

Product datasheet for SC318821

OriGene Technologies, Inc.

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Aspartate beta hydroxylase (ASPH) (NM_032468) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Aspartate beta hydroxylase (ASPH) (NM 032468) Human Untagged Clone

Tag: Tag Free Symbol: ASPH

Synonyms: AAH; BAH; CASQ2BP1; FDLAB; HAAH; JCTN; junctin

Mammalian Cell Neomycin

Selection:

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

Fully Sequenced ORF: >SC318821 representing NM_032468.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

TAA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

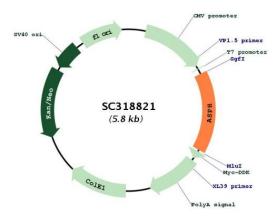
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul





Plasmid Map:



ACCN: NM_032468

Insert Size: 900 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

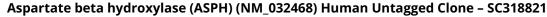
at the bottom.

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: NM 032468.4

RefSeq Size: 3642 bp RefSeq ORF: 900 bp





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UniProt ID: Q12797
Cytogenetics: 8q12.3

Protein Families: Druggable Genome, Transmembrane

444

MW: 33.8 kDa

Gene Summary: This gene is thought to play an important role in calcium homeostasis. The gene is expressed

from two promoters and undergoes extensive alternative splicing. The encoded set of proteins share varying amounts of overlap near their N-termini but have substantial variations in their C-terminal domains resulting in distinct functional properties. The longest isoforms (a and f) include a C-terminal Aspartyl/Asparaginyl beta-hydroxylase domain that hydroxylates aspartic acid or asparagine residues in the epidermal growth factor (EGF)-like domains of some proteins, including protein C, coagulation factors VII, IX, and X, and the complement factors C1R and C1S. Other isoforms differ primarily in the C-terminal sequence and lack the hydroxylase domain, and some have been localized to the endoplasmic and sarcoplasmic reticulum. Some of these isoforms are found in complexes with calsequestrin, triadin, and the ryanodine receptor, and have been shown to regulate calcium release from the sarcoplasmic reticulum. Some isoforms have been implicated in metastasis. [provided by RefSeq, Sep 2009]

Transcript Variant: This variant (2) represents use of an alternate promoter and 5' UTR, uses a distinct start codon, includes an alternate in-frame exon, and uses a distinct 3' splice pattern that lacks many coding exons, compared to variant 1. The encoded isoform (b) shares residues with the central region of isoform a. This isoform lacks the catalytic domain and enzymatic function of isoform a. Isoform b has also been referred to as junctate and is located in both the endoplasmic and sarcoplasmic reticulum. Calcium binding properties of this isoform suggest an active role in the calcium storage and release process in the endoplasmic reticulum. This variant is expressed in heart, brain, pancreas, placenta, lung, liver, kidney, and skeletal muscle tissues. 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.