

Product datasheet for SC333422

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OriGene Technologies, Inc.

ODAPH (NM 001257072) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ODAPH (NM_001257072) Human Untagged Clone

Tag: Tag Free Symbol: ODAPH

Synonyms: Al2A4; C4orf26

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >SC333422 representing NM_001257072.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

TATTTCCCCAGAAGAAGACTCCAGAGAGGAAGCTCATCTGAGGAAAGCTGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul

ACCN: NM_001257072

Insert Size: 258 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: <u>NM 001257072.1</u>

 RefSeq Size:
 1747 bp

 RefSeq ORF:
 258 bp

 Locus ID:
 152816

 UniProt ID:
 Q17RF5

 Cytogenetics:
 4q21.1

 MW:
 9.7 kDa

Gene Summary: Dental enamel forms the outer cap of teeth and is the hardest substance found in

vertebrates. This gene is thought to encode an extracellular matrix acidic phosphoprotein that has a function in enamel mineralization during amelogenesis. Mutations in this gene are associated with recessive hypomineralized amelogenesis imperfecta. [provided by RefSeq,

Oct 2012]

Transcript Variant: This variant (3) lacks an alternate exon that results in a frameshift in the central and subsequent coding region, and it lacks a segment in the 3' coding region,

compared to variant 1. The encoded isoform (3) has a distinct C-terminus and is shorter than

isoform 1.