

Product datasheet for MC215077

Cfap74 (NM 001166029) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Cfap74 (NM_001166029) Mouse Untagged Clone

Tag: Tag Free Symbol: Cfap74

Synonyms: 2010015L04Rik; A530045M11

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

Cell Selection: Neomycin

Fully Sequenced ORF: >MC215077 representing NM_001166029

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

AGCCGAGATCGCCACAGAGAAAGAAGCCAACAACATGTAG

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001166029

Insert Size: 390 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).



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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 001166029.1</u>, <u>NP 001159501.1</u>

RefSeq Size: 2451 bp
RefSeq ORF: 390 bp
Locus ID: 544678
Cytogenetics: 4 E2

Gene Summary: As part of the central apparatus of the cilium axoneme may play a role in cilium movement.

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) differs in the presence and absence of exons in its 3' coding

region and 3' UTR, compared to variant 1, resulting in an isoform (2) with a shorter C-

terminus, compared to isoform 1.