

## **Product datasheet for MC204689**

## Magohb (NM\_025564) Mouse Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Product Name: Magohb (NM 025564) Mouse Untagged Clone

Tag: Tag Free
Symbol: Magohb

Synonyms: 2010012C16Rik; Mago; Magoh; Magoh-rs1

**Mammalian Cell** 

Selection:

Neomycin

Vector: PCMV6-Kan/Neo (PCMV6KN)

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

Fully Sequenced ORF: >BC027505

**Restriction Sites:** EcoRI-NotI ACCN: NM\_025564

**Insert Size:** 441 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>BC027505</u>, <u>AAH27505</u>

RefSeq Size: 632 bp
RefSeq ORF: 441 bp
Locus ID: 66441
UniProt ID: Q9CQL1

**Cytogenetics:** 6

**Gene Summary:** Required for pre-mRNA splicing as component of the spliceosome. Plays a redundant role

with MAGOH in the exon junction complex and in the nonsense-mediated decay (NMD)

pathway.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) encodes the longer isoform (1). CCDS Note: The coding region has been updated to extend the N-terminus to one that is also supported by available conservation data. The use of an alternative upstream start codon would result in a protein

that is 2 aa longer.