

Product datasheet for **MG210487**

Large1 (NM_010687) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Large1 (NM_010687) Mouse Tagged ORF Clone
Tag:	TurboGFP
Symbol:	Large1
Synonyms:	BPFD#36; enr; fg; froggy; Gyltl1a; Large; Mbp-1; Mbp1; myd
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)



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ORF Nucleotide
Sequence:

>MG210487 representing NM_010687
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGCC**

ATGCTGGGAATCTGCAGAGGGAGAAGAAATTCCTGGCTGCCTCTTTGACTCTACTCTGCATACCGCCA
 TCACCTGGATTTACTTGTGGTGGGAGCTTTGAAGATGGGAAACCTGTGTCTGTCTCCCTGGAGTC
 CCAAGCCACAGTCCCGGTACACAGCCTCCAGCCAACGGGAGCGGGAGAGCCTGGAAGTGCCTGTTTGA
 GAAGTTGAGGAGGAAAACCGTGTCTGCGCAGGCAGCTCAGCCTGGCCAGGGTCAAGAGCCTGCTCACC
 ACCGTGGGAACCACTCCAAGACCTACTCCATGGAGGAGGGACAGGGGACAGTGAGAATCTGCGAGCTGG
 CATTGTGGCAGGTAACAGCTCTGAGTGTGGCAGCAGCCAGCTGTGAAAAAGTGTGAGACCATCCATGTG
 GCCATCGTCTGTCCGGATAACAATGCCAGCAGGGACGTTGTCACCCTGGTAAATCTGTCCTTTTCATA
 GCGGAACCTCTGCACTCCACCTCATTGCTGATTCTATTGCTGAGCAGATCCTGGCTACACTCTTCCA
 GACCTGGATGGTACCTGCAGTGCAGTAGACTTCTATAATGCTGATGAGCTCAAGTCTGAAGTGTCTGG
 ATTCACCAACAAACATTACTCTGGAATTTATGGCCTGATGAAGCTGGTTCTGACCAAAACTCTGCCTGCCA
 ACCTGGAGAGAGTCAATTGCTAGACACCGACATCACCTTTGCAACAGACATTGCAGAGCTGTGGCCGT
 GTTCCATAAGTTCAAAGGGCAACAGGTCCTGGGCTCGGTAGAGAACCAGAGTGACTGGTATCTTGAAAA
 CTCTGGAAAAACCACTCCATGGCCAGCTCTTGAAGGGGCTATAACACAGGAGTCATACTACTGCTTC
 TGGACAAGCTTCGGAAGATGAAATGGGAGCAGATGTGGAGACTGACAGCAGAGAGGGAACACATGGGCAT
 GTTGTCTACATCCTTGGCTGACCAGGATATTTCAATGCTGTTATCAAACAAAACCCCTTTCTGGTGTAT
 CAGCTTCCCTGCTTCTGGAATGTGCAGCTGTGAGACCACCCGCTCTGAGCAGTGTACAGAGATGTGT
 CTGATTTAAAGGTCATCCACTGGAACCTCCCAAGAAGCTCCGTGTGAAGAACAACATGTGGAATCTT
 CCGTAACCTCTACCTGACCTTCTTGGAGTATGATGGCAACCTCCTGCGCGGGGAGCTGTTTGGCTGCCCC
 AGTGAGACTGATGTCAACAATGAAAATCTTCAAGCAGCTGTCTGAGCTGGATGAGGATAAATTGTGCT
 ATGAGTTCGCGGAGAGCGCTTCACTGTCCACAGAACCACCTGTACTTCTGCACTACGAGTTTGAAGCC
 TTCTGCTGACAACACAGACGTTACCCTTGTGCTCAGCTCTCCATGGACAGACTACAGATGCTGGAGGCC
 ATCTGTAAACACTGGGAGGGACCCATCAGCCTGGCCCTTACCTGTGAGATGCAGAAGCCCAGCAGTTCC
 TCCGCTATGCCAGGGATCCGAGGTGCTCATGAGCCGGCAGAACGTGGGCTACCACATCGTGTACAAGGA
 GGGTCAGTTCTATCCTGTCAATCTTCTGCGAAACGTGGCTATGAAGCACATCAGCACACCCTACATGTT
 TTATCTGACATTGACTTCTGCCTATGTATGGGCTCTATGAGTACCTCAGGAAATCTGTCATCCAGCTTG
 ACCTTGCCAATACCAAGAAAGCAATGATTGTCCCAGCATTGAAACGCTGCGCTACAGACTCTTTTCCC
 CAAGTCAAAGCAGAGCTGCTGTCAATGTTGGACATGGGGACCCTATTCACATTCAGGTATCACGTATGG
 ACTAAAGGCCATGCACCCACAACTTTGCCAAGTGGCGGACTGCCACCACACCTTACCAGTTGAGTGGG
 AGGCTGATTTTGAAGCCATATGTTGTTGTGAGACGAGACTGCCCTGAGTATGATCGGAGATTTGTGGGCTT
 TGGCTGGAACAAAGTGGCTCACATCATGGAATTGGATGCACAGGAGTATGAGTTCACCGTGTGCCCAAT
 GCCTACATGATCCACATGCCCATGCCCTAGTTTCGACATCACTAAGTCCGTTCCAACAAGCAATACC
 GCATCTGTCTCAAACCTGAAGGAAGAGTTTCAACAAGACATGTCCCGACGATATGGCTTTGCTGCCTT
 GGAATATCTCACAGCTGAGAACAACAGC

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence: >MG210487 representing NM_010687
 Red=Cloning site Green=Tags(s)

MLGICRGRRKFLAASLTLLCIPAITWIYLFAGSFEDGKPVSLSPLESQAHSPTYASSQRERESLEVRVR
 EVEEENRALRRQLSLAQGQSPAHHHRGNHSTYSMEEGTGDSENLRAGIVAGNSSECGQPAVEKCEIIV
 AIVCAGYNASRDVVTLVKSVLFHRRNPLHFHLLIADSLAEQILATLFTQWMPAVRVDFYNADLKVSEVSW
 IPNKHYSGIYGLMKLVLTKTLPANLERVIVLDTDTITFADIAELWAVFHKFKGQQVLSGVENQSDWYLG
 LWKNHRPWPALGRGYNTGVILLLLDKLRKMKWEQMWRLTAEREHMGLSTSLADQDIFNAVIKQNPFLVY
 QLPCFVNVQLSDHTRSEQCVRDVSCLKVHWNSPKKLRVKNKHVEFFRNLYLTFLEYDGNLLRRELFGCP
 SETDVNNENLQKQLSELDENLCYEFRRERFTVHRTHLYFLHYEFEPSADNTDVTLVLAQLSMDRLQMLEA
 ICKHWEGPISLALYLSDAEAQQFLRYAQGSEVLMRQNVGYHIVYKEGQFYPVNLNRNVAMKHISTPYMF
 LSDIDFLPMYGLYEYLRKSVIQLDLANTKKAMIVPAFETLRYRLSFPKSKAELLSMLDMGTLFTFRYHVW
 TKGHAPTNAKAWRTATTPYQVEWADFEPYVVVRRDCPEYDRRFVGFGWNVVAHIMELDAQEYFTVLPN
 AYMIMPHAPSFDITKFRSNKQYRICLTKLKEEFQQDMSRRYGFAALEYLTAENNS

TRTRPLE - GFP Tag - V

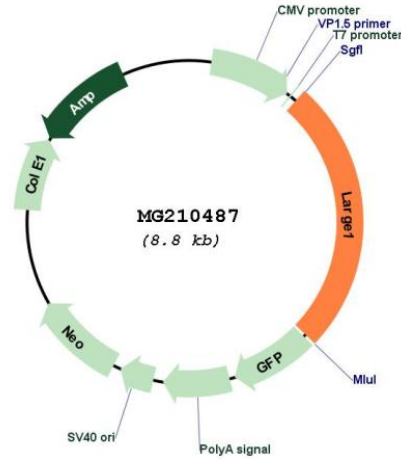
Restriction Sites: SgfI-MluI

Cloning Scheme:

Cloning sites used for ORF Shutting:



Plasmid Map:



ACCN: NM_010687

ORF Size: 2268 bp

OTI Disclaimer: 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](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

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: [NM_010687.1](#), [NP_034817.1](#)

RefSeq Size: 3669 bp

RefSeq ORF: 2271 bp

Locus ID: 16795

UniProt ID: [Q9Z1M7](#)

Cytogenetics: 8 35.08 cM

Gene Summary: Bifunctional glycosyltransferase with both xylosyltransferase and beta-1,3-glucuronyltransferase activities involved in the biosynthesis of the phosphorylated O-mannosyl trisaccharide (N-acetylgalactosamine-beta-3-N-acetylglucosamine-beta-4-(phosphate-6-)mannose), a carbohydrate structure present in alpha-dystroglycan (DAG1) (PubMed:23125099, PubMed:23135544). Phosphorylated O-mannosyl trisaccharid is required for binding laminin G-like domain-containing extracellular proteins with high affinity and plays a key role in skeletal muscle function and regeneration (PubMed:15184894, PubMed:24132234). LARGE elongates the glucuronyl-beta-1,4-xylose-beta disaccharide primer structure initiated by B3GNT1/B4GAT1 by adding repeating units [-3-Xylose-alpha-1,3-GlcA-beta-1-] to produce a heteropolysaccharide (By similarity).[UniProtKB/Swiss-Prot Function]