

## **Product datasheet for SC304135**

## VSX1 (NM 014588) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

Product Name: VSX1 (NM\_014588) Human Untagged Clone

Tag: Tag Free Symbol: VSX1

Synonyms: CAASDS; KTCN; KTCN1; PPCD; PPCD1; PPD; RINX

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >SC304135 representing NM\_014588.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGACCGGCCGGGACTCGCTTTCCGACGGGCGCACTAGCAGCAGGGCGCTGGTGCCTGGCGGTTCCCCT AGGGGCTCGCGCCCCCGGGGCTTCGCCATCACGGACCTGCTGGGCTTGGAGGCCGAGCTGCCGGCGCCC GCTGGCCCAGGACAGGGATCTGGCTGCGAGGGTCCGGCAGTCGCCCGTGCCCGGGCCCGGGGCTTGAC GGCTCCAGCCTGGCGCGTGGGGCCCTACCGCTGGGACTCGGCCTCCTCTGTGGCTTCGGCACGCAGCCG CCGGCGGCCGCTCGAGCACCCTGCCTGCTCCTAGCGGACGTGCCGTTCCTGCCGCCCAGGGGCCCCGAG CCCGCTGCCCGCTGGCTCCCAGCCGTCCGCCGCCTCGCCCCAGAAGCGCAGCGCCAGCAGCGTC TCCACGTCCGATGAGGACAGCCAGTCTGAAGACAGGAATGACCTAAAGGCATCCCCCACCTTGGGCAAG AGGAAGAGCGGCGCACAGGACAGTTTTCACTGCTCACCAGCTGGAAGAGGTTGGAGAAGGCATTCAGC ATACAGGTCTGGTTTCAAAACCGCAGGGCCAAATGGCGCAAGCGGGAGAAGCGCTGGGGCGGCAGCAGC GTGATGGCCGAGTACGGGCCTGTACGGGGCCATGGTGCGCCACTGCATCCCGCTGCCAGACTCCGTGCTC AACTCCGCCGAGGGCGGCCTGCTGGGCTCCTGCGCGCCCTGGCTCCTGGGGATGCATAAAAAATCCATG GGGATGATAAGGAAGCCAGGAAGTGAAGATAAGTTGGCAGGACTCTGGGGCTCTGACCACTTCAAAGAA GGTTCTAGCCAGAGTGAGTCAGGATCACAGAGAGGGCTCAGATAAAGTGAGCCCTGAGAATGGCTTGGAA GATGTGGCTATTGACCTCTCCAGCTCTGCCCGGCAGGAGACCAAGAAAGTGCACCCTGGGGCTGGTGCT 

**ACGCGTACGCGGCCGCTC**GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul ACCN: NM 014588



**OriGene Technologies, Inc.** 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com ORIGENE

**Insert Size:** 1098 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 014588.5</u>

 RefSeq Size:
 2216 bp

 RefSeq ORF:
 1098 bp

 Locus ID:
 30813

 UniProt ID:
 Q9NZR4

 Cytogenetics:
 20p11.21

 MW:
 38.4 kDa

**Gene Summary:** The protein encoded by this gene contains a paired-like homeodomain and binds to the core

of the locus control region of the red/green visual pigment gene cluster. The encoded protein may regulate expression of the cone opsin genes early in development. Mutations in this gene can cause posterior polymorphous corneal dystrophy and keratoconus. Alternatively spliced transcript variants encoding different isoforms have been described. [provided by

RefSeq, Jul 2008]

Transcript Variant: This variant (1) encodes the longest isoform (a). 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.