

## **Product datasheet for MC206782**

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## Rab22a (BC006596) Mouse Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

Product Name: Rab22a (BC006596) Mouse Untagged Clone

Tag: Tag Free Symbol: Rab22a

**Synonyms:** 3732413A17Rik; Al662177; AW319644; AW550514; E130120E14Rik; Rab22

**Mammalian Cell** 

Selection:

Neomycin

Vector: PCMV6-Kan/Neo (PCMV6KN)

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

Fully Sequenced ORF: >NCBI ORF sequence for BC006596, the custom clone sequence may differ by one or more

nucleotides

Restriction Sites: Rsrll-Notl
ACCN: BC006596
Insert Size: 585 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>BC006596</u>, <u>AAH06596</u>

RefSeq Size:1911 bpRefSeq ORF:585 bpLocus ID:19334

Cytogenetics: 2 97.26 cM

**Gene Summary:** Plays a role in endocytosis and intracellular protein transport (PubMed:19759177,

PubMed:27718357). Mediates trafficking of TF from early endosomes to recycling endosomes. Required for NGF-mediated endocytosis of NTRK1, and subsequent neurite outgrowth. Binds GTP and GDP and has low GTPase activity. Alternates between a GTP-bound active form and

a GDP-bound inactive form (By similarity).[UniProtKB/Swiss-Prot Function]