

Product datasheet for **MC205891**

Pigv (NM_178698) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Pigv (NM_178698) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Pigv
Synonyms:	B330013B03; D430024F16Rik
Mammalian Cell Selection:	Neomycin
Vector:	PCMV6-Kan/Neo (PCMV6KN)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >BC055060
 GTTGGAGCTCCGCGGGGGTGGTCTGGACGCGGGGCGGGAGGATCCGGCGCCAGCTCTGCAACTCTGGA
 GTCCTCCGGGGAGGGAAGCTGTCAAACGCGAGCCGAGAGAAATTAGGCGGAGGCTGTAGAGAAGCGAGGA
 GCCCCGCCCTCCGCCATAGCAGCCTCCGGCGCCGCTTTCCTTTGCAGCCTCCGGCCCTCGTCTGCTT
 CCGGCCCGTTCCTGGCCCGCCCGCGATTCCCCGGGACGAAGCCCTGGGCCGTGGCTCCCTGGGA
 ATTTAGAAGCCGGAGGAAGCTCAGTCTGGTGGTGGAGAGAGAGGATGGGACTCCTGGACCCATCCAGAA
 AGAGGTGCTGAGATTGCGAGTAACTGCCGCATCCTGACTCTAGTGCTACAGGCACTTCAACCTCATC
 ATCCCAGATCACCACGCAGATGCCTTTTGCCTCCCGCCTCGCCCTCGGGCTCTGCGGATCAGCTTG
 TGAAGGTCTTCTGGGTGGTCTGTCTCGATGGGATGCTGAACACTTCTGTTTATCGCTGAGCACGGCTA
 CCTTTATGAGCACAACCTTGCCTTCTTCCCTGGCTTTCCTTGGCCCTGCTGATGGGGACTGAGCTGCTG
 AGACCCTTGACAGGCCTTGTAGCCAGCGCAGTTGCCTGCTGGTCTCAGTAGCACTTCTCAACTGCTGT
 TCTCCGTGCTGGTGCAGTGGCACTCCATGACTTGGGTTGTCTGGTTCTGCACTGTCTCCGCGAGCCCT
 GTGTGCAGCGTCTTCTGCATCAGTCTGCCAACGTTTTCTGGCAGCCGGCTACTCAGAGGCTTTG
 TTTGCCTTCTGACGTTACGCGCATGGGGCAGCTGGAGAGGGCCGTGGCTGGGCTAGTGGGCTGCTCT
 TTGCTCTTGTCTGGGTGCGCTCCAATGGGCTGGTGGCTTGGCTTCTCTGCACTCCAGTGCAG
 AGGCTTTTGTTCCTCTCTGGCGGTGCTGAGCCCTGGAAGCCCTCGTGAAGCTGATGGCCTCTGTGTGC
 TTGTCCGTGCTCATAGTCAGCCTTCCCTTCGCCCTCTTTCAGTATCGTGCCTACATCCAGTTCTGCTCAC
 CAGGCTCAGCCCCCTCCATCCCTGAGCCCTTGCTGCAGCTAGTCCGACAAAGGCTACCGGCTTGACAGG
 AGAAAAATGCACCCCCATGGTGCTCCTGGGACCTTCTCTAATATAACAATAATTCAGGATGTCTACTGG
 AATGTTGGTCTTCTGCGATACTATGAGCTCAAACAGGTGCCCAATTTCTACTAGCTACACCAGTGACAG
 TACTGGTTGTGTGGGCAACCTGGACATATGTGACCACCCACCCTTGGCTGCGCTTACACTTGGGCTGCA
 AAGAACCAAGGACAGGGAGAACC CGGAGAAGCCCCACCGTGGCTTCTGAGTCCGAAGGTGTTTGTGTAC
 CTGGTCCACGCTGCAGCACTGCTGGTCTTTGGAGGTCTGTGCATGCATGCCAGTTCTCACGCGGTTTT
 TGGCCTTCCACTCCTATCATGTACTGGTTTCCAGCTCACCTGCTTTCAGGATCAAGAACCCTGCTGAG
 GTGTGTGGACACAGACCTGGGAAGCTTCTCAGGAGAAGTCCCCACCAGGACAGAAAGGCCCCAGAAAC
 TGTCTCATGAAACTCTTCTACAATTGGAAGAGATGCTCTCCAGTCAAGATGCGTTCTGGTCTACTTCC
 TGACCTACTGGCTTCTGGGACTAATTCTCCATTGCAACTTCTGCCTTGGACTTGACCTGGAGTGTCAAG
 GGTGAAAGCTCTAAGACGACACCCTGGCTGGGACAGCCCGCACACTCCGGAGAGCACTGTGCAGTAGGA
 TCTCCGGTCTCTTGGGGCCAGCTAAGAAGGGAAAAATGGGAGCACAGGCGCCCTTCTCATCCTGGGC
 GGGGCAGGTGTAGGGGAGTAACCTCAGTGCTCAGTGCAGGGTCTCAGATCTCCCTAGTCCACAAGTACT
 AGGCAATCTTAAACCTACTACTGATCCAGTGTA AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA AAAAAAAAAAAAAAAG

Restriction Sites: RsrII-NotI

ACCN: NM_178698

Insert Size: 1482 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).

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: [BC055060](#), [AAH55060](#)

RefSeq Size: 2115 bp

RefSeq ORF: 1482 bp

Locus ID: 230801

UniProt ID: [Q7TPN3](#)

Cytogenetics: 4 D2.3

Gene Summary: Alpha-1,6-mannosyltransferase involved in glycosylphosphatidylinositol-anchor biosynthesis. Transfers the second mannose to the glycosylphosphatidylinositol during GPI precursor assembly (By similarity).[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (1) encodes the longest isoform (1). Variants 1 and 4 encode the same isoform (1). 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.