

Product datasheet for MC226390

Lgals8 (NM_001291060) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Lgals8 (NM_001291060) Mouse Untagged Clone

Tag: Tag Free
Symbol: Lgals8

Synonyms: 1200015E08Rik; Al326142; D13Ertd524e; Lgals-8

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

Cell Selection: Neomycin

Fully Sequenced ORF: >MC226390 representing NM_001291060

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ATGCCCTTCAGAAAAGAAAAGTCCTTTGAGATCGTGTTCATGGTGCTCAAGAACAAATTCCAGGTGGCTG
TGAACGGAAGGCATGTTCTGCTGTACGCCCACAGGATCAGCCCGGAGCAGATCGACACAGTGGGCATCTA
CGGCAAAGTGAACATCCACTCCATCGGGTTCAGATTCAGCTCGGATTTACAGAGTATGGAAACATCTGCT
CTGGGACTGACACAGATAAACAGAGAGAATATACAAAAGCCAGGCAAGCTCCAGCTGAGCCTGCCATTTG
AAGCAAGGTTGAATGCCTCCATGGGTCCTGGACGAACCGTTGTCATTAAAAGGGGAAGTGAACACCAATGC
CCGAAGCTTTAATGTTGACCTAGTGGCAGGAAAAACAAGGGATATCGCTCTGCACTTGAACCCACGCCTC
AATGTGAAAGCATTTGTAAGAAATTCCTTTCTTCAGGATGCCTGGGGAGAAGAGGAGAAAATATTACCT
GCTTCCCATTTAGTTCTGGGATGTACTTTTGAGATGATAATCTACTGTGATGTCCGGGAATTCAAGGTTGC
TATAAATGGTGTGCACAGCCTGGAGTACAAACACAGATTTAAAGACCTAAGCAGTATTGATACACTATCA
GTCGATGGTGATATCCGTTTGCTGGATGTAAGGAGCTGGTAG

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001291060

Insert Size: 672 bp



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

Lgals8 (NM_001291060) Mouse Untagged Clone - MC226390

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).

OTI Annotation: Clone contains native stop codon, and expresses the complete ORF without any c-terminal

tag.

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 001291060.1</u>, <u>NP 001277989.1</u>

RefSeq Size:2664 bpRefSeq ORF:672 bpLocus ID:56048

Cytogenetics: 13 4.64 cM

Gene Summary: Beta-galactoside-binding lectin that acts as a sensor of membrane damage caused by

infection and restricts the proliferation of infecting pathogens by targeting them for autophagy. Detects membrane rupture by binding beta-galactoside ligands located on the lumenal side of the endosome membrane; these ligands becoming exposed to the cytoplasm

following rupture. Restricts infection by initiating autophagy via interaction with

CALCOCO2/NDP52. Required to restrict infection of bacterial invasion such as S.typhimurium. Also required to restrict infection of Picornaviridae viruses. Has a marked preference for 3'-O-

sialylated and 3'-O-sulfated glycans.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) differs in the 5' UTR and 5' coding region, initiates translation at a downstream start codon and uses an alternate in-frame splice site in the 5' coding region, compared to variant 1. Variants 4 and 5 encode the same isoform (3), which is shorter than isoform 2. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript

alignments.