

## Product datasheet for **MC226275**

### Galt (NM\_001302511) Mouse Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Galt (NM\_001302511) Mouse Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Galt  
**Synonyms:** AW553376  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >MC226275 representing NM\_001302511  
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**GCGATCGCC**

ATGATGGGCTGTTCTAACCCCATCCCCACTGCCAGGTTGGGCTAGCAGCTTCCTGCCAGATATCGCCC  
AGCGTGAAGAGCGATCCCAGCAGACCTATCACAGCCAGCATGGAAAACCTTTGTTATTGGAATATGGTCA  
CCAAGAGCTCCTCAGGAAGGAACGTCTGGTCTAACCAGTGAGCACTGGATAGTTCTGGTCCCCTTCTGG  
GCAGTGTGGCCTTCCAGACACTTCTGCTGCCCGGGCGGCACGTGCGGGCGCTACCTGAGCTGAACCCCG  
CTGAGCGTGATGATCTCGCTCCATCATGAAGAAGCTCTTGACCAAGTACGACAATCTATTTGAGACATC  
CTTTCCCTACTCCATGGGCTGGCATGGGGCTCCCACGGGATTAAGACTGGAGCCACCTGTGACCACTGG  
CAGCTCCACGCCCACTACTACCCCACTTCTGCGATCCGCAACTGTCCGGAAGTTCATGGTTGGCTATG  
AAATGCTTGCCAGGCCAGCGTGACCTCACTCCGAACAGGCCGAGAAAGATTAAGGGCGCTTCCCGA  
GGTACACTATTGCCTGGCGCAGAAAGACAAGGAAACGGCAGCCATTGCT**TGA**

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-MluI  
**ACCN:** NM\_001302511  
**Insert Size:** 612 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).



[View online »](#)

<b>OTI Annotation:</b>	Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_001302511.1</a> , <a href="#">NP_001289440.1</a>
<b>RefSeq Size:</b>	1544 bp
<b>RefSeq ORF:</b>	612 bp
<b>Locus ID:</b>	14430
<b>Cytogenetics:</b>	4 22.07 cM
<b>Gene Summary:</b>	<p>The protein encoded by this gene is the second enzyme in the Leloir pathway, the metabolic pathway for D-galactose catabolism. It catalyzes the conversion of galactose-1-phosphate and uridine diphosphate-glucose to glucose-1-phosphate and uridine diphosphate galactose. Deficiency of this enzyme causes the genetic metabolic disorder galactosemia. Mice lacking this protein accumulate high levels of galactose and galactose-1 phosphate but are viable and fertile. This protein is negatively regulated through signaling by the polypeptide hormone prolactin, specifically via the short isoform of the prolactin receptor and the transcription factor Forkhead box O3. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Oct 2014]</p> <p>Transcript Variant: This variant (2) uses an alternate exon structure at the 5' end and utilizes a downstream start codon compared to variant 1. The resulting isoform (2) has a shorter N-terminus compared to isoform 1. Variants 2, 7, 8, and 9 all encode the same isoform (2).</p> <p>Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.</p>