

GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCCGCCGCGGATCGCC

ATGAACGACCAGTACCACCGGGCAGCCCGGGATGGCTACCTGGAGCTCCTCAAGGAGGCCACCCGAAAG GAGCTGAATGCCCCCGACGAGGATGGCATGACCCCCACTCTCTGGGCTGCCTACCATGGCAACCTCGAG TCGCTGCGTCTCATTGTGAGCCGCGGGGGTGACCCGGACAAGTGTGACATCTGGGGCAACACCCCTG TACCTGGACTCCATCGCGGCCAAGCAGAGCAGCCTCAACCCCAAGCTGGTGGGTAAGCTGAAGGACAAG GCCTTCCGCGAGGCGGAGCGCGCATCCGCGAGTGCGCCAAGCTGCAGCGGAGGCACCACGAACGCATG GAGCGGCGATACCGGCGCGAGCTTGCCGAGCGTTCCGACACCCTCAGCTTCTCCAGCCTCACGTCCAGC ACCCTGAGCCGCCGGCTGCAGCATCTGGCGCTGGGCAGCCACCTGCCGTACTCTCAGGCCACGCTGCAC GGCACGGCCAGGGGCAAGACCAAGATGCAGAAGAAGCTGGAGCGGCGCAAGCAGGGCGGCGAAGGCACC ATGTTCGTGCGCCAGGGCACCTACGCCAATCCCAAGGAGTGGGGCCGAGCCCCGCTCCGGGACATGTTC CTCTCGGACGACGACAGCGTCTCCCGTGCCACGCTGGCGACCCTGCCCACTCGGAGGTCAGCACC GACTCAGGCCACGACTCCCTGTTTACCCGCCCCGGCCTGGGCACCATGGTGTTCCGCAGAAATTACTTG AGCAGTGGGCTGCACGGACTGGGCCGCGAGGATGGGGGTCTGGATGGGGTGGGAGCGCCGCGGGGTCGG CTGCAGAGCTCCCCCAGCCTGGACGATGACAGCCTGGGCAGTGCCAACAGCCTGCAGGACCGCAGCTGT GGGGAGGAGCTGCCCTGGGATGAGCTCGATTTAGGCTTGGACGAGGACCTGGAGCCCGAGACTAGCCCG CTGGAGACCTTCCTGGCCTCTCTGCACATGGAGGACTTTGCCGCCCTCCTGCGGCAGGAGAAGATCGAC CTCGAGGCTTTGATGCTGTGCTCTGACCTCGACCTCCGCAGCATCAGCGTCCCACTGGGGCCCCGAAAG **CTATAA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul
ACCN: NM_173477
Insert Size: 1386 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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 173477.4</u>

 RefSeq Size:
 3565 bp

 RefSeq ORF:
 1386 bp

 Locus ID:
 124590

 UniProt ID:
 Q495M9

 Cytogenetics:
 17q25.1

 MW:
 51.5 kDa

Gene Summary: This gene encodes a protein that contains three ankyrin domains, a class I PDZ-binding motif

and a sterile alpha motif. The encoded protein interacts with harmonin, which is associated with Usher syndrome type 1C. This protein plays a role in the development and maintenance of the auditory and visual systems and functions in the cohesion of hair bundles formed by inner ear sensory cells. Mutations in this gene are associated with Usher syndrome type 1G (USH1G). Alternative splicing results in multiple transcript variants. [provided by RefSeq, Sep

2013]

Transcript Variant: This variant (1) represents the longer transcript and encodes the longer 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.