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## Product datasheet for MR215685

## Fabp6 (NM_008375) Mouse Tagged ORF Clone

## Product data:

Product Type:
Product Name:

## Tag:

Symbol:
Synonyms:
Mammalian Cell
Selection:
Vector:
E. coli Selection:

ORF Nucleotide
Sequence:

Expression Plasmids
Fabp6 (NM_008375) Mouse Tagged ORF Clone
Myc-DDK
Fabp6
GT; I; I-1; I-15P; I-B; I-BABP; IL; ILBP; ILBP3; Illbp
Neomycin
pCMV6-Entry (PS100001)
Kanamycin ( $25 \mathrm{ug} / \mathrm{mL}$ )
>MR215685 representing NM_008375
Red=Cloning site Blue=ORF Green=Tags(s)

OriGene Technologies, Inc.
9620 Medical Center Drive, Ste 200
Rockville, MD 20850, US
Phone: +1-888-267-4436
https://www.origene.com techsupport@origene.com EU: info-de@origene.com
CN: techsupport@origene.cn

## TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC GCCGCGATCGCC

ATGGCCTTCAGTGGCAAATATGAATTTGAGAGTGAGAAGAATTACGATGAGTTCATGAAGCGCCTGGGTC TTCCAGGAGACGTGATTGAAAGGGGACGTAACTTCAAGATCATCACAGAGGTCCAGCAGGACGGACAGGA CTTCACCTGGTCCCAGTCTTACTCTGGGGGCAACATTATGAGCAACAAGTTCACCATTGGCAAAGAATGT GAAATGCAGACCATGGGGGGCAAGAAGTTCAAGGCTACCGTGAAGATGGAGGGTGGCAAGGTGGTGGCAG AGTTCCCCAACTATCACCAGACTTCGGAGGTCGTGGGTGACAAGTTGGTGGAGATCTCCACCATCGGGGA TGTGACCTATGAGCGCGTAAGCAAGAGGCTGGCT

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR215685 representing NM_008375
Red=Cloning site Green=Tags(s)

MAFSGKYEFESEKNYDEFMKRLGLPGDVIERGRNFKIITEVQQDGQDFTWSQSYSGGNIMSNKFTIGKEC EMQTMGGKKFKATVKMEGGKVVAEFPNYHQTSEVVGDKLVEISTIGDVTYERVSKRLA

TRTRPLEQKLISEEDLAANDILDYKDDDDKV
Chromatograms:
Restriction Sites:
https://cdn.origene.com/chromatograms/mm9042 a04.zip
Sgfl-Mlul

## Cloning Scheme:

Cloning sites used for ORF Shuttling:


 $\begin{array}{llllllllllllllllll}\mathrm{D} & \mathrm{L} & \mathrm{A} & \mathrm{A} & \mathrm{N} & \mathrm{D} & \mathrm{I} & \mathrm{L} & \mathrm{D} & \mathrm{Y} & \mathrm{K} & \mathrm{D} & \mathrm{D} & \mathrm{D} & \mathrm{D} & \mathrm{K} & \mathrm{V} & \text { stop }\end{array}$

* The last codon before the Stop codon of the ORF


## ACCN:

ORF Size:
OTI Disclaimer:

## OTI Annotation:

Components:

Reconstitution Method: 1. Centrifuge at $5,000 \mathrm{xg}$ for 5 min .
NM_008375
384 bp
The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. More info
This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
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).
2. Carefully open the tube and add 100 ul 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 5000 xg ) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at $-20^{\circ} \mathrm{C}$. The DNA is stable for at least one year from date of shipping when stored at $-20^{\circ} \mathrm{C}$.

## RefSeq:

NM 008375.2, NP 032401.1
RefSeq Size:
387 bp
RefSeq ORF: $\quad 387 \mathrm{bp}$
Locus ID: 16204
UniProt ID: P51162

Cytogenetics:

## MW:

Gene Summary:
1125.81 cM
14.9 kDa

The protein encoded by this gene is part of the fatty acid binding protein family (FABP). FABPs are a family of small, highly conserved, cytoplasmic proteins that bind long-chain fatty acids and other hydrophobic ligands and participate in fatty acid uptake, transport, and metabolism. This protein functions within the ileum, the distal $25-30 \%$ of the small intestine, and plays a role in enterohepatic circulation of bile acids and cholesterol homeostasis. In humans, it has been reported that polymorphisms in FABP6 confer a protective effect in obese individuals from developing type 2 diabetes. In mice deficiency of this gene affects bile acid metabolism in a gender-specific manner and was reported to be required for efficient apical to basolateral transport of conjugated bile acids. [provided by RefSeq, Jan 2013]

## Product images:



