

Product datasheet for SC201846

OriGene Technologies, Inc.

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Galectin 3 (LGALS3) (NM_002306) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Galectin 3 (LGALS3) (NM_002306) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: LGALS3

Synonyms: CBP35; GAL3; GALBP; GALIG; L31; LGALS2; MAC2

ACCN: NM_002306

Insert Size: 182 bp

Insert Sequence: >SC201846 3'UTR clone of NM_002306

The sequence shown below is from the reference sequence of NM_002306. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGTCATCTACTTAATAAATATTACAGTGAATTACCTGTCTCAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeg: NM 002306.4





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Summary:

This gene encodes a member of the galectin family of carbohydrate binding proteins. Members of this protein family have an affinity for beta-galactosides. The encoded protein is characterized by an N-terminal proline-rich tandem repeat domain and a single C-terminal carbohydrate recognition domain. This protein can self-associate through the N-terminal domain allowing it to bind to multivalent saccharide ligands. This protein localizes to the extracellular matrix, the cytoplasm and the nucleus. This protein plays a role in numerous cellular functions including apoptosis, innate immunity, cell adhesion and T-cell regulation. The protein exhibits antimicrobial activity against bacteria and fungi. Alternate splicing results in multiple transcript variants.[provided by RefSeq, Oct 2014]

Locus ID: 3958

MW: 7