

## **Product datasheet for SC325838**

## VPS37A (NM 001145152) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** VPS37A (NM\_001145152) Human Untagged Clone

Tag: Tag Free Symbol: VPS37A

Synonyms: HCRP1; PQBP2; SPG53

Mammalian Cell

Selection:

Neomycin

Vector: pCMV6-Entry (PS100001)

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

Fully Sequenced ORF: >SC325838 representing NM\_001145152.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGAGCTGGCTTTTTCCCCTGACCAAGAGCGCCTCCTCCTCCGCGGCTGGGTCCCCCGGTGGCCTCACC AGCCTCCAGCAGCAGCAGCAGCGCCTGATCGAGTCCCTCCGGAACTCACACTCCAGATTGCTTCCTCCA CAGTTTCCTCAGGAAAAACCAGTGATCAGTGTTTATCCACCAATACGACATCACTTAATGGATAAACAA GGAGTGTATGTTACCTCTCCATTAGTAAACAATTTTACAATGCACTCAGATCTTGGAAAAATTATTCAG AGTCTGTTGGATGAGTTTTGGAAGAATCCTCCAGTTTTAGCTCCTACTTCAACAGCATTTCCTTATCTA TACAGTAACCCAAGTGGGATGTCTCCTTATGCTTCTCAGGGTTTTCCATTTCTTCCTCCATATCCTCCA CAAGAAGCAAACAGGAGTATCACTTCTTTATCTGTTGCTGACACTGTTTCTTCTTCAACAACAAGTCAT ACCACAGCCAAGCCTGCCGCTCCTTCATTTGGTGTCCTTTCAAATCTGCCATTACCCATTCCCACAGTG GATGCTTCAATACCGACAAGCCAAAATGGTTTTGGGTACAAGATGCCAGATGTCCCTGATGCATTTCCA TTTCTGACTTTGCCTCAACTAAAACAAATTATTACCGACAAAGATGACTTAGTAAAAAAGTATTGAGGAA CTAGCAAGAAAAATCTCCTTTTGGAGCCCAGCTTGGAAGCCAAAAGACAAACTGTTTTAGATAAGTAT GAATTACTTACACAGATGAAGTCCACTTTCGAAAAGAAGATGCAAAGGCAGCATGAACTTAGTGAGAGC TGTAGTGCAAGTGCCCTTCAGGCAAGATTGAAAGTAGCTGCACATGAAGCTGAGGAAGAATCTGATAAT ATTGCAGAAGACTTCTTGGAGGGAAAGATGGAAATAGATGATTTTCTCAGTAGCTTCATGGAAAAGAGA ACAATTTGCCACTGTAGAAGAGCCAAGGAAGAGAAACTTCAGCAGGCGATAGCAATGCACAGCCAATTT

CATGCTCCACTATAG

 ${\color{blue} \textbf{ACGCGTACGCGGCCGCTC} \textbf{GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT} }$ 

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul



**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

## VPS37A (NM\_001145152) Human Untagged Clone - SC325838

**ACCN:** NM\_001145152

**Insert Size:** 1119 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 001145152.1</u>

 RefSeq Size:
 4510 bp

 RefSeq ORF:
 1119 bp

 Locus ID:
 137492

 UniProt ID:
 Q8NEZ2

 Cytogenetics:
 8p22

**Protein Pathways:** Endocytosis

**MW:** 41.5 kDa

**Gene Summary:** This gene belongs to the VPS37 family, and encodes a component of the ESCRT-I (endosomal

sorting complex required for transport I) protein complex, required for the sorting of ubiquitinated transmembrane proteins into internal vesicles of multivesicular bodies. Expression of this gene is downregulated in hepatocellular carcinoma, and mutations in this gene are associated with autosomal recessive spastic paraplegia-53. A related pseudogene has been identified on chromosome 5. Alternatively spliced transcript variants have been

found for this gene. [provided by RefSeq, Dec 2012]

Transcript Variant: This variant (2) lacks an exon in the coding region compared to variant 1. The encoded isoform (2) is shorter but has the same N- and C-termini 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.